

Are you...

Graduating with a degree in **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?

Intrigued about applying your coding skills to develop **cutting-edge tools** that will...

...solve **interdisciplinary problems**...

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

...and help to **revolutionize the way we do cancer research?**

If so...

Apply for the new position **Bioinformatics/Computational Associate**

at the Broad Institute of MIT & Harvard, located in Kendall Square, Cambridge MA.



How to apply:

- 1) Send cover letter, resume, and one letter of recommendation to:

Dr. Rachel Woodruff

Email: woodruff@broadinstitute.org

Fax: (617) 800-1145

- 2) Submit online application for Requisition #743 at:

<http://www.broadinstitute.org/careers/career-center>

Complete application is due:

March 30th, 2012

Characteristic Duties:

- Develop data analysis strategies and algorithms to share with the global cancer research community
- Conceive, implement, and test statistical models to analyze large cancer and sequencing data sets
- Explore novel data visualization tools, with emphasis on those that integrate diverse data types

Candidate Eligibility:

- Must hold a B.S. or M.S. degree in a relevant field by June 15th 2012
- Must have interest in applying your skills to research aimed at improving human health
- Must have familiarity with programming/analysis using Matlab and/or R (or similar)

Instructions for Application:

- Complete application must be received by **March 30th 2012**
- Ideally, candidates will begin work on **June 25th, 2012**
- Application must include a cover letter, your resume, and one letter of recommendation
- We prefer that ALL application materials be emailed or faxed
- If application materials must be mailed, postmark by **March 30th 2012**