Two Postdoc Positions Availble

The research group of Dr. Yongtao Guan (https://haplotype.org), recently revived at the Beijing Institute of Mathematical Sciences and Applications (BIMSA), is seeking two highly motivated postdoctoral researchers to work on cuttingedge projects in statistical genetics, Bayesian statistics, and computational methods, with applications in genomics and genetic diagnosis.

Key Research Areas

- Bayesian statistics and variable selection for genetic and genomics studies.
- Haplotype variation, local ancestry, and haplotype-based analyses.
- Development of computational tools for genomic studies and genetic diagnosis.

Qualifications

- Ph.D. in quantitative fieleds such as mathematics, statistics, and compute sciences, or
- Ph.D. in a non-quantitative field but with extensive experiences in austism or cancer genomics.
- Strong programming (R/Python/C++) and quantitative skills, or
- Demonstrated proficiency with genomic data (WGS, RNAseq, multiomics, single cell).
- Hands-on experiences with training deep learning models is a plus.

Application Process

Interested candidates should email Dr. Guan at ytguan@bimsa.cn with:

- An updated CV
- Names and contact information for three references (letters not required initially).

Shortlisted candidates will be invited for interviews and encouraged to submit a formal application via the BIMSA Postdoc Portal: https://www.bimsa.cn/join/join_postdocs.html

Why Join Us?

- Work in a dynamic, interdisciplinary team at BIMSA, a leading research institute in China.
- Collaborate on highly innovative projects with opportunities for methodological innovation.
- Collaborate with genetic diagnostic labs for projects with significant practical impacts.
- Competitive salary and benefits supported by BIMSA, not tied with a specific grant.
- Dr. Guan has an excellent track records in mentoring postdocs and graduate students.
- Dr. Guan's approach to mentoring is handson and detail oriented.

Application Deadline

• Rolling until positions are filled. Informal inquiries are welcome.