## **Advanced Application of Bioinformatics in Genetic Research.**

This special topic course is designed for students who took GMS6014 and want more hands-on experience in applying bioinformatics and computational approaches to solve real research problems. Unlike GMS6014, knowledge of programming (i.e. a college-level course in basic programming concepts or previous experience in writing scripts) is required. In lieu of this requirement, a strong commitment to learning the programming language Python in parallel to this course must be demonstrated by communicating with the course director.

In addition to learning basic scripting skills using Python and/or R, students will take a bioinformatics project pertinent to their research work (or assigned by the instructor) and develop it over 10 weeks. Examples of suitable projects include, but are not limited to, analyzing high throughput sequence data sets, comparative analysis of sequences, automated data extraction and integration, modeling a biological process, etc. Class will meet at 10:30-11:45 on every Thursday during modules 1 & 2 of the Fall semester (August 24 -October 26). In person meeting will be in CGRC-291. Zoom is optional for the second half.

There will be two types of class meetings. The *lectures* will be given by the instructors on topics related to the interest of this group. The *discussion classes* will be mediated by the faculty to discuss the resources, strategy, planning, and development of the projects in a learn-from-peer environment.

The class is limited to a maximum of 7 students. A laptop computer with a wireless connection is required for participating in discussions and sharing information.

## • Recommended reference book:

"Bioinformatics Data Skills: Reproducible and Robust Research with Open Source Tools." by Vince Buffalo (2015, O'Reilly; Amazon Link) – this book is more useful for those who already had some exposure with the Linux (Unix) system but want to be more efficient.

## • Course Directors:

Dr. Raad Gharaibeh (<u>Raad.Gharaibeh @medicine.ufl.edu</u>)
Dr. Lei Zhou (leizhou @ufl.edu)

Tentative schedule – based on the last time this course was offered. The actual schedule will be modified based on the projects involved.

| Date     | TOPIC   |   |             |          |
|----------|---|---|-------------|----------|
|          | Concept                                       | Practice  | Instructors | Location |
| Aug. 24  | Intro. How plan for a bioinformatics project. | Initiation of projects                                  | LZ, RG      | CGRC-291 |
| Aug. 31  | Intro to Python                               | Project Objectives and background                       | LZ, RG      | CGRC-291 |
| Sep.7    | RNA-Seq analysis                              | DEG identification                                      | RG, LZ      | CGRC-291 |
| Sept. 14 | Intro to R                                    | PCA, Volcano plot, Heat Maps for RNA-Seq.               | RG, LZ      | CGRC-291 |
| Sept. 21 | Project discussion                            |   | LZ ,RG      | CGRC-291 |
| Sept. 28 | Functional analysis of RNA-Seq data           | Pathway, GO, GSEA.                                      | RG, LZ      | CGRC-291 |
| Oct. 5   | Project discussion                            |   | RG, LZ      | CGRC-291 |
| Oct. 12  | Cross genome comparison                       | Comparing expression profiles from different organisms. | LZ, RG      | CGRC-291 |
| Oct. 19  | Trouble-shooting meeting                      |   | RG, LZ      | CGRC-291 |
| Oct. 26  | Final progress report                         |   | LZ, RG      | CGRC-291 |