Mohammad Shamshad

Post-Doctorate Research Associate, Department of Plant Breeding & Genetics, Punjab Agricultural University, Ludhiana, Punjab India 542/3 Ussman Khan Wala, Jamalpura, Malerkotla, Punjab, India Pin-148023 Email: shamshad.rattan@gmail.com website: https://shamshadm.github.io

Summary

Plant breeder and Geneticist with extensive experience in R-programming, SAS, bioinformatics, data analysis, trait phenotyping, genotyping, breeding methodology, quantitative techniques, germplasm development and genetic mapping.

Education

Ph.D. in Plant Breeding and Genetics; Sep. 2019

Punjab Agricultural University, Ludhiana, India.

- O Dissertation: "Genetic and physiological basis of nitrogen uptake and utilization in wheat (Triticum aestivum L.)"
- o Advisor: Prof. Dr. Achla Sharma
- Collaboration with University of Bristol, UK and University of Nottingham, United Kingdom

M.Sc. in Plant Breeding & Genetics; Nov. 2014

Punjab Agricultural University, Ludhiana, India

- O Dissertation: "Assessment of morphological and genetic diversity and its association with heterosis and F_1 performance in sunflower (Helianthus annuus L.)"
- o Advisor: Prof. Dr. Satwinder Kaur Dhillon

B.Sc. in Agriculture (Plant Science), June 2012

Punjab Agricultural University, Ludhiana, India

Research Experience:

Post Doctorate Research Associate (TIGR2ESS)

August 2020 to March 2022

Plant Breeding & Genetics, Punjab Agricultural University

• Responsibilities:

- Pre-breeding and germplasm characterization using novel genomics, phenomics and statistical tools for irrigated, rainfed and hybrid wheat breeding programs.
- o Development of core panels, identify potential trait gaps in terms of phenotypic variation and build novel strategies to enrich for missing phenotypic variations.
- o Conducted extensive field trials, crossing, phenoyping and data collections across multiple environments over the years
- o Double haploid breeding in wheat using wheat × Maize crosses

Research Fellow

August 2019 to July 2020

Plant Breeding & Genetics, Punjab Agricultural University

- Responsibilities:
 - o Quantitative genetic analysis of field trials.
 - Perform genome-wide analysis and genomic predictions for the nitrogen use efficiency
 - Making of crosses in wheat for various objectives viz. grain yield, heat, drought, salt and stripe rust resistance

Scientific Associate

June to August 2018

School of Biosciences, University of Nottingham, United Kingdom

- Hydroponics and shovelomics for the root system architecture
- Image's analysis using RootNav and ImageJ ObjectJ

Student Exchange

September to October 2018

School of Biosciences, University of Bristol, UK

- 35K array of genotyping
- SNPs Calling
- QTLS mapping for NUE

Ph. D (INEW International Fellow and Graduate Research Assistantship) January 2016 to July 2019

- Focused on the development and evaluation of RIL mapping population related to nitrogen uptake and utilization.
- Development of high-density linkage map based on Breeder array of 35K SNPs.
- Genome-wide QTL mapping for various agro-physiological, nitrogen uptake and utilization efficiency traits phenotyped across a wide range of environments.
- High-throughput phenotyping in wheat and soybean using newly developed phenotyping platforms.
- Genome selection/predictions in Punjab Agricultural University wheat breeding program.
- Gained proficiency in statistical analysis and programming including Unix operating system, perl language, GBS data analysis, and statistical analysis using R packages and SAS software.
- Conducted extensive field trials, crossing, phenoyping and data collections across multiple environments for the three years.
- Analyzed multi-environment phenotypic data in ASReml package in R and incorporating spatial corrections using mixed linear models and generate variance components.
- BLUP/BLUE estimates for genomic selections and QTL mapping.
- Molecular mapping of nitrogen uptake and utilization efficiency in wheat.

Research Assistant

Jan 2015 to Dec 2015

Department of Plant Breeding and Genetics, PAU, Ludhiana, India

- Responsibilities:
 - Development of the germplasm and core panel for the breeding objective for yield, quality and quantity oils
 - Conducted extensive field trials, crossing of A, B and R lines, and data collections of the Advanced variety trials

Graduate Research Assistant (M.Sc.)

July 2012 to Nov. 2014

Department of Plant Breeding and Genetics, PAU, Ludhiana, India

- Identification of potential restorers for newly developed temperate CMS lines and identify the heterotic combinations.
- Association of morphological and genetic diversity with heterosis and combining ability

Publications:

Peer-Reviewed Articles

- Padhy, A. K., Kaur, P., Singh, B., Kaur, R., Bhatia, S., **Shamshad, M.**, ... Sharma, A. (2022). In silico characterization of Thinopyrum elongatum-derived PsyE1 gene and validation in 7D/7E bread wheat introgression lines open avenues for carotenoid biofortification in wheat. *Cereal Research Communications*, (0123456789). Retrieved from https://doi.org/10.1007/s42976-022-00279-w
- Gudi, S., Saini, D. K., Singh, G., Halladakeri, P., Kumar, P., **Shamshad, M.,** ... Sharma, A. (2022). Unravelling consensus genomic regions associated with quality traits in wheat using meta-analysis of quantitative trait loci. *Planta*, 255(6), 1–19. Retrieved from https://doi.org/10.1007/s00425-022-03904-4
- Gupta, P. K., Balyan, H. S., Chhuneja, P., Jaiswal, J. P., Tamhankar, S., Mishra, V. K., ... Vishwakarma, M. K. (2022). Pyramiding of genes for grain protein content, grain quality, and rust resistance in eleven Indian bread wheat cultivars: a multi-institutional effort. *Molecular Breeding*, 42(4), 1–16. Retrieved from https://doi.org/10.1007/s11032-022-01277-w
- Kaur, Sarabjit, **Shamshad, M.**, Jindal, S., Kaur, A., Singh, S., sharma, A., & Kaur, S. (2022). RNA-Seq-Based Transcriptomics Study to Investigate the Genes Governing Nitrogen Use Efficiency in Indian Wheat Cultivars. *Frontiers in Genetics*, 13(March), 1–15. Retrieved from https://doi.org/10.3389/fgene.2022.853910
- Kaur, Sukhjit, Suhalia, A., Sarlach, R. S., **Shamshad, M.**, Singh, P., Grover, G., ... Sharma, A. (2022). Uncovering the Iranian wheat landraces for salinity stress tolerance at early stages of plant growth. *Cereal Research Communications*, (0123456789). Retrieved from https://doi.org/10.1007/s42976-022-00245-6
- Sharma, A., Ankita, **Shamshad, M.**, Kaur, S., Singh, A., Srivastava, P., ... Sohu, V. S. (2021). SALT STRESS GENOTYPIC RESPONSE: RELATIVE TOLERANCE OF WHEAT CULTIVARS TO SALINITY. *Agricultural Research Journal*, 58(6), 974–982. Retrieved from https://doi.org/10.5958/2395-146X.2021.00138.1
- **Shamshad, M.** & Dhillon, S. K. (2016). Morphological and molecular genetic diversity analyses in Helianthus annuus (L.). *Electronic Journal of Plant Breeding*, 7(4), 1216–1223. Retrieved from https://doi.org/10.5958/0975-928X.2016.00169.1
- Shamshad, M., Dhillon, S. K., & kaur, G. (2016). Heterosis for oil content and oil quality in sunflower

- (Helianthus annuus L.). *Current Advances in Agricultural Sciences(An International Journal)*, 8(1), 44. Retrieved from https://doi.org/10.5958/2394-4471.2016.00010.
- **Shamshad, M.**, Dhillon, S. K., Tyagi, V., Akhatar, J. (2014). Assessment of Genetic Diversity in Sunflower (Helianthus annuus L.) Germplasm. *International Journal of Agriculture and Food Science Technology*, 5(4), 267–272. Retrieved from httpx//www:ripublication:com/ ijafst:htm
- **Shamshad, M.**, Dhillon, S. K. (2014). Magnitude of combining ability for seed yield and oil content in sunflower (Helianthus annuus L.). In *National Symposium on Crop Improvement for Inclusive Sustainable Development* (Vol. 10, pp. 248–250). Retrieved from https://doi.org/10.5958/0975-928X.2017.00014.X

Book Chapters

- **Shamshad, M.** and Sharma, A. (2018). The usage of genomic selection strategy in plant breeding. In: Yelda Ozden C (ed) Next Generation Plant breeding. Pp 93-108. Intech Publisher, London, UK. Retrieved from http://dx.doi.org/10.5772/intechopen.76247
- Sharma, A., **Shamshad, M.,** Kaur, S., Srivastava, P., Mavi, G.S., Sohu, V.S. (2022). Tackling a Cereal Killer on the Run: Unending Fight Between Wheat Breeding and Foliar Rusts. In: GP, et al. New Horizons in Wheat and Barley Research. Springer, Singapore. https://doi.org/10.1007/978-981-16-4449-8 10.

Thesis PhD

https://krishikosh.egranth.ac.in/handle/1/5810137744

Posters and Conferences

- Shamshad M, Dhillon S K, Tyagi V and Singh S (2013) Studies on sunflower (*Helianthus annuus* L.) germplasm evaluation for drought tolerance. Pp 109-10, abstracted in Indraprastha International Conference on Biotechnology, October 22-25, 2013, Guru Gobind Singh Indraprastha University, New Delhi, India.
- Shamshad M, Tyagi V and Dhillon S K (2014) Estimation of drought susceptibility index in sunflower (*Helianthus annuus* L.) germplasm. Pp 97, abstracted in National Seminar on Plant Biotechnology: Challenges and Opportunities in 21st Century, March 03-04, 2014, Jamia Hamdard, New Delhi, India.
- Tyagi V, Dhillon S K, Shamshad M and Singh S (2013) Assessment of response of sunflower (Helianthus annuus L.) genotypes to water stress. Pp 108, abstracted in Indraprastha International Conference on Biotechnology, October 22-25, 2013, Guru Gobind Singh Indraprastha University, New Delhi, India.
- Tyagi V, Singh S, **Shamshad M** and Dhillon S K (2013) Effect of moisture stress on correlations between yield and component traits in sunflower. Pp 108-09, abstracted in Indraprastha International Conference on Biotechnology, October 22-25, 2013, Guru Gobind Singh Indraprastha University, New Delhi, India.
- Tyagi V, Dhillon S K, **Shamshad M** and Akhatar J (2014) Genetic diversity in sunflower (*Helianthus annuus* L.) germplasm under water stress environment. Pp 98, abstracted in National Seminar on Plant Biotechnology: Challenges and Opportunities in 21st Century, March 03-04, 2014, Jamia Hamdard, New Delhi, India.
- Shamshad M, Sharma A, Srivastava P, Kaur S, Brar A, Bains N S (2019) Genetic variability for root traits in relation to nitrogen use efficiency in structured population of wheat (*T. aestivum*)

- Pp 133, abstracted in International Conference Genomics and Breeding for Crop Improvement, December 04-06, 2019 Chaudhary Charan Singh University, Meerut, UP, India.
- Ankita, Sharma A, Srivastava P, Kaur S, Shamshad M, Kaur A, Sarlach R S, Sharma H, Singh S (2019) Morphological characterization of Mexican wheat landraces under simulated salinity stress, Pp 109, abstracted in International Conference Genomics and Breeding for Crop Improvement, December 04-06, 2019 CCS University, Meerut, UP, India.
- Kaur S, Sharma A, Ankita, Srivastava P, **Shamshad M**, Sarlach R S, Sharma H, Singh S (2019) Yield component and their association for enhancing grain yield in Iranian wheat landraces under saline field conditions, Pp 154, abstracted in International Conference Genomics and Breeding for Crop Improvement, December 04-06, 2019 CCS University, Meerut, UP, India.
- Kumar A, Sharma A, Sharma H, Shamshad M, Srivastava P, Mavi G S, Kaur S, Chhuneja P, Rajput R, Pandey A and Sohu V S (2022). Yellow pigment enriched biofortified wheat for tackling micronutrient deficiencies Pp 45. Abstracted in Ist International Symposium on "Cereals for Food Security and Climate Resilience". January 18-20, 2022, Indian Institute of wheat and Barley, Karnal, India
- Sharma A, Shamshad M, Sharma H, Tanin M J, Kumar P, Singh S, Srivastava P, Mavi G S, Bala R, Kaur J, Kaur S, Chhuneja P, Kaur A and Sohu V S (2022). Combining nutritional and quality traits in wheat: A newer mandate for wheat breeders. Pp 46. Abstracted in Ist International Symposium on "Cereals for Food Security and Climate Resilience". January 18-20, 2022, Indian Institute of wheat and Barley, Karnal, India
- Rushali, Srivastava P, Kumar S, Ayoubi H, Ankita, Sarao L, **Shamshad R**, Sharma H (2022) Evaluation and utilization of breeding lines having diverse dwarfing genes. Pp 50. Abstracted in Ist International Symposium on "Cereals for Food Security and Climate Resilience". January 18-20, 2022, Indian Institute of wheat and Barley, Karnal, India
- Gudi S, Sharma A, Shamshad M, Sharma H, Singh S, Srivastava P, Mavi G S and Sohu V S (2022).
 Field-based, low-cost speed breeding technique to develop wheat (*Triticum aestivum*) having enhanced water use efficiency and early heat stress tolerance. Pp 38. Abstracted in Ist International Symposium on "Cereals for Food Security and Climate Resilience". January 18-20, 2022, Indian Institute of wheat and Barley, Karnal, India.
- Sharma H, **Shamshad M**, Gudi S, Tanin M J, Kumar P, Sharma A, Singh S, Srivastava P, Mavi G S and Sohu V S (2022) Enhancing bioavailability of micronutrients by reducing phytic acid content in wheat. Pp 74. Abstracted in Ist International Symposium on "Cereals for Food Security and Climate Resilience". January 18-20, 2022, Indian Institute of wheat and Barley, Karnal, India.
- Shamshad M, Sharma A, Singh J, Srivastava P, Kaur R and Kaur H (2022). Agronomic/physiological characterization of spring wheat genotypes for various nitrogen use efficiency traits. Pp 80. Abstracted in Ist International Symposium on "Cereals for Food Security and Climate Resilience". January 18-20, 2022, Indian Institute of wheat and Barley, Karnal, India
- Kaur S, **Shamshad M**, Sharma A, Kaur A and Kaur S (2022). RNA-seq based transcriptomics analysis of roots and shoots of wheat cultivar in response to low nitrogen stress. Pp 173. Abstracted in Ist International Symposium on "Cereals for Food Security and Climate Resilience". January 18-20, 2022, Indian Institute of wheat and Barley, Karnal, India.

Talks

Hydroponics and shovelomics for the root system architecture

October 26, 2018 Invited Talk at PBG, PAU

 SNP chip based High-Density Linkage Map and its Application to QTL Mapping of nitrogen uptake and utilization in Bread Wheat

October 03, 2018

Invited Talk at Life Science, School of Bioscience, University of Bristol, UK

Breaking yield ceiling in wheat: Genomic and molecular approaches

April 20, 2017

Ph.D. Seminar talk at PBG, PAU

Evolution of free threshing habit in wheat

November 04, 2016

Ph.D. Seminar Talk at PBG, PAU

Does genomic selection strategy help to improve the crop plants?

May 16, 2016 Invited Talk at PBG, PAU

Correlation and its application in Crops

October 03, 2013 M.S Lecture Talk PBG, PAU

Overcoming the barriers to alien gene transfer in Crop Improvement

September 27, 2013 M.S seminar talk at PBG, PAU

Professional Trainings

- The series of next generation genomics and integrated breeding for crop improvement (VII-NGGIBCI) on genomics for food, health and nutrition, ICRISAT, Hyderabad. May 14, 2020 (Virtual)
- General assembly/ Training to PDRA and early carrier researchers, International Crop Research Institute for Semi-Arid Tropics (ICRISAT), Hyderabad, India. January 19-25, 2020,
- Scientific exchange to learn shovelomics and hydroponics for the roots system architecture School of Biosciences University of Nottingham, Sutton Bonington, United Kingdom (UK).
- Workshop on Precision nitrogen nutrition in wheat: integrating genetics and precision agronomy for improving nitrogen use efficiency, BISA, Ludhiana, India. March 1 -5 March 2018.

- Workshop on CerealsDB, Life Science, School of Bioscience, University of Bristol, United Kingdom. 18th -19 September 2018
- Scientific Student exchange University of Bristol, UK, Usages of advance generation sequence platforms and molecular work.
- workshop on Wheat genomic data analysis for improvement in nitrogen use efficiency ICAR- National Bureau of plant genic resources (NBPGR), New Delhi, India. February 27 to March 2, 2019
- Short term bioinformatics training (RNA-seq and DNA-seq). ArrayGen Technologies, India, Pune. 23 Oct-21 Nov. 2018.
- Experience in raising *Cajanus cajan* (Arhar) and *Triticum aestivum* (Wheat) crops at field scale during Practical Crop Production Training with first-hand information on crop production and protection technology.
- Work experience to make a cross in Wheat, Rice, Chickpea, Pigeon pea, Field pea, Maize and Sunflower.

Skills

Quantitative Skills

- Genotyping-by-sequencing data analysis and SNP calling
- Trained to perfom RNA-seq data anlysis
- Quantitative Genetics
 - o QTL Mapping: R/qtl, R/ASMap, IciMapping etc.
 - o Genome-wide Association Mapping: Plink, TASSEL, BGLR, rrBLUP, GAPIT, FarmCPU etc.
 - o Genomic Selections: BGLR, rrBLUP etc.
- Statistics
 - o R
 - o SAS
 - ASReml
 - o Python

Wet Lab Skills

• DNA extractions, PCR, real-time PCR, electrophoresis, SSR marker and KASP marker genotyping

Computational Skills

- Operating Systems: LINUX, OS X, Windows
- Latex and R Markdown

Field Experiment Skills

- Field designs
- Crossing

- Agronomic management
- DH production

References and additional information available upon request