

Evaluation of Western Ethiopian Sorghum Landraces for Resistance to *Striga hermonthica* (Del.) Benth

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Out of the initial 48 sorghum genotypes tested in pots, a total of 33 sorghum genotypes were selected for further evaluation in a sick-plot specifically designed to measure their resistance to *S. hermonthica*. Additionally, another resistant check, Framida variety, was included in this evaluation. The sick-plot trial was conducted at the Assosa Agricultural Research Center [7].

For further validation of the pot and sick-plot trials, seven sorghum genotypes, including resistant check varieties like Berhan and Framida, as well as promising resistant landraces such as ETS102969, ETS102970, and ETS102975, alongside susceptible check varieties like Assosa-1 and ETS102957 were selected and evaluated under hot-spot farmers' fields in three locations at Assosa, Benishangul Gumuz, Ethiopia [8].

The *Striga* seeds used in the study were collected over a period of three years, from 2019 to 2021. These seeds were obtained from sorghum fields that were highly infested with *S. hermonthica* in various districts of Assosa Zone, specifically Bambasi, Abramo, and Ura districts. After collection, the seeds were stored in glass jars, kept in a dark environment at room temperature until they were needed for infesting the pots and sick plots [9].

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A trial was established at the Assosa Agricultural Research Center (AsARC), situated in Assosa Zone, Benishangul Gumuz region, Ethiopia. The pot trials were laid out in randomized complete block design having two replications in 2020 and 2021 under the Lath-house condition of AsARC. Sand/peat/compost (1:3:1 v/v) mix soil was used to fill 96 pots. Each pot was infested with 4 mg *S. hermonthica* seeds at 5 cm depth and covered with a thin layer of the mix soil. After a 10 day delay to precondition the *Striga* seed, six sorghum seeds of each genotype were sown in the pot and later thinned to three plants per pot [10].

A total of 33 sorghum genotypes were also evaluated in *S. hermonthica* sick plot of AsARC in 2022. The trial site was ploughed twice and furrows with 70 cm spacing were prepared. The trial was laid out in RCBD having two replications. The furrows within the trial unit (2 m x 1.40 m) were uniformly infested by *S. hermonthica* seeds collected during 2021 cropping season, Ethiopia. The infested *Striga* seeds were covered with a thin layer of soil and preconditioned for 10 days. After preconditioning, sorghum genotypes were sown within the furrows at a rate of 10 kg ha⁻¹. Apart from *Striga*, other weeds were t si11k plots [9].

emergence compared to the resistant check variety Framida in the sick-plot trial. The results are consistent with the findings from a previous pot trial [16-20].

Result of this study revealed that the resistant sorghum genotypes were early maturing with maturity period of 125 days for the Berhan, 142 days for ETS102970, and 144 days for ETS102969. It is argued with the finding of Ayana et al. (2019) that reported early maturing sorghum genotypes showed resistance to *S. hermonthica*. Franke et al. (2006) was also reported that earlier maturing sorghum genotypes had positive response to Striga stress.

Moreover, *S. hermonthica* resistant sorghum genotypes of this study were shorter plant heights that ranged from 101.96 cm for ETS102969 to 139.79 cm for Berhan. Similarly, they have lower number of leaves per plant that ranged from 3.45 for ETS102969 to 5.87 for Berhan [21-23].

As illustrated the genotypes ETS102970, Berhan, and ETS102969 have lower ASNPC values compared to other genotypes, indicating a slower or less severe emergence of Striga plants in these resistant sorghum genotypes. On the other hand, genotypes ETS102957 and ETS102944 exhibit higher ASNPC values, suggesting a higher incidence and more rapid emergence of Striga plants in these susceptible sorghum genotypes. The resistant checks (Berhan and Framida) also demonstrate low ASNPC values, indicating their resistance to *S. hermonthica* infestation. These findings further

support the potential resistance of sorghum genotypes ETS102970, Berhan, and ETS102969 against *S. hermonthica* infestation in Assosa, Benishangul Gumuz region of Ethiopia (Figures 1-3).

Va da a a - fa e ' e d

As illustrated in the results of the validation trial confirmed that the resistant check Berhan and sorghum landrace ETS102969 have the lowest number of emerged *S. hermonthica* plants per 4m x4m plot. Additionally, sorghum landrace ETS102970 has a lower number of emerged *S. hermonthica* plants compared to the second resistant check Framida. The number of *Striga* per sorghum plant is also low for Berhan, ETS102969, and ETS102970 [24].

On the other hand, susceptible checks ETSL102957 and Assosa-1 variety have the highest count of emerged *S. hermonthica* plants. These findings indicate that Berhan ETSL102969, and ETSL102970 exhibit promising resistance against *S. hermonthica* infestation. Overall, this validation trial confirms that these sorghum landraces possess comparable or good resistance to *S. hermonthica* when compared to resistant checks like Berhan and Framida [25] (Table 3).

The promising sorghum landraces ETS102969 and ETS102970 have been found to have higher yield compared to resistant checks in a validation trial. Additionally, the bold seed size and white seed colour of sorghum landrace ETS102969 are desirable traits by the local farming communities. These traits can be advantageous for the breeding program, as they can be combined with the Striga resistance

trait to develop sorghum varieties that have both resistance to *Striga* infestation and the preferred seed characteristics. By incorporating these additional traits into the breeding program, there is a higher probability of obtaining F-generations that exhibit both white colour and bold sized seeds, along with resistance to *S. hermonthica*. is

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21. ~~DATA CITATION~~