



# The Laboratory for **RNA-Based Lifeforms**

## Computational Biologist - Virology (Postdoc or equal)

Join “[The Laboratory for RNA-Based Lifeforms](#)” in University of Toronto’s Donnelly Centre. Together with **Dr. Artem Babaian** the work will develop the state of the art for the detection and sequence analysis of RNA viruses and virus-like agents.

Building upon the *Serratus* project ([www.serratus.io](http://www.serratus.io)) we will explore the far limits of Earth’s Virome. Briefly, we developed an open-source AWS-cloud backed computing architecture to analyze 5.7 million sequencing datasets (10.2 petabytes) and discover >130,000 novel RNA viruses (only 15,000 were known previously)... in only 11 days. Learn more in our first [Nature](#) paper, or watch our ISMB22 talk, “[Serratus: Hacking Earth’s Virome](#)”.

We have a very collaborative and fast-paced research environment with international laboratories. Our work aims to also create free and open data resources to catalyze the field of virology globally.

### Who you are

Computational virology is undergoing a renaissance. We’re looking to support passionate, self-motivated scientists with a clear vision to take on high risk research ideas and move the field forward. In your application provide evidence for initiation and follow-through of scientific ideas. We’re open to applicants with expertise in the areas of computational biology.

Programming experience is required, although inversely weighted with biological experience. Advanced algorithm-level coding experience is not necessary, but you must be willing to learn/teach and converse with computer-science oriented scientists. Collaboration with and mentoring of junior scientists is expected.

**(cont’d)**

## Qualifications

### Minimum (*provide evidence please*)

- PhD in virology, ecology, epidemiology, genetics, biochemistry, molecular biology or computer science
- Fluency (show example code) in  $\geq 1$  programming language, such as R or Python
- Highly motivated to learn, teach and pursue science across disciplines
- Excellent written and verbal communication skills

### Preferred qualifications

- Innovative in computational virology, genomics, phylogenetics, ecoinformatics, RNA structure-function, and/or clinical/veterinary virology
- Experience with “big data” and cloud-computing infrastructures
- Demonstration of scientific mentorship, organizational and/or leadership skills

## Where we are

The [Donnelly Centre](#) is located in the downtown (St. George) University of Toronto campus in Canada. We are embedded amongst several world-class computational biology groups and the Donnelly offers a cross-disciplinary repertoire of biological laboratories for validation/collaboration. The University of Toronto is the top Canadian university, and offers countless opportunities for extending your academic experiences.

The start date is immediate/flexible, and we encourage applicants from diverse backgrounds. The position can be full-time or contracted consultant, an ideal contact will be Toronto-local or willing to relocate, but remote work is possible for an exceptional applicant.

We will be recruiting until this vacancy is filled.

Email your package to [artem@rRNA.ca](mailto:artem@rRNA.ca) with subject **Application id0001 - comp viro**