

Sources of Resistance to the Maize Weevil *Sitophilus zeamais* in Tropical Maize

Mwololo J. K.^{1,2}, Mugo S.³, Okori P.¹, Tefera T.³, Otim M.⁴ & Munyiri S. W.^{1,5}

¹ Makerere University, Kampala, Uganda

² Pwani University College, Kilifi, Kenya

³ International Maize and Wheat Improvement Center (CIMMYT), Village Market, Nairobi, Kenya

⁴ National Crops Resources Research Institute, Namulonge, Kampala, Uganda

⁵ Chuka University College, Chuka, Kenya

Correspondance: Stephen Mugo, International Maize and Wheat Improvement Center (CIMMYT), Village Market, Nairobi, Kenya. E-mail: s.mugo@cgiar.org

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Abstract

The maize weevil, *Sitophilus zeamais* (Motschulsky) (Coleoptera: Curculionidae), is among the major storage pests that enhance food insecurity among maize farmers. New sources of resistance to the maize weevil are critical in a successful breeding program to address grain damage by postharvest pests. The objective of the study was to evaluate resistance in maize genotypes to the maize weevil, and consequently their value for use in breeding programs. A total of 175 genotypes, including hybrids, landraces, open-pollinated varieties and checks, were tested for resistance to the maize weevil. The percentage grain damage, weight loss, flour weight and weight of damaged and undamaged grains were measured. Significant differences ($P < 0.001$) were observed among the genotypes for all the traits evaluated. The distribution of the genotypes among the different categories of resistance was an indication of the existence of genetic variation. The most resistant genotypes were CKPH08003 and BRAZ 2451 while the most susceptible were PH 3254 and BRAZ 4, among the hybrids and landraces respectively. Genotypes that were superior to the resistant checks were identified. The percentage weight loss and flour weight were identified as the most important insect-resistance traits for discriminating genotypes as evident from the canonical discriminant analysis. Correlation coefficients among the traits evaluated were highly significant. The resistant hybrids identified can be recommended for release and adoption by farmers, whereas the resistant landraces can act as sources of resistance for use in breeding programs.

Keywords: Maize weevil, *Sitophilus zeamais*, postharvest pests, pest resistance, tropical maize

1. Introduction

Over half of Africans earn their livelihood from agriculture (Pretty et al., 2011). Despite this, tropical African countries are among the most food-insecure in the world (Sasson, 2012). Food security could be achieved through enhancing agricultural productivity and by reducing field and postharvest crop losses. Agricultural production is seasonal, while the demand for agricultural commodities is evenly spread throughout the year. Crop storage for future use, therefore, becomes important (Thamaga-Chitja et al., 2004). Maize is an essential component of the global food security and forms a major part of the diet of millions of people. In Africa, maize is primarily grown by small-scale farmers for use as both food and feed. Its productivity is thus critical to raising rural incomes and stimulating broad-based economic growth (Byerlee et al., 1997). Maize grain storage is crucial and needed in order to maintain constant supply throughout the year. For small-scale farmers in Africa, the main purpose of storage is to ensure household food supplies and seed for planting (Adetunji, 2007).

Stored grains may suffer serious attack from pests (insects, rodents and birds) and pathogens (bacteria and fungi), especially when not protected and when storage hygiene is poor. Amidst other constraints of maize production, insect pests constitute a major threat, destroying approximately 20% of food produce (Pimentel, 2007). The damage caused by postharvest pests is much higher than that caused by other agents like rodents and micro-organisms. The maize weevils, *Sitophilus zeamais* (Motschulsky) (Coleoptera: Curculionidae), the Angoumois grain moth, *Sitotroga cerealella* (Olivier) (Lepidoptera: Gelechiidae) and the larger grain borer

Prostephanus truncatus (Horn) (Coleoptera: Bostrichidae) are the most important postharvest pests of stored cereal grains (Abate et al., 2000). In fact, in addition to direct destruction of grains through feeding and reproduction, the presence of insects has direct influence on grains, causing an increase in grain temperature and moisture contents. These lead to an increase in respiration and consequent loss in quantity and quality of the grain (Odogola, 1994). This pre-disposes the stored grain to secondary attack by disease-causing pathogens such as mycotoxin-producing fungi. The postharvest losses due to the *Sitophilus zeamais* have been recognized as an important constraint, with grain losses ranging from 20 - 90% being reported for stored untreated maize grains (Derera et al., 2001; Tefera et al., 2011a). The damage caused is irreversible and affects both farmers and traders. The adults and the larvae of the weevils are internal feeders on the maize grains and this affects seed viability and successful planting by small-scale farmers. Meeting the ever-increasing demand for sufficient quantity and acceptable quality of food, therefore, requires a reduction in damage from insects that attack the stored maize grain and seed.

Different technologies have proven effective in controlling storage insect pests, such as environmental manipulations to hinder their growth, maturation and reproduction. Such environmental manipulations can be attained by employing a number of control measures, including the use of pesticides, cultural and physical control measures (Pereira et al., 2009). Pesticides are effectively used against postharvest insect pests but are often associated with a number of drawbacks including high costs, environmental pollution and food safety, and the occurrence of resistance in the insect pests (Asawalam et al., 2006). Host-plant resistance through genetic improvements remains a cornerstone of an integrated pest-management strategy to minimize storage losses and impact on grain quality. Breeding for resistance to postharvest insect pests was initially ignored, probably due to the long duration from crop establishment to postharvest screening for resistance and the high cost involved. Genetic variation and the heritability of resistance maize weevil and other storage pests have been reported (Derera et al., 2001; Mwololo et al., 2010). This raises hope for the possibility of introducing weevil-resistance traits into elite maize germplasm. Effective breeding for resistance requires identifying resistant materials in the maize gene pool and understanding the mechanisms of resistance.

Progress has been made in developing maize cultivars resistant to post-harvest insect pests. However, the current resistance is based on a narrow source of an unimproved gene bank accession. Moreover, the current insect-resistant germplasm is not adequate since the varieties developed are lacking in one or more of the farmer-preferred agronomic traits. Breeding progress in relation to postharvest pests would depend on the extent of genetic variability within the maize germplasm. The higher the variability for a given trait, the higher the genetic gain would be expected from the selection process. There is need to search for more sources of resistance among adapted genotypes within the tropics. The use of resistant varieties against storage insect pests, when successful, would have a number of comparative advantages over other control measures, particularly the use of pesticides which have a range of shortcomings (Gemechu et al., 2011). The objective of the study was, therefore, to assess the reaction of maize hybrids, open pollinated varieties (OPVs) and landraces to maize weevil infestation and damage in order to identify sources of resistance for use in breeding programs and for deployment to farmers.

2. Materials and Methods

2.1 Germplasm

A total of 175 genotypes were used in the study, including hybrids, open-pollinated varieties (OPVs) and landraces. The 87 hybrids used comprised of commercial, drought-tolerant and quality protein maize genotypes from the International Maize and Wheat Improvement Center (CIMMYT) breeding program. The thirteen (13) OPVs and 75 landraces used in this study were sourced from the CIMMYT gene bank in Mexico. Two hybrid checks, CKPH09001 (resistant) and H513 (susceptible) checks were included in the study.

2.2 Experimental Sites

Field trials were established at the Kiboko and Embu centers of the Kenya Agricultural Research Institute (KARI). Kiboko is a dry, mid-altitude agro-ecological zone, located at 37° 75'E and 2° 15'S, situated at about 975 masl. Average rainfall in Kiboko is 530 mm, with mean monthly maximum temperature of 35.1°C and a minimum of 14.3 °C and has sandy clay soils. Embu is a moist, mid-altitude zone at 1,350 masl, located at 37° 42'E and 0° 49'S. The average rainfall in Embu is 1,200 mm with a mean monthly maximum and minimum temperature of 25 °C and 14.1 °C and has clay loam soils.

2.3 Experimental Design

The hybrids and OPVs were planted in one trial due to their similarity in architecture, while the landraces were

planted in a separate trial. The genotypes were replicated three times in a 20 x 5 and 15 x 5 alpha lattice design in 2 x 5 m row plots, for the hybrids and OPVs trial and for the landraces trial, respectively. Two seeds were planted per hill and later thinned to one, giving a plant density of 53,333 plants per hectare. In order to ensure a healthy crop, agronomic practices including weeding, fertilizer application, stem borer control and supplemental irrigation were done when necessary. At harvest, the maize was shelled and dried to obtain grains for insect-resistance bioassays.

2.4 Evaluation of Maize Germplasm for Resistance to the Maize Weevil

The insects used in the experiment were obtained from the KARI/CIMMYT Kiboko Maize Entomology Laboratory where they were reared on the grains of maize cultivar PH3253 under controlled conditions (28°C and 75% relative humidity (RH)). The maize grains were sun-dried to a moisture content of 12-13% and disinfested by fumigating with phostoxin tablets for seven days in a drum to eliminate field infestation. Samples of 100 grams of grain were taken from each plot of the three replications. The kernels were placed in 250 ml jars, infested with 50 unsexed 7-10 days old maize weevils, and incubated for 3 months at 26-28°C and 70-75% relative humidity in the Entomology Laboratory at KARI/CIMMYT Kiboko. The experimental design in the laboratory was a completely randomized design replicated three times.

The contents of each jar were sieved with mesh sieves (Endecotts Ltd, UK¹) 3 months after infestation to separate grains, insects and flour. The flour produced was measured with a weighing balance while the number of damaged kernels and adult insect progeny were counted. The grain weight loss was computed by subtracting the final from the initial weight of the grain sample and expressed as a percentage (Tefera et al., 2011b). The damaged kernels were separated from the undamaged ones using grain tunneling and holes as the criteria. These were counted and the percentage of damaged grain computed. The weight of the damaged and undamaged grains was measured. The parameters used for the maize weevil resistance assessment were the percentage weight loss, the grain damage percentage, the flour weight, and the number of insects recovered (Tefera et al., 2011b).

2.5 Statistical Analyses

A multivariate analysis of variance within a canonical variate analysis (Canonical Discriminant Analysis) was performed using statistical analysis system software (SAS) to determine the difference between genotypes and the most variable insect resistance trait among the genotypes. A univariate analysis of variance (ANOVA) for the individual resistance traits was done using the general linear model (GLM) procedure of SAS (SAS Institute, 2003). The number of insects was log transformed ($\log_{10}$), while flour produced, grain damage, weight of damaged and undamaged grains and weight loss were angular transformed to normalize their distribution before subjecting them to analysis of variance. The insect-damage traits were correlated using a canonical correlation to obtain error control and identify the strongest correlation.

3. Results

3.1 Discriminant Analysis

The ANOVA option specifies testing the hypothesis that the class means are equal using univariate statistics. The R^2 values ranged from 0.4 for the grain damage to 0.54 for the percentage flour weight and all the variables were significant at $P = 0.05$ (Table 1). The multivariate test for differences between the classes was also significant at $P = 0.05$ level, therefore, the multivariate tests of the hypothesis that the class mean vectors were equal indicate that they were not ($P < 0.001$) (Table 1). The linear combination of the centered variables were $(CAN1) 0.5438 \times \text{No. Insects} + 0.7323 \times \text{Flour weight} + 0.6272 \times \text{weight of damaged grains} - 0.8385 \times \text{weight of undamaged grains} + 0.7805 \times \text{weight loss} + 0.5863 \times \text{grain damage}$ which separates the entries most effectively (Table 1). In the first canonical variate, weight loss and/or flour weight were indicated as the most important variables for discriminating among genotypes for they had the largest coefficients. These were used to calculate the susceptibility indices as described by Tefera et al. (2011c): summing the ratio between the genotype mean values and grand means of the traits in question; divided by 2 (the number of variables). The genotypes with susceptibility index values less than 0.7 were regarded as highly resistant, 0.7-0.8 as resistant, 0.8-1.2 as moderately susceptible and >1.21 as highly susceptible. The coefficient of the weight of the undamaged grains was also high but negatively correlated to the other variables hence not suitable for discrimination of the genotypes in relation to resistance.

3.2 Resistance of Maize Hybrids and OPVs to the *Sitophilus Zeamais*

There were significant differences ($P < 0.01$) among the hybrids for grain damage, weight loss, flour weight, number of adult insects, weight of damaged and undamaged grains (Table 2). The genotype $\times$ location interaction was not significant for all the traits evaluated (Table 2). The genotypes were categorized as resistant, moderately resistant, moderately susceptible and highly susceptible based on the susceptibility index which was

computed based on the flour weight and weight loss as these were indicated as the most important traits for discrimination by the canonical discriminant analysis. The resistant genotypes had low percentage of flour weight, number of insects and high weight of undamaged grains (Table 2).

The classification shows that the genotypes were not normally distributed probably due to the fact that they were deliberately selected (Figure 1). The highly resistant and moderately resistant represented 39% of the hybrids evaluated, 36% were moderately susceptible and 25% highly susceptible. The most resistant genotype among the hybrids was CKPH08003 with susceptibility index of 0.40 which was five times lower than that of PH 3252 the most susceptible hybrid. The percentage weight loss and flour weight of the most resistant was three-fold lower than that of the most susceptible genotype and twice lower than the mean of all the genotypes. The numbers of insects were higher in the most susceptible and lower among the resistant genotypes. The resistant check CKPH9001 had a susceptibility index of 0.60 which was higher than that of the 11 genotypes categorized as the most resistant. Most of the resistant genotypes were the CIMMYT hybrids which had been bred for storage pest resistance. The most susceptible genotypes had a susceptibility index of 2.2 which was 1.9 times higher than that of the susceptible check (H513).

Table 1. Canonical discriminant analysis parameters for the number of emerged weevils, flour and grain weight, weight loss and grain damage in the 175 maize hybrids, OPVs and landraces

Variable	Pooled SD	Between SD	R-Square	F Value	Pr > F	Can 1	Can 2	Can 3
Number of insects	125.97	85.60	0.41	1.39	0.0273	0.5438	0.4247	0.1310
Flour weight (%)	1.47	1.32	0.54	2.42	<.0001	0.7323	0.5994	-0.1637
Weight-DG (%)	15.46	11.95	0.47	1.79	0.0003	0.6272	0.3225	0.5808
Weight-UDG (%)	20.87	17.97	0.52	2.23	<.0001	-0.8385	-0.1525	-0.3018
Weight loss (%)	10.03	8.08	0.49	1.94	<.0001	0.7805	-0.0894	-0.2350
Grain damaged (%)	14.87	9.96	0.40	1.35	0.0401	0.5863	-0.1465	0.3596

Note: Can = Canonical variable coefficient; SD=Standard deviation; GD=grain damage

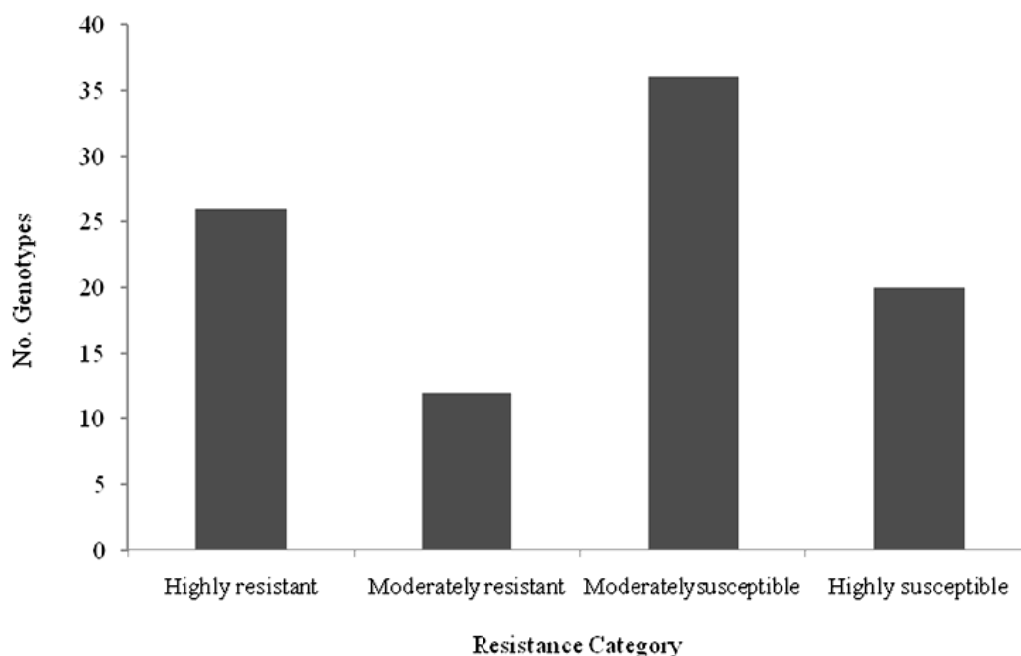


Figure 1. The distribution of genotypes according to resistance categories in the 87 maize hybrids and 13 OPVs

Table 2. Number of emerged weevils, flour and grain weight loss, grain damage and susceptibility indices for the 20 most resistant and 20 most susceptible maize hybrids and OPVs

Entry	Genotype	No. of Insects	Flour weight (%)	Weight Damaged grains (%)	Weight Undamaged grains (%)	Weight loss (%)	Grain damage (%)	Susceptibility Index
67	CKPH08003	68.92	0.48	14.28	79.48	6.24	23.50	0.40
69	CKPH09004	73.83	0.55	19.91	73.70	6.38	28.56	0.42
70	CKPH08009	83.00	0.55	15.91	77.25	6.84	25.89	0.44
71	CKPH08010	60.00	0.61	15.29	78.20	6.51	25.08	0.45
78	CKPH08028	84.67	0.68	14.25	79.73	6.02	26.48	0.46
77	CKPH08026	80.75	0.68	18.46	74.74	6.80	19.71	0.48
57	CKPH08035	83.83	0.62	20.39	71.82	7.79	32.29	0.50
73	CKPH08014	72.08	0.73	15.96	77.11	6.92	19.92	0.51
59	CKPH08037	83.17	0.66	15.93	76.27	7.80	29.62	0.52
72	CKPH08012	66.92	0.68	17.53	74.72	7.75	25.79	0.52
75	CKPH08024	62.17	0.79	14.73	78.39	6.88	20.73	0.52
76	CKPH08025	81.00	0.70	16.75	74.73	8.52	28.28	0.56
62	CKPH08040	71.42	0.80	20.93	70.24	8.83	29.84	0.60
23	500Q	92.50	0.89	17.69	74.16	8.21	33.63	0.61
54	CKPH09002	114.42	0.81	26.70	63.62	9.68	44.02	0.64
68	CKPH08004	96.17	0.90	17.65	73.31	9.04	23.76	0.64
49	CKIR06007	89.58	0.94	21.36	69.84	8.80	30.47	0.65
55	CKPH08033	103.08	0.92	25.68	65.12	9.21	42.54	0.66
53	CKPH08032	100.08	0.87	26.58	63.73	9.70	40.08	0.66
66	CKPH08002	101.25	0.95	23.84	67.01	9.15	26.34	0.66
63	CKPH08041	106.08	1.03	19.32	71.99	8.69	25.91	0.67
Mean Resistant check		84.52	0.75	19.01	73.10	7.89	28.69	0.55
	CKPH09001	78.08	0.72	24.21	66.26	9.53	38.56	0.60
	CKIR07017	155.42	2.42	41.54	44.17	14.29	45.48	1.35
	CKIR09001	234.67	2.33	39.39	43.62	16.99	53.88	1.42
46	CKIR06001	194.25	2.49	35.04	49.04	15.92	34.80	1.44
11	<i>DH04</i>	<i>230.83</i>	<i>2.26</i>	<i>39.69</i>	<i>42.06</i>	<i>18.25</i>	<i>72.79</i>	<i>1.45</i>
21	<i>531A</i>	<i>222.08</i>	<i>2.50</i>	<i>50.12</i>	<i>33.44</i>	<i>16.44</i>	<i>60.63</i>	<i>1.46</i>
14	<i>SC Duma 43</i>	<i>205.08</i>	<i>2.52</i>	<i>50.95</i>	<i>31.33</i>	<i>17.72</i>	<i>59.03</i>	<i>1.51</i>
44	CKIR09007	<i>211.17</i>	<i>2.56</i>	<i>38.57</i>	<i>44.01</i>	<i>17.42</i>	<i>48.93</i>	<i>1.52</i>
36	CKIR07008	<i>190.42</i>	<i>2.32</i>	<i>43.39</i>	<i>35.23</i>	<i>21.38</i>	<i>66.35</i>	<i>1.58</i>
18	<i>KH 600-15 A</i>	<i>192.25</i>	<i>2.22</i>	<i>42.40</i>	<i>35.27</i>	<i>22.33</i>	<i>67.27</i>	<i>1.59</i>
6	<i>H626</i>	<i>230.83</i>	<i>2.63</i>	<i>50.57</i>	<i>29.73</i>	<i>19.70</i>	<i>64.99</i>	<i>1.63</i>
33	CKIR07003	<i>198.00</i>	<i>2.76</i>	<i>35.41</i>	<i>45.41</i>	<i>19.18</i>	<i>54.92</i>	<i>1.65</i>
4	<i>H629</i>	<i>210.75</i>	<i>2.22</i>	<i>48.15</i>	<i>27.75</i>	<i>24.11</i>	<i>71.84</i>	<i>1.66</i>
3	<i>H6210</i>	<i>226.92</i>	<i>2.58</i>	<i>53.88</i>	<i>23.82</i>	<i>22.31</i>	<i>81.29</i>	<i>1.71</i>
24	<i>DK803 I</i>	<i>194.67</i>	<i>3.29</i>	<i>42.04</i>	<i>39.41</i>	<i>18.55</i>	<i>61.74</i>	<i>1.80</i>
5	<i>H628</i>	<i>196.67</i>	<i>2.72</i>	<i>48.68</i>	<i>22.38</i>	<i>27.08</i>	<i>69.91</i>	<i>1.94</i>
2	<i>H6212 SC</i>	<i>268.25</i>	<i>2.80</i>	<i>50.21</i>	<i>22.16</i>	<i>27.63</i>	<i>78.41</i>	<i>1.98</i>
13	<i>Duma 41</i>	<i>284.33</i>	<i>3.28</i>	<i>52.29</i>	<i>24.18</i>	<i>23.53</i>	<i>67.96</i>	<i>1.99</i>
17	<i>611D</i>	<i>285.00</i>	<i>3.20</i>	<i>49.14</i>	<i>26.54</i>	<i>24.32</i>	<i>69.58</i>	<i>1.99</i>
1	<i>H6213 PH</i>	<i>261.67</i>	<i>3.44</i>	<i>56.41</i>	<i>19.92</i>	<i>23.67</i>	<i>80.00</i>	<i>2.05</i>
16	<i>3253</i>	<i>255.75</i>	<i>3.96</i>	<i>47.43</i>	<i>29.76</i>	<i>22.80</i>	<i>73.35</i>	<i>2.19</i>
Mean Susceptible Check		222.45	2.72	45.76	33.46	20.68	64.16	
	<i>H513</i>	<i>190.08</i>	<i>2.17</i>	<i>46.26</i>	<i>36.73</i>	<i>17.01</i>	<i>66.88</i>	<i>1.40</i>
CV		19.09	32.79	29.74	24.18	27.49	34.8	
P Value		<0.001	<0.001	<0.001	<0.001	<0.001	<0.001	

Italicized genotypes are commercial maize hybrids

3.3 Resistance of Landrace Accessions to the *Sitophilus Zeamais*

There were significant differences ($P < 0.05$) among the landrace accessions for grain damage, weight loss, flour weight, number of adult insects, weight of damaged and undamaged grains (Table 3). The genotypes were categorized into resistant and susceptible based on the susceptibility indices adapted from the method of Tefera et al., (2011b). This showed great variation for resistance against the maize weevil among the landrace accessions.

The top four most resistant were ranked above one of the resistant check CKPH09002 which had a susceptibility index of 0.64 and the rest were categorized as moderately resistant (Figure 2). The most susceptible landraces were BRAZ 4 and RIGS GP10 with susceptibility index of 2.34 and 2.22, respectively. Their susceptibility indices were five times higher than that of the most resistant (BRAZ2451) which had a susceptibility index of 0.42. Among the landrace accessions evaluated, 23% were highly resistant to moderately resistant, 60% were moderately susceptible and 17% were highly susceptible.

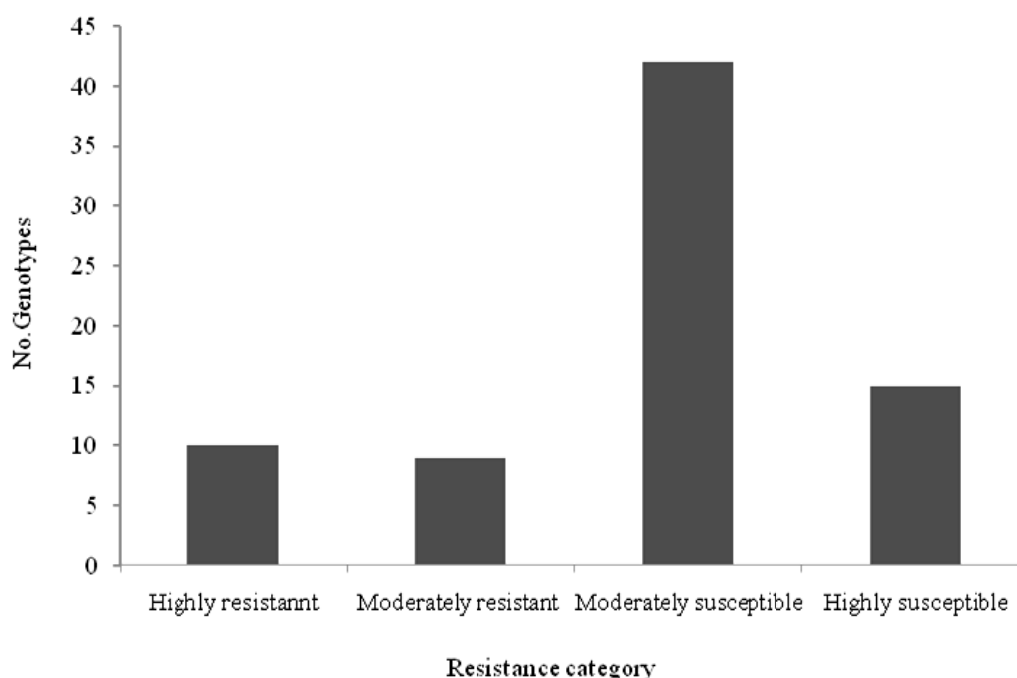


Figure 2. The distribution of genotypes according to resistance categories in the 75 maize landraces

3.4 Canonical Correlations

The probability level for the null hypothesis that all the canonical correlations were zero (0) among the variables was highly significant ($P < 0.0001$) hence the two sets of variables were highly correlated. There were great variations in the correlation coefficients between the two sets of variables (Table 4). All the set one variables (number of insects, weight of damaged and undamaged grains) accounted for a substantial amount of shared variance between the two sets of variables. The squared multiple correlation indicate that the first canonical variable of the set one variables has predictive power for number of insects ($r = 0.94$), weight of damaged grains (0.90) and undamaged grains (-0.95), weight loss (0.96), grain damage (0.95) and flour weight (0.94) (Table 4). The first canonical correlation which is the greatest possible multiple correlation with the classes that can be achieved using a linear combination of the quantitative variables was 0.9730. The likelihood ratio test for the hypothesis that the current canonical correlation and all the smaller ones are zero show that none of them was zero ($P < 0.005$) and as anticipated, the first line was equivalent to Wilk's Lambda multivariate test (0.0386). The R^2 between the first canonical variable (CAN1) and the class variable, 0.95, was much larger than the corresponding R^2 for the other two (Table 4). The number of insects and weight of damaged grains were highly and positively correlated to the percentage weight loss, flour weight and grain damage whereas the percentage weight of the undamaged grains was negatively correlated to these.

Table 3. Number of emerged weevils, flour and grain weight, weight loss, grain damage and susceptibility indices for the 20 most resistant and bottom 20 most susceptible maize landraces

Entry	Genotype	No. of Insects	Weight of damaged grains(%)	Weight undamaged grains (%)	Flour weight (%)	Weight loss (%)	Susceptibility index
59	BRAZ 2451	93.67	14.02	72.68	0.59	7.91	0.42
28	SINA 21	99.25	21.33	70.02	0.70	8.65	0.48
54	GUAN 84	111.75	19.59	69.14	0.99	7.41	0.51
11	GUAT 1162	131.92	22.98	67.94	1.01	9.08	0.57
2	GUAT 1008	131.50	27.31	62.70	1.15	9.99	0.64
53	GUAN 36	133.50	24.29	65.94	1.23	9.76	0.65
5	GUAT 1100	128.00	22.49	67.02	1.21	10.50	0.67
44	GUAT 79	145.70	21.43	67.73	1.21	10.80	0.68
42	VENE 897	122.58	21.74	67.44	1.29	10.82	0.70
27	BRAZ 1486	105.83	30.17	57.37	1.20	12.47	0.73
58	CHIS 94	112.92	19.80	70.12	1.53	10.08	0.73
4	GUAT 1014	144.17	23.77	64.46	1.36	11.76	0.75
3	GUAT 1010	128.58	32.07	56.22	1.38	11.71	0.75
52	GUAN 20	185.25	33.79	53.19	1.35	13.02	0.79
17	BRAZ 2454	152.67	24.02	63.44	1.42	12.53	0.79
66	BRAZ 1736	188.83	31.35	58.30	1.75	10.35	0.80
57	VALL 380	136.58	24.10	64.91	1.66	10.99	0.80
12	GUAT 1168	156.33	25.43	61.71	1.50	12.86	0.82
35	BRAZ 1832	166.25	30.24	56.68	1.51	13.08	0.83
	Mean	135.90	24.88	63.62	1.27	10.88	0.70
Resistant Check	CKPH0900 2	111.33	20.43	69.14	1.11	10.43	0.64
22	CHIS 114	205.42	27.93	57.93	2.24	14.14	1.05
20	SONO 72	129.75	34.73	51.97	2.46	13.30	1.08
33	VENE 352	172.96	30.59	53.23	2.15	16.18	1.10
19	BRAZ 2375	203.17	31.01	52.73	2.31	16.26	1.15
65	BRAZ 1495	188.33	33.97	49.83	2.33	16.20	1.15
7	GUAT1034	172.42	26.88	57.83	2.62	15.29	1.19
6	GUAT1030	218.50	36.96	45.73	2.41	17.31	1.21
43	VERA 197	164.88	35.12	46.59	2.46	18.30	1.26
32	BRAZ 2179	179.33	42.34	38.48	2.49	19.18	1.29
34	BRAZ 1364	215.00	39.42	42.96	2.74	17.62	1.30
61	NAYA 129	195.50	38.94	41.99	2.71	19.03	1.34
64	BRAZ 222	163.25	32.76	48.89	2.86	18.35	1.36
56	GUAN 34	195.67	27.95	50.11	2.44	21.94	1.38
36	NAYA 130	212.42	37.36	45.17	3.17	17.47	1.40
21	BRAZ 1470	224.08	33.70	44.84	3.81	21.47	1.71
41	BRAZ 1384	243.92	39.18	40.11	4.28	20.71	1.80
45	PARA GP3	323.02	44.74	31.85	4.62	23.44	1.98
55	PERU 397	253.83	42.60	33.40	4.74	24.00	2.03
23	RIGS GP10	294.83	41.49	32.93	5.29	25.58	2.22
26	BRAZ 4	265.25	44.39	33.83	6.27	21.79	2.34
	Mean	211.76	36.43	44.77	3.16	18.80	1.45
Susceptible Check	H513	190.08	46.26	36.73	2.20	17.01	1.40
	CV	17.87	31.97	30.30	32.50	33.80	
	P Value	P<0.01	P<0.01	P<0.01	P<0.01	P<0.01	

Table 4. Canonical correlations parameters between set 1 (No. of insects, weight of damaged and undamaged grains) and the set 2 (weight loss, flour weight and grain damage) variables

Can	Can corr	SE	Squared can corr	Eigen value	Variance	Cum variance	LR	F Value	Pr > F	No. of insects	Weight DG (%)	Weight UDG (%)	Weight loss (%)	Grain damage (%)	Flour weight (%)
1	0.9730	0.005	0.95	17.79	0.98	0.98	0.04	49.67	<.0001	0.9433	0.9040	-0.9580	0.962	0.949	0.958
2	0.4330	0.082	0.19	0.23	0.01	0.99	0.73	5.43	<.0001	0.0064	0.1318	0.0527	-0.225	0.242	-0.051
3	0.3256	0.090	0.11	0.12	0.01	1.00	0.89	5.63	0.0049	0.0793	-0.0567	0.0389	-0.153	-0.201	0.281

1. Test of H0: The canonical correlations in the current row and all that follow are zero;
2. Key: Can corr = Canonical correlation; SE=Standard error; Cum=Cumulative; LR=Likelihood ratio; DG=damaged grains; UDG=undamaged grains;
3. *Can: Canonical variable coefficient.

4. Discussion

The results from the study show that there is great variation in resistance to the maize weevil in the maize germplasm evaluated. The percentage weight loss was the most important variable for discriminating germplasm for resistance to the maize weevil. The flour weight, number of emerged insects at the end of screening period, and percentage grain damage in that order, were highly correlated as indicated by the canonical correlation analysis. Consequently, these four variables can be integrated to calculate a susceptibility index as reported by Tefera et al. (2011c). The resistant genotypes had low percentage of weight loss, grain damage and reduced number of insects. The low weight loss among the resistant genotypes indicates that mechanisms conferring resistance against the maize weevil exist. The resistance could either be due to physical factors such as grain hardness or antibiosis as a result of biochemical compounds which are toxic to the insects (García-Lara et al., 2004; Siwale et al., 2009). As reported by Abraham et al. (1991), the level of damage during storage depend on the number of emerging adults, and the grains permitting faster and high levels of adult emergence. The hybrids which showed low levels of susceptibility can be stored for relatively longer periods of time. Resistant varieties, therefore, can be adopted as an environmentally friendly way to reduce damage by the maize weevil under the widely used storage conditions and facilities in the tropics. The parental lines of the resistant hybrids (CKPH08003, CML202/CML204, CKPH09004, CKPH08009, CKPH08010, CKPH08028, CKPH08026, CKPH08035) identified can be used as sources of resistance in breeding for resistance to postharvest insect pests. The use of resistant maize varieties leading to low grain weight loss would reduce the negative impact of the maize weevil. Host plant resistance can, therefore, be used as a vital component of an integrated pest management strategy against postharvest insect pests.

The commercial cultivars which have been bred for the mid to high altitude were the most susceptible to maize weevil attack among the hybrids evaluated. This indicates that integration of resistance to storage pests has not been routinely used by many breeding programs, thereby leading to high postharvest losses despite increased yields. According to Derera et al. (1999), most breeding programs have been addressing yield increase at the expense of storage pest resistance. Consequently, it is critical that breeding for resistance to postharvest insect pests be prioritized as equally important to safeguard farmers from huge losses due to maize weevil attack. This would empower the farmers economically through reduced use of storage pesticides thus saving costs as well as avoiding negative environmental impacts associated with pesticides. Superior hybrids were identified among the genotypes for they performed far much better than the susceptible hybrid check. These can be improved further to incorporate the farmer preferred traits such as high yields and resistance to other diseases followed by release to the farmers. For instance, the CKPH08028 and CKPH08043 have been improved and released as MTPEH0701 and EMB0701 for the lowlands and midlands, respectively. These hybrids should be promoted for more adoption by farmers in those ecological zones.

The open pollinated varieties were categorized as moderately resistant to highly susceptible. The great variation observed in the germplasm evaluated forms a genetic resource base for further improvement to raise the levels of resistance to storage pest while conserving the farmer preferred traits. The classification of the landraces into resistant to highly susceptible followed a normal distribution, an indication of their variability. The great variation in response to the maize weevil attack gives evidence of the existence of genetic diversity. Hence a rich genetic resource base for breeding for resistance therefore exists. Landraces are enriched with vast genetic variability since they have not been subjected to selection over a long period of time; aiding in their adaptation to different agro-ecological niches which makes them important as genetic resources for useful traits (Yamasaki et

al., 2007). According to Dhillon and Pixley (2003), inheritance for resistance to the *Sitophilus zeamais* is mostly additive and non-additive to a lesser extent. Landraces are considered to be important genotypes, representing the original biological material created by the process of natural selection and adapted to local growing conditions. The highly resistant landraces identified: (BRAZ 2451, SINA 21, GUAN 84, and GUAT 1162) can be used in developing inbred lines and hybrid varieties which are resistant to the maize weevil.

5. Conclusion and Recommendations

Resistant maize hybrids and landraces were identified from the large number of germplasm screened. This indicated that breeding for resistance to the maize weevil is not only achievable, but has already resulted in resistant hybrids being released in Kenya. These could be grown to reduce damage by the maize weevil in storage. The resistant landraces could be used as new sources of resistance. It is recommended that proper links be effected between plant entomologists and plant breeders in understanding the mechanisms of resistance and the heritability of the traits, to enhance synergism in breeding varieties with improved resistance.

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