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Performance of sorghum genotypes under zero tillage conditions in rice fallows with reference to stem borer *Chilo partellus*

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A field experiment was carried out to screen the sorghum genotypes against stem borer in rice fallow under zero tillage condition. Based on mean stem tunnel length, the genotypes were categorized as least susceptible (0–5 cm), moderately susceptible (5–10 cm) and highly susceptible (>10 cm). The resistant check CSH 16 (C) was found least susceptible with 4.65 cm, whereas NTJ-2 (C), NLCW-6 and N-14 were found to be highly susceptible as they recorded 10.45, 10.46 and 11.44 cm mean stem tunnel length respectively. The

remaining genotypes were moderately susceptible with 6.60–9.84 cm mean stem tunnel length. There was non-significant positive correlation between the number of larvae and leaf damage, dead hearts, stem tunnelling, white ears and chaffy grains respectively, but it was negative for tiller damage.

Keywords: Genotypes, sorghum, stem borer, stem tunnel length, susceptible.

SORGHUM [*Sorghum bicolor* (L.) Moench] is the fifth major cereal crop after wheat, rice, maize and barley. It is the most important crop of Asia, Africa, Australia, America and is cultivated as a staple crop in the semi-arid tropics (SAT). In India, it is cultivated in an area of 6.18 m ha with 5.33 million tonnes (mt) production and productivity of 863 kg/ha (ref. 1). In general sorghum is cultivated during *kharif*, maghi (late *kharif*) and *rabi* seasons in Andhra Pradesh (AP) in an area of 287,000 ha with production of 546,000 tonne and productivity of 1904 kg/ha (ref. 2) as against normal area of 760,000 ha with production of 552,000 tonne and productivity of 730 kg*/*ha. The reasons for low productivity under normal type of cultivation might be due to shifting of jowar area to cultivation of commercial crops, high humidity in the coastal regions, and ravage due to pests and diseases in jowar-cultivating areas.

 Insect pest conditions are dynamic in nature and change with climate and farming practices – introduction of improved varieties has been known to result in pest outbreaks or changes in pest status³. Sorghum is attacked by more than 150 insect species causing 32% crop loss⁴. Losses in sorghum due to insect pests differ on a regional basis and have been estimated at US\$ 1089 million in the SAT, US\$ 250 million in USA and US\$ 80 million in Australia⁵. Among the insect pests, shoot fly, *Atherigona soccata* (Rondani) and stem borer, *Chilo partellus* (Swinhoe) are the major threats causing 75.6% and 24.3–36.3% yield loss respectively⁶.

 Management of the pests is being done using pesticides. However, due to the adverse effects of pesticides, it is imperative to seek for alternate integrated pest management methods like host plant resistance, which it not only cost-effective and does not require application skills in pest control techniques, but also enhances the effectiveness of natural enemies and reduces the need to use pesticides⁷. The effect of resistant genotypes on insect population is continuous and cumulative over time. Umakanth *et al.*⁸ reported 'SPV 1022', 'PKV809' and 'CO28' as promising sorghum cultivars in rice fallows.

 Performance of sorghum genotypes under zero tillage conditions in rice fallows with reference to stem borer was carried out during *rabi* 2014–15 in the southern block of Agricultural College Farm, Bapatla, Guntur district, AP. Studies were carried out to screen the sorghum genotypes against shoot fly in rice fallow under zero

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tillage condition. Twenty genotypes were procured from the Directorate of Sorghum Research, Hyderabad, Telangana, and Regional Agricultural Research Station (RARS), Nandyal, AP to be used as source material for the screening study. The experiment was laid out in randomized block design at Agricultural college Farm, Bapatla and the treatments were replicated twice. The crop was sown on 7 January 2015. The length of each line was 4 m and spacing between two lines of each genotype was 45 cm; also intra-row spacing adopted was 15 cm.

 Observations were recorded on number of larvae per plant by destructive sampling at vegetative, flowering – grain formation and harvesting stages, dead hearts caused by *C. partellus* (number of plants with dead hearts symptoms and total number of plants were recorded from each plot based on which per cent dead hearts was calculated from 30 to 60 DAS at weekly intervals), per cent damaged leaves (number of leaves with leaf injury symptoms like scraping, shot holes and total number of leaves were recorded from each plot based on which per cent leaf injury was calculated from 30 to 60 DAS at weekly intervals), damage caused by *C. partellus* in tillers (damaged tillers and total number of tillers per hill were recorded and per cent tiller damage was calculated) and stem tunnelling (at the time of harvesting, by destructive sampling, the main stems of plants infested with *C. partellus* were split open from the base to the apex, and cumulative tunnel length and stem length were measured; cm). The percentage tunnelling was calculated using the formula given below.

Length of tunnelling (cm) × 100.
Total length of stem (cm)

Genotypes were categorized based on stem tunnelling according to Rajasekhar and Srivastav⁹ (Table 1).

 The number of white ears/plot (number of white ears due to stem borer infestation per plot from the total number of plants sampled was recorded and percentage white ears was calculated), chaffy grain percentage (total number of grains from each ear head and chaffy or under developed grains in randomly selected five ear heads from each plot were counted and per cent chaffy grain was calculated) and number of larvae and pupae per stubble after harvest (to know the carry-over population of *C. partellus*, data on number of larvae and pupae per stubble after harvest were recorded) were recorded.

Table 1. Particulars of stem tunnel length range for the damage by stemborer

Range of mean tunnel length (cm)	Attribute		
$0 - 5$	Least susceptible		
$5 - 10$	Moderately susceptible		
>10	Highly susceptible		

 Genotypes exhibited significant variation pertaining to larval incidence during their crop growth period. The number of larvae per plant recorded at vegetative stage ranged from 5.00 to 8.90. The highest number of larvae was recorded in genotype CSH 23 (8.90) followed by CSV 15 (8.50), SSV 84 (7.80), CSH 14 (7.80) and CSH 20MF (7.50), whereas the lowest number of larvae was recorded in genotypes CSH 24MF, CSH 25, NTJ-4 (C) (5.00 each) followed by CSH 13 (5.10), N-14 (5.20) and BRJ-358 (5.60) compared to the resistant check CSH 16 (5.80) and Mahalaxmi 296 (6.80).

 At flowering stage, the number of larvae per plant ranged from 4.50 to 11.40. The highest number of larvae was recorded in genotype SSV 84 (11.40) followed by CSH 22SS (10.90), CSV 26 (9.90) and CSV 24SS (9.60). The lowest number of larvae was recorded in genotype N-14 (4.50) followed by NTJ-2 (C) (5.10), N-13 (5.90) and resistant check CSH 16 (C) (6.20), when compared to the popular check Mahalaxmi 296 (7.90) (Table 2).

 At harvesting stage, the number of larvae per plant ranged from 3.20 to 7.60. The highest number of larvae was recorded in NTJ-4 (C) (7.60) followed by NLCW-6 (6.70), NLCW-8 (6.60) and BRJ-358 (6.50), whereas the lowest number of larvae per plant was recorded in genotype CSV 15 (3.20) followed by CSH 23, CSH 30 (3.40 each), CSV 29R (3.50) and SSV 84 (3.70) compared to the resistant check CSH $16(4.0)$, NTJ-3 (4.40) and popular check Mahalaxmi 296 (4.80). Mohan *et al.*10 reported the highest seasonal incidence of *C. partellus* on variety HC-136 and JS-20 during *rabi*-summer and *kharif*, and high larval and pupal populations during *kharif* season crop than in *rabi*-summer. Adverse effect of resistant genotypes on insect development resulted in low larval mass due to nutritional abnormalities and/or because of poor food utilization by the larvae of resistant varieties 11 . Painter¹² suggested that with rare exceptions, the feeding of insects during the developmental stages on resistant varieties results in individuals that are smaller and having less weight. The sorghum varieties appear to possess some antibiotic factors, which exist either in the leaves or in the stem, or in both, and adversely influence larval duration¹³.

 Prolongation of larval period ultimately results in reduction of the number of generations in a season/year. The adverse effects on the post-embryonic development of stem borer might possibly be because of the antibiotic $factors¹⁴$. Thus, it can be concluded that the adverse effects of the resistant genotypes on larval and pupal mass, prolonged larval and pupal period and low population and adult emergence may be due to some nutritional abnormalities.

 At 42 DAE, the number of dead hearts and per cent dead hearts ranged from 0.00 to 0.02 and 0.00 to 2.22 respectively. The highest number of dead hearts and per cent dead hearts were observed in genotypes CSV 22 (0.02 and 2.22), and N-13 (0.02 and 2.18) followed by CSV

RESEARCH COMMUNICATIONS

216R (0.01 and 1.14), CSH 25 (0.01 and 1.00), N-14 and popular check Mahalaxmi 296, resistant check CSH 16 (C) (0.01 and 1.02) and CSV 24SS (0.01 and 1.14). No infestation was recorded in the remaining genotypes (Table 3).

 At 49 DAE, the number of dead hearts and per cent dead hearts ranged from 0.00 to 0.04 and 0.00 to 2.44 respectively. The highest number of dead hearts and per cent dead hearts were observed in genotype N-13 (0.04 and 4.37) followed by CSV 22 (0.02 and 2.44), CSV 216R, popular check Mahalaxmi 296 (0.02 and 1.61), CSV 216R (0.02 and 0.96) and N-14 and CSH 25 (0.01 and 1.00). No infestation was recorded in the remaining genotypes.

 Similar trend reaction was noticed at 56 and 63 DAE. The number of dead hearts and per cent dead hearts ranged from 0.00 to 0.05 and 0.00 to 4.92 as well as 0.00 to 0.06 and 0.00 to 5.59 respectively. Less number of dead hearts was recorded in the tested genotypes.

 The present study reveals that the per cent dead hearts range is very low. These findings are in conformity with those of Hussian *et al.*¹⁵, who recorded lowest stem borer dead hearts in genotype CSH 23 (4.87) and Vyas *et al.*¹⁶ who recorded 2.39% in CSV 21F and 3.58% in CSH 20MF in the *kharif* season. The third-instar larvae migrate to the base of the plant, bore into the shoot and damage the growing point resulting in the formation of dead heart.

 The harvested stems (stalk) of sorghum were stored for fodder purpose while stubbles (base of stem) were left in the field. The carryover of *C. partellus* was accomplished through the stages like larvae or pupae embedded in the stubbles and stem (stalk). The number of larvae per stubble ranged from 0.30 to 1.80. The highest number of larvae per stubble was recorded in genotype CSH 24MF (1.80) followed by N-14 (1.58), CSV 216R, NTJ-3 (C) (1.50 each) and CSV 29R (1.40), while the lowest number of larvae per stubble was recorded in the least susceptible check CSH 16 (C) (0.30). The genotypes BRJ-358 (0.40) and CSH 23 (0.50) were on par with CSH 16 compared to the popular local check Mahalaxmi 296 (1.3).

 The number of pupae per stubble ranged from 0.40 to 1.70. The highest number of pupae per stubble was recorded in genotype BRJ-358 (1.7) followed by CSV 17 (1.5), NLCW-8 (1.4) and CSH 16 (C), CSH 14, NLCW-12 (1.2 each). The lowest number of pupae per stubble was recorded in genotype CSV 26 (0.4) followed by CSH 22SS, CSH 30 (0.60 each), popular check Mahalaxmi 296, CSH 24MF, CSV 22, NTJ-3 (0.70 each) and CSV 24SS, CSV 23, CSV 216R, CSV 15, N-14 (0.80 each) (Table 2). These results indicate that the stem borer survives in crop residues during off-season. Patel and Purohit¹⁷ recorded 16.24% larvae in stalk and 11.25% pupae in the variety CSH 16.

The authors¹⁷ also recorded mean per cent hibernation of stem borer larva and pupa through stalk as 13.75 and 2.25 respectively, whereas in case of hibernation through stubble it was 8.96 and 4.17 respectively. Thus carryover of stem borer through larval stage was higher than that of pupal stage in the *rabi* season. The hibernation of stem borer through stalk was more (16.04%) than that of stubble (13.13%) in the *rabi* season. Overall, hibernation of stem borer (larva + pupa) through stubble was least in variety GJ 41 (20.0%) and highest in GJ 38 (33.75%). Also, hibernation of stem borer through stubble was more (26.88%) than that of stalk (20.21%) in *kharif* season.

 Leaf damage caused by *C. partellus* infestation is indicated by the appearance of small, elongated windows in young whorl leaves, where the young larvae have eaten the upper surface of the leaves due to stem borer scraping. The leaf damage at 42 DAE ranged from 0.00% to 33.86%. The highest per cent leaf damage was recorded in genotype CSH 22SS (33.86) followed by N-13 (29.5), CSH 25 (28.0) and NTJ-3 (C) (27.0) , while the lowest was recorded in genotype followed by NTJ-1 (C) (4.0), CSV 17 (5.0) and CSH 16 (C) (5.71) compared to the popular check Mahalaxmi 296 (23.52). NLCW-12 showed no leaf damage (Table 3).

 At 49 DAE leaf damage ranged from 1.75% to 18.86%. The genotypes CSV 22 (18.86), SSV 84 (16.67), CSH 23 (16.10) and NTJ-4 (C) (16.00) recorded highest per cent leaf damage, whereas CSH 25 (1.75), NTJ-2 (1.80) and NLCW-12 (1.85) recorded lowest per cent leaf damage compared to the checks CSH 16 (4.50), NTJ-1 (4.50) and NTJ-3 (14.33). Popular local check Mahalaxmi 296 recorded 10.67% leaf damage.

 At 56 DAE, leaf damage ranged from 4.29% to 21.19%. The highest per cent leaf damage was recorded in the genotype CSV 22 (21.19), popular check Mahalaxmi 296 (20.14) and CSH 14 (17.74), while the lowest was recorded in genotypes NLCW-12 (4.29), N-14 (5.54) and NLCW-6 (7.17) compared to the least susceptible check CSH 16 (8.86).

 At 63 DAE, leaf damage ranged from 4.11% to 18.87%. The highest per cent leaf damage recorded in the genotypes SSV 84 (18.87), CSV 24 SS (18.21), popular check Mahalaxmi 296 (18.17) and CSV 216R (16.71), while genotypes N-14 (4.11), NLCW-12 (5.18), BRJ-358 (5.36) and NLCW-6 (6.94) recorded the lowest per cent leaf damage compared to the resistant check CSH 16 (8.04). These findings are in conformity with those of Muturi *et al.*¹⁸, who reported 19.9% to 60.9% leaf damage by stem borer and Vyas *et al.*16 who reported 4.47% damaged leaves in the genotype CSH 20MF and 4.78% damaged leaves in genotype CSV 21F. The results of the present study reveal that leaf damage reduced with crop growth. Marulasiddesha *et al*. 19 reported that the genotype SSV-7073 showed significantly less dead heart, leaf scraping, pinhole, peduncle or stem tunnelling damage compared to all other genotypes.

 The number of damaged tillers and per cent tiller damage ranged from 1.60 to 2.30 and 30.77 to 46.10

CURRENT SCIENCE, VOL. 117, NO. 4, 25 AUGUST 2019 697 **Table 4.** Reaction of sorghum genotypes against *C. partellus* infestation at different crop growth stages Stem Mean No. of Mean Stem borer % Leaf No. of tillers/ Damaged % Tiller tunnelling stem tunnel white ears White chaffy grain Genotype DH%# damage# plant† tillers† damage# percentage# length (cm) Attribute per plot† ears %# percentage# CSV 24SS 3.41 (7.57) 18.21 (25.25) 5.10 (2.25) 2.00 (1.41) 39.22 (39.20) 5.56 (13.64) 8.78 MS 0.11 (1.84) 10.90 (19.15) 8.33 (16.58) CSH 22SS 1.96 (8.05) 15.36 (22.43) 5.60 (2.36) 2.40 (1.55) 42.86 (41.69) 4.78 (12.64) 9.45 MS 0.05 (1.38) 4.88 (12.72) 8.12 (16.32) CSV 23 1.96 (5.71) 8.97 (17.40) 5.70 (2.39) 2.00 (1.41) 35.09 (36.39) 5.71 (13.83) 9.13 MS 0.07 (1.63) 6.92 (15.23) 15.45 (23.13) CSH 20MF 1.00 (4.07) 11.37 (19.11) 6.50 (2.55) 2.40 (1.55) 36.92 (37.50) 5.12 (13.08) 8.98 MS 0.00 (1.00) 0.00 (0.00) 12.50 (20.60) CSH 24MF 4.92 (12.81) 6.96 (15.21) 6.50 (2.55) 2.40 (1.55) 36.92 (37.43) 3.53 (10.84) 6.10 MS 0.08 (1.64) 8.50 (16.95) 13.64 (21.63) CSV 17 1.96 (5.71) 6.79 (15.03) 6.00 (2.45) 2.30 (1.52) 38.33 (38.26) 8.53 (16.99) 8.37 MS 0.05 (1.51) 4.98 (12.85) 13.33 (21.25) SSV 84 0.96 (3.99) 18.87 (25.75) 6.50 (2.55) 2.50 (1.58) 38.46 (38.26) 5.81 (13.95) 9.35 MS 0.09(1.74) 8.65 (17.09) 5.42 (13.41) CSV 216R 5.59 (13.68) 16.71 (24.14) 5.00 (2.23) 1.80 (1.34) 36.00 (36.83) 4.39 (12.10) 8.87 MS 0.03 (1.37) 3.27 (10.39) 8.57 (16.89) CSV 15 1.92 (5.66) 12.80 (20.93) 5.10 (2.26) 2.30 (1.51) 45.10 (42.74) 5.74 (13.87) 8.07 MS 0.08 (1.74) 8.30 (16.70) 9.36 (17.72) CSH 14 4.42 (12.12) 13.51 (21.32) 5.70 (2.38) 2.10 (1.45) 36.84 (37.44) 5.99 (14.17) 8.34 MS 0.06 (1.63) 6.35 (14.52) 9.60 (17.98) CSV 22 4.66 (11.92) 14.15 (22.06) 4.00 (1.98) 1.60 (1.26) 40.00 (40.54) 4.39 (12.11) 8.57 MS 0.10 (1.75) 9.88 (18.33) 8.67 (17.06) CSV 26 0.00 (0.00) 12.22 (20.47) 5.80 (2.41) 2.40 (1.55) 41.38 (40.27) 3.38 (10.60) 6.60 MS 0.08 (1.64) 7.57 (15.97) 7.28 (15.47) CSH 23 0.00 (0.00) 12.34 (20.55) 6.80 (2.60) 2.70 (1.64) 39.71 (39.20) 5.16 (13.14) 7.83 MS 0.00 (1.00) 0.00 (0.00) 15.45 (23.13) CSV 29R 3.36 (10.50) 9.80 (18.24) 5.60 (2.36) 2.40 (1.55) 42.86 (40.98) 4.76 (12.60) 9.19 MS 0.07 (1.63) 6.71 (14.94) 6.53 (14.64) CSH 30 0.00 (0.00) 9.46 (17.87) 6.60 (2.57) 2.10 (1.45) 31.82 (34.39) 5.83 (13.98) 8.65 MS 0.04 (1.22) 4.13 (11.31) 12.50 (20.21) CSV 14R 3.56 (10.61) 10.42 (18.79) 4.60 (2.14) 2.00 (1.41) 43.48 (41.21) 4.50 (12.25) 8.60 MS 0.09 (1.74) 8.85(17.28) 19.17 (25.97) CSH 13 1.04 (4.15) 14.19 (21.68) 5.30 (2.30) 1.90 (1.37) 35.85 (36.72) 4.42 (12.14) 9.14 MS 0.05 (1.51) 4.97 (12.84) 9.58 (18.02) CSH 25 1.00 (4.07) 11.53 (19.86) 5.80 (2.41) 2.50 (1.58) 43.10 (41.01) 5.55 (13.63) 9.48 MS 0.03 (1.22) 3.13 (10.02) 7.93 (16.34) Mahalaxmi 296 (C) 1.02 (4.11) 18.17 (25.24) 5.30 (2.26) 2.30 (1.51) 43.40 (43.53) 5.01 (12.94) 9.84 MS 0.09 (1.75) 8.89 (17.34) 10.83 (19.16) CSH 16 (C) 3.23 (7.36) 8.04 (16.48) 5.40 (2.31) 2.20 (1.48) 40.74 (40.10) 6.70 (15.01) 4.65 LS 0.04 (1.67) 3.54 (22.09) 11.25 (19.57) N-13 4.37 (1.51) 10.60 (2.45) 2.60 (2.51) 5.46.72, 1.57.69.72, 1.57.69.72, 1.51.69.72, 1.51.69.72.69 (1.57.726
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RESEARCH COMMUNICATIONS

et 1.43* 5.68* 0.28* 0.29* 2.38* 2.38* 2.38* 2.38* 2.30* 2.30* 2.38* 2.30* 2.30* 2.30* 2.30* 2.30* 2.50* 4.29* CV% 11.80 14.66 8.65 8.77 9.16 17.0 12.8 19.7 12.5 10.6 **†**Numbers in parenthesis are square root-transformed values; **#**Numbers in parenthesis are arcsine-transformed values. *Significant. MS, Moderately susceptible, HS, Highly susceptible and LS, Least

susceptible.

RESEARCH COMMUNICATIONS

Table 5. Correlation between stem borer infestation and larval population in sorghum

Parameter	% Damaged leaves	% Tiller damage	% Dead hearts	% Stem tunnelling % White ears % Chaffy grain		
No. of larvae per plant	0.1316	-0.0181	-0.2690	0.2212	0.0745	0.1292

*Significant at 5%; *r* table value = 0.361 ; Number of observations = 30 .

respectively. The highest per cent tiller damage was recorded in genotype CSV 15 (45.10) followed by CSV 14R (43.48), popular check Mahalaxmi 296 (43.40) and CSH 25 (43.10), while the lowest in the genotypes NTJ-1 (C) (30.77), CSH 30 (31.82), NTJ-4 (C) (34.92) and CSV 23 (35.09) compared to the resistant check CSH 16 (40.74) (Table 4). Mane *et al.*20 reported that 3.16 tillers/plant was formed, out of which 2.48 tillers (78.5%) were damaged due to stem borer. Tiller production due to deadhearts, serves as a component of tolerance.

 After causing damage to the growing point of the plant, *C. partellus* continues to feed inside the stem throughout the crop growth and makes tunnels inside the stem. The mean stem tunnel length ranged from 6.60 to 10.46 cm with 3.38–7.80% tunnelling (Table 3). Marulasiddesha *et al.*19 recorded 32.57% stem tunnelling in genotype SSV 84 and 49.15% in CSH 14. Based on mean stem tunnel length the genotypes were categorized as least susceptible (0–5 cm), moderately susceptible (5–10 cm) and highly susceptible (>10 cm). The genotype resistant check CSH 16 (C) was found least susceptible with 4.65 cm, whereas, NTJ-2 (C), NLCW-6 and N-14 were found to be highly susceptible as they recorded mean stem tunnel length of 10.45, 10.46 and 11.44 cm respectively. The remaining genotypes were found to be moderately susceptible with 6.60–9.84 cm mean stem tunnel length.

 With regard to white ears, there was significant variation among the genotypes; it ranged from 0% to 13.39% (Table 4). The genotypes NLCW-12 (13.39) followed by CSV 24SS (10.90) and NTJ-1 (C) (10.43) recorded the highest per cent white ears, whereas genotypes CSH 25 (3.13), CSV 216R (3.27) and CSH 30 (4.13) showed lesser value compared to the popular check Mahalaxmi 296 (8.89) and resistant check CSH 16 (3.54). CSH 20MF and CSH 23 did not record any white ears.

 The chaffy grains recorded from the 30 genotypes ranged from 5.00% to 40.00%. The genotype BRJ-358 significantly differed from the others by recording the highest chaffy grain 40.00% followed by NLCW-8 (22.86%), CSV 14R (19.17%) and NLCW-12 (18.00%), whereas the lowest was in the genotype NTJ-1 (C) (5.00%) followed by SSV 84 (5.42%), NTJ-2 (C) (7.27%) and CSV 26 (7.28%) when compared to the popular check Mahalaxmi 296 (10.83%) and CSH 16 (11.25%).

 There was non-significant positive correlation between the number of larvae per plant and per cent damaged leaves, stem tunnelling, white ears, chaffy grain and dead hearts respectively and negative correlation with per cent tiller damage (Table 5).

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698 CURRENT SCIENCE, VOL. 117, NO. 4, 25 AUGUST 2019

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Burrow morphology of the ocypodid crab *Ocypode ceratophthalma* **at Chandipur Coast, Eastern India and its implications**

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Several burrow morphological features of crab *Ocypode ceratophthalma* **including burrow diameter, orientation, inclination, branching and volume were studied from Chandipur, a sedimentologically and biologically diverse beach on the eastern Indian coast. Burrow morphologies (e.g.** *I***,** *J***,** *Y***) were independent of their positions with respect to the coast line. In addition, no correlation between burrow morphology and burrow diameter was observed; however, diameter of burrow appeared to be a good proxy of the total amount of sediment excavated. Burrow diameters are significantly smaller in the foreshore compared to that of the backshore, suggesting that larger individuals reside along the backshore, where they excavate deeper and large-diameter burrows to minimize chances of desiccation. Smaller burrows are more or less vertical, whereas larger burrows are inclined towards the land, probably to stabilize their domicile from tidal** **activities, as well as to minimize energy required to excavate sediments. All these suggest that different types of abiotic factors determine the** *Ocypode* **burrow morphology and their habitat segregation.**

Keywords: Chandipur, crab burrow morphology, crab energetic, Ocypodidae.

THE semi-terrestrial, burrowing crabs of the family Ocypodidae are most common along almost all the tropical and subtropical coastlines of the world^{1,2}. Among them, ghost crabs of the genus *Ocypode*³ are common members of sandy beaches all over the world. They live across a wide range of coastal zones and make characteristic burrows^{1,2,4,5}. Burrows made by these crabs mainly act as shelters from natural environmental stress, a refuge from both aerial and terrestrial predators, as well as a place for reproduction and moulting6,7. The most characteristic feature of burrows produced by *Ocypode* spp. is that their burrows are morphologically diverse and vary from simply straight, unbranched, I-shaped burrows grading to more complex, gently curved and branched U, V, and Y-shaped ones to multi-branched, spiral burrows^{2,8–12}. Moreover, burrow morphology of these ghost crabs does not vary with $sex^{8,13}$. Secondly, the spatial distribution of these different types of burrows along the sandy, coastal beaches is non-random depending on burrow diameter, vertical depth, length and orientation. Their complexity increases from the foreshore to the backshore region^{5,8,9,12,13}.

 These large arrays of burrow morphologies produced by members of the family Ocypodidae are speciesspecific^{2,7,8,11}, although they may also depend upon several abiotic/environmental factors, such as geography and geomorphology (slope, vegetation cover, salinity, groundwater table and tidal cycle) and substrate properties (compactness and composition)^{5,13–17}. Other biological factors, like size and ontogeny, can also affect the overall morphology of these burrows^{5,8,14,15}. Unfortunately, to-date, there is no consensus regarding the major causes influencing the three-dimensional morphology of *Ocypode* crab burrows. Although there are plenty of studies on burrow morphologies, no studies have yet studied the energetics of the burrow excavation as a potential factor influencing the burrow morphology, at least for the ghost crabs. In the present study, we collected data of *Ocypode* crab burrow morphologies from Chandipur, the eastern part of India to identify the nature of morphological complexity (diameters, inclinations, orientation, length and depth, and three-dimensional morphology) and interpret how the burrow morphology is affected by several abiotic factors at Chandipur coast.

 The present study was conducted at Chandipur (21°27′45.17″N, 87°3′21.57″E) near the confluence of the river Burahbalang with the Bay of Bengal (Figure 1). The coastal area of Chandipur is characterized by a wide

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