

SAQIB ALI

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Chiniot, Punjab, Pakistan

About Me

Plant pathologist and bioinformatician (M.Sc. Hons.) specializing in comparative genomics, phylogenomics, and molecular evolution. Combines advanced wet-lab skills (molecular diagnostics, SSR genotyping, marker-assisted selection, stress screening) with computational expertise in genome-wide analysis, synteny mapping, protein structure prediction, molecular dynamics simulations, and population genetics. Proficient in integrated R/Python workflows for plant-pathogen interaction studies and stress-responsive gene discovery.

Education

University of Agriculture Faisalabad M.Sc. (Hons.) Agriculture Plant Pathology CGPA: 3.55/4.00	Sep 2022 – Sep 2024 <i>Faisalabad, Pakistan</i>
University of Agriculture Faisalabad B.Sc. (Hons.) Agriculture Plant Pathology CGPA: 3.64/4.00	Aug 2018 – Jul 2022 <i>Faisalabad, Pakistan</i>

Research Experience

Postgraduate Thesis Research Sept 2022 – Jul 2024

Department of Plant Pathology, University of Agriculture Faisalabad Faisalabad, Pakistan.

Thesis: "Incidence of *Meloidogyne graminicola* in Maize Crops and Its Integrated Management"

Supervisor: [Prof. Dr. Sajid Aleem Khan](#)

- Conducted **field surveys** and greenhouse **experiments** to assess root-knot nematode incidence in **maize**, implementing **integrated management** strategies through biological, chemical, and cultural practices in replicated field trials
- Developed **proficiency in nematology** and **plant pathology techniques** including nematode extraction and identification, pathogen isolation and culturing, media preparation, and Koch's postulates validation
- Collaborated on field experiments for **cotton** Verticillium wilt, **vegetable crop** nematode screening (**carrot, grapevine**), and **cereal pathology trials** (rice, wheat), contributing to experimental design, data collection, and statistical analysis.
- Applied **novel agricultural technologies** including **laser-assisted nanoparticle** treatments on **rice** and **wheat** for disease management and crop improvement research.
- Mentored undergraduate students** through laboratory demonstrations and training in plant pathology techniques and experimental protocols.

Research Fellow

Nuclear Institute for Agriculture and Biology - (PAEC-IAEC)

Sep 2025 – Present

Faisalabad, Pakistan

Supervisor: Dr. Muhammad Rashid

Project 1: In-Silico Analysis of WRKY Genes in Rice Subspecies (Sep–Oct 2025)

- Identified and characterized **252 WRKY transcription factors** across *O. sativa indica*, *japonica*, and *O. glaberrima* using PlantTFDB and SMART databases
- Constructed maximum likelihood **phylogenetic trees** (ClustalX, MEGA12) to classify genes into evolutionary subgroups based on conserved domain architecture
- Analyzed **spatio-temporal expression patterns** across **42 tissue types** using RiceXPro and constructed **co-expression networks** with RiceFRIEND to identify functional modules
- Generated publication-quality visualizations using **R** (gplots, heatmap.2, ComplexHeatmap) and **TBtools** for multi-dimensional expression profiling

Project 2: Comparative Genomics Across Seven Cereal Genomes (Nov 2025–Jan 2026)

- Conducted genome-wide identification of **547 WRKY genes** using HMMER and BLAST with **redundancy filtering** (CD-HIT); performed **multiple sequence alignment** (MAFFT) and phylogenetic reconstruction (iTOL)
- Characterized **gene structure**, **conserved motifs** (MEME Suite), and protein domains (InterProScan) to establish **evolutionary relationships** across cereals
- Executed comparative **synteny analysis** using MCScanX and Circa, mapping **700+ orthologous gene pairs** and visualizing chromosomal relationships with TBtools and circlize
- Calculated **evolutionary selection pressure** (KaKs_Calculator 2.0), demonstrating strong **purifying selection** ($Ka/Ks = 0.08–0.12$) across WRKY gene families
- Applied **multivariate statistical methods** (PCA, t-SNE) and codon usage bias analysis (seqinr) to compositional datasets; developed **reproducible analytical workflows** using Python (BioPython) and R for statistical testing and high-dimensional visualization

Wet Laboratory & Physiological Screening (Sep–Oct 2025)

- Conducted comprehensive **stress resistance screening in rice germplasm** using **physiological** (SPAD, IRGA, porometer, osmometer, pressure chamber) and **biochemical assays** (MDA, POD, EC, ion quantification via spectrophotometry and flame photometry)
- Applied molecular techniques for **genetic characterization**: DNA quantification, PCR, gel electrophoresis, **SSR marker genotyping**, marker-assisted selection (MAS), and primer design (**Primer3**)
- Performed **plant pathology assays** including inoculum preparation, pathogen staining, varietal screening, and plant-pathogen interaction studies.

Research Intern

PepsiCo Roshan Kal for Pakistan

Jul 2024 – Aug 2024

Faisalabad, Pakistan

- Contributed to **Bonsucro-certified model farm development** by validating **sustainable agriculture practices** and conducting **supply chain sustainability** assessments in collaboration with industry and academic partners.

Research Intern

Vegetable Research Institute, Ayub Agricultural Research Institute Faisalabad, Pakistan

Feb 2022 – May 2022

- Designed and conducted **field trials** in **walk-in tunnels** to study **disease prevalence** and **fungicide efficacy** in **vegetable crops** (tomatoes, cucumbers, watermelons, onions, garlic), with emphasis on tomato early blight and grey mold management.
- Performed **plant pathology diagnostics** including pathogen isolation, **ELISA-based virus identification**, and **PCR-based detection** for bacterial and nematode pathogens.
- Contributed **disease resistance data** to **vegetable** and **cereal breeding programs** through **statistical analysis** (RStudio, Python), supporting crop improvement strategies.

Professional Experience

Agriculture Officer Extension (Intern)

Department of Agriculture Extension, Government of Punjab

Sep 2024 – Sep 2025

Chiniot, Pakistan

- Provided **technical advisory** support to **1,500+ farmers** on **integrated pest and disease management**, translating **research-based recommendations** into practical, cost-effective control strategies to improve crop yields.
- Conducted **field-based pest scouting** and **crop health assessments** through regular **farmer visits**, diagnosing disease/pest issues and providing targeted intervention recommendations.
- Assisted in **seed and agrochemical quality assessments** to ensure product efficacy and **regulatory compliance**, supporting farmer access to certified inputs.

Research & Knowledge Management Intern

World Agriculture Forum

May 2025 – July 2025

Remote

- Collaborated with **policy research** team to **analyze global agricultural trends** and compile technical reports **promoting sustainable agricultural practices** and stakeholder dialogue.

Publications

Publication Summary: 8 peer-reviewed articles (6 first author), 4 conference presentations

Research Articles (Submitted/Under Review):

1. Ali, S. (2026). Evolutionary Origins and Structural Mechanism of V2-Mediated Host Immunity Suppression in Soybean Stay-Green Associated Virus (SoSGV). **Submitted to Archives of Virology** (Springer Nature). 
2. Ali, S. (2026). Systematic Characterization of Glycine max IQD Genes Reveals Insights into Drought Stress Adaptation and Hormonal Regulation of Plant Architecture. **Submitted to 3 Biotech** (Springer Nature). 
3. Ali, S., Hanjra, Z., Saeed, M., Fatima, T., Khalil, S., Iltaf, B., & Rashid, M. (2025). In-silico analysis of WRKY transcription factors (WRKY) gene family in rice. **Submitted to BMC Plant Biology**. 
4. Hanjra, Z., Ali, S., Fatima, T., Saeed, M., Khalil, S., & Rashid, M. (2026). High-resolution comparative genomics and compositional analysis of WRKY transcription factors across seven cereal and model plant genomes. **Submitted to BMC Plant Biology**. 

Research Articles (Published/Accepted):

5. Ali, S., Khan, S. A., Abbas, H., Khadim, U., Aslam, A. & Anwaar, M. (2025). Incidence of *Meloidogyne Graminicola* on Maize and Its Integrated Management. [Accepted](#) in [Pakistan Journal of Zoology](#).
6. Tufail, M. M., Abbas, H., Khadim, U., Bashir, M. R., Kamran, M., Khan, S. A., Zafar, M., Shahbaz, M. U., ul Haq, M. E., & Ali, S. (2024). Management of *Macrophomina phaseolina* infecting sesame germplasm. [Journal of Phytobiology](#), 4(1), 9–17.

Review Articles:

7. Ali, S., Mithal, U., Bibi, J., Hanif, A., Zohaib, M., Asghar, J., ... & Sohail, A. (2025). Revolutionizing Cotton Leaf Curl Disease Detection: CRISPR-Based Diagnostics for Rapid, Field-Deployable Solutions. [Planta Animalia](#), 4(4), 347-362.
8. Ali, S., Anwaar, M., Khadim, U., & Bakhsh, B. P. (2025). CRISPR-based biosensors for plant disease diagnostics: Innovations, applications, and future prospects. [Annual Methodological Archive Research Review](#), 3(8).

Conference Abstracts:

9. Ali, S. (2025). Multi-Omics and Machine Learning for Predicting Biocontrol Efficacy under Climate-Driven Pathogen Shifts. [\[Abstract\]](#) 9th ICCPS PMAS UAAR, Pakistan
10. Aslam, A., Khan, S. A., Ali, S., Abbas, H., Anwar, M., & Ul Haq, U. (2023). Plant growth-promoting rhizobacteria as a biocontrol tool for root-knot nematode killing in lab [\[Abstract\]](#). 8th ICCPS IUB, Pakistan.
11. Ali, S., Khan, S. A., Abbas, H., Aslam, A., Abbas, W., & Khadim, U. (2023). Evaluation of resistant sources of tomato against early blight under natural field conditions and its management through newly introduced chemicals [\[Abstract\]](#)—8th ICCPS IUB, Pakistan.
12. Ali, S. (2025). Harnessing Host-Induced and Exogenous RNAi for Precision Nematode Control. [\[Abstract\]](#) VIRTUAL NEMATOLOGY CONFERENCE 2025

Professional Service:

- Peer Reviewer, *Journal of Plant Diseases and Protection* | Springer Nature (2025).
- Peer Reviewer, [Journal of Advances in Biology & Biotechnology](#) (2025).

Honors & Awards

Ehsaas Undergraduate Scholarship

HEC Pakistan

PEEF Scholarship

[Punjab Educational Endowment Fund](#)

Sep 2019 – Jul 2022

Faisalabad, Pakistan

Sep 2016 – Jun 2018

Faisalabad, Pakistan

Training & Certifications

BONSUCRO Sugarcane Sustainability Training Workshop

Aug 2024

PepsiCo Pakistan

Lahore, Pakistan

International Conference Climate-Smart Agriculture for Food Security

July 2024

University of Poonch Rawalakot

Azad Jamu Kashmir

University-Level Certifications (2025)

- Python for Everybody Specialization | University of Michigan (via [Coursera](#))
- Bioinformatic Specialization | University of California, San Diego (via [Coursera](#))
- Data Visualization & Dashboarding with R | Johns Hopkins University (via [Coursera](#))
- Geographic Information Systems (GIS) | University of California, Davis (via [Coursera](#))
- Plant Bioinformatic Methods Specialization | University of Toronto (via [Coursera](#))
- Agroforestry Specialization | University of Florida (via [Coursera](#))
- Agribusiness & Sustainable Food Production Economics Specialization | University of Illinois Urbana Champaign (via [Coursera](#))
- Agriculture, Economics and Nature | University of Western Australia (via [Coursera](#))
- DNA Decoded | McMaster University (via [Coursera](#))

Fellowships

Forward Program 	Mar 2025 – Jul 2025
<i>McKinsey.org</i>	Remote
Career-Prep Fellowship 	Aug 2023 – Oct - 2023
<i>Amal Academy (Stanford Seed-funded)</i>	<i>Faisalabad, Pakistan</i>
Foster Fellowship 	Mar 2019 – May - 2019
<i>Foster Learning Pakistan</i>	<i>Faisalabad, Pakistan</i>

Skills

- **Molecular Biology & Plant Pathology:** Pathogen isolation & identification, **PCR**, Gel Electrophoresis, **SSR Genotyping**, Marker-Assisted Selection (**MAS**), Primer Design. **Physiological Assays** (SPAD, IRGA, Porometer, Osmometer, Spectrophotometry). Biochemical profiling (MDA, POD activity).
- **Genome-Wide & Comparative Genomics:** HMM-based gene family identification (HMMER 3.0), protein domain verification (SMART, Pfam), gene structure analysis (GSDS 2.0), motif discovery (MEME Suite). **Synteny analysis** (MCScanX, Circos, TBtools-II), promoter cis-element analysis (PlantCARE), subcellular localization prediction (WoLF PSORT).
- **Phylogenomics & Molecular Evolution:** **Phylogenetic reconstruction** (MEGA 11/12; Maximum Likelihood, Neighbor-Joining; 1,000 bootstrap replicates). **Sequence alignment** (MAFFT, ClustalW). **Recombination detection** (RDP5, 7 algorithms). Positive selection analysis (MEME). **Evolutionary metrics** (KaKs_Calculator, DnaSP 6). **Tree visualization** (iTOL, ggtree).
- **Protein Structure & Docking:** Structure prediction (ColabFold), Protein-protein docking (HDOCK), **Molecular dynamics simulations** (GROMACS). PPI network analysis (STRING database).
- **Data Analysis & Visualization:** **R Programming** (tidyverse, ggplot2, ComplexHeatmap, pheatmap, circlize, seqinr), **Python** (BioPython, Pandas), **Statistical analysis** (ANOVA, Kruskal-Wallis, PCA, t-SNE), **RNA-seq analysis** (expression profiling, heatmap clustering), **GIS & Remote Sensing**.

- **Field & Agricultural Research:** Greenhouse & field trial management (RCBD, factorial design), Screening for biotic & abiotic stress tolerance in rice and other crops, Agroecological data collection, IPM/IDM practices, sustainable agriculture frameworks
- **Academic & Professional:** Scientific writing, conference presentations, Fundraising.

Recommendations

Dr. Muhammad Rashid

Deputy Chief Scientist Rice Group, Plant Breeding and Genetics Division,
Nuclear Institute for Agriculture and biology (NIAB) Faisalabad, Pakistan

Email: mrashid_niab@yahoo.com

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Relationship: *Fellowship Supervisor*

Prof. Dr. Sajid Aleem Khan

Professor,

Department of Plant Pathology, University of Agriculture, Faisalabad

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Relationship: *Master's Supervisor*

Dr. Muhammad Amjad Ali

Associate Professor,

Department of Plant Pathology, University of Agriculture, Faisalabad

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Relationship: *Bachelors and Master Instructor*