This page is now deprecated. All scripts used by the lab are maintained on github or the HPC

HPC

- logging onto the HPC
- align_things.pbs

R

- facets.R
- ggplot_facets.R
- ggplot.intro.R

Perl

- Perl - Biobricking Primer Designer - interactive version
- Perl - Biobricking Primer Designer Source Code
- Perl - Overlap Oligos Designer - interactive
- Perl - Overlap Oligos Source Code
- Perl - Convert SNPs to amino acid changes