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Characterization of a Classical W14 Albino Mutation and a Putative New Robertson's Mutator-Induced Allele in Maize (*Zea Mays*)

By Ngoc (Josie) Huynh

Midwestern State University

Abstract- The newly identified albino mutation, w^*-5200 , arose in a maize (*Zea mays*) population derived from Robertson's Mutator transposon tagging crosses (Cook, 1988). Due to municipal water restrictions lasting three growing seasons, classical genetic allelism testing was not practical. Instead a molecular approach was pursued in an attempt to verify the genetic locus causing the mutant phenotype. Sequences of w^*-5200 and $616B w14-N335$ were compared to identify defects that would lead to inactivation of the gene in each, yielding evidence of neither a large-scale insertion nor any other obvious block to gene expression. Ultimately the lifting of water restrictions allowed a traditional allelism test, which verified that the novel mutation affects the $w14$ locus.

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CHARACTERIZATION OF A CLASSICAL W14 ALBINO MUTATION AND A PUTATIVE NEW ROBERTSON'S MUTATOR-INDUCED ALLELE IN MAIZE (ZEAMAYS)

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Characterization of a Classical W14 Albino Mutation and a Putative New Robertson's Mutator-Induced Allele in Maize (*Zea Mays*)

Ngoc (Josie) Huynh

Abstract- The newly identified albino mutation, *w*5200*, arose in a maize (*Zea mays*) population derived from Robertson's Mutator transposon tagging crosses (Cook, 1988). Due to municipal water restrictions lasting three growing seasons, classical genetic allelism testing was not practical. Instead a molecular approach was pursued in an attempt to verify the genetic locus causing the mutant phenotype. Sequences of *w*5200* and *616B w14-N335* were compared to identify defects that would lead to inactivation of the gene in each, yielding evidence of neither a large-scale insertion nor any other obvious block to gene expression. Ultimately the lifting of water restrictions allowed a traditional allelism test, which verified that the novel mutation affects the *w14* locus.

I. INTRODUCTION

Albino mutations are well known in plant and animal species. These mutations are caused by various recessive alleles which, when homozygous, lead directly or indirectly to a reduction in normal pigmentation in the organism. Genes that cause albinism in maize are identified by a number of different designations (Vancetovic et al., 2010): *w1*, *w2*, *w3* represent three white-albino loci; *wd1* the white deficiency locus; *lw1*, *lw2*, *lw3*, *lw4* four lemon white loci; *vp2*, *vp5*, *vp9* three viviparous loci; and *y10* the yellow endosperm locus (Neuffer et al., 1997).

A common feature of these genetic defects is that among their phenotypic effects is a depletion of carotenoid pigmentation in some or all tissues. The concomitant loss of chlorophyll and other biomolecules in affected tissues is due to the loss of photo-protective carotenoids. The first comprehensive review of albino maize mutants was by Robertson (1971). Since then, studies of these mutants have focused on developmental and metabolic features. Most studies have only described the mutations phenotypically, there are no molecular assays for the different genes that produce albino mutants in maize (Vančetović, et al., 2004; Vančetović, et al., 2010).

Robertson divided albino mutations into two classes. Class I albino mutations produce white-endosperm and white seedlings, whereas class II albino mutants produce both white and yellow endosperm and off-white seedlings (Robertson, 1975). In the albino mutant that is the focus of this research, designated *w**-

5200, trace amounts of chlorophyll are formed under dim light conditions. Since these albino seedlings grow from both white kernels and yellow kernels, they bear a Class II albino mutation, most likely in the *w14* locus (Stinard, 2013, and personal communication). Based on chromosome mapping of previously characterized *w14* alleles (chromosome 6L), and putative gene identification within the B73 draft genomic sequence, the likely identity of the *w14* locus is *Dxs1*, which encodes 1-deoxy-D-xylulose-5-phosphate synthase.

The newly identified mutation was derived from a population carrying Robertson's Mutator transposable elements. To determine whether the new albino mutation was due to a lesion at the *w14* locus, it was compared with another genetically confirmed *w14* allele: *616B w14-N335* (generated by EMS mutagenesis). Since most studies on albinism in maize have only described these mutations phenotypically, there was no comprehensive molecular assay for the potential genes that produce these albino mutants or to identify the mutation causing the new *w*5200* phenotype. The DNA sequences of *w*5200* and *w14* allele *616B w14-N335* were compared following amplification of gene segments by Polymerase Chain Reaction (PCR). We hypothesized that the new albino phenotype was caused by failure of the mutant to produce normal levels of carotenoids due to a mutation that affects the *w14* gene product. Specifically, we pursued the hypothesis that the defect was due to the insertion of a Robertson's Mutator transposable element into the *w14* locus.

Typically, allelism of a new lethal mutation is confirmed by field crosses between plants heterozygous for the new mutation and plants heterozygous for a mutation in the target locus. However, at the outset of this project municipal water restrictions, which continued for three successive summers, prohibited field work, which would have included allelism testing. Given the availability of a draft genome sequence, confirmed alleles of the putative target gene, and the facilities to perform the required molecular operations, the sequences of the new mutation and a known *w14* allele were determined and compared to identify the lesions resulting in their mutant phenotypes.

II. LITERATURE REVIEW

a) *Maize as a Model Organism*

Maize has historically been an important model organism for classical genetic research. However, it has a long reproductive cycle (four-months) and tall stature that are not easily accommodated in a greenhouse setting, as well as a large (2.3 x 10⁹ bp) haploid genome. By comparison, *Arabidopsis thaliana*, which has become the preferred flowering plant model, boasts diminutive size, a short (~6-week) reproductive cycle, and a small (1.35 x 10⁸ bp) haploid genome size (Johnston et al., 2007). Although *Arabidopsis* is superior for many studies, maize remains an attractive model organism for the study of mutations that result in seedling lethal phenotypes. *Arabidopsis* seeds provide too little stored energy to support the growth of homozygous photosynthesis-defective seedlings to a size useful for biochemical analysis, even when grown on sucrose-supplemented medium (Koorneef and Meinke, 2010). In contrast, typical maize kernels store adequate starch to produce 3- or 4-leaf seedlings, which provides sufficient tissue for many biochemical analyses. Furthermore, the mechanics of classical genetic manipulation of maize remain as straightforward as ever, including simple and reliable self- and outcrosses.

b) *Transposable Elements in Maize*

In maize, transposable elements (TE) make a big contribution to the genome (about 85% of the genomic material). Although most of the transposable elements are silenced most of the time to maintain genome stability, TEs still play an important role in plant evolution and environmental adaptation (Tenailon et al., 2010). Robertson's Mutator (Mu) has been frequently utilized for mutant induction (Vollbercht et al., 2010). The Mutator family of elements includes both an autonomous (master) element, Mu9/MuDR, and nonautonomous components, Mu1-Mu8 (Chomet, 1994). The elements routinely insert to unlinked sites in high numbers, making the family very mutagenic (McCarty et al., 2013). There are six classes within the Mu family, all of which share a comparable ~200 bp terminal inverted repeats (Bennetzen et al., 1993).

Miniature inverted-repeat transposable elements (MITE) are class II transposable elements. MITEs are short (80-500 bp) non-autonomous DNA transposons that are present near or within plant genes. Most MITEs are AT rich and produce target-site duplications of between 2 and 9 bp (Charrier et al., 1999; Patel et al., 2004).

c) *Effect of Albino Mutations on Leaves*

Chloroplasts degenerate in sunlight if they lack photo-protective carotenoids. Besides their photosynthetic role, chloroplasts perform a variety of other essential biochemical functions, such as biosynthesis of

amino acids, vitamins, and storage proteins. In the absence of carotenoids, the variety of compounds that are synthesized by chloroplasts, are reduced or absent. Plants bearing such mutations are only able to grow to maturity as heterozygotes or with sugar in their growth media (Wallis, 1963). In particular, carotenoids play an important role in protection of chlorophyll from photochemical degradation. Without carotenoids, chlorophyll is destroyed under normal light conditions resulting in a white (albino) leaf phenotype. Different albino mutations produce phenotypes that differ developmentally depending on the step in carotenoid biosynthesis affected and on the expression pattern of the gene that is mutated.

The w^{*}-5200 mutation produces unpigmented white albino seedlings in ambient sunlight. In very dim light, mutant seedlings produce very low levels of chlorophyll. Furthermore, there is no linkage between the white seedling phenotype and white endosperm as is the case for certain carotenoid-deficient mutations. Based on this phenotype, w^{*}-5200 is hypothesized to encode DXS (1-deoxy-D-xylulose-5-phosphate synthase), which catalyzes the first step in the 2-C-methyl-D-erythritol-4-P (MEP), isoprenoid biosynthetic pathway (Fig. 1). In maize, DXS is encoded by three different genes: Dxs1 on chromosome 6, Dxs2 on chromosome 7, and Dxs3 on chromosome 9 (Cordoba et al., 2011). Dxs1 is expressed primarily in young leaves and at lower levels in husks, tassel and mature leaves. Dxs2 is expressed primarily in mature leaves and at lower levels in yellow kernels and roots. Dxs3 is expressed in all vegetative and reproductive tissues but at the highest level in mature leaves. In addition to tissue-specific differences in expression, the three genes differ in the magnitude of their responses to light.

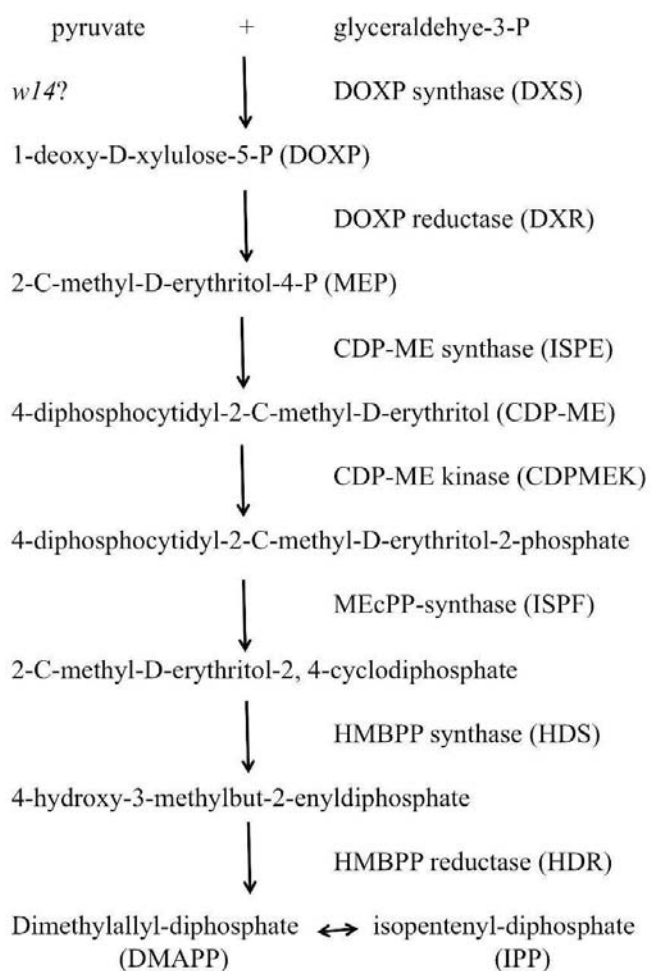


Figure 1: Chloroplast MEP Isoprenoid Biosynthetic Pathway (adapted from Stinard, 2013)

III. MATERIALS AND METHODS

a) Genetic Seed Sources

The maize (*Zea mays*) seed stock bearing w*-5200 arose from a self-pollination in a family that segregated another photosynthetic mutation independent of the w14 locus. The immediate source of experimental seed was ear 5200-a5 from the 2012 field season, which was the last field season before water use restrictions interrupted field production. The 616B w14-N335 EMS-induced stock was obtained from the Maize Genetics Cooperation Stock Center at the University of Illinois.

b) Growth Conditions

Zea mays kernels were grown in potting soil in the greenhouse at 21 °C - 25 °C (70 °F - 77 °F) under ambient light (360-650 $\mu\text{mol photons m}^{-2} \text{s}^{-1}$). For phenotypic verification of low light chlorophyll production, seedlings were germinated at 22 °C - 23°C at a light intensity of approximately 6 $\mu\text{mol photons m}^{-2} \text{s}^{-1}$.

c) Mutant Screening

Mutant seedlings were screened visually and recognized by their paper-white leaf color under ambient natural light in the greenhouse. Sprouting w*-5200 mutant seedlings were screened for the w14 albino phenotype using cool-yellow fluorescent lamps (approximately 6 $\mu\text{mol photons m}^{-2} \text{s}^{-1}$ 100-fold lower photon flux compared to the midday green house) in the dark. The use of a low photon flux permitted the light dependent conversion of barely detectable amounts of protochlorophyllide to chlorophyll without its subsequent destruction by photo-bleaching. The weakly pigmented leaves subsequently turned to paper-white under greenhouse lighting. To distinguish between homozygous and heterozygous phenotypically normal 616B seedlings with or without a MITE insertion in intron 3, genomic DNA was amplified using PCR primers A5F and A5R to produce products of one size in homozygotes (~800 bp) and products of two different sizes (~650 bp and ~800 bp) in heterozygotes (Table 1).

d) DNA Extraction

For Polymerase Chain Reaction (PCR) screens, genomic DNA was extracted from albino mutant seedlings and normal green seedlings (modified from Doyle and Doyle, 1987). Fresh, young leaves of albino and normal plants were collected. The tips (3 cm² of fresh leaves were cut into small pieces. Leaf tissue was ground thoroughly with a plastic pestle in 1.5 ml microfuge tubes containing 250 μl of extraction buffer (2% cetyl trimethylammonium bromide [CTAB], 20 mM EDTA (pH 8), 280 mM Tris-HCl (pH 8), 1.4 M NaCl, 8.2 mM 2-mercaptoethanol and 10 $\mu\text{g ml}^{-1}$ RNase A). An additional 750 μl extraction buffer were added to the test tube and the mixture was heated at 55 °C in a shaking water bath for 10 min then left at room temperature (25 °C) for 5 min to cool. To the cooled homogenate, 400 μl chloroform/isoamyl alcohol (24:1) were added and shaken gently to form an emulsion. Samples were centrifuged at 13,000g for 10 min at 5°C. The aqueous phase was transferred to a new microfuge tube and 2/3 volume of isopropanol (-20 °C) was added. Samples were placed at -20 °C for 10 min then centrifuged (13,000g) for 10 min to precipitate the DNA. The supernatant was removed and the pellet was washed with 76% EtOH and centrifuged (13,000g) for 10 min at 4 °C. The supernatant was removed and the pellet was air dried at room temperature. The DNA pellet was resuspended in 100 μl deionized H₂O (dH₂O).

e) Polymerase Chain Reaction (PCR) Amplification

The quality of the DNA samples was verified by PCR amplification using a primer pair unrelated to the w14 locus. Screening PCR was performed using Phusion Master Mix with HF Buffer (NEB). The PCR reactions of 20 μl contained 50 ng template DNA, 1X

Phusion High-Fidelity buffer mix and forward and reverse primers, each at 500nM. PCR amplification was carried out in an Applied Biosystems thermal cycler (v.2.09) with the following profile: initial denaturation for 1 min at 98 °C, followed by 35 cycles each of 10 s at 98 °C (denaturation), 30 s at 66 °C (annealing), 2.0-3.0 min depending on product size at 72°C (extension), followed by a final extension of 5 min at 72 °C. Screening PCR was also performed using Stratagene 1X master mix. The thermal profile for using Stratagene master mix was 5 min at 94°C for initial denaturation, followed by 35 cycles each of 1 min at 94 °C (denaturation), 1 min at 60 °C (annealing), 3.0-4.0 min depending on product size at 72 °C (extension), followed by a final extension of 5 min at 72 °C.

f) *PCR Product Analyses*

The amplification products were electrophoresed in 1% agarose gels in 1× TAE and stained with ethidium bromide (0.5 µg ml⁻¹ for detection. PCR product sizes were determined by comparison to Hi-Lo DNA size markers (Minnesota Molecular). Selected PCR products were cloned in vector pHSG299 by blunt-end ligation in reactions containing 0.1 µg vector and 3 µl PCR product in 1x T4 DNA ligase reaction mix (New England Biolabs)

incubated overnight in water at 10 °C and allowed to warm to room temperature through the night. Three microliters of the ligation reaction were added to NEB 5-alpha competent E. coli cells, mixed by flicking, and incubated on ice for 30 min. Cells were heat shocked at 42 °C for exactly 30 s, then incubated on ice for 5 min before addition of 950 µl of room temperature SOC. The mixture was incubated in a shaking incubator (260 rpm) at 37 °C for 60 min. The transformation was diluted 10-fold in SOC and 100 µl were plated on LB-Kanamycin plates supplemented with 50 µl X-gal (40 µg ml⁻¹) and 50µl IPTG(100 µg ml⁻¹) and dried just before plating. Colonies with recombinant plasmids were selected by blue white screening, checked for insert size by PCR, and sequenced by Eurofins Genomics. Alternatively, some PCR products were sequenced directly by the same commercial lab. Sequences were analyzed manually using SerialCloner® software (SerialBasics) by comparison to the Dxs1 region of maize inbred line B73 chromosome 6, RefGen_v4, whole genome shotgun sequence (NCBI accession number: NC_024464) to identify potentially mutagenic lesions. Putative mutagenic regions were sequenced with 3X or better coverage.

Table 1: Tm for selected primer pairs at 500 nM in 1X Master Mix

1F4 Primer	GCACACTCTCTCCCCGGC Primer Sequence	Tm (°C)	
		Phu	Stra
1F4	GCACACTCTCTCCCCGGC		
1R4	CCACCGCCATCCCGA	66	59
2F4	GAGTACGACAGCTTCGGCACG		
2R4	GAATGGGCCGGTCAAACCTAG	69	60
3F4	GGATCTCAGGTCGCAGCAAGTT		
3R4	ACGACGTCGATCTGCAGAAGCTA	69	60
4F4	GGTCCTCGACTGACGCCG		
4R4	CGGTAAGTGTGTTCCGGCGC	67	60
A5F	ATCCTCAACGACAACAAGCA		
A5R	AGAGTCAACTTGCTGCGACC	67	60
6BF	ACGTCGGGATCGCGGAGCAG		
6BR	CAACGGGACGCCAACGCCGT	69	60
1Falig	CCAACATGGTCGTCATG		
1Ralig	AAGTTCAGACACTCTAG	69	60
2Falig	GCACACTCTCTCCCCGG		
2Ralig	GGTGGTTAATTAGCTAG	65	60
3Falig	ATGGCTCTGGGTAACGT		
3Ralig	ACAGTCTGGAAATTTGA	69	60
UpSF	CATGGGGCTTTAGGAGCATAGGTCT		

UpSR	TGCGAGCAATGGGTGTCCTACCAAT	69	60
1021UpSR	GTCAGCGGTGGCAAAGTGAAGATTA		
UpSR	TGCGAGCAATGGGTGTCCTACCAAT	68	60
DSF	GCCAAACGCGTAGAACTTGTGCTGA		
DSR	TTCCAGAAATGGAGAAATTGGATCT	69	60

g) *Allelism Test*

Three years into the project municipal outdoor watering restrictions were lifted permitting a return to field propagation and the opportunity for a genetic test of allelism between w^*-5200 and 616B in the summer of 2016. Reciprocal crosses were performed between individuals heterozygous for the w^*-5200 allele and one individual heterozygous for 616B. When present on cross participants, a second viable ear shoot was self-pollinated in order to distinguish between heterozygous and homozygous-normal individuals. Progeny of allelism

crosses were screened for segregation of the albino phenotype indicating heterozygous parentage.

IV. RESULTS

a) *Seeding Screening*

Albino mutants were initially identified by their white leaf phenotypes. In subsequent screens albino seedlings were examined for low-light greening using 15W cool-yellow fluorescent lamps (approximately $6 \mu\text{mol photons m}^{-2} \text{s}^{-1}$ in a darkened room before transfer to the greenhouse for photo-bleaching (Fig. 2).

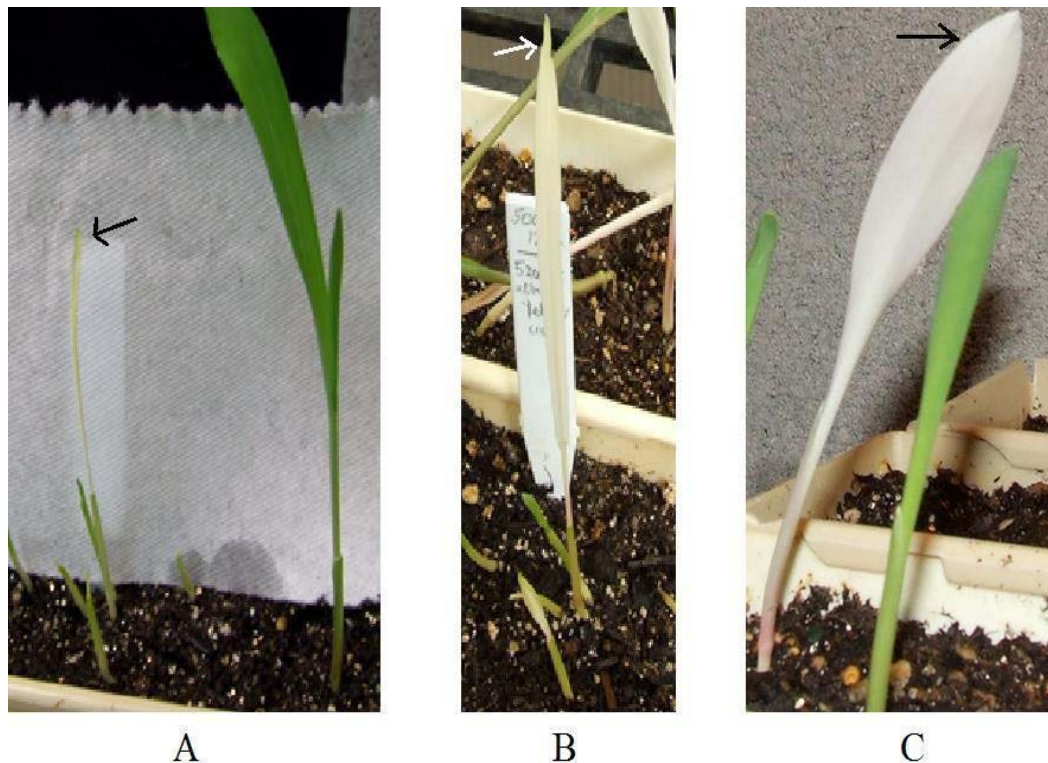


Figure 2: Pale green pigmentation in w14 mutant leaves. A Chlorophyll (black arrow), following low-light treatment; B Chlorophyll (white arrow) on day 4 of the greenhouse growth before sunrise; C No trace of chlorophyll following growth in full sunlight

b) *Allelism Test*

Allelism crosses confirmed that w^*-5200 is allelic to 616B w14-N335 (Table 2). After a nearly complete crop failure, one individual of the 616B stock survived. This individual served as pollen donor for one cross to w^*-5200 heterozygotes and as pollen recipient

in addition to supporting a self-pollination to verify heterozygosity.

Table 2: Allelism test $w^*-5200 \times 616B$ w14-N335

Cross	Phenotype		Ratio	Result
	Albino	Normal		
$w^*-5200 \otimes$	82	247	1:3.01	heterozygous
$w^*-5200-sib \times w^*-5200-sib$	37	102	1:2.76	heterozygous
$616B \otimes$	17	53	1:3.12	heterozygous
$w^*-5200 \times 616B$	15	37	1:2.47	allelic

\otimes indicates self-pollination

c) Sequence Analysis of DXS from w^*-5200 and 616B

To identify the mutations causing the w^*-5200 and 616B phenotype, their normal and mutant Dxs1 alleles were sequenced. The sequences of the two mutants were compared to each other, their normal siblings, and to the reference genome sequence of B73.

Figures 3 and 4 illustrate typical results from the amplification of the Dxs1 locus. Primers pairs 1F-2R, and 3F-4R gave similar results in that both amplified products of approximately the same sizes in mutant and normal samples.

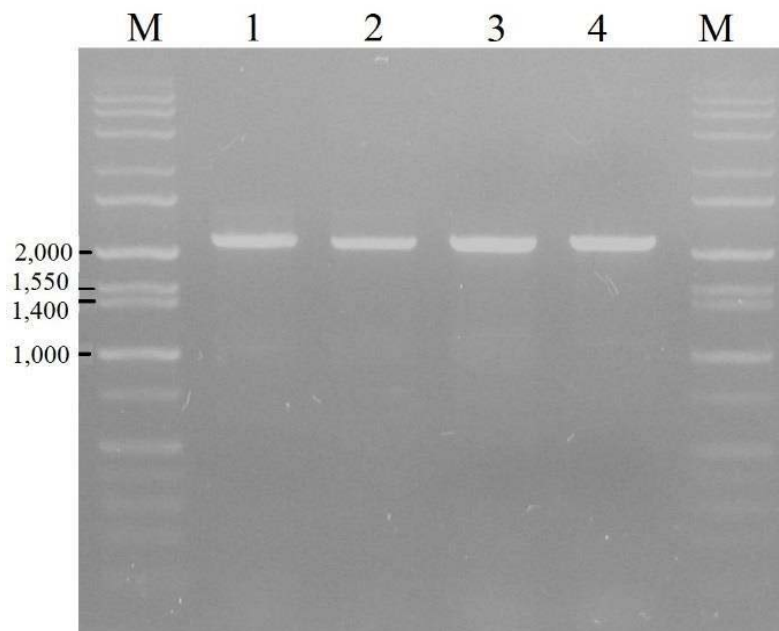


Figure 3: Agarose gel electrophoresis (1 %) of PCR products amplified with primers 1F and 2R. Lane 1: w^*-5200 albino; lane 2: w^*-5200 normal; lane 3: 616B albino; lane 4: 616B normal. M: Hi-Lo DNA markers (bp).

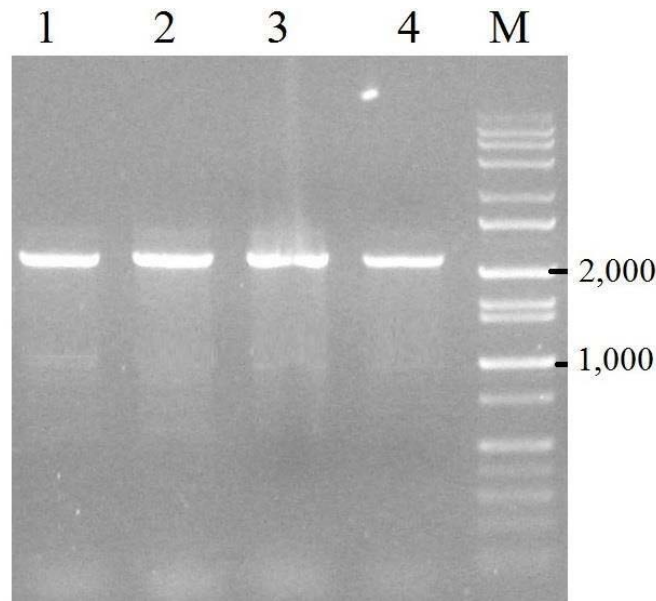


Figure 4: Agarose gel electrophoresis (1 %) of PCR products amplified with primers 3F and 4R. Lane 1: *w*5200* albino; lane 2: *w*5200* normal; lane 3: 616B albino; lane 4: 616B normal. M: Hi-Lo DNA markers (bp).

17 The sequence alignment of the DNA from the four *w14* alleles is shown in Figure 5. The alignment identifies no consequential difference between *w*5200* and 616B *w14*.

There are a few difference between the *Dxs1* reference sequence and *w*5200* and 616B *w14* (Table

3). Most of sequence differences occur in non-coding regions: in 5' or 3' untranslated regions (UTR) (G15T and C50T) or within the introns (A1291C, C1322A, T1338A, A1396G) in locations unlikely to affect splicing. Among mutations within exons, most were silent or consevative mutations (G1183T).

Table 3: Consequential differences between B73 and 616B albino, *w*5200* albino, *w*5200* normal, 616B normal and effects on protein sequence

Difference Relative to B73	Effect on Amino Acid Sequence	616B Albino	<i>w*5200</i> Albino	<i>w*5200</i> Normal	616B Normal
A294C	N31H	√	√	√	
G2769A	S436N	√	√	√	√
G3063A	D484N	√		√	
G3314A	G568D	√	√	√	
C3452T	A586V	√		√	

The sequences of the *Dxs1* locus from the two mutants were more than 90% identical. In comparison to the *Dxs1* sequence from B73, *w*5200* albino, *w*5200* normal and 616B normal, the 616B albino sequence includes a 140-bp deletion from intron 3 (Fig.

5). Where present, the 140-bp sequence is flanked by 3-bp direct repeats and contains 15-bp inverted repeats, each a signature of a miniature inverted-repeat transposable element (MITE) of the PIF/Harbinger class (Wessler et al., 1995).

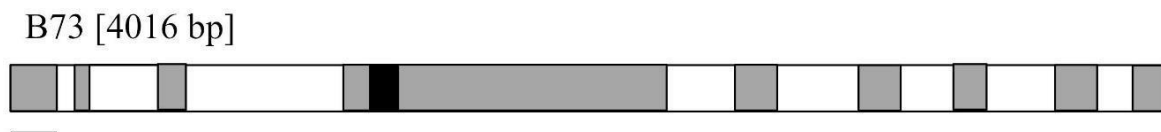


Figure 5: Map of maize *Dxs1* transcribed region. Grey: 5', 3' UTR or intron; white: exon; black: MITE insert when present. Segment lengths drawn to scale. Scale bar= ~200 bp.

As the MITE was inserted into intron 3 of the 616B normal allele and in both w*-5200 albino and normal alleles, the possibility that the deletion of the element from 616B was the cause of the albino phenotype was evaluated. The region surrounding the site of the MITE insertion was amplified in order to

determine the presence or absence of the element in 616B and three other genetically confirmed w14 alleles: 612A, 612N, and 612M using A5F and A5R (Table 1). To determine whether the MITE elements, when present, were inserted in the same location in all cases, the amplification products were sequenced.

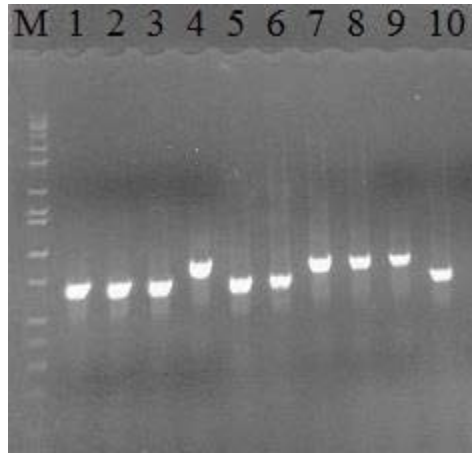


Figure 6: Agarose gel electrophoresis (1 %) of PCR products amplified with primers A5F and A5R. Lane 1: 612A normal; lane 2: 612A albino; lane 3: 612M normal; lane 4: 612M normal; lane 5: 612N normal; lane 6: 612N albino; lane 7: w*-5200 normal, lane 8: w*-5200 albino; lane 9: 616B normal; lane 10: 616B albino. M: Hi-Lo DNA markers (bp).

The results in both PCR products (Fig. 6) and sequences (appendix A3) provide evidence that the MITE is present in either or both normal and mutant DNAs among the five different allele-bearing stocks and that in each case where it is present, the MITE insertion is in the same location.

An additional mutation was considered as the possible cause of one or more of the albino phenotypes. A mutation that affects a critical upstream and

downstream regulatory region could prevent expression of the gene and result in the observed phenotypes. To evaluate this possibility, the 2,000 bp regions immediately upstream and downstream of the coding regions were amplified and the products compared to detect any large-scale insertions, deletions or other rearrangement that might affect gene expression (Fig. 7, 8).

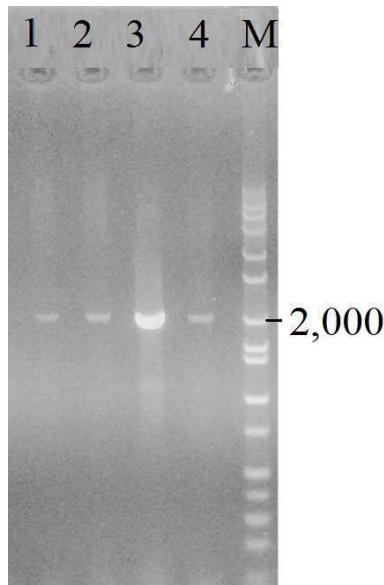


Figure 7: Agarose gel electrophoresis (1 %) of PCR products amplified downstream of Dxs1 with primers DSF and DSR. Lane 1: w*-5200 normal, lane 2: w*-5200 albino; lane 3: 616B normal; lane 4: 616B albino. M: Hi-Lo DNA markers (bp).

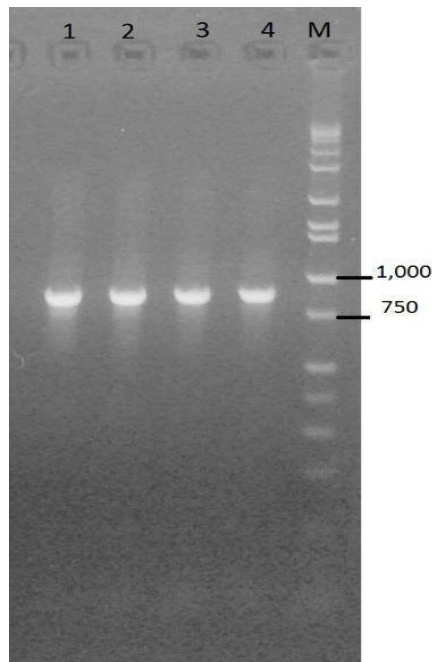


Figure 8: Agarose gel electrophoresis (1 %) of PCR products amplified upstream of *Dxs1* with primers 1021UpSF and UpSR. Lane 1: *w**-5200 normal, lane 2: *w**-5200 albino; lane 3: 616B normal; lane 4: 616B albino. M: Hi-Lo DNA markers (bp).

V. DISCUSSION AND CONCLUSION

The allelism test confirmed that the newly isolated albino mutation affects the *w14* locus. The mutations in *w**-5200 and 616B were determined to affect the same locus because the segregation ratio among progeny of the allelism crosses approximated the 1 (albino): 3 (normal) expected for a recessive trait (Table 1).

Although several sequence differences were identified among the samples that were examined, none was likely to inactivate *DXS* (Table A1). The MITE insert that is present in *w**-5200 albino, *w**-5200 normal, and 616B normal samples is absent from the 616B albino sequence (Fig. 6), which suggests that the excision may be the cause of the albino mutation in 616B. However, the MITE element was absent from both 612A samples, both 612N samples, and from 612M normal, but present in the 612M albino sequence. The presence or absence of the MITE element is independent of the albino phenotype and not its cause.

The *w**-5200 *Dxs1* sequence included neither a large-scale disruption, such as the hypothesized Robertson's Mutator transposable element insertion, nor any smaller variations that would obviously result in failure to produce a functional gene product.

The unexpected absence of obvious molecular evidence for the cause of the mutation in either *w**-5200 or 616B may be due to one of two possible explanations. The first is that the albino mutation that affects the *w14* gene product is not present within the transcribed sequence of the locus. Gene expression

may be affected by 5' or 3' sequence features outside the coding region that was examined. Although the MITE element that was detected in this study was present in the transcribed region of the locus, it raises the question of whether another MITE element may be inserted outside the transcribed region in a location able to affect expression. Reduced expression of *ZmRAP2.7*, a flowering time repressor gene, is associated with increased methylation in a regulatory element (*Vgt1*) that bears a highly methylated MITE element although a causative relationship between the two phenomena has not been demonstrated (Castelletti et al., 2014; Salvi et al., 2007). An ancient transposable element insertion ~60 kb upstream of *tb1* locus contributes to the enhanced expression of the gene and the resulting apical dominance seen in modern maize (Studer et al., 2011). A MITE insertion in the 3'UTR of *TaHSP16.9* (wheat) affects gene expression by stabilizing transcripts after exposure to heat stress (Li et al., 2014).

Amplification of the 5' and 3' UTRs identified no large-scale insertions or deletions that might prevent expression of *Dxs1* (Figs. 7 and 8). Although no large-scale changes were detected within 2,000 bp of the gene, smaller scale changes may be present that would have to be detected by more focused sequence analysis as was done for the transcription unit in this study. Modifications to more distant elements that may affect transcription of the *Dxs1* would be detected only by more elaborate genomic analyses. While disruption of a regulatory region is a plausible explanation for altered gene expression in individual cases, the likelihood that both mutation events (*w**-5200 and

616B), each generated in a mutagenesis project of known mechanism, would result from such disruptions seems small.

The second possible explanation for the lack of evidence of mutations in this locus is that Dxs1 is not the locus affected in w14 mutations. Although the circumstantial evidence on which the hypothesis of w14 involvement was based seems solid, to date it has not been confirmed by molecular means.

While it has been impossible to identify a sequence change that would obviously affect the expression of the w14 gene, it may be that a change eluded detection by sequencing that can be detected by other means. In future studies isolation and comparison of mRNA from mutant and normal sibling individuals should be performed. Northern analysis would provide information about the presence or absence of the transcripts from w14 and their sizes. In the case that mRNA is detected by Northern analysis, Real Time PCR should be employed to more precisely determine the relative levels of transcript accumulation between mutant and normal siblings.

While it seems unlikely that two independent mutations affecting the same locus would leave no molecular evidence in the transcribed sequence, it is possible. Fortunately, due to the long and active history of mutational analysis of maize, several other independent mutations affecting the w14 locus exist. Three of these will be propagated during the 2017 crop season to prepare DNA for sequence analysis. While this study did not yield a satisfying answer regarding the identity of the w14 locus or the nature of the mutations affecting it, data from additional w14 mutations and results from further experiments described above may shed light on the question.

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APPENDIX

Table A1: List of all sequence differences between B73 and 616B albino, w*-5200 albino, w*-5200 normal, 616B normal and effects on protein synthesis

Difference Relative to B73	Effect on Amino Acid Sequence	616B Albino	w*-5200 Albino	w*-5200 Normal	616B Normal
G15T	5'UTR	√	√		
C50T	5'UTR	√	√		
A236G	silent	√	√		
C293T	silent	√	√		
A294C	N31H	√	√	√	
A1396G	intron		√	√	
C2674A	silent		√	–	
G2769A	S436N	√	√	√	√
T2904C	intron	√	√	√	
A2951C	intron	√	√	√	
C2952G	intron	√	√	√	
T2953A	intron	√	√	√	
C2954T	intron	√	√	√	
G2955C	intron	√	√	√	
A2956G	intron	√	√	√	
ins 2956TTT	intron	√	√	√	
G3063A	D484N	√	–	√	
G3083A	silent	√	√	√	
C3084A	silent	√	√	√	
T3140G	silent	√	√	√	
C3239A	silent	√	√	√	
G3314A	G568D	√	√	√	
C3333T	intron	√	√	√	
C3335A	intron	√	√	√	
C3452T	A586V	√	–	√	
T3492C	silent	√	√	√	
C3633A	silent	√	√	√	
C3680T	silent	√	√	√	
C3707A	intron	√	√	√	
del 3753GA	intron	√	√	√	
A3761C	intron	√	√	√	
del 3787AA	intron	√	√	√	
A3975T	silent	√	√	√	
A4015G	3'UTR	√	√	√	

<p>w5200 616B w5200normal Dxs1 616Bnormal</p>	<pre>gcacactctctcccctgccacttcccaaatccgcccgcattcatgcaactcttctgtgca gcacactctctcccctgccacttcccaaatccgcccgcattcatgcaactcttctgtgca gcacactctctcccggccacttcccaaatccgcccgcattcatgcaaccttctgtgca gcacactctctcccggccacttcccaaatccgcccgcattcatgcaaccttctgtgca</pre>
<p>w5200 616B w5200normal Dxs1 616Bnormal</p>	<pre>ctgtcagcgccaccattagctcgcagctcaagctcgccactaccatTTTGGTCGGTtctt ctgtcagcgccaccattagctcgcagctcaagctcgccactaccatTTTGGTCGGTtctt ctgtcagcgccaccattagctcgcagctcaagctcgccactaccatTTTGGTCGGTtctt ctgtcagcgccaccattagctcgcagctcaagctcgccactaccatTTTGGTCGGTtctt</pre>
<p>w5200 616B w5200normal Dxs1 616Bnormal</p>	<pre>gaggaaatcgatcgaaccgTTGGAGTgcccaccactggcagaggctgTTgcattcttgagt gaggaaatcgatcgaaccgTTGGAGTgcccaccactggcagaggctgTTgcattcttgagt gaggaaatcgatcgaaccgTTGGAGTgcccaccactggcagaggctgTTgcattcttgagt gaggaaatcgatcgaaccgTTGGAGTgcccaccactggcagaggctgTTgcattcttgagt</pre>
<p>w5200 616B w5200normal Dxs1 616Bnormal</p>	<pre>tgagcaggaagaggagaggaagcaATGGCTCTGTGACGTTCTCTGTCCCAAGGGGTTTC tgagcaggaagaggagaggaagcaATGGCTCTGTGACGTTCTCTGTCCCAAGGGGTTTC tgagcaggaagaggagaggaagcaATGGCTCTGTGACGTTCTCTGTCCCAAGGGGTTTC tgagcaggaagaggagaggaagcaATGGCTCTGTGACGTTCTCTGTCCCAAGGGGTTTC</pre>
<p>w5200 616B w5200normal Dxs1 616Bnormal</p>	<pre>CTCGGCGTGCCGGCTCAGGACTCCCATTTGCTTCGGCGGTGAGCTCCATGTTACAAG CTCGGCGTGCCGGCTCAGGACTCCCATTTGCTTCGGCGGTGAGCTCCATGTTACAAG CTCGGTGTCGGCTCAGGACTCCCATTTGCTTCGGCGGTGAGCTCCATGTAACAAG CTCGGTGTCGGCTCAGGACTCCCATTTGCTTCGGCGGTGAGCTCCATGTAACAAG</pre>
<p>w5200 616B w5200normal Dxs1 616Bnormal</p>	<pre>CTGCTCCAGGCCAGGCCTATCAATCTCAAGgtaagttcctctgcccagttgtacgcaagc CTGCTCCAGGCCAGGCCTATCAATCTCAAGgtaagttcctctgcccagttgtacgcaagc CTGCTCCAGGCCAGGCCTATCAATCTCAAGgtaagttcctctgcccagttgtacgcaagc CTGCTCCAGGCCAGGCCTATCAATCTCAAGgtaagttcctctgcccagttgtacgcaagc</pre>
<p>w5200 616B w5200normal Dxs1 616Bnormal</p>	<pre>taaattttctcagttccggttccggttagttgatggccaatgctgctgagCCTCGGCG taaattttctcagttccggttccggttagttgatggccaatgctgctgagCCTCGGCG taaattttctcagttccggttccggttagttgatggccaatgctgctgagCCTCGGCG taaattttctcagttccggttccggttagttgatggccaatgctgctgagCCTCGGCG</pre>
<p>w5200 616B w5200normal Dxs1 616Bnormal</p>	<pre>GAGGCCGGCATGCGTGTGCGGCTCGTGTGCTCGGAGCGCGAGGCCGAGTACTACTCGCA GAGGCCGGCATGCGTGTGCGGCTCGTGTGCTCGGAGCGCGAGGCCGAGTACTACTCGCA GAGGCCGGCATGCGTGTGCGGCTCGTGTGCTCGGAGCGCGAGGCCGAGTACTACTCGCA GAGGCCGGCATGCGTGTGCGGCTCGTGTGCTCGGAGCGCGAGGCCGAGTACTACTCGCA</pre>
<p>w5200 616B w5200normal Dxs1 616Bnormal</p>	<pre>GAGGCCGCCACGCCGCTGCTGGACACCATCAACTACCCCGTCCACATGAAGAACCTGTCT GAGGCCGCCACGCCGCTGCTGGACACCATCAACTACCCCGTCCACATGAAGAACCTGTCT GAGGCCGCCACGCCGCTGCTGGACACCATCAACTACCCCGTCCACATGAAGAACCTGTCT GAGGCCGCCACGCCGCTGCTGGACACCATCAACTACCCCGTCCACATGAAGAACCTGTCT</pre>
<p>w5200 616B w5200normal Dxs1 616Bnormal</p>	<pre>TGTGAAGGAGCTGCGGCAGCTGGCCGACGAGCTCCGGTCCGACGTCATCTTCCACGTCTC TGTGAAGGAGCTGCGGCAGCTGGCCGACGAGCTCCGGTCCGACGTCATCTTCCACGTCTC TGTGAAGGAGCTGCGGCAGCTGGCCGACGAGCTCCGGTCCGACGTCATCTTCCACGTCTC TGTGAAGGAGCTGCGGCAGCTGGCCGACGAGCTCCGGTCCGACGTCATCTTCCACGTCTC</pre>

<p>w5200 616B w5200normal Dxs1 616Bnormal</p>	<p>CAAGACCGGCGGCCACCTCGGGTCCAGCCTCGGCGTGGTGGAGCTCACCGTCGCGCTGCA CAAGACCGGCGGCCACCTCGGGTCCAGCCTCGGCGTGGTGGAGCTCACCGTCGCGCTGCA CAAGACCGGCGGCCACCTCGGGTCCAGCCTCGGCGTGGTGGAGCTCACCGTCGCGCTGCA CAAGACCGGCGGCCACCTCGGGTCCAGCCTCGGCGTGGTGGAGCTCACCGTCGCGCTGCA -----</p>
<p>w5200 616B w5200normal Dxs1 616Bnormal</p>	<p>CTACGTCTTCAACGCGCCGCGAGACCGCATCCTCTGGGACGTCGGCCACCAGgtacgctg CTACGTCTTCAACGCGCCGCGAGACCGCATCCTCTGGGACGTCGGCCACCAGgtacgctg CTACGTCTTCAACGCGCCGCGAGACCGCATCCTCTGGGACGTCGGCCACCAGgtacgctg CTACGTCTTCAACGCGCCGCGAGACCGCATCCTCTGGGACGTCGGCCACCAGgtacgctg -----</p>
<p>w5200 616B w5200normal Dxs1 616Bnormal</p>	<p>atgcgccatgggcccgcgcgcgcggcatggctctgggtaacgtgcgctccatgtgagcgctg atgcgccatgggcccgcgcgcgcggcatggctctgggtaacgtgcgctccatgtgagcgctg atgcgccatgggcccgcgcgcgcggcatggctctgggtaacgtgcgctccatgtgagcgctg atgcgccatgggcccgcgcgcgcggcatggctctgggtaacgtgcgctccatgtgagcgctg -----</p>
<p>w5200 616B W5200normal Dxs1 616Bnormal</p>	<p>ccgggcaggtcgcggacaggctagctaattaaccaccccgaccggggttttgtttgtct ccgggcaggtcgcggacaggctagctaattaaccaccccgaccggggttttgtttgtct ccgggcaggtcgcggacaggctagctaattaaccaccccgaccggggttttgtttgtct ccgggcaggtcgcggacaggctagctaattaaccaccccgaccggggttttgtttgtct -----</p>
<p>w5200 616B w5200normal Dxs1 616Bnormal</p>	<p>gattcgcgcgcagTcagTCGTACCCGCACAAGATCCTGACGGGGCGGCGGACAAGATGC gattcgcgcgcagTcagTCGTACCCGCACAAGATCCTGACGGGGCGGCGGACAAGATGC gattcgcgcgcagTcagTCGTACCCGCACAAGATCCTGACGGGGCGGCGGACAAGATGC gattcgcgcgcagTcagTCGTACCCGCACAAGATCCTGACGGGGCGGCGGACAAGATGC -----</p>
<p>w5200 616B w5200normal Dxs1 616Bnormal</p>	<p>CGACGATGCGGCAGACCAACGGCCTGGCGGGCTTCACCAAGCGCGCCGAGAGCGAGTACG CGACGATGCGGCAGACCAACGGCCTGGCGGGCTTCACCAAGCGCGCCGAGAGCGAGTACG CGACGATGCGGCAGACCAACGGCCTGGCGGGCTTCACCAAGCGCGCCGAGAGCGAGTACG CGACGATGCGGCAGACCAACGGCCTGGCGGGCTTCACCAAGCGCGCCGAGAGCGAGTACG -----</p>
<p>w5200 616B w5200normal Dxs1 616Bnormal</p>	<p>ACAGCTTCGGCACGGGCCACAGTCCACCACCATCTCCGCGGCGCTCGGGATGGCGGTGG ACAGCTTCGGCACGGGCCACAGTCCACCACCATCTCCGCGGCGCTCGGGATGGCGGTGG ACAGCTTCGGCACGGGCCACAGTCCACCACCATCTCCGCGGCGCTCGGGATGGCGGTGG ACAGCTTCGGCACGGGCCACAGTCCACCACCATCTCCGCGGCGCTCGGGATGGCGGTGG -----</p>
<p>w5200 616B w5200normal Dxs1 616Bnormal</p>	<p>GCCGGGACCTCAAGGGCGGCAAGAACAACGTGGTCGCGGTGATCGGCGACGGCGCCGATGA GCCGGGACCTCAAGGGCGGCAAGAACAACGTGGTCGCGGTGATCGGCGACGGCGCCATGA GCCGGGACCTCAAGGGCGGCAAGAACAACGTGGTCGCGGTGATCGGCGACGGCGCCATGA GCCGGGACCTCAAGGGCGGCAAGAACAACGTGGTCGCGGTGATCGGCGACGGCGCCATGA -----</p>
<p>w5200 616B w5200normal Dxs1 616Bnormal</p>	<p>CGCCCCGGCAGGCGTACGAGGCCATGAACAACGCCGGGTACCTGGACTCCGACATGATCG CGGCCGGCAGGCGTACGAGGCCATGAACAACGCCGGGTACCTGGACTCCGACATGATCG CGGCCGGCAGGCGTACGAGGCCATGAACAACGCCGGGTACCTGGACTCCGACATGATCG CGGCCGGCAGGCGTACGAGGCCATGAACAACGCCGGGTACCTGGACTCCGACATGATCG -----</p>

<i>w5200</i>	TCATCCTCAACGACAACAAGCAGGTGTCCTTGCCACGGCGACTCTCGACGGGCCGGTGC
<i>616B</i>	TCATCCTCAACGACAACAAGCAGGTGTCCTTGCCACGGCGACGCTCGACGGGCCGGTGC
<i>w5200normal</i>	TCATCCTCAACGACAACAAGCAGGTGTCCTTGCCACGGCGACGCTCGACGGGCCGGTGC
<i>Dxs1</i>	TCATCCTCAACGACAACAAGCAGGTGTCCTTGCCACGGCGACGCTCGACGGGCCGGTGC
<i>616Bnormal</i>	ACGACAACAAGCAGGTGTCCTTGCCACGGCGACGCTCGACGGGCCGGTGC
<i>w5200</i>	CGCCCGTGGGCGCGCTCAGCAGCGCCCTCAGCAAGCTGCAGTCAAGCAGGCCGCTCAGGG
<i>616B</i>	CGCCCGTGGGCGCGCTCAGCAGCGCCCTCAGCAAGCTGCAGTCAAGCAGGCCGCTCAGGG
<i>w5200normal</i>	CGCCCGTGGGCGCGCTCAGCAGCGCCCTCAGCAAGCTGCAGTCAAGCAGGCCGCTCAGGG
<i>Dxs1</i>	CGCCCGTGGGCGCGCTCAGCAGCGCCCTCAGCAAGCTGCAGTCAAGCAGGCCGCTCAGGG
<i>616Bnormal</i>	CGCCCGTGGGCGCGCTCAGCAGCGCCCTCAGCAAGCTGCAGTCAAGCAGGCCGCTCAGGG
<i>w5200</i>	AGCTCAGGGAGGTGGCCAAGGtacgtccgcaaaaaccttggatgggacgctacatcacc
<i>616B</i>	AGCTCAGGGAGGTGGCCAAGGtacgtccgcaaaaaccttggatgggacgctacatcacc
<i>w5200normal</i>	AGCTCAGGGAGGTGGCCAAGGtacgtccgcaaaaaccttggatgggacgctacatcacc
<i>Dxs1</i>	AGCTCAGGGAGGTGGCCAAGGtacgtccgcaaaaaccttggatgggacgctacatcacc
<i>616Bnormal</i>	AGCTCAGGGAGGTGGCCAAGGtacgtccgcaaaaaccttggatgggacgctacatcacc
<i>w5200</i>	tattgacagcccggccggataggaagcgccacgtaagggcttgttcggttattcccaat
<i>616B</i>	tcttgacagcccggccggtaggaagcgccacgtaagggcttgttcggttattcccaat
<i>W5200normal</i>	tcttgacagcccggccggtaggaagcgccacgtaagggcttgttcggttattcccaat
<i>Dxs1</i>	tcttgacagcccggccggtaggaagcgccacgtaagggcttgttcggttattcccaat
<i>616Bnormal</i>	tcttgacagcccggccggtaggaagcgccacgtaagggcttgttcggttattcccaat
<i>w5200</i>	acacatggattggatgggattggaaaaaattatgaagaagtttgagctgtttgggattca
<i>616B</i>	-----
<i>w5200normal</i>	acacatggattggatgggattggaaaaaattatgaagaagtttgagctgtttgggattca
<i>Dxs1</i>	acacatggattggatggatgggattggaaaaaattatgaagaagtttgagctgtttgggattca
<i>616Bnormal</i>	acacatggattggatggatgggattggaaaaaattatgaagaagtttgagctgtttgggattca
<i>w5200</i>	aaccatccaatcccgtcaatccacatggattgagagctaaccgaacaagccctaacac
<i>616B</i>	-----cac
<i>w5200normal</i>	aaccatccaatcccgtcaatccacatggattgagagctaaccgaacaagccctaacac
<i>Dxs1</i>	aaccatccaatcccgtcaatccacatggattgagagctaaccgaacaagccctaacac
<i>616Bnormal</i>	aaccatccaatcccgtcaatccacatggattgagagctaaccgaacaagccctaacac
<i>w5200</i>	gtcaaatttccagactgtcctcgttctcacggaagcgctagattttctggaatcttgat
<i>616B</i>	gtcaaatttccagactgtcctcgttctcacggaagcgctagattttctggaatcttgat
<i>W5200normal</i>	gtcaaatttccagactgtcctcgttctcacggaagcgctagattttctggaatcttgat
<i>Dxs1</i>	gtcaaatttccagactgtcctcgttctcacggaagcgctagattttctggaatcttgat
<i>616Bnormal</i>	gtcaaatttccagactgtcctcgttctcacggaagcgctagattttctggaatcttgat
<i>w5200</i>	tagtagtaattcctatagttgcacatattttttatcaaatcattaagataaataataa
<i>616B</i>	tagtagtaattcctatagttgcacatattttttatcaaatcattaagataaataataa
<i>W5200normal</i>	tagtagtaattcctatagttgcacatattttttatcaaatcattaagataaataataa
<i>Dxs1</i>	tagtagtaattcctatagttgcacatattttttatcaaatcattaagataaataataa
<i>616Bnormal</i>	tagtagtaattcctatagttgcacatattttttatcaaatcattaagataaataataa
<i>w5200</i>	ccttgcctttggaattaaatggaaaactgtcaacagtcacatcgggcagcagcagcgtaca
<i>616B</i>	ccttgcctttggaattaaatggaaaactgtcaacagtcacatcgggcagcagcagcgtaca
<i>w5200normal</i>	ccttgcctttggaattaaatggaaaactgtcaacagtcacatcgggcagcagcagcgtaca
<i>Dxs1</i>	ccttgcctttggaattaaatggaaaactgtcaacagtcacatcgggcagcagcagcgtaca
<i>616Bnormal</i>	ccttgcctttggaattaaatggaaaactgtcaacagtcacatcgggcagcagcagcgtaca
<i>w5200</i>	tgacgcgagctatggagcttcttgaatctactgcaacgaaagcgtctgaatgaaacttgtt
<i>616B</i>	tgacgcgagctatggagcttcttgaatctactgcaacgaaagcgtctgaatgaaacttgtt
<i>W5200normal</i>	tgacgcgagctatggagcttcttgaatctactgcaacgaaagcgtctgaatgaaacttgtt
<i>Dxs1</i>	tgacgcgagctatggagcttcttgaatctactgcaacgaaagcgtctgaatgaaacttgtt
<i>616Bnormal</i>	tgacgcgagctatggagcttcttgaatctactgcaacgaaagcgtctgaatgaaacttgtt

w5200	tagtcttagcgaacatgccaaaataggctgcgtcagatgcagagtgcactaatct
616B	tagtcttagcgaacatgccaaaataggctgcgtcagatgcagagtgcactaatct
w5200normal	tagtcttagcgaacatgccaaaataggctgcgtcagatgcagagtgcactaatct
Dxs1	tagtcttagcgaacatgccaaaataggctgcgtcagatgcagagtgcactaatct
616Bnormal	tagtcttagcgaacatgccaaaataggctgcgtcagatgcagagtgcactaatct
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616B	gcttctagggggcatgcatggacacatggtcaccagaaaggccgatgacgaagcgtatac
w5200normal	gcttctagggggcatgcatggacacatggtcaccagaaaggccgatgacgaagcgtatac
Dxs1	gcttctagggggcatgcatggacacatggtcaccagaaaggccgatgacgaagcgtatac
616Bnormal	gcttctagggggcatgcatggacacatggtcaccagaaaggccgatgacgaagcgtatac
w5200	gagctagcattctagcaacagtcctagatccgtagatccgagtaatgccactactaggcaaacttt
616B	gagctagcattctagcaacagtcctagatccgtagatccgagtaatgccactactaggcaaacttt
w5200normal	gagctagcattctagcaacagtcctagatccgtagatccgagtaatgccactactaggcaaacttt
Dxs1	gagctagcattctagcaacagtcctagatccgtagatccgagtaatgccactactaggcaaacttt
616Bnormal	gagctagcattctagcaacagtcctagatccgtagatccgagtaatgccactactaggcaaacttt
w5200	gtataaacaagctactcaaagcatggatgatggatctcaggtcgcagcaagttgactct
616B	gtataaacaagctactcaaagcatggatgatggatctcaggtcgcagcaagttgactct
w5200normal	gtataaacaagctactcaaagcatggatgatggatctcaggtcgcagcaagttgactct
Dxs1	gtataaacaagctactcaaagcatggatgatggatctcaggtcgcagcaagttgactct
616Bnormal	gtataaacaagctactcaaagcatggatgatggatct-----
w5200	agaaatagtttatcatgctactcgagctgtatcca-gtttgactgacattggttcatctt
616B	agaaatagtttatcatgctactcgagctgtatccaagtttgactgacattggttcatctt
w5200normal	agaaatagtttatcatgctactcgagctgtatccaagtttgactgacattggttcatctt
Dxs1	agaaatagtttatcatgctactcgagctgtatccaagtttgactgacattggttcatctt
616Bnormal	-----
w5200	ctactggactggtcataatattacctgggctagggttttgaccggcccattcttgttgggc
616B	ctactggactggtcataatattacctgggctagggttttgaccggcccattcttgttgggc
w5200normal	ctactggactggtcataatattacctgggctagggttttgaccggcccattcttgttgggc
Dxs1	ctactggactggtcataatattacctgggctagggttttgaccggcccattcttgttgggc
616Bnormal	-----
w5200	cgacaaggattgtggaaggaatgggcccagggtttgcagctctgagctctctgaccgat
616B	cgacaaggattgtggaaggaatgggcccagggtttgcagctctgagctctctgaccgat
w5200normal	cgacaaggattgtggaaggaatgggcccagggtttgcagctctgagctctctgaccgat
Dxs1	cgacaaggattgtggaaggaatgggcccagggtttgcagctctgagctctctgaccgat
616Bnormal	-----
w5200	cgctggcgtgcgtgcttgtgcttgtgcaggGAGTGACGAAGCAGATCGGTGGCTCAGTGC
616B	cgctggcgtgcgtgcttgtgcttgtgcaggGAGTGACGAAGCAGATCGGTGGCTCAGTGC
w5200normal	cgctggcgtgcgtgcttgtgcttgtgcaggGAGTGACGAAGCAGATCGGTGGCTCAGTGC
Dxs1	cgctggcgtgcgtgcttgtgcttgtgcaggGAGTGACGAAGCAGATCGGTGGCTCAGTGC
616Bnormal	-----
w5200	ACGAGCTGGCGGCGAAGGTGGACGAGTACGCCCGCGGCATGATCAGCGGGCCCGGCTCCT
616B	ACGAGCTGGCGGCGAAGGTGGACGAGTACGCCCGCGGCATGATCAGCGGGCCCGGCTCCT
w5200normal	ACGAGCTGGCGGCGAAGGTGGACGAGTACGCCCGCGGCATGATCAGCGGGCCCGGCTCCT
Dxs1	ACGAGCTGGCGGCGAAGGTGGACGAGTACGCCCGCGGCATGATCAGCGGGCCCGGCTCCT
616Bnormal	-----

<i>w5200</i>	CGCTCTTCGAGGAGCTCGGTCTCTACTACATCGGCCCGTCGACGGCCACAACATCGACG
<i>616B</i>	CGCTCTTCGAGGAGCTCGGTCTCTACTACATCGGCCCGTCGACGGCCACAACATCGACG
<i>w5200normal</i>	CGCTCTTCGAGGAGCTCGGTCTCTACTACATCGGCCCGTCGACGGCCACAACATCGACG
<i>Dxs1</i>	CGCTCTTCGAGGAGCTCGGTCTCTACTACATCGGCCCGTCGACGGCCACAACATCGACG
<i>616Bnormal</i>	-----
<i>w5200</i>	ACCTCATCACCATCCTCAACGACGTCAAGAGCACCAAGACCACCGGCCCGTCCTCATCC
<i>616B</i>	ACCTCATCACCATCCTCAACGACGTCAAGAGCACCAAGACCACCGGCCCGTCCTCATCC
<i>w5200normal</i>	ACCTCATCACCATCCTCAACGACGTCAAGAGCACCAAGACCACCGGCCCGTCCTCATCC
<i>Dxs1</i>	ACCTCATCACCATCCTCAACGACGTCAAGAGCACCAAGACCACCGGCCCGTCCTCATCC
<i>616Bnormal</i>	-----
<i>w5200</i>	ACGTCGTCACCGAGAAGGGCCGCGGTACCCCTACGCCGAGCGAGCCGCCGACAAGTACC
<i>616B</i>	ACGTCGTCACCGAGAAGGGCCGCGGTACCCCTACGCCGAGCGAGCCGCCGACAAGTACC
<i>w5200normal</i>	ACGTCGTCACCGAGAAGGGCCGCGGTACCCCTACGCCGAGCGAGCCGCCGACAAGTACC
<i>Dxs1</i>	ACGTCGTCACCGAGAAGGGCCGCGGTACCCCTACGCCGAGCGAGCCGCCGACAAGTACC
<i>616Bnormal</i>	-----
<i>w5200</i>	ACGGTacaacgcaccacatgagctagcatatgtgccactgtttgctgttgacagcaca
<i>616B</i>	ACGGTacaacgcaccacatgagctagcatatgtgccactgtttgctgttgacagcaca
<i>w5200normal</i>	ACGGTacaacgcaccacatgagctagcatatgtgccactgtttgctgttgacagcaca
<i>Dxs1</i>	ACGGTacaacgcaccacatgagctagcatatgtgccactgtttgctgttgacagcaca
<i>616Bnormal</i>	-----
<i>w5200</i>	gatcgtaccccgaccggaatctgtgcgatcattggctctgttgtttgatgcggtgctgc
<i>616B</i>	gatcgtaccccgaccggaatctgtgcgatcattggctctgttgtttgatgcggtgctgc
<i>W5200normal</i>	gatcgtaccccgaccggaatctgtgcgatcattggctctgttgtttgatgcggtgctgc
<i>Dxs1</i>	gatcgtaccccgaccggaatctgtgcgatcattggctctgttgtttgatgcggtgctgc
<i>616Bnormal</i>	-----
<i>w5200</i>	gtgcaggtGTCGCCAAGTTTGATCCGGCGACCGGGAAGCAGTTCAAGTCCCCGCCAAGA
<i>616B</i>	gtgcaggtGTCGCCAAGTTTGATCCGGCGACCGGGAAGCAGTTCAAGTCCCCGCCAAGA
<i>w5200normal</i>	gtgcaggtGTCGCCAAGTTTGATCCGGCGACCGGGAAGCAGTTCAAGTCCCCGCCAAGA
<i>Dxs1</i>	gtgcaggtGTCGCCAAGTTTGATCCGGCGACCGGGAAGCAGTTCAAGTCCCCGCCAAGA
<i>616Bnormal</i>	-----
2641	
<i>w5200</i>	CGCTGTCTACACCAACTACTTCGCCGAGGCGCTAATCGCCGAGGCGGAGCAGGACAGCA
<i>616B</i>	CGCTGTCTACACCAACTACTTCGCCGAGGCGCTAATCGCCGAGGCGGAGCAGGACAGCA
<i>w5200normal</i>	CGCTGTCTACACCAACTACTTCGCCGAGGCGCTAATCGCCGAGGCGGAGCAGGACAGCA
<i>Dxs1</i>	CGCTGTCTACACCAACTACTTCGCCGAGGCGCTAATCGCCGAGGCGGAGCAGGACAGCA
<i>616Bnormal</i>	-----
<i>w5200</i>	AGATCGTGGCCATCCACGCGGCCATGGGCGGCGGCACGGGGCTCAACTACTTCTCCGCC
<i>616B</i>	AGATCGTGGCCATCCACGCGGCCATGGGCGGCGGCACGGGGCTCAACTACTTCTCCGCC
<i>w5200normal</i>	AGATCGTGGCCATCCACGCGGCCATGGGCGGCGGCACGGGGCTCAACTACTTCTCCGCC
<i>Dxs1</i>	AGATCGTGGCCATCCACGCGGCCATGGGCGGCGGCACGGGGCTCAACTACTTCTCCGCC
<i>616Bnormal</i>	-----
<i>w5200</i>	GCTTCCCGAACCGGTGCTTCGACGTCCGGATCGCGGAGCAGCAGCCGTCACGTTCCGCGG
<i>616B</i>	GCTTCCCGAACCGGTGCTTCGACGTCCGGATCGCGGAGCAGCAGCCGTCACGTTCCGCGG
<i>w5200normal</i>	GCTTCCCGAACCGGTGCTTCGACGTCCGGATCGCGGAGCAGCAGCCGTCACGTTCCGCGG
<i>Dxs1</i>	GCTTCCCGAACCGGTGCTTCGACGTCCGGATCGCGGAGCAGCAGCCGTCACGTTCCGCGG
<i>616Bnormal</i>	-----

w5200 CCGGCCTGGCCTGCGAGGGCCTCAAGCCCTTCTGCGCCATCTACTCGTCTTTTCCTGCAGC
616B CCGGCCTGGCCTGCGAGGGCCTCAAGCCCTTCTGCGCCATCTACTCGTCTTTTCCTGCAGC
w5200normal CCGGCCTGGCCTGCGAGGGCCTCAAGCCCTTCTGCGCCATCTACTCGTCTTTTCCTGCAGC
Dxs1 CCGGCCTGGCCTGCGAGGGCCTCAAGCCCTTCTGCGCCATCTACTCGTCTTTTCCTGCAGC
616Bnormal

w5200 GCGGCTACGACCAGGTgcgcaagcgcgcccgtgtgcccgcggccgggcccgttcttcgcatt
616B GCGGCTACGACCAGGTgcgcaagcgcgcccgtgtgcccgcggccgggcccgttcttcgcatt
w5200normal GCGGCTACGACCAGGTgcgcaagcgcgcccgtgtgcccgcggccgggcccgttcttcgcatt
Dxs1 GCGGCTACGACCAGGTgcgcaagcgcgcccgtgtgcccgcggccgggcccgttcttcgcatt
616Bnormal

w5200 tgcttgctgctcgatcgtttcgttttcttcttttgctgccccgcggctcctcgactgacgc
616B tgcttgctgctcgatcgtttcgttttcttcttttgctgccccgcggctcctcgactaacgc
W5200normal tgcttgctgctcgatcgtttcgttttcttcttttgctgccccgcggctcctcgactgacgc
Dxs1 tgcttgctgctactcgcttttcttcttttgctgccccgcggctcctcgactgacgc---
616Bnormal

W5200 cgtacgcacgtcgccgatgggcccgtgtgggtggtggcgaggtcgTGCACG
616B cgtacgcacgtcgccgatgggcccgtgtgggtggtggcgaggtcgTGCACG
W5200normal cgtacgcacgtcgccgatgggcccgtgtgggtggtggcgaggtcgTGCACG
Dxs1 cgtacgcacgtcgccgatgggcccgtgtgggtggtggcgaggtcgTGCACG
616Bnormal

w5200 ACGTCAATCTGCAGAAGCTACCGGTAAGGTTCCGCATGGACAGGGCCGGCTGGTCGGCG
616B ACGTCAATCTGCAGAAGCTACCGGTAAGGTTCCGCATGGACAGGGCCGGCTGGTCGGCG
w5200normal ACGTCAATCTGCAGAAGCTACCGGTAAGGTTCCGCATGGACAGGGCCGGCTGGTCGGCG
Dxs1 ACGTCAATCTGCAGAAGCTACCGGTAAGGTTCCGCATGGACAGGGCCGGCTGGTCGGCG
616Bnormal

w5200 CGGACGGGCCGACCCACTGCGGGGCGTTTCGACGTTCGCGTACATGGCCTGCCTGCCCAACA
616B CGGACGGGCCGACCCACTGCGGGGCGTTTCGACGTTCGCGTACATGGCCTGCCTGCCCAACA
w5200normal CGGACGGGCCGACCCACTGCGGGGCGTTTCGACGTTCGCGTACATGGCCTGCCTGCCCAACA
Dxs1 CGGACGGGCCGACCCACTGCGGGGCGTTTCGACGTTCGCGTACATGGCCTGCCTGCCCAACA
616Bnormal

w5200 TGGTCGTCATGGCCCCGTCGACGAGGCCGAGCTCTGCCACATGGTCGCCACAGCCGCGG
616B TGGTCGTCATGGCCCCGTCGACGAGGCCGAGCTCTGCCACATGGTCGCCACAGCCGCGG
w5200normal TGGTCGTCATGGCCCCGTCGACGAGGCCGAGCTCTGCCACATGGTCGCCACAGCCGCGG
Dxs1 TGGTCGTCATGGCCCCGTCGACGAGGCCGAGCTCTGCCACATGGTCGCCACAGCCGCGG
616Bnormal

w5200 CAATCGACGACCCCGCTCCTGCTTCCGCTACCCGAGAGGCAACGGCGTTGGCGTCCCGT
616B CAATCGACGACCCCGCTCCTGCTTCCGCTACCCGAGAGGCAACGGCGTTGGCGTCCCGT
w5200normal CAATCGACGACCCCGCTCCTGCTTCCGCTACCCGAGAGGCAACGGCGTTGGCGTCCCGT
Dxs1 CCATCGACGACCCCGCTCCTGCTTCCGCTACCCGAGAGGCAACGGCGTTGGCGTCCCGT
616Bnormal

w5200 TGCCGCCCAACTACAAAGACTCTCCCTCGAGGTATGTATAACATCGATCGATGGTGTATT
616B TGCCGCCCAACTACAAAGACTCTCCCTCGAGGTATGTATAACATCGATCGATGGTGTATT
w5200normal TGCCGCCCAACTACAAAGACTCTCCCTCGAGGTATGTATAACATCGATCGATGGTGTATT
Dxs1 TGCCGCCCAACTACAAAGACTCTCCCTCGAGGTATGTATAACATCGATCGATGGTGTATT
616Bnormal

w5200 TTACTCATTFFFFFFATCAATCGGGGATCTTCAGTTTTGGTAATCGGCCATATAGGTCGGC
616B TTACTCATTFFFFFFATCAATCGGGGATCTTCAGTTTTGGTAATCGGCCATATAGGTCGGC
w5200normal TTACTCATTFFFFFFATCAATCGGGGATCTTCAGTTTTGGTAATCGGCCATATAGGTCGGC
Dxs1 TTACTCATTFFFFFFATCAATCGGGGATCTTCAGTTTTGGTAATCGGCCATATAGGTCGGC
616Bnormal

w5200 AAAGGCAGGATCCTGCTGGAGGGCGACCGGGTGGCTGCTGGGGTACGGGTCGGCAGTG
616B AAAGGCAGGATCCTGCTGGAGGGCGACCGGGTGGCTGCTGGGGTACGGGTCGGCAGTG
w5200normal AAAGGCAGGATCCTGCTGGAGGGCGACCGGGTGGCTGCTGGGGTACGGGTCGGCAGTG
Dxs1 AAAGGCAGGATCCTGCTGGAGGGCGACCGGGTGGCTGCTGGGGTACGGGTCGGCAGTG
616Bnormal

w5200 CAGTACTGCCTGACCGCCGCTCCCTGGTGCAGCGCCACGGCCTCAAGGTCACCGTCGCC
616B CAGTACTGCCTGACCGCCGCTCCCTGGTGCAGCGCCACGGCCTCAAGGTCACCGTCGCC
w5200normal CAGTACTGCCTGACCGCCGCTCCCTGGTGCAGCGCCACGGCCTCAAGGTCACCGTCGCC
Dxs1 CAGTACTGCCTGACTGCCCGCTCCCTGGTGCAGCGCCACGGCCTCAAGGTCACCGTCGCC
616Bnormal

w5200 GACGCGAGGTTCTGCAAGCCGCTGGACCACGCCCTGATCAGGAGCCTGGCCAAGTCCCAC
616B GACGCGAGGTTCTGCAAGCCGCTGGACCACGCCCTGATCAGGAGCCTGGCCAAGTCCCAC
w5200normal GACGCGAGGTTCTGCAAGCCGCTGGACCACGCCCTGATCAGGAGCCTGGCCAAGTCCCAC
Dxs1 GACGCGAGGTTCTGCAAGCCGCTGGACCACGCCCTGATCAGGAGCCTGGCCAAGTCCCAC
616Bnormal

w5200 GAGGTGCTCATCACCGTGGAGGAAGGCTCCATCGGAGGGTTCGGCTCGCACGTCGCCAG
616B GAGGTGCTCATCACCGTGGAGGAAGGCTCCATCGGAGGGTTCGGCTCGCACGTCGCCAG
w5200normal GAGGTGCTCATCACCGTGGAGGAAGGCTCCATCGGAGGGTTCGGCTCGCACGTCGCCAG
Dxs1 GAGGTGCTCATCACCGTGGAGGAAGGCTCCATCGGAGGGTTCGGCTCGCACGTCGCCAG
616Bnormal

w5200 TTCATGGCCCTGGACGGCCTTCTTGACGGCAAACCTCAAGgcaagtctcacactagctagc
616B TTCATGGCCCTGGACGGCCTTCTTGACGGCAAACCTCAAGgcaagtctcacactagctagc
w5200normal TTCATGGCCCTGGACGGCCTTCTTGACGGCAAACCTCAAGgcaagtctcacactagctagc
Dxs1 TTCATGGCCCTGGACGGCCTTCTTGACGGCAAACCTCAAGgcaagtctcacactagctagc
616Bnormal

w5200 tgctcggtcgcccctaatgataacgagagagagagagaaaaaactccgaactccatcttt
616B tgctcggtcgcccctaatgataacgagagagagagagaaaaaactccgaactccatcttt
w5200normal tgctcggtcgcccctaatgataacgagagagagagagaaaaaactccgaactccatcttt
Dxs1 tgctcggtcgcccctaatgataacgagagagagagagaaaaaactccgaactccatcttt
616Bnormal

w5200 agctgacaagtgatgaactcgacttttatttgggtgggtgcagTGGCGACCGCTGGTG
616B agctgacaagtgatgaactcgacttttatttgggtgggtgcagTGGCGACCGCTGGTG
w5200normal agctgacaagtgatgaactcgacttttatttgggtgggtgcagTGGCGACCGCTGG
Dxs1 agctgacaaaagtgatgaactcgacttttatttgggtgggtgcagTGGCGACCGCTGGTG
616Bnormal

w5200 CTTCTGACAGGTACATCGACCATGGATCGCCGGCCGATCAGCTGGCCGAGGCTGGGCTG
616B CTTCTGACAGGTACATCGACCATGGATCGCCGGCCGATCAGCTGGCCGAGGCTGGGCTG
w5200normal CTTCTGACAGGTACATCGACCATGGATCGCCGGCCGATCAGCTGGCCGAGGCTGGGCTG
Dxs1 CTTCTGACAGGTACATCGACCATGGATCGCCGGCCGATCAGCTGGCCGAGGCTGGGCTG
616Bnormal

w5200 ACGCCGTCACACATCGCCGCGTGGTGTCAACATCCTGGGGCAGAACAGGGAGGCTCTT
616B ACGCCGTCACACATCGCCGCGTGGTGTCAACATCCTGGGGCAGA---- GGAGGCTCTT
w5200normal ACGCCGTCACACATCGCCGCGTGGTGTCAACATCCTGGGGCAGAACAGGGAGGCTCTT
Dxs1 ACGCCGTCACACATCGCCGCGTGGTGTCAACATCCTGGGGCAGAACAGGGAGGCTCTT
616Bnormal

w5200	GCCATCATGGCAGTGCCTAACGCGTAGaacttgtgctgatcttggcctatagagatggtt
616B	GCCATCATGGCAGTGCCTAACGCGTAGaacttgtgctgatctggcctatagagatggtt
w5200normal	GCCATCATGGCAGTGCCTAACGCGTAGaacttgtgctgatcttggcctatagagatggtt
Dxs1	GCCATCATGGCAGTGCCTAACGCGTAGaacttgtgctgatctggcctatagagatggtt
616Bnormal	-----
w5200	gtacatTTTgtcgTTaactagagtgtctgaacttgggagattagtcttcttTggatgaaa
616B	gtacatTTTgtcgTTaactagagtgtctgaacttgggagattagtcttcttTggatgaaa
W5200normal	gtacatTTTgtcgTTaactagagtgtctgaacttgggagattagtcttcttTggatgaaa
Dxs1	gtacatTTTgtcgTTaactagagtgtctgaacttgggagattagtcttcttTggatgaaa
616Bnormal	-----
W5200	gtgtcgccggaacaacagttaccg
616B	gtgtcgccggaacaacagttaccg
W5200normal	gtgtcgccggaacaacagttaccg
Dxs1	gtgtcgccggaacaacagttaccg
616Bnormal	-----

Figure A1: Multiple sequence alignment of Dxs1 alleles from normal and mutant maize seedlings. Dxs1: B73 sequence. Intron sequences are in small cap and exon sequences are in all caps. Grey shading highlights the differences between sequences. Dashes (-) indicate gaps inserted to optimized the alignment

w5200	tattgacagcccggccggataggcaagcgcacgtaagggcttgttcggttattcccaat
616B	tcttgacagcccggccggtaggcaagcgcacgtaa-----
W5200normal	tcttgacagcccggccggtaggcaagcgcacgtaagggcttgttcggttattcccaat
Dxs1	tcttgacagcccggccggtaggcaagcgcacgtaagggcttgttcggttattcccaat
616Bnormal	tcttgacagcccggccggtaggcaagcgcacgtaagggcttgttcggttattcccaat
612A	tcttgacagcccggccggtaggcaagcgcacgtaa-----
612Anormal	tcttgacagcccggccggtaggcaagcgcacgtaa-----
612M	tcttgacagcccggccggtaggcaagcgcacgtaa-----
612Mnormal	tcttgacagcccggccggtaggcaagcgcacgtaagggcttgttcggttattcccaat
612N	tcttgacagcccggccggtaggcaagcgcacgtaa-----
612Nnormal	tcttgacagcccggccggtaggcaagcgcacgtaa-----
w5200	acacatggattggatgggattggaaaaaattatgaagaagtttgagctgttgggattca
616B	-----
w5200normal	acacatggattggatgggattggaaaaaattatgaagaagtttgagctgttgggattca
Dxs1	acacatggattggatgagattggaaaaaattatgaagaagtttgagctgttgggattca
616Bnormal	acacatggattggatgagattggaaaaaattatgaagaagtttgagctgttgggattca
612A	-----
612Anormal	-----
612M	-----
612Mnormal	acacatggattggatgggattggaaaaaattatgaagaagtttgagctgttgggattca
612N	-----
612Nnormal	-----
w5200	aacctccaatcccgcTcaatccacatggattgagagctaaccgaacaagccctaacac
616B	-----cac
w5200normal	aacctccaatcccgcTcaatccacatggattgagagctaaccgaacaagccctaacac
Dxs1	aacctccaatcccgcTcaatccacatggattgagagctaaccgaacaagccctaacac
616Bnormal	aacctccaatcccgcTcaatccacatggattgagagctaaccgaacaagccctaacac
612A	-----cac
612Anormal	-----cac
612M	-----cac
612Mnormal	aacctccaatcccgcTcaatccacatggattgagagctaaccgaacaagccctaacac
612N	-----cac
612Nnormal	-----cac

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w5200          gtcaaatttcagactgtcctcgttctcacggaagcgcgtagatTTTTCTGGAATCTTGAT
616B          gtcaaatttcagactgtcctcgttctcacggaagcgcgtagatTTTTCTGGAATCTTGAT
W5200normal   gtcaaatttcagactgtcctcgttctcacggaagcgcgtagatTTTTCTGGAATCTTGAT
Dxs1          gtcaaatttcagactgtcctcgttctcacggaagcgcgtagatTTTTCTGGAATCTTGAT
616Bnormal    gtcaaatttcagactgtcctcgttctcacggaagcgcgtagatTTTTCTGGAATCTTGAT
612A          gtcaaatttcagactgtcctcgttctcacggaagcgcgtagatTTTTCTGGAATCTTGAT
612Anormal    gtcaaatttcagactgtcctcgttctcacggaagcgcgtagatTTTTCTGGAATCTTGAT
612M          gtcaaatttcagactgtcctcgttctcacggaagcgcgtagatTTTTCTGGAATCTTGAT
612Mnormal    gtcaaatttcagactgtcctcgttctcacggaagcgcgtagatTTTTCTGGAATCTTGAT
612N          gtcaaatttcagactgtcctcgttctcacggaagcgcgtagatTTTTCTGGAATCTTGAT
612Nnormal    gtcaaatttcagactgtcctcgttctcacggaagcgcgtagatTTTTCTGGAATCTTGAT
    
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Figure A2: Multiple sequence alignment of MITE region in Dxs1 alleles from normal and mutant maize seedlings. Dxs1: B73 sequence. Intron sequences are lower case and exon sequences are upper case. Dashes (-) indicate the deletion of MITE transposable element sequences leaving a TAA footprint.





Amphimermis thezamica Sp. N. (Nematoda: Mermithidae) a New Species of Nematode from Georgia

By Oleg Gorgadze, Manana Lortkipanidze & Giorgi Bakhtadze

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Abstract- Female and male individuals of *Amphimermis thezamica* sp. n. (Nematoda: Mermithidae) have been described. The host organism is unknown. The material was collected from the soil of private gardening plot, located in village Tezami of Mtskheta-Mtianeti Region (East Georgia). A new species is characterized by the combination of the following features: amphids cup-shaped, less oval and with average size, vagina prolonged, S-shaped, round eggs with smooth surface and thick envelope, 56 papillae, arranged in three rows in genital part, long double spicule, intertwined in some sections. By morphological and morphometric data a new species is close to the group of *A. bogongae*, especially to *A. litoralis*. New species resembles *A. litoralis* by the shape of amphids, S-shaped vagina, ending of a tail and twisted spicule. It differs from *A. litoralis* by the length of vagina, structure of spicule and twisted parts, presented in its different sections, by the length of twisted and untwisted parts; by the shape of stoma.

Keywords: *amphimermis thezamica* sp.n., *nematoda*, *mermithidae*, *parasitic*, *Georgia*.

GJSFR-C Classification: FOR Code: 060899



Strictly as per the compliance and regulations of :



Amphimermis thezamica Sp. N. (Nematoda: Mermithidae) a New Species of Nematode from Georgia

Oleg Gorgadze ^α, Manana Lortkipanidze ^σ & Giorgi Bakhtadze ^ρ

Abstract- Female and male individuals of *Amphimermis thezamica* sp. n. (Nematoda: Mermithidae) have been described. The host organism is unknown. The material was collected from the soil of private gardening plot, located in village Tezami of Mtskheta-Mtianeti Region (East Georgia). A new species is characterized by the combination of the following features: amphids cup-shaped, less oval and with average size, vagina prolonged, S-shaped, round eggs with smooth surface and thick envelope, 56 papillae, arranged in three rows in genital part, long double spicule, intertwined in some sections. By morphological and morphometric data a new species is close to the group of *A. bogongae*, especially to *A. litoralis*. New species resembles *A. litoralis* by the shape of amphids, S-shaped vagina, ending of a tail and twisted spicule. It differs from *A. litoralis* by the length of vagina, structure of spicule and twisted parts, presented in its different sections, by the length of twisted and untwisted parts; by the shape of stoma. We present the list of species of the genus *Amphimermis*, distributed in Holarctic, with brief information on morphological characters, hosts and places of distribution.

Keywords: *amphimermis thezamica* sp.n., nematoda, mermithidae, parasitic, Georgia.

I. INTRODUCTION

Nematodes, united in the genus *Amphimermis* are polyphages; they are of economic importance as agents of biological control of harmful insects (Poinar & Welch, 1981; Chen & Yang, 1985). They parasitize on both, terrestrial and soil insects belonging to orders Orthoptera, Lepidoptera, Coleoptera, Hymenoptera and cause their death (Poinar, 1975; Rubtsov, 1977, 1978; Ipatieva & Pimenova, 1985; Poinar et al., 2006). Development of nematodes of this group takes place mainly in the soil. They differ from mermatids of other genera by a very long, twisted spicule and bent S-shaped long vagina (Rubtsov, 1978). More than 20 species are described from this genus worldwide. In Georgia, in particular in West Georgia, only one species of the genus *Amphimermis* - *Amphimermis lagidzae* Rubzov (Rubtsov, 1975) was registered so far. Morphological, anatomic and morphometric analysis of nematodes, isolated from soil has shown that the described nematode is a new form and it belongs to the genus *Amphimermis*.

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II. MATERIAL AND METHODS

Mermithid nematodes have been collected using the shovel from the soil of private gardening plot, located in village Tezami [41°53'2"C, 44°57'28" B] of Mtskheta-Mtianeti Region (East Georgia) in April of 2015. The nematodes were found at 20-25 cm depth from the soil surface. Adult female and male nematodes were observed alive and then killed by immersion in 60°C distilled water for 2 min, fixed in TAF (7 ml formalin, 2 ml triethanolamine, and 91 ml distilled water) (Poinar, 1975) and processed to glycerol by Seinhorst's method for taxonomic studies (Polozhentsev & Artyukhovski, 1963; Curran & Hominick, 1980). Drawings and pictures have been done using both alive and dead individuals of nematodes. Measurements were made using the light microscope (Motic-DMB1) on 4x, 10x, 40x and 100x magnification. Some small-size details of nematodes, significant for the diagnosis (stoma, amphids, etc.) were examined using the immersion objective of 100x magnification. Drawings and photos of nematodes mounted on slides were performed using a digital video camera Genius (G-Shot) DV 1110. Host of the nematode is unknown. The range of the characters is given following the mean value. The range of the characters is given following by the mean value. Type material deposited in the collection of the museum 16 of the Institute of Zoology of Ilia State University, Georgia

III. RESULTS

The description of the new species is based on the forms of nematodes that were taken in the East Georgia region in the village of Tezami. This is the second record of this kind in Georgia. New species of mermithid are morphologically different from *Amphimermis lagidzae* which was recorded in Georgia, as well as from all species united in this genus.

a) *Diagnosis of genus Amphimermis Kaburaki & Imamura, 1932*

= *Complexomermis* Filipjev, 1934, Diagnosis (Kaiser, 1991 emend.)

Mermithids are of medium and big size – from 13 to 260 mm. Body is thin and long, frontal part is significantly narrowed, while the posterior part is less narrowed. The six head papillae are arranged in the

same circle. Stoma opening is terminal; sometimes slightly displaced to the abdominal side. Stoma tube is long; it makes 60% of the body length. The distal part of adult, sexually mature forms does not extend beyond the inner head capsule. Amphids are big in size, located at the level of papillae or are bent backwards. Cavity of amphids is whether sharply prolonged or barrel-shaped. Cuticle is thick, with well-defined fibrous crossed filaments. Hypodermal chordae six in number. Vagina is S-shaped by shape, with three nodes and very long; very often the frontal part is widened. Vulva has a shape of an oblique fissure. Eggs numerous, of medium size and round, with thick envelope, without villi. Spicules long (from 1.0 mm to 3.6 mm), partly twisted. Mermithids have three orders of sexual papillae: middle papillae doubles in the region of sexual sphere. At the end of rounded tail under the cuticle they have uneven margins. Terminus of a tail of parasite and postparasite larvae is equipped with mucro.

b) Taxonomy

i. *Amphimermis thezamica* sp. n. (Figures 1- 4).

Type Host and locality: Host of the described nematode is unknown. It was collected from the soil of private gardening plot, located in village Tezami of Mtskheta-Mtianeti Region (East Georgia) [41°53'2"C, 44°57'28" B]. The species *Amphimermis thezamica* sp.n. 4 was named according to the place of collection.

Type material: slide of the holotype (adult female and male): Mtskheta-Mtianeti Region (East Georgia), village Tezami, April 2015 (the code villi. Tezami -2015 0-01); paratypes (18 females -16 slides; 7 males- 5 slides and 20 invasive larvae - 6 slides) have the same characters as the holotypes. Holotypes of a new species are preserved in the collection of the museum of the Institute of Zoology of the Ilia State University.

ii. Measurements

Allotype (female): length=205 mm; head diameter at the level of cephalic papillae: 68 (µm; body diameter at the level of the nerve ring: 144 (µm; maximum body diameter at vulva: 372 (µm; body diameter at the posterior end of trophosome: 253 (µm; body diameter at vulva: 405 (µm; distance from the anterior end to the nerve ring: 322 (µm; V%=47.7; length of vagina: 2.1

mm; width of vagina: 114 µm.

Female (paratypes; n=18): length=242±77 (71-282) mm; diameter of head at the level of cephalic papillae: 68±5 (64-79) (µm; body diameter at the level of nerve ring: 148±7 (133-152) µm; maximum body diameter at vulva: 390±24 (349-420) µm; body diameter at the posterior end of trophosome: 247±22 (228-260) µm; distance from the anterior end to the nerve ring: 360±24 (292-360) µm; distance from the end of trophosome to the end of tail 480±96 (315-580) µm;V%=47.5±3 (41.8-47.7); length of vagina: 2.2 ±0,8

(1.8-2.7) mm; width of vagina: 120±11 (100-140) µm.

Holotype (male): Length=35 mm; head diameter at the level of cephalic papillae: 60 µm; body diameter at the level of the nerve ring: 114 µm; maximum body diameter:178 µm; body diameter at the level of anus: 159 µm; distance from the anterior end to the nerve ring: 228 µm; length of the spicule: 1.773 mm; width of the spicule: 16µm.

Male (paratypes; n=7):Length=40±8 (29-54) mm; head diameter at the level of cephalic papillae: 62±3 (56-68) µm; body diameter at the level of the nerve ring: 112±4 (107-118) µm; maximum body diameter: 182±12 (151-188) µm; body diameter at the level of anus: 169±23 (132-195) µm; distance from the anterior end to the nerve ring: 232±3 (224-232) µm; length of the spicule: 1.770±0.4(1.668-1.776) mm; width of the spicule: 15±0.8 (14-16) µm.

Invasive larva: (n=20). Body length 1.1±0.1 (1-1.3) mm; maximum width 22±2 (19-22) µm; larva has a style, which length fluctuates from 25 to 26 µm. Body diameter at the level of the head papillae 10±1 (9-11) µm; at the nerve ring - 19±3 (15-19) µm; distance from the apical end of the head to the nerve ring makes 53±4 (53-57) µm, while from the rectum from 83±6 (79-86) µm length till the end of tail larva is spindle-shaped. It is characterized by the movement of plectoidnematodes. Anterior part of the body is wide, but is very narrow at the terminal part and ends with micro bulb-like tail. Tail terminus is used by the larva to attach to the substrate.

iii. Description

a. Morphology

Female. Nematode is of white colour. Body is thin and long. Body diameter increases at the 45.5 mm distance from the apical end of anterior part and then it is of nearly even width till the end of tail. Head is rounded and slightly flattened (Figures 1A, 2A,B). Diameter of the head capsule 9.2 µm. Cuticle is thick, it contains crossed fibrous threads; stoma is symmetrically situated at the end of the head. The 6 head papillae are well defined. Each papilla has 2-3 sensils. Amphids have shape of cup, are less oval and middle size (length of amphids 15.2 µm, width 9.6 µm). Hollows of amphids open behind the papillae at 15 and 19 µm distance. Stoma is narrow (1.5 µm wide); length of stoma 19.4 µm. Stoma is without collar. Oesophagus 5.4mm long, diameter 5.5 µm. Channel of vagina starts obliquely. It is long1.5 mm. (Figures 1C, 2C) cylindrical and bent S-shaped, nearly of even width along the whole length. Eggs round (length fluctuates from 83 to 87 µm, width 79-87 µm); with smooth surface; envelope thick - 4.2 µm wide. Size of cells of trophosome fluctuates from 27 to 140µm. Lateral chordae narrow, size of chordae, located at dorsal and ventral sides fluctuates from 15 to 19µm. Tail is slightly conical (Figures 1B, 2D,E); its terminal part is rounded and unevenly immersed under the cuticle.

Male. In comparison with female, male is smaller in size. Oesophagus is 1.5 mm long. Long, twisted, double spicule is characteristic of the male (Figure 3D). It contains twisted and non-twisted parts (Figures 3B, C, D, E, 4C,E,F). Length of distal untwisted part of spicule is 117.5 μm (see Figure 3B), length of the next, twisted part is 428.7 μm , length of the central, untwisted part is 252.1 μm ; the next, twisted part is 663.2 μm long, and length of the proximal, untwisted part makes 312.4 μm . Terminal part of spicule is conical and its end is rounded. In the terminal part of spicule 2 notches are visible (Figure 3E), which is characteristic to this species. Cloaca diameter 159.6 μm . In postanal part 2 well defined nucleoli are visible (38 μm long and 30 μm wide), which are not found in other Amphimermitids. Sexual papillae are arranged in three orders; in preanal part 10 quite big papillae are situated medially, though on each of lateral parts 7-7 small papillae are presented. In postanal part 20 papillae are situated medially (among them the first, second, nineteenth and twentieth papillae are small in size, though the rest papillae are big in size) and on each of its lateral part 6-6 papillae are situated. Total 56 papillae are situated in the genital part. Size of spermatocytes fluctuates from 7 to 11 μm . Tail is 266.5 μm long. It is slightly conical (Figures 3C, 4D), rounded at the end.

c) *Diagnosis and Relationships*

By its morphological and morphometric characters a newly described species - *A. thezamica* sp. n. belongs to the genus *Amphimermis*. By the character of twisting of spicule and shape of amphids the nematodes, united in the genus *Amphimermis* are divided into four groups (Bacer & Poinar, 1994): 1. *volubilis* (*A. volubilis*); 2. *avoluta* (*A. avoluta*, *A. acridiorum*, *A. buraki*); 3. *bogongae* (*A. bogongae*, *A. maritime*, *A. litoralis*, *A. tinui*, *A. bonaerensis*, *A. mirabinda*); 4. *elegans* (*A. elegans*, *A. zuimushi*, *A. tongaensis*, *A. artyukhovskii*, *A. australoelegans*).

A new species, which is characterized by a twisted spicule, was compared with nematodes of all four groups. It turned out that, by morphological and morphometric characters it is the most close to the group of *A. bogongae*, especially to the species *A. litoralis*.

A. thezamica sp. n. Resembles *A. litoralis* by the shape of amphids (both species have cup-shaped amphids), S-shaped vagina and end of tail; (both *A. thezamica* sp. n. and *A. litoralis* are characterised by the uneven deepening under the cuticle at the end of the tail, twisted spicule and diametric dimensions of the body.

A. thezamica sp. n. differs from *A. litoralis* by the following characters: size of the body (in a new species male $L=40$ (29-54) mm, in *A. litoralis* – 57 (34-76) mm; in female – correspondingly 242 mm and 136

mm; by the diameter of the body - (in a new species diameter of the body is 182 (151-188) μm , though that of *A. litoralis* 267 (232-471) μm ; by the length of spicule - in a new species length of spicule is 1.773 (1.668-1.776) mm – in *A. litoralis* - correspondingly 3.200 (2.900-3.600) mm; by the length of twisted parts of spicule and their positioning: length of the first untwisted part of spicule in a new species is 117 μm ; in *A. litoralis* it is equal to 194 (41-336) μm . Length of the first twisted part of spicule in a new species is 428 μm , in *A. litoralis* – 647 μm ; length of a central untwisted part in a new species is 253 μm ; in *A. litoralis* – 302 μm ; length of the next twisted part in a new species is 663 μm , though in *A. litoralis* it is 646 μm long. Terminal part of spicule in a new species is untwisted (length 312.4 μm), though in *A. litoralis* it is twisted till the end. Vagina of the female of a new species is longer 1.5 mm, than that of *A. litoralis* (960 μm). Part of stoma, which from the apical part of head connects to the esophagus, is slightly widened in *A. thezamica* sp.n., though this is not a case in *A. litoralis*. Terminal part of a head is rounded in *A. litoralis*, though in a new species it is flattened. These two species differ from each other by the positioning of vulva towards the body: $V\%$ for *A. litoralis* is 50, though vulva of a new species is located slightly ahead ($V\%=47$). *A. thezamica* sp.n., and *A. litoralis* differ also by the thickness of the cuticle: thickness of cuticle at the head capsule - 13 μm ; at vulva - 12 μm ; at the end of tail - 51 μm ; in a new species these indexes are correspondingly 30, 38 and 41 μm . Comparison has shown that despite the fact that by morphological and morphometric characters new species, described by us, is quite close to *A. bogongae*, by the key diagnostic characters it differs from the latter: by the length of vagina, structure of spicule and presence of twisted segments at different places; by the length of twisted and untwisted parts; by the shape of stoma and thickness of the cuticle. Based on the above we consider that the mermithid, described by us can be regarded as a new species.

IV. DISCUSSION

Female individual of *A. thezamica* sp. n. is of white colour. Body is thin and long. At the 45.3 mm distance from the apical end of the body its diameter increases, and then body is nearly of the same width till the tail. Head is rounded and slightly flattened. Cuticle is thick and it contains crossed fibrous threads. Stoma is situated symmetrically at the end of the head. Number of head papillae – 6. Amphids are cup-shaped. Stoma is narrow, without collar. Channel of vagina starts obliquely. It is long, cylindrical, bent S-shaped, and has the same width along the whole length. Eggs are round and have smooth surface. Tail is slightly conical. Its terminal part is rounded and unevenly deepened under the cuticle.

Male individual is smaller in size than the female one. Male is characterized by the long, double spicule. It is composed of twisted and untwisted parts. The 2 notches are well defined at the end of the spicule, which is characteristic to this species. Total 56 papillae present in the genital part. Tail is slightly conically, rounded at the end.

Despite the resemblance of morphological and morphometric characters of *A. thezamica* sp.n. with those of species, belonging to the genus *Amphimermis*, by the structures of spicule and presence of twisted segment at its different parts, length of twisted and untwisted parts, shape of stoma, a new species differs from Holarctic species, united in this genus, such as: *A. zuimushi* Kaburaki & Imamura, 1932; *A. acridiorum* Baker & Poinar, 1994; *A. artyukhovskii* Artyukhovski & Karchenko, 1965; *A. australoelegans* Baker & Poinar, 1994; *A. avoluta* Rubtsov & Koval, 1975; *A. bogongae* Welch, 1963; *A. bonaerensis* Miralles & Camino, 1982; *A. buraki* Baker & Poinar, 1994; *A. elegans* (Hagmeier, 1912) Welch, 1963; *A. lagidzae* Rubtsov, 1975; *A. litoralis* Artyllkhovski & Karchenko, 1971; *A. longiscapus* Rubtsov, 1976; *A. maritime* Rubstov, 1971; *A. mirabinda* Baker & Poinar, 1994; *A. mongolica* Rubtsov, 1976; *A. polaris* Spiridonov & Lantsov, 1996; *A. tinyi* Nickle, 1972; *A. tongaensis* Spiridonov, 1987; *A. volubilis* Rubtsov & Koval, 1975;

A. zuimushi - parasitizes on Lepidoptera: Noctuidae-*Agrotis infusa* (Boisd.), Pyralidae- *Chilo simplex* (Butler). Cavity of amphidin a lateral position is extended along the longitudinal axis of the body. It differs from all genera by a very long and stretched spicule and long, S-shaped bent vagina, Spicules 1.02-1.45 mm long. Distribution: Japan.

A. acridiorum- parasitizes on Orthoptera: Acrididae-*Phaulacridium vittatum* (Sjostedt); *Oedaleus australis* (Saussure); *Chortoicetes terminifera* (Walker)., Amphid opening posterior to lateral cephalic papilla. Vagina short and broad (in lateral view), vagina long and narrow (in lateral view). Distribution: Australia.

A. artyukhovskii - parasitizes on Lepidoptera: Lymantriidae - *Lymantria dispar* (L), Geometridae - *Operopthera brumata* (L)., Spicule 1.6-1.8 mm. Body length 26-53 mm. Tail conoid, rounded. Female body length 36-190 mm. Distribution: Voronezh region, (Russia).

A. australoelegans- parasitizes on Orthoptera: Acrididae-*Phaulacridium vittatum* (Sjostedt), *Chortoicetes terminifera* (Walker); Coleoptera : Scarabaedae - *Sericesthis* sp., Tail conoid, pointed terminus. Vagina long (1.5 mm), thin walled. Distribution: Australia.

A. avoluta - parasitizes on Coleoptera: Chrysomelidae-*Leptinotarsa decemlineata* Say. Proximal half of spicule untwisted (=avoluta group), vagina medium (0.5 mm). Distribution: Ukraine.

A. bogongae - parasitizes on Lepidoptera: Noctuidae-*Agrotis infusa* (Boisd.). Tail bluntly rounded, distance of proximal papillae from cloaca=20% of spicule length; medium sized amphids in relation to head diameter, thick-walled. Distribution: Australia.

A. bonaerensis -parasitizes on Orthoptera: Acrididae-*Laplatacris dispar* Rhen., Tail pointed, distance proximal papillae to cloaca≤spicule length. Vulva with flanges. Distribution: Australia.

A. buraki - parasitizes on Orthoptera:Tettigoniidae-*Conocephalus* sp. ; Body short (30 mm), spicule length 750-975 μm, terrestrial, amphids thick - walled, situated anterior to neck region with opening immediately posterior to lateral head papillae. Distribution: Australia.

A. elegans - parasitizes on Orthoptera: Acrididae: *Stenobothrus* sp. Tail bluntly rounded. Body length 195-260 mm. Distribution: Germany.

A. lagidzae - the host organism and male individuals are not known. Amphids situated in neck region with opening well posterior to lateral head papillae. Body 74 mm. Distribution: Georgia.

A. litoralis - unknown. Spicule long (2.9-3.6 mm) in relation to the body length (45-70 mm). Body length 136 mm, terrestrial. Distribution: Voronezh Region (Russia).

A. longiscapus - parasitizes on Lepidoptera: Lymantriidae: *Lymantria dispar* (L). Male individuals are not known. Amphids situated anterior to the neck region with opening, immediately posterior to the lateral head papillae. Body 43 mm. Distribution: Kirgizstan.

A. maritime - parasitizes, unknown. Spicule 2.2 mm. Body 53 mm. Junction of vagina and uterus at an obtuse anterior-ventral angle. Distribution: Primorsk Region (Russia).

A. mirabinda - parasitizes on Orthoptera: Acrididae: *Phaulacridium vittatum* (Sjostedt). Spicule 1.2-1.5 mm. Body 35-42 mm. large amphids in relation to the head diameter, thin-walled. Distribution: Australia;

A. mongolica - parasitizes, unknown. Amphids thin-walled, situated in neck region with opening well posterior to lateral head papillae. Distribution: Mongolia;

A. polaris- parasitizes on Diptera: Tipulidae -*Tipula (Savtshenkia) glaucocinerea* Lundtsr. Body length of male - 6.065 mm, length of spicule – 1.648 mm, length of females – 10.740 mm; usually on the tail terminus is developed 12 μm long cuticular appendix. Distribution: Taimyr, Dolgano-Nenetskyi Autonomous region.

A. tinyi - parasitizes on Odonata: Coenagrionidae: *Ischuraposita* (Hagen), *Annomalagrion bastatum* (Say). Male body short (11-17 mm), spicule 700-860 μm, aquatic. Female body length 30 mm, aquatic. Distribution: USA.

A. tongaensis - parasitizes, unknown. Diameter of amphid is greater than two thirds of head width. Female unknown. Distribution: Tonga.

A. volubilis - parasitizes on Coleoptera: Chryso-melidae: *Leptinotarsa decemlineata* Say. Spicule twisted entire length (= volubilis group), vagina long (0.7 mm).

V. ACKNOWLEDGEMENTS

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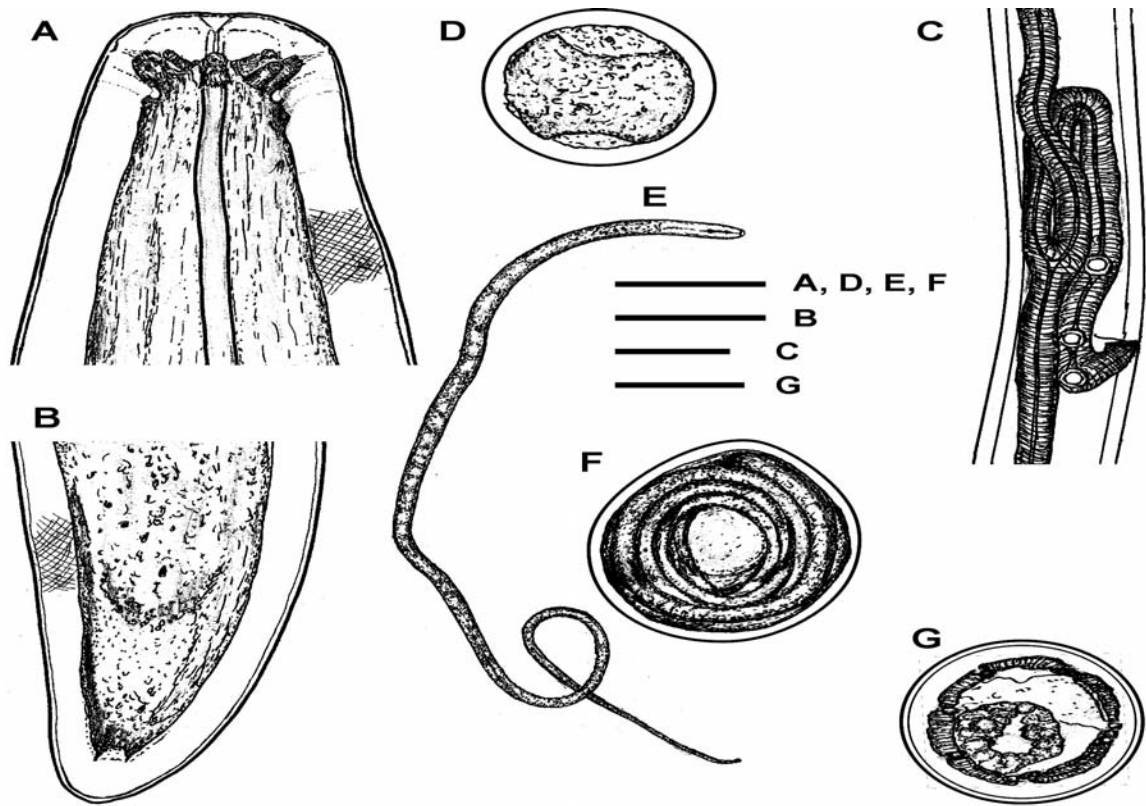


Figure 1: *Amphimermis thezamica* sp. n., female, holotype. A: Anterior region of the body, lateral view; B: Posterior end, lateral view; C: Vagina, lateral view; D: Uterine egg; E: stage -2 (infective), whole body; F: Last stage of larva in egg; G: Mid-body, cross section. Scale for A, D, E, F = 50 μ m; B = 100 μ m; C = 400 μ m; G = 200 μ m.

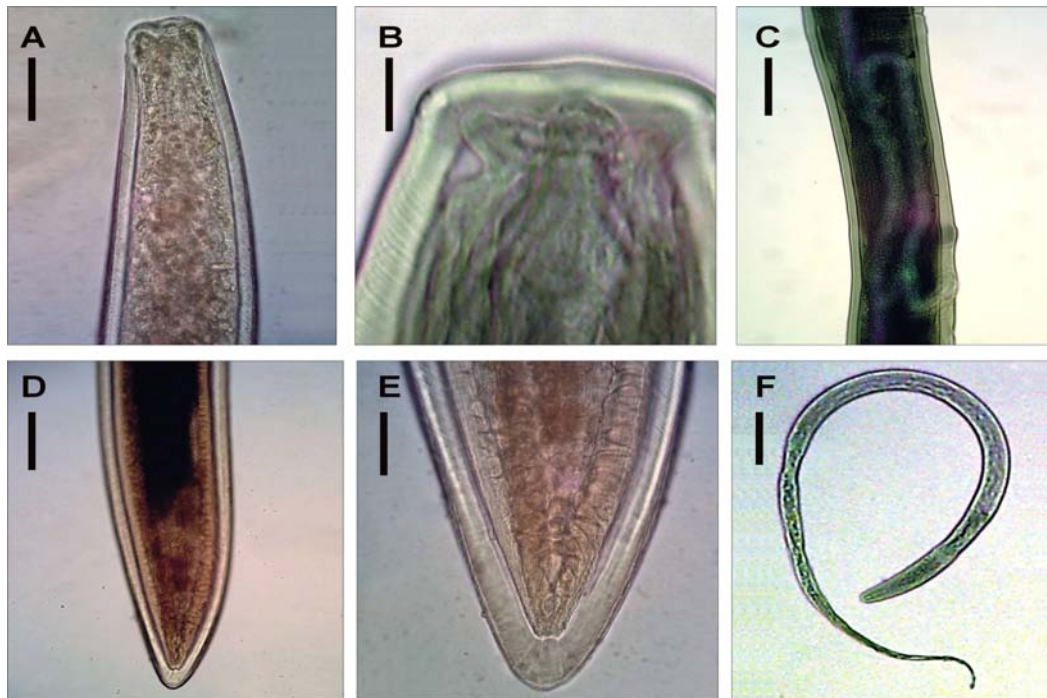


Figure 2: *Amphimermis thezamica* sp. n., female, holotype. A: Anterior region of the body, nerve ring, and pharyngeal tube, lateral view; B: Head, lateral view; C: Vagina, lateral view; D: Posterior region of the body, lateral view; E: Tail end; F: stage -2 (infective juveniles) whole body. Scale for A, E = 25 μ m; for B = 30 μ m; for C, D = 100 μ m; for F = 100 μ m.

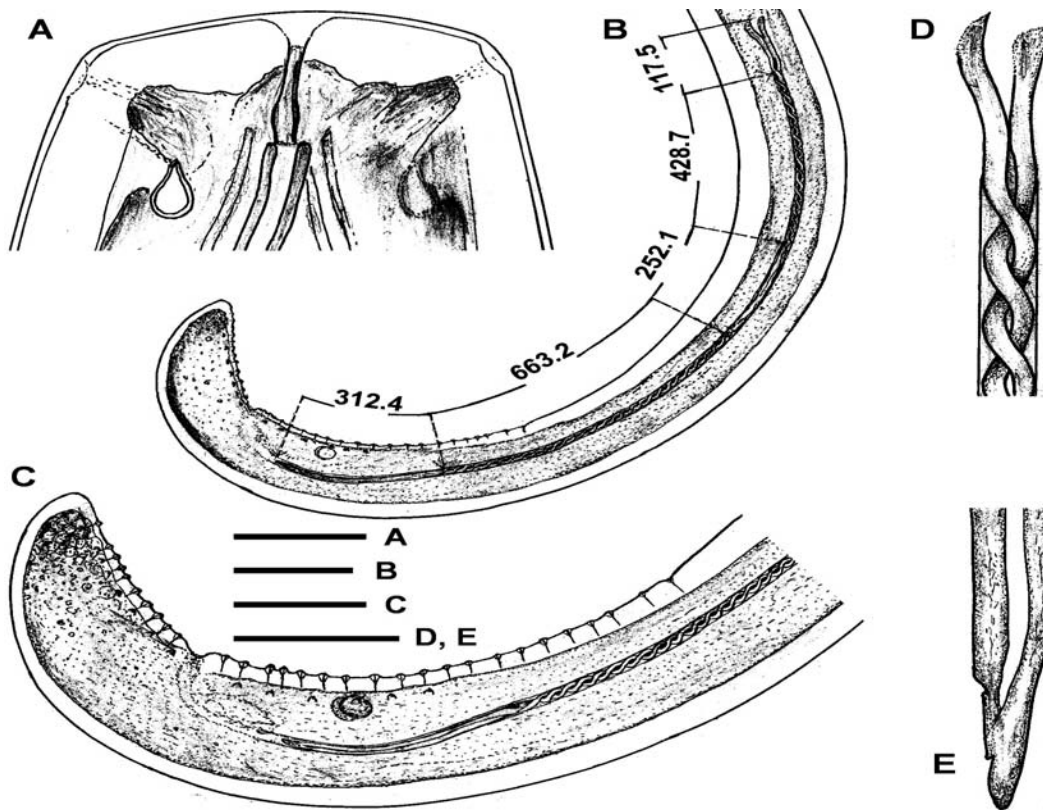


Figure 3: *Amphimermis thezamica* sp. n., male, holotype. A: Head, wide of the cuticle, lateral view; B: Tail, with whole spicula, lateral view; C: An enlarged tail with a half spicula; D: Non twisted and twisted head of spicule; E: Last part of non-twisted spicule. Scales for A=20 μ m; B=150 μ m; C=100 μ m; D,E=25 μ m.

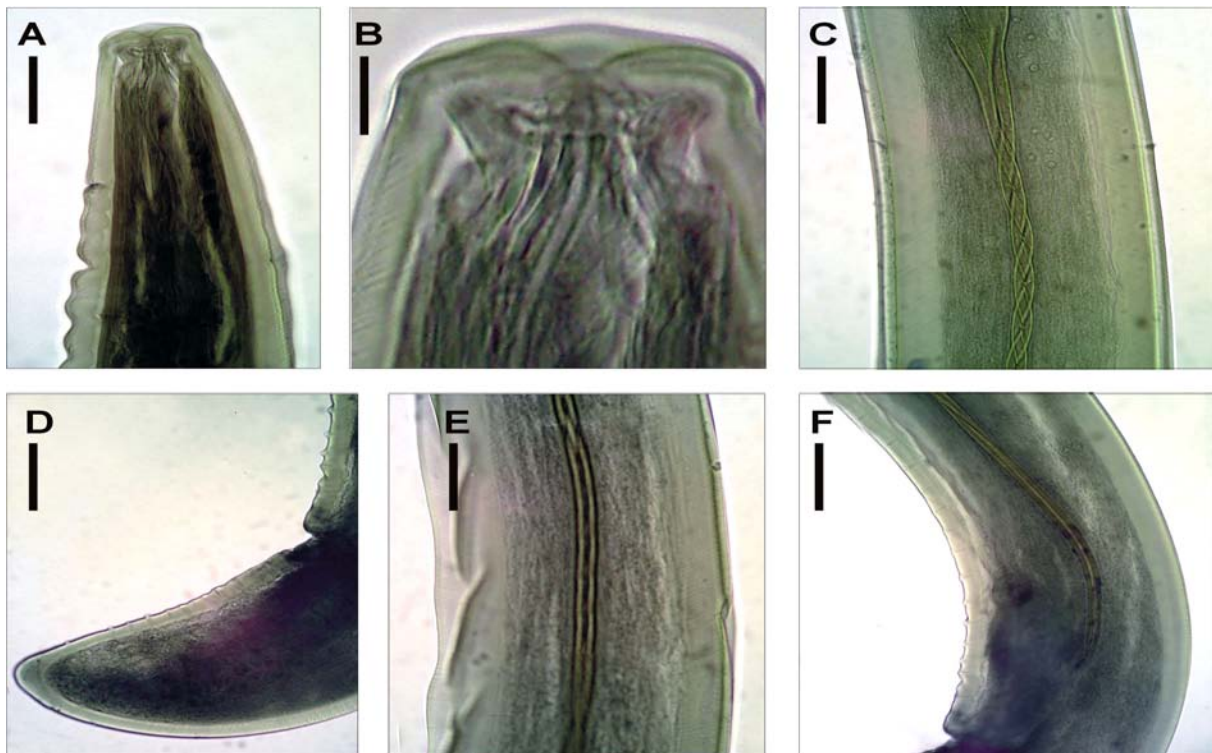


Figure 4: *Amphimermis thezamica* sp. n., male, holotype. A: Anterior region of body, lateral view; B: Head, lateral view; C: Non - twisted and twisted part of spicule head; D: Tail, lateral view; E: Middle part of non-twisted spicule; F: End of non-twisted spicule, lateral view; Scales for A,C,D,E,F=25 μ m; for B=30 μ m.

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Effects of Common Bean Varieties and Densities Intercropped with Rice on the Performance of Associated Components in Kaffa and Benchi Maji Zones, Southwestern Ethiopia

By Mitiku Woldesenbet & Getachew Mekonnen

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Abstract- Intercropping of cereals and legumes is important for sustainable food production in the tropics aimed at minimizing risks associated with monoculture. A field experiment was conducted at kuja site in Guraferda Woreda of Bechi Maji zone and Gojeb site in Ghimbo woreda of Kaffa zone, southwestern Ethiopia, during 2016 cropping season to determine the effects of the density and varieties of common bean intercropped with rice on growth, yield components and yield of the associated crops and productivity of the system. Rice variety 'NARICA-4' was intercropped with three varieties of common bean (Red Wolaita, Awash Melka and Nasir) in a factorial combination of *three* population densities of 25% (62,500 plant ha⁻¹), 50% (125,000 ha⁻¹) and 75% (187,500 plants ha⁻¹) of the recommended population density along with sole crops of the respective varieties of common bean and rice in randomized complete block design with three replications.

Keywords: *economic feasibility, land equivalent ratio, gross monetary value and productivity.*

GJSFR-C Classification: *FOR Code: 060799*



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Mitiku Woldesenbet ^a & Getachew Mekonnen ^o

Abstract- Intercropping of cereals and legumes is important for sustainable food production in the tropics aimed at minimizing risks associated with monoculture. A field experiment was conducted at kuja site in Guraferda Woreda of Bechi Maji zone and Gojeb site in Ghimbo woreda of Kaffa zone, southwestern Ethiopia, during 2016 cropping season to determine the effects of the density and varieties of common bean intercropped with rice on growth, yield components and yield of the associated crops and productivity of the system. Rice variety 'NARICA-4' was intercropped with three varieties of common bean (Red Wolaita, Awash Melka and Nasir) in a factorial combination of *three* population densities of 25% (62,500 plant ha⁻¹), 50% (125,000 ha⁻¹) and 75% (187,500 plants ha⁻¹) of the recommended population density along with sole crops of the respective varieties of common bean and rice in randomized complete block design with three replications. The results of the study showed that days to 50% heading, days to 90% maturity, were significantly affected by common bean density. The shortest days to 50% heading (95.58 days) and the shortest days to 90% maturity (120.33 days) of the associated rice were recorded at 75% planting density of common bean and significantly increased with the decrease in density of common bean to 25%. The plant height of rice was significantly affected by the main effect of variety, density and their interaction. Accordingly, the highest plant height (78 cm) was observed when Nasir variety was intercropped with rice at 75% planting density. It was observed that either the main effect or the interaction of common bean varieties and planting densities were not significant on the grain yield and harvest index of rice. Though it was not significant, the highest grain yields (3042 and 2855 kg ha⁻¹) were obtained from intercropping Awash Melka variety at 50% planting population, respectively. It was observed that the highest dry biomass (12556 kg /ha) of the rice was recorded when intercropped with common bean at 25% planting density and both parameters decreased significantly with increase in planting density of common bean to 75%. The main effect of common bean varieties had significant (<0.05) effect on dry bio mass and highly significant (p<0.01) effect on days to 90% maturity, leaf area index, plant height, number of seed per pods hundred seed weight and harvest index of common bean. The shortest days to 90% maturity (83.6 days) and the highest plant height (53.00 cm) were recorded for the Nasir variety. Conversely, the highest plant height (57.33cm) and the highest number seeds

per pod (4.7) were recorded in responses to 25% planting density and significantly increased as the bean planting density increased to 75%. The highest grain yield (1842 kg /ha) was recorded for variety Awash Melka at 75% planting density. The highest total LER of 2.38 and GMV of 30,883 ETB/ha were recorded when rice was intercropped with bean variety Awash Melka at planting density of 75%. Therefore, based on the above agronomic and economic evaluations, rice (100%) intercropped with common bean variety Awash Melka at planting density of 75% of the common bean can be recommended for intercropping of rice with common bean in the study area. However, the experiment has to be repeated across over seasons with consideration of farmers preference of the common bean varieties to reach at conclusive recommendation.

Keywords: economic feasibility, land equivalent ratio, gross monetary value and productivity.

I. INTRODUCTION

Poor soil fertility management, poor crop husbandry and effects of climate change are the major challenges and contribute for low crop productivity. Agro-ecological intensification of land use is a prerequisite for increased agricultural productivity, natural resource conservation and sustainable development (CCRP, 2009). The limited land areas are facing pressure to meet basic demands, such as food, since most growers own very small plots of land, especially in Africa (Rezaei-Chianeh *et al.*, 2011). In view of this, there is need for increased production in small areas through intercropping, which utilize common limiting resources better than the species grown separately, as an efficient resource use method (Ghosh *et al.*, 2006; Sobkiewicz, 2006).

The bulk of agricultural output in Ethiopia comes from 13.3 million small scale subsistence households, each owning, on average about 0.93 ha of land and produces a number of different food and cash crops (CSA, 2008). Tef, maize, wheat, rice and sorghum are among the cereal crops used to be the staple food crops and target of most of the food security programs in Ethiopia (CSA, 2014). Rice was introduced and evaluated initially at different areas of Ethiopia such as Gambella, Pawe, and Fogera in the late 1960s.

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However, attention was not given to rice research prior mid 1990s. Since 1990, seven upland rice varieties including two NERICAs (New Rice for Africa) have been released. The average productivity of these varieties ranges from 2.5 to 4 t/ha on farmers fields (Wolelaw, 2005). Ethiopia has different rice agro-ecologies that can grow rain fed upland rice, rain fed lowland rice, and irrigated rice with a total potential land mass of 1 million hectare (Sewagegne, 2011). However, this yield is very low as compared to the world production due to different constraints among which soil fertility problem is the first.

The current trend in global agriculture is to search for highly productive, sustainable and environmentally friendly cropping systems (Crew and Peoples, 2004). With increasing economic and environmental costs associated with fertilizer use, the need for low-input agro-ecological systems is rising (Meighen and Marney, 2012). Traditional agriculture, as practiced through the centuries, has always included different forms of intercropping (Lithourgidis *et al.*, 2011). As in most tropical countries, in Ethiopia, traditional cropping systems are based on resource poor farmers' subsistence requirements, and are not necessarily the most efficient ones (Tesfa *et al.*, 2002). In southwestern Ethiopia, small scale farmer uses a combination of crops grown on the same land in such a way that cereals, pulses, and oil seeds are represented. However, this cereal-legume intercropping study was not scientifically justified for rice in southwestern Ethiopia, though the area is potential area for rice production. Therefore, this study was initiated with the objective of determining the effects of the varieties and densities of common bean intercropped with rice on performance of the components.

II. MATERIALS AND METHODS

a) Description of the Study Site

The experiment was conducted in two locations namely, Ghimbo district Gojeb site in Kaffa Zone and Guraferda district Kuja site in Bench Maji Zone, Southwestern Ethiopia, during 2016 main cropping season. These sites were selected because of that they are the most potential areas of southwestern Ethiopia. The study sites at Gojeb and Kuja are located at 8° 06' N, 36°29' E, 1490 m.a.s.l. and 9° 07' N, 37° 35' E, 1238 m.a.s.l., respectively. The rainfall pattern of these areas is characterized by bimodal distribution with small rainy season *belg* (March-June) and main rainy season *meher* (July-November).

b) Description of Experimental Materials

An improved variety of rice named as *NERICA-4* (WAB-450-IB-P-9/1), which is currently grown extensively by the model farmers in the study areas, was used as a test crop. The variety was released in 2006 by the Pawe

Agricultural Research Center for its high yield and promising agronomic performances (MoARD, 2007).

c) Treatments and Experimental Design

The treatments consisted of planting of rice at the density of 100% of sole population with three common bean varieties (Red Wolaita, Awash Melka and Nasir) at the density of 25% (62500 plants ha⁻¹), 50% (12500 plants ha⁻¹), and 75% (187500 plants ha⁻¹). The sole rice was planted at spacing of 70 cm between rows by drilling. The sole common beans were planted at spacing of 40 x 10 cm (250000 plants ha⁻¹), respectively. For intercropping, common bean was planted inside rice rows at intra-row spacing of 7 cm, 11 cm and 21 cm representing 75%, 50% and 25% of sole population density of common bean, respectively. The experiment was laid out in a Randomized Complete Block Design (RCBD) in factorial arrangement in three replications. There were sole crop rice and the three common bean varieties.

The plot size and spacing of the experiment for sole common bean was 4.2 m length, 40 cm inter-row spacing, 10 cm intra- row spacing with a gross plot size of 4.2 m x 0.4 m x 6 = 10.08 m² and the central four rows of three meters length (4 rows x 0.4 m x 3 m = 4.8 m²) were harvested while for the intercrops four rows inside the rice with a plot size of (4 rows x 0.70 m x 3 m = 8.4 m²) were harvested. Sole rice was planted in 4.2 m length, 70 cm inter rows spacing by drilling and with 6 rows and had a gross plot size of 4.2 m x 0.70 m x 6 rows = 17.64 m² and three central rows of three meters (3 rows x 0.70 m x 4.2 m = 8.82 m²) were harvested both from sole and intercropped rice.

d) Experimental Procedure

Land preparation was done in mid June 2016 using daily labour and the rice seeds were sown in rows spaced 20 cm apart by hand drilling at the seed rate of 100 kg ha⁻¹. The sources of N and P were urea (46% N) and triple super phosphate (TSP, 46% P₂O₅), respectively. All P and half of the N fertilizer sources for the respective inorganic N and P₂O₅ treatments were applied at planting. The remaining half of the inorganic N fertilizer was applied at tillering stage by side drilling. Weeds were removed manually three times *i.e.* at early tillering, maximum tillering and booting stages. No insecticide or fungicide was applied as there was no serious incidence of insect pests or diseases. Harvesting was done manually using hand sickles. The harvested product was sun-dried to a constant weight and threshing and winnowing were done subsequently.

e) Soil Analysis

Prior to planting, surface (0 - 30 cm) soil samples, from five spots across the experimental fields, were collected in a zigzag pattern, composited, and analyzed for soil physico-chemical properties and the

results are depicted in Table 1. The soil physico-chemical analysis of the study areas revealed that the soils of the experimental field were clay and clay loam in texture both at Gojeb and Kuja, respectively. The results also indicated that the soil of Gojeb and Kuja are moderately and slightly acidic with pH of 6.31 and 5.66,

respectively. The soils have medium organic carbon (1.46) and total N (0.09%) at Kuja and low organic carbon (0.99) and total N (0.06%) at Gjebo. Available P is low both at Kuja (6.30 ppm) and Gojeb (5.90 ppm) (Table 1).

Table 1: Physico-chemical characteristics of soil of the experimental sites

Soil parameters	Districts	
	Gojeb	Kuja
Textural composition (%)		
Sand	22.00	16.30
Silt	25.00	24.80
Clay	53.00	58.90
Textural class	Clay	Clay loam
pH	5.66	6.31
Organic Carbon (%)	0.99	1.46
Total N (%)	0.06	0.09
Available P (mg/kg)	5.90	6.30
CEC (cmol/kg)	16.22	23.41

f) Data Collection

i. Rice Component

Days to heading were recorded when the ears or panicles were fully visible on 50% of the plants from each plot by visual observation and days to physiological maturity were recorded when 90% of the plants reached maturity in each plot, i.e. when grains were difficult to break with thumb nail. Number of productive tillers m² was counted from two random 1m X 1m areas (5 rows of 1m length) within the net plot area at physiological maturity and the average was recorded as number of productive tiller m². Plant height (cm) was determined from measurements of 10 randomly pre-tagged mother shoots from ground level to the top of the spike excluding the awns at physiological maturity. Likewise, the spikes in the pre-tagged 10 plants were collected and the total grains were counted to record the number of grains per spike. Thousand grains were counted in each plot using electronic seed counter from a bulk of threshed grain and their weight was measured using a sensitive balance at harvest and the weight was adjusted to 12.5% moisture content.

The total aboveground dry biomass yield including straw and spikes of plants in a net plot area was measured using spring balance after sun drying to a constant weight. Then threshed and the grain yield per net plot was weighed and adjusted to 12.5% moisture content. Harvest index was calculated as the ratio of grain yield to total aboveground dry biomass and expressed in percentage.

g) Common Bean Component

i. Phenological and Growth Parameters

Days to 50% flowering, 90% physiological maturity, Leaf area, Number of primary branches, Dry biomass, Stand count, Number of pods per plant, Number of seeds per pod, Hundred seed weight (g), Grain yield (kg/ha) and Harvest index.

ii. Productivity and Economic Evaluation of the Intercropping System

Productivity of the intercropping system was determined by calculating the Land Equivalent Ratio (LER) (Willey, 1979). Land Equivalent Ratio (LER) was a relative land area required as sole crop to produce the same yield as an intercrop system and calculated as

$$LER = \frac{Y_{ab}}{Y_{aa}} + \frac{Y_{ba}}{Y_{bb}}$$

Where Y_{ab} is yield per ha of rice in intercrop with common bean; Y_{aa} yield per ha of sole rice; Y_{ba} were grain yield per ha of common bean in intercrop with rice; Y_{bb} was grain yield per ha of sole common bean. LER values >1.00 indicate an agronomic advantage of intercropping over sole cropping.

The economic evaluation was done using Gross Monetary Values (GMV) as described by Willey (1979). To calculate the GMV of component crops, the prevailing prices at local market (18.00 Birr/kg at Gojeb

and 19.50 Birr/kg at Kuja for rice and 8.0 Birr per kg for common bean at Gojeb and 18.00 Birr/kg) were used.

h) Statistical Data Analysis

The agronomic data were subjected to analysis of variance (GLM procedure) using SAS software program version 9.2 (SAS Institute, 2003). Homogeneity of variances was evaluated using the F-test as described by Gomez and Gomez (1984) and since the F-test has showed homogeneity of the variances of the two locations for most of the agronomic parameters, the average data analysis was used for the two locations. The Fisher's protected least significant difference (LSD) test at 0.05 probability level was employed to separate treatment means where significant treatment differences existed.

III. RESULTS AND DISCUSSION

a) Rice Component

i. Rice Crop Phenology

The result of this study showed that the main effect of planting density of common bean had a highly significant ($P < 0.01$) effect on days to 50% heading and days to 90% maturity of rice crop.

ii. Rice Growth Parameters

The result of this study indicated that the main effects of varieties and planting density of common bean as well as the interactions had no significant effect on leaf area and leaf area index of rice crop. This result is in agreement negatively with the finding of Wogayehu (2005) who reported that the leaf area index of maize was significantly ($P < 0.05$) affected by intercrops of the associated bean varieties. Though it was not significant, the highest leaf area (1593 cm²) of rice was obtained

when Nasir variety was intercropped with rice compared to Awash Melka variety (1585 cm²) and Red Wolaita variety (1424 cm²) (Table 3). This might be due to less competition between rice and the associated Nasir variety for growth factors as it was early in maturity as compared to the two varieties, and may have required less nutrients during growth than the other variety.

The result of this study also indicated that Common bean variety and planting density had a highly significant ($P < 0.01$) effect and the interaction had a significant ($P < 0.05$) effect on plant height of rice. As a result, the highest plant height (78 cm) was obtained when rice was intercropped with common bean variety Nasir at planting density of 75% while the shortest plant height (64.7 cm) was obtained when common bean variety Red Wolaita was intercropped with rice at planting density of 25% (Table 2). In general as the planting density of common bean increased, the height of the associated rice increased indicating increased competition from the associated common bean for the limited resources. This result is in agreement with previous studies conducted by Sarma (1994) on *Sesamum indicum* which indicated that in narrow spacing plants compete more for available resources especially for light and result in more height than widely spaced plants. As sesame plants compete for light, high populations grow taller and faster than low populations (Langham, 2007). Unlike to the results of the present study, Demesew (2002) on maize/bean intercropping, Sisay (2004) on sorghum/green gram intercropping and Wogayehu (2005) on maize/common bean intercropping reported that the plant height of the cereal component was not significantly affected by the associated legume components.

Table 2: The average results of the interaction effect of varieties and planting density of common bean on plant height (cm) of rice intercropped with common bean

Common bean Varieties (V)	Planting Density of Common bean (PD)		
	25%	50%	75%
Red Wolaita	64.7 ^d	74.6 ^{ab}	74.7 ^{ab}
Awash Melka	74.7 ^{ab}	71 ^{bc}	65 ^d .1
Nasir	72.2 ^{cd}	69.7 ^c	78 ^a
Intercrop mean	74.3		
Sole rice mean	261.3		
	V × PD	Sole vs Intercrop	
LSD _(0.05)	4.88	NS	
CV (%)	4.7	6.6	

NS= Non-significantly different at 5% probability level; Means followed by the same letters are not significantly different at 5% level of significance according to LSD test.

b) Yield component and Yield of Rice

Form this study it was observed that the number of grains per spike and 100 seed weights (g) were not significantly affected by either the main effects of the varieties and planting density of common bean or their interaction. In agreement to the result of this study,

Tilahun (2002) reported non-significant effect of planting densities and planting patterns on maize 1000-kernel weight. Similarly, Wogayehu (2005) reported non-significant effect of the associated bean varieties on thousand-kernel weight of maize.

It was observed that, the number of grains per spike and 100 seed weights (g) decreased as the planting density of common bean increased from 25% to 75% (Table 1). This might be due to increased competition for growth resources from the associated common bean as its density increased.

The hundred seed weight recorded from intercropped rice was lower than that from sole crop

although the difference was not significant (Table 3). In line with this result, Tamado and Eshetu (2002) from intercropping of maize and haricot bean reported that 1000-kernel weight of maize was not significantly affected by the cropping system. On the other hand, Bandyopadhyay and De (1986) attributed the highest sorghum grain yield in intercrops to greater panicle and 1000-grain weight.

Table 3: The average result of main effects of varieties and planting density of common bean on number of grains per spike and 100 seed of rice intercropped with common bean and grown under sole crop

Treatment	Number of grains per spike	100 seed weight(g)
Common bean Varieties		
R + Red Wolaita	16.06	37.1
R + Awash Melka	13.00	38.2
R + Nasir	17.04	39.7
LSD (0.05)	NS	NS
Common bean Planting Density		
R + 25 % B	1.13	42.2
R + 50 %B	1.05	40.5
R + 75 %B	1.05	38.0
LSD (0.05)	NS	NS
CV (%)	17.6	6
Cropping system		
Intercrop	1.23	172.7
Sole crop	1.13	193.3
LSD(0.05)	NS	NS
CV (%)	6	5.8

NS= Non-significantly different at 5% probability level; Means followed by the same letters are not significantly different at 5% level of significance according to LSD test.

It was also observed that the main effect as well as interaction effect of planting density and variety of common bean did not significantly affect the grain yield and harvesting index of rice. This could be due to stronger competitiveness of the rice component as it

was planted early at full population as compared to common bean. Amare (1992) also found that intercropping different haricot bean varieties did not significantly affect maize grain yield.

Table 4: The average result of the main effects of varieties and planting density of common bean on grain yield (kg ha⁻¹), dry bio mass (kg ha⁻¹) and harvest index (%) of rice intercropped with common bean and grown under sole crop

Treatment	Grain yield (kg ha ⁻¹)	Dry biomass (kg ha ⁻¹)	Harvest index (%)
Common bean Varieties			
R + Red Wolaita	2811	12586	25.41
R + Awash Melka	3042	11968	26.51
R + Nasir	3011	11354	25.32
LSD(0.05)	NS	NS	NS
Common bean Planting Density			
R+ 25 % B	2732	12556 ^a	21.75
R + 50 % B	2855	11224 ^b	25.43
R + 75 % B	2733	11223 ^b	24.35
LSD (0.05)	NS	366	NS
CV (%)	7.6	3.8	7.7

Cropping system			
Intercrop mean	2622	11586	22.63
Sole crop mean	3125	11568	27.01
LSD(0.05)	NS	NS	NS
CV (%)	4.8	6.2	9.8

NS= Non-significantly different at 5% probability level; Means followed by the same letters are not significantly different at 5% level of significance according to LSD test.

However, among the intercropped common bean varieties, rice intercropped with variety Awash Melka gave the highest grain yield (3042 kg ha⁻¹) (Table 4). In agreement with this results of this study, Davis and Garcia (1987), Harwood *et al.* (2000), Tolessa *et al.* (2002) and Tolera *et al.* (2003) reported that planting beans in association had no appreciable effect on the grain yield of maize.

Dry biomass was highly significantly affected by the planting density of the associated common bean and non significantly affected by variety and the interaction of variety with planting density. The highest dry bio mass yield (12556 kg ha⁻¹) was obtained when rice was intercropped with common bean at plant population of 25% and the dry bio mass decreased as the planting density increase to 75% (Table 4). The reduction in dry biomass production in intercropped rice could be due to shading effect of the common bean during the early growth stage and inter-specific competition. Biscoe and Gallagher (1972) reported that the rate of dry bio mass production in crops depend up on the efficiency of the interception of photosynthetically active radiation (PAR).

The harvest index of rice was non significantly affected by the main effect of common bean varieties, population and their interaction. In this study, though the difference was statistically non-significant, relatively higher harvest index was recorded from sole crop (27.01%) than the intercropped rice (22.63%) (Table 4). In conformity to this Karikari *et al.* (1999) in Bambara groundnut + maize and Bambara groundnut + sorghum intercropping, reported significantly higher harvest indices for sole maize (0.599) and sole sorghum (0.386) than those in intercrops.

c) Common Bean Component

i. Crop Phenology

The main effect of common bean varieties and planting density had significantly affected days to 50% emergence, days to 50% flowering and days to maturity.

ii. Growth Parameters

The main effect of varieties of common bean was highly significant ($P < 0.01$) on leaf area, leaf area index, and plant height. Similarly the effect of planting density on leaf area index and plant height was highly significant and the interaction effect of varieties and planting density was significant on leaf area.

Common bean variety Nasir has the highest LAI (2.9) while the lowest was for variety Awash Melka (LAI=1.4). The variation in leaf area and leaf index observed due to varieties might be due to the difference in inherent characters of the varieties. Similarly, Wogayehu (2005) found significant difference in the leaf area index of common bean among the intercropped common bean varieties.

With regards to the common bean planting density, the highest leaf area index (2.9) was recorded when common bean was intercropped with rice at planting density of 75% and then the leaf area index decreased as planting density was decreased (Table 5). In line with this result, Sisay (2004) reported the highest LAI (2.321) when green gram was sown at its full rate (100%) while the lowest LAI (0.266) from the treatment containing the lowest rate (20%) of green gram. Although the difference is not statistically significant, higher leaf area index (2.53) of common bean was recorded from sole crop than the intercrop with LAI of 2.20 (Table 5).

Table 5: The average result of the main effects of varieties and planting density of common bean on days to maturity, leaf area index and plant height of common bean intercropped with rice and grown as sole crop

Treatment	Days to 90 % maturity	LAI	Plant height (cm)
Common bean varieties			
Red Wolaita	113.4b	2.5 ^{ab}	52.56 ^b
Awash Melka	117.3a	1.4 ^c	46.44 ^c
Nasir	83.6 ^c	2.9 ^a	53.00 ^b
LSD (0.05)	0.92	0.87	0.72
Common bean Planting Density			
R + 25%	98.1c	1.5 ^b	53.83 ^c
R + 50%	99.9b	2.2 ^{ab}	55.75 ^b
R + 75%	101.6a	2.9 ^a	57.33 ^a
LSD (0.05)	0.79	0.75	0.62
CV (%)	0.9	35.8	1.3
Inter-crop comparison			
Intercrop mean	116.3	2.03	55.64
Sole crop mean	100.0	2.53	55.83
LSD (0.05)	9.05	NS	NS
CV%	1.5 2.3	9.9	5.4

NS= Non-significantly different at 5% probability level; Means followed by the same letters are not significantly different at 5% level of significance according to LSD test.

With regards to the effect of planting density the highest plant height (57.33 cm) was recorded at common bean planting density of 75% and the height was significantly increased as planting density of common bean was increased (Table 5). The reduction in plant height of common bean with increase in planting density of common bean might be due to increased competition for growth resources such as radiation, soil moisture and nutrients with increased population of the intercropping system.

iii. Yield Components and Yield

In the present study, the main effect of common bean varieties had a highly significant ($P < 0.01$) effect on number pods per plant, number seeds per pod, hundred seed weight, grain yield and harvest index. Moreover, the effect of planting density was highly significant on the above parameters except on hundred seed weight and harvest index which was non-significant. The interaction effect of varieties and planting density of common bean was significant ($P < 0.05$) on number pods per plant and highly significant ($P < 0.01$) on grain yield of common bean.

The highest number of pods per plant (16.33) was obtained from Nasir common bean variety at planting density of 25% and the lowest number of pods per plant (3.0) was obtained from Awash Melka variety at planting density of 75% (Table 6). In general, the number of pods per plant decreased with the increase in planting density for all the varieties. This decrease in number of pods per plant at higher density could be attributed to increased competition among plants for growth factors. In line with this, in sorghum + mung

bean and sorghum + pigeon pea intercropping, Subramanian and Rao (1988) reported that decrease in grain number per unit area was responsible for lower grain yields in intercrops than in sole crops.

Above ground dry biomass (kg ha^{-1}) was significantly ($P < 0.05$) affected by the common bean varieties and highly significantly ($P < 0.05$) by the planting density. The highest above ground dry biomass (4466 kg ha^{-1}) was recorded for variety Awash Melka while the lowest (3121 kg ha^{-1}) was recorded for variety Red Wolaita (Table 6). With regards to the planting density, the highest above ground dry biomass (5292 kg ha^{-1}) was recorded at common planting density of 75% and it was decreased significantly with the decrease in planting density of common bean to 25% (Table 6). This decrease might be decrease in population of common bean in the intercropping system. In agreement with this result, Sisay (2004) recorded the highest above ground dry biomass from 100% green gram broadcast with sorghum followed by 80% green gram broadcast with sorghum. Similarly, intercropping of full density of barley with five planting densities of faba bean (100:12.5%, 100:25%, 100:37.5%, 100:50% and 100:62.5%) showed significant increment on dry biomass yield of intercropped faba bean from 653 kg/ha to 2494 kg/ha as plant density of faba bean increased from 12.5% to 62.5% (Getachew *et al.*, 2006). Though the difference was not statistically significant sole crop bean gave higher above ground dry biomass than the intercrop.

Table 6: Main effects of varieties and planting density of common bean on growth parameters and yield components of common bean intercropped with rice and grown as sole crop

Treatment	Number of branches	Dry bio mass (kg/ha)	No. of seed per pod	100 seed wt (g)	Harvest index (%)
Common bean varieties					
Red Wolaita	2.93	3121 ^b	5.0 ^a	22.6 ^c	17.7 ^{bc}
Awash Melka	3.36	4416 ^a	3.5 ^c	22.3 ^a	27.7 ^a
Nasir	2.78	3166 ^b	4.8 ^a	20.3 ^c	18.7 ^b
LSD (0.05)	NS	839.2	0.35	8.3	3.1
Common bean Planting Density					
R + 25%	2.98	2323 ^c	4.7 ^a	31.7	18.7
R + 50%	2.92	3292 ^b	4.6 ^a	30.8	20.4
R + 75%	2.93	5292 ^a	3.9 ^b	30.6	20.2
LSD (0.05)	NS	726.8	0.3	NS	NS
CV (%)	25	23	7.7	25.6	16.9
Intercrop mean					
	2.9	3491	4.47	23.67	23.3
Sole crop mean					
	3.2	4147	4.47	32.84	23.7
LSD (0.05)	NS	NS	NS	8.1	NS
CV (%)	4.1	19	6.6	7.7	5.2

NS= Non-significantly different at 5% probability level; Means followed by the same letters are not significantly different at 5% level of significance according to LSD test.

The highest number of seeds per pod (5.0) was obtained from common bean variety Red Wolaita while the lowest number of seed per pod (3.5) was for variety Awash Melka (Table 6). This difference in number of seeds per pod might be due to the inherent behavior of the varieties where Awash melka with large seed size had the smallest number of seeds per pod.

The main effects of common bean varieties, planting density and the interaction had a highly significant ($P < 0.01$) effect on grain yield of the intercropped common bean (Table 7). The highest grain yield (1842 kg ha⁻¹) was recorded for common bean variety Awash Melka at planting density of 75% while the

lowest grain yield (180 kg ha⁻¹) was recorded for bean variety Red Wolaita at 25% planting density (Table 7). In general variety Awash Melka gave higher grain yield than the other varieties possibly due to its large seed size. Moreover, the grain yield of common bean was increased as the planting density increased which might be due to the increased population. In agreement with this result, Sisay (2004) reported the highest seed yield of green gram when it was intercropped with sorghum with full rate and the lowest seed yields of green gram from intercrop combinations containing 20% and 40% populations of green gram.

Table 7: The interaction effect of varieties and planting density of common bean on grain yield (kg ha⁻¹) of common bean intercropped with rice

Common bean Varieties (V)	Planting Density of Common bean (PD)		
	25%	50%	75%
Red Wolaita	180 ^g	407 ^e	712 ^c
Awash Melka	546 ^d	1142 ^b	1842 ^a
Nasir	182 ^g	421 ^d	691 ^g
Intercrop mean			708
Sole common bean mean			845
	V × PD		Sole vs Intercrop
LSD _(0.05)	133		NS
CV (%)	10.8		13.7

NS= Non-significantly different at 5% probability level; Means followed by the same letters are not significantly different at 5% level of significance according to LSD test.

In this study, there was no significant difference in grain yield of the intercropped common bean sole cropped bean. However, this is in contrast to results by Huxley and Maigu (1978) who reported that in cereals and legumes intercropping system, the grain yield of the legume component declined, on average, by about 50% of the sole crop yield, where as the cereals yield was reduced by only 11 %. In agreement with this finding, Pal *et al.* (1993) reported that grain yield of sole cropped maize; sorghum and bean were significantly higher than the intercropped yield of these crop

d) *Productivity and Economic Evaluation of Rice and Common Bean Intercropping*

i. *Land Equivalent Ratio*

The agronomic productivity of this experiment was evaluated by calculating total land equivalent ratio (LER) by summing up the partial land equivalent ratio (PLER) of common bean and rice as described by Willey (1979). The main effect of variety and planting density of common bean and their interactions had a highly significant ($P < 0.01$) effect on total Land Equivalent Ratio (Appendix Table 7).

The highest total LER of 2.38 was recorded when rice was intercropped with bean variety Awash Melka at planning density of 75% while the lowest total

LER of 1.01 was obtained when bean variety Red Wolaita was intercropped at planting density of 25% (Table 17). In general, as the planting density of common bean increased, the total LER was increased indicating the importance of increased bean population in the intercropping system in improving the agronomic efficiency of the intercropping system. The value of LER above 1 indicates that the intercropping system utilizes the available growth resources more efficiently than sole cropped. In this study all of the total LER values of the intercropping system were greater than one, indicating that intercropping of rice and common bean was productive and had yield advantage over growing either rice or common bean in sole. This could be due to the efficient utilization of resources by the intercropped crops. In line with the results of this study, Ofori and Stern (1987) pointed out that the value of LER follow the density of legume component. Similarly, Eyob (2007) reported the highest LER of 1.94 from intercropping of faba bean with 75% plant density of sorghum. In contrast to this, Yesuf (2003) reported that the LER decreased with the increase planting density. Intercropping had higher mean LER (1.80) than sole crop (1.0) .

Table 8: Interaction effect of varieties and planting density of common bean on total land equivalent ratio of the intercropped rice and common bean

Common bean Varieties (V)	Planting Density of Common bean (PD)		
	25%	50%	75%
Omo	1.01 ^f	1.23 ^{de}	1.47 ^b
Ibbando	1.40 ^{bc}	1.92 ^a	2.38 ^a
Hawassa Bume	1.10 ^{ef}	1.26 ^{cde}	1.49 ^b
Intercrop mean	1.7		
Sole mean	1.0		
	V × PD	Sole vs Intercrop	
LSD _(0.05)	0.16	NS	
CV (%)	6.3	10.5	

NS= Non-significantly different at 5% probability level; Means followed by the same letters are not significantly different at 5% level of significance according to LSD test.

ii. *Gross Monetary Value*

The main effect of variety and planting density of common bean and their interactions had significant ($P < 0.05$) effect on Gross Monetary Value (Appendix Table 7). The highest Gross Monetary Value of 30,883 ETB/ha was obtained from common bean variety Awash

Melka intercropped with rice at planting density of 75% while the lowest Goss Monetary Value of 17,356 ETB/ha was obtained when bean variety Red wolaita was intercropped with rice at planting density of 25% (Table 18).

Table 9: Interaction effect of varieties and planting density of common bean on Gross Monetary Value of the intercropped rice and common bean

Common bean Varieties (V)	Planting Density of Common bean (PD)		
	25%	50%	75%
Red Wolaita	17356	19584	21821
Awash Melka	21623 ^b	26787 ^a	30883 ^a
Nasir	18884 ^{de}	20130 ^{bcd}	22454 ^b

Intercrop mean	28984
Sole rice mean	17169
Sole common bean	7493

NS= Non-significantly different at 5% probability level; Means followed by the same letters are not significantly different at 5% level of significance according to LSD test.

As it was in total LER, as the planting density of common bean increased, the Gross Monetary Value increased indicating the importance of increased bean population in the intercropping system in increasing the economic efficiency of the intercropping system as rice had high price (18 Birr/kg) than bean (5.6 Birr/kg) in the local market. In this study all the intercrops gave higher gross monetary value than either of sole rice or sole bean (Table 18). In agreement with the result of this study, Tesfay (2012) reported the highest GMV (46375.2 ETB/ha) from additive mixture of faba bean and wheat variety HAR 2501 at the seed rate of 75% while the lowest GMV (32222.1 ETB/ha) was obtained from intercropping of faba bean and variety HAR 2501 with seed rate of 50%.

IV. CONCLUSION

Intercropping is an important option for efficient utilization of resource especially under gradually decreasing cultivated land. Even though there is practice of intercropping cereals and legumes in the study area, the practice of intercropping rice with different densities of common bean is not common. Therefore, this study was initiated with the objective of determining the effect of density and varieties of common bean intercropped with rice on performance of the associated crops at Kuja and Gojeb, southwestern Ethiopia. The treatment consisted of three improved common bean varieties (Red Wolaita, Awash Melka and Nasir) and three planting densities (25%, 50% and 75%) of the recommended seed rate of sole common bean laid out in randomized complete block design (RCBD) in factorial arrangement and replicated three times.

The results of the study showed that days to 50% heading, days to 90% maturity, leaf area index, number of seeds per spike, 100 seed weight, grain yield, dry bio mass and harvest index were not significantly affected by the varieties of the associated common bean. However, days to 50% heading, days to 90% maturity and dry bio mass were significantly affected by common bean density. The shortest days to 50% heading (95.58 days) and the shortest days to 90% maturity (120.33 days) of the associated rice were recorded at 25% planting density of common bean and significantly increased with the increase in density of common bean to 75%. The highest dry biomass (12556 kg/ha) of the rice crops were recorded when intercropped with common bean at 25% planting density and both parameters decreased significantly with increase in planting density of common bean to 75%.

The interaction effect of varieties and planting density was highly significant ($P < 0.01$) on plant height of the rice intercropped with common bean varieties. The highest plant height (78 cm) was recorded when rice was intercropped with common bean variety Nasir at planting density of 25%. In general as the planting density of common bean increased the height of the associated rice was increased.

The main effect of common bean varieties had significant (< 0.05) effect on dry bio mass and highly significant ($p < 0.01$) effect on days to 90% maturity, leaf area index, plant height, number of seed per pods, hundred seed weight and harvest index. The shortest days to 90% maturity (83.6 days) and the highest plant height (53 cm) were recorded for the Nasir variety. Common bean variety Awash Melka was found to have the highest days to maturity (117.3 days), dry bio mass (4466 kg/ha), hundred seed weight (52.3 g) and harvest index (27.7%).

The main effect of common bean density had significant effects on days to 90% maturity, leaf area index, plant height, dry bio mass and number of seeds per pod. The shortest days to 90% maturity (83.6 days) and the lowest leaf area index (1.4) were recorded at common bean planting density of 25% and significantly increased as the bean planting density increased to 75%. Conversely, the highest plant height (57.33cm) and the highest number seeds per pod (5.0) were recorded in responses to 25% planting density and significantly increased as the bean planting density increased to 75%.

Days to 50% emergence, stand count difference, leaf area and number of pods per plant of common bean were significantly affected by the interaction effect of rice intercropped with common bean. Grain yield of common bean was also significant affected by the interaction effect of rice intercropped with common bean. The shortest days to 50% emergence (10 days) of common bean was recorded at the Awash Melka and Nasir at planting density of 25% and Red Wolaita at 75% planting density intercropped with rice crop. The highest leaf area (2.9 cm²) and the highest number of pods per plant (16.33) were recorded for varieties Awash Melka and Red Wolaita at 25% planting density intercropped with rice, respectively.

The main effect of variety and planting density of common bean and their interactions had a highly significant ($P < 0.01$) effect on total Land Equivalent Ratio. The highest total LER of 2.38 was recorded when rice was intercropped with bean variety Awash Melka at

planning density of 75% while the lowest total LER of 1.01 was obtained when bean variety Red Wolaita was intercropped at planting density of 25%. In general, as the planting density of common bean increased, the total LER was increased. As it was for LER, the main effect of variety and planting density of common bean and their interactions had significant ($P < 0.05$) effect on Gross Monetary Value. The highest Gross Monetary Value of 30,883 ETB/ha was obtained from common bean variety Awash Melka intercropped with rice at planting density of 75% while the lowest Goss Monetary Value of 17,356 ETB/ha was obtained when bean variety Red Wolaita was intercropped with rice at planting density of 25%). In this study all the intercrops gave higher gross monetary value than either of sole rice or sole bean. Therefore, based on the above agronomic and economic evaluations, rice (100%) intercropped with common bean variety Awash Melka at planting density of 75% of the common bean can be recommended for intercropping of rice with common bean in the study area. However, as this is one field experiment, the experiment has to be repeated over seasons with consideration of farmers preference of the common bean varieties to reach at conclusive recommendation.

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Evaluation of Glyphosate Toxicity on Arabian killifish, *Aphanius dispar* Collected from Southwestern Saudi Arabia

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Abstract- Glyphosate (Tiller 480 SL), which is used extensively to control and inhibit weeds in terrestrial and aquatic ecosystems, has been blamed of harming non-targeted species like fish. This current study evaluated the acute and chronic impacts of glyphosate on the Arabian killifish (*Aphanius dispar*) collected from Southwestern Saudi Arabia at different levels of biological organization including behavioral and histopathological responses. Glyphosate toxicity (96 h LC50) to *Aphanius dispar* was determined at 115.25 mg/l after exposure to 60, 90, 120, 150, 180, 210, 240 mg glyphosate/l. Fish during the 96 h displayed abnormal behavioral changes: erratic movements, hyperactivity, rapid opercula and mouth movements, surfacing, hypoactivity, exhaustion and mortality. While, exposure of fish to 1/4th of the 96-h LC50 for two weeks, the gills and liver organs displayed histopathological alterations.

Keywords: *glyphosate, fish toxicity, aphanius dispar, behavioral changes, histology, acute toxicity, environmental health.*

GJSFR-C Classification: FOR Code: 069999



Strictly as per the compliance and regulations of :



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Abstract- Glyphosate (Tiller 480 SL), which is used extensively to control and inhibit weeds in terrestrial and aquatic ecosystems, has been blamed of harming non-targeted species like fish. This current study evaluated the acute and chronic impacts of glyphosate on the Arabian killifish (*Aphanius dispar*) collected from Southwestern Saudi Arabia at different levels of biological organization including behavioral and histopathological responses. Glyphosate toxicity (96 h LC50) to *Aphanius dispar* was determined at 115.25 mg/l after exposure to 60, 90, 120, 150, 180, 210, 240 mg glyphosate/l. Fish during the 96 h displayed abnormal behavioral changes: erratic movements, hyperactivity, rapid opercula and mouth movements, surfacing, hypoactivity, exhaustion and mortality. While, exposure of fish to 1/4th of the 96-h LC50 for two weeks, the gills and liver organs displayed histopathological alterations. The gills revealed: epithelial uplifting, edema, hyperplasia associated with fusion of secondary lamellae, clubbing, thinning and shortening of secondary lamellae. Whereas, the liver exhibited: necrosis, deterioration, hypertrophy of hepatocytes with loss of determined peripheries, pyknotic nuclei, and cytoplasmic vacuolization with a foamy appearance.

In conclusion, this current study results revealed that glyphosate is very toxic leading not only to abnormal behavioral responses and tissue alterations, but might cause mass extinction of fish species. Therefore, glyphosate should be used carefully in/or near aquatic systems to avoid extinctions of life forms, particularly *Aphanius dispar*. Thence, protecting species diversity, which is a key issue for stability and resiliency of aquatic ecosystems.

Keywords: glyphosate, fish toxicity, *aphanius dispar*, behavioral changes, histology, acute toxicity, environmental health.

1. INTRODUCTION

Glyphosate herbicide (Tiller 480 SL), which is a broad spectrum non-selective herbicide is used excessively to control and inhibit a great variety of annual, biennial and perennial grasses, sedges, broad leaved weeds and woody shrubs in agricultural, industrial, urban, forestry, aquatic ecosystems, fish ponds, lakes, and canals (Cavas and Konen, 2007; Langiano and Martinez, 2008; Sani and Idris, 2016; Tsui and Chu, 2008). Glyphosate has been reported to be the most important herbicide ever developed (WHO,

1994) to particularly be applied in plant varieties that are genetically modified to be better able to tolerate glyphosate treatment during weed control (Langiano and Martinez, 2008), without affecting crops (Sani and Idris, 2016). Glyphosate is not only used directly to control noxious weeds in aquatic systems, but also reported to reach aquatic systems after application in agricultural fields, thus affect non-targeted organisms indirectly like invertebrate, fish and other life forms from the first level up higher the food chains (Jofre et al., 2013), thence reduce species diversity, community structure affecting the stability and resilience of aquatic ecosystems (Perez et al., 2011).

Glyphosate mode of action is through competitive inhibition of phosphoenolpyruvate (PEP) on the active site of 5-enolpyruvylshikimate-3-phosphatethensate, an enzyme involved in the biosynthesis of aromatic amino acids (phenylalanine, Tyrosine, and tryptophan), which are essential for protein synthesis (Mallory-Smith, 2013, Tu et al., 2001). Glyphosate and its formulations, especially those containing surfactants are considered hazardous to the aquatic environment, which showed higher toxicity to most aquatic organisms than the active ingredient itself, which has been classified into very slight to high toxicants to aquatic organisms including fish species (WHO, 1994; Perez et al., 2011) due to its higher solubility varying from 10,000 to 15,000 mg/l at 25oC (Nwani et al., 2013). Glyphosate and its formulations acute and chronic effects on aquatic organisms including fish species have been reported to involve behavioral, histopathological, biochemical, and physiological changes (Langiano and Martinez, 2016; Jiraungkoorskul et al., 2002; Thanomsit et al., 2016), reflecting slight to severe concentration-related alterations over a short period of time (Perez et al., 2011). Several studies on fish species have reported variations for the acute toxicity of glyphosate concentrations of which: 10 mg/l for Asian sea bass, *Lates Calcarifer*, 13.69 mg/l for the Neotropical fish, *Prochilodus lineatus* (Langiano and Martinez, 2008), and 8.3 mg/l for *Oncorhynchus mykiss*, (Waynon, 1980), 16.8 mg/l for Nile tilapia, *Oreochromis niloticus* (Jiraungkoorskul et al., 2002), 97.47 mg/l for catchama blanca, *Piaractus brachypomus* (Ramirez-Duarte et al.,

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2008), and 211.80 mg/L for *Tilapia zilli* (Nwani et al., 2013). *Aphanius dispar* juveniles has been reported to respond differently to different pesticides upon 24h acute exposure to different concentrations of synthetic pyrethroid pesticides, fenpropathrin and fenvalerate (Shoaib et al., 2013) expressing differently low LC50 values, indicating high sensitivity of fish juveniles to different pesticides as many environmental factors influence the bioassay results (Shoaib et al., 2012).

Fish behavior has been indicated to be the most sensitive indicators upon exposure to environmental stressors particularly in fish species (Banace et al., 2011; Zarei et al., 2013). Fish were observed in previous studies to express various abnormal behavioral changes such as hyperactivity, loss of schooling, overcrowding, hypoactivity, breathing difficulties, jumping out of water, surfacing, jerky swimming, rapid opercula movements, loss of righting response convulsion, loss of balance, after exposure to a number of environmental toxicants, thus their living standards is affected overwhelmingly before extinctions occur (Banace et al., 2011; Ba-Omar and Al-Jardani, 2011; Kumar et al., 2015; Zarei et al., 2013; Sani and Idris, 2016; Nwani et al., 2013). *Aphanius dispar* exposed to temphos, which is an organophosphate pesticide expressed common behavioral changes such as restlessness, erratic swimming, convulsion, and loss of balance (Ba-Omar and Al-Jardani, 2011). Furthermore, the reported effects of glyphosate exposure in inducing histopathological changes of gills and liver tissues of fish included epithelial uplifting, interlamellar hyperplasia, hypertrophy of epithelial cells, shortening and folding of lamellae, necrosis of lamellar epithelium, lamellar fusion, aneurism, hyperplasia of chloride cells and mucus cells in the interlamellar spaces, clubbing, edema, and degeneration of filaments of the gills (Akinsorotan and Olele, 2013; ayoola, 2008; Deivasigamani, 2015; Jiraungkoorskul et al., 2002). while, they observed histopathological changes of the liver to include cytoplasmic and nuclear degeneration, hyperplasia, vacuolization of the cytoplasm, mild to severe infiltration of leukocytes, pyknotic nuclei, hypertrophy of hepatocytes, necrosis and bile stagnation. Ba-Omar and Al-Jardani, (2011) found lamellar damages including degradation of chloride cells, desquamation, epithelial uplifting, sloughing of epithelial cells, hypertrophy of lamellae, fusion of secondary lamellae, curling, tearing and collapsing of the lamellae after exposure of *Aphanius dispar* to the organophosphate temphos.

Aphanius dispar (Ruppell 1829), which also is known by the common name Arabian killifish has a wide distribution throughout Africa, Asia, and coast line of the red sea including Saudi Arabia. However, Arabian killifish population represents a single species, but with many color variations and patterns depending on locality. In Saudi Arabia, *Aphanius dispar* can breed all

year round with vivid coloration of the males that attract females and can tolerate in their habitats a wide range of temperature, salinity, and other factors making *Aphanius dispar* able to tolerate stressors, which might jeopardize their existence due to reduced food availability, habitat degradation, exotic species, chemical contamination, and exploitation.

Since *Aphanius dispar* existence is threatened not only by their habitat degradation and food availability, but also by various environmental stressors (Saeed et al., 2015), the focus of this current study was to determine the acute toxicity of the commercially formulated glyphosate and its effects on behavior and histology of the gills and liver tissues in light of the excessive use of glyphosate in agriculture.

II. MATERIALS AND METHODS

a) Chemicals

A commercial formulation of glyphosate (480 g/l glyphosate-isopropylamine salt) with trade name (Tiller 480 SL) manufactured by Astra Chem., Tabuk, KSA, was used in this current study.

b) Experimental Fish

Male and female juvenile Arabian killifish, *Aphanius dispar* were collected from southwestern Saudi Arabia, specifically from sadder Weila valley, through netting using a hand net. The juveniles were transported in a clean-aerated freshwater to the laboratory with care to lessen stress. *Aphanius dispar* juveniles mean weight was 1.5 ± 0.3 g and 4.5 ± 0.5 cm of length were allocated to aquaria randomly and left to be acclimatized under laboratory conditions for two weeks before running the static bioassay. Fish were fed commercial diet (flaked- food) once daily. The average values of water quality were (temperature 22 ± 1.0 °C, pH 7.2 ± 0.1 , dissolved oxygen $7.03 \pm .02$ mg/L, and total hardness 220 ± 2 mg/l). The light and dark cycle of 12 h: 12 h was maintained throughout the whole study duration.

c) Acute Toxicity Test

The bioassay test was conducted according to the US EPA guidelines 712-C-6-118 (1996) to determine the 96-h LC50 values of commercial formulation of glyphosate (Tiller 480 SL). Fish were starved for 24 hours prior to and during the bioassay. The test was conducted in plastic aquaria (30 + 30 + 15 cm) containing 8 L of static water (10 fish per aquarium). Seven different concentrations of glyphosate (60, 90, 120, 150, 180, 210 and 240 mg/L) with two replicates plus the control were used for running the acute toxicity test. Fish mortality in each aquarium was recorded and dead fish were removed immediately throughout the bioassay duration. The LC50 value of the fish was determined using the Probit analysis method (Finney,1971), and their behavior was monitored daily

for any abnormal behavioral changes throughout the bioassay.

d) *Sub-Acute Toxicity Test*

In order to investigate the histopathological effect of glyphosate on the gill and liver tissues, fish were exposed to 1/4 th of the 96-h LC50 glyphosate dose for 14 days. At the end of the exposure, fish were sacrificed and the gill and liver tissues of the control and treated fish were immediately excised and fixed in Bouin’s solution for 48 h at room temperature. After fixation, the tissues were washed with tap water, dehydrated through a graded ethanol series, cleared in xylene and embedded in paraffin wax. Sections of 6 μm were cut using a microtome (American Optical Co., USA), and stained with hematoxylin and eosin (Bancroft J and Steven A 1996). The stained sections were then examined for histopathological changes and photographed using Olympus light microscope

(Olympus, Tokyo, Japan) equipped with a digital camera.

e) *Statistical Analysis*

The 96-h LC50 value of the *Aphanius dispar* was calculated using the Probit analysis method (Finney, 1971). One way ANOVA was performed using SPSS software to detect significant differences among groups. P value < 0.05 was considered statistically significant.

III. RESULTS

a) *Acute Toxicity Test*

The 96-h LC50 value (Fig.1) of glyphosate upon exposure of *Aphanius dispar* to different glyphosate concentrations (60, 120, 150, 200, 180, 210 and 240 mg/L, was determined to be 115.25 mg/L using the probit analysis method.

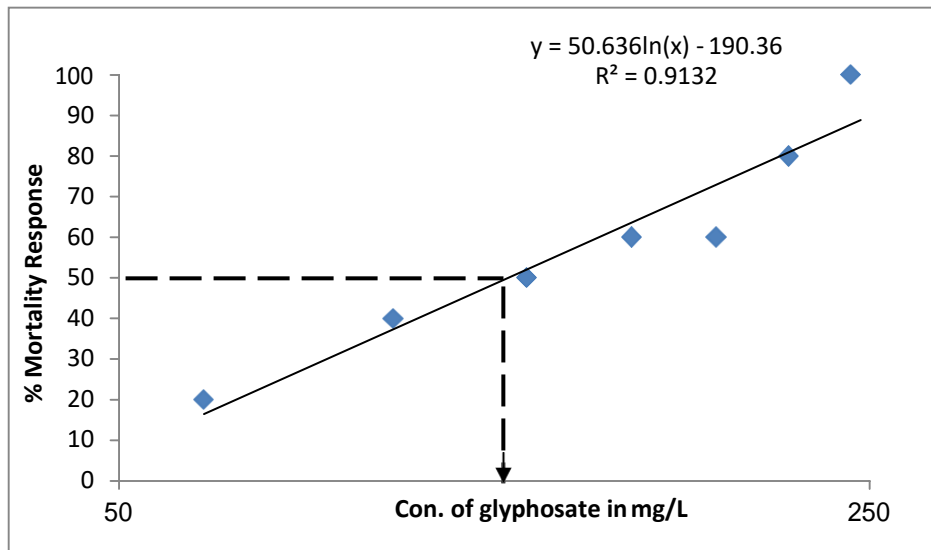


Fig. 1: The relationship between glyphosate concentrations and mortality percentages

b) *Fish behavior*

Unexposed fish group to glyphosate did not exhibit any adverse abnormal behavioral responses or any mortality throughout the duration of the bioassay as

compared to the treated fish. However, exposed fish to glyphosate exhibited various abnormal behavioral responses, which were concentration-related as shown in Table1.

Table 1: Behavioral responses of the Arabian killifish, *Aphanius dispar* after acute exposure to different concentrations (mg/L) of glyphosate.

Concentration mg/L	Behavioral responses of <i>Aphanius dispar</i>
0	Normal activity to mild hyperactivity on the first day, Normal feeding behavior
60-89	Mild hyperactivity associated with loss of schooling into schools
90-119	Hyperactive surfacing to the tank top as an avoidance response associated with rapid opercula movements
120-149	Erratic movements associated with opercula movements and rapid mouth movement rate gulping for air
150-179	Schooling on and off, frequent surfacing and jumping outside the aquaria, cannibalism, loss of balance associated with hanging vertically in the water column head-up tail-down, swirling with rapid speed

180-209	Exhaustion associated with hypoactivity, settling on the bottom of the tanks with less opercula and mouth movements
210-240	Exhaustion associated with hypoactivity and mortality

c) *Histopathological Study*

Mild focal changes of the Arabian killifish, *Aphanius dispar* gill lamellae were observed in the control group (Fig. 2-a), while there were no observed histopathological changes in the liver tissues. However, exposed fish to the herbicide glyphosate (1/4 96 hLC50) concentration exhibited a wide range of mild to excessive histopathological alterations of the gills (Fig. 2, b-e) and Liver (Fig. 3, b-d) (Fig. 3-a.).

The gills of exposed fish exhibited histopathological changes of which epithelial uplifting,

edema, epithelial hyperplasia, fusion of lamellae, clubbing of the tips of secondary lamellae, and thickening of lamellar epithelium (Figure 2, b-e). While, the liver organ exhibited histopathological changes of significance including deterioration and necrosis of the liver hepatocytes, hypertrophy of hepatocytes, eccentric nuclei and pyknosis, as well as mild to extensive vacuolization of hepatocytes with a foamy appearance (Figure 3, b-d).

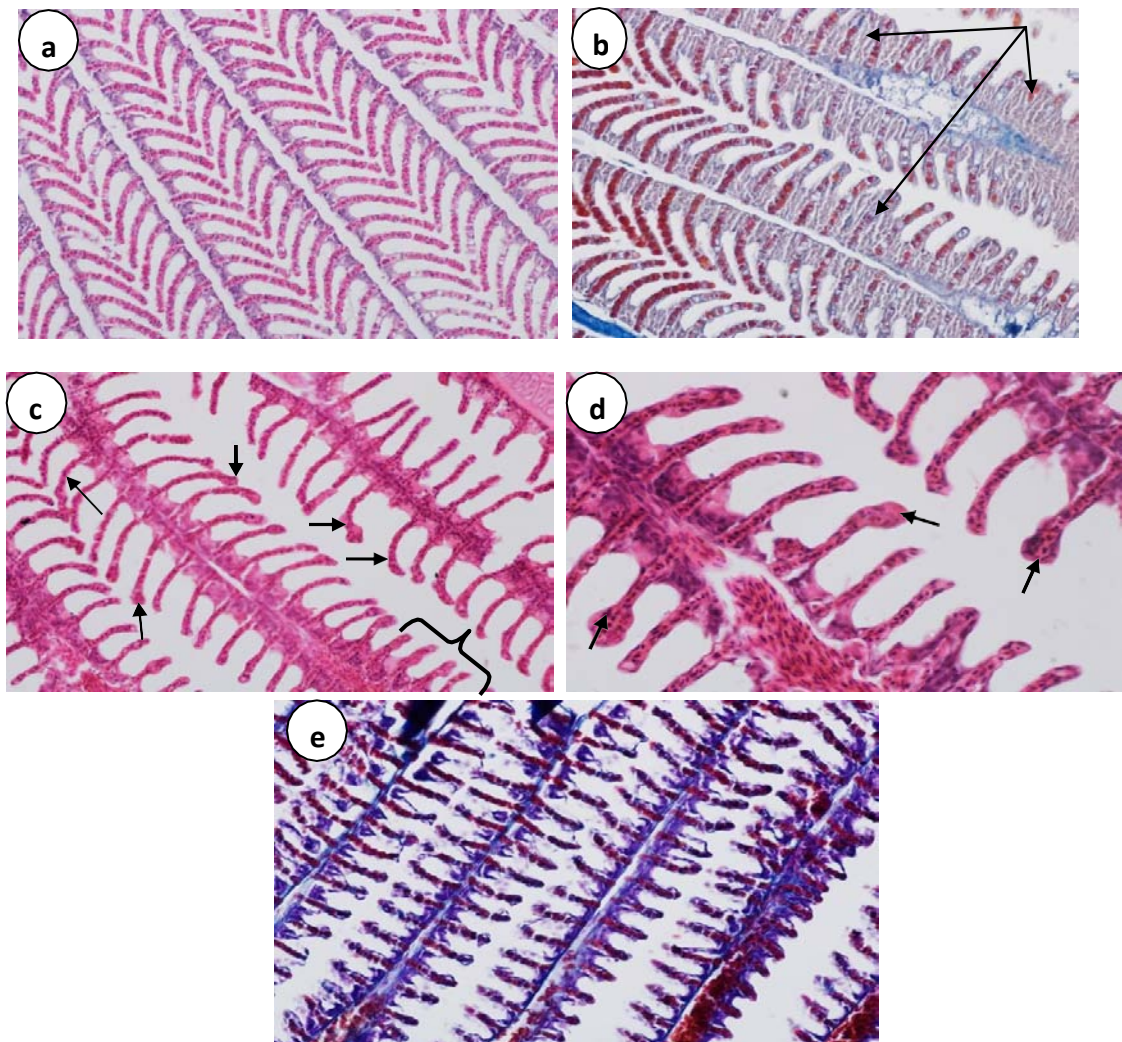


Fig. 2 (a-e): Histopathological changes of *Aphanius dispar* gills: Control group (a) and glyphosate treated (b-e). a: normal gill filaments and secondary lamellae. b: interlamellar hyperplasia of the filaments leading to fusion of secondary lamellae (arrows), c: shortening of secondary lamellae (bracket), elongation associated with curling, clubbing and elongation of secondary lamellae ((small arrows), d: clubbing of the secondary lamellae (arrows). e: gills showing generalized excessive epithelial uplifting from the basement membrane. (H & E) X 20.

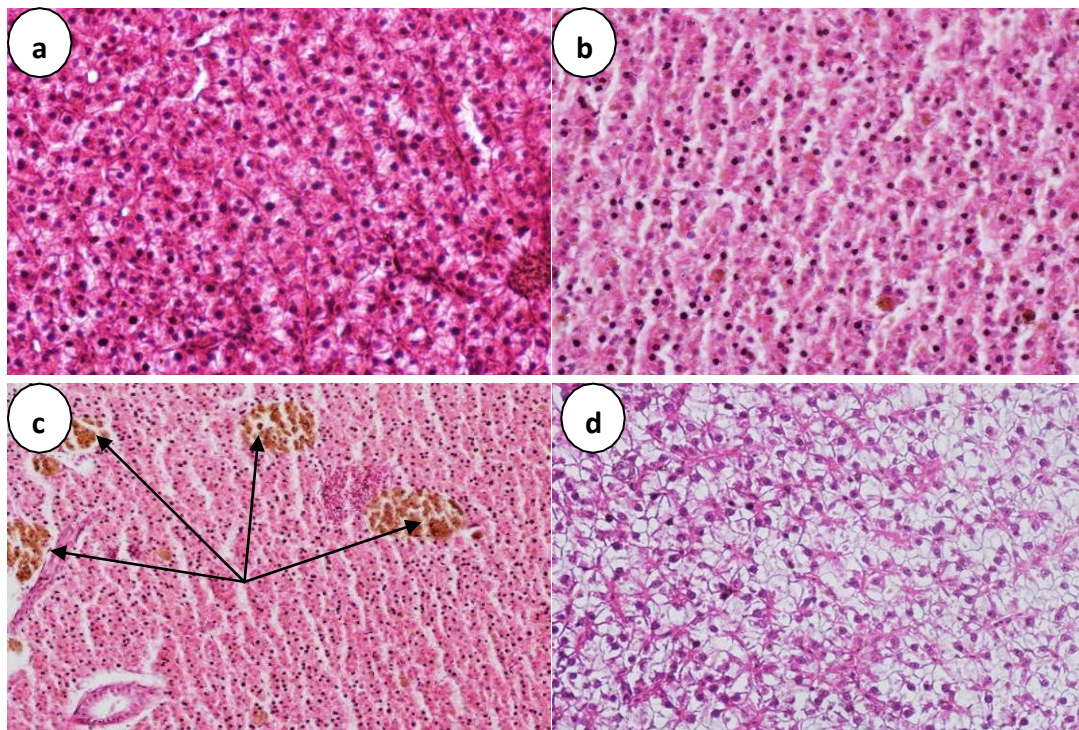


Fig. 3: Histopathological changes of *Aphanius dispar* liver: Control group (a) and glyphosate treated (b-d). a: normal liver of the control group with normal polygonal hepatocytes associated with very mild deterioration. b: deterioration of the hepatocytes and pyknosis of nucleolus. c: extensive deterioration and excessive bile stagnation (arrows). d: excessive cytoplasmic vacuolization with a foamy appearance. (H & E) X 20.

IV. DISCUSSION

a) Acute Toxicity

Mortality and extinction of many life forms is imposed by myriads of chemical pollutants including herbicides. The impacts of chemical pollutants reflect their concentrations, duration of exposure, environmental factors, and sensitivity of life forms. Thus determination of the lethal dose (LC50), the dose that might kill half of any population within a short period of time, is considered the first step prior to any physiological studies as with regard to chemical pollutants. In this current study, glyphosate formulation (Tiller 480 SL) 96 h LC50 of the Arabian killifish, *Aphanius dispar* juveniles valued at 115.25 mg/l (Figure. 1). The 96h LC50 values of glyphosate have been investigated on different fish species at different environmental conditions, indicating variation in concentrations (Neskovic et al., 1996; Jiraungkoorskul et al., 2002). Previously very limited studies have stated that *Aphanius dispar* juveniles responded differently to different pesticides (Shoab et al., 2013), indicating variability in sensitivity to different pesticides being influenced by various environmental factors (Shoab et al., 2012; Shoab et al., 2013). Furthermore, Nwani et al. (2013) reported the 96 h LC50 for *Tilapia Zilli* at 211.80 mg/l upon exposure to glyphosate formulation (Forceup). The reported LC 50 values which are close to the results of this current study upon exposure to

glyphosate for 96 h were found to be 97.47 mg/L for cachama blanca, *Piaractus brachipomus* (Ramirez-Duarte et al., 2008) and 86 mg/l for the common carp, *Cyprinus carpio* (Deivasigamani, 2015). Additionally, lower 96 h LC50 values were recorded: 43.65 mg/L for the African catfish, *Clarias gariepinus* (Akinsorotan, 2013), 13.69 mg/l for Neotropis fish, *Pochilodn lineatus* (Landgiano and Martinez, 2008), 10.0 mg/l for the Asian Bass, *Lates calcarifer* (Thunomsit et al. 2016), 1.05 mg/l for Nile tilapia, *Oreochromis niloticus* (Ayoola 2008) and 0.05 mg/l for the African catfish, *Clarias gariepinus* (Ayanda et al., 2015).

Therefore, from the determined 96 h LC50 values (115.25 mg/l), we do believe that *Aphanius dispar* juveniles are very sensitive to glyphosate and its toxicity might be exacerbated further upon exposure under harsher environmental factors. Thus, as we compare this current study 96 h LC50 value to the previously reported studies, 96 h LC50 higher and lower values on different fish species and different environmental conditions, we believe that the determined 96 h LC50 value in our study on might be influenced by the ambient environmental conditions for this species.

b) Fish behavior

Behavioral responses have been indicated to be the most sensitive indicators upon exposure to potential toxic effects in fish species (Banace et al., 2011; Ba-Omar and Al-Jardani 2011). Our results

showed that the unexposed fish group did not reveal any adverse abnormal behavioral responses or mortality throughout the duration of the bioassay as compared to the treated fish groups other than mild hyperactivity at the onset of the experimental execution, which might be attributed to fish handling during allocation. On the contrary, exposed fish to glyphosate exhibited various abnormal behavioral responses (Table1) and were concentration-related similar to the previously reported observations after exposure of different fish species to the herbicide glyphosate (Akinsorotan et al., 2013; Ayoola 2008; Okayi et al., 2010). Abnormal behavioral changes such as mild to moderate erratic swimming, rapid rate of opercular and mouth movements, infrequent surfacing were observed at low to moderate concentrations, while at higher concentrations *Aphanius dispar* exhibited rapid swimming associated with frequent surfacing and jumping outside of the aquaria, hanging head-up tail-down position, and hypoactivity before the fish became weak, hypoactive, and settled at the bottom followed by exhaustion and death. These observations were consistent with the previously reported abnormal behavioral changes after exposure of various fish species to the herbicide glyphosate: *Clarias gariepinus* adult (Akinsorotan et al., 2013), *Clarias gariepinus* fingerlings (Okayi et al., 2010), juvenile African catfish, *Clarias gariepinus* (Ayoola, 2008), the common carp, *Cyprinus carpio* (Deivasigamani, 2015), and Asian sea bass, *Lates calcarifer* (Thanomsit et al., 2016). Similarly, *Aphanius dispar* juveniles exposed to the organophosphate temphos expressed abnormal behavioral changes (Ba-Omar and Al-Jardani 2011). These previously mentioned authors reported such abnormal behavioral changes and mortality to occur after acute and chronic toxicity indicating respiratory failure inflicted by the effects of the glyphosate on the gills. Thus, fish mortalities observed in this study could be due to the destruction of gill tissues and impairment of gas-exchange capacity after fish became very lethargic and exhausted. Furthermore, fish respiratory failure might be an indication of physiological distress on juveniles resulting from potential progressive energy expenditure with time preceding mortality of fish.

Furthermore, according to Kumar et al. (2015), Zarei et al. (2013), Nwani et al., (2013), and Okayi et al., (2010) mucus secretion observed in this current study at the water surface at higher concentrations might suggest excessive impacts of glyphosate on fish gills forming a mucus film on the gills interrupting gaseous exchange and causing death of fish following exhaustion and lethargic responses. While, Pandey et al., (1990) attributed the secretion of mucus to dysfunction of the endocrine gland under toxic stress thus changes in the number and area of mucus glands and chromatophores. On the other hand, Sani and Idris, (2016) the previously reported behavioral changes have been indicated to occur as a result of not only metabolic

dysfunction but also due to nervous disorder upon exposure to toxic glyphosate. Thus we do believe that fish exhibiting such abnormal behavioral changes and mucus secretion in *Aphanius dispar* in this current study might have been due to the toxic effects of glyphosate on gill tissues and respiration impairment.

c) *Histopathology*

Aphanius dispar upon exposure to glyphosate revealed gill and liver tissue alterations which were concentration-and-time related. Literatures on the impacts of noxious chemicals on fish histopathology of *Aphanius dispar* gills, liver, kidney and all other levels of biological organizations are scarce and very current. For example, the acute and chronic impacts of 3,4-dichloroaniline (DCA), sodium dodecyl sulfate, and zinc sulfate and chlorine on *Aphanius dispar* embryos development were studied (Saeed et al., 2015). While, the effects of the organophosphate temphos on the gills of *Aphanius dispar* revealed various concentration-related gill damages including hemorrhage of lamellae, epithelial uplifting, epithelial hypertrophy, swelling at the base and tips of lamellae, fusion of lamellae, tears in the filaments, sloughing of epithelial cells from the filaments and lamellae (Ba-Omar and Al-Jardani, 2011).

The gill organs of fish, which are very complex structure essential for gaseous exchange, acid-base balance, excretion, and osmoregulatory function are in contact with the outside environment. There was no recognizable alterations in the gills of the control fish. However fish exposed to 1/4th 96 h LC50 glyphosate exhibited various tissue alterations including fusion of secondary lamellae, epithelial uplifting, minor clubbing of the secondary lamella tips, hyperplasia of the primary filaments and secondary lamellae, and curling of the secondary lamellae. Similarly, previous studies have reported wide spectrum of gill histopathological changes after exposure of fish species to variety of noxious agents including glyphosate. Fish exposure to glyphosate for 96 h of *Clarias gariepinus* (Akinsorotan et al., 2013), Juveniles African catfish, *Clarias gariepinus* (Ayoola, 2008), common carp, *Cyprinus carpio* (Deivasigaman, 2015), *Cyprinus carpio* (Neskovic et al., 1996), Nile tilapia, *Oreochromis niloticus* (Jiraungkoorskul et al., 2002) and *Aphanius dispar* to temphos (Ba-Omar and Al-Jardani (2011) caused wide spectrum of gill histopathological changes of which epithelial hyperplasia, edema, lifting of epithelium, epithelial hyperplasia thickening of primary lamellar epithelium, clubbing, fusion of lamellae and secretion of mucus etc. The pronounced variety of insults of the gill organs which have been observed in this current study followed by exhaustion, lethargy and death of fish are clear indication of the impairment of gaseous exchange and reduced functional efficiency of the gills before suffocation of fish occurred. Therefore, the histopathological changes of fish gills can impair the

respiratory function by reducing the total surface area available for oxygen uptake and increase the diffusion distance between the external environment and the blood inside the lamellae preventing gaseous exchanges, which cause suffocation of fish and death (Ayoola 2008; Ba-Omar and Al-Jardani, 2011; Jiraungkoorskul et al., 2002;).

On the other hand, liver, which is an organ performing various functions associated with the metabolism of xenobiotics (Langiano et al., 2008) exhibited various histopathological alterations after exposure to glyphosate used in this current study including vacuolization of the cells cytoplasm, hypertrophy of hepatocytes, degeneration of hepatocytes, and bile stagnation. Similar findings on the impacts of glyphosate on fish species at various concentrations were reported (Langiano et al., 2008; Ayola 2008; Akinsorotan et al., 2013; Jiraungkoorskul et al., 2002; Deivasigamani 2015; Neskovic et al., 1996). It is believed that the vacuolization of liver cells might indicate evidence of fatty degeneration (Deivasigamani 2015; Jiraungkoorskul et al., 2002; Ayoola 2008). The localized necrosis of hepatocytes suggest excessive work required by the fish to get rid of the toxicant from its body during the process of detoxification as well as failure of the liver cells to regenerate due to continuous exposure to noxious agents (Deivasigamani 2015; Ayoola 2008). Bile stagnation was observed in the control and treated Neotropical fish *Prochilodus lineatus* as manifestation of a physiopathological conditions caused by a lack of bile metabolism and secretion (Langiano et al., 2008; Ayoola 2008; Deivasigamani 2015), whereas we found that bile secretion by hepatocytes observed within the cells as yellowish droplets in the exposed fish, however, bile droplets were very intense and the degree of intensity was observed in the treated fish with higher glyphosate concentrations, which might indicate glyphosate effects as opposed to suggested nutritional deficiency by Langiano et al., (2008) since we were prompt on feeding the fish once daily and on time.

In conclusion, the results of this current study asserts the toxic impacts of glyphosate on fish behavior and histopathology of the gill and liver tissues. Thus, impairment of gills and liver functional efficiencies before exhaustion, suffocation, and death occurred. Therefore, we recommend regulating glyphosate usage in/or near aquatic environment and the importance of establishing environmental monitoring commission guidelines to regulate or discourage the use of glyphosate and other harmful chemicals. Thence, protecting not only life forms like fish from extinction, but also ensures biological diversity and stability for healthier ecosystems.

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On the Investigation of Awareness Level of Family Planning among Rural Dwellers in Nigeria (Principal Component Analysis Approach)

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Abstract- This study examined the awareness level and attitude of family planning among rural dwellers in the South western part of Nigeria and its consequences on population/economy of Nigeria. Data were gathered from 300 respondents who were randomly selected from Seriki rural communities of western Nigeria using proportional allocation of the stratified random sampling technique. The research considered to know the level of awareness between the traditional family planning methods and modern methods of contraception. The modern methods were found to be more significantly embraced than traditional family planning methods based on the analysis of the result obtained which shows that 175% prefer modern methods while 125% preferred traditional methods.

Keywords: *family planning, principal component, eigen values and vectors, marital status and contraceptives.*

GJSFR-C Classification: FOR Code: 279999



ON THE INVESTIGATION OF AWARENESS LEVEL OF FAMILY PLANNING AMONG RURAL DWELLERS IN NIGERIA (PRINCIPAL COMPONENT ANALYSIS APPROACH)

Strictly as per the compliance and regulations of :



RESEARCH | DIVERSITY | ETHICS

On the Investigation of Awareness Level of Family Planning among Rural Dwellers in Nigeria (Principal Component Analysis Approach)

Ayoola, F. J. ^α, Adeboye, N. Olawale ^σ & Kayode Balogun ^ρ

Abstract- This study examined the awareness level and attitude of family planning among rural dwellers in the South western part of Nigeria and its consequences on population/economy of Nigeria. Data were gathered from 300 respondents who were randomly selected from Seriki rural communities of western Nigeria using proportional allocation of the stratified random sampling technique. The research considered to know the level of awareness between the traditional family planning methods and modern methods of contraception. The modern methods were found to be more significantly embraced than traditional family planning methods based on the analysis of the result obtained which shows that 175% prefer modern methods while 125% preferred traditional methods. The factors considered in examining these are educational background, economic factors, marital status, social factor, ethnicity and illiteracy level; the data collected based on these factors was analyzed using principal component analysis technique in order to determine the most prevalent factor that causes attitudinal problems. The results give rise to the Eigen values and Eigen vectors of the components, whereby the variance proportion for each is given as 0.4844, 0.2391, 0.1480, 0.0597, 0.0461 and 0.0227 for economics, educational background, ethnicity, illiteracy, marital status and social status respectively; thereby qualified the factors as first PC, second PC, third PC, fourth PC, fifth PC and the sixth PC respectively. Thus, economic factor which is the first PC is the factor that is predominantly responsible for the discovered low level of awareness and poor attitudinal behavior towards family planning.

Keywords: family planning, principal component, eigen values and vectors, marital status and contraceptives.

I. INTRODUCTION

Family planning implies the ability of individuals and couples to anticipate and attain their desired number of children by spacing and timing their births. It is achieved through the use of contraceptive methods and the treatment of involuntary fertility. The availability of family planning does more than enable women and men to limit family size. It safeguards

individual health and right and improves the quality of life of couples and their children.

Family planning has attracted attentions all over the world due to its relevance in decision making, population growth and development. Family planning is defined as birth spacing, preventing unwanted pregnancies or secure wanted pregnancy (W.H.O, 1995). Beekle and McCabe (2006) defined family planning as the practice that helps individuals or couples to attain certain objectives such as avoiding unwanted pregnancies, bringing about wanted babies at the right time, regulating, the interval between pregnancies, controlling the time at which birth occurs in relation to the ages of parents and determining the number of children in the family. Family planning is a means of reproductive health. In spite of the hue and cry in and outside Nigeria about family planning or birth control, many people are still confused about its meaning, the methods involved, the advantages and disadvantages and the factors hindering its wide application in Nigeria, especially among the rural communities (Iffih and Ezeah, 2004). Women's education enhances their capability and also their reproductive rights to decide freely and responsibly the number, spacing and timing of their children and to have other necessary information regarding reproductive rights. Studies have shown that education is a determinant of awareness of family planning practices in Nigeria, for instance; Anyanwuh et.al (2013) investigated the extent of family planning, the methods and contraceptive devices in use and the influence of education on family planning among couples in Nkanu Local Government Area of Enugu State. The findings revealed that educational background of the couples significantly influenced the choice of family planning in the community.

W.H.O (2013) found out that women's education is in line with lower fertility which constitutes management of reproductive resources. Maternal education has once been linked with reduction of child mortality among rural dwellers. Recent studies have also shown that religion is a good determinant of family planning practices. UNFPA (2013) identified some

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factors such as cultural background, and religions beliefs which place the women at disadvantaged position in reproductive health issues. Ezea and Iffih (2004) asserted that Catholic Church is rigid in their views of family planning. Catholics hold the view that the application of artificial method is wrong and should not be allowed. The Catholic Church is said to be comfortable with the use of Billings's ovulation method which is rather natural. Igbudu et. al (2011) conducted a study on the relationship between religious beliefs and family planning practices of married women in zone 5 barracks of the Nigeria police, comprising Edo, Delta, and Bayelsa state commands. The findings of the study revealed that attitudinal factor such as the strong religious desire for more children prevented women in these barracks from using family planning. Also noted were the insufficient knowledge on contraceptives methods, their fears and anxieties, rumor of others using contraceptives getting deleterious effect and poor delivery of health services (Hamau R.K et.al, 1996). Traditional methods such as coitus interruptus is described in Bible, periodic abstinence was used in ancient India and the precursor to the condom was used by the Egyptians back in 1350 BC (Odimegwu, 1999). Of recent, Ghulam Mustafa et al (2015) conducted a similar research in among rural dwellers in Pakistan and findings reveal that the majority knew about some modern contraceptive methods, but the overall contraceptive use was very low. Knowledge and use of any contraceptive method were particularly low. Reasons for not using family planning and modern contraception included incomplete family size, negative perceptions, in-laws' disapproval, religious concerns, side-effects, and lack of access to quality services.

The practices of modern contraceptives method offer many advantages compared to traditional approaches, in health and economy of the couples in the country. Family planning programs that stimulate modern approaches, policies and methods have become increasingly important in the last decades as a result of the social-economic problems influencing rapid population growth, as well as public health problems, especially control of sexually transmitted diseases (STDs) such as AIDS (UN, 1994).

The unwelcomed attitude of most rural dwellers to modern family methods has constitutes a continuous growth in African population size keeping it almost permanently underdeveloped, There is general believe that economic stagnation is often linked with over population and poor investment in health services and with Nigeria's population currently put at over 140 million people compared with Gross Domestic Product (GDP) that is not too encouraging and inflation still skyrocketing, there is the fear that future living standards may substantially depreciate (Olatayo and Adeboye, 2013). In recent years, population growth has begun to be discussed in terms of its effect on global stability.

Some of the supporters of this view have suggested that in post-cold war order, the growth of population has the potential to undermine the global stability. Thus, understanding of family planning scenario among different societies and communities, which by and large reside in urban slum and rural areas, might prove useful in increasing family planning acceptance by them and decreasing population growth (Sharma, V et.al, 2012). Within the context of above literatures, it has therefore becomes imperative to carry out this study in Seriki community; a rural community located in Ogun state, western region of Nigeria. The factors considered in examining this scenario are educational background, economic factors, marital status, social factor, ethnicity and illiteracy level; Principal component analysis technique shall be employed to fit model for each of the factor in order to determine the most prevalent among them that cause attitudinal problems to family planning among rural dwellers.

II. RESEARCH DESIGN AND METHODOLOGY

The study area of this research is Seriki community in Ogun State and is accessible by land. The community comprises of different ethnic groups namely Yoruba's, Igbo, Hausa, Calabar, Isoko, Ijaws, just to mention but few. The Yoruba's are considered to be the more than any other tribes in the city, because it is located at the western part of Nigeria. According to the unpublished census figure of 2006, its population size was estimated at a total of 2,075. The community was divided into nine (9) strata and a sample size of 335 estimated through Taro Yanmeni's formula was randomly selected among focus group of age 15 – 30+ comprises of single ladies, married couple and the divorced people using proportional allocation of Stratified random sampling technique.

Designed questionnaire was examined on the selected sample size under three (3) different categories. The instrument sought to solicit responses on area of residence, age at marriage, and highest educational qualification, duration or age of marriage, and number of children owned. The second category of the questionnaire covers the factors that encourage large families and their general perception on family planning. The major factors considered here are educational background, economic factors, marital status, social factor, ethnicity and illiteracy level. The third part purely seek information on family planning acceptance in both modern and traditional way. At the end of the survey, a cronbach's alpha validity rest was run on a sample of 10 respondents for reliability measures and the survey was found to be 81.2% reliable.

III. DATA ANALYSIS

The analytical technique adopted for this research is a Multivariate analysis technique known as Principal Component Analysis.

It is a statistical procedure that uses an orthogonal transformation to convert a set of observations of possibly correlated variables into a set of values of linearly uncorrelated variables called principal components. This transformation is defined in such a way that the first principal component has the largest possible variance (that is, accounts for as much of the variability in the data as possible), and each succeeding component in turn has the highest variance possible under the constraint that it is orthogonal to the

preceding components. The principal components are orthogonal because they are the Eigen vectors of the covariance matrix, which is symmetric.

If the variables Y_1, Y_2, \dots, Y_p in Y are correlated, the ellipsoidal swarm of point is not oriented parallel to any of the axis. The axes can be rotated by multiplying each y_i by an orthogonal matrix A and thus obtain

$$Z_i = A Y_i \tag{1}$$

Since A is an orthogonal, $A'A=I$ and the distance to the origin is unchanged. Thus we want to sample covariance matrix of $Z S Z = A S A'$ to be diagonal.

$$Z S Z = A S A' = \begin{bmatrix} S^2 z_1 & 0 & \dots & 0 \\ 0 & S^2 z_1 & \dots & 0 \\ 0 & 0 & \dots & S^2 z_1 \end{bmatrix} \tag{2}$$

Thus the orthogonal matrix A that diagonalizes S is the transpose of the matrix C .

$$A = C' = \begin{bmatrix} a_1 \\ a_2 \\ a_p \end{bmatrix}$$

a) Eigenvalues and Eigenvectors

For every square matrix A , a scalar λ and a nonzero vector x can be found such that $Ax = \lambda x$. It should be known that λ is called an eigenvalue of A , and x is an *eigenvector* of A corresponding to λ . To find λ and x , we write the equation as $(A - \lambda I)x = 0$.

If $|A - \lambda I| \neq 0$, then $(A - \lambda I)$ has an inverse and $x = 0$ is the only solution. Hence, in order to obtain nontrivial solutions, we set $|A - \lambda I| = 0$ to find values of λ that can be used to find corresponding values of x .

Thus, in $(A - \lambda I)x = 0$, the matrix $A - \lambda I$ must be singular in order to find a solution vector x that is not 0.

Suppose we have a 3×3 matrix A with eigenvectors X_1, X_2, X_3 , and eigenvalues $\lambda_1, \lambda_2, \lambda_3$ so that

$$A X_1 = \lambda_1 X_1$$

$$A X_2 = \lambda_2 X_2$$

$$A X_3 = \lambda_3 X_3$$

thus,

$$A \begin{bmatrix} X_1 & X_2 & X_3 \end{bmatrix} = \begin{bmatrix} X_1 & X_2 & X_3 \end{bmatrix} \begin{bmatrix} \lambda_1 & 0 & 0 \\ 0 & \lambda_2 & 0 \\ 0 & 0 & \lambda_3 \end{bmatrix} \quad (4)$$

The eigenvalues will then be used to compute the variance of the principal components and we can speak of the proportion of variance explained by the first K component.

b) *Eigenvectors of Covariance Matrix*

We will derive our first algebraic solution to PCA using linear algebra. This solution is based on an important property of eigenvector decomposition. The

sample correlation between the *j*th and *k*th variables is defined as $CovariancesX^T X$ itself can be recognized as proportional to the empirical sample covariance matrix of the dataset X.

IV. RESULT AND DISCUSSION

a) *Results*

Table 1: Ordinary correlations

	E.F	E.B	ETHNICITY	ILLITERACY	MARRIAGE-STATUS	SOCIAL-FACTOR
ECONOMIC_FACTOR	1.000000					
E.B	0.853992	1.000000				
ETHNICITY	-0.260151	-0.174099	1.000000			
ILLITERACY	-0.688533	-0.683219	0.134808	1.000000		
MARRIAGE_STATUS	0.353807	0.234607	-0.719295	-0.217871	1.000000	
SOCIAL_FACTOR	-0.247535	-0.224509	0.137374	0.281727	-0.132131	1.000000

Table 2: Eigenvectors

Variable	PC 1	PC 2	PC 3	PC 4	PC 5	PC 6
E.F	0.525305	0.192413	0.174055	0.349704	0.099909	-0.724197
E.B	0.498518	0.289330	0.192530	0.413398	-0.079352	0.673428
ETHNICITY	-0.295534	0.653014	-0.005069	-0.029402	0.695535	0.039670
ILLITERACY	-0.463585	-0.295967	-0.043416	0.825845	0.116089	-0.010534
M.S	0.339941	-0.601188	0.065349	-0.117261	0.696298	0.141992
S.F	0.240255	-0.061762	0.962526	-0.100871	-0.040569	-0.013651

Table 3: Sum of the Variance Proportion

Eigenvalues: (Sum = 6, Average = 1)					
Number	Value	Difference	Proportion	Cumulative Value	Cumulative Proportion
1	2.906611	1.472286	0.4844	2.906611	0.4844
2	1.434325	0.546397	0.2391	4.340936	0.7235
3	0.887928	0.529765	0.1480	5.228865	0.8715
4	0.358163	0.081266	0.0597	5.587028	0.9312
5	0.276897	0.140822	0.0461	5.863925	0.9773
6	0.136075	---	0.0227	6.000000	1.0000



Observed Matrix Scree Plot

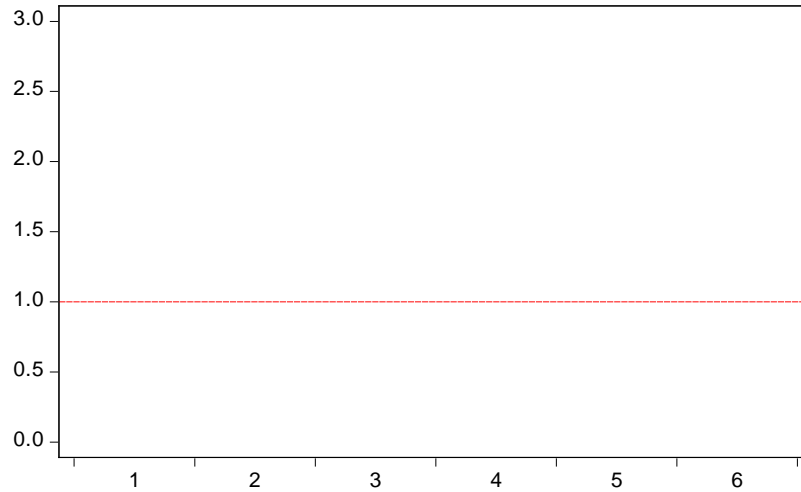


Figure 1: Observed matrix Scree plot

Table 4: Estimate of the Orthogonal

	Unrotated Loadings		Communality	Uniqueness
	F1	F2		
ECONOMIC_FACTOR	0.888934	0.188602	0.825774	0.174226
E.D	0.840691	0.301768	0.797825	0.202175
ETHNICITY	-0.436040	0.651089	0.614048	0.385952
ILLITERACY	-0.711991	-0.246761	0.567822	0.432178
MARRIAGE_STATUS	0.511711	-0.620111	0.646385	0.353615
SOCIAL_FACTOR	-0.300303	-0.013913	0.090376	0.909624

Factor	Variance	Cumulative	Difference	Proportion	Cumulative
F1	2.546057	2.546057	1.549884	0.718772	0.718772
F2	0.996173	3.542231	---	0.281228	1.000000
Total	3.542231	3.542231		1.000000	

Orthonormal Loadings Biplot

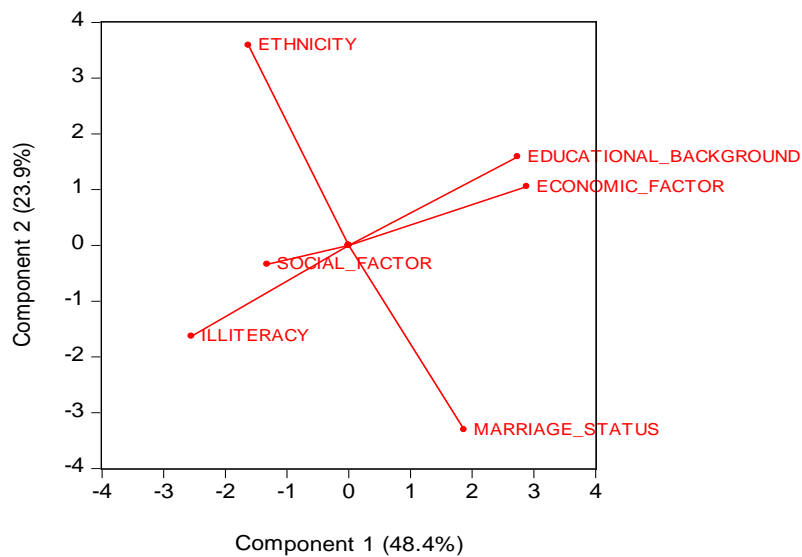


Figure 2: Orthogonal loadings

b) Discussion

In table 1, correlations among the causative factors were considered. The most prevalent dependence was exhibited between Educational background and Economic factor with a high positive correlation of 0.85 and that shows at what level the The model for the equation above are:

$$PC1 = 0.525305 (E.F) + 0.498518 (E.D) - 0.295534 (\text{ethnicity}) - 0.463585 (\text{illiteracy}) + 0.339941(M.S) - 0.240255 (S.F). \quad (5)$$

$$PC2 = 0.192413(E.F) + 0.289330 (E.B) + 0.653014 (\text{ethnicity}) - 0.295967(\text{illiteracy}) -0.601188 (M.S) -0.061762 (S.F). \quad (6)$$

$$PC3 = 0.174055 (E.F) + 0.192530 (E.B) - 0.005069 (\text{ethnicity}) - 0.043416 (\text{illiteracy}) +0.065349 (M.S) +0.962526 (S.F). \quad (7)$$

$$PC4 = 0.349704 (E.F) + 0.413398(E.B) -0.029402 (\text{ethnicity}) + 0.825845 (\text{illiteracy}) -0.117261 (M.S) -0.100871 (S.F). \quad (8)$$

$$PC5 = 0.099909 (E.F) -0.079352 (E.B) + 0.695535(\text{ethnicity}) +0.116089 (\text{illiteracy}) + 0.696298(M.S) -0.040569 (S.F). \quad (9)$$

$$PC6 = - 0.724197(E.F) + 0.673428 (E.B) + 0.039670 (\text{ethnicity}) -0.010534 (\text{illiteracy}) + 0.141992 (M.S) - 0.013651 (S.F). \quad (10)$$

In line with equations 5 to 10, the first principal component is positively correlated with three out of six of the original variables while others experienced negative correlation. Thus, the first principal component increases with the increase in economic factor, educational background and marriage status and this suggest that these three criteria vary together. If one increases, then the remaining two also increase. This component can be viewed as a measure of the quality of economic factor, educational background and marriage status. Furthermore, we see that the first principal component correlate moderately with the economic factor. In fact, we could state that based on the correlation of 0.525 that this principal component is a primary measure of the economic factor. It would follow that communities with high values would tend to have a lot of economic factor available around them in terms of economic status, type of family and this shows that economic factor will affect family planning in the community.

The second principal component also increases with ethnicity, educational background and economic factor. This component can be viewed as a measure of the degree of ethnicity in the community with a positive value of 0.65. Thus, this principal component is the primary measure of ethnicity which indicates that ethnicity also affect the attitudes to family planning in the community.

The third principal component is highly correlated with social factor with value of 0.962 follow by educational background, economic factor and marriage status. It implies that this component is the primary measure of social factor in the community. Thus, a degraded level of social factor within the community

correlation value will be of importance. Some of the factors equally exhibited negative correlation with one another while others exhibited weak positive correlation. According to Table 2, the main component variables are defined as linear combinations of the original variables.

have had serious effect on their family planning attitudes.

The fourth principal component also strongly correlated with illiteracy level in the community with a value of 0.825, follow by educational background and economic factor. This component is the primary measure of illiteracy in the community. Which indicate that there is high level of illiteracy in the community which is also affecting the family planning attitudes in the area.

The fifth principal component is the primary measure of marital status with a value of 0.696. This implies that marriage status of the focus group have strongly impacted on the community's attitudes to family planning in the area.

The sixth principal component is the primary measure of educational background with value of 0.67 follow by marriage status. Which indicate that educational background contributes a lot to the attitudes of family planning in the area.

Table3 shows that Economic factor has the highest variance proportion of 0.484 follows by Educational background with a variance proportion of 0.239 and it thus appears that the poor attitudes of the community dwellers are mostly affected by the first two principal components, and together they explained 72.35% of the total variation in the acceptability of family planning attitudes under study. This leads us to the conclusion that a two factor solution will probably be sufficient with the eigenvalues of 2.906611, difference of 1.472286, variance of 0.4844 and 1.434, difference of 0.5463, variance of 0.2391. The remaining other four components jointly explained 27.65 % of the attitudes.

Figure 2 Scree plot is a principal component analysis chart to determine how many important components are present in the data, the scree graph plot the Eigen values against the component number. It can be easily observed that only the first two components crossed above the cut-off line, meaning that the remaining components account for smaller amounts of the total variance. Generally we are interested in keeping only those principal component with Eigen values greater than one. However, two factor were extracted based on significance of Kaiser Criterion (Kaiser, 1960).

Table 4 shows a section displaying the estimates of the unrotated orthogonal loading, communalities and uniqueness estimates obtained from the estimates. we see that veliger MAP method has retained two factors, labeled F1 and F2 a brief examination of the unrotated loadings indicate that economic factor and educational background load on the first factor while ethnicity load on the second factor. To the right of the loadings are communality and uniqueness which apportion the diagonals of the correlation matrix into common (explained) and individual (unexplained) components therefore, the communalities explained what each variable can be accounted for with their weight(uniqueness). Thus, all the variables (factors) considered in this research are correlated since the ellipsoidal swarm of point is not oriented parallel to any of the axis as exhibited in figure 2.

V. CONCLUSION/RECOMMENDATION

This research provides insights into the local contexts related to family planning knowledge, attitudes, perceptions, and practices and also highlights the need for contraceptives, both modern and traditional methods. In the wake of changing attitudes towards family planning and desired family size among rural dwellers, more women and couples will be seeking family planning services. Addressing obstacles such as access, affordability, and availability will help meet these needs and ensure that women and couples can meet their childbearing and reproductive health goals. In addition, a very low perceived need for contraception was found amongst the respondents wishing to bear more children. The result shows that the most important factors that affect people awareness are economic factor and educational background and they jointly have the highest variance proportion and the highest Eigenvalue which is greater than 1.

In view of the above, the following recommendations were however made:

1. That government should provide social economic factors that would increase the knowledge of education among the people which will improve their level of understanding of family planning.

2. In Nigeria, the bulk of the population lives in the rural areas. The government should therefore emphasize modern methods of family planning to complement the traditional methods, provide adequate fund for family planning officials to enable them procure the necessary devices and reach the rural dwellers.
3. Due to the fact that economic factors in the country is very poor, it is advisable for the people to reduce their child bearing in order to reduce the population of the country so that the provision of government amenities will be sufficient enough for community.
4. The study also recommendstrong need for involving men in healthcare programs designed to improve women's and newborns' health as they mostly influence decision-making at the household level and this will also result in active male participation and community ownership. Young, especially first time, fathers need support and empowerment. Encouraging communication between wife and husband about family planning and birth spacing should also be part of such campaigns to promote mutual decision-making between wife and husband and make husbands responsible partners in family planning/birth spacing decisions and ease the burden of decision-making on women.

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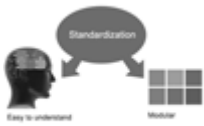
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Title: The title page must carry an instructive title that reflects the content, a running title (less than 45 characters together with spaces), names of the authors and co-authors, and the place(s) wherever the work was carried out. The full postal address in addition with the e-mail address of related author must be given. Up to eleven keywords or very brief phrases have to be given to help data retrieval, mining and indexing.

Abstract, used in Original Papers and Reviews:

Optimizing Abstract for Search Engines

Many researchers searching for information online will use search engines such as Google, Yahoo or similar. By optimizing your paper for search engines, you will amplify the chance of someone finding it. This in turn will make it more likely to be viewed and/or cited in a further work. Global Journals Inc. (US) have compiled these guidelines to facilitate you to maximize the web-friendliness of the most public part of your paper.

Key Words

A major linchpin in research work for the writing research paper is the keyword search, which one will employ to find both library and Internet resources.

One must be persistent and creative in using keywords. An effective keyword search requires a strategy and planning a list of possible keywords and phrases to try.

Search engines for most searches, use Boolean searching, which is somewhat different from Internet searches. The Boolean search uses "operators," words (and, or, not, and near) that enable you to expand or narrow your affords. Tips for research paper while preparing research paper are very helpful guideline of research paper.

Choice of key words is first tool of tips to write research paper. Research paper writing is an art. A few tips for deciding as strategically as possible about keyword search:



- One should start brainstorming lists of possible keywords before even begin searching. Think about the most important concepts related to research work. Ask, "What words would a source have to include to be truly valuable in research paper?" Then consider synonyms for the important words.
- It may take the discovery of only one relevant paper to let steer in the right keyword direction because in most databases, the keywords under which a research paper is abstracted are listed with the paper.
- One should avoid outdated words.

Keywords are the key that opens a door to research work sources. Keyword searching is an art in which researcher's skills are bound to improve with experience and time.

Numerical Methods: Numerical methods used should be clear and, where appropriate, supported by references.

Acknowledgements: Please make these as concise as possible.

References

References follow the Harvard scheme of referencing. References in the text should cite the authors' names followed by the time of their publication, unless there are three or more authors when simply the first author's name is quoted followed by et al. unpublished work has to only be cited where necessary, and only in the text. Copies of references in press in other journals have to be supplied with submitted typescripts. It is necessary that all citations and references be carefully checked before submission, as mistakes or omissions will cause delays.

References to information on the World Wide Web can be given, but only if the information is available without charge to readers on an official site. Wikipedia and Similar websites are not allowed where anyone can change the information. Authors will be asked to make available electronic copies of the cited information for inclusion on the Global Journals Inc. (US) homepage at the judgment of the Editorial Board.

The Editorial Board and Global Journals Inc. (US) recommend that, citation of online-published papers and other material should be done via a DOI (digital object identifier). If an author cites anything, which does not have a DOI, they run the risk of the cited material not being noticeable.

The Editorial Board and Global Journals Inc. (US) recommend the use of a tool such as Reference Manager for reference management and formatting.

Tables, Figures and Figure Legends

Tables: Tables should be few in number, cautiously designed, uncrowned, and include only essential data. Each must have an Arabic number, e.g. Table 4, a self-explanatory caption and be on a separate sheet. Vertical lines should not be used.

Figures: Figures are supposed to be submitted as separate files. Always take in a citation in the text for each figure using Arabic numbers, e.g. Fig. 4. Artwork must be submitted online in electronic form by e-mailing them.

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TECHNIQUES FOR WRITING A GOOD QUALITY RESEARCH PAPER:

1. Choosing the topic: In most cases, the topic is searched by the interest of author but it can be also suggested by the guides. You can have several topics and then you can judge that in which topic or subject you are finding yourself most comfortable. This can be done by asking several questions to yourself, like Will I be able to carry our search in this area? Will I find all necessary recourses to accomplish the search? Will I be able to find all information in this field area? If the answer of these types of questions will be "Yes" then you can choose that topic. In most of the cases, you may have to conduct the surveys and have to visit several places because this field is related to Computer Science and Information Technology. Also, you may have to do a lot of work to find all rise and falls regarding the various data of that subject. Sometimes, detailed information plays a vital role, instead of short information.

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3. Think Like Evaluators: If you are in a confusion or getting demotivated that your paper will be accepted by evaluators or not, then think and try to evaluate your paper like an Evaluator. Try to understand that what an evaluator wants in your research paper and automatically you will have your answer.

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21. Arrangement of information: Each section of the main body should start with an opening sentence and there should be a changeover at the end of the section. Give only valid and powerful arguments to your topic. You may also maintain your arguments with records.

22. Never start in last minute: Always start at right time and give enough time to research work. Leaving everything to the last minute will degrade your paper and spoil your work.

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26. Go for seminars: Attend seminars if the topic is relevant to your research area. Utilize all your resources.



27. Refresh your mind after intervals: Try to give rest to your mind by listening to soft music or by sleeping in intervals. This will also improve your memory.

28. Make colleagues: Always try to make colleagues. No matter how sharper or intelligent you are, if you make colleagues you can have several ideas, which will be helpful for your research.

29. Think technically: Always think technically. If anything happens, then search its reasons, its benefits, and demerits.

30. Think and then print: When you will go to print your paper, notice that tables are not be split, headings are not detached from their descriptions, and page sequence is maintained.

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33. Report concluded results: Use concluded results. From raw data, filter the results and then conclude your studies based on measurements and observations taken. Significant figures and appropriate number of decimal places should be used. Parenthetical remarks are prohibitive. Proofread carefully at final stage. In the end give outline to your arguments. Spot out perspectives of further study of this subject. Justify your conclusion by at the bottom of them with sufficient justifications and examples.

34. After conclusion: Once you have concluded your research, the next most important step is to present your findings. Presentation is extremely important as it is the definite medium through which your research is going to be in print to the rest of the crowd. Care should be taken to categorize your thoughts well and present them in a logical and neat manner. A good quality research paper format is essential because it serves to highlight your research paper and bring to light all necessary aspects in your research.

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- Fundamental goal
- To the point depiction of the research
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- Significant conclusions or questions that track from the research(es)

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Approach:

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Approach

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- Try to present substitute explanations if sensible alternatives be present.
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- Recommendations for detailed papers will offer supplementary suggestions.

Approach:

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<i>Abstract</i>	Clear and concise with appropriate content, Correct format. 200 words or below	Unclear summary and no specific data, Incorrect form Above 200 words	No specific data with ambiguous information Above 250 words
<i>Introduction</i>	Containing all background details with clear goal and appropriate details, flow specification, no grammar and spelling mistake, well organized sentence and paragraph, reference cited	Unclear and confusing data, appropriate format, grammar and spelling errors with unorganized matter	Out of place depth and content, hazy format
<i>Methods and Procedures</i>	Clear and to the point with well arranged paragraph, precision and accuracy of facts and figures, well organized subheads	Difficult to comprehend with embarrassed text, too much explanation but completed	Incorrect and unorganized structure with hazy meaning
<i>Result</i>	Well organized, Clear and specific, Correct units with precision, correct data, well structuring of paragraph, no grammar and spelling mistake	Complete and embarrassed text, difficult to comprehend	Irregular format with wrong facts and figures
<i>Discussion</i>	Well organized, meaningful specification, sound conclusion, logical and concise explanation, highly structured paragraph reference cited	Wordy, unclear conclusion, spurious	Conclusion is not cited, unorganized, difficult to comprehend
<i>References</i>	Complete and correct format, well organized	Beside the point, Incomplete	Wrong format and structuring



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