

# Tentative Project Design

1. Develop selection lines by
  - A. Direct selection for long or short horn size
  - B. Eliminate sexual antagonism (no-SA) by eliminating fecundity selection
    - a. Breeding happens in mass culture, but all females contribute equal offspring to subsequent generation
    - b. Expect horn size to increase due to sexual selection on males
2. Assemble the genome and map trait loci using combination GWAS / linkage mapping.
  - A. Generate recombinant lines by crossing high and low selection lines
  - B. Sequence whole genomes of selection lines to phase SNPs in recombinants
  - C. Use linkage info from recombinants to scaffold assembly and map SNP associations with horn size
3. Compare loci from no-SA regime to those mapped by direct selection

# Courses Taken So Far

| COURSE                    | INSTRUCTOR                                | SEMESTER   |
|---------------------------|---|------------|
| BIOMETRY                  | Dr. Walter Schargel                       | Fall '16   |
| BIOINFORMATICS            | Dr. Jeff Demuth                           | Fall '16   |
| EVOLUTION                 | Dr. Paul Chippindale                      | Spring '17 |
| ESSENTIALS OF<br>GENOMICS | Dr. Matthew Fujita<br>and Dr. Todd Castoe | Spring '17 |