

Supplementary Table S1. Traits measured, their description and unit

S/N	Traits	Traits Abb.	Trait description	Unit
1	Days to anthesis	Anthesis	The number of days from planting to the time when 50% of the plants have anther shedding pollen	Count
2	Days to silking	D.S.	The number of days from planting to the time when 50% of the plants have emerged silk	Count
3	Anthesis silking interval	ASI	The interval in days between dates of silking and anthesis	Count
4	Plant height	PLHT	The distance from the base of the plant to the height of the first tassel branch	cm
5	Ear aspect	EASP	It is scored on a scale of 1-5 , where 1= clean, uniform , large and well-filled ears and 5 = rotten small partially filled ear	Scale
6	Plant aspect	PASPUN	It is scored on a scale of 1-9 , where 1= clean, uniform , large and well-filled ears and 9 = rotten small partially filled ear	Scale
7	Ear per plant	EPP	Total number of ears with at least one fully developed grain divided by the number of harvested plants	Count
8	<i>Striga</i> emergence count	STRCO 1 & 2	Number of emerged striga plant per plot at 8 and 10 weeks after planting [WAP]	Count
9	<i>Striga</i> damage rate	STRRAT 1 & 2	It is scored on a scale of 1-9 , where 1= no visible host plant damage symptom and 9 = all leaves are completely scorched and finally dead plants. Taken at 8 and 10 WAP	Scale
10	Grain yield	Yield/G.Y.	Calculated from ear weight and grain moisture	kg

Supplementary Table S2. Mean squares from the combined ANOVA for traits recorded under infestation condition for 2years evaluated in 2 locations

Source	Df	Plant height	Ear Aspect	Ear per plant	Striga rate 8 WAP	Striga rate 10 WAP	Emerged Striga 8 WAP	Emerged Striga 10 WAP	Grain Yield [kg ha ⁻¹]
Geno	149	1347.57 [§]	1.27 [§]	0.25 [§]	7.04 [§]	7.65 [§]	1529.72 [§]	2433.63 [§]	3029370.5 [§]
Env	3	24923.75 [§]	7.35 [§]	4.26 [§]	174.40 [§]	427.67 [§]	5672.25 [§]	39063.45 [§]	55554413.3 [§]
Rep[Env]	4	1542.45 [§]	1.18 [§]	0.19 [§]	1.08	2.86**	91.15	282.23	2934078.5 [§]
Blk[Rep*Env]	112	305.96 [§]	0.32 [§]	0.06 [§]	1.07 [§]	1.40 [§]	454.29 [§]	1117.56 [§]	685188.6 [§]
Geno*Env	446	140.25***	0.29 [§]	0.03	1.18 [§]	1.24 [§]	269.12***	531.70**	546418.6 [§]
Error	483	106.25	0.18	0.03	0.63	0.72	202.1	421.76	356115
CV[%]		8.26	14.22	22.25	19.62	14.35	60.95	58.84	33.28

*Significant at $p \leq 0.05$ level,

**Significant at $p \leq 0.01$ level,

***Significant at $p \leq 0.001$ level,

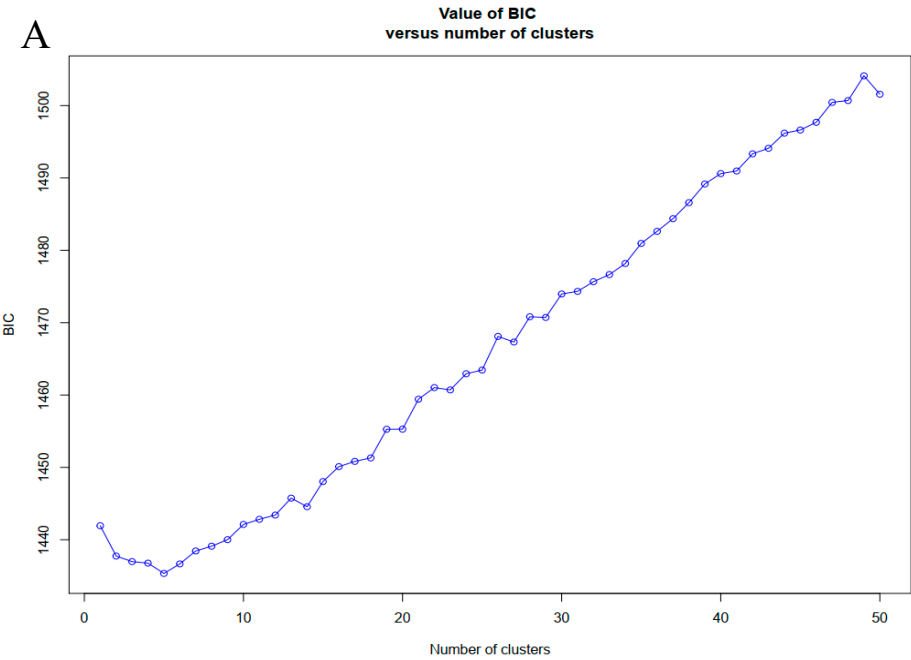
[§]Significant at $p \leq 0.0001$ level respectively. Blk = Block, Env= Environment, Rep = Replication, CV = Coefficient of Variation, WAP = weeks after planting. ns = data not significant

Supplementary Table S3. Mean and standard errors of variables under *Striga* infestation and *Striga* non-infestation across the entire four environments

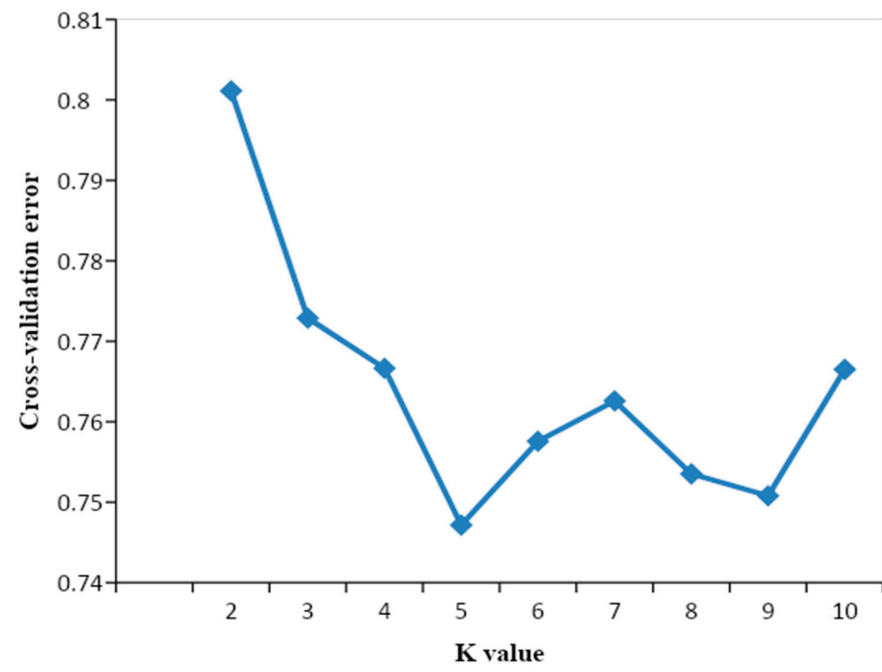
Variable	Min	Max	Mean	Std error	Heritability
Grain yield infested	13.00	3299	1580	270.5	0.83
Plant height (cm)	89.25	151	124.75	4.25	0.90
Ear per plant (number)	0.00	1.20	0.78	0.06	0.88
Ear aspect (1-5)	2.00	4.75	3.00	0.19	0.78
Striga damage rating 8 WAP	3.00	8.00	4.05	0.39	0.85
Striga damage rating 10 WAP	4.00	9.00	5.91	0.4	0.85
Emerged Striga count 8 WAP	1.00	73.00	23.31	5.85	0.85
Emerged Striga count 10 WAP	3.00	88.00	34.88	8.29	0.8

WAP=weeks after planting

Supplementary Figure S1: Graph [A] Bayesian information criterion versus the number of clusters and [B] cross-validation error versus K value



B



Supplementary Table S4. Grouping of the 150 inbred lines using 16,735 SNPs

Groups	Inbred lines	Grain color	Source population
1	22	White	ZDIP
2	27	White	IWDS
3	8	White	TZEC
4	54	White and Yellow	Mixed
5	39	White and Yellow	TZLC

Supplementary Fig S2: Mantel correlation test performed between the different dissimilarity matrices alongside the combined matrix. Sim = similarity coefficient, beacon = Mantel threshold indicating the correlation.

