

**Genetic variability on leaf morpho-anatomical traits in relation to sterility mosaic disease (SMD) resistance in pigeonpea**

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(Supplementary Table 1)-

<b>Percent Disease Incidence (%)</b>	<b>No of genotypes</b>	<b>Genotypes falling in different SMD response groups</b>
0-10% (Resistant)	13	*BRG 3 ( <i>C. cajan</i> ), *ICP 7035 ( <i>C. cajan</i> ), *ICP 15770 ( <i>C. volublis</i> ), *ICP 817 ( <i>R. bracteata</i> ), *ICP 15815 ( <i>R. bracteata</i> ), ICP 15853 ( <i>R. rothii</i> ), ICP 15890 ( <i>R. rothii</i> ), Badnapur-1 ( <i>R. sublobata</i> ), Badnapur-2 ( <i>R. minima</i> ), *Badnapur-3 ( <i>R. rothii</i> ), *BNG 1 ( <i>C. scarabaeoides</i> ), *BNG 3 ( <i>C. scarabaeoides</i> ) and *Badnapur-4 ( <i>C. albicans</i> )
10.1-30% (Moderately resistant)	7	BRG 1 ( <i>C. cajan</i> ), HY 3C ( <i>C. cajan</i> ), ICP 15698 ( <i>C. scarabaeoides</i> ), ICP 15718 ( <i>C. scarabaeoides</i> ), ICP 15922 ( <i>C. scarabaeoides</i> ), ICP 15817 ( <i>R. bracteata</i> ) and ICP 15724 ( <i>C. scarabaeoides</i> ).

30.1-100% (Susceptible)	50	<p><i>C. scarabaeoides</i> - ICP 15683, ICP 15685, ICP 15687, ICP 15688, ICP 15689, ICP 15692, *ICP 15701, *ICP 15703, ICP 15710, ICP 15711, ICP 15719, ICP 15722, ICP 15725, ICP 15727, ICP 15731, ICP 15739, ICP 15748, ICP 15882 and BNG-2.</p> <p><i>C. platycarpus</i> - *ICP 15661, ICP 15662, ICP 15663, ICP 15664, ICP 15666, ICP 15667, ICP 15921, *ICPW 61, ICPW 67 and ICPW 71.</p> <p>Others species - ICP 15621 (<i>C. albicans</i>), ICP 15642 (<i>C. lineatus</i>), ICP 15761 (<i>C. sericeus</i>), ICP 15900 (<i>C. viscosa</i>), ICP 15855 (<i>R. rothii</i>), ICP 15857 (<i>R. rothii</i>), *ICP 15681 (<i>R. rufescens</i>) and ICP 15799 (<i>F. macrophylla</i>).</p> <p>Cultivars (<i>C. cajan</i>) - *TTB 7, IPA 8F, GT 101, JKM 189, BSMR 736, *ICPL 87119, *WRP 1, *TS 3R, *ICP 8863, *GRG 333, *ICP 2376, *BRG 2 and *GRG 811.</p>
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Note\*- *C- Cajanus, R- Rynchosia, F-Flemingia*

\* Indicates twenty three extreme pigeonpea genotypes selected for correlation study