DOSSIER DE CANDIDATURE POUR LA PROMOTION AU GRADE DE :

CHARGE DE COURS

Présenté par

DESOBGO ZANGUE Steve Cary

Assistant

Comité Consultatif des Institutions Universitaires (CCIU) Commission VI : Sciences de l'Ingénieur Section 2

Sous-section 2 : Biologie et Biochimie appliquées

1- IDENTIFICATION



DESOBGO ZANGUE Steve Carly ; Ph.D ; Assistant

UNIVERSITE DE NGAOUNDERE; INSTITUT UNIVERSITAIRE DE TECHNOLOGIE

Département de Génie Alimentaire et Contrôle Qualité

B. P. 455 – Ngaoundéré, CAMEROUN ; Tel : 77909391/95134285 ; E-mail : desobgo@yahoo.fr/stevecarly@hotmail.fr

A Monsieur le Ministre de l'Enseignement Supérieur S/c Monsieur le Secrétaire Permanent du CCIU S/c Monsieur le Recteur de l'Université de Ngaoundéré S/c Monsieur le Directeur de l'IUT

Objet : Candidature au grade de Chargé de Cours

Monsieur le Ministre,

J'ai l'honneur de soumettre, pour examen à la prochaine session du Comité Consultatif des Institutions Universitaires (CCIU), mon dossier de candidature au grade de Chargé de Cours des Universités du Cameroun.

Je suis Assistant au département Génie Alimentaire et Contrôle Qualité de l'IUT de l'Université de Ngaoundéré, depuis le 16 avril 2008 (voir Décision N° 2008/229/UN/R/VR-EPDTIC/SG/DAAC/DEPE/SSPE).

Vous remerciant par avance de l'intérêt que porteront vos services à l'examen de mon dossier, et me tenant à votre disposition pour toute information complémentaire, Je vous prie de croire, Monsieur le Ministre, à mes sentiments les plus respectueux.

P.J. : Sept exemplaires du dossier de candidature dont 1 original et 6 copies.

DESOBGO ZANGUE Steve Carly

COMITE CONSULTATIF DES INSTITUTIONS UNIVERSITAIRES FICHE DE SYNTHESE DE CANDIDATURE

UNIVERSITE : NGAOUNDERE ETABLISSEMENT : INSTITUT UNIVERSITAIRE DE TECHNOLOGIE DEPARTEMENT : GENIE ALIMENTAIRE ET CONTROLE QUALITE GRADE ACTUEL : ASSISTANT depuis le 16/04/2008

GRADE POSTULE : *CHARGE DE COURS* **SOUS-SECTION 2 :** *BIOLOGIE ET BIOCHIMIE APPLIQUEES* **SECTION : 2 COMMISSION VI :** *SCIENCES DE L'INGENIEUR*

Identification du candidat	Cursus et Titres Universitaires	Expérience Professionnelle/Publications	Avis motivé du département	Avis de la commission
Nom : DESOBGO ZANGUE	1996 : Baccalauréat « D » (Institut	2008 à ce jour : Assistant / IUT de		
Prénom : Steve Carly	Privé Polyvalent de Bonamoussadi,	Ngaoundéré		
Né le : 02 Janvier 1977	Douala)	-		
A : Douala	1998 : DUT en Génie Agro-	Publications dans le grade		
Matricule : 652117-M	Industriel (Institut Universitaire de	d'assistant	Avis motivé de	Avis du Comité
	Technologie, Université de		l'établissement	
Pièces fournies	Ngaoundéré)	5 publications dans des journaux à		
Sept exemplaires de dossier de changement de	2001 : Ingénieur des Industries	comité de lecture		
grade dont 1 original comprenant chacun :	Agricoles et Alimentaires (Ecole			
1. Demande de changement de grade	Nationale Supérieure des Sciences	Participation à la 13 ième chair J.		
2. Curriculum Vitae	Agro-Industrielles, Université de	DE CLERCK. 2008	Avis motivé de la	Avis du Conseil
3. Photocopie de l'acte de naissance	Ngaoundéré)		Sous-section	d'Université
4. Photocopie des diplômes depuis le BACC	2003 : DEA en Génie des Procédés	Participation aux 1ères Journées		
5. Actes Administratifs	(Ecole Nationale Supérieure des	Scientifiques (JS) du réseau Génie		
6. Actes Académiques	Sciences Agro-Industrielles, Université	des Procédés Appliqués à l'Agro-		
7. Liste des productions scientifiques	de Ngaoundéré)	alimentaire (GP3A). 2008		
8. Une liste d'encadrement des thèses, mémoires et	2012 : Doctorat/Ph.D en Génie des		Avis motivé de la	Décision du
rapports	Procédés (Ecole Nationale Supérieure	Encadrement	Section	Conseil
9. Une liste des experts	des Sciences Agro-Industrielles,	• Vingt-sept (27) DUT		d'Administration
10. Sept (07) exemplaires de publications avec	Université de Ngaoundéré)	soutenus		
comité de lecture		• Six (6) Licences		
Rapport pédagogique (07)		Professionnelles soutenues		
Rapport administratif (07)				

DESOBGO ZANGUE Steve Carly

(Assistant à l'IUT de l'Université de Ngaoundéré, CAMEROUN)

Curriculum Vitae

Information personnelle

Adresse(s) *Téléphone(s) Courrier(s) électronique(s) Nationalité(s)* Date et lieu de naissance Sexe Statut Matrimonial

Domaine de compétence

Expérience professionnelle

Dates *Nom de l'employeur* Type ou secteur d'activité Fonction ou poste occupé Activités et responsabilités

Dates Nom de l'employeur Type ou secteur d'activité Fonction ou poste occupé Activités et responsabilités

Activités et responsabilités

Dates Nom de l'employeur Type ou secteur d'activité Fonction ou poste occupé Activités et responsabilités

Dates

Mai – Septembre 2001

Nom de l'employeur Type ou secteur d'activité Fonction ou poste occupé Activités et responsabilités

Doctorat/Ph.D en Génie des Procédés **DEA en Génie des Procédés** Ingénieur en Industries Agricoles et Alimentaires **DUT en Génie Agro-Industriel**

BP: 455 IUT Ngaoundéré, Cameroun Portable : 77 90 93 91 / 95 13 42 85 desobgo@yahoo.fr / stevecarly@hotmail.fr Camerounais 02 Janvier 1977 à Douala Masculin Marié et père de 2 enfants

Génie des Procédés et Ingénierie

2008- A nos jours Université de Ngaoundéré/Institut Universitaire de Technologie (IUT) Enseignement supérieur Enseignant-Chercheur

- Responsable Pédagogique
- Enseignements, Travaux Dirigés, Travaux Pratiques, Recherches

2006-2007

Collège de Mazenod (Ngaoundéré) Education (Enseignement secondaire) Vacataire

- Enseignant de Mathématiques
- Enseignant d'Informatique -
- TP Génie et Procédés des Industries Alimentaires
- **TP** Biodynamique _

2004-2005

ENSAI de Ngaoundéré

Agro-alimentaire

Moniteur

- TP Génie et Procédés des Industries Alimentaires
- **TP** Biodynamique

CAM Assistance

Agro-alimentaire (Fabrication des machines, mise au point des nouveaux produits) Stagiaire (Stage Fin d'Etudes Ingénieur)

- Elaboration des procédés de fabrication des jus et vins d'ananas
- Mise au point des jus et vins d'ananas
- -Elaboration d'un test de dégustation
- Rétention du meilleur procédé pour le jus et le vin d'ananas _

Juin – Août 2000 Dates Cameroon Development Corporation (CDC) Nom de l'employeur

5

Type ou secteur d'activité Fonction ou poste occupé Activités et responsabilités

Dates

Agro-alimentaire (Hévéa, Thé,.....) Stagiaire (Stage Agent de Maîtrise 2)

- Investigations sur les problèmes de qualité du caoutchouc
- Mise au point d'un équipement permettant l'évacuation des eaux issues du lavage du latex coagulé vers les égouts pour prétraitement

Juillet - Août 1999

Nom de l'employeur Société des Provenderies du Cameroun Agro-alimentaire (Provenderies) *Type ou secteur d'activité* Fonction ou poste occupé Stagiaire (Stage Agent de Maîtrise 1) Activités et responsabilités

- Contribution à la fabrication de la provende pour volaille, porcs, lapins,....
- -Contribution à la gestion de stocks de provende et maintenance

Dates Nom de l'employeur Type ou secteur d'activité Fonction ou poste occupé Activités et responsabilités

Education et formation

Juin – Septembre 1998 Guinness Cameroon S.A.

Agro-alimentaire (Brasseries)

Stagiaire (Stage Fin d'Etudes Technicien Supérieur)

- Analyse des casses bouteilles au service Packaging
- Investigation sur les points critiques des chocs thermiques et des casses de _ bouteilles sur la chaîne d'embouteillage plus défauts de fabrication.
- Evaluation des casses de bouteilles et coûts

Juin – Juillet 1997 Dates Nom de l'employeur Guinness Cameroon S.A. Agro-alimentaire (Brasseries) Type ou secteur d'activité Fonction ou poste occupé Stagiaire (Stage Ouvrier) Activités et responsabilités Apprentissage du fonctionnement de la brasserie, responsable stagiaire

2001 - 2003Dates Intitulé du diplôme délivré Diplôme d'Etudes Approfondies en Génie des Procédés (D.E.A - GP) Compétences couvertes Domaine général : Génie des Procédés, Mathématique, Statistiques Domaine spécifique : Modélisation, Optimisation des procédés des Industries Agricoles et Alimentaires (Boissons) Établissement École Nationale Supérieure des Sciences Agro-Industrielles (ENSAI), Ngaoundéré, Cameroun 1998 - 2001Dates Intitulé du diplôme délivré Diplôme d'Ingénieur Compétences couvertes Domaine général : Génie des Procédés et chimique, Ingénierie, Boissons Etablissement Ecole Nationale Supérieure des Sciences Agro-Industrielles (ENSAI), Ngaoundéré, Cameroun 1996 - 1998 Dates Intitulé du diplôme délivré Diplôme Universitaire de Technologie (DUT) Compétences couvertes

Dates Intitulé du diplôme délivré Compétences couvertes Etablissement

Établissement

Domaine général : Génie Agro-Industriel Institut Universitaire de Technologie (IUT), Ngaoundéré, Cameroun

1995 - 1996Baccalauréat de l'Enseignement Secondaire (Série D) Domaine général : Mathématiques et Sciences de la nature Institut Privé Polyvalent de Bonamoussadi (IPPB), Douala, Cameroun

Aptitudes et compétences personnelles

Langue(s) maternelle(s)	Français				
Auto évaluation	Comprendre		Parler	Parler	
	Écouter	Lire	conversation	oralement	
Anglais	Bien	Bien	Assez Bien	Assez Bien	Bien
Aptitudes sociales		t d'équipe, ada apacités de con	ptation facile aux	environnements	multiculturels
Aptitudes organisationnelles	 Responsable Pédagogique de la mention Génie Biologique niveau (2009-2011) Intérimaire régulier du chef de département Génie Alimentaire et Contré Qualité (GACQ) en cas de déplacement. Président de la mutuelle des Moniteurs et Doctorants de l'Université Ngaoundéré (2005) Moniteur à l'ENSAI de Ngaoundéré 2004/2005 et 2005/2006 Membre du Club Junior Entreprise section de Ngaoundéré (1998 – 2000) Secrétaire général de la Commission des Projets dans l'Association de Etudiants en Formation Doctorale de l'ENSAI de Ngaoundéré (2002004) Commissaire aux comptes du Club Génie Agro-industriel, IUT (1991997) 			aire et Contrôle l'Université de 06 (1998 – 2000) Association des oundéré (2001-	
Compétences informatiques	- Bonne Maîtrise des logiciels : Word, Excel, PowerPoint, SigmaPlot, Statistica, Mathcad, Matlab, Mathematica, Workplace, Statgraphics, SPSS, Systat, S Plus 2000, Derive Photoshop, fortran, etc			ica, Scientific	
Compétences artistiques	- Pratique of	du dessin et de	la poésie		
Autres compétences	- Pratique of	du Football, ter	nnis, tennis de table	e	
Permis de conduire	- Permis B				
Complément	Valeur e (Recteur	 Distinction Spéciale décernée pour avoir validé toutes les Unités de Valeur en session normale par le Recteur Pr. Maurice TCHUENTE (Recteur de l'Université de Ngaoundéré) (1998) Major de la Promotion DEA Génie des Procédés (2003) 			
Thèses, Mémoires	 physico-c de Docto Ngaounde Etude de de sorgho en Génie Mise au (2001). M Ngaounde Investigat Agent de Contribut 	chimiques des brat Ph.D en (éré. l'effet de quelo (<i>Madjeru, Sa</i> des Procédés. I point du jus e Mémoire de fin éré. tion sur les pro Maîtrise II. El ion à la fabric apport de stag	on des hydrolases moûts de deux cu Génie des Procéd ques hydrolases su <i>frari, S.35</i>) non m ENSAI de l'Univer t du vin d'ananas n d'études Ingénie oblèmes de caoute NSAI de l'Univers cation de la prove ge Agent de Maîtri	ltivars de sorghe és. ENSAI de r la filtrabilité d altés (2003). Me rsité de Ngaound pour une unité eur. ENSAI de chouc (2000). R ité de Ngaoundé nde pour volaill	b (2012). Thèse l'Université de es moûts a base émoire de DEA déré. semi artisanale l'Université de apport de stage rré. le, porcs, lapins

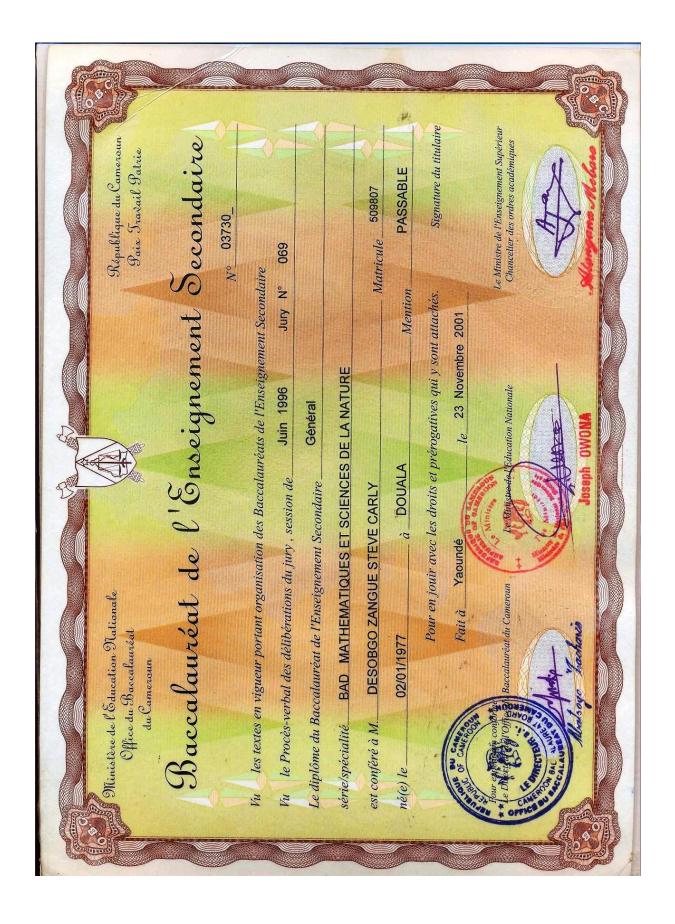
- Analyse des casses de bouteilles au service packaging chaîne n°2 (1998).
 Mémoire de fin d'études DUT. IUT de l'Université de Ngaoundéré.
- Rapport de stage ouvrier (1997). IUT de l'Université de Ngaoundéré.

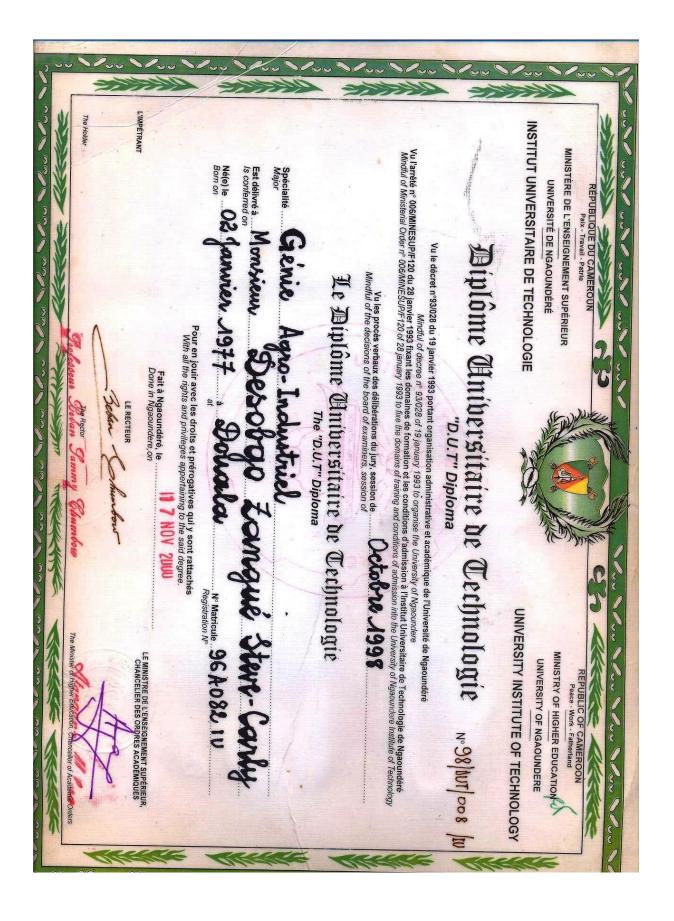
Publications et colloques

- **Desobgo, Z.S.C.**, Nso, E.J. & Tenin, D., 2011. The response surface methodology as a reliable tool for evaluating the need of commercial mashing enzymes for alleviating the levels of reducing sugars of worts of malted sorghum: Case of the *Safrari* cultivar. *Journal of Brewing and Distilling*, 2, 5-15.
- **Desobgo, Z.S.C.,** Nso, E.J. & Tenin, D., 2011. Use of the response surface methodology for optimizing the action of mashing enzymes on wort reducing sugars of the *Madjeru* sorghum cultivar. *African Journal of Food Science*, 5, 91-99.
- Desobgo, Z.S.C., Nso, E.J. & Tenin, D., 2011. Modeling the action of technical mashing enzymes on extracts and free-amino nitrogen yields of the *Madjeru* sorghum cultivar. *Journal of Brewing and Distilling*, 2, 29-44.
- **Desobgo, Z.S.C.**, Nso, E.J. & Tenin, D., 2011. Optimisation of the Action of Commercial Mashing Enzymes on Wort Extracts and Free Amino Nitrogen of the *Safrari* Sorghum Cultivar. *Technical Quarterly Master Brewers Association of the Americas*, 48, 77-86.
- **Desobgo, Z.S.C.**, Nso, E.J., Tenin, D. & Kayem, G.J., 2010. Modelling and optimizing of mashing enzymes-effect on yield of filtrate of unmalted sorghum by use of response surface methodology. *Journal of the Institute of Brewing*, 116, 62-69.
- The polyphenol paradox in alcoholic beverages (a beer and wine paradox ?). 13th Chair J. DE CLERCK. 7-10 september 2008. Louvain-la-Neuve, Belgium. <u>www.declerckchair.com</u>
- Desobgo Z.S.C., 2008. Modélisation et optimisation de l'effet des enzymes industrielles sur le rendement en filtrat des maïsches de sorgho. 1ères Journées Scientifiques (JS) du réseau Génie des Procédés Appliqués à l'Agro-alimentaire (GP3A). 11 et 12 septembre 2008, Louvain-la-Neuve, Belgique. <u>http://www.gp3a.auf.org</u>

DEPARTEMENT REPUBLIQUE UNIE DU CAMEROU Paix-Travail-Potrie DIVISION . UNITED REPUBLIC OF CAMEROON du Moun Peace-Work-Fatherland ARRONDISSEMENT SUBDIVISION de Dougla 1 CENTRE D'ETAT CIVIL CIVIL STATUS REGISTRATION CENTRE de - Of Audo Mun rod Acte d e Naissance Nº 9/22 BIRTH CERTIFICATE Nom de l'enfant.m. 212MGML S Name of the child Le - On the Alune Est he a. Canada Was born at Nom de l'enfant 🛋 Name of the child De sexe - Sex.m. De-Of Just Né à - Born at le - On the de Vomicilië à... Resident at Profession - Occupation U. Hutowsbelles henne El de - And of Née à - Born at: 1917 T Le - On the Domiciliée à Resident at · Profession - Occupation Dressé le huits to Drawn up on the pline oght Sur la déclaration de 1992 et In accordance with the declaration of Tillares mase Lesquels ont certifié la sincérité de la présente déclaration. Who attested to the truth of this declaration, 10 21983 Par Nous Beun By Us. dein Signature de l'Officier de l'Etat-Civil Signature of Registrar, Le Déclarant, The declarant,

2- DIPLOMES





	REPUBLIC OF CAMEROON Paso - WON - Fathenand MINISTRY OF HIGHER EDUCATION UNIVERSITY OF NGAOUNDERE NATIONAL ADVANCED SCHOOL OF	AGRO-INDUSTRIAL SCIENCES N ^g 01/J1NG/028/EN Indéré en Universités Industrielles de l'Université de Ngaoundéré e University of Ngaoundere D1	Contrast	98A005 E Le MINSTRE DE LE CHANCELLER DE	Maurice CHUENTE Minister of Higher Education, Chancellor of Academic Octaes
		AGRO-INDUSTRIELLES AGRO-INDUSTRIELLES Biplome by addition of decree of 92/074 du 13 avril 1992 portant transformation des Centres Universitaires de Buea et de Ngaoundéré en Universités Mindiul of decree of 92/074 of 13 April 1992 to transformation des Centres Universities de Buea et de Ngaoundéré en Universités Mindiul of decree of 92/074 of 13 April 1992 to transformation des Centres Universities de Buea et de Ngaoundéré en Universités Mindiul of decree of 92/074 of 13 April 1995 to organise the University of Ngaoundéré Mindiul of decree of 92/061 of 3 April 1995 to organise the Universite de Sciences of the Université de Ngaoundéré Wu le decree of 92/061 of 3 April 1995 to organise the National Advanced Science of Agro-Industrielles de l'Université de Ngaoundéré Wu le Beroce of the decree of 92/061 of 3 April 1995 to organise the National Advanced Sciences of the Université de Ngaoundéré Wu les procés - verbaux des delibérations du Jury, session de	Le Biplôme D'Ingénieur The Master of Engineering Degree Aricoles et Alimentoires Eargue Steve-Carl	And Connectors se droits et prérogatives qui y sont rattachés privileges appendaring to the said degree 1 9 MARS 2004 Le RECTEUR Le RECTEUR	R. Smerader Lotto Read Herrer
~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~	RÉPUBLIQUE DU CAMEROUN Par-Tendi - Parie MINISTÈRE DE L'ENSEIGNEMENT SUPÉRIEUR UNIVERSITÉ DE NGAOUNDÉRÉ ÉCOLE NATIONALE SUPÉRIEURE DES	SCIENCES AGRO-INDUSTRIELLES	A -	Is conterted on Né(e) le. 02 – 01 – 1977 a Bom on Bom on Pour en jouir avec te With all the rights and Fait à Yaoundé, le. Done in Yaoundé, le.	holder



## UNIVERSITE DE NGAOUNDERE

THE UNIVERSITY OF NGAOUNDERE

ECOLE NATIONALE SUPERIEURE DES SCIENCES AGRO-INDUSTRIELLES THE NATIONAL SCHOOL OF AGRO-INDUSTRIAL SCIENCES B.P.: 455 Ngaoundéré – Cameroun Tél//Fax: (237) 225-27-51 E-mail : enais: stages@yahoo.fr Site Internet : http://ensai-iut.minesup.gov.cm



## DIPLOME D'ETUDES APPROFONDIES EN GENIE DES PROCEDES CERTIFICAT DE REUSSITE / SUCCESS CERTIFICATE N° 003/003 /UN/D.ENSAI/DAACRS

Les soussignés

Vu le proce	ès verbal du jury en date du	19 juillet 2003	
M./Mlle	<b>DESOBGO ZANGUE St</b>	eve Carly	
Né(e) le	02 janvier 1977		

attestent que : Matricule n° 98A005EN

Douala

Titulaire du Diplôme des Ingénieurs des Industries Agricoles et Alimentaires

Ayant suivi les enseignements de DEA de la filière de Génie des Procédés (GP) pendant l'année académique

2001-2002 a validé la totalité des Modules / Unités de Valeurs avec les performances suivantes :

Code Module/UV	Intitulé des Modules / Unités de Valeurs	Note / 20	Mention	Année
MIGP6011	Mathématiques - Informatique pour le Génie des Procédés	18,0	TB	2002
PTGP6012	Phénomènes de Transport	15,0	B	2002
OU6014GP	Opérations unitaires de séparation et de Conservation	15,50	B	2002
TEGP6015	Techniques et Procédés nouveaux pour les IAA	13,50	B	2002
GA6013GP	Propriétés Phys., Physico-chimiques et Technol. des aliments	13,46	AB	2002
TA6113GP	Techniques Analytiques avancées	11,45	AB	2002
GBGP6312	Génie Biochimique et Biotechnologique	16,00	B	2002
MRGP6016	Méthodologie de la Recherche	15,60	B	2002
	Moyenne écrit	14,81	В	
IRGP6017	Stage d'Initiation à la recherche	15,50	В	2003
STGP6018	Mémoire de Recherche	14,00	B	2003
	Moyenne recherche :	15,50		
-/	MOYENNE GENERALE :	15,15		
	MENTION : Bien	and the second		

En foi de quoi ce certificat lui est délivré pour servir et valoir ce que de droit./.

Ngaoundéré le :

27 DEC. 2004

Le Chef de Division des Affaires Académiques de la Coopération, de la Recherche et de la Scolarité,

Pr. Hamga

Le Directeur,

# **3- ACTES ADMINISTRATIFS**

			DESILIPLIC OF CAMEROON
	REPUBLIQUE DU CAMEROUN		REPUBLIC OF CAMEROON Peace-Work-Fatherland
	Paix – Travail - Patrie		THE UNIVERSITY OF NGAOUNDERE
	UNIVERSITE DE NGAOUNDERE		*******
	BP: 454	Statement of the	P.O Box : 454
	RECTORAT		RECTOR'S OFFICE
4	VICE -RECTORAT CHARGE DES ENSEIGNEMENTS, D PROFESSIONNALISATION ET DU DEVELOPPEMENT TECHNOLOGIES DE L'INFORMATION ET DE LA COMMUN	DES	THE VICE RECTORATE IN CHARGE OF TEACHING, PROFESSIONALIZATION AND DEVELOPMENT OF INFORMATION AND COMMUNICATION TECHNOLOGIES
	******	PERATION O O	DIRECTION OF ACADEMIC AFFAIRS AND CO-OPERATION
	20 Décision N°	0 8 / 2 2 9 V /UN/R/VR-EPDTIC/S	S/DAAC/DEPE/SSPE
			alité d'Assistant au Département de Génie
	Alimentaire et Contrôle Qua	ité de l'Institut Universitaire de Techno	logie de l'Université de Ngaoundéré.
	LE RI RAI	CTEUR DE L'UNIVERSITE DE NG. PPORTEUR DU CONSEIL D'ADMINIS	AOUNDERE, STRATION
	Vu la Constitution ;		
	Vu 1a Loi nº2007/005 du 2	6 décembre 2007 portant Loi de Finances de la	République du Cameroun pour l'exercice 2008 ;
	subséquents :		nancier du Cameroun et les textes modificatifs
	Vii le décret nº 2005/142	lu 29 avril 2005 portant organisation du Ministé	ère de l'Enseignement Supérieur ;
	Vu le décret n° 92/74 du 1	3 avril 1992 transformant les Centres Universit	aires de Ngaoundéré et Buéa en Universités ;
	Vu le décret n°93/026 du	19 janvier 1993 portant création des Université 19 janvier 1993 portant dispositions communes	s ; applicables aux Universités, modifié et complété
	Par le décret nº 2005/3	42 du 10 septembre 2005:	
	Vu le décret n° 93/035 d	a 19 janvier 1993 portant statut spécial des p	ersonnels de l'Enseignement Supérieur, modifié et
		n° 2000/048 du 15 mars 2000; 19 janvier 1993 portant organisation administra	tive et académique
	de l'Université de Nga	oundéré ;	
	Vu le décret nº 93/028 du	19 janvier 1993 modifiant la rémunération des 1 15 mars 2000 fixant les modalités de rémunér	Fonctionnaires et Agents de l'Etat ;
	Vu l'arrêté n° 2000/050 d Vu le décret n° 2000/209	du 27 juillet 2000 fixant les modalités de reindrei du 27 juillet 2000 fixant la valeur du point d'inc	lice des Fonctionnaires de l'Etat
	Vu le décret n°2000/209 Vu le décret n°2000/212	du 27 juillet 2000 modifiant certaines disposit	ions du décret n°91/324 du 09 juillet 1991
	fixant les conditions d	attribution des logements administratifs :	
	Vu le décret n° 2008/	099 du 07 mars 2008 portant revalorisation	de la rémunération mensuelle de base des
	vu personnels civils et mi Vu le décret n° 2008/100	litaires; ) du 07 mars 2008 portant revalorisation du	taux de l'indemnité de non logement servie aux
	personnels civils et mi	litaires :	
	Vu le décret nº 2003/049	du 16 septembre 2003 portant nomination des R	tecteurs dans les Universités d'Etat ;
		ommission Consultative de Recrutement des As	ssistants de l'Université de Ngaoundéré en sa séance
	du 23 janvier 2008 ; Vu la décision du Conseil	de l'Université de Ngaoundére lors de Da Min	deuxième session en date du 24 janvier 2008 ;
	Vu l'avis favorable émis	par le Conseil d'Administration de l'Universite	en Nesoundéré lors de sa vingt unième session en
	date du 25 janvier 20	08;	Veneral Finan
	Vu la lettre nº 08/01866	/L/MINESUP/SG/DDES/SGE/CEA1/ne.ou.31 ITER recrutés lors de la CORActin 23 janvier 24 res de l'exercice 2008.	treuxième session en date du 24 janvier 2008 ; de l'insequendéré lors de sa vingt unième session en de la fil masse de la constant autorisation de mise en service 908 ; 200
	Vu les prévisions budgétai	res de l'exercice 2008	200 000981
	vu les previsions oudgoui	res de l'exercice 2008. VISA	200: 7 /
			d de .
		DECIDE . DOG	TAIGAOUN
	Article 1er : Monsieur DESOBGO Z	ANGUE Steve Carly titulaire d'un Diplôme	d'Effetles Approfondies (DEA) en Génie des
	Procédés est pour compter du 16 avril	2008, date effective de sa prise de service, rec	⁹ de Weadour Ta Weadour Ta Martine Appprofondies (DEA) en Génie des ruté en qualité d'Assistant, indice 1E/465, au e Technologie de l'Université de Ngaoundéré.
	Département de Génie Alimentaire et C	controle Quante de l'Institut Universitaire de	rechnologie de l'oniversité de righounderd
	Article2: L'Intéressé sera pris en c	harge par le budget de l'Etat Chap. 18 Art 1	02 Par 000 (18-102-000).
	Article 3: La présente décision ser	a enregistrée et communiquée partout où be	esoin sera.
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REPUBLIQUE DU CAMEROUN

Paix – Travail - Patrie

UNIVERSITE DE NGAOUNDERE

REPUBLIC OF CAMEROON

Peace - Work -- Fatherland

THE UNIVERSITY OF NGAOUNDERE

INSTITUT UNIVERSITAIRE DE TECHNOLOGIE DE NGAOUNDERE

B.P. 455 NGAOUNDERE - Tél. 77 11 22 20 & 99 85 13 82

Le Directeur

Ngaoundéré, le 16 AVR 2009

20_N 8/011_{/UN/D.IUT}

## CERTIFICAT DE PRISE DE SERVICE

Le Directeur de l'Institut Universitaire de Technologie (IUT) de l'Université de Ngaoundéré soussigné, certifie que Monsieur DESOBGO ZANGUE Steve Carly né le 02 Janvier 1977 à Douala, actuellement en cours de recrutement, a effectivement pris service à l'Institut Universitaire de Technologie le 16 Avril 2008 en qualité d'Assistant au Département de Génie Alimentaire et Contrôle Qualité.

En foi de quoi ce certificat est délivré à l'intéressé pour servir et valoir ce que de droit.

Ampliation: -Chrono/Archives.

LE DIRECTEUR MER

IUT DE NGAOUNDERE - B.P. 455 - Tél. 77 11 22 20 & 99 85 13 82

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<ul> <li>Vu les Résolutions du Conseil d'Administration de l'Université de Ngaoundéré, en sa session du 10 juillet</li> <li>Vu les prévisions budgétaires de l'exercice 2010.</li> </ul>	2010;
DÉCIDE :	
<u>Article 1</u> ^{er} : Monsieur DESOBGO ZANGUE Stève Carly, Matricule 652 117-M, Assistant 1 ^E /465 depuis est à compter du 16/04/2010, avancé au grade d'Assistant 2 ^E /530.	le 16/04/2008
Article 2 : L'Intéressé percevra à ce titre une rémunération mensuelle calculée sur la base de l'indice 530 co	orrespondant à
son grade et pourra prétendre aux avantages accordés aux Enseignants du même grade.	
Imputation : Budget de l'Etat, Exercice 2010 Chap.18, Art.102, Para. 000.	
Article 3 : La présente décision sera enregistrée et publiée partout où besoin sera./-	
Fait à Ngaoundéré, le 09 DEC. 2010	
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- DIRECTEUR- IUT/ CF	
- INTÉRESSÉ.	

# 4- ACTES ACADEMIQUES

4-1) Note de présentation soulignant la contribution à l'enseignement, la recherche, au développement de la science et de la culture ainsi qu'à la gouvernance de l'Université, de l'Enseignement Supérieur et de la Nation

## A. Contribution à l'Enseignement

Je suis enseignant à l'Université de Ngaoundéré depuis le 16 avril 2008, date de ma prise de service à l'Institut Universitaire de Technologie (IUT). Depuis lors, sous instruction de l'administration de l'IUT de Ngaoundéré, j'ai contribué aux enseignements. Dans un premier temps, j'ai été employé à la réalisation et à l'exécution des travaux pratiques pour finir par obtenir des cours au courant de l'année académique suivante. J'ai été mis en contribution dans les cours de technologies alimentaires au niveau du département Génie Alimentaire et Contrôle Qualité (GACQ). Ces cours étaient globalement dans le domaine des boissons et des céréales et amidonnerie. Par la suite, il m'a été confié des cours dans le domaine de l'huilerie, la sucrerie et fruits et légumes.

J'ai été choisi pour être responsable pédagogique sur deux ans. En effet, la tache consistait en l'élaboration des emplois de temps et à m'assurer tenue effective des enseignements. J'ai également assuré plusieurs intérims ponctuels du Chef de Département GACQ lors de ses déplacements pour les colloques. J'ai également participé en tant qu'encadreur lors des déplacements des étudiants dans le cadre des voyages d'études (visites d'entreprises)

Par la suite, j'ai été associé à la commission de refonte des programmes du département Génie Alimentaire et Contrôle Qualité dans le cadre de la cohérence des programmes du LMD. Depuis plusieurs années déjà, je participe à la recherche des stages académiques auprès des industriels, et également aux encadrements mémoires de fins d'études DUT et Licence Professionnelle.

## **B.** Contribution à la Recherche

Le point focal de la recherche que j'ai menée gravite autour de la valorisation du sorgho camerounais dans le domaine brassicole.

Le sorgho est la céréale par excellence de substitution à l'orge or, paradoxalement au Cameroun, l'orge reste exclusivement la céréale utilisée pour la fabrication de la bière. Pourtant, une politique ambitieuse de culture intensive du sorgho permettrait, sur le plan économique une rentrée de devises dans la mesure où cette matière première pourra être utilisée dans le secteur brassicole et des spiritueux. Politique que le Nigéria a impulsée depuis 1985.

### B-1) Les Problématiques du sorgho pour la fabrication de la bière

Le sorgho est indéniablement la céréale prépondérante dans la région de l'extrême nord du Cameroun. L'une des utilisations de cette céréale est la filière *Bili-Bili* (bière traditionnelle de sorgho) qui, prend des ampleurs de plus en plus importantes. Ainsi, la culture de cette céréale n'a plus pour objectif une alimentation sous forme de couscous et autres. Il a été démontré que cette filière *Bili-Bili* est une source indéniable de revenu pour les femmes qui sont à la base de la prolifération de la filière. Les qualités d'une telle boisson seraient indéniablement liées à la qualité du malt, du moût et de la fermentation. Malgré le fait que les schémas traditionnels soient suffisamment éprouvés, les faibles rendements et les qualités approximatives des bières qui y sont produites traduisent des lacunes dans lesquelles, la science pourrait apporter des éléments de solution. Il est donc apparu au vue de la maltabilité insuffisante de certains cultivars qu'un pan de la recherche sur l'amélioration de la filtrabilité de ces maïsches doit être fait. Par la suite, il a paru nécessaire de modéliser et optimiser l'utilisation des enzymes pour pallier à cette carence de synthèse enzymatique au cours du maltage, afin d'obtenir des moûts de qualité.

## **B-2)** Actes posées

Afin de mieux cerner les contours liés à la production du *Bili-Bili*, j'ai effectué dans le cadre d'un groupe de travail des descentes sur le terrain, à la rencontre des productrices de cette boisson dans les villes de Ngaoundéré et Garoua. Le travail de recherche se poursuit dans l'utilisation des plantes amères pour amériser les bières à base de ces céréales locales. La publication de 5 articles est la parfaite illustration du travail déjà entamé.

## C. Résumé des publications dans le grade d'Assistant

1. Desobgo, Z. S. C., Nso, E. J., Tenin, D. and Kayem, G. J.,

Modelling and optimizing of mashing enzymes-effect on yield of filtrate of unmalted sorghum by use of response surface methodology. *Journal of the Institute of Brewing*, 2010, 116, 62-69.

The effect of commercial enzymes on liquefaction of starch from unmalted sorghum was studied. The effects which these enzymes had on rates of filtration were evaluated. Models were developed, validated and optimized to establish the actions of enzymes, either alone or in combination. Preliminary studies on the sorghum cultivars Safrari, Madjeru and S.35 showed that  $\alpha$ -amylase was the backbone enzyme for starch liquefaction among the enzymes used ( $\alpha$ -amylase, Filtrase, protease and  $\beta$ -amylase). Models confirmed this observation as  $\alpha$ -amylase individually in its first order (X₁) contributed 25, 11 and 17 %, and in its sum of first and second orders  $(X_1 + X_1^2)$  contributed a 29, 31 and 36 % yield of filtrate for Safrari, Madjeru and S.35 respectively. The ease of starch liquefaction, assessed by summing the first and second orders of individual intervention of all enzymes, was found to be in the order of Madjeru, S.35 and Safrari (79, 70 and 56 % of yield of filtrate respectively). The importance of the enzyme combination in starch liquefaction in Safrari, S.35 and Madjeru was shown to be 44, 30 and 21 % respectively. Enzyme combinations giving maximal starch liquefaction, as identified from a Doehlert experimental matrix, displayed a similar yield of filtrate (Safrari: 85 ml, Madjeru: 84 ml and S.35: 81 ml) after filtration of a 130 ml mash during 1 h. Validation of the models revealed the model developed for Madjeru was the most reliable ( $R^2 = 0.994$ ), while those developed for Safrari ( $R^2 = 0.987$ ) and S.35 (R² = 0.976) were slightly less reliable. Model optimization gave theoretical enzyme (Brewers Amyliq TS, Filtrase NLC, Brewers Protease and β-amylase) combinations of 25 mg, 5.68 mg, 100 mg and 67.4 U for Safrari, 15.06 mg, 0.51 mg, 24.32 mg and 53.8 U for Madjeru and 19.01 mg, 6.36 mg, 58.76 mg and 43.48 U for S.35, with a resulting yield of filtrate of 94, 87.7 and 83.8 ml respectively.

**Key words**: Mashing enzymes, model validation, modeling, optimization, unmalted sorghum, yields of filtrates.

The response surface methodology as a reliable tool for evaluating the need of commercial mashing enzymes for alleviating the levels of reducing sugars of worts of malted sorghum: Case of the *Safrari* cultivar. *Journal of Brewing and Distilling*, 2011 Vol. 2(1) 5-15.

A three factor Doehlert design was used to develop a statistical model to optimize the action of three commercial mashing enzymes (Hitempase 2XL, Bioglucanase TX and Brewers protease) on reducing sugars content of the worts of unmalted and malted Safrari sorghum. The response surface methodology revealed that increasing amounts of Hitempase considerably increased reducing sugars content during mashing of unmalted Safrari sorghum grist to about 90 g/L. Limited amounts of reducing sugars were obtained with increasing concentrations of both Bioglucanase ( $\approx 40$  g/L) and Brewers Protease ( $\approx 30$  g/L). The milling process facilitated the dissolution of about 10 g/L in yields of reducing sugars for the unmalted Safrari sorghum mash type without the help of enzyme. None of the three enzymes as sole mashing enzyme appeared to be of use in mashing malted Safrari, as reducing sugars yields were at maximum (168 g/L) after dissolving the grist in water and rather decreased with increasing amounts of enzyme supplements. Optimizing the concerted actions of the three enzymes for reducing sugars content of unmalted Safrari sorghum mash gave a combination of 2163 U, 937.5 BGU and 0 mg for Hitempase, Bioglucanase and Brewers Protease respectively. This gave a maximal reducing sugars content of 126.57 g/L. This combination was 0 U, 137.48 BGU and 0 mg for malted Safrari sorghum mash, giving a maximal reducing sugars yield of 168.56 g/L.

Key words: Response surface methodology, optimization, mashing enzymes, *Safrari*, reducing sugars

Use of the response surface methodology for optimizing the action of mashing enzymes on wort reducing sugars of the *Madjeru* sorghum cultivar. *Journal of Brewing and Distilling*, 2011 Vol 2(1), 5-15.

A three factor Doehlert design was used to develop a statistical model to optimize the action of three commercial mashing enzymes (Hitempase 2XL, Bioglucanase TX and Brewers protease) on reducing sugars content of the worts of unmalted and malted *Madjeru* sorghum. The response surface methodology revealed that increasing amounts of Hitempase considerably increased reducing sugars content during mashing of unmalted and malted *Madjeru* sorghum grist to about 105.39 g/L and 132.25 g/L respectively. The milling process contributed to about 22 g/L and 54 g/L for the unmalted and malted mash types respectively. Increasing amounts of Bioglucanase was virtually insignificant, while for Brewers protease, reducing sugar yields rather decreased to nil for both the unmalted and malted mash types. Optimization of the concerted actions of the three enzymes for reducing sugars content of unmalted *Madjeru* sorghum mash gave a combination of 1995 U, 89.31 BGU and 28.86 mg for Hitempase, Bioglucanase and Brewers Protease respectively. This gave a maximal reducing sugars content of 108.78 g/L. This combination was 3000 U, 0 BGU and 49.69 mg for malted *Madjeru* sorghum mash, giving a maximal reducing sugars yield of 153.15 g/L.

Key words: Response surface methodology, optimization, mashing enzymes, Madjeru, reducing sugars.

Modeling the action of technical mashing enzymes on extracts and free-amino nitrogen yields of the Madjeru sorghum cultivar. *Journal of Brewing and Distilling*, 2011 Vol 2(3), 29-44.

The action of three technical mashing enzymes (Hitempase 2XL, Bioglucanase-TX and Brewers Protease) on yields of extract and free amino nitrogen (FAN) of the worts of mashes of unmalted and malted Madjeru sorghum was modeled and analyzed using the response surface methodology. The analysis showed that increasing amounts of Hitempase 2XL considerably increased yields of extract during mashing of unmalted Madjeru sorghum The use of Bioglucanase-TX was not indispensable, while Brewers Protease grist. contributed very little. Increasing amounts of Hitempase contributed approximately 45 % of the free amino nitrogen, while Brewers Protease influence amounted to not more than 15 %. Bioglucanase's action was globally nil. Addition of the three enzymes into malted Madjeru sorghum mashes had no significant effect on the yields of extracts and FAN, but the milling operation singularly liberated more than 50 % of FAN for both mash types. Optimization of the concerted actions of the three enzymes for extract yield for unmalted Madjeru sorghum mash gave a combination of (1960.5 U; 132.61 BGU and 28.86 mg) for Hitempase, Bioglucanase and Brewers Protease respectively). This gave a maximal extract yield of 16.55 °P. This combination was: 2610 U; 0 BGU and 40.44 mg for malted Madjeru sorghum mash, giving a maximal extract yield of 16.35 °P. Optimization for free amino nitrogen for unmalted Madjeru sorghum mash gave a combination of: 3000 U; 0 BGU and 100 mg for Hitempase, Bioglucanase and Brewers Protease respectively). This gave maximal FAN of 93.55 mg/L. The combination was: 3000 U; 0 BGU and 100 mg for malted Madjeru sorghum mash, giving a maximal FAN of 144.48 mg/L.

**Key words**: Modeling, technical mashing enzymes, yields of extract, free-amino-nitrogen, *Madjeru*, optimization.

Optimisation of the Action of Commercial Mashing Enzymes on Wort Extracts and Free Amino Nitrogen of the Safrari Sorghum Cultivar. *MBAA TQ*, 2011 Vol 48(3), 77-86.

The influence of three commercial mashing enzymes (Hitempase 2XL, Bioglucanase TX and Brewers Protease) used as sole mashing enzymes on yields of extract and free amino nitrogen (FAN) of the worts of the mashes of unmalted and malted Safrari sorghum was studied using the response surface methodology. The study revealed that increasing amounts of Hitempase considerably increased yields of extract during mashing of unmalted Safrari grist, while the effect of Bioglucanase was smaller, and that of Brewers Protease was insignificant. Extract yields decreased with increasing amounts of the three enzymes during the mashing of malted Safrari. This decrease was least expressed in the case of Brewers Protease. Yields in FAN amounted to less than 50 %, with increasing amounts of both Hitempase and Brewers Protease, but constantly decreased to nil for Bioglucanase's action in both unmalted and malted Safrari mashes. The milling operation singularly liberated more than 50 % of FAN for both mash types and for each of the enzymes. Optimisation of the concerted actions of the three enzymes for extract yields gave a combination of 2,098.5 U, 937.5 BGU and 0 mg (for Hitempase, Bioglucanase and Brewers Protease, respectively) for This gave a maximal extract yield of 18°P. unmalted Safrari sorghum mash. The combination was 0 U, 28.68 BGU and 0 mg for malted Safrari sorghum mash, giving a maximal extract yield of 18.82°P. Optimisation for FAN gave a combination of 2,434.5 U, 0 BGU and 100 mg (for Hitempase, Bioglucanase and Brewers Protease, respectively) for unmalted Safrari sorghum mash. This gave maximal FAN of 144.77 mg/L. The combination was 2,191.5 U, 0 BGU and 100 mg for malted Safrari sorghum mash, giving a maximal FAN of 196.73 mg/L.

Keywords: Commercial mashing enzymes, extract, free amino nitrogen, optimisation, *Safrari*.

## **Perspectives**

Les travaux menés sur l'amélioration des caractéristiques physico-chimiques des moûts montrent des possibilités d'élaboration d'une bière à base de sorgho camerounais. Il sera par la suite question de scruter les aléas liés à la fermentation de ces moûts. On pourra déterminer les profils qualitatifs et quantitatif des sucres et acides aminés de ces moûts. Ceci permettra de pouvoir appréhender les déterminants de cette fermentation.

## D. Rayonnement Scientifique et Culturel

Suite à la qualité des travaux effectués, j'ai été invité à la 13^{ième} Chair J. DE CLERCK sur le thème «The polyphenol paradox in alcoholic beverages (a beer and wine paradox ?)» du 7 au 10 september 2008, puis aux 1ères Journées Scientifiques (JS) du réseau Génie des Procédés Appliqués à l'Agro-alimentaire (GP3A), du 11 et 12 septembre 2008 à Louvain-la-Neuve en Belgique où, j'y ai présenté une communication intitulée «Modélisation et optimisation de l'effet des enzymes industrielles sur le rendement en filtrat des maïsches de sorgho.»

4-1) Liste des experts

Carl M. F. MBOFUNG (Pr); (Biochimie et Nutrition) Département de Sciences Alimentaire et Nutrition, ENSAI, Université de Ngaoundéré ;

2. TENIN DZUDIE (pr) ; (Génie Alimentaire) Département de Génie des Procédés et Ingénierie, ENSAI, Université de Ngaoundéré ;

3. Robert NDJOUENKEU (pr); (Chimie et Biochimie Alimentaire) Département de Sciences Alimentaire et Nutrition, ENSAI, Université de Ngaoundéré ;

4. TCHIEGANG Clergé (pr) ; (Biochimie Alimentaire) Département de Sciences Alimentaire et Nutrition, IUT, Université de Ngaoundéré ;

5. ETOA François Xavier (pr) ; (Microbiologie Alimentaire) Département de Biochimie,

FS, Université de Yaoundé 1;

6. ESSIA NGANG Jean Justin (MC) ; (Microbiologie Industrielle) Département de Biochimie, FS, Université de Yaoundé 1 ;

7. FONTEH Florence (MC); (Biochimie Alimentaire), FASA, Université de Dschang

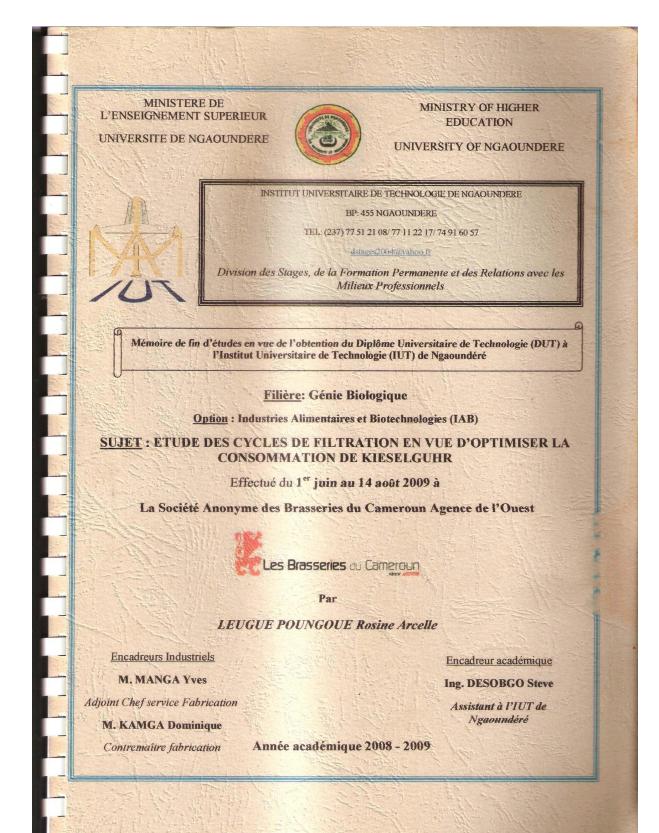
4-2) Liste des enseignements dispensés et dûment approuvée par le Chef de Département 4-2) Liste des rapports et mémoires dirigés dûment validée par le Chef de Département

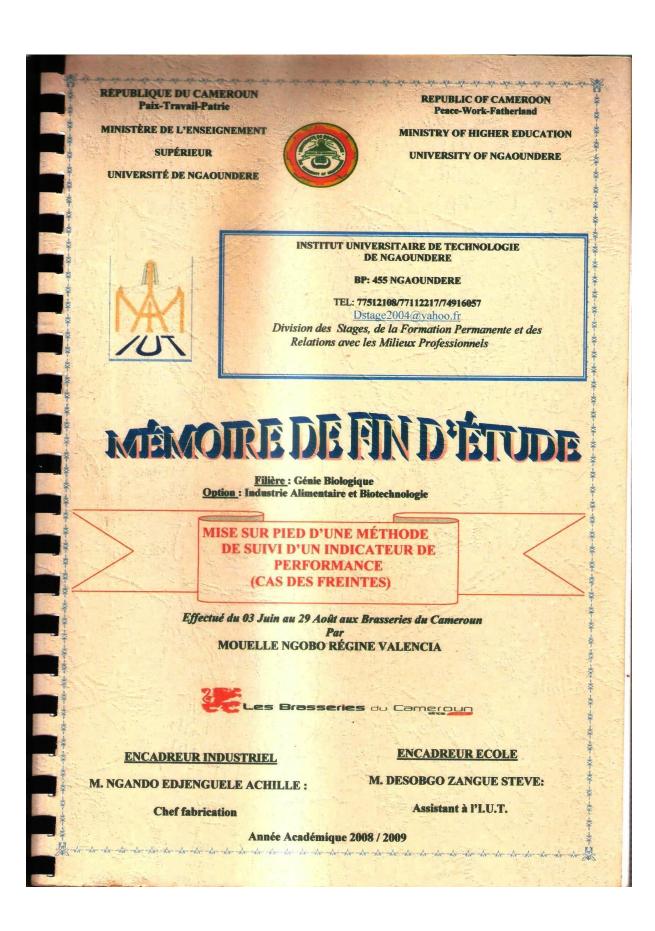
# Liste des mémoires de fins d'études DUT et LICENCE PROFESSIONNELLE encadrés par Dr. DESOBGO ZANGUE Steve Carly depuis 2009

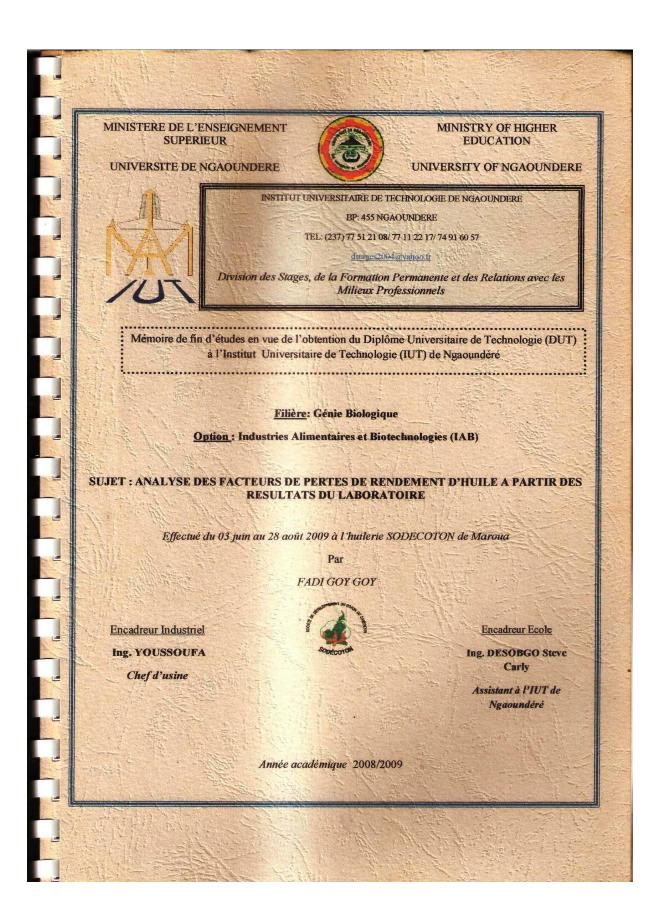
N°	Noms et Prénoms	Sujet	Année	Type de Mémoire
1	LEUGUE Rosine	Etude des cycles de filtration en vue d'optimiser la consommation du Kieselguhr	2009	DUT/IAB
2	MOUELLE NGOBO	Mise sur pied d'une méthode de suivi d'un indicateur de performance (cas des freintes)	2009	DUT/IAB
3	FADI GOY GOY	Analyse des facteurs de perte de rendement d'huile à partir des résultats du laboratoire	2009	DUT/IAB
4	ZENGUE EVINA Laurent Patrice	Etude de faisabilité pour la création d'une usine de pâtes alimentaires	2009	DUT/IAB
5	JOUBOUNE NGOMSI Christelle	Contribution à l'amélioration de la qualité du Palm'or	2009	DUT/IAB
6	ELOUNGOU ZOGO Salomon	Etude des freintes sur les chaines de production des jus de fruits «TAMPA» et «FRESCO»	2009	DUT/IAB
7	NNOMO MANGA Franck	Gestion de début et fin de filtration de la bière	2009	DUT/IAB
8	TAWEDI Robert Elvis	Gestion des déchets : traitement, recyclage et essais de valorisation, cas de la communauté urbaine d'Edéa.	2009	DUT/IAB
9	TIYOU PADRIK	Etude de l'effet de quelques hydrolases (a- amylase, b-glucanase, protéase et amyloglucosidase) sur l'extrait et la fermentescibilité des moûts de <i>Safrari</i> malté	2009	DUT/IAB
10	SUH QUEENIVA LUM	Contribution to the improvement of the physico-chemical characteristics of malted sorghum wort	2009	DUT/IAB

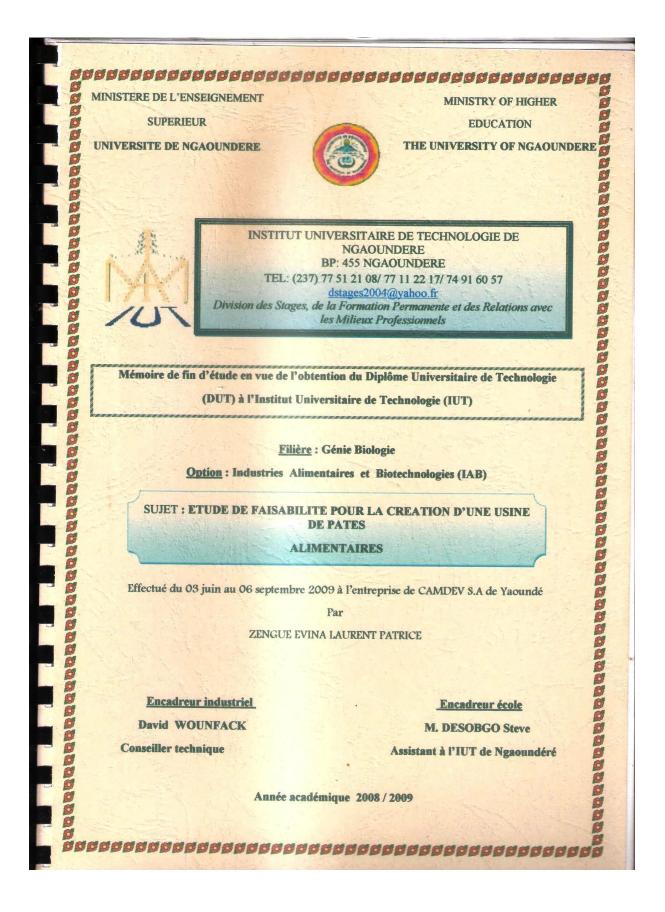
11	AZOULINE MICHEL	Influence des auxiliaires technologiques sur le rendement et la qualité de l'huile Diamaor	2010	DUT/IAB
12	DAOUDA YAOUBA	Suivi des paramètres physico-chimiques de l'oléine du CCIC en fonction du temps	2010	DUT/IAB
13	DJIZANNE FOFOU Gérard	Optimisation du fonctionnement des filtres à maïsche en salles à brasser	2010	DUT/IAB
14	KANA Simon	Etude du process d'extraction du jus de soja, en vue de l'amélioration du rendement et du brix	2010	DUT/IAB
15	LEUGOUE TCHAGO	Evaluation de l'efficacité de chaque thimonnier	2010	DUT/IAB
16	MEMANA	Evaluation des pertes sur la ligne «Riverr»	2010	DUT/IAB
17	MPON ETSIKE Luc Michel	Evaluation des pertes sur les lignes Bidon et Sachet à ADIC	2010	DUT/IAB
18	NIDJO SAHA	Optimisation du filtre MEVRA 2001	2010	DUT/IAB
19	BONBO DIBE	Etude sur la gestion des déchets d'une savonnerie et proposition d'amélioration (cas d'ISF)	2010	LICENCE PRO/GEN
20	ELOUNGOU ZOGO SALOMON	Etude comparative du pouvoir fermentaire de trois levures (levures sauvages, Saccharomyces carlsbergensis, CLIB 655) sur les moûts de Safrari malté	2010	LICENCE PRO/IAB
21	NJIAH SUSAN FOFEYIN	Comparative study on the fermentation of malted <i>Madjeru</i> by the use of three differents yeasts, <i>Saccharomyces</i> <i>carlsbergensis</i> , <i>Clib</i> 655 and Indigenous yeast)	2010	LICENCE PRO/IAB
22	SUH QUEENIVA LUM	Proposal of a production unit and the recommendation of the land dimensions for the manufacturing of baby food (kukulu) SERDIF project	2010	LICENCE PRO/IAB

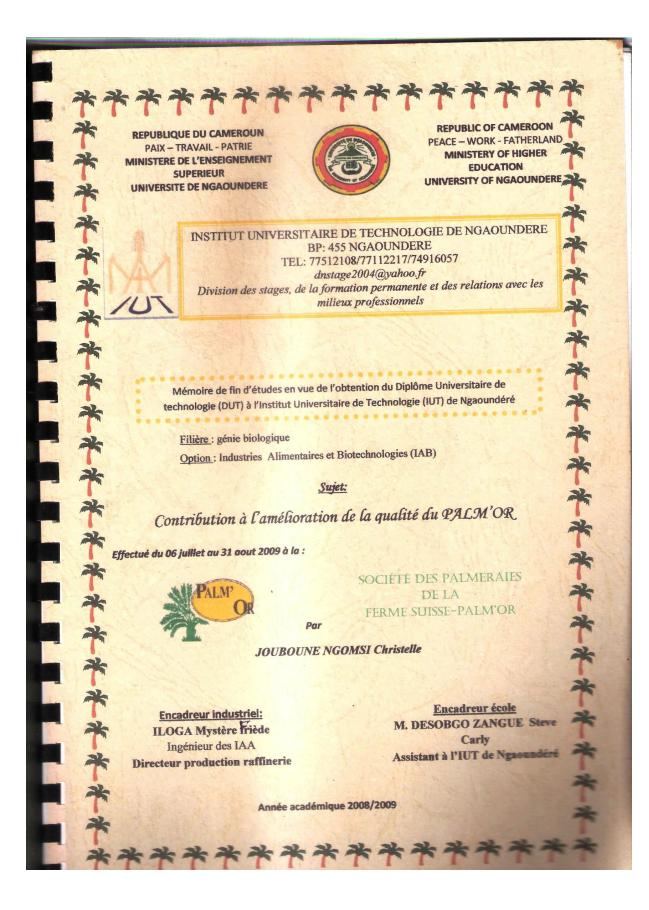
23	LUM Linda	An environmental audit of International Soap Factory	2011	DUT/GEN
24	MENIAPI Berthe	Suivi de la méthode de gestion du kieselguhr	2011	DUT/GEN
25	BATCHEP BEKIMA Bill	Evaluation de la cadence des ensacheuses volumétriques, respectivement sur les lignes de conditionnement «Nouriss» et «Jovino»	2011	DUT/IAB
26	BOUN LIKENG Joseph	Suivi de la viabilité des levures types SABC : axes d'amélioration	2011	DUT/IAB
27	MBELLI Juliet ABONG	Ameliorating the ageing time of P14 ageing oven using Cameroon Natural Rubber (CNR) 10	2011	DUT/IAB
28	MBETMI KELLEU Charles Ghislain	Etude des freintes des boites sur la chaine de production des conserves de haricots verts	2011	DUT/IAB
29	NDOOH Samuel	Optimisation de la production en yaourterie	2011	DUT/IAB
30	NGNOKAM NZOUKOU Calixte	Optimisation du temps d'execution aux postes d'analyses physico-chimiques au laboratoire : cas poste de physico-chimie bière	2011	DUT/IAB
31	TJEEGA Jean Elisée	Mise au point d'un yaourt aromatisé à la noix de coco	2011	DUT/IAB
32	YOMI YOMI Brice Herman	Mise au point d'un Fromage enrichi aux épices locales	2011	DUT/IAB
33	KENMOGNE YONY Blaise	Suivi de la fermentation d'une bière de sorgho et essai de clarification à l'aide d'extraits de <i>Moringa oleifera</i>	2011	DUT/IAB
34	MORNADJI Félix	Etude de la capabilité du filtre à maïsche usine SABC de Garoua	2011	LICENCE PRO/IAB

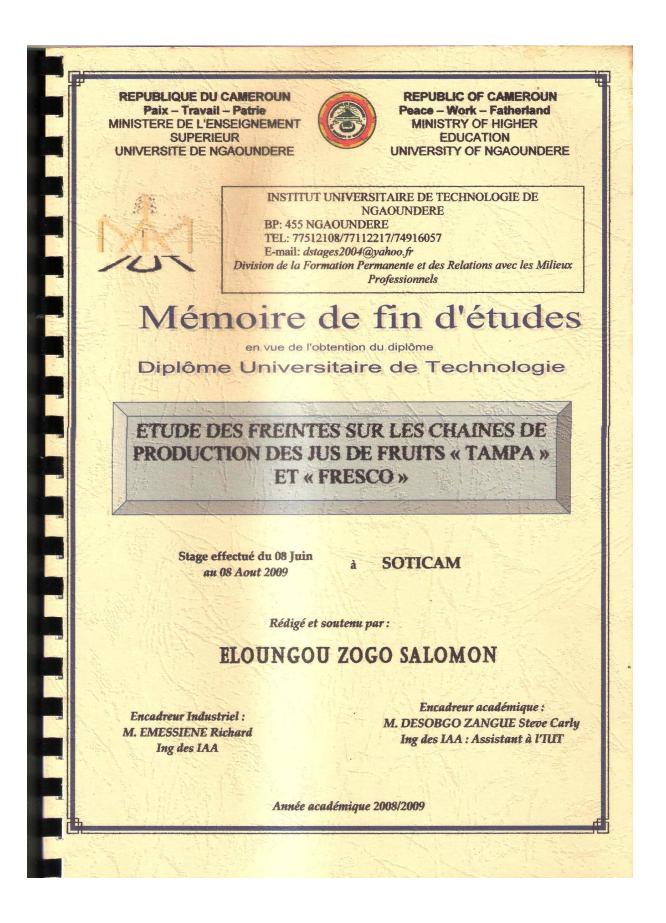














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En vue de l'obtention du Diplôme Universitaire de Technologie (DUT),

Filière : Génie Biologique.

**Option : Industries Alimentaires et Biotechnologies (IAB)** 

Sujet :

GENTION DE DRIEUT ET DE FIN DE FILTRATION DE LA BIERE

Effectué du 08 Juin au 31 Août 2009 à la S.A. B.C. Direction d'usine De Yaoundé

es Brasseries du Cameroun

Par NNOMO MANGA FRANK

Encadreur industriel

Mr. FELIX MBANGTENG Chef Service Fabrication

MILE NATHALIE NOAH **Chef Atelier Caves** 

Encadreur école :

M& DESOBGO STEVE C. Assistant chargé de cours IUT

Année académique 2008/2009

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Mémoire de fin d'études en vue de l'obtention du Diplôme Universitaire de Technologie (DUT) à l'Institut Universitaire de Technologie (IUT) de Ngaoundéré

Filière: Sénie Biologique

Option : Génie de l'environnement (GEN)

SUJET : GESTION DES DECHETS : TRAITEMENT, RECYCLAGE ET ESSAIS DE VALORISATION, CAS DE LA COMMUNAUTE URBAINE D'EDEA

Effectué du 1^{er} Juillet au 28 Août 2009 à la Communauté Urbaine d'Edéa

Par:

TAWEDI Robert Elvis

Encadreur industriel

M. NYEMBE Roger

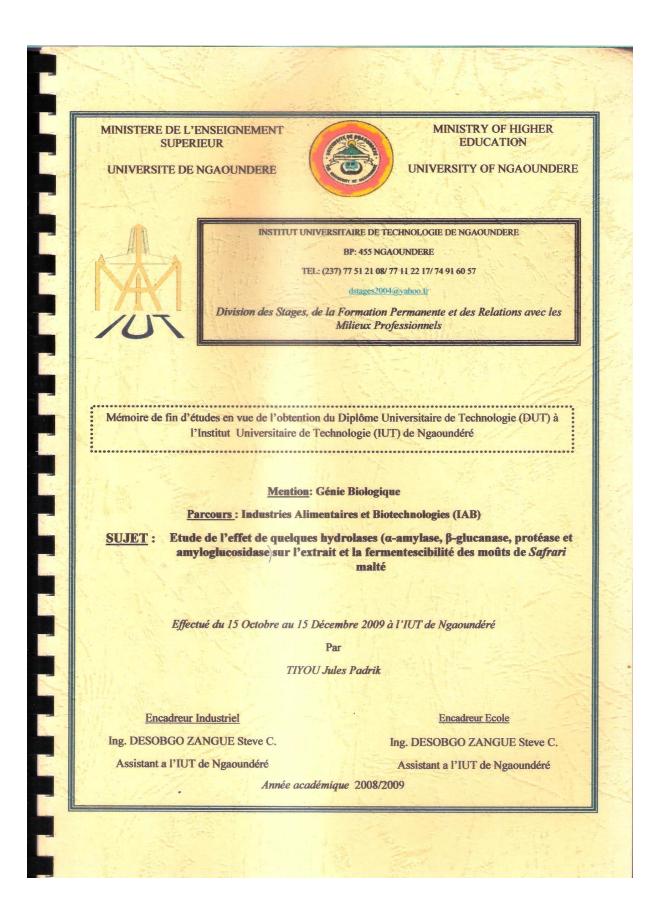
Chef service hygiène et assainissement

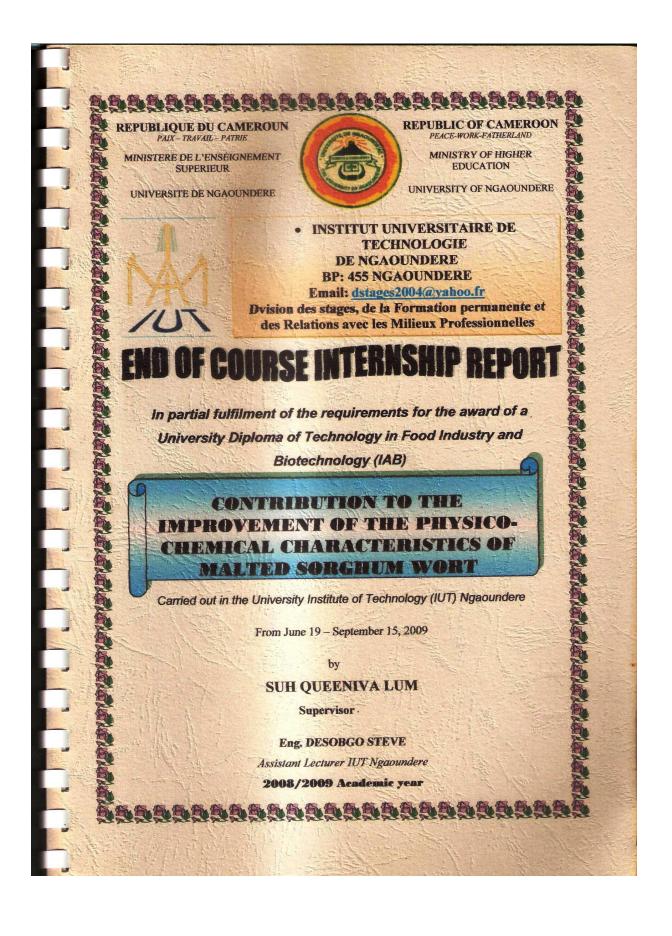
Encadreur école

M. DESOBGO Steve

Assistant à l'IUT de Ngaoundéré

Année académique : 2008/2009







# **REPUBLIQUE DU CAMEROUN**

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Université de Ngaoundéré

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# REPUBLIC OF CAMEROON

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Mémoire fin d'études en vue de l'obtention du Diplôme Universitaire de Technologie

### (D.U.T.)

Parcours : Génie Biologie

Mention : Industries Alimentaires et Biotechnologies (I.A.B.)

Sujet :

INFLUENCE DES AUXILIAIRES TECHNOLOGIQUES SUR LE RENDEMENT ET LA QUALITE D'HUILE DIAMAOR

Effectué du 01 juin au 27 Aout 2010 à l'huilerie de la SODECOTON de GAROUA

Par



# AZOTEINNE MICHEL

Matricule : 08B007IU

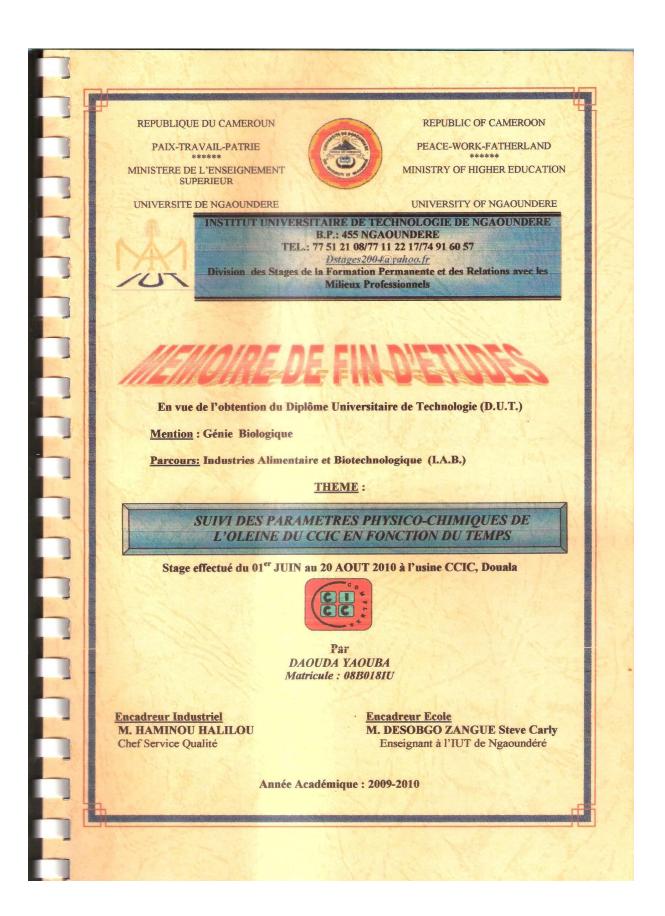
Encadreur industriel :

Ing. Bobo Yaya

Encadreur Ecole :

Ing. Desobgo Steve Carly

Année académique 2009 /2010



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Filière : Génie Biologique

**Option : Industries Alimentaires et Biotechnologies (IAB)** 

SUJET : OPTIMISATION DU FONCTIONNEMENT DES FILTRES

À MAISCHE EN SALLES À BRASSER



Effectué du 15 octobre au 15 décembre 2009 à Pusine SABC de Koumassi

Par DJIZANNE FOFOU Gérard

THEATERS

Encadreurs Industriels: M. ZONTSOP Etienne Directeur des Usines Adjoint

SABC Littoral

M. NSANGOU ABOUBAKAR Adjoint au Chef Service Fabrication

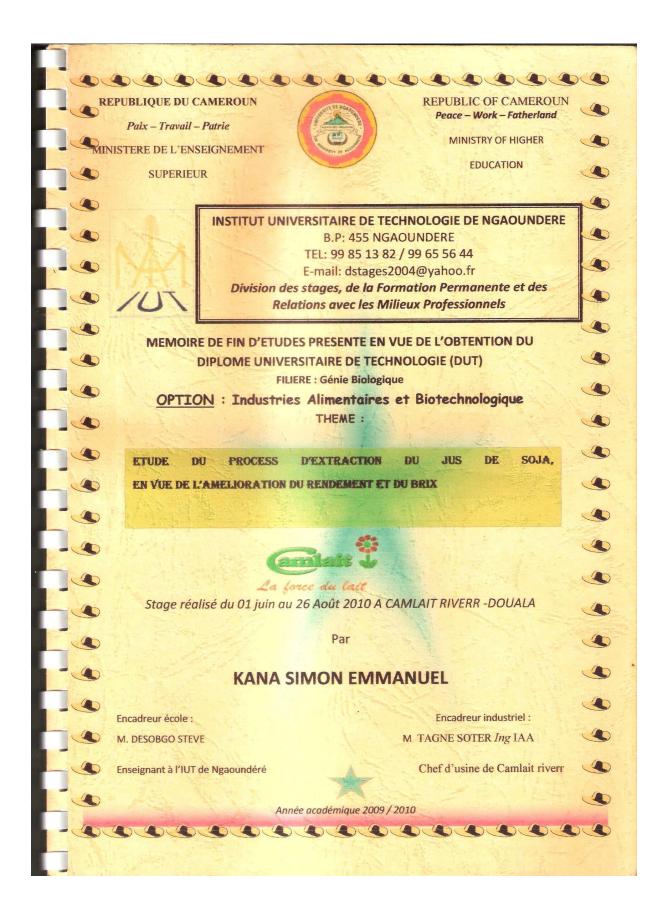
à l'usine de Koumassi

Année académique : 2008 / 2009

M. DESOBGO Steve

Encadreur école:

à l'IUT de Ngaoundéré



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Filière : Génie Biologique

Mention : Industries Agroalimentaires et Biotechnologiques



Stage effectué du 02 juin au 26 Août 2010 à CAMLAIT NDOKOTTI-DOUALA

Par

TAILER

Ca force du lait

# MEMANA GEINGA Salomé Flora

08B044IU

Encadreur industriel

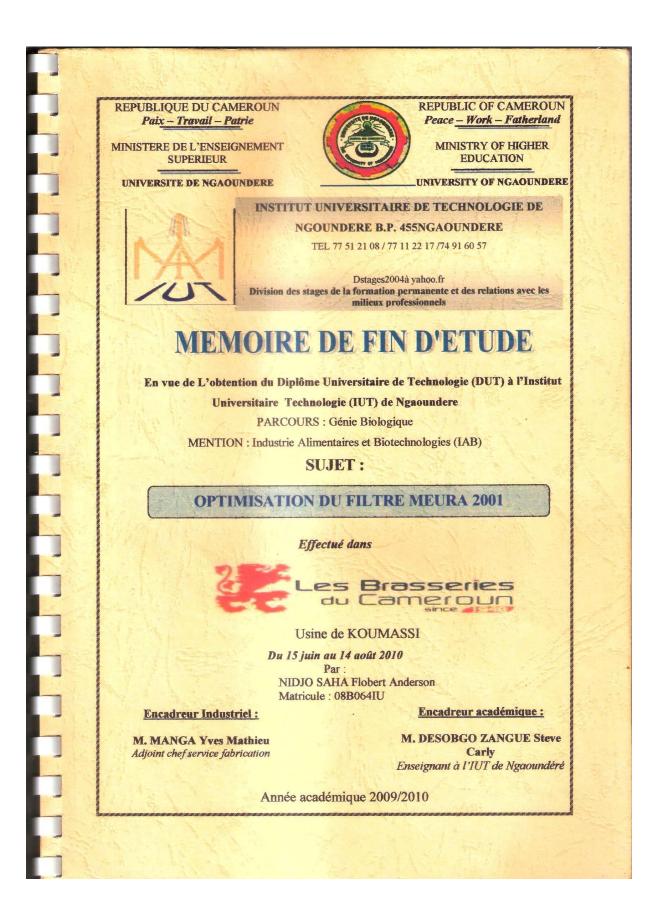
Encadreur école

**M. TAGNE SOTER Ledoux** 

Chef d'usine de CAMLAIT NDOKOTTI

**M. DESOBGO ZANGUE Steve Carly** Enseignant à l'I.U.T. de Ngaoundéré

Année académique 2009 / 2010



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**DEPARTEMENT DE GENIE BIOLOGIQUE** 

Mémoire de fin d'études en vue de l'obtention de la licence professionnelle à l'Institut Universitaire de Technologie (IUT) de Ngaoundéré

**MENTION** : Génie Biologique

PARCOURS : Génie de l'Environnement (GEN)

# Sujet : ETUDE SUR LA GESTION DES DECHETS D'UNE SAVONNERIE ET PROPOSITION D'AMELIORATION (cas d'ISF)

Effectué du 05 aout au 29 octobre 2010 à International Soap Factory BAMENDA

Par BONBO DIBE Pierre Pascal

Encadreur industriel :

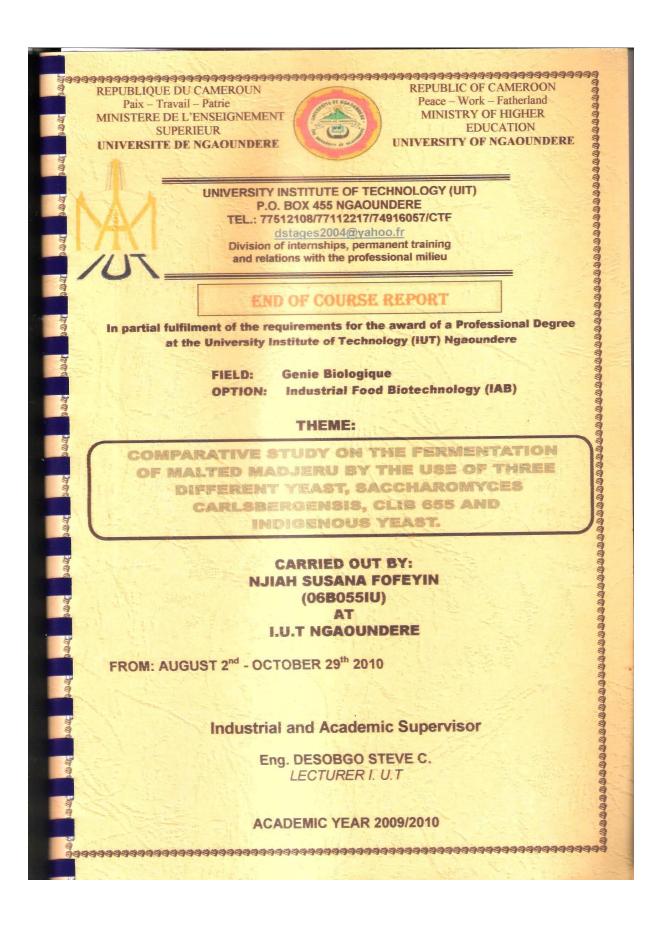
M.SITCHEU ELI

**Directeur Technique d'ISF** 

Encadreur école :

M.DESOBGO ZANGUE Steve. C Assistant à l'IUT de Ngaoundéré

Année académique 2009-2010



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# END OF COURSE INTERNSHIP REPORT

IN PARTIAL FULFILMENT OF THE REQUIREMENTS FOR THE AWARD OF A PROFESSIONAL BACHELOR'S DEGREE

IN

Food Processing and Biotechnology (IAB)

TOPIC

PROPOSAL OF A PRODUCTION UNIT AND THE RECOMMENDATTION OF THE LAND DIMENSIONS FOR THE MANUFACTURING OF BABY FOOD (kukulu) SERDIF/ PROJECT

> Done from 5th August to 29th October 2010 AT

> > SERDIF

Presented by:

# SUH QUEENIVA LUM

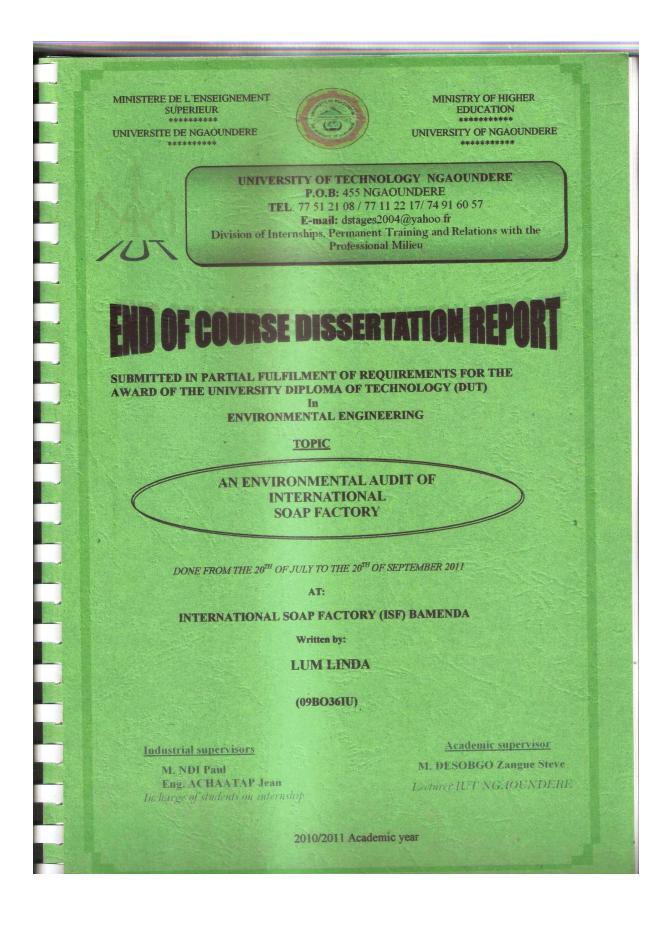
Industrial Supervisor Eng .KITO Guy Technical Director Academic Supervisor Eng.DESOBGO STEVE Lecturer IUT Ngaoundéré

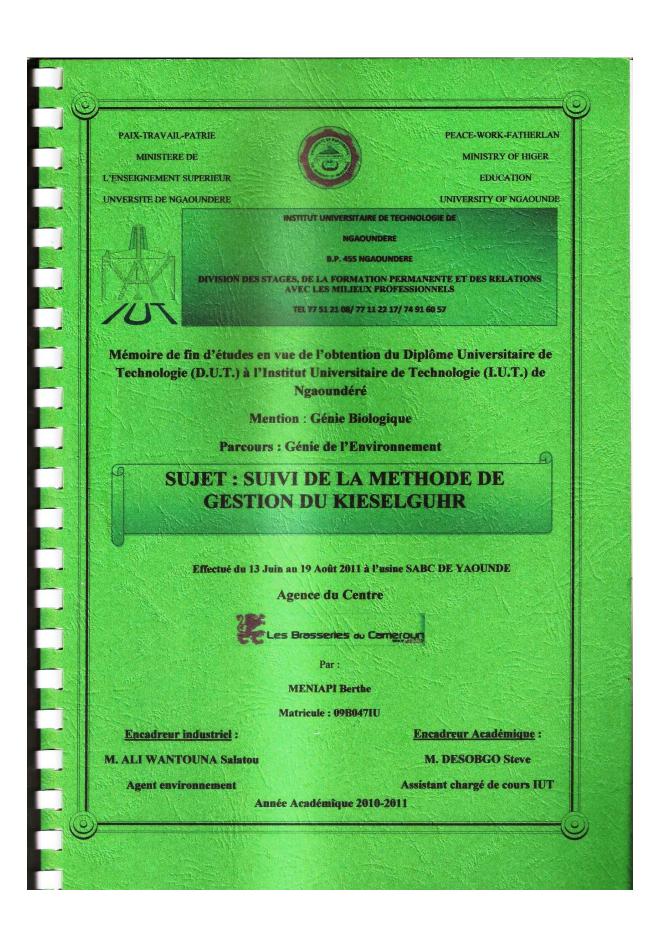
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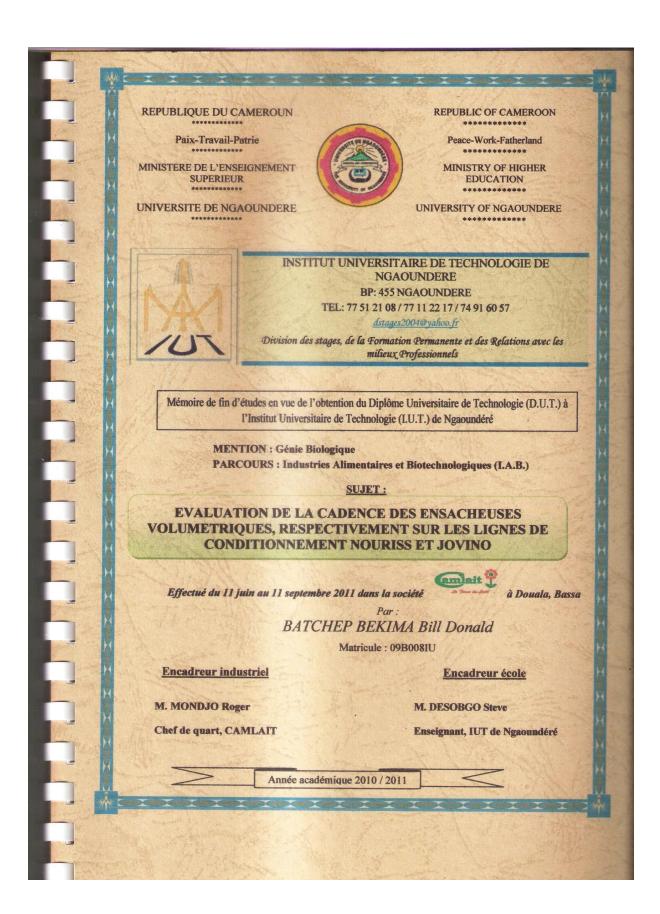
UNIVERSITY

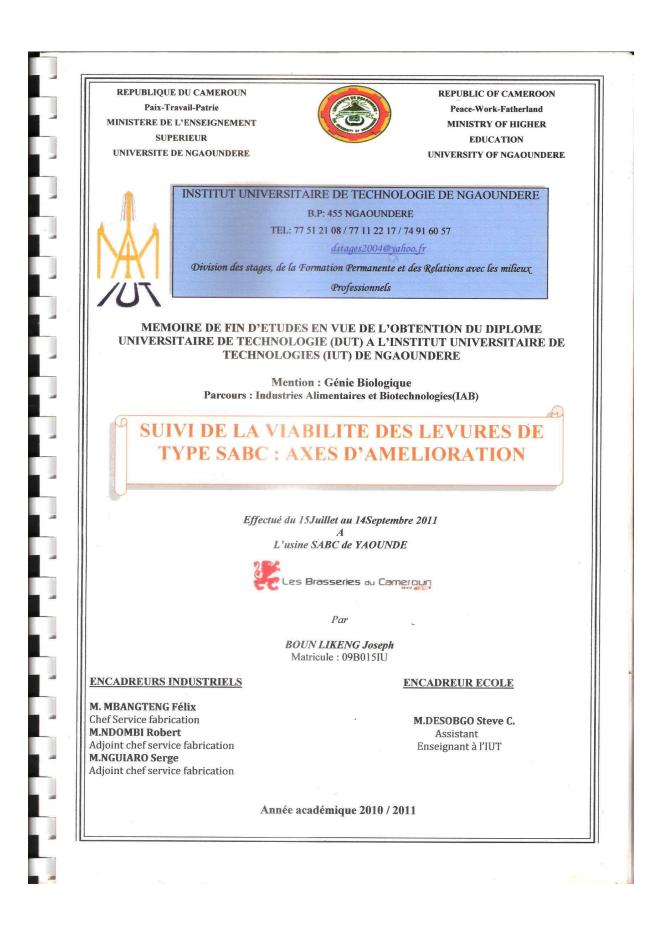
OF NGAOUNDERE

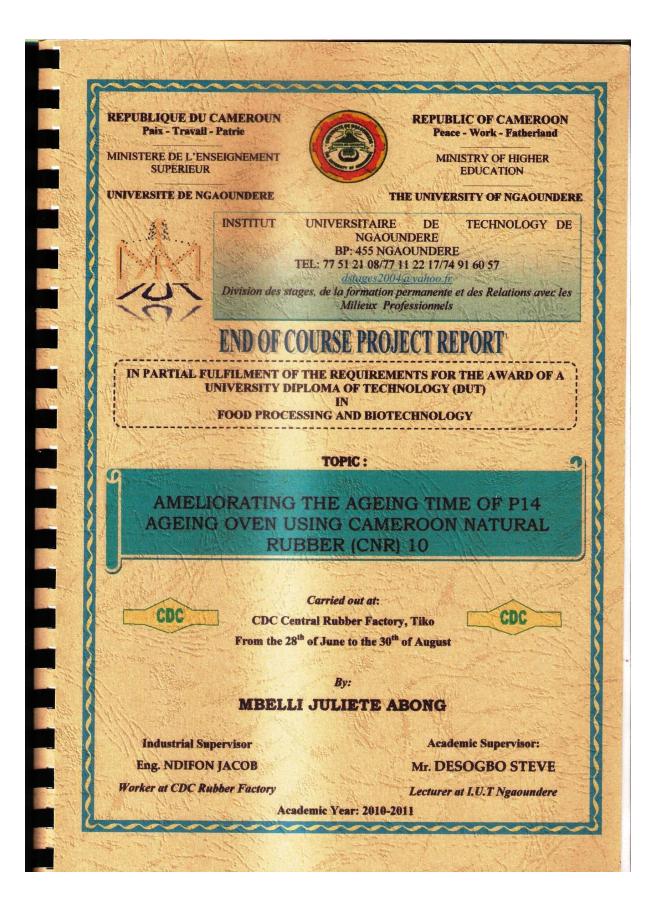
Academic year 2009/2010











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Mémoires de fin d'études en vue de l'obtention du diplôme universitaire de technologie (DUT) à l'Institut Universitaire de Technologie (IUT) de Ngaoundéré

MENTION : Génie Biologie PARCOURS : Industries Alimentaires et Biotechnologies (IAB)

Sujet :

# ETUDES DES FREINTES DES BOITES SUR LA CHAINE DE PRODUCTION DES CONSERVES DE HARICOT VERT

Effectué du 15 Juin au 01 Septembre 2011 à l'usine PROLEG S.A Par

MBETMI Charles Ghislain Matricule : 09BO42IU

Encadreur industriel COUAYA Sandrine Responsable qualité Encadreur académique M. DESOBGO Steve Enseignant à l'1.U.T

Année académique 2010 - 2011

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REPUBLIQUE DU CAMEROUN

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SUNTERIAN SUNTERI

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Mémoire de fin d'études en vue de l'obtention du Diplôme Universitaire de Technologie (DUT) à l'Institut Universitaire de Technologie (IUT) de Ngaoundéré

**MENTION : Génie Biologique** 

PARCOURS : Industries Alimentaires et Biotechnologies (IAB)

Usine de l'Ouest

SUJET : OPTIMISATION DU TEMPS D'EXECUTION AUX POSTES DES ANALYSES PHYSICOCHIMIQUES AU LABORATOIRE. CAS : POSTE DE LA PHYSICOCHIMIE BIERE

Effectué du 13 juin au 09 septembre 2011 à la

SOCIETE ANONYME DES BRASSERIES DU CAMEROUN

LES BRASSERIES DU CAMERGUN

Par

NGNOKAM NZOUKOU Claude Calixte

Matricule : 09B059IU

Encadreurs industriels

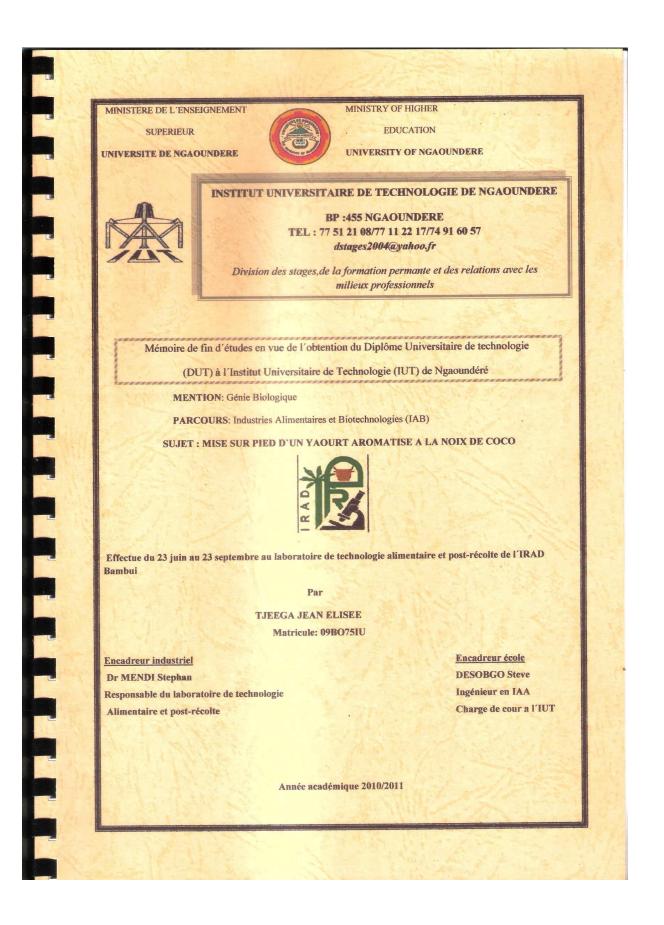
Mr MAYOH Jacques Fidèle Chef Laboratoire et Qualité

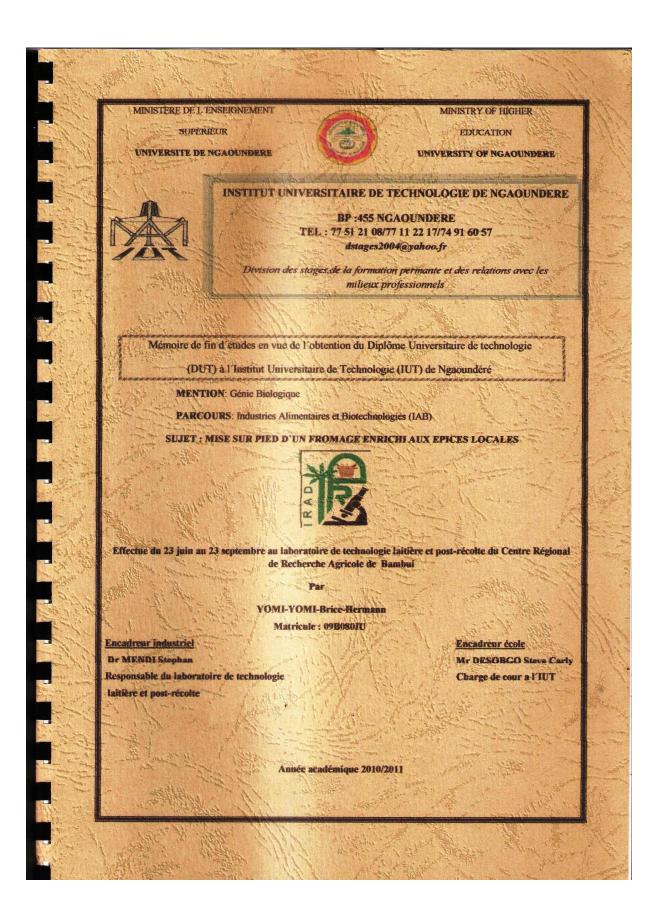
Mr BIBILA Georges Chef de groupe

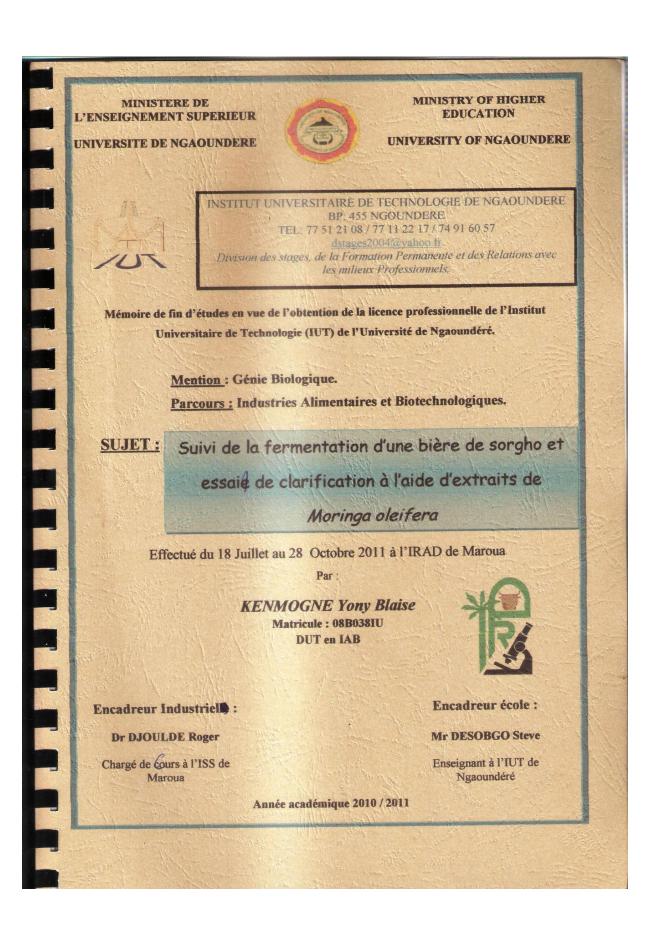
Année Académique 2010/2011

Encadreur école

Ing. DESOBGO Steve Assistant à l'IUT de Ngaoundéré







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Mémoire de fin d'études en vue de l'obtention de la Licence Professionnelle à l'Institut Universitaire de Technologie (IUT) de Ngaoundéré

MENTION : Génie Biologique PARCOURS : Industries Alimentaires et Biotechnologies (IAB)

SUJET : ETUDE DE CAPABILITE DU FILTRE A MAISCHE USINE SABC DE GAROUA

Effectué du 18 Juillet 2011 au 28 Octobre 2011 à l'usine SABC de GAROUA

Par MORNADJI Félix

Matricule : 93A009UT

Encadreur industriel NGANDO EJENGUELE Achille Ingénieur IAA Chef Service Fabrication

Encadreur école Ing. DESOBGO Steve

Enseignant IUT Ngaoundéré

Année académique 2010 / 2011

4-3) Publications dans le Grade d'Assistant

# Modelling and Optimizing of Mashing Enzymes – Effect on Yield of Filtrate of Unmalted Sorghum by Use of Response Surface Methodology

# Zangué S. C. Desobgo, Emmanuel J. Nso^{*}, Dzudie Tenin and G. J. Kayem

#### ABSTRACT

### J. Inst. Brew. 116(1), 62-69, 2010

The effect of commercial enzymes on liquefaction of starch from unmalted sorghum was studied. The effects which these enzymes had on rates of filtration were evaluated. Models were developed, validated and optimized to establish the actions of enzymes, either alone or in combination. Preliminary studies on the sorghum cultivars Safrari, Madjeru and S.35 showed that  $\alpha$ amylase was the backbone enzyme for starch liquefaction among the enzymes used ( $\alpha$ -amylase, Filtrase, protease and  $\beta$ amylase). Models confirmed this observation as α-amylase individually in its first order  $(X_1)$  contributed 25, 11 and 17%, and in its sum of first and second orders (X1+X12) contributed a 29, 31 and 36% yield of filtrate for Safrari, Madjeru and S.35 respectively. The ease of starch liquefaction, assessed by summing the first and second orders of individual intervention of all enzymes, was found to be in the order of Madjeru, S.35 and Safrari (79, 70 and 56% of yield of filtrate respectively). The importance of the enzyme combination in starch liquefaction in Safrari, S.35 and Madjeru was shown to be 44, 30 and 21% respectively. Enzyme combinations giving maximal starch liquefaction, as identified from a Doehlert experimental matrix, displayed a similar yield of filtrate (Safrari: 85 mL, Madjeru: 84 mL and S.35: 81 mL) after filtration of a 130 mL mash during 1 h. Validation of the models revealed the model developed for Madjeru was the most reliable ( $R^2 = 0.994$ ), while those developed for Safrari ( $R^2 = 0.987$ ) and S.35 ( $R^2 = 0.976$ ) were slightly less reliable. Model optimization gave theoretical enzyme (Brewers Amyliq TS, Filtrase NLC, Brewers Protease and  $\beta$ -amylase) combinations of 25 mg, 5.68 mg, 100 mg and 67.4 U for Safrari, 15.06 mg, 0.51 mg, 24.32 mg and 53.8U for Madjeru and 19.01 mg, 6.36 mg, 58.76 mg and 43.48 U for S.35, with a resulting yield of filtrate of 94, 87.7 and 83.8 mL respectively.

**Key words:** Mashing enzymes, model validation, modeling, optimization, unmalted sorghum, yields of filtrates.

# INTRODUCTION

The addition of enzymes to increase fermentable sugars and free amino acids, and to facilitate filtration when

Ecole Nationale Supérieure des Sciences Agro-industrielles (ENSAI), Université de Ngaoundéré, BP 455 Ngaoundéré, Cameroun. *Corresponding author. E-mail: nso_emmanuel@yahoo.fr

Publication no. G-2010-0316-1048 © 2010 The Institute of Brewing & Distilling mashing with poorly malted or unmalted cereals, is an established practice in beer brewing^{1,2,6,7,10,12,15}. One of the most important technological parameters to which brewers pay attention during mashing is the ease with which the mash filters^{8,10,17,19,20}. Studies carried out on malts of three sorghum cultivars of North Cameroon showed that, whether malted traditionally or under controlled laboratory conditions to brew the traditional beer Bili-Bili, the mashes of Madjeru cultivar filtered slower than the cultivars Safrari and S.3517. Although the proportions of fermentable sugars of worts of the three cultivars were comparable, that of maltose was less than 50% for Madjeru, when compared with Safrari and S.3517. These observations were partly attributed to limited amounts of starch hydrolyzing enzymes such as  $\alpha$ -amylase and  $\beta$ -amylase present in the Madjeru malts¹⁷. This is indeed one of the major problems encountered in mashing with some malted sorghum cultivars^{3-5,8,9,13,14,17,18,21}. However, it is known that acceptable worts for beer brewing can be obtained from 100% unmalted sorghum by supplementing the mashes with optimal amounts of thermostable  $\alpha$ -amylase, fungal  $\alpha$ -amylase and bacterial proteases¹¹. In contrast, what is not known is the effect of singular or combined contributions of these enzymes on starch liquefaction of unmalted sorghum grains. In this paper, we report validated and optimised mathematical models for mashing unmalted grains of the Safrari, S.35 and Madjeru sorghum cultivars using as mashing enzymes the following: Brewer Amyliq (α-amylase), Filtrase NLC (β-glucanase and hemicellulase), Brewers protease and  $\beta$ -amylase in order to assess the singular or combined contributions of these enzymes in mash liquefaction.

# MATERIALS AND METHODS

### Chemicals

The characteristics of the commercial enzymes used (Brewers Amyliq TS from *Baccillus licheniformis*, Filtrase NLC, Brewers protease from *Bacillus amyloliquefaciens* and  $\beta$ -amylase type II-B from crude barley) are presented in Table I. The amounts used for the first three ranged from 0–25 mg, 0–10 mg and 0–100 mg respectively, while for  $\beta$ -amylase the range used was 0–80 U.

	Organism of origin	Activity	Description	Temperature optima	pH optimum	Recommended application level in adjuncts	Form	
Brewers Amyliq TS	Bacillus licheni- formis	27.7 ± 6.5 U/mg of solid	α-amylase	93–95°C	5.5-6.5	0.3 g/Kg	Powder	
β-amylase (E 3.2.1.2)	Type raw II-B of barley	23-80 U/mg of protein	β-amylase	*NI ^b	*NI	*NI	Powder	
Filtrase NLC	*NI	*NI	β-glucanase and hemicellulase	*NI	*NI	0.15–0.2 g/Kg	Solution	
Brewers Protease	Bacillus amylo- liquefaciens	1842.2 ± 1.8 mg FAN/min/mL	Protease	45–50°C (de- natured at 85°C)	6.5–7.5	0.4–2 g/Kg	Solution	

^a All the commercial enzymes used in this study were obtained from DSM Food Specialities, Cedex France, apart from β-amylase which was sourced from SIGMA CHIMIE, Cedex, France.

^bNot indicated by DSM Food Specialities France/SIGMA.

Coded values			Transformed experimental values													
α-amy- lase	Filtrase	Pro- teases	β-amy- lase	α-amy- lase (mg)	Fil- trase (mg)	Pro- teases (mg)	β-amy- lase (U)	Madjeru			Safrari			S.35		
$\mathbf{X}_1$	$X_2$	X ₃	$X_4$	$\mathbf{X}_{1}$	$X_2$	X ₃	$X_4$	Exp ^a	Theo ^b	Res ^c	Exp	Theo	Res	Exp	Theo	Res
1.000	0.000	0.000	0.000	25	5	50	40	68	65	3	85	83	2	76	76.1	-0.1
-1.000	0.000	0.000	0.000	0	5	50	40	24	27	-3	36	38	-2	35	34.9	0.1
0.500	0.866	0.000	0.000	18.75	10	50	40	61	60.405	0.595	67	70.001	-3	77	78.404	-1.4
-0.500	-0.866	0.000	0.000	6.25	0	50	40	64	64.596	-0.59	60	56.999	3.001	60	58.602	1.398
0.500	-0.866	0.000	0.000	18.75	0	50	40	84	83.097	0.903	71	71.5	-0.5	69	67.702	1.298
-0.500	0.866	0.000	0.000	6.25	10	50	40	40	40.905	-0.9	40	39.502	0.498	45	46.304	-1.3
0.500	0.289	0.816	0.000	18.75	6.66	100	40	55	57.088	-2.08	75	75.698	-0.69	80	79.993	0.007
-0.500	-0.289	-0.816	0.000	6.25	3.33	0	40	63	60.885	2.115	59	58.297	0.703	61	60.991	0.009
0.500	-0.289	-0.816	0.000	18.75	3.33	0	40	74	76.885	-2.88	71	69.789	1.211	77	72.579	4.421
0.000	0.577	-0.816	0.000	12.5	8.33	0	40	61	60.29	0.71	64	62.805	1.195	70	69.897	0.103
-0.500	0.289	0.816	0.000	6.25	6.66	100	40	38	35.088	2.912	41	42.19	-1.19	46	50.381	-4.38
0.000	-0.577	0.816	0.000	12.5	1.66	100	40	60	60.685	-0.68	66	67.205	-1.2	69	69.097	-0.09
0.500	0.289	0.204	0.791	18.75	6.66	62.5	80	55	56.459	-1.45	75	73.301	1.699	81	79.483	1.517
-0.500	-0.289	-0.204	-0.791	6.25	3.33	37.5	0	35	33.458	1.542	47	48.7	-1.7	45	46.483	-1.48
0.500	-0.289	-0.204	-0.791	18.75	3.33	37.5	0	63	63.97	-0.97	65	67.693	-2.69	67	72.583	-5.58
0.000	0.577	-0.204	-0.791	12.5	8.33	37.5	0	42	42.372	-0.37	53	51.701	1.299	65	62.394	2.606
0.000	0.000	0.612	-0.791	12.5	5	87.5	0	49	49.078	-0.07	54	50.894	3.106	67	62.491	4.509
-0.500	0.289	0.204	0.791	6.25	6.66	62.5	80	50	48.971	1.029	50	47.293	2.707	70	64.383	5.617
0.000	-0.577	0.204	0.791	12.5	1.66	62.5	80	70	69.565	0.435	62	63.299	-1.29	73	75.592	-2.59
0.000	0.000	-0.612	0.791	12.5	5	12.5	80	76	75.876	0.124	55	58.093	-3.09	72	76.489	-4.48
0.000	0.000	0.000	0.000	12.5	5	50	40	78	77.25	0.75	63	64.25	-1.25	79	78.25	0.75
0.000	0.000	0.000	0.000	12.5	5	50	40	79	77.25	1.75	65	64.25	0.75	77	78.25	-1.25
0.000	0.000	0.000	0.000	12.5	5	50	40	75	77.25	-2.25	63	64.25	-1.25	77	78.25	-1.25
0.000	0.000	0.000	0.000	12.5	5	50	40	77	77.25	-0.25	66	64.25	1.75	80	78.25	1.75

^a Experimental result values.

^bTheoretical values (values coming from mathematical models).

c Residue.

### Sorghum cultivars

The *Safrari, Madjeru* and *S.35* sorghum cultivars were obtained from the Institute of Research and agronomic development (IRAD) Maroua, Cameroon.

### Mashing

Sorghum cultivar grains were milled to particle sizes of 0.7 mm or less using a hammer mill Polymix PX-MFC 90D apparatus type (VWR International S.A.S. Le Périgas 201, rue Carnot, 94126 Fontenay-sous-Bois Cedex, France). Twenty five grams of unmalted sorghum was weighed, placed in a 600 mL beaker and 150 mL of distilled water added. The suspension was homogenised at 24°C by stirring with a glass rod. It was then heated to boiling temperature, at which starch gelatinization was

allowed to take place for 20 min, with intermittent stirring at intervals of 5 min, before cooling to 60°C. Mashing was carried out at 60°C for 1 h taking into consideration the conditions indicated in Table I. The mash was cooled to 25°C and filtered for 1 h using Whatman grade 1 filter paper.

#### Mathematical modelling

The establishment of an experimental matrix was necessary to develop the mathematical model. The Doelhert (number of factors, k = 4)¹⁶ matrix was adopted for this work. Transformation of the matrix of coded variables to the experimental matrix was automatic (Table II), while the quantities of mashing enzymes used were maintained at fixed levels. The amounts of selected enzymes ranged as follows: Brewers Amyliq TS (0–25 mg); Filtrase NLC (0–10 mg); Protease (0–100 mg); and  $\beta$ -amylase (0–80 U) ( $\beta$ -amylase activity was 31 U/mg of solid).

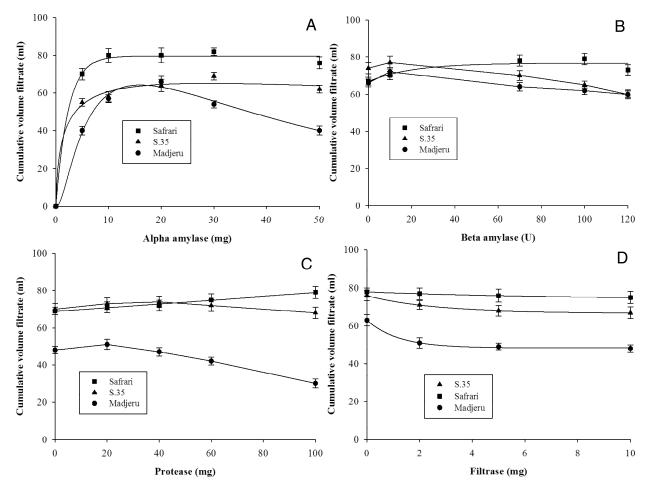
Doelhert's experiment design (having a homogenous distribution in space) was used to establish the experiment matrix (Table II). This matrix has coded variables and must be converted into an experimental matrix¹⁶ having real variables directly usable in the laboratory. The results of the phenomenon to study are then established and the data considered as " $y_{exp}$ ". With the help of the matrix of coded variables and " $y_{exp}$ ", the coefficients of the model are obtained. The model is obtainable only from the coded variable matrix because, in this case, there is no need for a counterbalancing effect of the factors in study¹⁶. The coefficients of the model obtained will thus be effectively linked to the impact of each factor. These coefficients and the model were obtained with the help of the Systat version 12 software (Systat Software, Inc., San Jose, USA). This software also gives a statistical analysis on the model. Lastly, the curves are plotted using Sigmaplot version 11 build 11.0.0.77 software (WPCubed, GmbH, Germany).

Validation of the model was conducted after assays using several enzyme combinations found in the experimental domains not explored by the experimental matrix. Another method consisted of tracing the theoretical results against the experimental results, as the coefficient of correlation  $R^2$  gives an appreciation of the reliability of the model.

Model optimization used Mathcad version 14 software (Parametric Technology Corporation, Massachusetts, USA). Optimal combination was obtained by initially entering the model and then specifying the starting point of each factor. The sweeping interval by the software was given for each factor. Once the data was entered, the software gave a response for a maximum or minimum combination as requested for. This theoretical optimum was then explored for the confirmation of the optimal point.

# **RESULTS AND DISCUSSION**

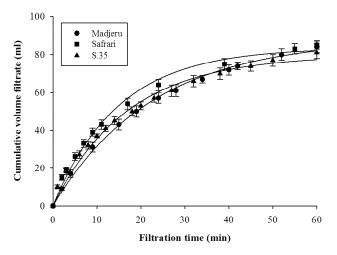
Before establishing the mathematical model for mashing, trials were carried out to investigate the action of each of the enzyme preparations in starch liquefaction



**Fig. 1.** (A) Effect of  $\alpha$ -amylase concentration on filtration of unmalted sorghum mash. The  $\beta$ -amylase, Protease, and Filtrase being constant at 80 U, 60 mg and 10 mg respectively. (B) Effect of  $\beta$ -amylase concentration on filtration of unmalted sorghum mash. The  $\alpha$ -amylase, Protease and Filtrase being constant at 20 mg, 60 mg and 10 mg respectively. (C) Effect of Protease concentration on filtration of unmalted sorghum mash. The  $\alpha$ -amylase, Filtrase and  $\beta$ -amylase were constant at 20 mg, 10 mg and 80 U respectively. (D) Effect of Filtrase concentration on filtration of unmalted sorghum mash. The  $\alpha$ -amylase, Filtrase and  $\beta$ -amylase, Protease and  $\beta$ -amylase were constant at 20 mg, 10 mg and 80 U respectively. (D) Effect of Filtrase and  $\beta$ -amylase were constant at 20 mg, 60 mg and 80 U respectively.

during mashing. Among the enzyme preparations used in this study, it was observed (Fig. 1a) that  $\alpha$ -amylase was the principal enzyme of starch liquefaction during mashing^{1,7,12}. Figures 1b, 1c and 1d show that the actions of  $\beta$ amylase, filtrase and protease on starch liquefaction were only noticeable in the presence of  $\alpha$ -amylase. This is justified by the fact that filtration was not possible after mashing in the presence of these enzymes, but in the absence of  $\alpha$ -amylase (Fig. 1a). These results indicated that mathematical modeling would provide more precise information on the contribution of each enzyme component and combination of enzyme components required to liquefy the mash optimally.

Previous studies on malting and mashing of sorghum cultivars used in this study showed that Madjeru mashes filtered slower than those of Safrari and S.3517,18. This slow filtration rate was suggested to be due to the low amounts of  $\alpha$ -amylase and  $\beta$ -amylase in *Madjeru* malts compared to Safrari and S.35. Experimental conditions for the three cultivars in this study were standardized by using unmalted grains and several combinations of the mashing enzymes as shown in Table II. The best enzyme combinations to mash with were identified and the rates of filtration compared under those conditions (Fig. 2). The results showed that the filtration rates, and thus the yields of filtrates for the three varieties were comparable as 85 mL, 84 mL and 81 mL of filtrate was obtained after filtration of 130 mL for 1 h of the Safrari, Madjeru and S.35 mash, respectively. These results confirmed the hypothesis that the slow filtration of mashes of Madjeru malts during mashing could be partly attributed to insufficient quantities of  $\alpha$ -amylase and  $\beta$ -amylase as suggested previously¹⁷. A mathematical model was established for each cultivar using a precise matrix. Models obtained by the response surface method established the following equations for mashing the three cultivars.



**Fig. 2.** The effect of best combinations of enzymes on mash filtration: *Madjeru* [α-Amylase (18.75 mg), Filtrase (0 mg), Protease (50 mg), β-amylase (40 U)]; *Safrari* [α-amylase (25 mg), Filtrase (5 mg), Protease (50 mg), β-amylase (40 U)]; *S.35* [α-amylase (18.75 mg), Filtrase (6.66 mg), Protease (62.5 mg), β-amylase (80 U)].

### Safrari

$$\begin{split} Y &= 64.25 + 22.5X_1 - 5.484X_2 - 1.182X_3 + 3.636X_4 + \\ 9.237X_1X_2 + 10.219X_1X_3 - 1.576X_1X_4 - 5.437X_2X_3 + \\ 3.470X_2X_4 + 8.655X_3X_4 - 3.75X_1^2 - 5.083X_2^2 - 0.169X_3^2 - \\ 8.796X_4^2 \end{split}$$

### Madjeru

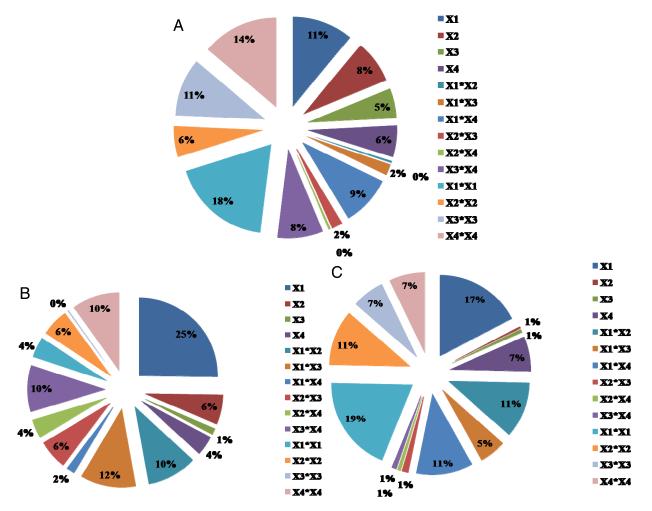
 $\begin{array}{l} Y = 77.25 \ + \ 19X_1 \ - \ 13.39X_2 \ - \ 9.226X_3 \ + \ 9.801X_4 \ + \\ 0.577X_1X_2 \ + \ 3.472X_1X_3 \ - \ 15.66X_1X_4 \ + \ 3.427X_2X_3 \ - \\ 0.638X_2X_4 \ - \ 14.23X_3X_4 \ - \ 31.25X_1^2 \ - \ 9.583X_2^2 \ - \ 17.96X_3^2 \ - \ 23.87X_4^2 \end{array}$ 

### S.35

 $\begin{array}{l} Y = 78.25 + 20.6X_1 - 0.461X_2 - 0.816X_3 + 8.217X_4 + \\ 13.279X_1X_2 + 6.342X_1X_3 - 13.44X_1X_4 - 1.536X_2X_3 - \\ 0.822X_2X_4 + 1.147X_3X_4 - 22.75X_1^2 - 13.08X_2^2 - 7.692X_3^2 - \\ - 8.508X_4^2 \end{array}$ 

The three equations were structurally identical and all of second order with interactions. In the models, the factors  $X_1$  and  $X_4$ , corresponding to  $\alpha$ -amylase and  $\beta$ -amylase respectively had positive coefficients. These indicated that the two enzymes were directly involved in the liquefaction of starch and were thus instrumental contributors to yields of filtrates. Figures 3a, 3b and 3c detail the individual and combined contributions of the various mashing enzyme components on the total yield of filtrates of the liquefied starch. Thus,  $\alpha$ -amylase, in its singular expression as first order in the model, contributed 25%, 11% and 17% of the yield of filtrate for the mashes of Safrari, *Madjeru* and S.35 respectively. Similarly,  $\beta$ -amylase accounted for 4%, 6% and 7% respectively. The factors  $X_2$ and X₃, corresponding to filtrase and protease respectively, had negative coefficients. This suggested that these enzymes were not directly responsible for starch liquefaction, and cannot, a priori, be considered as directly contributing to yield of filtrate during mashing. Filtrase, in its singular expression as first degree in the model, indirectly contributed to 6%, 8% and 1% in yields of filtrates, while protease contributed to 1%, 5% and 1% in Safrari, Mad*jeru* and S.35 mashes respectively. The factors  $X_1^2$ ,  $X_2^2$ ,  $X_3^2$  and  $X_4^2$  corresponding to the second order of the components of  $\alpha$ -amylase, filtrase, protease and  $\beta$ -amylase respectively, showed that the singular action of all these enzymes expressed to their second order in the model, had a negative coefficient. Their absolute contribution to the yield of filtrate was about 4%, 6%, 0% and 10% for Safrari, 20%, 5%, 10% and 14% for Madjeru and 19%, 11%, 7% and 7% for S.35 mashes respectively. Table III summarises these results, with the sum totals in percent, of the contribution in yield of filtrate by these enzymes in their first and second orders as expressed by the model. The results corroborated those shown in Fig. 1, wherein  $\alpha$ -amylase is evidently the sole enzyme of starch liquefaction during mashing. Indeed, this enzyme in its first and second orders in the model, singularly contributed 29%, 31% and 36% to filtrate yield of the cultivars Safrari, *Madjeru* and S.35 respectively.

Table III also summarises the results for the contributions of the interactions between various enzyme components used. Thus, the interaction  $X_1X_2$ , corresponding to the combined action of  $\alpha$ -amylase and filtrase, had a pos-



**Fig. 3.** (A) Contribution of the model factors on the total effect for *Safrari*. (B) Contribution of the model factors on the total effect for *Madjeru*. (C). Contribution of the model factors on the total effect for *S.35*.

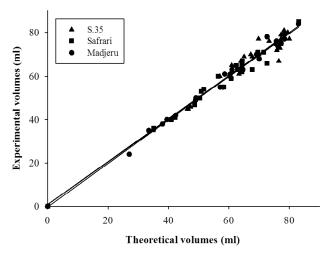
itive coefficient, suggesting that the second enzyme contributed to reducing the viscosity of the mash and thus facilitated its filtration. This positive interaction between  $\alpha$ -amylase and filtrase roughly contributed to 10%, 0% and 11% in yield of filtrate for Safrari, Madjeru and S.35 mashes respectively. The interaction  $X_1X_3$ , corresponding to the combined action of  $\alpha$ -amylase and protease, also had a positive coefficient. Similarly to the  $\alpha$ -amylase/filtrase interaction, protease facilitates mash hydrolysis by  $\alpha$ -amylase. This positive interaction between the two enzymes contributed to about 12%, 2% and 5% in yield of filtrate for Safrari, Madjeru and S.35 mashes respectively. The interaction  $X_1X_4$ , corresponding to the combined action of  $\alpha$ -amylase and  $\beta$ -amylase had a negative coefficient. Since these two enzymes are starch hydrolysing enzymes, they independently and directly attack free convertible starch granules in suspension without any prerequisites. Whereas starch granules embedded in the protein matrix and surrounded by  $\beta$ -glucano-hemicellulosic cell walls must absolutely be liberated by the action of proteases and  $\beta$ -glucanases in order to allow  $\alpha$ -amylase and  $\beta$ -amylase to act on starch²¹. The contribution by this combined effect in yield of filtrate was 2%, 9% and 11% for Safrari, Madjeru and S.35 mashes respectively. Upon adding the contributions of  $\alpha$ -amylase's action in its singular first and second orders, to those of its interactions with the other enzymes ( $\alpha$ -amylase/filtrase,  $\alpha$ -amylase/ protease and  $\alpha$ -amylase/ $\beta$ -amylase) in the model, we obtained the direct and indirect contribution to starch liquefaction. These contributions amounted to 53%, 42% and 63% for Safrari, Madjeru and S.35 respectively. The interaction X₂X₃, corresponding to the combined action of filtrase and protease, had a negative coefficient for the mashes of the cultivars Safrari and S.35, but was positive for Madjeru. The contribution by this combined effect in yield of filtrate was roughly 6%, 2% and 1% for Safrari, Madjeru and S.35 mashes respectively. These two enzymes are not starch hydrolysing enzymes, and only help to liberate starch granules embedded in the protein matrix and surrounded by  $\beta$ -glucano-hemicellulosic cell walls. The low contribution of this interaction is therefore predictable. Interaction  $X_2X_4$ , corresponding to the combined action of filtrase and  $\beta$ -amylase, had a positive coefficient for the cultivar Safrari, but was negative for Madjeru and S.35. The contribution by this combined effect in yield of filtrate was about 4%, 0% and 1% for Safrari, Madjeru and S.35 mashes respectively. Once more, the low contribution of this interaction for the three cultivars suggested that the filtrase/*β*-amylase combination was not important for filterability. This can be explained by the fact that the

	Single component as 1st,		Cultivar typ	e
Enzyme	2nd degree and sum	Safrari	Madjeru	S.35
α-amylase	$\mathbf{X}_1$	25	11	17
	$X_{1}^{2}$	4	20	19
	$X_1 + X_1^2$	29	31	36
Filtrase	$\mathbf{X}_2$	6	8	1
	$X_{2}^{2}$	6	5	11
	$X_2 + X_2^2$	12	13	12
Protease	$X_3$	1	5	1
	$X_{3}^{2}$	0	10	7
	$X_3 + X_3^2$	1	15	8
β-amylase	$\mathbf{X}_4$	4	6	7
	$X_4^2$	10	14	7
	$X_4 + X_4^2$	14	20	14
Total		56	79	70
Enzymes interacting	Combined component	Safrari	Madjeru	S.35
α-amylase / Filtrase	$X_1 X_2$	10	0	11
α-amylase / Protease	$X_1 X_3$	12	2	5
α-amylase / β-amylase	$X_1 X_4$	2	9	11
Filtrase / Protease	$X_2 X_3$	6	2	1
Filtrase / β-amylase	$X_2 X_4$	4	0	1
Protease / β-amylase	$X_3 X_4$	10	8	1
Total		44	21	30

 Table III. Contributions of single (1st and 2nd orders), their sums and combined enzyme components to filtrate yields.

role of filtrase was limited to unmasking starch granules embedded in cells by hydrolysing the  $\beta$ -glucans and hemicelluloses of the walls of these cells, while  $\beta$ -amylase, a saccharifying enzyme, had only limited action on the molecular size of starch material. Finally, the interaction  $X_3X_4$  corresponding to the combined action of protease/ $\beta$ amylase had a positive coefficient for the cultivars Safrari and S.35, but negative for the cultivar Madjeru. The contribution of this combined effect on yield of filtrate was about 10%, 8% and 1% for Safrari, Madjeru and S.35 respectively. The impact of this enzyme combination on the filtration of mash will depend not only on the extent of hydrolysis of the protein matrix surrounding starch granules, but also on that of the  $\beta$ -amylase component, which will be assessed by the extent of hydrolysis of convertible starch granules. Thus, if proteolysis is effective, but the starch type is more of amylopectin than amylose, the rate of mash filtration would be slower, as the saccharifying action of  $\beta$ -amylase would be reduced. This could explain the disparity in the yield of filtrate observed for these three cultivars of sorghum, with respect to the enzyme combinations.

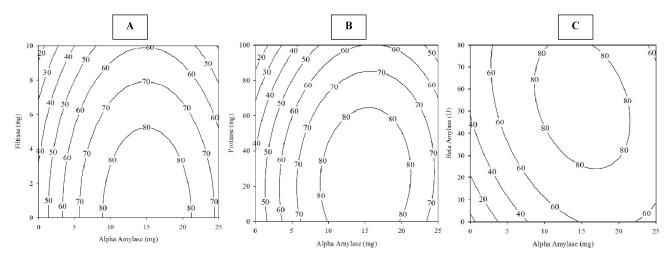
With respect to the observed effects of the first and second orders as a sum for each of the enzyme components in the model, 56%, 79% and 70% for Safrari, Mad*jeru* and S.35 respectively, it is clear that the ease of mash liquefaction was in the order of Madjeru, S.35 and Safrari (Table III). Also, the model showed that the ease of mash liquefaction on the basis of enzyme combined components was in the order of Safrari, S.35 and Madjeru (44%, 30% and 21% respectively) (Table III). These results suggest that the action of filtrase and protease in liberating protein embedded starch granules surrounded by  $\beta$ -glucano-hemicellulosic cell walls, was of great importance in liquefying and saccharifying starch by  $\alpha$ -amylase and  $\beta$ amylase respectively for Safrari, followed by S.35 and then by Madjeru. This also suggests that Safrari was richer in proteins and/or β-glucans than S.35. This has in-



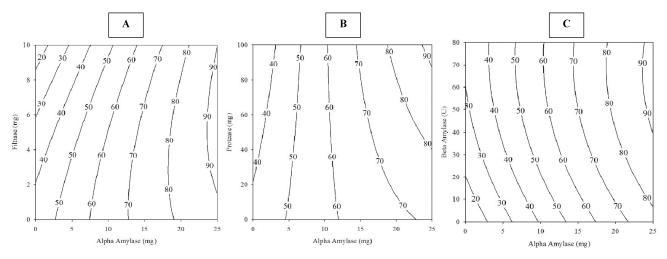
**Fig. 4.** Validation of mathematical models for *Madjeru, Safrari* and *S.35* with  $R^2 = 0.994$ , 0.987 and 0.976 respectively.

deed been shown with respect to the total protein contents of these three cultivars¹⁷.

The mathematical models were validated using two methods. Firstly, tests were carried out on several combinations of the experimental space which were not explored within the framework of the experimental matrix. The results were then compared with the theoretical results and the errors statistically evaluated. These errors were between 2% and 3.7%, 1.7% and 3%, and 0.1% and 3.4% for *Safrari, Madjeru* and *S.35* respectively. The global error thus was 0.1% to 3.7%. As the highest error limit is approximately 3.7%, it could be concluded that the mathematical models established, satisfactorily describe the observed phenomena. It is however necessary to determine this error and reliability. This can be done by a second method consisting of plotting the theoretical results against the experimental results and determining the



**Fig. 5.** Response surface curves for *Madjeru* yield in filtrates [all other factors were fixed at optimal quantity: (A) Protease: 24.32 mg;  $\beta$ -amylase: 53.8 U; (B) Filtrase: 0.51 mg;  $\beta$ -amylase: 53.80U; (C) Filtrase: 0.51 mg; Protease: 24.32 mg].



**Fig. 6.** Response surface curves for *Safrari* yield in filtrates [all other factors were fixed at optimal quantity: (A) Protease: 100 mg;  $\beta$ -amylase: 67.40 U; (B) Filtrase: 5.68 mg;  $\beta$ -amylase: 67.40 U; (C) Filtrase: 5.68 mg; Protease: 100 mg].

coefficient of correlation  $R^2$ , in order to appreciate the reliability of the models. The second method of validating the models allowed for the classification of reliability in the following order: first *Madjeru* ( $R^2 = 0.994$ ), followed by *Safrari* ( $R^2 = 0.987$ ) and then *S.35* ( $R^2 = 0.976$ ) (Fig. 4).

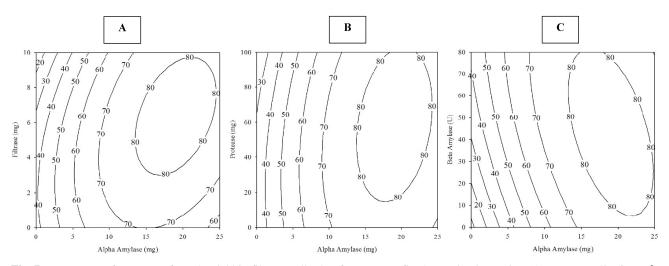
Optimization of the outputs in filtrates was logically the final step in this work after validating the models. This was conducted using Mathcad version 14 software (Parametric Technology Corporation, Massachusetts, USA). The theoretical maxima for the combinations in amounts of dispensable mashing enzyme preparations, in order to obtain optimal filtrates in our working conditions, were for  $\alpha$ -amylase, filtrase, protease and  $\beta$ -amylase, as follows: 25 mg, 5.68 mg, 100 mg and 67.4 U (or in coded values: 1, 0.118, 0.816, 0.542) respectively for Safrari, 15.06 mg, 0.51 mg; 24.32 mg and 53.80 U (or in coded values: 0.205, -0.777, -0.419, 0.273) respectively for Madjeru and: 19.01 mg, 6.36 mg, 58.76 mg and 43.48 U (or in coded values: 0.521, 0.236, 0.143, 0.069) respectively for S.35. Figs. 5, 6 and 7 show these results in their response surface representations. Thus, the theoretical optimal vol-

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umes calculated for the three models were 94, 87.7 and 83.8 mL for the three cultivars respectively. It is however important to note that the optima obtained were with respect to cooking, cooling and then mashing at 60°C as described in the Materials and Methods. The response surface methodology will obviously give different results if other temperature regimes taking into consideration the high thermostability of the  $\alpha$ -amylase (temperature optimum at 93–95°C) are applied.

## CONCLUSIONS

The  $\alpha$ -amylase was the main enzyme component responsible for starch liquefaction during the mashing of unmalted sorghum. The  $\beta$ -amylase, Filtrase and protease served basically as supporting enzymes. The release and hydrolysis (liquefaction) of starch were facilitated by the sequential actions of filtrase, which hydrolysed cell wall materials, and protease, which facilitated the breakdown of released protein materials and thus enhanced starch release. The  $\beta$ -amylase complements the dominant action of  $\alpha$ -amylase in starch liquefaction and extract develop-



**Fig. 7.** Response surface curves for *S.35* yield in filtrates [all other factors were fixed at optimal quantity: (A) Protease: 58.76 mg;  $\beta$ -amylase: 43.48 U. (B) Filtrase: 6.36 mg;  $\beta$ -amylase: 43.48 U. (C) Filtrase: 6.36 mg; Protease: 58.76 mg].

ment. In general, the response surface methodology appears to be a very reliable tool in assessing the scope and the actions of mashing enzymes, as single or combined components, in starch liquefaction during mashing. This methodology should also be helpful in predicting the development of important wort parameters such as soluble nitrogen, free  $\alpha$ -amino nitrogen, extract content, fermentability and viscosity.

#### ACKNOWLEDGEMENTS

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Full Length Research Paper

# The response surface methodology as a reliable tool for evaluating the need of commercial mashing enzymes for alleviating the levels of reducing sugars of worts of malted sorghum: Case of the *Safrari* cultivar

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A three factor Doehlert design was used to develop a statistical model to optimize the action of three commercial mashing enzymes (Hitempase 2XL, Bioglucanase TX and Brewers protease) on reducing sugars content of the worts of unmalted and malted *Safrari* sorghum. The response surface methodology revealed that increasing amounts of Hitempase considerably increased reducing sugars content during mashing of unmalted *Safrari* sorghum grist to about 90 g/L. Limited amounts of reducing sugars were obtained with increasing concentrations of both Bioglucanase ( $\approx 40$  g/L) and Brewers Protease ( $\approx 30$  g/L). The milling process facilitated the dissolution of about 10 g/L in yields of reducing sugars for the unmalted *Safrari* sorghum mash type without the help of enzyme. None of the three enzymes as sole mashing enzyme appeared to be of use in mashing malted *Safrari*, as reducing sugars yields were at maximum (168 g/L) after dissolving the grist in water and rather decreased with increasing amounts of enzyme supplements. Optimizing the concerted actions of the three enzymes for reducing sugars content of unmalted *Safrari* sorghum mash gave a combination of 2163 U, 937.5 BGU and 0 mg for Hitempase, Bioglucanase and Brewers Protease respectively. This gave a maximal reducing sugars content of 126.57 g/L. This combination was 0 U, 137.48 BGU and 0 mg for malted *Safrari* sorghum mash, giving a maximal reducing sugars yield of 168.56 g/L.

Key words: Response surface methodology, optimization, mashing enzymes, Safrari, reducing sugars

## INTRODUCTION

Sorghum (*Sorghum bicolor (L.) Moench*) is a vital caloriebased food component in human nutrition in some parts of Africa (Taylor, 2004). Besides this fundamental function, sorghum is used in the production of beer, traditional opaque beer and non-alcoholic drinks in developing countries, and also industrial beer (Palmer, 1989; Taylor and Dewar, 2001). The poor developmental profile of the principal hydrolytic enzymes during malting of this cereal is however a limiting factor to easy mashing of its malts as compared to barley malt (EtokAkpan and Palmer, 1990; EtokAkpan, 1992). This was ascribed to the malting procedures and varietal types of sorghum

used. Work on some popular sorghum cultivars of Northern Cameroon used in brewing the traditional beer Bili-Bili confirmed that the profile of hydrolytic enzymes and the levels of fermentable sugars during mashing were indeed cultivar-dependent. The Safrari and S.35 sorohum cultivars were shown to be poorer in starch than the Madjeru cultivar, but had higher hydrolytic enzymes profiles and fermentable sugars potentials (Nso et al., 2003; Nso et al., 2006). The use of commercial enzymes when mashing with sorghum in order to obtain better wort specifications for beer brewing has however become a common practice (MacFadden and Clayton, 1989;Dale et al., 1990; Bajomo and Young, 1992; Agu and Palmer, 1998; Goode et al., 2002; Goode and Arendt, 2003; Goode et al., 2003). However, it is not clear whether the use of mashing enzyme supplements in Safrari sorghum

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cultivar mashes are indispensable in alleviating the levels of fermentable sugars in its worts. In this work, the action of three principal commercial mashing enzymes (Hitempase 2XL, Bioglucanase TX and Brewers Protease) on the reducing sugars content of worts from the *Safrari* sorghum cultivar (unmalted and malted) was modeled and optimized using the response surface methodology (RSM) to verify the necessity of mashing enzymes in mashing with this cultivar.

#### MATERIALS AND METHODS

#### Enzymes

The characteristics of the commercial mashing enzymes used, hitempase  $2xI_{\mbox{\tiny B}}$ , a thermo stable  $\alpha$ -amylase from Baccillus licheniformis, Brewers protease from bacillus amyloliquefaciens and bioglucanase tx, from an enzymatic composition of  $\beta$ -glucanase and hemicellulases from trichoderma reesei) and their sources are presented in Table 1.

#### Sorghum cultivar

The *Safrari* sorghum cultivar was obtained from the Institute of Research and Agronomic Development (IRAD) Maroua, Cameroon.

#### Modeling

Modeling was carried out as previously described (Desobgo et al., 2010). A Doehlert matrix design of 3 factors representing Hitempase 2XL ( $X_1$ ), Bioglucanase TX ( $X_2$ ) and Brewers Protease ( $X_3$ ) at ranges of [0 to 3000 U], [0 to 937.5 BGU] and [0 to 100 mg] respectively, was used. The transformed matrix of coded variables to an experimental matrix and the desired response (reducing sugars) are shown in Table 2. Mathematical models describing the relationships among the process dependent variable and the independent variables in a second-order equation were developed (Giovanni, 1983). Design-based experimental data were matched according to the following second-order polynomial Equation [1].

$$y = \beta_0 + \sum_{j=1}^k \beta_j x_j + \sum_{j=1}^k \beta_{jj} x_j^2 + \sum_{i< j} \beta_{ij} x_i x_j + \varepsilon$$
[1]

where, I and j, are linear and quadratic coefficients respectively, while ' $\beta$ ', the regression coefficient, k the number of factors studied and optimized in the experiment and ' $\epsilon$ ' is random error.

The coefficients of the models and the models were obtained using the Systat version 12 software (Systat Software, Inc., San Jose, USA). This software also gives a statistical analysis on the model. Lastly, the curves were plotted using Sigmaplot version11 build 11.0.0.77 software (WPCubed, GmbH, Germany).

#### Validation of models

The quality of fit of the second order equations was expressed by the coefficient of determination  $R^2$ . The models were validated using two differents methods. The first method was the Absolute Average Deviation (AAD) method (Bas and Boyac, 2007), while the

second method consisted in applying the bias factor and accuracy factor (Ross, 1996; Baranyi et al., 1999).

#### Malting

One kilogram of Safrari sorghum cultivar grains was washed three times using 3 L of distilled water to remove dirt and foreign bodies. The grains were steeped in 3 L of distilled water for 48 h at room temperature ( $\approx$ 25 °C) with 3 changes of water at intervals of 12 h before steep out. Germination was carried out for 4 days in a Heraeus type incubator (D-63450 Hanau, Germany) at a temperature of 25 °C with water sprinkled on the grains on daily basis. The malt was then air dried at 40 °C for 4 days using a CKA 2000 AUF-type dryer Ngaoundere; Cameroon. The malt was rubbed-off of its rootlets and stored in plastics sachets at -18 °C until further use.

#### Mashing

Two hundred and fifty millilitres of distilled water were put into a 600 ml beaker and 50 g of sorghum (malted or unmalted) flour ( $\emptyset$ < 1 mm) added with continuous stirring until a homogenous mixture was obtained. This mixture was incubated at 45 °C for 1 h in a water bath with intermittent stirring at intervals of 5 min. The mix was allowed to decant and 50 ml of the supernatant was withdrawn and kept aside. The temperature of the mash was then raised to boiling so as to gelatinize sorghum starch for 40 min with intermittent stirring at intervals of 5 min of supernatant to the which commercial enzyme/s is/are added according to the Doehlert matrix design of 3 factors, were added to the mash and allowed to incubate (at 65 °C) for 1 h 30 min with intermittent stirring at intervals of 10 min. The mash was filtered at 25 °C for one 1 h 30 min using a Whatmann paper NO 42.

Determination of the content of Reducing Sugars

The reducing sugars content was determined using DNS reagent (Miller, 1959).

#### **Optimization of models**

Models were optimized as previously described (Desobgo et al., 2010). The intersection of the curves, representing the optimal zone, was highlighted.

#### **RESULTS AND DISCUSSION**

#### Modeling and validation of results

Optimisation of the action of mashing enzymes on the reducing sugars yields was carried out by modeling the experimental design required for laboratory purposes. Table 2 shows the results obtained after mashing unmalted and malted *Safrari* using the commercial mashing enzymes Hitempase ( $\alpha$ -amylase), Bioglucanase ( $\beta$ -glucanase) and Brewers Protease (Protease). The mathematical models obtained for reducing sugars after mashing unmalted and malted *Safrari* were as follows respectively:

	Organism of origin	Activity	Description	Optimum temperature	Optimum pH	Recommended application level in adjuncts	Form
Hitempase 2XL	Bacillus licheniformis	4416.29 ± 19.34 U/ml	α-amylase	60 – 95 ℃	4 – 8	60 U/g	Solution
BioglucanaseTX	Trichodermareesei	750 BGU/ml	β-glucanase	60 °C	4.5 – 6.5	0.01 et 0.025 % (v/w)	Solution
Brewers protease	Bacillus amyloliquefaciens	1842.2 ± 1.8 mg FAN/min/ml	Protease	45 – 50 ℃ (denatured at 85 ℃)	6.5 – 7.5	0.4 – 2 g/Kg	Solution

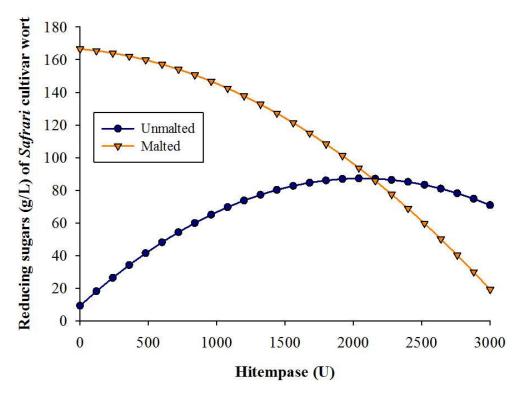
Table 1. Characteristics of commercial enzyme preparations used in mashing.

Hitempase 2XL and Bioglucanase TX were obtained from Kerry Bioscience; Kilnagleary, Carrigaline, Co. Cork, Ireland. Brewers Protease was obtained from DSM Food Specialities, Cedex France.

Table 2. Matrices of Doehlert coded and transformed experimental values.

(	Coded va	alues	Trans	formed expe	rimental values		Reducing s	ugars conte	ent ( <i>Safrari</i> w	ort cultivar)	
Hit	Bio	Brew Prot	Hit (U)	Bio (BGU)	Brew Prot (mg)		Unmalted			Malted	
<b>X</b> ₁	<b>X</b> 2	<b>X</b> 3	<b>X</b> 1	<b>X</b> ₂	X ₃	Exp ^a	Theo ^b	Res ^c	Ехр	Theo	Res
0.000	0.000	0.000	1500	468.75	50	99.00	99.80	-0.80	130.50	131.98	-1.48
1.000	0.000	0.000	3000	468.75	50	94.00	84.36	9.64	110.50	109.91	0.59
0.500	0.866	0.000	2250	937.5	50	110.00	110.47	-0.47	108.73	104.90	3.83
-0.500	-0.866	0.000	750	0.00	50	71.00	63.88	7.12	108.91	113.31	-4.40
0.500	-0.866	0.000	2250	0.00	50	89.54	89.97	-0.43	107.93	105.45	2.48
-0.500	0.866	0.000	750	937.5	50	80.00	78.54	1.46	75.90	78.35	-2.45
0.500	0.289	0.816	2250	615.18	100	95.00	96.45	-1.45	141.50	146.27	-4.77
-0.500	-0.289	-0.816	750	312.32	0.0	79.62	68.95	10.67	161.67	157.74	3.93
0.500	-0.289	-0.816	2250	312.32	0.0	96.71	101.72	-5.01	92.50	95.54	-3.04
0.000	0.577	-0.816	1500	781.07	0.0	114.00	113.95	0.05	109.13	110.48	-1.35
-0.500	0.289	0.816	750	615.18	100	77.00	71.20	5.80	68.50	65.38	3.12
0.000	-0.577	0.816	1500	156.43	100	92.00	94.86	-2.86	109.07	107.42	1.65
0.000	0.000	0.000	1500	468.75	50	96.20	99.80	-3.60	131.00	131.98	-0.98
-1.000	0.000	0.000	0.000	468.75	50	24.00	26.35	-2.35	90.50	91.23	-0.73
-1.000	-0.866	-0.816	0.000	0.0	0.0	05.00	06.21	-1.21	166.98	166.53	0.45
0.000	0.000	0.000	1500	468.75	50	98.70	99.80	-1.10	133.05	131.98	1.07
0.000	0.000	0.000	1500	468.75	50	99.35	99.80	-0.45	134.00	131.98	2.02

with:  $Y_{SafSRed}(X_1, X_2, X_3)$ , representing the mathematical model for unmalted *Safrari*;  $Y_{SafMSRed}(X_1, X_2, X_3)$ , the model for malted *Safrari*;  $X_1$ , Hitempase;  $X_2$ , Bioglucanase and X3



**Figure 1A.** Effect of concentration of Hitempase ( $\alpha$ -amylase) as sole mashing enzyme on reducing sugars content (g/L) of the worts of *Safrari* sorghum cultivar.

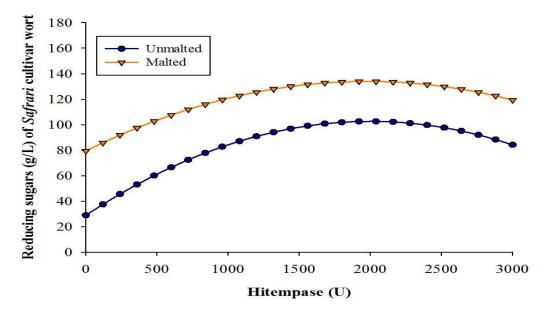
## Brewers Protease.

These mathematical models are polynomials having several variables with correlation coefficients R2 = 0.975 for unmalted Safrari and R2 = 0.990 for malted Safrari. These coefficients, coupled to AAD values of 0.101 and 0.021 (Bas and Boyac, 2007) for unmalted and malted Safrari respectively, allowed for the validation of the models for the wort reducing sugars yields. In addition, bias factors of 1.04 and 1.00 for unmalted and malted Safrari respectively, coupled to accuracy factors of 1.08 and 1.02 for unmalted and malted Safrari respectively, also allowed for validation of the models according to the method described (Ross, 1996; Baranyi et al., 1999). The factors of the models were linear or of first degree (X1, X2 and X3), quadratic or of the second degree (X12, X22 and X32) and of interacting form (X1X2, X1X3, X2X3). They were statistically considered significant or not if the probability (P) of increasing reducing sugars yields was ≤  $0.05 \text{ or} \ge 0.05 \text{ respectively}$  (Table 3).

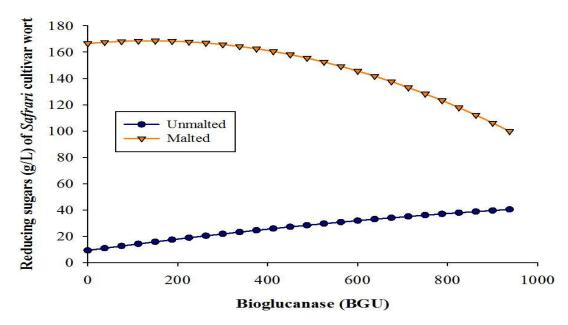
# Effect of Hitempase 2XL on reducing sugars production

The impact of Hitempase as the sole mashing enzyme on the yields of reducing sugars of unmalted and malted Safrari is shown in Figure 1A. Reducing sugars yields

increased with increasing amounts of enzyme for unmalted Safrari mash to attain a maximum level (≈ 90 g/L) at about 2000 U, followed by a slight and steady decrease thereafter. The reducing sugars yield was maximal (≈ 168 g/L) even in absence of Hitempase for malted Safrari mash. Indeed, increasing concentrations of enzyme instead resulted into a steady decrease of reducing sugars content for malted Safrari mash. The maximal reducing sugars content observed for malted Safrari mash in absence of Hitempase was however attributed to the natural virtues that the malting process imparts to mashing cereals in terms of synthesis and development of hydrolytic enzymes. This explains why though known to be the principal mashing enzyme, supplements of Hitempase ( $\alpha$ -amylase) seemed to be of no use in producing reducing sugars as the natural enzyme synthesized during malting seemed to be in sufficient amounts to ensure starch hydrolysis. When the mathematical models were applied to predict the impact of supplements of Bioglucanase and Brewers Protease at concentrations of 400 BGU and 60 mg respectively as accompanying mashing enzymes to Hitempase, it was observed that the profile of reducing sugars increased with increasing amounts of Hitempase for malted Safrari mash though the amounts in the absence of enzyme dropped by roughly two folds (Figure 1B). In both mash types, the milling operation facilitated the dissolution of reducing sugars as could be seen in Figure 1A. This



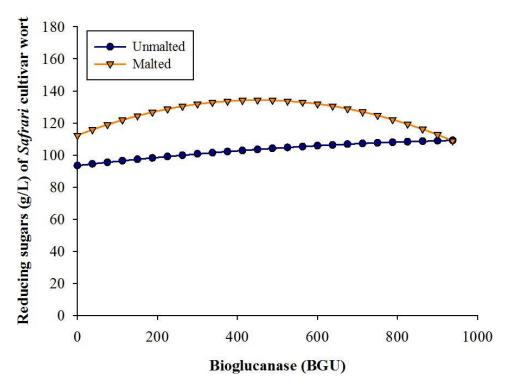
**Figure 1B.** Effect of concentration of Hitempase ( $\alpha$ -amylase) in the presence of fixed concentrations of Bioglucanase (400 BGU) and Brewers protease (60 mg) on reducing sugars content (g/L) of the worts of *Safrari* sorghum cultivar.



**Figure 2A.** Effect of concentration of Bioglucanase as sole mashing enzyme on reducing sugars content (g/L) of the worts of *Safrari* sorghum cultivar.

dissolution accounted for total reducing sugars contents (168 g/L) for malted *Safrari* mash and for 10 g/L for unmalted Safrari mash. Starch is indeed the main macromolecule of cereals and the main substrate of  $\alpha$ -amylase. It is therefore expected that Hitempase contributes to the greatest amounts of reducing sugars in resulting worts due to its action on starch (Goode et al.,

2003; Phiarais et al., 2006; Desobgo et al., 2010). The steady decrease in yields of reducing sugars of malted mash could be attributed to Milliard reactions between soluble nitrogenous compounds and reducing sugars obtained earlier due to the action of hydrolytic enzymes synthesized during malting (Figure 1A) (Hough et al., 1982). From the mathematical models, it was shown that



**Figure 2B**. Effect of concentration of Bioglucanase ( $\beta$ -glucanase) in the presence of fixed concentrations of Hitempase (2000 U) and Brewers protease (60 mg) on reducing sugars content (g/L) of the worts of *Safrari* sorghum cultivar.

in its linear form (X₁), Hitempase's impact on reducing sugars yields was significant (P = 0.000 and 0.002) (Table 3) for both unmalted and malted *Safrari* mash types. This action contributed to 26and 4% for unmalted and malted *Safrari* respectively (Table 3). In its quadratic form (X₁²), Hitempase's action remained statistically significant for both mash types (P = 0.000) (Table 3). Its contribution to increasing reducing sugar yields in this quadratic form (X₁²) (excess of  $\alpha$ -amylase in principle) is indeed 37 and 15% for unmalted and malted *Safrari* respectively (Table 3).

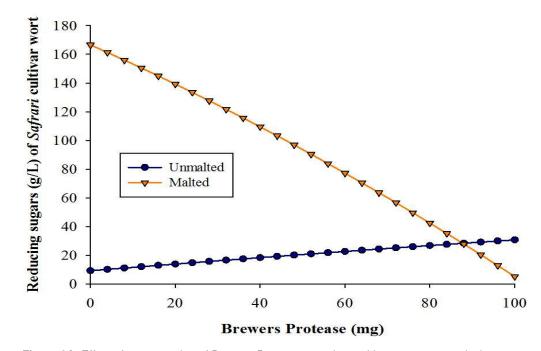
# Effect of Bioglucanase TX on reducing sugars production

Figure 2A shows the effect of mashing unmalted and malted *Safrari* using Bioglucanase as only mashing enzyme on reducing sugars yields. There was a slight but constant increase in reducing sugars yield ( $\approx$  40 g/L) as enzyme concentration increased for unmalted *Safrari* mash. This small yield of reducing sugars due to Bioglucanase's action on unmalted *Safrari* mash can be attributed to its ability to hydrolyse  $\beta$ -glucans into glucose and other soluble carbohydrates. Contrary to Hitempase, Bioglucanase was therefore not a backbone enzyme for reducing sugars yield for malted *Safrari* mash was

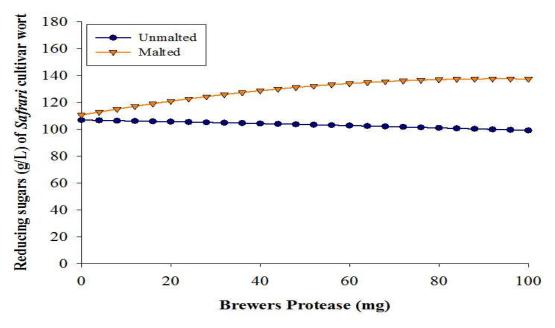
maximal even in the absence of Bioglucanase, and steadily decreased with increase in enzyme concentration. A similar application of the mathematical models as carried out above for Hitempase's action, using 60 mg of Brewers Protease and 2000 U Hitempase as accompanying enzymes, predicted that supplements of these two key mashing enzymes could provide similar results in reducing sugar yields for both unmalted and malted Safrari mashes (Figure 2B). Hitempase once more demonstrated that it was the backbone enzyme contributing to most of the reducing sugars. Figure 2B indeed showed that in the absence of Bioglucanase, but in the presence of Hitempase, yields of reducing sugars became comparable for both unmalted and malted Safrari mashes. In addition, the facilitator role of dissolving reducing sugars by the milling process was considerably accentuated. These observations were all statistically confirmed. Indeed, in its linear form (X₂), Bioglucanase's action was significant (P = 0.018 and 0.001 for unmalted and malted Safrari respectively (Table 3)). Table 3 showed that this enzyme contributed for 9 and 5% of reducing sugars for unmalted and malted Safrari respectively. In its quadratic form  $(X_2^2)$  (excess of enzyme in principle), Bioglucanase contributed for 3and 15% of reducing sugars for unmalted and malted Safrari respectively (Table 3). These contributions were not statistically significant for unmalted Safrari mash, but were for malted safrari mash (P = 0.492 and 0.000)

Effects	Coeffic	eients	Standard	Standard Error		t		oility	Enzyme Cor	ntributions (%)
Ellecis	Unmalted	malted	Unmalted	malted	Unmalted	malted	Unmalted	malted	Unmalted	malted
Constant	99.804	131.976	3.197	1.938	31.217	68.108	0.000	0.000		
X ₁	29.007	9.344	3.289	1.993	8.82	4.688	0.000	0.002	26	4
X2	10.153	-10.252	2.868	1.738	3.066	-5.108	0.018	0.001	9	5
X ₃	-4.522	-9.124	2.71	1.642	-1.362	-4.533	0.216	0.003	4	4
$X_{1}^{2}$	-41.448	-31.406	5.638	3.417	-7.351	-9.19	0.000	0.000	37	15
$X_2^2$	-3.968	-31.5	4.103	2.487	-0.725	-9.499	0.492	0.000	3	15
$X_3^2$	-0.818	-11.548	3.54	2.145	-0.154	-3.584	0.882	0.009	1	5
$X_1 X_2$	3.376	19.873	6.437	3.901	0.454	4.411	0.663	0.003	3	9
$X_2 X_3$	-13.731	10.301	5.665	3.434	-1.713	2.12	0.130	0.072	12	5
$X_1^*X_3$	-5.8	80.639	6.707	4.065	-0.706	16.186	0.503	0.000	5	38

Table 3. Estimation of regression coefficients and enzyme contributions for the reducing sugars content of Safrari.



**Figure 3A**. Effect of concentration of Brewers Protease as sole mashing enzyme on reducing sugars content (g/L) of the worts of *Safrari* sorghum cultivar.



**Figure 3B**. Effect of concentration of Brewers protease in the presence of fixed concentrations of Hitempase (1875 U) and Bioglucanase (750 BGU) on reducing sugars content (g/L) of the worts of *Safrari* sorghum cultivar.

(Table 3). This 1 to 5 ratio difference in reducing sugars content between the unmalted and malted *Safrari* mash types could once more be attributed to the natural virtues of the malting process.

# Effect of Brewers Protease on reducing sugars production

The effect of mashing unmalted and malted Safrari on vields of reducing sugars using sole mashing enzyme, Brewers Protease, is shown in Figure 3A. A small but constant increase in reducing sugars content with increasing enzyme concentrations was observed (≈ 30 g/L) for unmalted Safrari mash. The milling process once more contributed to about 10 g/L of these reducing sugars amounts. As observed for Bioglucanase, the maximal reducing sugars amounts were obtained when mashing with malted Safrari in the absence of supplements of Brewers Protease. This could once more be attributed to the virtues known to the malting process as earlier explained above. A high rate of decrease of reducing sugars content with increasing concentrations of Brewers Protease for the malted Safrari mash was observed. This could be explained by the fact that this enzyme released into the medium soluble nitrogenous materials (amino acids and others) which reacted with the reducing sugars (produced during the malting process), displaying as such the rapid decrease in the amounts of the sugars as observed in Figure 3A (Hough et al., 1982). The mathematical models were once more used to

predict the yield in reducing sugars as carried out above for Hitempase and Bioglucanase actions. Thus, using 2000 U of Hitempase and 400 BGU of Bioglucanase as accompanying mashing enzymes with increasing amounts of Brewers Protease, similar results in reducing sugars yields for both unmalted and malted Safrari mashes were once more observed (Figure 3B). These observations were statistically confirmed. In its first degree form (X₃), the impact of Brewers Protease was not significant for unmalted Safrari mash but significant for malted Safrari mash (P = 0.216 and 0.003 respectively) (Table 3). Its contribution to reducing sugars was barely 4% for both mash types (Table 3). The impact of the enzyme in its quadratic form  $(X_3^2)$ , was not significant for unmalted Safrari mash but significant for malted Safrari mash (P = 0.882 and 0.009 respectively) (Table 3). Its contribution to reducing sugars was 1 and 5% respectively (Table 3).

# Effect of enzyme interactions on the production of reducing sugars

The models were further exploited to predict the impacts of the interactions  $(X_1X_2, X_1X_3 \text{ and } X_2X_3)$  of these enzymes on yields of reducing sugars. The results are shown in Table 3. Globally, they were statistically not significant for unmalted *Safrari* mashes (P = 0.339), but were for malted *Safrari* mashes (P = 0.000) (Table 4). The interaction  $X_1X_2$  (Hitempase/Bioglucanase) had no significant impact on unmalted *Safrari* mash, but had for

Courses	dl	Sum s	Sum square		Mean square			Probability	
Source	ai	Unmalted	Malted	Unmalted	Malted	Unmalted	Malted	Unmalted	Malted
Regression	9	12650.765	11623.844	1405.641	1291.538	30.804	77.047	0.000	0.000
Linear	3	8527.165	2482.605	2842.388	827.535	62.29	49.367	0.000	0.000
Quadratic	3	3941.669	583.306	1313.89	194.435	28.793	11.599	0.000	0.004
Interactions	3	181.932	8557.933	60.644	2852.644	1.329	170.175	0.339	0.000
Residual error	7	319.423	117.341	45.632	16.763				
Total error	16	12970.188	11741.185						

Table 4. ANOVA for the reducing sugars content of Safrari.

Table 5. ANOVA for comparing reducing sugars content of unmalted and malted Safrari worts.

Source	dl	Sum square	Mean square	F	Р
Inter-groups	1	9200.22	9200.22	11.91	0.0016
Intra-groups	32	24711.4	772.23		
Total	33	33911.6			

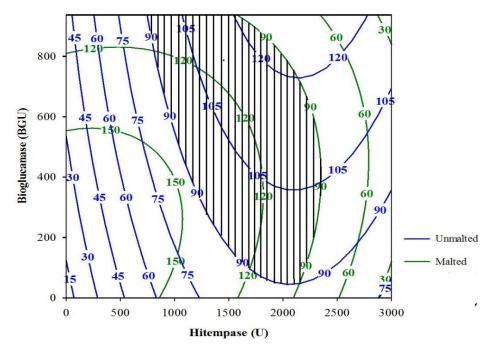
malted Safrari (P = 0.663 and 0.003 respectively (Table 3). It contributed for merely 3% of reducing sugars for unmalted Safrari and 9% for malted Safrari mash (Table 3). The interaction  $X_1X_3$ , corresponding to the couple Hitempase/Brewers Protease, also had no significant impact on yields of reducing sugars of unmalted Safrari, but had a significant impact on those of malted Safrari mashes (P = 0.503 and 0.000 respectively) (Table 3). Its contribution to reducing sugar yields was merely 5% for unmalted Safrari, but as much as 38% for malted Safrari (Table 3). This remarkable difference was once more to be attributed to the virtues of the malting process and not simply to supplements of the Hitempase/Brewers Protease couple as such. Efficient starch hydrolysis by αamylase indeed occurs only after the breakdown of cereal grain cell walls by β-glucanase, followed by liberation of starch granules due to proteolysis of the protein matrix enrobing them. This should be the natural sequence of events which accounted for the results obtained for the malted sample type as compared to the unmalted Safrari sample type. The interaction Bioglucanase/Brewers Protease (X₂X₃) had no significant impact on reducing sugars yields for both the unmalted Safrari and malted Safrari mashes (P = 0.130 and 0.072 respectively) (Table 3). Its contribution to reducing sugars yields was 12 and 5% respectively for both mash types (Table 3). The Bioglucanase/Brewers Protease combination plays a supporting role in starch hydrolysis during mashing (Desobgo et al., 2010). Table 4 statistically confirmed the observation that Safrari sorghum malted type samples were more potential mashing materials than unmalted Safrari sorghum adjuncts to which commercial enzymes are supplemented for the production of worts of higher reducing sugars yields (P =

0.001).

# Optimization of the concerted mashing enzymes' action on the production of reducing sugars

The results obtained for the action of the enzymes on reducing sugars yields after mashing on the basis of the models, were optimized to define a satisfactory domain of compromise for the action of the mashing enzymes. This domain was obtained for a reducing sugars content  $\geq$  90 g/L. The theoretical optimal combination of enzyme action for unmalted *Safrari* gave the following triplet of coded variables for reducing sugars content: 0.442, 0.866 and – 0.816 (2163 U, 937.5 BGU and 0 mg real variables) for Hitempase, Bioglucanase and Brewers Protease respectively.

This triplet allowed for a maximal reducing sugars content of 126.57 g/L. The triplet for malted Safrari was -1, - 0.612 and - 0,816 (0 U, 137.48 BGU and 0 mg real variables). It allowed for a maximal reducing sugars content of 168.56 g/L. The optimal enzyme combinations were thus different and gave different results for the two mash types. These results showed that maximal reducing sugars contents could be obtained using Hitempase and Bioglucanase but not Brewers Protease supplements when mashing unmalted Safrari. whereas only Bioglucanase, and in limited amounts, could be used for mashing malted Safrari. The significant difference (P = 0.001) between reducing sugars content of unmalted and malted Safrari worts is shown in Table 5. Minimal reducing sugars content (90 g/L) could be obtained under these conditions permitting the highlight of the optimal domain (Figure 4).



**Figure 4.** Response surface curves for the enzyme combinations providing for optimal reducing sugars content (g/L) for unmalted and malted sorghum worts of the cultivar *Safrari*.

#### Conclusion

This work clearly showed that though Hitempase 2XL was the most important enzyme component for producing significant amounts of reducing sugars during mashing of unmalted *Safrari* grist, it appeared to be indispensable for mashing malted *Safrari* grist. Bioglucanase TX and Brewers Protease were merely supporting enzymes to Hitampase 2XL only for mashing unmalted *Safrari* grist, as they too showed little or no participatory role in increasing reducing sugars amounts when used as supplements when mashing with malted *Safrari* grist - in general, the malting process seems to be auto-sufficient in accounting for the required natural mashing enzymes in terms of both amounts and quality, for producing the amounts of reducing sugars obtainable when mashing malted *Safrari* grist.

#### ACKNOWLEDGEMENTS

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Full Length Research Paper

# Use of the response surface methodology for optimizing the action of mashing enzymes on wort reducing sugars of the *Madjeru* sorghum cultivar

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A three factor Doehlert design was used to develop a statistical model to optimize the action of three commercial mashing enzymes (Hitempase 2XL, Bioglucanase TX and Brewers protease) on reducing sugars content of the worts of unmalted and malted *Madjeru* sorghum. The response surface methodology revealed that increasing amounts of Hitempase considerably increased reducing sugars content during mashing of unmalted and malted *Madjeru* sorghum grist to about 105.39 g/L and 132.25 g/L respectively. The milling process contributed to about 22 g/L and 54 g/L for the unmalted and malted mash types respectively. Increasing amounts of Bioglucanase was virtually insignificant, while for Brewers protease, reducing sugar yields rather decreased to nil for both the unmalted and malted mash types. Optimization of the concerted actions of the three enzymes for reducing sugars content of unmalted *Madjeru* sorghum mash gave a combination of 1995 U, 89.31 BGU and 28.86 mg for Hitempase, Bioglucanase and Brewers Protease respectively. This gave a maximal reducing sugars content of 108.78 g/L. This combination was 3000 U, 0 BGU and 49.69 mg for malted *Madjeru* sorghum mash, giving a maximal reducing sugars yield of 153.15 g/L.

Key words: Response surface methodology, optimization, mashing enzymes, Madjeru, reducing sugars.

## INTRODUCTION

Sorghum (*Sorghum bicolor* (L.) Moench) is an important tropical cereal-bearing monocotyledonous plant found in the semi-arid areas of the world. It is a vital calorie-based food component in humane nutrition in some parts of Africa (Taylor, 2004).

Besides this fundamental function, sorghum is used in the production of traditional opaque beer and nonalcoholic drinks in developing countries and, until recently, in the production of industrial beer (Palmer, 1989; Taylor and Dewar, 2001). The low synthesis and development of principal hydrolytic enzymes during malting of this cereal is however a limiting factor for easy mashing of its malts as compared to barley malt (EtokAkpan and Palmer, 1990; EtokAkpan, 1992). This was ascribed to the malting procedures and varietal types of sorghum used. Work on some popular sorghum cultivars of Northern Cameroon used in brewing the traditional beer *Bili-Bili* confirmed that the profile of hydrolytic enzymes and the levels of fermentable sugars during mashing were indeed cultivar-dependent. The *Madjeru* sorghum cultivar was shown to be richer in starch than the *Safrari* and *S.35* cultivars, but had poorer hydrolytic enzymes and fermentable sugar potentials (Nso et al., 2003; 2006).

The use of commercial enzymes when mashing with sorghum in order to obtain better wort specifications for

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**Table 1.** Characteristics of commercial enzyme preparations used in mashing.

	Organism of origin	Activity	Description	Optimum temperature	Optimum pH	Recommended application level in adjuncts	Form
Hitempase 2XL BioglucanaseTX	Bacillus licheniformis Trichodermareesei	4416.29 ± 19.34 U/ml 750 BG U/ml	α-amylase β-glucanase	60 − 95 ℃ 60 ℃	4 – 8 4.5 – 6.5	60 U/g 0.01 et 0.025 % (v/w)	Solution Solution
Brewers Protease	Bacillus amyloliquefaciens	1842.2 ± 1.8 mg FAN/min/mL	Protease	45 – 50 ℃ (denatured at 85 ℃)	6.5 – 7.5	0.4 – 2 g/Kg	Solution

Hitempase 2XL and Bioglucanase TX were obtained from Kerry Bioscience; Kilnagleary, Carrigaline, Co. Cork, Ireland. Brewers Protease was obtained from DSM Food Specialities, Cedex France.

beer brewing has however become common practice (Bajomo and Young, 1992, Agu and Palmer, 1998; Goode et al., 2002; 2003; Goode and Arendt, 2003,).

It is not however clear how far mashing enzyme supplements in *Madjeru* sorghum cultivar mashes could help alleviate the levels of fermentable sugars in its worts. In this work, the action of three principal commercial mashing enzymes (Hitempase 2XL, Bioglucanase TX and Brewers Protease) on the reducing sugars content of worts from the *Madjeru* sorghum cultivar (unmalted and malted) was modeled and optimized using the response surface methodology (RSM).

#### MATERIALS AND METHODS

#### Enzymes

The characteristics of the commercial enzymes used (Hitempase 2XL, a thermo stable  $\alpha$ -amylase from *Baccillus licheniformis*, Brewers Protease from *Baccillus amyloliquefaciens* and Bioglucanase TX, from an enzymatic composition of  $\beta$ -glucanase and hemicellulases from *Trichoderma reesei*) and sources are presented in Table 1.

#### Sorghum cultivar

The *Madjeru* sorghum cultivar was obtained from the Institute of Research and Agronomic Development (IRAD) Maroua, Cameroon.

#### Modeling

Modeling was carried out as previously described (Desobgo et al., 2010). A Doehlert matrix design of 3 factors representing Hitempase 2XL (X₁), Bioglucanase TX (X₂) and Brewers Protease (X₃) at ranges of [0-3000 U], [0-937.5 BGU] and [0-100 mg] respectively, was used. The transformed matrix of coded variables to an experimental matrix and the desired response (reducing sugars) are shown in Table 2. Mathematical models describing the relationships among the process dependent variable and the independent variables in a second-order equation were developed (Giovanni, 1983).

Design-based experimental data were matched according to the following second-order polynomial Equation 1.

$$y = \beta_0 + \sum_{j=1}^k \beta_j x_j + \sum_{j=1}^k \beta_{jj} x_j^2 + \sum_{i < j} \beta_{ij} x_i x_j + \varepsilon$$
(1)

Where, I and j, are linear and quadratic coefficients respectively, while ' $\beta$ ', the regression coefficient, k the number of factors studied and optimized in the experiment

and ' $\epsilon$ ' is random error. The coefficients of the models and the models were obtained using the Systat version 12 software (Systat Software, Inc., San Jose, USA). This software also gives a statistical analysis on the model. Lastly, the curves were plotted using Sigmaplot version11 build 11.0.0.77 software (WPCubed, GmbH, Germany).

#### Validation of models

The quality of fit of the second order equations was expressed by the coefficient of determination  $R^2$ . The models were validated using two differents methods. The first method was the absolute average deviation (AAD) method (Bas and Boyac, 2007), while the second method consisted in applying the bias factor and accuracy factor (Ross, 1996; Baranyi et al., 1999).

#### Malting

One kilogram of *Madjeru* sorghum cultivar grains was washed three times using 3 L of distilled water to remove dirt and foreign bodies. The grains were steeped in 3 L of distilled water for 48 h at room temperature ( $\approx 25 \,^{\circ}$ C) with 3 changes of water at intervals of 12 h before steep out. Germination was carried out for 4 days in a Heraeus type oven (D-63450 Hanau, Germany) at a temperature of 25  $^{\circ}$ C with water sprinkled on the grains on daily basis. The malt was then air dried at 40  $^{\circ}$ C for 4 days using a CKA 2000

C	oded valu	Jes	Transform	ed experime	ental values	Reduc	ing sugar	s content	(g/L)( <i>Madje</i>	<i>ru</i> wort cu	ltivar)
Hit	Bio	Brew Prot	Hit (U)	Bio (BGU)	Brew Prot (mg)		Unmalted			Malted	
<b>X</b> ₁	<b>X</b> ₂	<b>X</b> 3	<b>X</b> ₁	<b>X</b> ₂	<b>X</b> 3	Exp ^a	Theo [⊳]	Res ^c	Ехр	Theo	Res
0.000	0.000	0.000	1500	468.75	50	100.81	102.57	-1.76	124.00	123.08	0.92
1.000	0.000	0.000	3000	468.75	50	85.55	78.61	6.94	137.59	132.66	4.93
0.500	0.866	0.000	2250	937.5	50	87.56	87.51	0.05	106.21	109.81	-3.60
-0.500	-0.866	0.000	750	0.00	50	80.34	73.74	6.60	76.30	77.63	-1.33
0.500	-0.866	0.000	2250	0.00	50	104.64	105.30	-0.66	140.30	142.90	-2.60
-0.500	0.866	0.000	750	937.5	50	76.98	76.93	0.05	99.27	96.23	3.04
0.500	0.289	0.816	2250	615.18	100	79.65	82.49	-2.84	114.47	118.89	-4.42
-0.500	-0.289	-0.816	750	312.32	0.0	88.61	75.79	12.82	93.99	96.95	-2.96
0.500	-0.289	-0.816	2250	312.32	0.0	94.01	100.67	-6.66	115.48	117.55	-2.07
0.000	0.577	-0.816	1500	781.07	0.0	89.28	90.00	-0.72	88.89	87.90	0.99
-0.500	0.289	0.816	750	615.18	100	70.97	65.23	5.74	63.37	60.65	2.72
0.000	-0.577	0.816	1500	156.43	100	80.02	82.93	-2.91	79.34	77.66	1.68
0.000	0.000	0.000	1500	468.75	50	97.32	102.57	-5.25	126.00	123.08	2.92
-1.000	0.000	0.000	0.000	468.75	50	31.20	36.47	-5.27	47.65	53.82	-6.17
-1.000	-0.866	-0.816	0.000	0.0	0.0	17.40	22.85	-5.45	58.95	54.94	4.01
0.000	0.000	0.000	1500	468.75	50	100.03	102.57	-2.54	125.00	123.08	1.92
0.000	0.000	0.000	1500	468.75	50	104.39	102.57	1.82	123.00	123.08	-0.08

**Table 2.** Matrices of Doehlert coded and transformed experimental values.

^aExperimental result values. ^bTheretical values (values coming from mathematical models). ^c Residue. Hit: Hitempase 2XL, Bio: Bioglucanase TX, Brew Pro: Brewers Protease.

AUF-type dryer Ngaoundere, Cameroon. The malt was rubbed-off of its rootlets and stored until further use.

#### Mashing

Two hundred and fifty (250) ml of distilled water were put into a 600 ml beaker and 50 g of sorghum (malted or unmalted) flour ( $\emptyset < 1$  mm) added with continuous stirring until a homogenous mixture was obtained. This mixture was incubated at 45 °C for 1 h in a water bath with intermittent stirring at intervals of 5 min. The mix was allowed to decant and 50 ml of the supernatant withdrawn and kept aside. The temperature of the mash was then raised to boiling so as to gelatinize sorghum starch during 40 min with intermittent stirring at intervals of 5 °C. The 50 ml of supernatant to the which commercial enzyme/s is/are added according to the Doehlert matrix design of 3 factors, were added to the mash and allowed to incubate for 1 h 30 min with intermittent stirring at intervals of 10 min. The mash was filtered at 25 °C during 1 h 30 min using Whatmann paper N° 42.

#### Determination of reducing sugars contents

The reducing sugars content was determined using DNS reagent (Miller, 1959).

#### Optimization of models

Models were optimized as previously described (Desobgo et al.,

2010). The intersection of the curves, representing the optimal zone, was highlighted.

## **RESULTS AND DISCUSSION**

Optimization of the action of mashing enzymes on reducing sugars was carried out by modeling the experimental design required for manipulation in laboratory. Table 2 shows the results obtained for reducing sugars after mashing unmalted and malted *Madjeru* using Hitempase 2XL ( $\alpha$ -amylase), Bioglucanase TX ( $\beta$ -glucanase) and Brewers Protease (Protease).

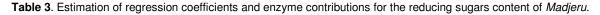
#### Modeling and validation of results

The mathematical models (Equations 2 and 3) obtained for reducing sugars after mashing unmalted and malted *Madjeru* were as follows respectively:

 $Y_{MadSRed}(X_1, X_2, X_3) = 102.573 + 21.070X_1 - 4.217X_2 - 7.312X_3 - 12.118X_1X_2 - 0.378X_1X_3 + 5.688X_2X_3 - 45.031X_1^2 - 7.261X_2^2 - 16.534X_3^2$ (2)

$$\begin{split} Y_{\text{MadMSRed}}(X_1, X_2, X_3) &= 123.077 + 39.421X_1 - 4.181X_2 - 9.230X_3 - 29.849X_1X_2 + \\ 33.635X_1X_3 + 28.583X_2X_3 - 29.835X_1^2 - 11.971X_2^2 - 34.318X_3^2 \end{split} \tag{3}$$

<b>-</b> <i>H</i>	Coefficient	ts	Standard erre	or	t		Probability		Enzyme conti	ributions(%)
Effects	Unmalted	malted	Unmalted	malted	Unmalted	malted	Unmalted	malted	Unmalted	malted
Constant	102.573	123.077	3.79	2.304	27.066	53.413	0.000	0.000		
<b>X</b> 1	21.07	39.421	3.898	2.37	5.405	16.631	0.001	0.000	18	18
X2	-4.217	-4.181	3.399	2.067	-1.074	-1.752	0.318	0.123	3	2
X3	-7.312	-9.23	3.212	1.953	-1.858	-3.856	0.106	0.006	6	4
X ₁ ²	-45.031	-29.835	6.683	4.064	-6.738	-7.342	0.000	0.000	38	13
X ₂ ²	-7.261	-11.971	4.864	2.958	-1.12	-3.036	0.300	0.019	6	5
X ₃ ²	-16.534	-34.318	4.196	2.551	-2.624	-8.957	0.034	0.000	14	16
$X_1 X_2$	-12.118	-29.849	7.63	4.639	-1.375	-5.572	0.211	0.001	10	14
$X_2 X_3$	5.688	28.583	6.715	4.083	0.599	4.947	0.568	0.002	0	15
X1*X3	-0.378	33.635	7.951	4.834	-0.039	5.677	0.970	0.001	5	13



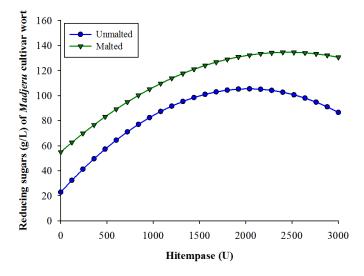


Figure 1A. Effect of concentration of Hitempase ( $\alpha$ -amylase) as sole mashing enzyme on reducing sugars content (g/L) of sorghum wort cultivar *Madjeru*.

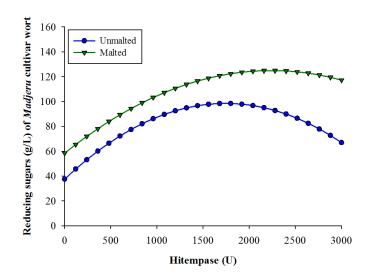
With  $:Y_{MadSRed}(X_1, X_2, X_3)$  representing the mathematical model for unmalted *Madjeru*;  $Y_{MadMSRed}(X_1, X_2, X_3)$ , the model for malted *Madjeru*:  $X_{1,}$  Hitempase;  $X_{2,}$  Bioglucanase and  $X_{3,}$  Brewers Protease.

These mathematical models are polynomials having several variables with correlation coefficients  $R^2 = 0.951$  for unmalted *Madjeru* and  $R^2 = 0.987$  for malted *Madjeru*. These coefficients, coupled to AAD values of 0.067 and 0.032 obtained by the method (Bas and Boyac, 2007) for unmalted and malted *Madjeru* respectively, allowed for the validation of the models for the wort reducing sugars. In addition, a bias factor (1.01 and 1 for unmalted and malted *Madjeru*), coupled to accuracy factors of 1.07 and 1.03 for both unmalted and malted *Madjeru* respectively,

also allowed for validation of the models according to the method described (Ross, 1996; Baranyi et al., 1999). The factors of the models were linear or of first degree ( $X_1$ ,  $X_2$  and  $X_3$ ), quadratic or of the second degree ( $X_1^2$ ,  $X_2^2$  and  $X_3^2$ ) and of interaction form ( $X_1X_2, X_1X_3, X_2X_3$ ). They were statistically considered significant or not if the probability (P) of increasing reducing sugars content was  $\leq 0.05$  or  $\geq 0.05$  respectively (Table 3).

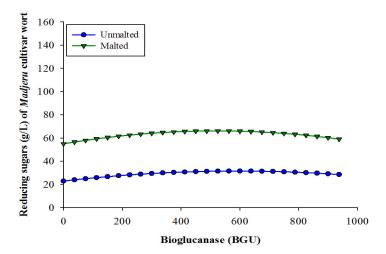
# Effect of Hitempase 2XL on reducing sugars production

The impact of Hitempase 2XL ( $\alpha$ -amylase) as sole mashing enzyme on the reducing sugars content of unmalted and malted Madjeru is shown in Figure1A. Reducing sugars content increased with increasing concentration of enzyme for unmalted and malted Madieru mash types to attain maxima of 105.39 g/L and 132.25 g/L respectively, at an enzyme concentrations of 2031 U and 2451 U respectively. This was followed by a slight and steady decrease thereafter for the unmalted Madjeru mash. The measurable reducing sugar contents could not however be ascribed entirely to the action of Hitempase supplements, as in the absence of the enzyme, Figure 1A clearly showed that the contents for the unmalted and malted Madjeru mash types were about 22 g/L and 54 g/L respectively. These amounts of reducing sugars could be attributed to the milling process in the case of the unmalted Madjeru mash type, and to milling and malting processes for the malted Madjeru mash type. When the mathematical models were applied to predict the impact of a combination of the three enzymes (Bioglucanase and Brewers protease added at fixed amounts of 750 BGU and 60 mg, respectively) on the reducing sugar contents, the same trends as described above in Figure 1A were observed with increasing amounts of Hitempase for both Madjeru mash types



**Figure 1B.** Effect of concentration of Hitempase ( $\alpha$ -amylase) in the presence of fixed concentrations of Bioglucanase (750 BGU) and Brewers protease (60 mg) on reducing sugars content (g/L) of sorghum wort cultivar *Madjeru*.

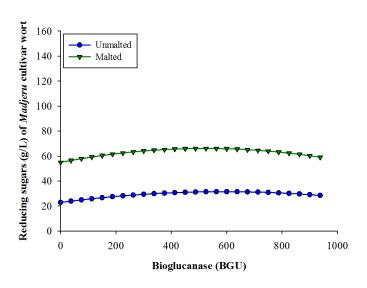
(Figure 1B). In fact, starch is the main macromolecule of sorghum and the main substrate of  $\alpha$ -amylase. It could therefore be expected that Hitempase contributes for the greatest amount of soluble materials in the form of reducing sugars that could be found in resulting worts due to its action on starch (Goode et al., 2003, Phiarais et al., 2006, Desobgo et al., 2010). The reducing sugars content for the unmalted Madjeru mash type slightly reduced beyond the maxima concentrations of the enzyme (Figures 1A and 1B). This could be explained by the fact that traces of amino-acids and other soluble nitrogenous materials present in the wort reacted with some of the reducing sugars (Hough et al., 1982), thus decreasing the measurable content. These results also demonstrated that the presence of the key mashing enzyme (a-amylase) developed during malting boosted the mashing efficiency of this cereal (Figure 1A). From the mathematical model, it was shown that in its linear form  $(X_1)$ , the action of Hitempase contributed for 18% of reducing sugars for both unmalted Madjeru and malted Madjeru (Table 3). Statistical analyses also showed that this contribution was significant (P = 0.001 and 0.000 for the two mash types, respectively) (Table 3). In its quadratic form  $(X_1^2)$ , the action of Hitempase remained statistically significant (P = 0.000) for both unmalted and malted Madjeru. This confirmed the above biological observation according to which supplements of this enzyme in both the unmalted and malted mashes of *Madjeru* are important. Its contribution to reducing sugars in its quadratic form  $(X_1^2)$  (excess of  $\alpha$ -amylase in principle) is indeed 38 and 13% for unmalted and malted Madjeru respectively (Table 3).



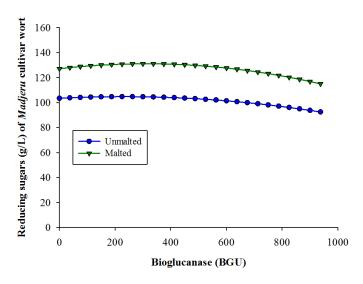
**Figure 2A.** Effect of concentration of Bioglucanase as sole mashing enzyme on reducing sugars content (g/L) of sorghum wort cultivar *Madjeru*.

# Effect of Bioglucanase TX on reducing sugars production

Figure 2A shows the effect of mashing unmalted and malted Madjeru using Bioglucanase TX as sole mashing enzyme on reducing sugars content. There was no significant effect of Bioglucanase on the reducing sugars content of both unmalted and malted Madjeru mashes (Table 3). Reducing sugars were present at 22.85 and 54.94 g/L for the two mash types respectively in the absence of this mashing enzyme. Almost all of the measurable soluble reducing sugars therefore seemed to be due to the milling operation. In the presence of Bioglucanase, reducing sugars contents were maximal at 22.86 and 54.95 g/L at concentrations of 590.24 and 521.74 BGU for unmalted and malted Madjeru respectively with increasing enzyme concentration. The small reducing sugars yields due to Bioglucanase's action can be attributed to its hydrolysis of β-glucans into glucose and other soluble carbohydrates. Bioglucanase was therefore not a backbone enzyme for production of reducing sugars during mashing (Phiarais et al., 2006, Desobgo et al., 2010). A similar application of the carried out above mathematical models as for Hitempase's action, using 60 mg of Brewers protease and 1875 U of Hitempase, predicted that supplementing these two key mashing enzymes could provide higher reducing sugars content for both unmalted and malted Madjeru mashes (Figure 2B). Hitempase once more demonstrated that it was the backbone enzyme that contributed most in reducing sugars' yields. Figure 2B showed that even in the absence indeed of Bioglucanase, but in the presence of Hitempase, the reducing sugars content increased 5 folds for unmalted



**Figure 2A.** Effect of concentration of Bioglucanase as sole mashing enzyme on reducing sugars content (g/L) of sorghum wort cultivar *Madjeru*.



**Figure 2B.** Effect of concentration of Bioglucanase ( $\beta$ -glucanase) in the presence of fixed concentrations of Hitempase (1875 U) and Brewers protease (60 mg) on reducing sugars content (g/L) of sorghum wort cultivar *Madjeru*.

*Madjeru* mashes and 3 folds for malted *Madjeru* mashes. These observations were all statistically confirmed. Indeed, in its linear form (X₂), Bioglucanase's action was not significant (P = 0.318 and 0.123 for unmalted and malted *Madjeru* respectively) (Table 3). Table 3 showed that this enzyme contributed to merely 3 and 2% of reducing sugars content for unmalted and malted *Madjeru* mashes respectively. In its quadratic form  $(X_2^2)$  (excess of enzyme in principle), Bioglucanase contributed to 6 and 5 % of reducing sugars for unmalted and malted *Madjeru* mashes respectively (Table 3). This contribution was not statistically significant for unmalted *Madjeru* mash, but significant for malted *Madjeru* mash (P = 0.300 and 0.019) (Table 3).

Once more, the highest values of reducing sugars obtained for malted *Madjeru* mash, as compared to unmalted *Madjeru* mash, could be attributed to the impact of hydrolytic enzymes developed during malting.

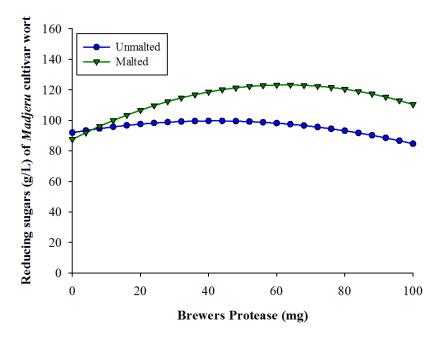
# Effect of Brewers protease on reducing sugars production

The effect of mashing unmalted and malted *Madjeru* on reducing sugars content using as sole mashing enzyme, Brewers protease, is shown in Figure 3A. There was no contribution of reducing sugars for both unmalted and malted *Madjeru* by this enzyme.

All the measurable soluble reducing sugars seemed to be due to the milling operation for the unmalted mash type, and the malting and milling operations, for the malted mash type. The decrease of yields of reducing sugars with increase of Brewers protease for unmalted and malted Madjeru mashes could be attributed to reactions between nitrogenous functions and some of the reducing sugars (Hough et al., 1982). The mathematical model was once more used to predict the reducing sugars content in the presence of fixed amounts of the two other key mashing enzymes. Thus, using 1875 U of Hitempase and 750 BGU of Bioglucanase with increasing amounts of Brewers protease, the results showed that higher amounts of reducing sugars could be obtained for both unmalted and malted Madjeru mashes (Figure 3B). These observations were statistically confirmed using the mathematical model. In its first degree form  $(X_3)$ , the impact of Brewers protease was not significant for unmalted Madjeru mash but significant for malted Madjeru mash (P= 0.106 and 0.006 respectively) (Table 3). Its contribution to reducing sugars content was 6 and 4% respectively for both mashes (Table 3). The impact of the enzyme in its quadratic form  $(X_3^2)$ , was significant for both mashes (P= 0.034 and 0.000 respectively). Its contribution to reducing sugars content was 14 and 16% respectively (Table 3).

## Effect of enzymes interactions on reducing sugars production

The models were further exploited to predict the impacts of the interactions  $(X_1X_2, X_1X_3 \text{ and } X_2X_3)$  of these enzymes on reducing sugars content. The results are shown in Table 3. They were statistically not significant



**Figure 3B.** Effect of concentration of Brewers protease in the presence of fixed concentrations of Hitempase (1875 U) and Bioglucanase (750 BGU) on reducing sugars content (g/L) of sorghum wort cultivar *Madjeru*.

Table 4. ANOVA for the reducing sugars content of Madjeru.

Courses		Sum s	quare	Mean	square	F		Р	
Source	DF	Unmalted	Malted	Unmalted	Malted	Unmalted	Malted	Unmalted	Malted
Regression	9	8675.21	12655.782	963.912	1406.198	15.034	59.323	0.001	0.000
Linear	3	4174.58	8636.535	1391.53	2878.845	21.704	121.45	0.001	0.000
Quadratic	3	4337.98	2342.886	1446	780.962	22.553	32.946	0.001	0.000
Interactions	3	162.641	1676.361	54.214	558.787	0.846	23.574	0.511	0.000
Residual error	7	448.802	165.928	64.115	23.704				
Total error	16	9124.01	12812.71						

DF: Degree of freedom

for unmalted *Madjeru* mashes (P = 0.511), but were for malted *Madjeru* mashes (P=0.000) (Table 4). The interaction  $X_1X_2$  (Hitempase/Bioglucanase) had no significant impact on unmalted *Madjeru* mash but had for malted *Madjeru* (P = 0.211 and 0.001) respectively (Table 3). It contributed to 10% of reducing sugars content for unmalted *Madjeru* mash and to 14% for malted *Madjeru* mash (Table 3). Though known to be the backbone starch hydrolyzing enzyme, the action of Hitempase is best exploited when the cell walls of cereal grains are broken down by  $\beta$ -glucanases, hemicellulases and cellulases, to liberate starch granules. This sequence of events during malting was confirmed by the mathematical models above.

The interaction  $X_1X_3$ , corresponding to the couple

Hitempase/Brewers protease, also had no significant impact on the reducing sugars content of unmalted *Madjeru* mash (P = 0.970), but had a significant impact on the reducing sugars content of malted *Madjeru* mash (P = 0.001) (Table 3). Its contribution to reducing sugars content was 0 and 15% respectively (Table 3). This result was once more in conformity with the biological sequence occurring during malting. Efficient starch hydrolysis by  $\alpha$ amylase indeed occurs only after the breakdown of cell walls by  $\beta$ -glucanase, followed by liberation of starch granules due to proteolysis of the protein matrix enrobing them. The interaction Bioglucanase/Brewers protease (X₂X₃) had no significant impact on reducing sugars content for unmalted *Madjeru* mashes, but had for malted *Madjeru* mashes (P = 0.568 and 0.002 respectively)

Table 5. ANOVA for comparing reducing sugars content of unmalted and malted Madjeru worts.

Source	DF	Sum square	Mean square	F	Р
Inter-groups	1	3224.25	3224.25	4.7	0.037
Intra-groups	32	21945.7	685.804		
Total	33	25170			

DF : Degree of freedom.

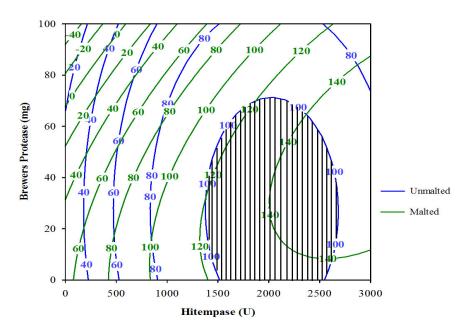


Figure 4. Response surface curves for the enzyme combinations providing for optimal reducing sugars content (g/L) for unmalted and malted sorghum wort cultivar *Madjeru*.

(Table 3). Its contribution to reducing sugars content was 5 and 13% respectively for both mash types (Table 3). These low contributions by the couple (Bioglucanase/ Brewers protease) were expected, as the two enzymes only play a supporting role in starch hydrolysis during mashing (Desobgo et al., 2010).

# Optimization of the concerted mashing enzymes action on the production of reducing sugars

The results obtained for the action of the enzymes on reducing sugars yields after mashing on the basis of the models, were optimized to define a satisfactory domain of compromise for the action of the mashing enzymes. This domain was obtained for a reducing sugars content  $\geq$  100 g/L. The theoretical optimal combination of enzyme action for unmalted *Madjeru* gave the following triplet of coded variables for reducing sugars content: 0.330, – 0.701 and – 0.345 (1995 U, 89.31 BGU and 28.86 mg real variables) for Hitempase, Bioglucanase and Brewers protease respectively. This triplet allowed for a maximal

reducing sugars content of 108.78 g/L. The triplet for malted Madjeru was 1, - 0.866 and - 0.005 (3000 U, 0 BGU and 49.69 mg real variables). It allowed for a maximal extract of 153.15 g/L. The optimal enzyme combinations were thus different and gave different results. These results once more confirmed the fact that commercial enzymes supplements for mashing of unmalted and malted Madjeru were necessary to obtain maximum reducing sugars contents. The significant difference (P = 0.037) between reducing sugars content of unmalted and malted Madjeru worts is shown in Table 5. By fixing Bioglucanase concentration at 0 BGU (according to the optimization), the minimal reducing sugars content (100 g/L) could be obtained and the necessary enzyme combination could therefore permit highlight the optimal domain (Figure 4).

### Conclusion

Optimizing the amounts of the enzyme needed for alleviating the levels of reducing sugars during mashing

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demonstrated that Hitempase 2XL was the most important enzyme component in terms of both amounts and efficiency for the unmalted and malted mash types of the Madjeru sorghum cultivar. Optimizing studies also showed that Brewers Protease was the next important enzyme in increasing reducing sugars yields, as its need in both mash types was demonstrated. Bioglucanase TX was not indispensable for the malted mash type as shown by the optimal triplet of mashing enzymes obtained for this sample. The role of the milling process also independently facilitated the dissolution of some of these reducing sugars during mashing of both unmalted and malted Madjeru grist. Finally, the response surface methodology used in this work permitted to confirm the specific virtues always allotted to the malting process as an unavoidable route in providing the best malt types that can be used in mashing and beer brewing.

## ACKNOWLEDGEMENTS

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Full Length Research Paper

# Modeling the action of technical mashing enzymes on extracts and free-amino nitrogen yields of the *Madjeru* sorghum cultivar

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The action of three technical mashing enzymes (hitempase 2XL, bioglucanase-TX and brewers protease) on yields of extract and free amino nitrogen (FAN) of the worts of mashes of unmalted and malted Madjeru sorghum was modeled and analyzed using the response surface methodology. The analysis showed that increasing amounts of hitempase 2XL considerably increased yields of extract during mashing of unmalted Madjeru sorghum grist. The use of bioglucanase-TX was not indispensable, while Brewers' protease contributed very little. Increasing amounts of hitempase contributed approximately 45% of the free amino nitrogen, while Brewers' protease influence amounted to not more than 15%. Bioglucanase's action was globally nil. Addition of the three enzymes into malted Madjeru sorghum mashes had no significant effect on the yields of extracts and FAN, but the milling operation singularly liberated more than 50% of FAN for both mash types. Optimization of the concerted actions of the three enzymes for extract yield for unmalted Madjeru sorghum mash gave a combination (1960.5 U; 132.61 BGU and 28.86 mg) for hitempase, bioglucanase and of brewers protease respectively). This gave a maximal extract yield of 16.55 °P. This combination was: 2610 U; 0 BGU and 40.44 mg for malted Madjeru sorghum mash, giving a maximal extract yield of 16.35 °P. Optimization for free amino nitrogen for unmalted Madjeru sorghum mash gave a combination of: 3000 U; 0 BGU and 100 mg for hitempase, bioglucanase and brewers protease respectively). This gave maximal FAN of 93.55 mg/L. The combination was: 3000 U; 0 BGU and 100 mg for malted Madjeru sorghum mash, giving a maximal FAN of 144.48 mg/L.

**Key words:** Modeling, technical mashing enzymes, yields of extract, free-amino-nitrogen, *Madjeru*, optimization.

## INTRODUCTION

Sorghum in its malted form or as adjuncts has become a potential brewing cereal particularly in the tropics where barley is not grown (Taylor, 1983; Aisien and Muts, 1987; Arri, 1989; Palmer, 1989; Adejemilua, 1995). Its low contents of potential mashing enzymes (Aisien, 1982; EtokAkpan and Palmer, 1990; Nso et al., 2003, 2006) due to their poor development during malting has often

triggered the use of technical mashing enzymes as supplements (Agu and Palmer, 1998; Goode et al., 2002, 2003; Goode and Arendt, 2003) to achieve higher yields in extracts and other important wort specifications in beer brewing. Modeling and optimization approaches to ameliorate wort properties of sorghum grist and buckwheat malts has recently thrown more light into the precise role played by supplements of technical mashing enzymes (Goode et al., 2003, Phiarais et al., 2006, Desobgo et al., 2010). It is not however clear whether the use of technical mashing enzymes to obtain optimal mashing and brewing specifications for worts when using malted

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sorghum is indispensable. A series of work using two popular sorghum cultivars used in northern Cameroon for the production of the traditional beer Bili-Bili is currently taking place. This will help to have a clear understanding on the necessity of applying or not, technical enzymes when mashing with these cultivars. Some of the findings were recently reported for yields of reducing sugars for these cultivars (Desobgo et al., 2011a, b).

As a follow up, the effect of the same three technical mashing enzymes (Hitempase 2XL, Bioglucanase B-10L and Brewers protease) were modeled and optimized in this work for yields of extracts and free amino nitrogen during mashing of *Madjeru*, one of the popular sorghum cultivars of Northern Cameroon, using the response surface methodology (RSM).

#### MATERIALS AND METHODS

#### Enzymes

The characteristics of the technical enzymes used (Hitempase 2XL, a thermo stable  $\alpha$ -amylase from *Baccillus licheniformis*, Brewers protease from *Bacillus amyloliquefaciens* and Bioglucanase TX, an enzymatic composition of  $\beta$ -glucanase and hemicellulases from *Trichodermareesei*) and their sources are presented in Table 1.

#### Sorghum cultivar

The *Safrari* sorghum cultivar was obtained from the Institute of Research and agronomic development (IRAD) Maroua, Cameroon.

#### Modelling

Modelling was carried out as previously described (Desobgo et al., 2011a, b).

#### Validation of models

The models were validated as previously described (Desobgo et al., 2011a, b).

#### Mashing

Mashing was carried out as described (Desobgo et al., 2011a, b).

#### **Determination of extract**

Extract was determined as described in analytica-EBC, 1998.

#### Determination of free amino nitrogen

Free amino nitrogen (FAN) was determined as described in analytica-EBC, 1998.

#### **Optimization of models**

Models were optimized as previously described (Desobgo et al., 2011a, b).

## **RESULTS AND DISCUSSION**

The modeling and optimization of the action of mashing enzymes on the two key wort properties: extract and free amino nitrogen (FAN), was carried out for the experimental design required for manipulation in the laboratory. Table 2 shows the results obtained for extracts and free amino nitrogen (FAN) after mashing unmalted and malted *Madjeru* using the technical enzymes hitempase 2XL ( $\alpha$ amylase), bioglucanase TX ( $\beta$ -glucanase) and brewers protease (protease).

## Modeling and validation of results of yields of extract

The mathematical models obtained for extracts after mashing unmalted and malted *Madjeru* were as follows respectively:

With:  $Y_{MadEX}$  (X₁, X₂, X₃) representing the mathematical model for unmalted *Madjeru*;  $Y_{MadMEX}$  (X₁, X₂, X₃), the model for malted *Madjeru*; X₁, Hitempase; X₂, Bioglucanase and X₃ Brewers Protease.

The mathematical models were polynomials having several variables with coefficients of determination  $R^2$  = 0.940 for unmalted *Madjeru* and  $R^2 = 0.980$  for malted Madjeru. These coefficients, coupled to AAD values of 0.091 and 0.006 for unmalted and malted Madjeru respectively, allowed for the validation of the models for vields of extract of the worts. In addition, a bias factor of 1.05 and 1 for unmalted and malted *Madjeru* respectively, coupled to exactitude factors of 1.19 and 1.01 for both unmalted and malted Madjeru respectively, also allowed for validation of the models according to the method described (Ross, 1996). The factors of the models were linear or of first degree (X₁, X₂ and X₃), quadratic or of the second degree (X₁², X₂² and X₃²) and of interaction form (X1X2, X1X3, X2X3). They were statistically considered significant or not if the probability (P) of increasing yields of extracts was  $\leq 0.05$  or  $\geq 0.05$  respectively (Table 3).

## Effect of hitempase 2XL on yields of extract

The impact of hitempase 2XL as sole mashing enzyme on the yield of extract of unmalted and malted *Madjeru* is shown in Figure 1A. Extract yield increased from 2 °P with increasing concentration of enzyme for unmalted *Madjeru* mash to attain a maximum level (15.72 °P) at **Table 1.** Characteristics of commercial mashing enzyme preparations.

Commercial mashing enzyme	Organism of origin	Activity	Description	Temperature optima (°C)	pH optima	Recommended application level in adjuncts	Form
Hitempase 2XL	Bacillus licheniformis	4416.29 ± 19.34 U/ml	α-amylase	60 – 95	4 – 8	60 U/g	Solution
Bioglucanase TX	Trichoderma reesei	750 BGU/ml	β-glucanase	60	4.5 – 6.5	0.01 et 0.025% (v/w)	Solution
Brewers Protease	Bacillus amyloliquefaciens	1842.2 ± 1.8 mg FAN/min/mL	Protease	45 – 50 (denatured at 85)	6.5 – 7.5	0.4 – 2 g/Kg	Solution

Hitempase 2XL and bioglucanase TX were obtained from Kerry bioscience; Kilnagleary, Carrigaline, Co. Cork, Ireland. Brewers protease was obtained from DSM Food Specialities, Cedex France.

Table 2. Matrices of Doehlert coded and transformed experimental values.

	Coded value		Transformed experimental value			Madjeru											
Hit	Bio	Brew Prot	Hit	Bio	Brow Brot (mg)	Unmalted malted											
пі			(U)	(BGU)	Brew Prot (mg)	E	xtract (°P)		F	AN (mg/L)		E	xtract (°F	<b>)</b>		FAN (mg/l	L)
<b>X</b> 1	X ₂	<b>X</b> 3	<b>X</b> 1	<b>X</b> 2	<b>X</b> 3	Expa	Theo⁵	Res⁰	Exp	Theo	Res	Exp	Theo	Res	Exp	Theo	Res
0.000	0.000	0.000	1500	468.75	50	15.07	15.13	-0.06	64.34	63.50	0.84	15.60	15.57	0.03	94.00	94.81	-0.81
1.000	0.000	0.000	3000	468.75	50	13.93	12.10	1.83	71.79	71.17	0.62	15.70	15.49	0.21	107.70	107.32	0.38
0.500	0.866	0.000	2250	937.5	50	13.10	13.99	-0.89	98.00	94.42	3.58	15.08	15.17	-0.09	147.00	141.67	5.33
-0.500	-0.866	0.000	750	0.00	50	13.61	11.93	1.68	65.38	64.31	1.07	14.56	14.47	0.09	98.10	95.82	2.28
0.500	-0.866	0.000	2250	0.00	50	15.09	15.56	-0.47	83.08	83.01	0.07	16.12	16.24	-0.12	126.00	125.54	0.46
-0.500	0.866	0.000	750	937.5	50	9.05	8.66	0.39	81.79	82.30	-0.51	14.97	14.86	0.11	122.70	123.85	-1.15
0.500	0.289	0.816	2250	615.18	100	10.00	10.45	-0.45	84.78	86.08	-1.30	15.14	15.27	-0.13	130.00	130.96	-0.96
-0.500	-0.289	-0.816	750	312.32	0.0	11.25	9.60	1.65	44.00	35.71	8.29	14.97	14.85	0.12	66.00	53.61	12.39
0.500	-0.289	-0.816	2250	312.32	0.0	14.14	15.55	-1.41	48.00	48.97	-0.97	15.39	15.49	-0.10	72.00	73.27	-1.27
0.000	0.577	-0.816	1500	781.07	0.0	13.20	12.79	0.41	49.07	52.58	-3.51	14.76	14.80	-0.04	73.60	78.48	-4.88
-0.500	0.289	0.816	750	615.18	100	8.74	7.45	1.29	68.86	68.53	0.33	13.93	13.84	0.09	103.30	103.07	0.23
0.000	-0.577	0.816	1500	156.43	100	10.73	11.58	-0.85	73.80	72.84	0.96	14.56	14.53	0.03	110.70	109.97	0.73
0.000	0.000	0.000	1500	468.75	50	14.56	15.13	-0.57	62.27	63.50	-1.23	15.60	15.57	0.03	93.40	94.81	-1.41
-1.000	0.000	0.000	0.000	468.75	50	1.52	3.15	-1.63	40.88	40.35	0.53	13.20	13.42	-0.22	61.30	59.78	1.52
-1.000	-0.866	-0.816	0.000	0.0	0.0	1.20	1.86	-0.66	33.31	37.13	-3.82	13.52	13.52	0.00	50.00	56.24	-6.24
0.000	0.000	0.000	1500	468.75	50	14.90	15.13	-0.23	62.00	63.50	-1.50	15.60	15.57	0.03	93.00	94.81	-1.81
0.000	0.000	0.000	1500	468.75	50	15.08	15.13	-0.05	60.00	63.50	-3.50	15.46	15.57	-0.11	90.00	94.81	-4.81

^aExperimental result values. ^bTheoretical values (values coming from mathematical models). ^c Residue.

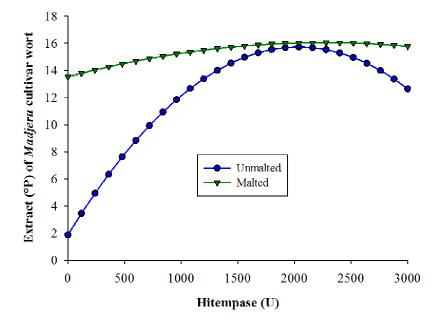
about 2038 U, followed by a slight and steady decrease thereafter. It was already high (13.52 °P) in the absence of Hitempase for malted *Madjeru* mash and only increased slightly to attain

a maximal level (16.04 °P) at 2268 U of enzyme concentration. When the mathematical models were applied to predict the impact of supplements of bioglucanase and brewers protease at

concentrations of 750 BGU and 60 mg respectively and in the presence of Hitempase, it was observed that the profile of extract yields remained similar to that in Figure 1a (compare with

	Coeffic	cient	Std. dev	viation	t-stati	stics	P-value	
Effect	Unmalted	Malted	Unmalted	Malted	Unmalted	Malted	Unmalted	Malted
Constant	15.134	15.566	0.766	0.08	19.768	195.176	0.000	0.000
X ₁	4.477	1.036	0.788	0.082	5.684	12.63	0.001	0.000
X ₂	-1.397	-0.197	0.687	0.072	-1.762	-2.387	0.121	0.048
X ₃	-1.729	-0.304	0.649	0.068	-2.174	-3.67	0.066	0.008
$X_1^2$	-7.507	-1.11	1.35	0.141	-5.56	-7.895	0.001	0.000
$X_2^2$	-0.964	-0.14	0.983	0.102	-0.736	-1.027	0.486	0.339
$X_3^2$	-3.732	-0.843	0.848	0.088	-2.932	-6.357	0.022	0.000
X ₁ *X ₂	0.98	-0.842	1.541	0.161	0.55	-4.543	0.599	0.003
$X_{2}^{*}X_{3}$	0.296	0.623	1.357	0.141	0.154	3.115	0.882	0.017
X ₁ *X ₃	-2.155	0.784	1.606	0.167	-1.095	3.823	0.310	0.007

Table 3. Estimation of regression coefficients for the extracts of umalted and malted Madjeru.

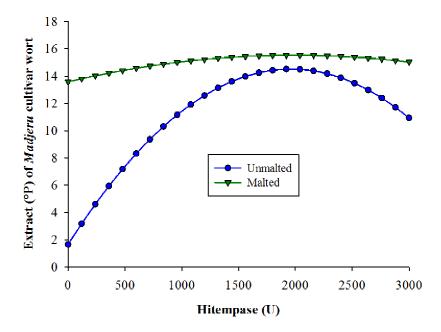


**Figure 1a.** Effect of concentration of hitempase ( $\alpha$ -amylase) as sole mashing enzyme on yield of wort extract of sorghum cultivar *Madjeru*.

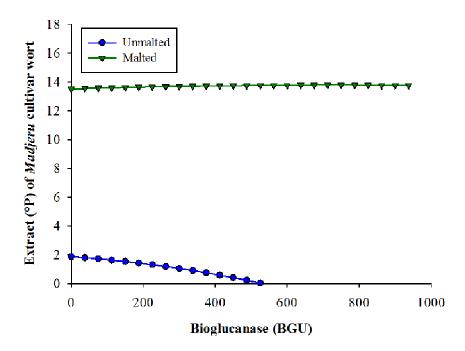
Figure 1b). Starch is indeed the main macromolecule of cereals and the main substrate of  $\alpha$ -amylase. It is therefore expected that this enzyme contributes to the greatest amount of soluble materials that could be found in resulting worts due to its action on starch (Goode et al., 2003, Phiarais et al., 2006, Desobgo et al., 2010). Figures 1 A and B also showed that supplements of hitempase in unmalted *Madjeru* mash was by far more useful than for the malted *Madjeru* mash type. The important role of milling in obtaining instantaneous dissolution of soluble materials at the beginning of mashing was better displayed for the malted *Madjeru* mash type (93 and 14% of extract yields respectively). The soluble nitrogenous compounds and reducing sugars (Hough et al., 1982) in

the medium (Figures 1a and b).

From the mathematical models, it was shown that in its linear form (X₁), hitempase contributed 19 and 18% of extract yields for unmalted and malted *Madjeru* respectively (Figures 4a and b). This more or less equitable contribution of extracts in the worts of the two mash types, pairs with the observations made earlier whereby extract levels were virtually the same for both. Moreover, statistical analyses also showed that this contribution was significant (P = 0.001 and 0.000 for unmalted and malted *Madjeru* respectively (Table 3). In its quadratic form (X₁²), hitempase remained statistically significant for mashing both unmalted *Madjeru* and malted *Madjeru* (P = 0.001 and 0.000 respectively). This confirmed the earlier biological observation according to which supplements of this



**Figure 1b.** Effect of concentration of hitempase ( $\alpha$ -amylase) in the presence of fixed concentrations of bioglucanase (750 BGU) and brewers protease (60 mg) on yield of wort extract of sorghum cultivar *Madjeru* 

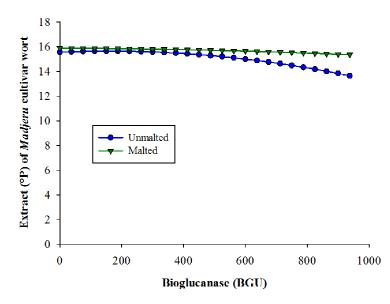


**Figure 2a.** Effect of concentration of bioglucanase ( $\beta$ -glucanase) as sole mashing enzyme on yield of wort extract of sorghum cultivar *Madjeru*.

enzyme in malted mashes of *Madjeru*, was also necessary. Its contribution in increasing extract yields in its quadratic form  $(X_1^2)$  (excess of  $\alpha$ -amylase in principle) was indeed 33 and 19% for unmalted and malted *Madjeru* respectively (Figures 4a and b).

## Effect of bioglucanase TX on yields of extract

Figure 2a shows the effect of mashing unmalted and malted *Madjeru* using bioglucanase, as sole mashing enzyme, on yields of extract. There was a progressive



**Figure 2b.** Effect of concentration of bioglucanase ( $\beta$ -glucanase) in the presence of fixed concentrations of hitempase (1875 U) and brewers protease (60 mg) on yield of wort extract of sorghum cultivar *Madjeru*.

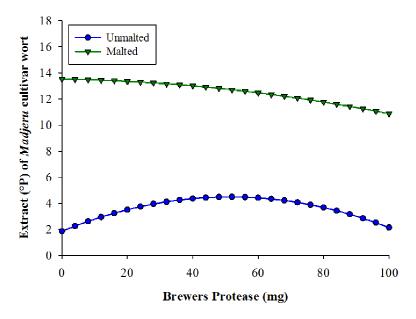
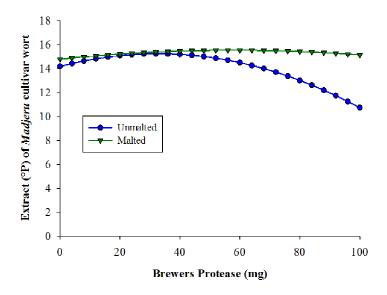
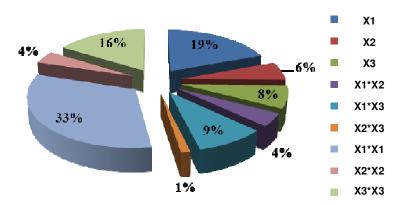


Figure 3a. Effect of concentration of brewers protease as sole mashing enzyme on yield of wort extract of sorghum cultivar *Madjeru*.

decrease of yield of extract from 1.86 °P to nil as enzyme concentration increased for unmalted *Madjeru* wort. This figure also shows that the extract obtained (1.86 °P) was probably completely due to milling, suggesting that the enzyme plays no important role in production of extract from the grist of the unmalted *Madjeru* cultivar during mashing. The yield of extract for malted *Madjeru* mash was virtually at its maximal level (13.78 °P) even in the absence of bioglucanase and remained virtually constant with increasing enzyme concentration. Bioglucanase was therefore not a backbone enzyme for extract production during mashing. A similar application of the mathematical models as carried out earlier for hitempase's action, using 60 mg of brewers protease and 1875 U hitempase, predicted that the supplementation of these two mashing enzymes could provide similar results in extract yields for both unmalted and malted *Madjeru* mashes (Figure 2B). Hitempase once more revealed that it was the real



**Figure 3b.** Effect of concentration of brewers protease in the presence of fixed concentrations of hitempase (1875 U) and bioglucanase (750 BGU) on yield of wort extract of sorghum cultivar *Madjeru*.



**Figure 4a.** Contribution to yield of wort extract (°P) of each factor in its linear, quadratic and interaction (combined) forms for unmalted sorghum cultivar *Madjeru*.

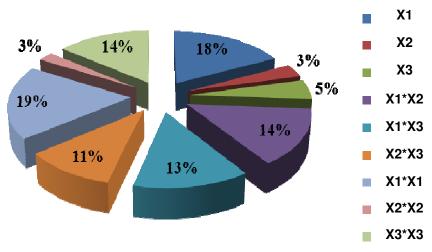
backbone enzyme that contributed for most of the yields of extract. The level of extract yields reached almost 15.54 °P for unmalted *Madjeru* and 15.87 °P for malted *Madjeru* mash types. From the two figures, it is also important to underline the natural virtues of the malting procedure in rendering the Madjeru grains potentially mash-able to reasonable extract yields in the absence of all these enzymes. These observations were all statistically confirmed. Indeed, in its linear form (X₂), bioglucanase's action was not significant (P = 0.121) for unmalted but significant for malted *Madjeru* (P = 0.048) (Table 3).

Figures 4A and B showed that this enzyme contributed barely for 6 and 3% of extract yields for unmalted and

malted *Madjeru* respectively. In its quadratic form  $(X_2^2)$  (excess of enzyme in principle), bioglucanase contributed 4 and 3% of extract yield for unmalted and malted *Madjeru* respectively (Figures 4a and b). Similarly, these contributions were statistically not significant for both unmalted and malted *Madjeru* mash types (p = 0.486 and 0.0.339 respectively) (Table 3).

## Effect of brewers protease on yields of extract

The effect of mashing unmalted and malted *Madjeru* on yields of extract using as sole mashing enzyme, "brewers protease", is shown in Figure 3A. Yields of extracts



**Figure 4b.** Contribution to yield of wort extract (°P) of each factor in its linear, quadratic and interaction (combined) forms for malted sorghum cultivar *Madjeru*.

 Table 4. ANOVA for the extracts of umalted and malted Madjeru.

0		Sum sq	uare	Mean square		F-val	ue	P-val	ue	
Source	Ddl	Unmalted	Malted	Unmalted	Malted	Unmalted	Malted	Unmalted	Malted	
Regression	9	285.91	9.965	31.768	1.107	12.14	38.993	0.002	0.000	
Linear	3	156.256	6.507	52.085	2.169	19.905	76.379	0.001	0.000	
Quadratic	3	126.512	2.496	42.171	0.832	16.116	29.299	0.002	0.000	
Interactions	3	3.141	0.963	1.047	0.321	0.400	11.302	0.757	0.004	
Residual error	7	18.317	0.199	2.617	0.028					
Total error	16	304.227	10.164							

gradually increased from 1.86 °P with increasing amounts of enzyme to reach a maximum of 4.48 °P at enzyme concentration of 51.39 mg. This small yield of extract due to the action of this enzyme could be attributed to its capacity of hydrolyzing proteins into soluble amino acids and peptides (Briggs et al., 2004) during which additional soluble sugars could also be liberated. Figure 3a however shows once more that in the absence of supplements of "brewers protease", maximal extract was obtained when mashing with malted Madieru. This could be once more attributed to the virtues well known to the malting process as explained earlier. These results confirmed earlier observations (Desobgo et al., 2011a, b). The gradual and slight decrease of extract yields with increasing amounts of "brewers protease" for malted Madjeru and unmalted Madjeru (as from 51.39 mg concentration) mash types could once more be attributed to reactions between soluble nitrogenous functions and some of the soluble sugars. The mathematical models were once more used to predict the yield in extract as carried out earlier for hitempase and bioglucanase actions. Thus, using 1875 U hitempase and 750 BGU of bioglucanase with increasing amounts of "brewers protease", the models once more showed that the adding of these mashing enzymes could provide similar results in extract yields for both unmalted and malted Madjeru

mashes (Figure 3b). The aforementioned observations were statistically confirmed using the mathematical model. In its first degree form (X₃), the impact of "brewers protease" was not significant for unmalted *Madjeru* mash but was for malted *Madjeru* mash (P = 0.066 and 0.008 respectively) (Table 3). Its contribution to extract yield was 8 and 5% respectively for both unmalted and malted mashes (Figures 4a and b).

The impact of the enzyme in its quadratic form  $(X_3^2)$  was significant for both unmalted and malted mashes (P = 0.022 and 0.000 respectively) (Table 3). Its contribution to extract yield was 16 and 14% respectively (Figures 4a and 4).

## Effect of enzymes' interactions on yields of extract

The models were further exploited to predict the impacts of enzyme interactions  $(X_1X_2, X_1X_3 \text{ and } X_2X_3)$  on yields of extract. The results are shown in Figures 4a and b. They were globally not statistically significant for unmalted *Madjeru* mashes (P = 0.757), but were for malted *Madjeru* mashes (P = 0.004) (Table 4). The interaction  $X_1X_2$  (hitempase/bioglucanase) had no significant impact on unmalted *Madjeru* mash, but had for malted *Madjeru* (P = 0.599 and 0.003 respectively (Table 3). It contributed

Source	DF	Sum of squares	Mean of squares	F-value	P-value
Inter-groups	1	102.344	102.344	10.420	0.003
Intra-groups	32	314.391	9.824		
Total	33	416.735			

**Table 5.** ANOVA for comparing extracts of unmalted and malted Madjeru worts.

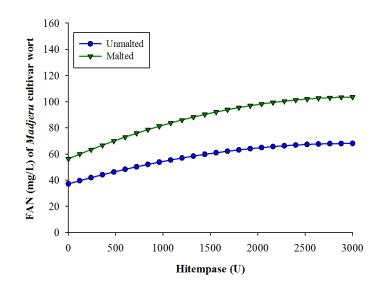


Figure 5a. Effect of concentration of hitempase ( $\alpha$ -amylase) as sole mashing enzyme on yield of wort free amino nitrogen of sorghum cultivar *Madjeru*.

for merely 4% of extract for unmalted Madjeru mash, but up to 14% for malted Madjeru mash (Figures 4a and b). It is however significant to underline that this important contribution could be attributed to the intrinsic virtues that malting offers when mashing with malted Madjeru and not to the hitempase/bioglucanase interaction as such. Though known to be the backbone starch hydrolyzing enzyme, the action of hitempase is best exploited when the cell walls of cereal grains are broken down by βglucanases, hemicellulases and cellulases to liberate starch granules. This sequence of events during malting was confirmed by the mathematical models aforementioned (Desobgo et al., 2010). The interaction  $X_1X_3$ corresponding to the couple hitempase/brewers protease. also had no significant impact on extract yields of unmalted Madjeru mash (P = 0.310), but had on malted *Madjeru* mash (P = 0.007) (Table 3). Its contribution to extract yield was 9 and 13% respectively (Figures 4a and b). This result was once more in conformity with the biological sequence occurring during malting. Efficient starch hydrolysis by a-amylase indeed occurs only after the breakdown of cell walls by  $\beta$ -glucanase followed by liberation of starch granules due to proteolysis of the protein matrix enrobing them. The interaction bioglucanase/ brewers protease  $(X_2X_3)$  had no significant impact on extract yields for unmalted *Madjeru* mash (P = 0.882), but

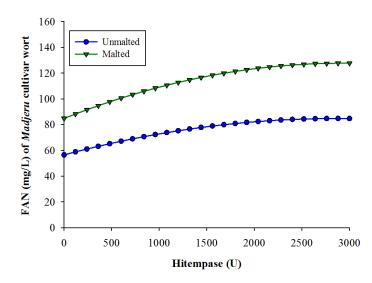
had for malted *Madjeru* mashes (P = 0.017) (Table 3). Its contribution to extract yields was 1 and 11% respectively for both mash types (Figures 4a and b). These low contributions by the couple (bioglucanase/brewers protease) were expected, as the two enzymes only play a supporting role in starch hydrolysis during mashing (Desobgo et al., 2010).

Table 5 statistically confirms the observation that malted *Madjeru* worts associated with the technical enzymes have better yields of extracts than unmalted *Madjeru* worts (P = 0.003).

## Modeling and validation of results of free amino nitrogen (FAN)

The mathematical models obtained for free amino nitrogen (FAN) for mashing unmalted and malted *Madjeru* were as follows respectively:

$$Y_{MadMAAL}(X_1, X_2, X_3) = 94.807 + 23.771X_1 + 12.745X_2$$



**Figure 5b.** Effect of concentration of hitempase ( $\alpha$ -amylase) in the presence of fixed concentrations of bioglucanase (750 BGU) and brewers protease (60 mg) on yield of wort free amino nitrogen of sorghum cultivar *Madjeru*.

Table 6. Estimation of regression coefficients for free amino nitrogen of umalted and malted Madjeru.

<b>F</b> He etc	Coeffic	cient	Std. dev	viation	t-statis	stics	P-value	
Effects	Unmalted	Malted	Unmalted	Malted	Unmalted	Malted	Unmalted	Malted
Constant	63.504	94.807	2.047	3.024	31.026	31.347	0.000	0.000
X ₁	15.411	23.771	2.106	3.111	7.319	7.641	0.000	0.000
X ₂	8.487	12.745	1.836	2.713	4.003	4.069	0.005	0.005
X ₃	18.418	28.311	1.735	2.563	8.663	9.012	0.000	0.000
$X_1^2$	-7.747	-11.26	3.61	5.334	-2.146	-2.111	0.069	0.073
$X_2^2$	25.922	36.639	2.627	3.882	7.4	7.658	0.000	0.000
$X_{3}^{2}$	-8.636	-11.98	2.266	3.348	-2.538	-2.383	0.039	0.049
X ₁ *X ₂	-3.798	-6.875	4.121	6.089	-0.798	-0.978	0.451	0.361
$X_{2}^{*}X_{3}$	7.807	12.315	3.627	5.359	1.521	1.624	0.172	0.148
X ₁ *X ₃	3.976	7.478	4.294	6.345	0.756	0.962	0.475	0.368

 $+28.311X_{3} - 6.875X_{1}X_{2} + 7.478X_{1}X_{3} + 12.315X_{2}X_{3} - 11.261X_{1}^{2} + 39.639X_{2}^{2} - 11.983X_{3}^{2}$ 

With:  $Y_{MadAAL}$  (X₁, X₂, X₃) representing the mathematical model for unmalted *Madjeru*;  $Y_{MadMAAL}$  (X₁, X₂, X₃) for malted *Madjeru*; X₁, hitempase; X₂, bioglucanase and X₃, brewers protease (protéase). These mathematical models were once more polynomials having several variables with determination coefficients of R² = 0.973 for unmalted *Madjeru* and R² = 0.974 for malted *Madjeru*. These coefficients, coupled to AAD values of 0.037 and 0.036 for unmalted and malted *Madjeru* respectively, allowed for the validation of the models for assessment of the wort free amino nitrogen content. In addition, a bias factor of 1 for both unmalted and malted *Madjeru* mash types, coupled to exactitude factors of 1.00 for both mash types, also allowed for validation of the models according to the method described (Ross, 1996). The factors of the models were once more linear or of first degree ( $X_1$ ,  $X_2$  and  $X_3$ ), quadratic or of the second degree ( $X_1^2$ ,  $X_2^2$  and  $X_3^2$ ) or of interaction form ( $X_1X_2$ ,  $X_1X_3$ , and  $X_2X_3$ ). They were statistically considered significant or not if the probability (P) in increasing yields of FAN was  $\leq$  0.05 or  $\geq$  0.05 respectively (Table 6).

# Effect of hitempase 2XL on yields of free amino nitrogen (FAN)

The impact of hitempase as sole mashing enzyme on wort FAN for unmalted and malted *Madjeru* is shown in Figure 5a. Free amino nitrogen content of wort gradually increased from 37.12 mg/L with increasing enzyme

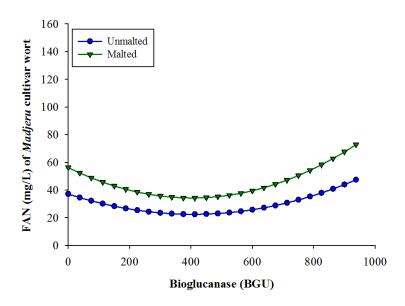
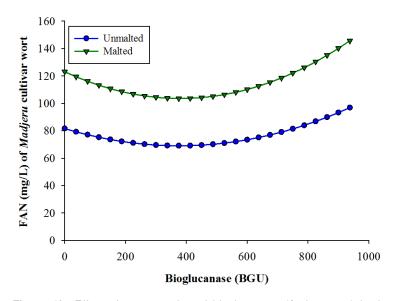


Figure 6a. Effect of concentration of bioglucanase ( $\beta$ -glucanase) as sole mashing enzyme on yield of wort free amino nitrogen of sorghum cultivar *Madjeru*.



**Figure 6b.** Effect of concentration of bioglucanase ( $\beta$ -glucanase) in the presence of fixed concentrations of hitempase (1875 U) and brewers protease (60 mg) on yield of wort free amino nitrogen of sorghum cultivar *Madjeru*.

concentration to reach a maximum of 68.04 mg/L for both unmalted mashes and from 56.26 mg/L to reach a maximum of 103.51 mg/L for malted *Madjeru* mashes. The Figure also showed that for both mash types, the yields in FAN in the absence of the enzyme represented more than 50% of the final yields. This suggests that the milling operation was at the basis of the instantaneous dissolution of these considerable amounts of free amino nitrogen at the start of mashing. Although hitempase is not a protein hydrolyzing enzyme, it exposes more free amino nitrogen functions upon acting on globular proteins and starch granules. This could explain the increase in FAN observed with increase in enzyme concentration. The higher FAN content for malted *Madjeru* mash compared to unmalted *Madjeru* mash is once more to be attributed to the natural virtues that the grains incur during

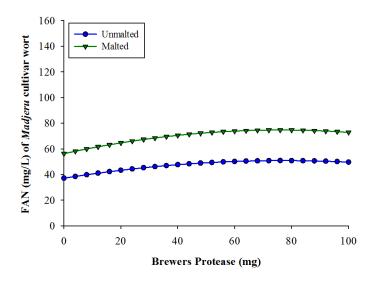


Figure 7a. Effect of concentration of brewers protease as sole mashing enzyme on yield of wort free amino nitrogen of sorghum cultivar *Madjeru*.

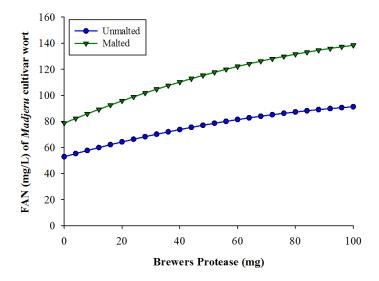


Figure 7b. Effect of concentration of brewers protease in the presence of fixed concentrations of hitempase (1875 U) and bioglucanase (750 BGU) on yield of wort free amino nitrogen of sorghum cultivar *Madjeru*.

during the malting process. Use of the models to predict the profile of FAN content of worts if the mashing enzymes bioglucanase (at 750 BGU) and brewers protease (at 60 mg) were coupled to hitempase's action, showed a profile of increases in yields of FAN similar to that observed in Figure 5b with increments contributing to amounts equivalent to roughly 15%. The models also showed that hitempase (X₁), in its first degree form, contributed for 15% of the FAN content of both the unmalted and malted *Madjeru* mashes (Figures 8a and b). This contribution was statistically significant for the two mash types (P = 0.000) (Table 6). Similarly, in its quadratic form  $(X_1^2)$ , hitempase's effect was not significant for the two mash types (P = 0.069 and 0.073 respectively) (Table 6). Its contribution in this form was 8 and 7% respectively (Figures 8a and b).

# Effect of bioglucanase TX on yields of free amino nitrogen (FAN)

Figure 6a shows the effect of bioglucanase on the FAN content as sole mashing enzyme in unmalted and malted *Madjeru*. This content initially decreased with increasing enzyme concentration in both mash types, dropping from

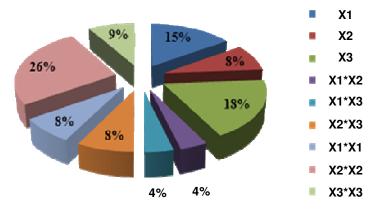
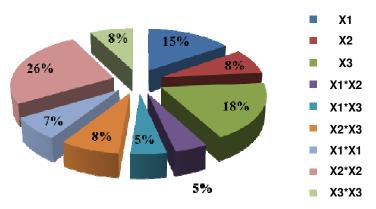


Figure 8a.Contribution to yield of wort free amino nitrogen (FAN) of each factor in its linear, quadratic and interaction (combined) forms for unmalted sorghum cultivar *Madjeru*.



**Figure 8b.** Contribution to yield of wort free amino nitrogen (FAN) of each factor in its linear, quadratic and interaction (combined) forms for malted sorghum cultivar *Madjeru*.

37.12 to a minimal of 22.47 mg/L for unmalted Madjeru, and from 56.26 to 34.25 mg/L for malted Madjeru, and henceforth gradually increased with increases in enzyme concentration as from 407 BGU. This observation indicated that bioglucanase is not a protein hydrolyzing enzyme. The initial decrease in FAN content, followed by a subsequent increase, could be explained by the fact that, the free amino nitrogen instantaneously dissolved in the mash after the milling process, reacted with soluble sugars, after which the hydrolyzing action of cell wall components by bioglucanase, kinetically become perceptible to permit observing the liberation of extra FAN molecules. Upon using the model to predict the amounts of FAN if mashed in the presence of hitempase (at 1875) U) and brewers protease (at 60 mg), the same profile as in Figure 6A were observed, but with the levels of free amino nitrogen contents for both mash types increasing 2 times as compared to the original contents (Figure 6b). This confirmed the need to have all mashing enzymes present in appropriate proportions during mashing to permit obtaining substantial amounts of FAN in worts. These observations once more displayed the role of the malting process in guaranteeing worts of higher brewing

#### properties.

According to the models, bioglucanase (X₂), in its linear form, contributed to 8% of the FAN content for both the unmalted and malted *Madjeru* mash types (Figures 8A and B). This contribution was statistically significant for the two mash types (P = 0.005) (Table 6). Similarly, in its quadratic form (X₂²), the effect of the enzyme remained significant for both mash types (P = 0.000) (Table 6). Its contribution in this form was 26% for both mash types (Figures 8a and b).

## Effect of brewers protease on yields of free amino nitrogen (FAN)

The effect of brewers protease as sole mashing enzyme on FAN content for unmalted and malted *Madjeru* is presented in Figure 7A. Free amino nitrogen content increased very slightly from 37.12 to 49.66 mg/L for unmalted *Madjeru* worts and from 56.26 to 72.86 mg/L for malted *Madjeru* worts with increasing enzyme concentrations. These levels were maintained virtually constant as from about 60 mg of enzyme input thereof. The models predicted that coupling hitempase (at 1875 U)

Courses	DE	Sum of	squares	Mean of	squares	F-val	ue	P-val	ue
Source	DF	Unmalted	Malted	Unmalted	Malted	Unmalted	Malted	Unmalted	Malted
Regression	9	4697.665	10845.612	521.963	1025.068	27.908	29.51	0.000	0.000
Linear	3	3323.582	7688.072	1107.861	2562.691	59.235	62.756	0.000	0.000
Quadratic	3	1311.862	2987.332	437.287	995.777	23.381	24.385	0.001	0.000
Interactions	3	62.221	170.208	20.74	56.736	1.109	1.389	0.407	0.323
Residual error	7	130.919	285.848	18.703	40.835				
Total error	16	4828.585	11131.46						

Table 7. ANOVA for free amino nitrogen of umalted and malted Madjeru.

Table 8. ANOVA for comparing free amino nitrogen of unmalted and malted Madjeru worts.

Source	DF	Sum of squares	Mean of squares	F-value	P-value
Inter-group	1	8813.2	8813.2	17.67	0.000
Intra-group	32	15960	498.751		
Total	33	24773.2			

and bioglucanase (at 750 BGU) to the action of brewers protease would induce a steady increase in FAN content (Figure 7b). This once more indicates the need of having all mashing enzymes present in order to obtain higher FAN yields. The additional FAN content observed for malted *Madjeru* mash as compared to unmalted *Madjeru* mash could once more be attributed to the natural virtues that the malting process contributed to the grains used. The mathematical model statistically showed that the action of brewers protease was in its linear form significant for both mash types (P = 0.000 for both (Table 6). Its contribution to FAN content in this form (X₃) was 18% for both the unmalted and malted *Madjeru* worts (Figures 8a and b).

The effect of the enzyme in its quadratic form  $(X_3^2)$  however remained significant for both mash types (P = 0.039 and 0.049 respectively) (Table 6). Its contribution was 9% for unmalted *Madjeru* worts and 8% for malted *Madjeru* worts (Figures 8a and b).

# Effect of enzymes interactions on yields of free amino nitrogen (FAN)

The global action of the enzymes' interaction or as coupled forms  $(X_1X_2, X_1X_3 \text{ and } X_2X_3)$  on the FAN content was statistically not significant (P = 0.407 for unmalted *Madjeru* and P = 0.323 for malted *Madjeru*) (Table 7). Their contributions of FAN content are shown in Figures 8a and b. The effect of the  $X_1X_2$  (hitempase 2XL/bioglucanase TX) interaction was not significant for both wort types (P = 0.451 for unmalted *Madjeru* and P = 0.361 for malted *Madjeru*) (Table 6). Its contribution of FAN content in both mash types was 4 and 5% respectively) (Figures 8a and b). Similarly, the action of the couple  $X_1X_3$  (hitempase 2XL/brewers protease) was also

not significant for both mash types (P = 0.475 for unmalted *Madjeru* and P = 0.368 for malted *Madjeru*) (Table 6). Its contribution of FAN content in both mash types was 5 and 4% respectively (Figures 8a and b). Finally, for the couple  $X_2X_3$  (BIOGLUCANASETX/brewers protease), its action was also not significant for both wort types (P = 0.172 for unmalted *Madjeru* and P = 0.148 for malted *Madjeru*) (Table 6). The contribution of FAN content in both wort types was 8% for each (Figures 8a and b).

Table 8 statistically confirmed the biological assertion that malted type samples were more potential raw materials for mashing in terms of FAN contents than the combination of unmalted grains and commercial enzymes (P = 0.000).

# Optimization of the concerted mashing enzymes' action on yields of extracts and free amino nitrogen

The results obtained for the action of the enzymes on extract and free amino nitrogen yields after mashing on the basis of the models were optimized to define satisfactory domains of compromise for the mashing enzymes. These domains were obtained for the two key brewing parameters by fixing the wort conditions at: extract  $\geq$  12 °P and free amino nitrogen  $\geq$  80 mg/L. The theoretical optimal combination of enzyme action for unmalted Madjeru gave the following triplet of real variables for extract: 1960.5 U; 132.61 BGU and 28.86 mg for hitempase 2XL. Bioglucanase TX and brewers protease respectively. This triplet allowed for a maximal extract of 16.55 °P. The triplet for malted Madjeru was: 2610 U; 0 BGU and 40.44 mg. It allowed for a maximal extract of 16.35 °P. The optimal enzyme combinations were thus different particularly with regards to

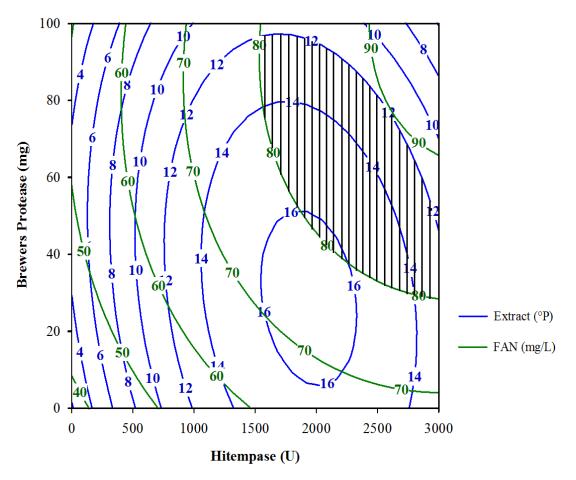


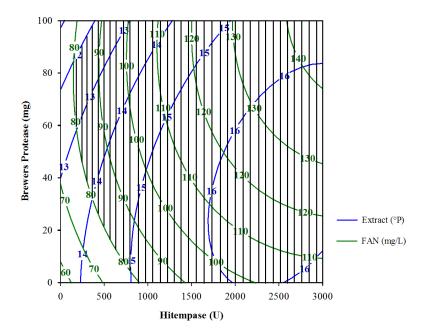
Figure 9a. Response surface curves for the enzyme combinations providing for optimal yields in extract and free amino nitrogen for unmalted sorghum cultivar *Madjeru*.

bioglucanase TX but both gave comparable yields of extracts. These results show that bioglucanase TX supplements for mashing malted Madjeru is not indispensable for obtaining required yields in extract, whereas the three enzymes are needed as supplements when mashing unmalted Madjeru. Figure 9a shows the response surface areas exploitable for efficient mashing capable of giving optimal results of yields of extract. For free amino nitrogen, the theoretical optimal combination of enzymes' action for unmalted Madjeru gave as triplet of real variables: 3000 U; 0 BGU and 100 mg for hitempase 2XL. bioglucanase ТΧ and brewers protease respectively. This triplet allowed for maximal free amino nitrogen of 93.55 mg/L. The triplet for malted Madjeru was: 3000 U; 0 BGU and 100 mg. It allowed for maximal free amino nitrogen content of 144.48 mg/L. These results show that bioglucanase TX is not an indispensable mashing enzyme when seeking for appreciable amounts of free amino nitrogen in the worts of Madjeru, be it malted or not.

Figure 9b shows the response surface areas exploitable for efficient mashing giving optimal results of free amino nitrogen content.

# Conclusions

The effects of three technical mashing enzymes (hitempase 2XL, bioglucanase TX and brewers protease) on yields of extract and free amino nitrogen were studied during the mashing of unmalted and malted Madjeru grist. Hitempase 2XL was principally responsible for extract yields in unmalted Madjeru mash but its impact on malted Madjeru mash type was mild. Bioglucanase TX played no role, while brewers protease showed limited contributions to yields of extract. Hitempase 2XL and brewers protease individually contributed to yields in free amino nitrogen in both unmalted and malted Madjeru mashes, though the milling operation contributed to FAN yields for more than 50% in both mashes. This study shows that proper malting and mashing of this sorghum cultivar could lead to satisfactory worts properties in terms of extract and free amino nitrogen for brewing purposes. Supplements of technical mashing enzymes to boost their yields of extract in particular, are thus not indispensable when mashing with malted Madjeru. Optimization of mashing properties through models clearly describing the actions of individual technical mashing



**Figure 9b.** Response surface curves for the enzyme combinations providing for optimal yields in extract and free amino nitrogen for malted sorghum cultivar *Madjeru*.

enzymes, as displayed in this study using the response surface methodology is however of interest particularly when mashing with high amounts of sorghum adjuncts. Further studies on the fermentability of worts obtained after such studies would be of importance in order to assess the exploitability of the results for improved brewing practices with this sorghum cultivar.

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# Optimisation of the Action of Commercial Mashing Enzymes on Wort Extracts and Free Amino Nitrogen of the *Safrari* Sorghum Cultivar

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#### ABSTRACT

The influence of three commercial mashing enzymes (Hitempase 2XL, Bioglucanase TX and Brewers protease) used as sole mashing enzymes on yields of extract and free amino nitrogen (FAN) of the worts of the mashes of unmalted and malted Safrari sorghum was studied using the response surface methodology. The study revealed that increasing amounts of Hitempase considerably increased yields of extract during mashing of unmalted Safrari grist, while the effect of Bioglucanase was smaller, and that of Brewers protease was insignificant. Extract yields decreased with increasing amounts of the three enzymes during the mashing of malted Safrari. This decrease was least expressed in the case of Brewers protease. Yields in FAN amounted to less than 50%, with increasing amounts of both Hitempase and Brewers protease, but constantly decreased to nil for Bioglucanase's action in both unmalted and malted Safrari mashes. The milling operation singularly liberated more than 50% of FAN for both mash types and for each of the enzymes. Optimisation of the concerted actions of the three enzymes for extract yields gave a combination of 2,098.5 U, 937.5 BGU and 0 mg (for Hitempase, Bioglucanase and Brewers protease, respectively) for unmalted Safrari sorghum mash. This gave a maximal extract yield of 18°P. The combination was 0 U, 28.68 BGU and 0 mg for malted Safrari sorghum mash, giving a maximal extract yield of 18.82°P. Optimisation for FAN gave a combination of 2,434.5 U, 0 BGU and 100 mg (for Hitempase, Bioglucanase and Brewers protease, respectively) for unmalted Safrari sorghum mash. This gave maximal FAN of 144.77 mg/L. The combination was 2,191.5 U, 0 BGU and 100 mg for malted Safrari sorghum mash, giving a maximal FAN of 196.73 mg/L.

Keywords: commercial mashing enzymes, extract, free amino nitrogen, optimisation, Safrari.

#### SÍNTESIS

Se estudiaron, utilizando el método de superficie de respuesta, tres enzimas comerciales de maceración (Hitempase 2XL, Bioglucanase TX y "Brewers protease") con respecto al rendimiento de extracto y de amino nitrógeno libre (FAN) en mostos de macerados de sorgo Safrari malteado y no malteado. El estudio reveló que un aumento en la cantidad de Hitempase aumentó considerablemente el rendimiento de extracto de un macerado de Safrari no malteado, mientras que el efecto de la Biogluconase era menor y el efecto de la "Brewers protease" era insignificante. El rendimiento de extracto disminuyó con un aumento en la cantidad de las tres enzimas en el macerado de Safrari malteado, siendo menor la disminución con el "Brewers protease". El rendimiento de FAN fue menor del 50 por ciento con el aumento en las cantidades tanto de Hitempase y de "Brewers protease" y con Biogluconase se fue bajando hasta cero, tanto en macerados de Safrari malteado como no malteado. La molienda liberó por sí solo más de 50 por ciento del FAN para los dos tipos de macerado y para cada uno de las enzimas. La optimización del uso conjunto de las tres enzimas para aumentar el rendimiento de extracto resultó en la combinación 2.098,5 U, 937,5 BGU y 0 mg (para Hitempase, Bioglucanase y "Brewers protease", respectivamente) para macerados de sorgo Safrari no malteado, resultando en un extracto máximo de 18°P. La combinación óptima para Safrari malteado fue de 0 U, 28,68 BGU y 0 mg dando un extracto máximo de 18,82°P. La combinación óptima para FAN fue de 2.434,5 U, 0 BGU y 100 mg para Safrari no malteado, dando un FAN máximo de 144,77 mg/L, mientras que para Safrari malteado fue de 2.191,5 U, 0 BGU y 100 mg, para un FAN máximo de 196,73 mg/L.

Palabras claves: enzimas comerciales de maceración, extracto, amino nitrógeno libre (FAN), optimización, sorgo *Safrari*.

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#### Introduction

Sorghum in its malted form or as adjuncts has become a potential brewing cereal, particularly in the tropics where barley is not commonly grown^{6,14,24,25,26,30,31,32,33,34}. Its low contents of potential mashing enzymes due to their poor development during malting has often triggered the use of commercial mashing enzymes as supplements to achieve higher yields in extracts and other important wort specifications in beer brewing^{1,2,3,4,5,7,8,16,17,19,20,28,29,37,38}. The inclusion of commercial mashing enzymes when mashing sorghum malts and adjuncts is important^{9,10,11,14,15,22,23,24,27}, but the amount of enzymes needed for optimal mashing may vary from one sorghum cultivar to the next. It is also not clear whether the use of commercial mashing enzymes for optimal mashing of malted sorghum is indispensable. In this work, the influence of three commercial mashing enzymes (Hitempase 2XL, Bioglucanase TX and

Table 1. Characteristics of co	mmercial mashing	r enzyme preparations
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Commercial mashing enzyme	Organism of origin	Activity	Description	Temperature optima	pH optima	Recommended application level in adjuncts	Form
Hitempase 2XL ^a Bioglucanase TX ^a	Bacillus licheniformis Trichodermareesei	4,416.29 ± 19.34 U/ml 750 BGU/ml	α-amylase β-glucanase	60–95°C 60°C	4–8 4.5–6.5	60 U/g 0.01 and 0.025% (v/w)	Solution Solution
Brewers protease ^b	Bacillus amyloliquefaciens	1,842.2 ± 1.8 mg FAN/min/ml	Protease	45–50°C (denatured at 85°C)	6.5–7.5	0.4–2 g/kg	Solution

^a Hitempase 2XL and Bioglucanase TX were obtained from Kerry Bioscience; Kilnagleary, Carrigaline, Co. Cork, Ireland.

^b Brewers protease was obtained from DSM Food Specialities, Cedex France.

Brewers protease) was modelled and optimised for yields of extracts and free amino nitrogen during mashing of unmalted and malted *Safrari* sorghum cultivar using the response surface methodology (RSM).

# **Materials and Methods**

#### Enzymes

The characteristics of the commercial enzymes used (Hitempase 2XL, a thermo stable  $\alpha$ -amylase from *Baccillus licheniformis;* Bioglucanase TX, an enzymatic composition of  $\beta$ -glucanase and hemicellulases from *Trichodermareesel;* and Brewers protease from *Bacillus amyloliquefaciens*) are presented in Table 1.

#### Sorghum Cultivar

The *Safrari* sorghum cultivar was obtained from the Institute of Research and Agronomic Development (IRAD) Maroua, Cameroon.

#### Modelling

Modelling was carried out as previously described¹⁵. A Doehlert matrix design with three factors representing Hitempase ( $X_1$ ), Bioglucanase ( $X_2$ ) and Brewers protease ( $X_3$ ) at ranges of 0–3,000 U, 0–937.5 BGU and 0–100 mg, respectively, was used. The transformed matrix of coded variables to an experimental matrix and desired responses (extract and free amino nitrogen) are shown in Table 2. The coefficients of the models and the models were obtained using the Systat version 12 software (Systat Software, Inc., San Jose, USA). This software also gives a statistical analysis on the model. Lastly, the curves were plotted using Sigmaplot version 11 build 11.0.0.77 software (WPCubed, GmbH, Germany).

#### Validation of Models

The models were validated using two procedures. The first consited of coupling the method earlier described¹⁵ to the absolute average deviation (AAD) method¹³. The second procedure consisted of applying the method described^{12,36}.

#### Malting

About 1 kg of *Safrari* sorghum cultivar grains were washed three times using 3 litres of distilled water to remove dirt and other foreign bodies. The grains were steeped in 3 litres of distilled water for 48 h at room temperature ( $\approx 25^{\circ}$ C) with three changes of water at intervals of 12 h before steep out. Germination was carried out for 4 days in a Heraeus type oven (D-63450 Hanau, Germany) at a temperature of 25°C with water sprinkled on the grains on daily basis. The malt was then air dried at 40°C for 4 days using a CKA 2000 AUF-type dryer (Ngaoundere, Cameroon). The malt was rubbed-off of its rootlets and stored until further use.

## Mashing

Two hundred and fifty ml of distilled water were put into a 600 ml beaker and 50 g of sorghum (malted or unmalted) flour  $(\emptyset < 1 \text{ mm})$  added with continuous stirring until a homogenous mixture was obtained. This mixture was incubated at 45°C for 1 h in a water bath with intermittent stirring at intervals of 5 min. The mix was allowed to decant and 50 ml of the supernatant withdrawn and kept aside. The temperature of the mash was then raised to boiling so as to gelatinise sorghum starch during 40 min with intermittent stirring at intervals of 5 min before cooling to 65°C. The 50 ml of supernatant, to which commercial enzymes are added according to the Doehlert matrix design of three factors, were added to the mash and mashing continued for 1 h and 30 min with intermittent stirring at intervals of 10 min. The mash was cooled and filtered at 25°C for 1 h and 30 min using Whatmann paper no. 42

#### **Determination of Extract**

Extract was determined as described by analytica-EBC¹⁸.

#### **Determination of Free Amino Nitrogen**

Free amino nitrogen (FAN) was determined using the Ninhydrin method as described by analytica-EBC¹⁸.

#### **Optimisation of Models**

Models were optimised as previously described¹⁵. The optimal zone of intersection of the curves was highlighted.

### **Results and Discussion**

Optimisation of the action of mashing enzymes on the two key mashing parameters, extract and FAN, was carried out by modelling the experimental design required for laboratory purposes. Table 2 shows the results obtained for extracts and free amino nitrogen (FAN) after mashing unmalted and malted *Safrari* using the commercial mashing enzymes Hitempase ( $\alpha$ amylase), Bioglucanase ( $\beta$ -glucanase) and Brewers protease (protease).

The mathematical models obtained for extracts after mashing unmalted and malted *Safrari* were as follows, respectively:

$$\begin{split} Y_{\text{SafEX}}(X_1, X_2, X_3) &= 14.62 + 5.478X_1 + \\ 1.226X_2 - 0.881X_3 + 0.527X_1X_2 + 0.589X_1X_3 - \\ 2.008X_2X_3 - 6.569X_1^2 - 0.935X_2^2 - 0.496X_3^2 \\ Y_{\text{SafMEX}}(X_1, X_2, X_3) &= 16.566 + 1.076X_1 - \\ 1.377X_2 - 0.469X_3 + 4.774X_1X_2 + 2.338X_1X_3 - \\ 1.326X_2X_3 - 0.901X_1^2 - 3.117X_2^2 - 0.180X_3^2 \end{split}$$

With:  $Y_{SafEX}(X_1, X_2, X_3)$  representing the mathematical model for unmalted *Safrari;*  $Y_{SafMEX}(X_1, X_2, X_3)$ , the model for

C	Coded valu	es		ransform rimental v							Saj	frari					
								Unn	nalted			Malted					
Hit	Bio	Brew prot	Hit (U)	Bio (BGU)	Brew prot (mg)		Extract			FAN			Extract			FAN	
X ₁	X ₂	X ₃	X ₁	(DGC) X ₂	X ₃	Exp ^a	Theo ^b	Res ^c	Exp	Theo	Res	Exp	Theo	Res	Exp	Theo	Res
0.000	0.000	0.000	1,500	468.75	50	14.81	14.62	0.19	95	91.78	3.22	16.43	16.57	-0.14	140	136.97	3.03
1.000	0.000	0.000	3,000	468.75	50	14.21	13.53	0.68	88	90.76	-2.76	16.64	16.74	-0.10	132	131.32	0.68
0.500	0.866	0.000	2,250	937.5	50	15.89	16.31	-0.42	110	97.91	12.09	15.60	15.42	0.18	165	151.42	13.58
-0.500	-0.866	0.000	750	0.00	50	8.75	8.70	0.05	100	106.23	-6.23	16.02	16.72	-0.70	149	154.06	-5.06
0.500	-0.866	0.000	2,250	0.00	50	13.40	13.73	-0.33	135.60	128.72	6.88	14.56	13.67	0.89	189	182.34	6.66
-0.500	0.866	0.000	750	937.5	50	10.73	10.37	0.36	63	70.41	-7.41	9.36	10.21	-0.85	94.50	101.93	-7.43
0.500	0.289	0.816	2,250	615.18	100	14.29	14.79	-0.50	118.33	124.00	-5.67	16.64	17.05	-0.41	177.50	186.44	-8.94
-0.500	-0.289	-0.816	750	312.32	0.0	9.98	10.04	-0.06	66.33	51.87	14.46	17.16	17.53	-0.37	99.50	77.78	21.72
0.500	-0.289	-0.816	2,250	312.32	0.0	14.55	14.88	-0.33	61.67	66.12	-4.46	14.56	15.32	-0.76	92.50	100.32	-7.82
0.000	0.577	-0.816	1,500	781.07	0.0	16.44	16.35	0.09	38.33	43.54	-5.21	16.33	15.62	0.71	57.50	64.42	-6.92
-0.500	0.289	0.816	750	615.18	100	9.06	8.68	0.38	91.92	88.27	3.65	13.52	12.69	0.83	137.88	131.22	6.66
0.000	-0.577	0.816	1,500	156.43	100	13.61	13.50	0.11	126	123.99	2.01	16.02	16.44	-0.42	178	175.72	2.28
0.000	0.000	0.000	1,500	468.75	50	14.50	14.62	-0.12	85.33	91.78	-6.45	16.64	16.57	0.07	128	136.97	-8.97
-1.000	0.000	0.000	0	468.75	50	1.80	2.57	-0.77	45	40.78	4.22	14.56	14.59	-0.03	55	53.55	1.45
-1.000	-0.866	-0.816	0	0.0	0.0	1.02	0.72	0.30	54	58.80	-4.80	19.24	18.81	0.43	81	87.97	-6.97
0.000	0.000	0.000	1,500	468.75	50	14.60	14.62	-0.02	91.67	91.78	-0.11	16.64	16.57	0.07	137.50	136.97	0.53
0.000	0.000	0.000	1,500	468.75	50	15.00	14.62	0.38	88.33	91.78	-3.45	17.16	16.57	0.59	132.50	136.97	-4.47

^a Experimental result values.

^b Theoretical values (values coming from mathematical models).

^c Residue.

Table 3. Estimation of regression coefficients for the extracts of umalted and malted Safrari.

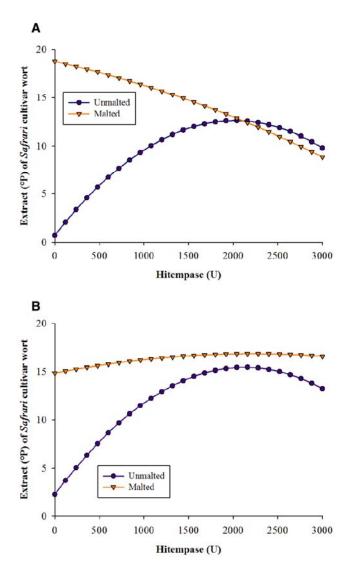
	Coeffi	cient	Std. de	viation	t-stati	istics	P-value	
Effects	Unmalted	Malted	Unmalted	Malted	Unmalted	Malted	Unmalted	Malted
CONSTANT	14.62	16.566	0.271	0.396	54.039	41.882	0.000	0.000
X ₁	5.476	1.076	0.278	0.407	19.683	2.644	0.000	0.033
X ₂	1.226	-1.377	0.243	0.355	4.374	-3.361	0.003	0.012
X3	-0.881	-0.469	0.229	0.335	-3.136	-1.141	0.016	0.291
$X_{1}^{2}$	-6.569	-0.901	0.477	0.698	-13.768	-1.291	0.000	0.238
$X_{2}^{2}$	-0.935	-3.117	0.347	0.508	-2.020	-4.604	0.083	0.002
$X_{3}^{2}$	-0.496	-0.18	0.300	0.438	-1.103	-0.274	0.307	0.792
X1*X2	0.527	4.774	0.545	0.796	0.839	5.192	0.429	0.001
$X_2 * X_3$	-2.008	-1.326	0.479	0.701	-2.960	-1.336	0.002	0.223
X1*X3	0.589	2.338	0.568	0.83	0.846	2.299	0.425	0.055

malted *Safrari;*  $X_1$ , Hitempase;  $X_2$ , Bioglucanase and  $X_3$  Brewers protease.

These mathematical models are polynomials having several variables with correlation coefficients  $R^2 = 0.993$  for unmalted *Safrari* and  $R^2 = 0.930$  for malted *Safrari*. These coefficients, coupled to AAD values of 0.026 and 0.031 for unmalted and malted *Safrari*, respectively, allowed for the validation of the models for the wort extract yields. In addition, a bias factor of 1, coupled to exactitude factors of 1.06 and 1.03 for both unmalted and malted *Safrari*, respectively, also allowed for validation of the models according to the method described^{12,36}. The factors of the models were linear or of first degree (X₁, X₂ and X₃), quadratic or of the second degree (X₁², X₂² and X₃²) and of interacting form (X₁X₂, X₁X₃ and X₂X₃). They were statistically considered significant or not if the probability (P) of increasing yields of extracts was ≤0.05 or ≥0.05, respectively (Table 3).

The impact of Hitempase as sole mashing enzyme on the yields of extract of unmalted and malted *Safrari* is shown in Figure 1A. Extract yield increased with increasing concentration of enzyme for unmalted *Safrari* mash to attain a maximum level ( $\approx$ 13°P) at about 2,000 U, followed by a slight and steady decrease thereafter. It was already maximal ( $\approx$ 18.5°P) even in absence of Hitempase for malted *Safrari* mash. Indeed,

increasing amounts in enzyme concentration instead induced a steady decrease of extract yield for malted Safrari mash. However, when the mathematical models were applied to predict the impact of supplements of Bioglucanase and Brewers protease at concentrations of 400 BGU and 60 mg, respectively, it was observed that the profile of extract remained fairly constant with increasing amounts of Hitempase for malted Safrari mash (Fig. 1B). Starch is indeed the main macromolecule of cereals and the main substrate of  $\alpha$ -amylase. It is therefore expected that this enzyme contributes the greatest amount of soluble materials that could be found in resulting worts due to its action on starch^{15,35}. Figures 1A and B also showed that supplements of Hitempase in malted Safrari mash was of no importance. The steady decrease in yields of extract of malted Safrari mash could be attributed to Maillard reactions between soluble nitrogenous compounds and reducing sugars earlier produced during malting (Fig. 1A). These results also indicated that efficient mashing was obtainable in the presence of the key mashing enzymes developed during malting (Fig. 1A). From the mathematical model, it was shown that in its linear form  $(X_1)$ , Hitempase contributed 29% and 7% of extract yields for unmalted and malted Safrari, respectively (Figs. 2A and B). Statistical analyses also showed that this contribution was significant (P = 0.000 and 0.033 for unmalted and malted



**Figure 1. A,** Effect of concentration of Hitempase ( $\alpha$ -amylase) as sole mashing enzyme (concentrations of Bioglucanase and Brewers protease set at 0) on yield of wort extract of sorghum cultivar *Safrari*. **B**, Effect of concentration of Hitempase ( $\alpha$ -amylase) in the presence of fixed concentrations of Bioglucanase (400 BGU) and Brewers protease (60 mg) on yield of wort extract of sorghum cultivar *Safrari*.

Safrari, respectively) (Table 3). In its quadratic form  $(X_1^2)$ , Hitempase remained statistically significant for mashing unmalted Safrari, but not for malted Safrari (P = 0.000 and 0.238, respectively). This confirmed the above biological observation according to which supplements of this enzyme in malted mashes of Safrari were of no importance. Its contribution to increasing extract yields in its quadratic form  $(X_1^2)$ (excess of  $\alpha$ -amylase in principle) is indeed 35% and 6% for unmalted and malted Safrari, respectively (Figs. 2A and B).

Figure 3A shows the effect of mashing unmalted and malted *Safrari* using Bioglucanase as sole mashing enzyme on yields of extract. There was a slight and constant increase in yields of extract as enzyme concentration increased for unmalted *Safrari* mash. The yields of extract for malted *Safrari* mash was maximal even in the absence of Bioglucanase and steadily decreased with an increase in enzyme concentration. The small yields of extract due to Bioglucanase action in unmalted *Sa* 

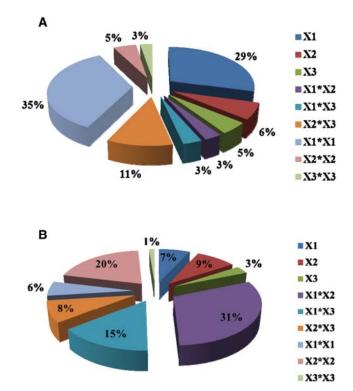
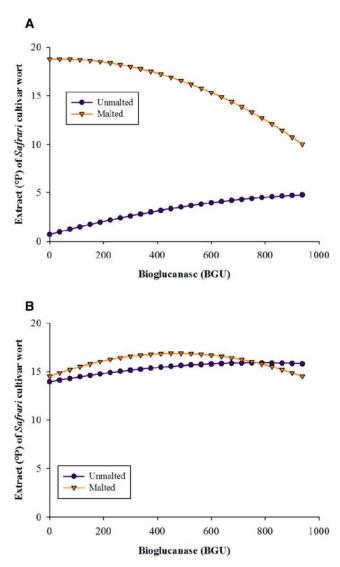


Figure 2. A, Contribution to yield of wort extract (°P) of each factor in its linear, quadratic and interaction (combined) forms for unmalted sorghum cultivar *Safrari*. B, Contribution to yield of wort extract (°P) of each factor in its linear, quadratic and interaction (combined) forms for malted sorghum cultivar *Safrari*.

frari mash can be attributed to its ability to hydrolyse β-glucans into glucose and other soluble carbohydrates. Bioglucanase was therefore not a backbone enzyme for extract production during mashing. A similar application of the mathematical models as carried out above for Hitempase's action, using 60 mg of Brewers protease and 2,000 U of Hitempase, predicted that supplements of these two key mashing enzymes could provide similar results in extract yields for both unmalted and malted Safrari mashes (Fig. 3B). Hitempase once more demonstrated that it was the backbone enzyme contributing to most of the extract yields. Figure 3B indeed showed that in the absence of Bioglucanase, but in the presence of Hitempase, yields of extract became comparable for both unmalted and malted Safrari mashes. These observations were all statistically confirmed. Indeed, in its linear form  $(X_2)$ , Bioglucanase's action was significant (P = 0.003 and 0.012 for unmalted and malted Safrari, respectively) (Table 3). Figures 2A and B showed that this enzyme contributed for 6% and 9% of extract yield for unmalted and malted Safrari, respectively. In its quadratic form  $(X_2^2)$  (excess of enzyme in principle), Bioglucanase contributed for 5% and 20% of extract yield for unmalted and malted Safrari, respectively (Figs. 2A and B). These contributions were not statistically significant for unmalted Safrari mash, but were for malted Safrari mash (P = 0.083 and 0.002) (Table 3). Once more, the observed 1:4 ratio difference of extract yield between unmalted Safrari mashes as to malted Safrari mashes could be attributed to the impact of enzymes developed during malting and not to Bioglucanase supplements as such.



**Figure 3. A,** Effect of concentration of Bioglucanase ( $\beta$ -glucanase) as sole mashing enzyme (concentrations of Hitempase and Brewers protease set at 0) on yield of wort extract of sorghum cultivar *Safrari*. **B**, Effect of concentration of Bioglucanase ( $\beta$ -glucanase) in the presence of fixed concentrations of Hitempase (2,000 U) and Brewers protease (60 mg) on yield of wort extract of sorghum cultivar *Safrari*.

The effect of mashing unmalted and malted Safrari on yields of extract using as sole mashing enzyme, Brewers protease, is shown in Figure 4A. This enzyme contributed little or no extracts in mashes of unmalted Safrari as compared to Hitempase and even Bioglucanase. Once more, the maximal extract observed when mashing with malted Safrari even in the absence of Brewers protease supplements is due to the virtues known to the malting process as explained earlier. These results confirmed earlier observations¹⁵. The steady and slight decrease of extract yield with increase of Brewers protease for the malted Safrari mash could once more be attributed to reactions between nitrogenous functions and some of the soluble sugars resulting from the malting process. The mathematical model was once more used to predict the yield in extract as carried out above for Hitempase and Bioglucanase actions. Thus, using 2,000 U of Hitempase and 400 BGU of Bioglucanase with increasing amounts of Brewers protease,

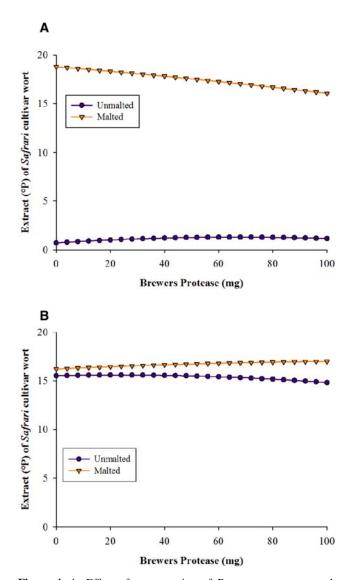


Figure 4. A, Effect of concentration of Brewers protease as sole mashing enzyme (concentrations of Bioglucanase and Hitempase set at 0) on yield of wort extract of sorghum cultivar *Safrari*. B, Effect of concentration of Brewers protease in the presence of fixed concentrations of Hitempase (2,000 U) and Bioglucanase (400 BGU) on yield of wort extract of sorghum cultivar *Safrari*.

the results once more showed that adding these mashing enzymes could provide similar results in extract yields for both unmalted and malted *Safrari* mashes (Fig. 4B). These observations were statistically confirmed using the mathematical model. In its first degree form (X₃), the impact of Brewers protease was significant for unmalted *Safrari* mash but not for malted *Safrari* mash (P = 0.016 and 0.291, respectively) (Table 3). Its contribution to extract yields was barely 5% and 3%, respectively, for both mash types (Figs. 2A and B). The impact of the enzyme in its quadratic form (X₃²) was not significant for both mashes (P = 0.307 and 0.792, respectively) (Table 3). Its contribution to extract yield was 3% and 1%, respectively (Figs. 2A and B).

The models were further exploited to predict the impacts of the interactions  $(X_1X_2, X_1X_3 \text{ and } X_2X_3)$  of these enzymes on yields of extract. The results are shown in Figures 2A and B. Globally, they were statistically not significant for unmalted

		Sum se	quare	Mean s	Mean square		lue	P-value	
Source	DDL	Unmalted	Malted	Unmalted	Malted	Unmalted	Malted	Unmalted	Malted
Regression	9	333.038	65.429	37.004	7.27	113.241	10.408	0.000	0.003
Linear	3	240.289	18.625	80.096	6.208	245.112	8.888	0.000	0.009
Quadratic	3	89.659	7.736	29.886	2.579	91.459	3.692	0.000	0.070
Interactions	3	3.090	39.069	1.030	13.023	3.152	18.645	0.095	0.001
Residual error	7	2.287	4.889	0.327	0.698				
Total error	16	335.326	70.319						

Table 4. ANOVA for the extracts of umalted and malted Safrari.

Table 5. ANOVA for comparing extracts of unmalted and malted *Safrari* worts.

Source	DF	Sum of squares	Mean of squares	F-value	P-value
Inter-groups	1	130.693	130.693	9.21	0.004
Intra-groups	32	454.199	14.193		
Total	33	584.892			

Safrari mashes (P = 0.095), but were for malted Safrari mashes (P = 0.001) (Table 4). The interaction  $X_1X_2$  (Hitempase/Bioglucanase) had no significant impact on unmalted Safrari mash, but did for malted Safrari (P = 0.429 and 0.001, respectively) (Table 3). It contributed merely 3% of extract for unmalted Safrari mash but up to 31% for malted Safrari mash (Figs. 2A and B). However, it is important to underline that this significant contribution could be attributed to the intrinsic virtues that malting offers when mashing with malted Safrari, and not to the Hitempase/Bioglucanase interaction as such. Though known to be the backbone starch hydrolysing enzyme, the action of Hitempase is best exploited when the cell walls of cereal grains are broken down by  $\beta$ -glucanases, hemicellulases and cellulases to liberate starch granules. This sequence of events during malting was confirmed by the mathematical models above. The interaction X1X3, corresponding to the couple Hitempase/Brewers Protease, also had no significant impact on both the extract yields of unmalted Safrari and malted Safrari mashes (P = 0.425 and 0.055, respectively) (Table 3). Its contribution to extract yields was 3% and 15%, respectively (Figs. 2A and B). This result was once more in conformity with the biological sequence occurring during malting. Efficient starch hydrolysis by  $\alpha$ -amylase indeed occurs only after the breakdown of cell walls by  $\beta$ -glucanase, followed by liberation of starch granules due to proteolysis of the protein matrix enrobing them. The interaction Bioglucanase/Brewers protease  $(X_2X_3)$  had a significant impact on extract yields for unmalted Safrari mash but not for malted Safrari mashes (P = 0.002 and 0.223, respectively) (Table 3). Its contribution to extract yields was 11% and 8%, respectively, for both mash types (Figs. 2A and B). These low contributions by the couple (Bioglucanase/Brewers protease) were expected, as the two enzymes only play a supporting role in starch hydrolysis during mashing¹⁵. Table 5 statistically confirmed the observation that Safrari sorghum malted type samples were more potential mashing materials than unmalted Safrari sorghum adjuncts to which commercial enzymes are supplemented for the production of worts of higher extract yields (P = 0.004).

The mathematical models obtained for FAN for mashing unmalted and malted *Safrari* were as follows, respectively:

$$\begin{split} Y_{SafAAL}(X_1, X_2, X_3) &= 91,781 + 24,992X_1 - \\ 19,234X_2 + 35,693X_3 + 2,892X_1X_2 + 12,142X_1X_3 + \\ 14,822X_2X_3 - 26,014X_1^2 + 20,723X_2^2 - 11,921X_3^2 \end{split}$$

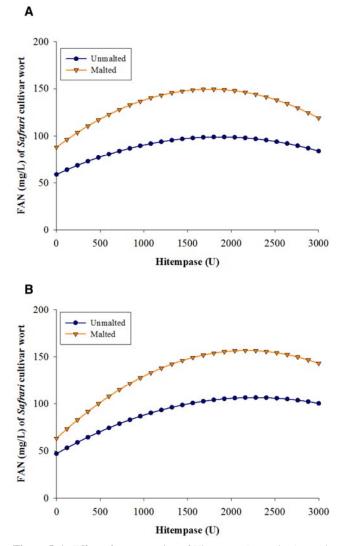
 $\begin{array}{l} Y_{SafMAAL}(X_1,X_2,X_3) = 136,969 + 38,884X_1 - \\ 23,975X_2 + 51,247X_3 + 12,243X_1X_2 + 15,682X_1X_3 + \\ 31,393X_2X_3 - 44,536X_1{}^2 + 28,806X_2{}^2 - 17,579X_3{}^2 \end{array}$ 

With:  $Y_{SafAAL}$  (X₁, X₂, X₃) representing the mathematical model for unmalted Safrari; Y_{SafMAAL} (X₁, X₂, X₃) for malted Safrari; X₁, Hitempase; X₂, Bioglucanase and X₃, Brewers protease. These mathematical models were once more polynomials having several variables with correlation coefficients  $R^2$ = 0.944 for unmalted *Safrari* and  $R^2$  = 0.955 for malted *Sa*frari. These coefficients, coupled to AAD values of 0.073 and 0.060 for unmalted and malted Safrari, respectively, allowed for the validation of the models for assessment of the wort free amino nitrogen content. In addition, a bias factor of 1, coupled to exactitude factors of 1.08 and 1.06 for both unmalted and malted Safrari, respectively, also allowed for validation of the models according to the method described^{12,36}. The factors of the models were once more linear or of first degree  $(X_1, X_2 and$  $X_3$ ), quadratic or of the second degree  $(X_1^2, X_2^2 \text{ and } X_3^2)$  or of interaction form  $(X_1X_2, X_1X_3 \text{ and } X_2X_3)$ . They were statistically considered significant or not if the probability (P) of increasing yields of FAN was ≤0.05 or ≥0.05, respectively (Table 6).

The impact of Hitempase as sole mashing enzyme on yields of FAN for unmalted and malted Safrari is shown in Figure 5A. FAN content of wort gradually increased with increasing enzyme concentration to reach maxima of about 85 and 150 mg/L at about 1,750 U for both unmalted and malted Safrari mashes, respectively. This was followed by slight decreases for the unmalted Safrari mash as compared with malted Safrari mash. The curves showed that either for unmalted Safrari or malted Safrari mash types, the FAN content at origin was well above zero (60 mg/L and 90 mg/L, respectively). This suggests that the milling operation was at the basis of a good amount of the FAN present at the beginning of mashing. Although Hitempase is not a protein hydrolyzing enzyme, it exposes more free amino nitrogen functions upon acting on starch granules. This could explain the slight increase in FAN observed with increase in enzyme concentration. The higher FAN content for malted Safrari mash as compared to unmalted Safrari mash is once more to be attributed to the natural virtues that the grains incur during the malting process. Use of the models to predict the profile of FAN content of worts if the mashing enzymes Bioglucanase (at 400 BGU) and Brewers protease (at 60 mg) were coupled to Hitempase's action showed no remarkable difference. A balance between the amounts of enzymes needed to obtain maximal FAN contents and its disappearance in the medium due to reactions with sugars needs to be known to clearly understand this indifference. The models also showed that Hitempase  $(X_1)$ , in its first degree form, contributed 15% of the FAN content of both the unmalted and malted Safrari mashes (Figs. 6A and B). This contribution was statistically significant for the two mash types (P = 0.001 and 0.000), reTable 6. Estimation of regression coefficients for free amino nitrogen of umalted and malted Safrari.

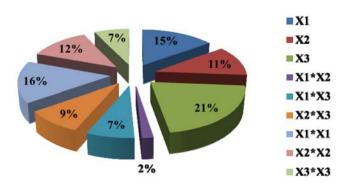
Effects	Coeff	icient	Std. de	viation	T-stat	istics	P-value	
	Unmalted	Malted	Unmalted	Malted	Unmalted	Malted	Unmalted	Malted
CONSTANT	91.781	136.969	4.75	6.148	19.321	22.278	0.000	0.000
X ₁	24.992	38.884	4.886	6.325	5.115	6.148	0.001	0.000
$X_2$	-19.234	-23.975	4.261	5.515	-3.909	-3.765	0.006	0.007
$\overline{X_3}$	35.693	51.247	4.026	5.211	7.234	8.025	0.000	0.000
$X_{1}^{2}$	-26.014	-44.536	8.377	10.843	-3.105	-4.107	0.017	0.005
$X_{2}^{2}$	20.723	28.806	6.097	7.891	2.549	2.738	0.038	0.029
$\tilde{X_3^2}$	-11.921	-17.579	5.259	6.807	-1.509	-1.72	0.175	0.129
X1*X2	2.892	12.243	9.563	12.378	0.262	0.857	0.801	0.420
$X_{2}^{*} X_{3}^{-}$	14.822	31.393	8.418	10.895	1.244	2.036	0.253	0.081
X1*X3	12.142	15.682	9.966	12.899	0.994	0.992	0.353	0.354

Α



**Figure 5. A,** Effect of concentration of Hitempase ( $\alpha$ -amylase) as sole mashing enzyme (concentrations of Bioglucanase and Brewers protease set at 0) on yield of wort free amino nitrogen of sorghum cultivar *Safrari.* **B,** Effect of concentration of Hitempase ( $\alpha$ -amylase) in the presence of fixed concentrations of Bioglucanase (400 BGU) and Brewers protease (60 mg) on yield of wort free amino nitrogen of sorghum cultivar *Safrari.* 

spectively (Table 6). Similarly, in its quadratic form  $(X_1^2)$ , Hitempase's effect remained significant for the two mash types (P = 0.017 and 0.005, respectively) (Table 6). Its contribution in this form was 16% and 17%, respectively (Figures 6A and



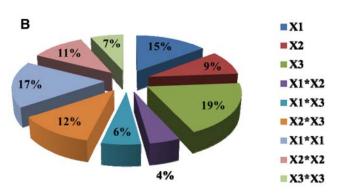
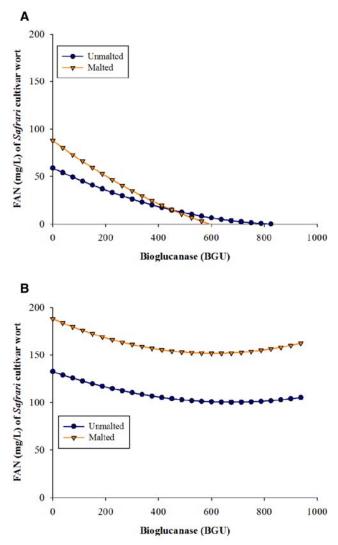


Figure 6. A, Contribution to yield of wort free amino nitrogen (FAN) of each factor in its linear, quadratic and interaction (combined) forms for unmalted sorghum cultivar *Safrari*. B, Contribution to yield of wort free amino nitrogen (FAN) of each factor in its linear, quadratic and interaction (combined) forms for malted sorghum cultivar *Safrari*.

B). The small difference in FAN content contribution  $(\pm 2\%)$  between its linear and quadratic (enzyme supplemented in excess) forms suggests that supplementing the enzyme in excess is of no technological use.

Figure 7A shows the effect of Bioglucanase on the FAN content as sole mashing enzyme in unmalted and malted *Sa-frari*. This content decreased to zero with increasing enzyme concentration in the mashes, indicating that Bioglucanase is not a protein hydrolysing enzyme. Mashing with the enzyme rather seemed to be of no use as the FAN was exposed to reactions with reducing sugars. Upon using the model to predict the amounts of FAN if mashed in the presence of Hitempase (at 2,000 U) and Brewers protease (at 60 mg), it was observed



**Figure 7. A,** Effect of concentration of Bioglucanase ( $\beta$ -glucanase) as sole mashing enzyme (concentrations of Hitempase and Brewer's Protease set at 0) on yield of wort free amino nitrogen of sorghum cultivar *Safrari.* **B,** Effect of concentration of Bioglucanase ( $\beta$ -glucanase) in the presence of fixed concentrations of Hitempase (2000 U) and Brewers protease (60 mg) on yield of wort free amino nitrogen of sorghum cultivar *Safrari.* 

that the decrease of FAN levels to zero could be stopped and controlled (Fig. 7B). This confirms the need to have all mashing enzymes present in appropriate proportions during mashing to permit obtain sustainable amounts of FAN contents. According to the models, Bioglucanase ( $X_2$ ), in its linear form, contributed 11% of the FAN content of the unmalted *Safrari* mash and 9% of the malted *Safrari* mash (Figures 6A and B). This contribution was statistically significant for the two mash types (P = 0.006 and 0.007, respectively) (Table 6). Similarly, in its quadratic form ( $X_2^2$ ), the effect of the enzyme remained significant for the two mash types (P = 0.038 and 0.029, respectively) (Table 6). Its contribution in this form was 12% and 11%, respectively (Figures 6A and B).

The effect of Brewers protease as sole mashing enzyme on FAN content for unmalted and malted *Safrari* is presented in Figure 8A. Free amino nitrogen content increased very slightly from 60 mg/L to 75 mg/L for unmalted *Safrari* mash and from 90 mg/L to 110 mg/L for malted *Safrari* mash with an increase

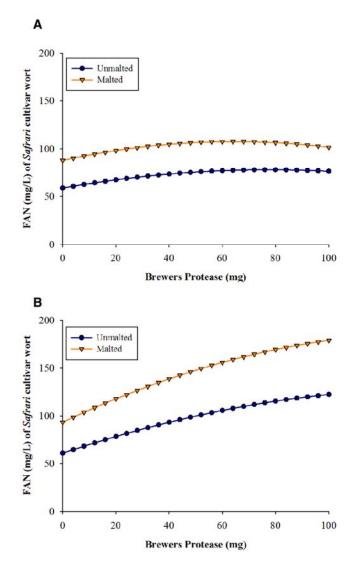


Figure 8. A, Effect of concentration of Brewers protease as sole mashing enzyme (concentrations of Bioglucanase and Hitempase set at 0) on yield of wort free amino nitrogen of sorghum cultivar *Safrari*. B, Effect of concentration of Brewers protease in the presence of fixed concentrations of Hitempase (2000 U) and Bioglucanase (400 BGU) on yield of wort free amino nitrogen of sorghum cultivar *Safrari*.

in enzyme concentration. This level was maintained constant from about 60 mg of enzyme input thereof. The model predicted that coupling Hitempase (at 2,000 U) and Bioglucanase (at 400 BGU) to the action of Brewers protease induced a steady increase in FAN content (Fig. 8B). This once more indicates the need of all mashing enzymes in order to obtain higher FAN yields. The additional FAN content observed for malted Safrari mash as compared to unmalted Safrari mash could once more be attributed to the natural and additional virtues of the malting process. The mathematical models statistically showed that the action of Brewers protease was in its linear form significant for both mash types (P = 0.000 for both) (Table 6). Its contribution to FAN content in this form  $(X_3)$  was 21% and 19% for both the unmalted and malted Safrari mashes, respectively (Figs. 6A and B). The effect of the enzyme in its quadratic form  $(X_3^2)$  is however not significant for the two mash types (P = 0.175 and 0.129, respectively) (Table 6). Its contribution decreased to as low as 7% for both

		Sum of squares		Mean of	squares	F-va	lue	P-value	
Source	DF	Unmalted	Malted	Unmalted	Malted	Unmalted	Malted	Unmalted	Malted
Regression	9	11,814.795	25,319.50	1,312.755	2,813.278	13.032	16.671	0.001	0.001
Linear	3	9,794.014	20,169.46	3,264.671	6,723.156	32.408	39.841	0.000	0.000
Quadratic	3	1,577.067	3,427.379	525.689	1,142.46	5.218	6.77	0.033	0.018
Interactions	3	443.714	1,722.655	147.905	574.218	1.468	3.403	0.303	0.083
Residual error	7	705.153	1,181.253	100.736	168.75				
Total error	16	12,519.94	26,500.75						

Table 7. ANOVA for free amino nitrogen of umalted and malted Safrari.

 
 Table 8. ANOVA for comparing free amino nitrogen of unmalted and malted Safrari worts.

Source	DF	Sum of squares	Mean of squares	F-value	P-value
Inter-groups	1	13,916.3	13,916.3	11.41	0.001
Intra-groups	32	39,020.7	1,219.4		
Total	33	52,937			

mash types (Figs. 6A and B). Excess doses of this enzyme during mashing therefore appear to be of no technological importance.

The global action of these enzymes in their interaction or coupled forms  $(X_1X_2, X_1X_3 \text{ and } X_2X_3)$  on the FAN content was statistically not significant (P = 0.303 for unmalted Safrari and P = 0.083 for malted *Safrari*) (Table 7). Their contributions of FAN content are shown in Figures 6A and B. The effect of the X1X2 (Hitempase/Bioglucanase) interaction was not significant for both mash types (P = 0.801 for unmalted Safrari and P =0.420 for malted Safrari) (Table 6). Its contribution of FAN content in both mash types was 2% and 4%, respectively. Similarly, the action of the couple X1X3 (Hitempase/Brewers protease) was also not significant for both mash types (P = 0.353for unmalted Safrari and P = 0.354 for malted Safrari) (Table 6). Its contribution of FAN content in both mash types was 7% and 6%, respectively (Figs. 6A and B). Finally, for the couple  $X_2X_3$  (Bioglucanase/Brewers protease) (P = 0.253 for unmalted Safrari and P = 0.081 for malted Safrari) (Table 6), the contribution of FAN content in both mash types was 9% and 12%, respectively (Figs. 6A and B). Once more, Table 8 statistically confirmed the observation that malted Safrari sorghum type samples were more potential mashing materials than unmalted Safrari sorghum adjuncts to which commercial enzymes were supplemented for production of worts of higher FAN content (P = 0.001).

The results obtained for the action of the enzymes on extract and FAN yields after mashing on the basis of the models were optimised to define satisfactory domains of compromise for the mashing enzymes. These domains were obtained for the two key brewing parameters by fixing the wort conditions at extract  $\geq 12^{\circ}P$  and FAN  $\geq 80$  mg/L. The theoretical optimal combination of enzyme action for unmalted Safrari gave the following triplet of coded variables for extract: 0.399, 0.866 and -0.816 (2,098.5 U, 937.5 BGU and 0 mg real variables) for Hitempase, Bioglucanase and Brewers protease, respectively. This triplet allowed for a maximal extract of 18°P. The theoretical optimal enzyme combination for maximal contribution of FAN for unmalted Safrari mash gave as triplet of coded variables: 0.623, -0.866 and 0.816 (2,434.5 U, 0 BGU and 100 mg real variables). This triplet allowed for maximal free amino nitrogen of 144.77 mg/L (Fig. 9A). This triplet for maximal extract yields for malted Safrari was -1, -0.813 and -0.816 (0 U, 28.68 BGU and 0 mg real variables) for Hitempase, Bioglu-

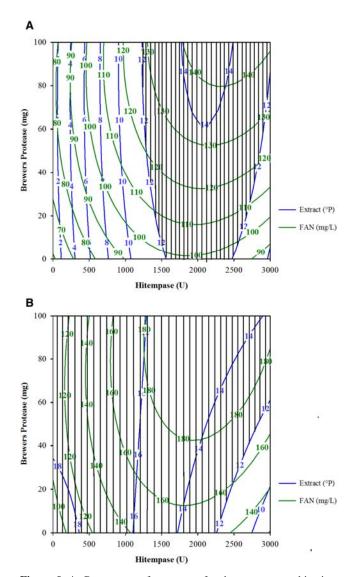


Figure 9. A, Response surface curves for the enzyme combinations providing for optimal yields in extract and free amino nitrogen for unmalted sorghum cultivar *Safrari*. B, Response surface curves for the enzyme combinations providing for optimal yields in extract and free amino nitrogen for malted sorghum cultivar *Safrari*.

canase and Brewers protease, respectively. It allowed for a maximal extract of 18.82°P. The optimal enzyme combinations were thus different for the two mash types, but both gave comparable yields of extracts. The triplet for maximal FAN content for malted *Safrari* was 0.461, -0.866 and 0.816 (2,191.5 U, 0 BGU and 100 mg real variables). It allowed for maximal FAN of 196.73 mg/L (Fig. 9B). These results once more confirmed

that commercial enzyme supplements for mashing malted *Sa-frari* are not indispensable for obtaining maximal yields of extract as compared to unmalted *Safrari*. The enzymes supplements however proved useful for obtaining maximal FAN.

### Conclusions

The effects of three commercial mashing enzymes (Hitempase 2XL, Bioglucanase TX and Brewers protease) on yields of extract and FAN were studied during the mashing of unmalted and malted Safrari grist. Hitempase 2XL was principally responsible for extract yields in unmalted Safrari mash but had no impact on the malted Safrari mash type. Bioglucanase TX barely played a supporting role in these yields, while Brewers protease showed no significant role. Hitempase 2XL and Brewers protease individually contributed to yields in FAN in both unmalted and malted Safrari mashes, though the milling operation contributed to FAN yields for more than 50% in both mashes. This study shows that proper malting and mashing of this sorghum cultivar could lead to satisfactory wort properties in terms of extract and FAN for brewing purposes. Supplements of commercial mashing enzymes to boost their yields of extract in particular are thus not indispensable when mashing with malted Safrari. Optimisation of mashing properties through models clearly describing the actions of individual commercial mashing enzymes, as displayed in this study using the response surface methodology, is however of interest, particularly when mashing with high amounts of sorghum adjuncts. Further studies on the fermentability of worts obtained after such studies would be of importance in order to assess the exploitability of the results for improved brewing practices with this sorghum cultivar.

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