# Mohammad Sayeef Alam

Trondheim, Norway | alam.sayeef@gmail.com | +4748671602 | LinkedIn

### **Key Qualifications**

- PhD in Genetic Epidemiology, dual master's in Biostatistics, and bachelor's in Statistics.
- Six years of experience in data analysis in various statistical methods and datasets related to health.
- Knowledge of methods related to epidemiology, biostatistics, clinical trials, and genetics with hands-on experience of its implementation in both R and Excel.
- Technically minded with the ability to break down a complex solution to ensure understanding at multiple levels and heterogeneous audience.
- Collaborative and supportive team player, conscious of my role as a contributor to a good working environment.
- Ability to take on new challenges in a fast-paced environment and make decisions under pressure.
- Good communication skills in English.

# **Professional Experience**

Associate, Parexel - Remote, India

July 2021 – February 2022

- Supported the development of interactive platforms (e.g., RShiny) to conduct economic evaluations (e.g., cost-effectiveness, budget impact) under senior guidance.
- Performed statistical analyzes (e.g., Bayesian network meta-analysis, MAIC, STC) aligned with health technology assessment (HTA) guidelines (e.g., TSD, NICE) across multiple disease areas.
- Automated dashboards (Excel/VBA) to visualize key clinical endpoints (e.g. survival curves, demographics) for regional stakeholder reporting.
- Supported client engagements by helping to coordinate projects, manage their expectations, and ensured the timely delivery of analytical output.
- Contributed to statistical analysis plans (SAPs) by programming scripts, conducting feasibility assessments, and validating data sets for accuracy.
- Assisted in presenting analytical results to internal and external stakeholders, incorporating feedback to refine deliverables.

## **Research Experience**

**PhD**, NTNU – Trondheim, Norway

April 2022 – Current

- Applied advanced statistical methods to analyze high-dimensional genetic and real-world registry data, generating actionable insights for large-scale studies.
- Designed and optimized analysis pipelines, reducing computational time and resource usage, and increasing team productivity.
- Presented research findings at national and international conferences and published results in peer-reviewed journals, contributing to scientific knowledge and visibility.
- Fostered and expanded collaborations with external university research groups, driving successful joint projects.
- Built a supportive academic network during an international research stay, facilitating new research opportunities.
- Secured more than 150,000 NOK in competitive external funding from three agencies, supporting innovative research initiatives.

## **Project**

## **Nordic Shortfall Calculator**

Link provided on request

- Developed a proof-of-concept model that calculates the absolute and proportional shortfall of 4 Nordic countries, i.e., Denmark, Finland, Norway, and Sweden
- Provides up-to-date estimates of quality-adjusted life expectency of respective Nordic population based on the EQ-5D or 3D values.
- Features comparative options between different value sets for each country, discount rates, updates automatically with dynamic input, provides HRQoL, Cummulative Survival and Cummulative QALYs.

#### **Skills and Interests**

Software: R, SAS, Python, Git, Microsoft Excel, Powerpoint, and Word.

Transferable: Communicator, Detail-oriented, Problem solver, Relationship builder, Results-driven,

Strategic

Hobbies: Hiking, swimming and cooking for others

#### **Education**

MPhil in Biostatistics and Demography International Institute for Population Sciences	June 2020 – May 2021
MSc in Biostatistics and Demography International Institute for Population Sciences	July 2018 – June 2020
BSc in Statistics University of Calcutta	April 2014 – May 2018

## Publications (3 out of 4)

**Alam MS**, Thomas L, Brumpton B, Hveem K, Lundin KE, Withoff S, Jonkers IH, Sollid LM, Hjort R, Ness-Jensen E. *Population screening of adults identifies novel genetic variants associated with celiac disease.* Scientific Reports. 2025 Jun 5;15(1):19764.

Singh M, Alam MS, Majumdar P, Tiwary B, Narzari H, Mahendradhata Y. *Understanding the spatial predictors of malnutrition among 0–2 years children in India using path analysis*. Frontiers in Public Health. 2021 Jul 30;9:667502.

Sen K, Sinha A, Sen S, Chakraborty S, **Alam MS**. *Thyroid function test in COVID-19 patients: a cross-sectional study in a tertiary care hospital*. Indian Journal of Endocrinology and Metabolism. 2020 Nov 1;24(6):532-6.

## Conferences (3 out of 4)

A genome wide association study reveals novel locus at 5p15.33 to be associated with celiac disease: results from population-based screening of adults, HUNT4. **20th International Celiac Disease Symposium – Sheffield, UK (oral)**. September 2024

Assessing the susceptibility of celiac disease by polygenic risk scores: analysis of a population-based cohort, the HUNT study. **Nordic Conference on Future Health – Trondheim, Norway (oral)**. September 2024

Prevalence of communicable disease and its epidemiological correlates in major urban states in India: Evidence from a nationally representative sample. **16th International Conference on Urban Health – Xiamen, China (oral)**. November 2020

#### References

References can be provided on request.