09-02-25_weekly-update

Cappy Pugliese

9/2 – 9/12

This week (9/2 - 9/5)

Weekend

- worked on some grading for bio 1108
- read book for grad 6001

Tuesday

- read book for grad 6001
- submitted assignments for popgen

Friday

- it worked, but I realized there was actually more missing than I originally thought
- merged the ones that did run into the folder with the others
- created a new script to run the rest of the missing files
- put it running

Week 2 (9/8 - 9/12)

Monday

- everything worked, except for a handful of files
  - those files did not exist in the bam folder to begin with so it couldn't work anyway
  - unsure what to do about the missing files
- lab meeting, stuff to do for next week:
  - finish running the data

- genomics dv
- data analysis with pixy or pcangst
- vcfr for figures
- start playing around with the data
- maybe build a tree out of the data

Tues-Thurs

- had some difficulties getting the consolidate gvcfs script to work
- needed an interval list, which I needed to create
- was eventually able to figure it out and put the script running
- script worked and generated a genomicsDBI workspace

Friday

- made script for the joint-calling step
- put it running
- failed at first, had to trouble shoot (there was a typo)
- successfully ran
- put another one going specifiying ploidy = 1