



*** Are you...** Graduating with a degree in biology, computer science, physics, math, or engineering? **Interested in working as a computational analyst with Harvard & MIT scientists?** Seeking on-the-job training alongside a small group of your peers? **Want to apply your coding skills to develop cutting-edge tools** that will...

...solve interdisciplinary problems...

...help to revolutionize the way we do cancer research...

...identify meaningful patterns in large sets of emerging biomedical data...

*** If so...** Apply for the new position **Bioinformatics/Computational Associate** at the Broad Institute of MIT & Harvard, located in Kendall Square, Cambridge MA.

Characteristic Duties:

- Develop data analysis algorithms to share with the global biomedical research community
- Conceive, implement and test statistical models to analyze genomic datasets
- Explore novel data visualization tools and big data approaches in biology

Candidate Eligibility:

- BS or MS degree in biology, computer science, physics, math, engineering, or a relevant field by Spring 2013
- Interest in applying your skills to research aimed at improving human health
- Programming skills in MATLAB, R, Python or similar language
- Ideally, candidates will begin work on July 8, 2013

How to Apply:

- Submit resume and cover letter for Requisition #905 at: www.broadinstitute.org/careers/career-center
- Complete application must be received by February 1, 2013
- For questions or concerns, please email ATP@broadinstitute.org

For more information, visit: www.broadinstitute.org/atp

Complete application is due: February 1, 2013