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**
- facs.R
- ggplot_facs.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