This course is designed for two audiences: graduate students in biology and graduate students in statistics. Biology students will learn about using aster models in research. Areas of life history analysis covered include comparison of fitness between groups, estimation of fitness landscapes and response to selection, estimation of population growth rates, quantitative genetics of fitness, and Fisher’s fundamental theorem of natural selection. Topics include graphical models, exponential families, parameterizations of aster models (six of them), sufficiency, maximum entropy, multivariate monotonicity, interpretation of aster models, aster models with random effects, aster models with subsampling, parametric bootstrap. All of these will be discussed with emphasis on their practical effects on aster analysis.

Statistics students will learn about algorithms for generalized linear models and aster models with random effects and about exponential family theory relevant to aster models, including conditions for existence (in the conventional sense) of maximum likelihood estimates in exponential families and how to do valid hypothesis tests and confidence intervals when they do not exist (in the conventional sense).

We hope mixing graduate students in biology and statistics will lead to each group learning from the other. Biology students may not understand all the theory but should learn enough to gain important insights about issues that arise in practice (for example, what to do when maximum likelihood estimates do not exist and the summary function in the aster package complains about “directions of recession”). Statistics students may not understand all the biology but should learn about important issues that arise in application of statistics in real science (which are very different from toy examples in textbooks and statistics journals).