

RESUME

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Research Interests: Genomics and Molecular Breeding in Vegetable Crops

SUMMARY

Working mainly on vegetable crops viz., peas, cucumber, common bean, chilli, garlic and ginger. We (Katochet *et al.*, 2010), for the first time reported the genetic mapping of *er2* (gene for powdery mildew resistance) to linkage group III of pea, thus clarifying the prevailing confusion in the literature regarding the existence and genomic location of *er2*. Other research interests include Linkage Mapping, GWAS, NGS data analysis and bioinformatics.

EDUCATION

| S. No. | Degree | Year | Subject | University |
|--------|--------------------|------|------------------------|----------------|
| 1 | B.Sc. (Agri.) | 2001 | Agriculture and Allied | GNDU, Amritsar |
| 2 | M. Sc. (Veg. Sci.) | 2003 | Vegetable Breeding | HPKV, Palampur |
| 3 | Ph.D. (Veg. Sci.) | 2008 | Vegetable Breeding | HPKV, Palampur |
| 4 | NET | 2005 | Vegetable Science | ASRB |

PROJECTS/GRANTS

| S. No. | Project | Funding Agency (Budget in lakhs) | Status |
|--------|--|----------------------------------|-----------|
| 1 | Collection and DNA profiling of nutritionally important underutilized vegetables of North Western Himalayas (Co-PI) | DST SERB (Rs. 38.61) | Ongoing |
| 2 | SSR based germplasm characterization for resistance to powdery mildew in cucumber (<i>Cucumissativus</i> L.) (PI) | DBT, New Delhi (Rs. 32.52) | Completed |
| 3 | Molecular marker assisted introgression of powdery mildew resistance genes into the elite cultivar of pea | DST (SERB) | Completed |

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|----------|--|---------------------------------|-----------|
| | (<i>Pisum sativum</i> L.) (PI) | (Rs. 19.00) | |
| 4 | Farmers' participatory collection, characterization and conservation of endangered genetic diversity of ginger (<i>Zingiber officinale</i> Rosc.) (PI) | DST, SARTHI (Rs. 16.69) | Completed |
| 5 | Transcriptome analysis, gene annotation and mining of a golden herb-garlic (<i>Allium sativum</i> L.) expressing resistance to destructive purple blotch (Mentor) | NPDF- SERB (DST) (Rs. 19.20) | Completed |
| 6 | Development of SNPs for <i>Brassica juncea</i> (Co-PI) | DBT, N. Delhi (Rs. 52.43) | Completed |
| 7 | Synthesis of new gene pool following introgression of disease resistance and drought tolerance genes from secondary (<i>Phaseolus coccineus</i> L) and tertiary (<i>Phaseolus acutifolius</i> L) gene pools into cultivated <i>Phaseolus vulgaris</i> L (Co-PI) | DST-SERB (Rs. 33.10) | Completed |

Best Ten Publications

1. Katoch, V., **Sharma, S.**, Pathania, S., Banyal, D.K., Sharma, S.K. and Rathour, R. 2010. Molecular mapping of pea powdery mildew resistance geneer2 to pea linkage group III. *Molecular Breeding* 25(2): 229-237. doi:10.1007/s11032-009-9322-7.
2. Dar, A.A., **Sharma, S.**, Mahajan. R., Mushtaq, M., Salathia, A., Ahamad, S., and Sharma, J.P. 2020. Overview of purple blotch disease and understanding its management through chemical, biological and genetic approaches. *Journal of Integrative Agriculture* 19(0): 2–13 doi: 10.1016/S2095-3119(20)63285-3
3. Dar, A.A., Mahajan, R., Lay, P and **Sharma, S.** 2017. Genetic diversity and population structure of *Cucumissativus* L. by using SSR markers. *3 Biotech* 7(5): 307 10.1007/s13205-017-0944-x.
4. **Sharma S**, Dar AA, Gupta S, Singh R. 2021. Evaluation of resistant genotypes and their characterization using molecular markers linked for powdery mildew resistance in cucumber (*Cucumissativus* L.). *Plant Genetic Resources: Characterization and Utilization* 1–6. <https://doi.org/10.1017/S1479262121000605>
5. Sudan, J., **Sharma, S.**, Salgotra, R.K., Pandey, R.K., Neelam, D. and Singh R. 2023. Elucidating the process of SNPs identification in non-reference genome crops, *Journal of Biomolecular Structure and Dynamics*, DOI: 10.1080/07391102.2023.2194002
6. Shah, R.A., Bakshi, P., Jasrotia, A., Wali, V.K., **Sharma, S.**, Gupta, M., Gupta, R.K., and Jamwal, M. 2023. Comparative morpho-molecular characterization of elite walnut variety Parbat (JWSP-06) with local selections of north-western Himalayan region of Jammu and Kashmir, India. *Scientia Horticulturae*. Vol. 319 <https://doi.org/10.1016/j.scienta.2023.112176>
7. Sudan J, Singh R ,**Sharma S**, Salgotra RK, Sharma V, Singh G, Sharma I, Sharma S, Gupta SK and Zargar SM. 2019. ddRAD sequencing-based identification of inter-genepool SNPs and association analysis in *Brassica juncea*. *BMC Plant Biology*

19:594.

8. Sharma, A., Sekhon, B., **Sharma, S.**, and Kumar, R. 2020. Newly isolated intervarietal garden pea (*Pisum sativum* L.) progenies (F7) under north western Himalayan conditions of India. *Experimental Agriculture*, 56(1), 76-87. doi:10.1017/S0014479719000115
9. Sharma, R., Dar, A.A., Mahajan, R. and **Sharma, S.** 2019. Molecular and Biochemical Characterisation of Indian Germplasm of *Pisum sativum* L. *Proc. Natl. Acad. Sci., India, Sect. B Biol. Sci.* (<https://doi.org/10.1007/s40011-018-01069-3>)
10. Sharma, P. **Sharma, S.**, Singh, J., Saha, S. and Baranwal, V.K. 2016. Incidence of Lettuce mosaic virus in lettuce and its detection by polyclonal antibodies produced against recombinant coat protein expressed in *Escherichia coli*. *Journal of Virological Methods* 230: 53–58. <http://dx.doi.org/10.1016/j.jviromet.2016.01.014>