

Name of the project	Exploitation of heterosis in Okra [<i>Abelmoscnus esculentus</i> (L.) Moench] using conventional and Biotechnological tools
Funding agency	Council of Science and Technology, Lucknow
Year of start	2015-16
Year of completion	2018-19
Name of PI and Co-PI	Dr. Bijendra Singh Dr. Vaishali
Total budget	09.00 lakhs

Significant achievements:

- Fifty okra genotypes were collected and screened on the basis of morphological, molecular, abiotic (salt) stress and biotic (YVMV) stress for further study.
- Three genotypes were found highly resistance to YVMV infection.
- Disease incidence ranged from 10.58 to 53.20 percent. Cross (hybrid) i.e. C-7801 x AB – 3 exhibited minimum percentage while cross (hybrid) KS – 310 x AB 1 exhibited maximum percentage of disease infestation.
- Out of 45 cross hybrids, 9 were highly resistance against YVMV, 6 were resistance, 21 moderately resistance, 8 moderately susceptible and 1 was highly susceptible.
- Three treatments were tested for salinity stress @ 25, 50 and 100 mmol respectively. After 20 days, it was noted that morphological characteristics affected with increase in salinity levels.
- The diversity or genetic similarities between varieties varied for all the 50 genotypes of okra germplasm.

Recommendations:

- The cross C-7801xAB-3 may be used for further study to develop the disease resistant genotype.

Publications:

- B. Singh, Aakansha Goswami and Vaishali (2017). Study of Genetic Diversity in Okra [*Abelmoschus esculentus* (L.) Moench]. *Vegetos*; 30 (Special) : 109-116.
- Kumar, A., Kumar, M., Sharma, V.R., Singh, M.K., Singh B. and Chand, P. (2019). Character association and path coefficient analysis of yield and yield related traits in Okra [*Abelmoschus esculentus* (L.) Moench]. *Progressive Agriculture*, 19 (1): 140-145.
- Kumar, A., Kumar, M., Sharma, V.R., Singh, M.K., Singh ,B. and Chand, P. (2019). Genetic variability, heritability and genetic advance studies in genotypes of Okra [*Abelmoschus esculentus* (L.) Moench]. *Journal of Pharmacognosy and Phytochemistry*, 8(1):1285-1290