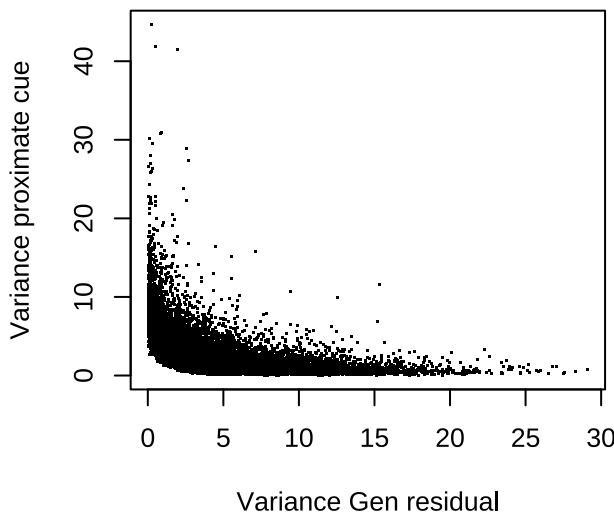
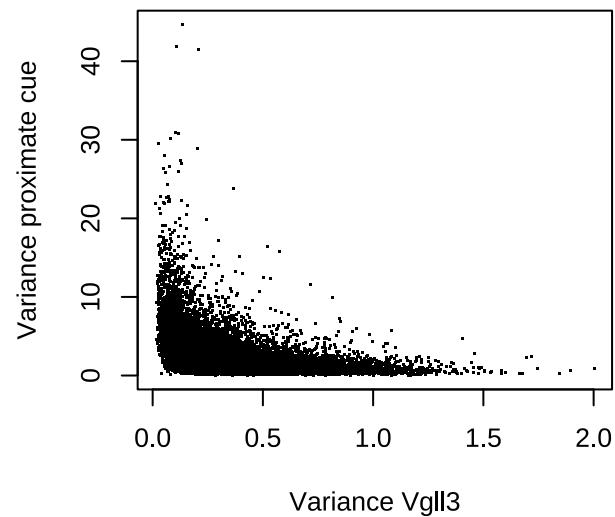
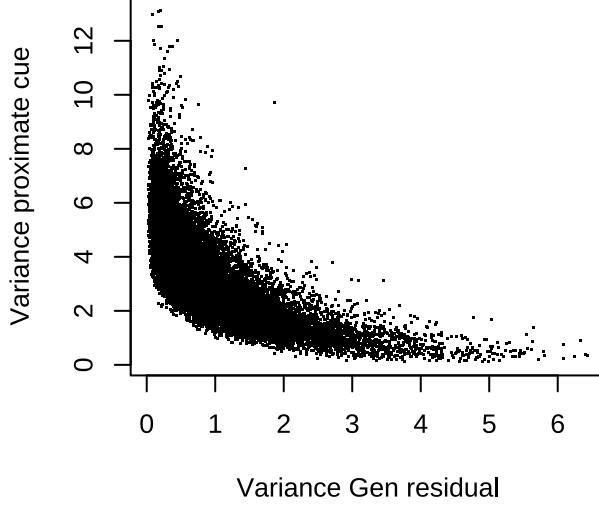
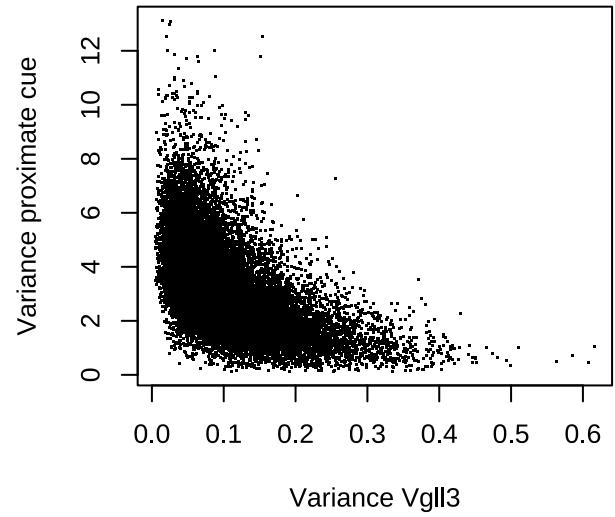
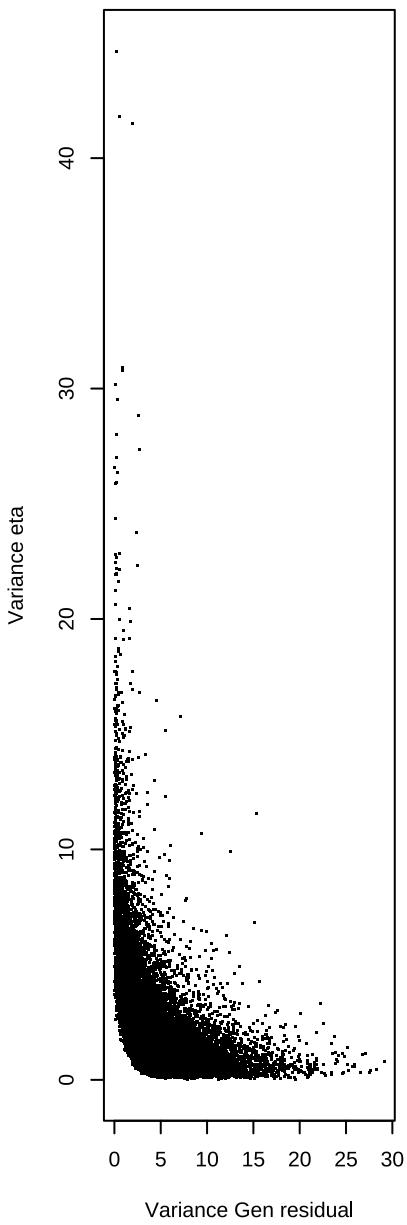
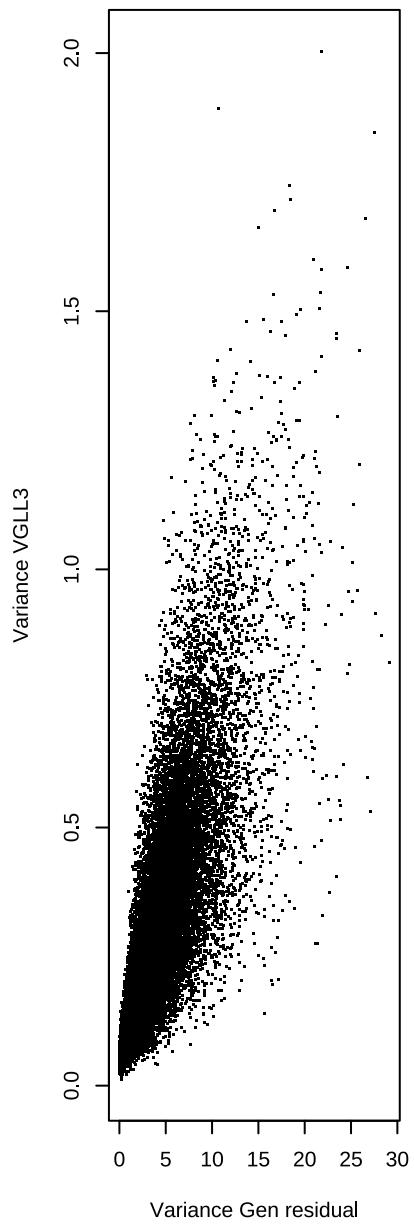
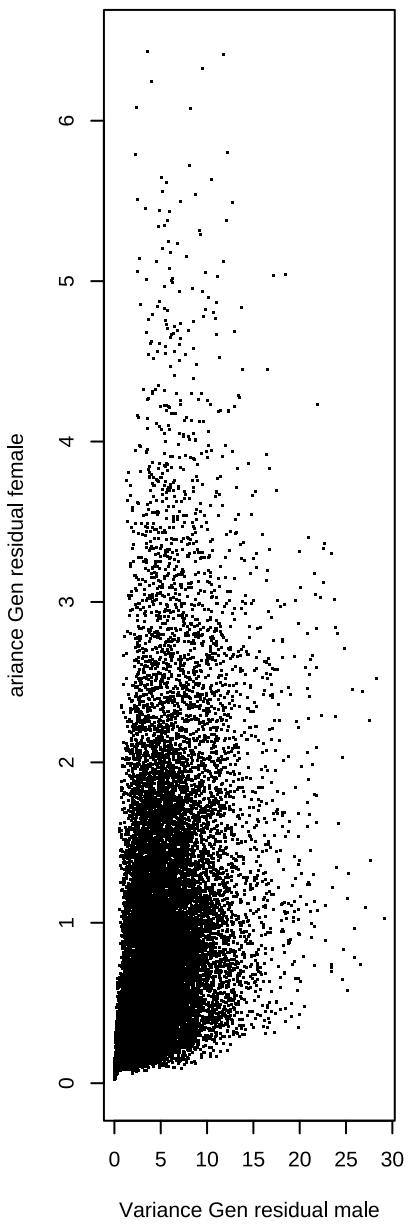
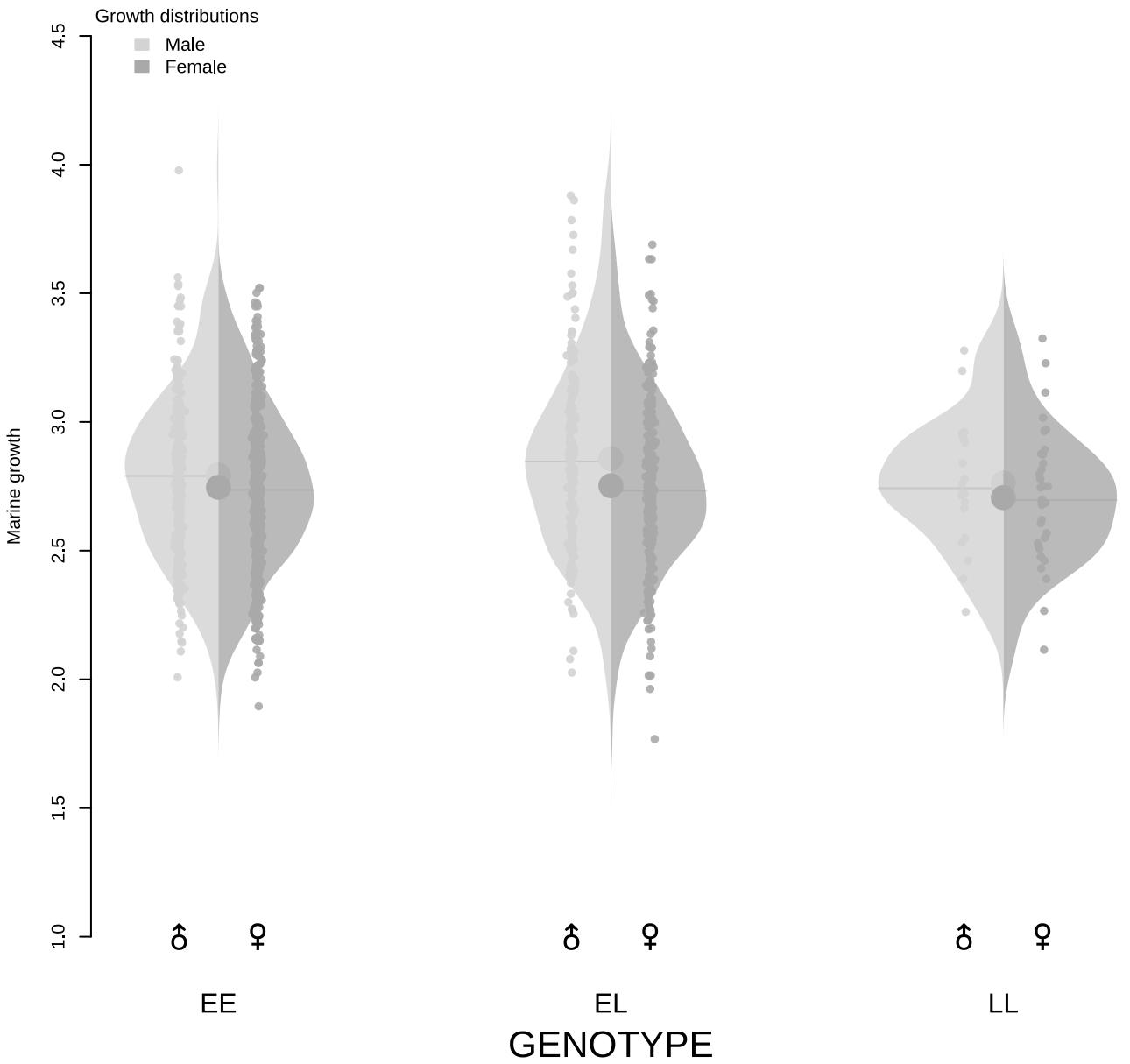
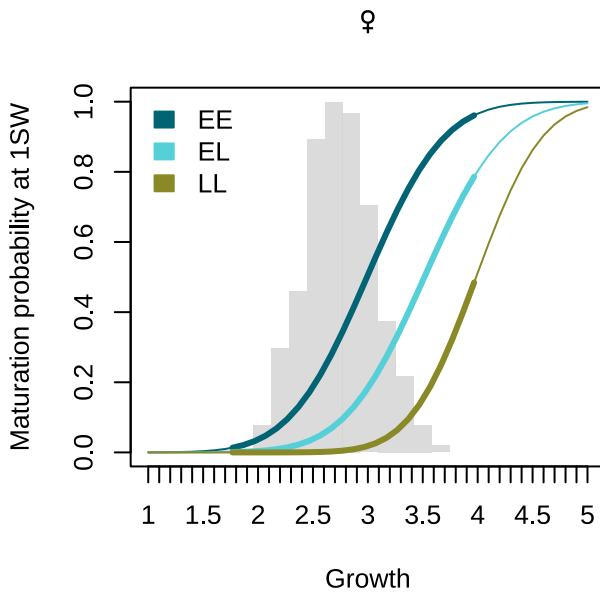
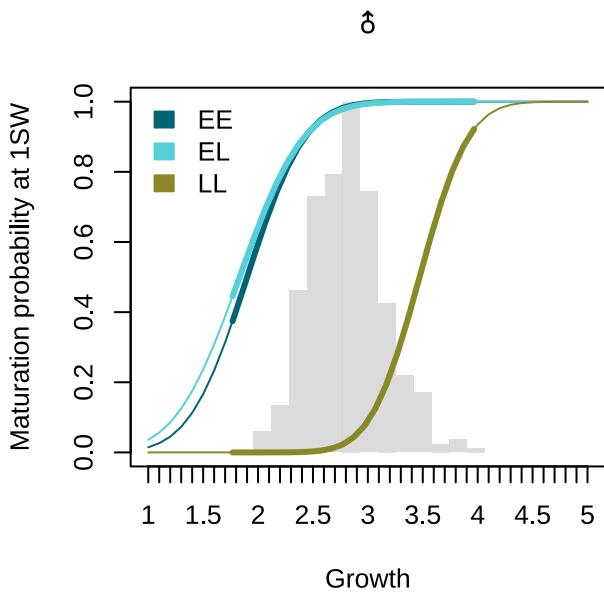
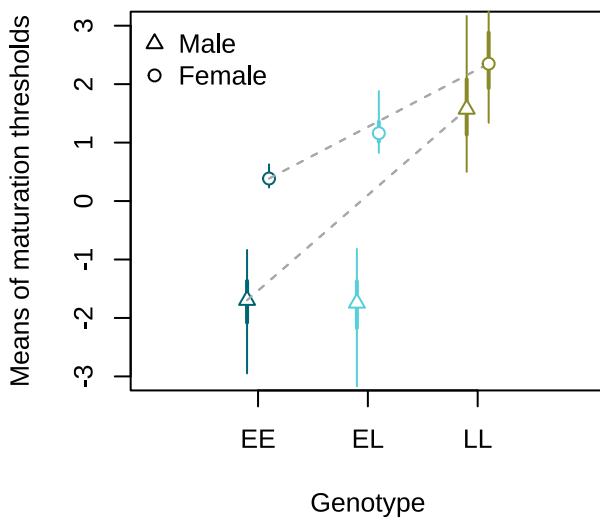
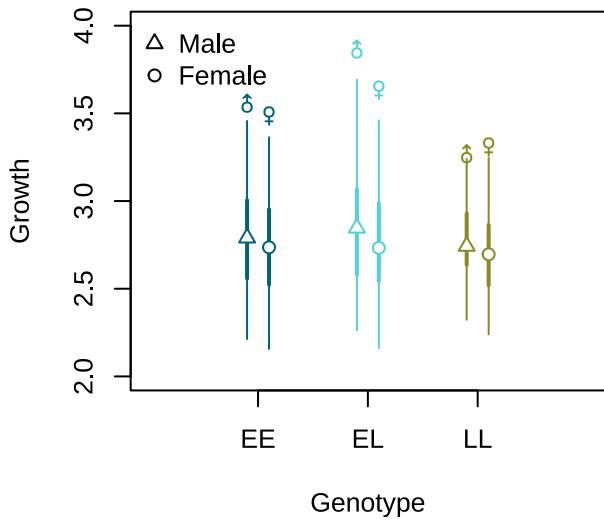
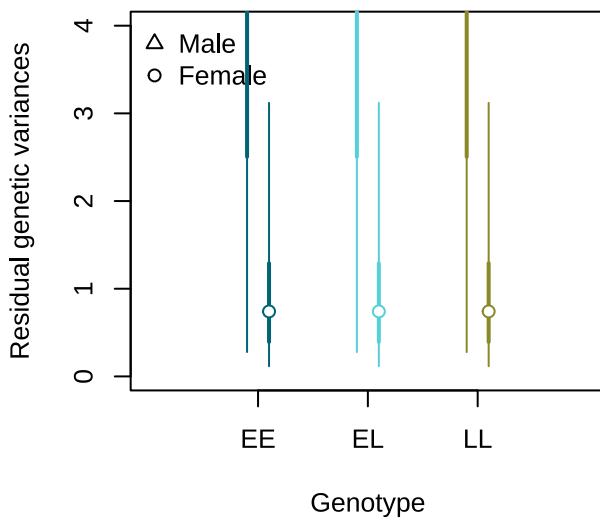
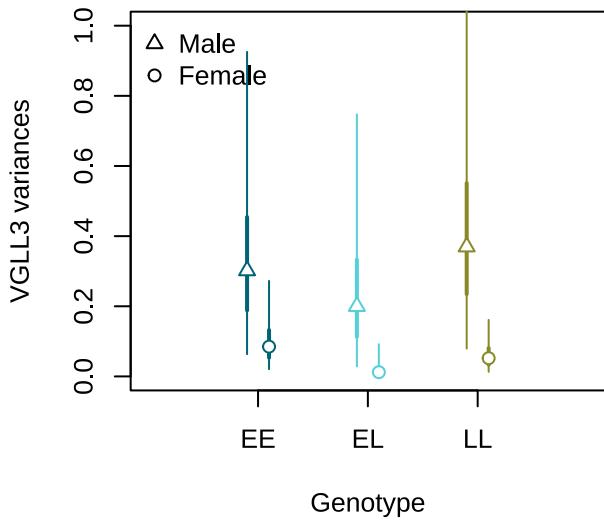


Male**Male****Female****Female**

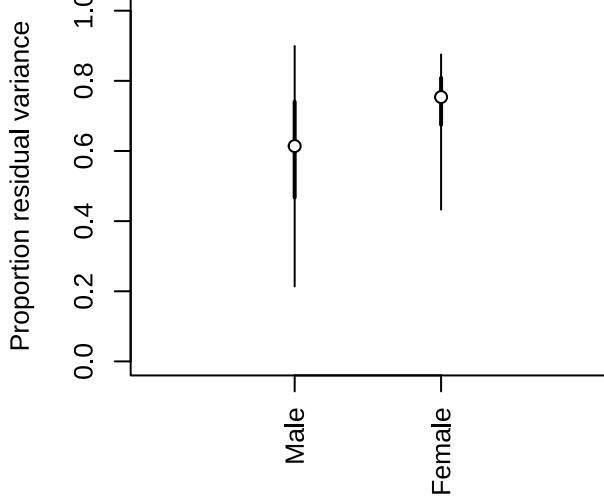




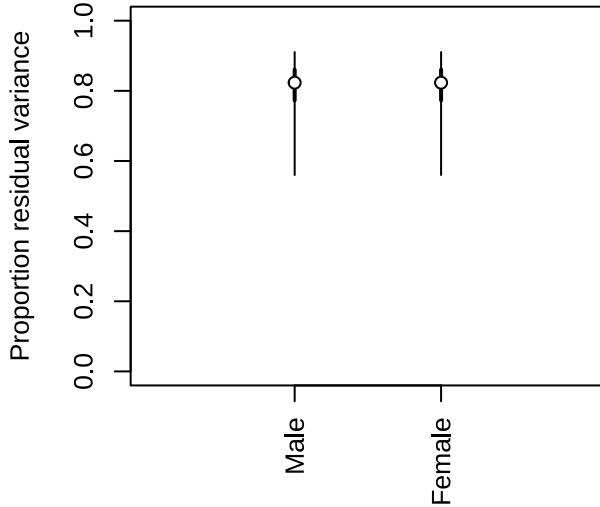




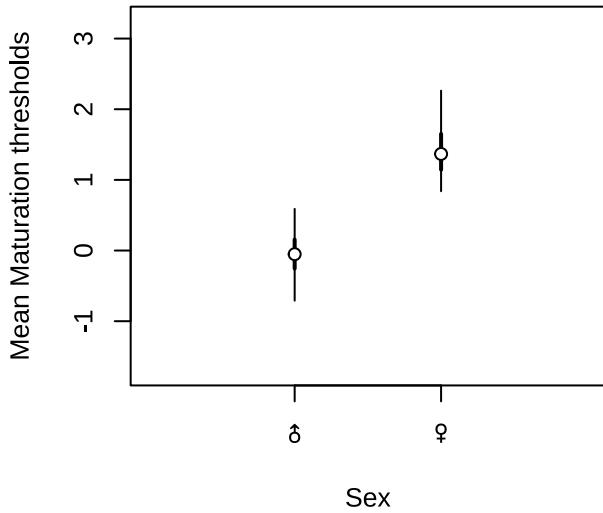
Environment



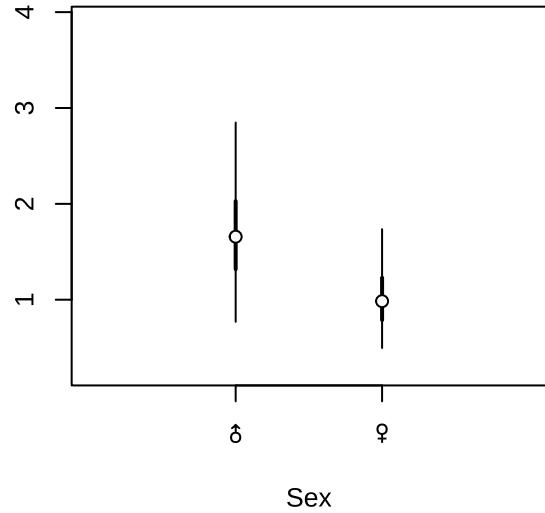
Genetic



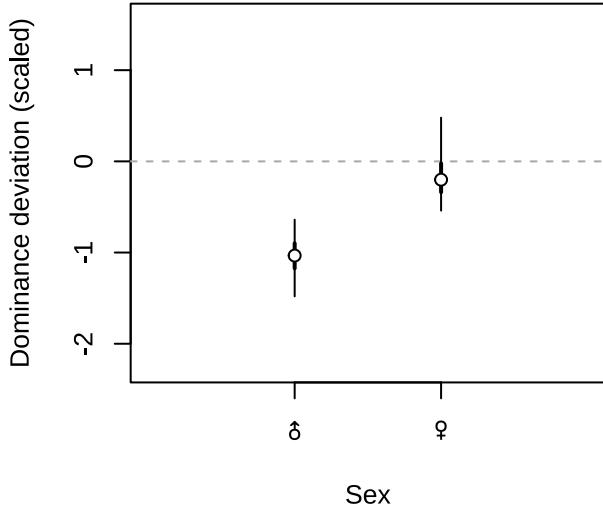
Mean Maturation thresholds



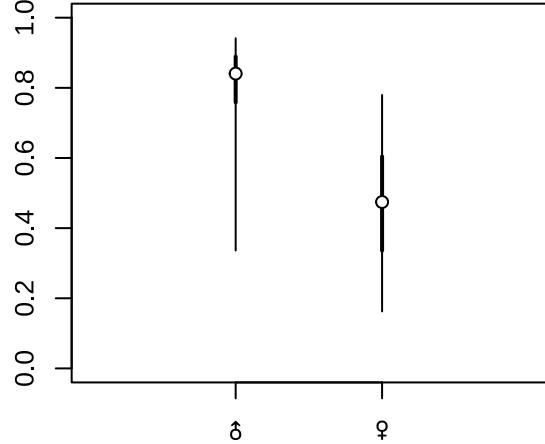
Genotypic value

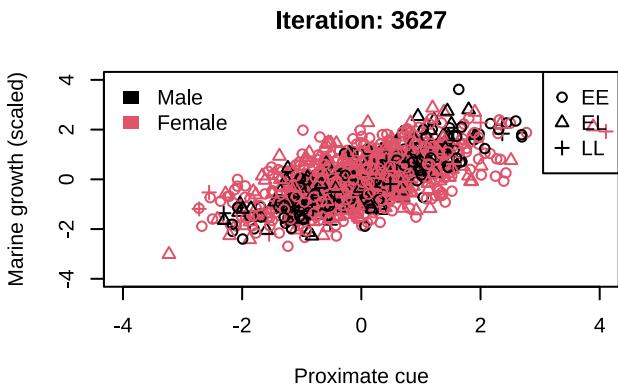
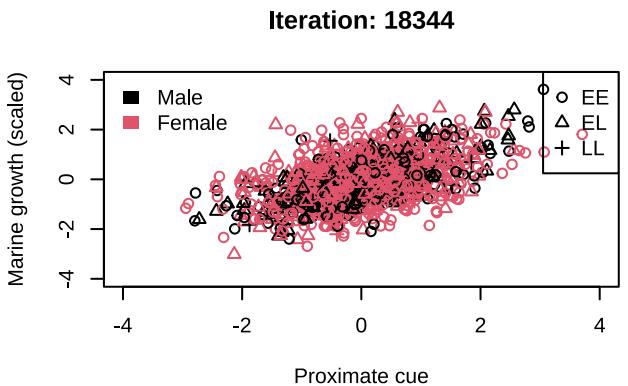
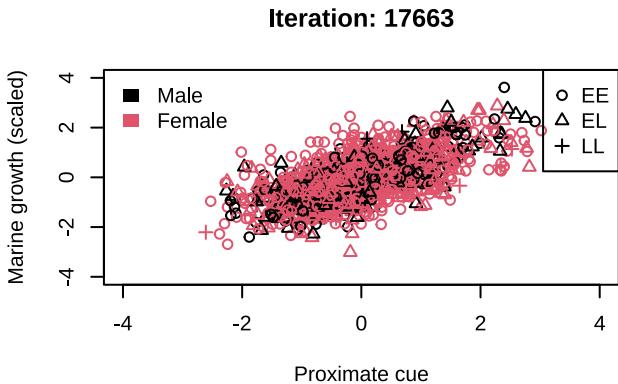
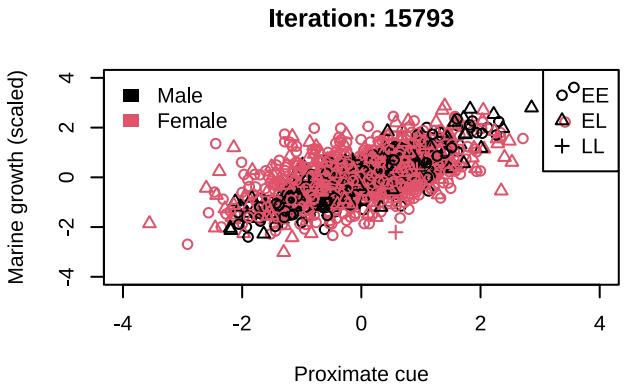
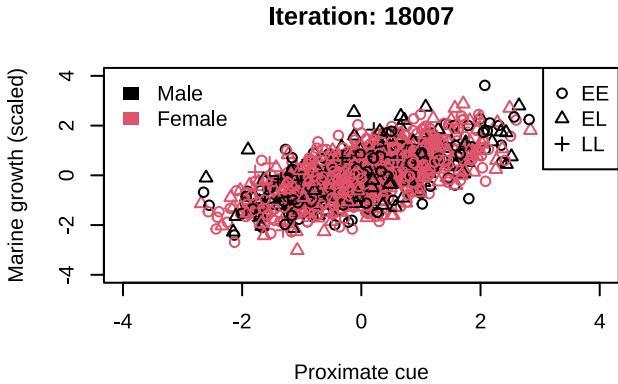
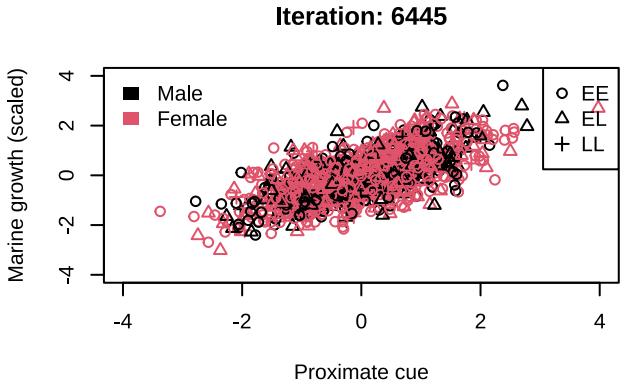


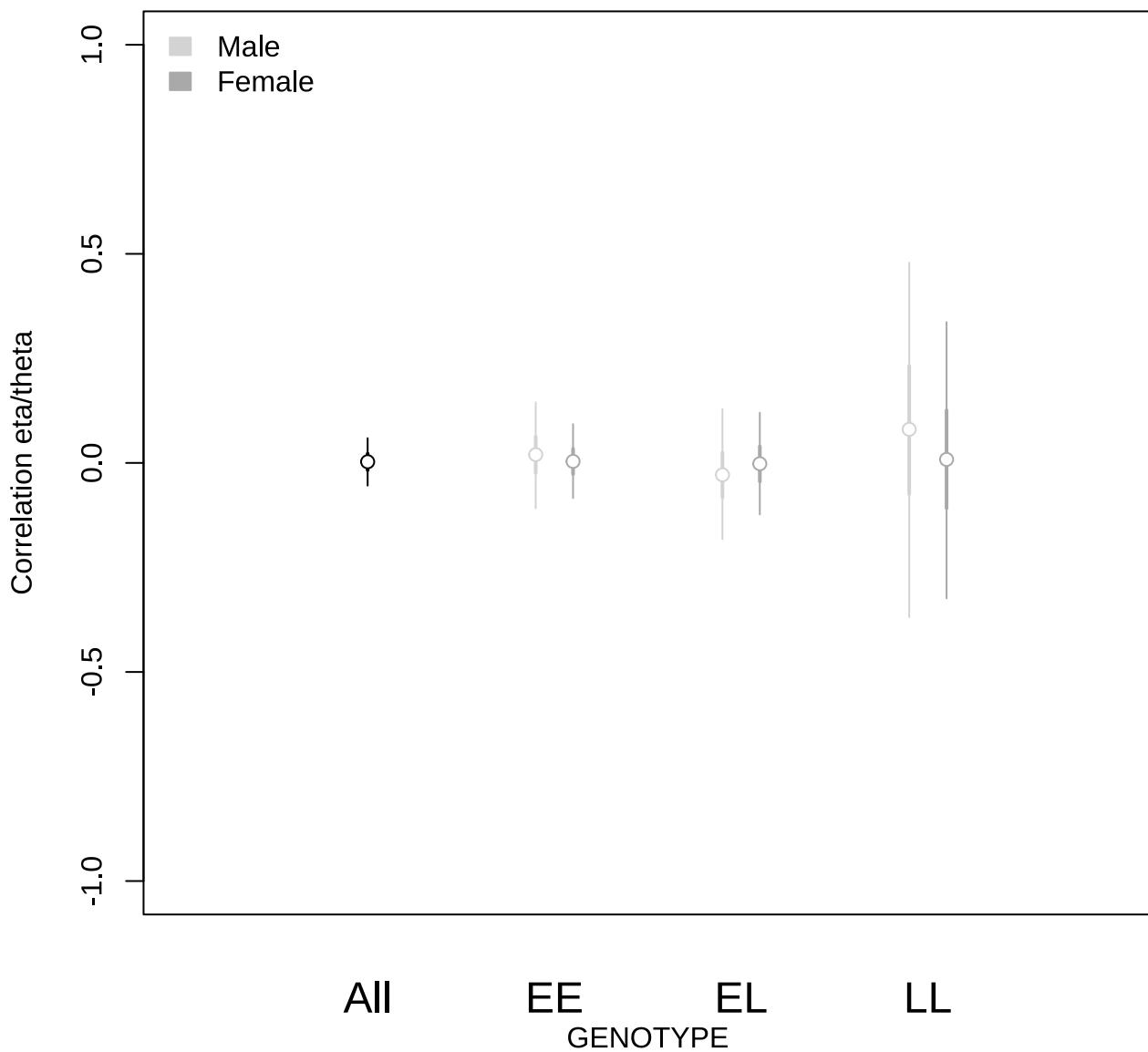
Dominance deviation (scaled)



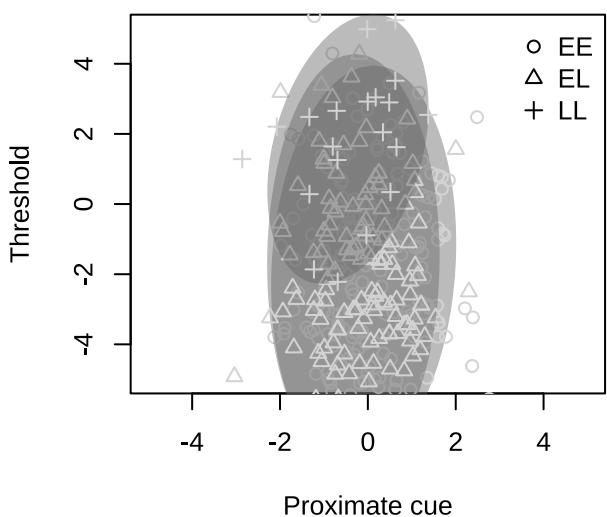
Total variance contribution



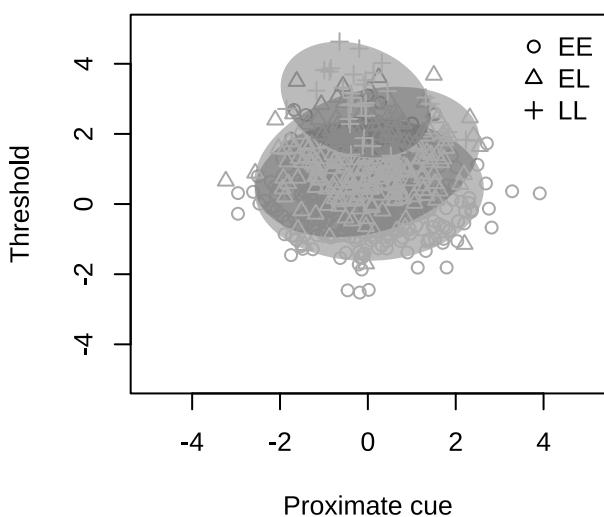




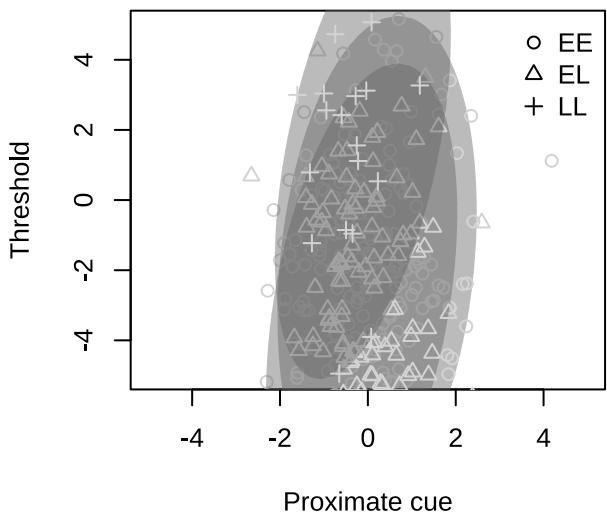
Male (iteration: 1098)



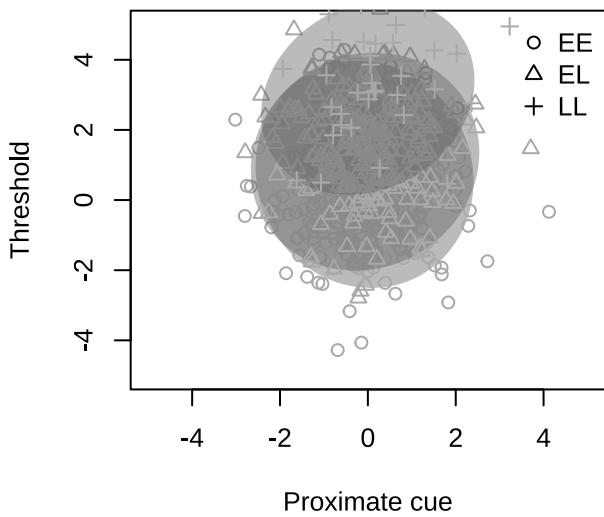
Female (iteration: 1098)



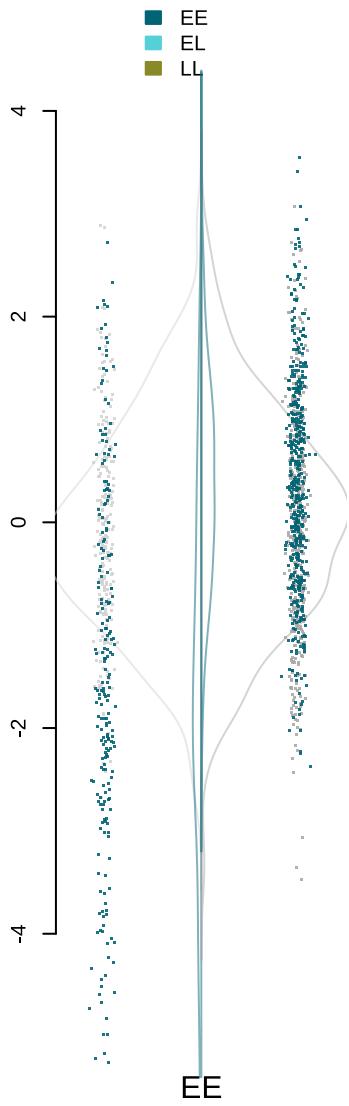
Male (iteration: 9506)



Female (iteration: 9506)



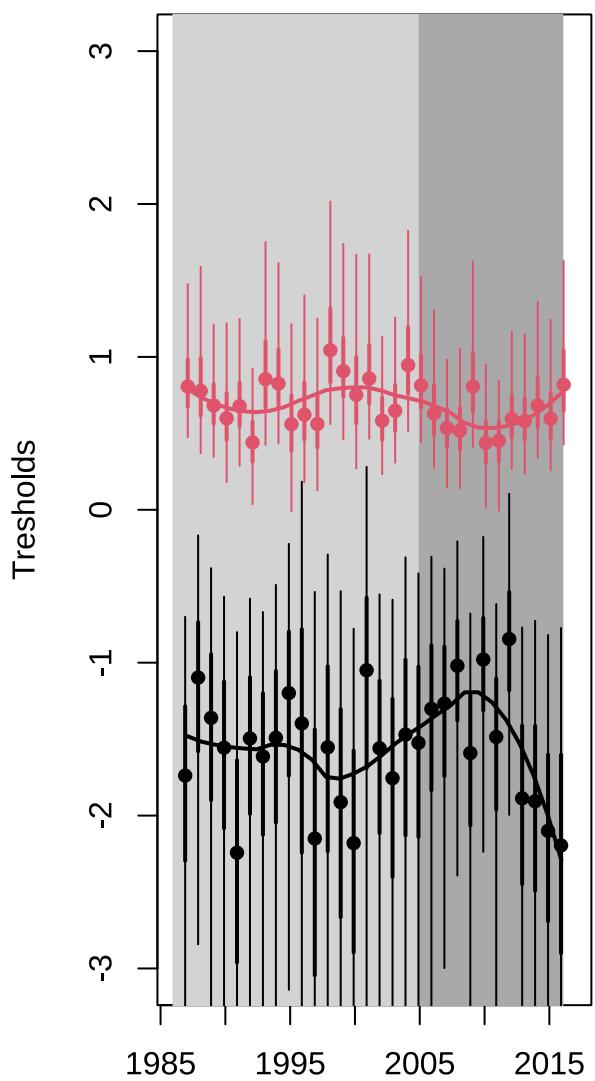
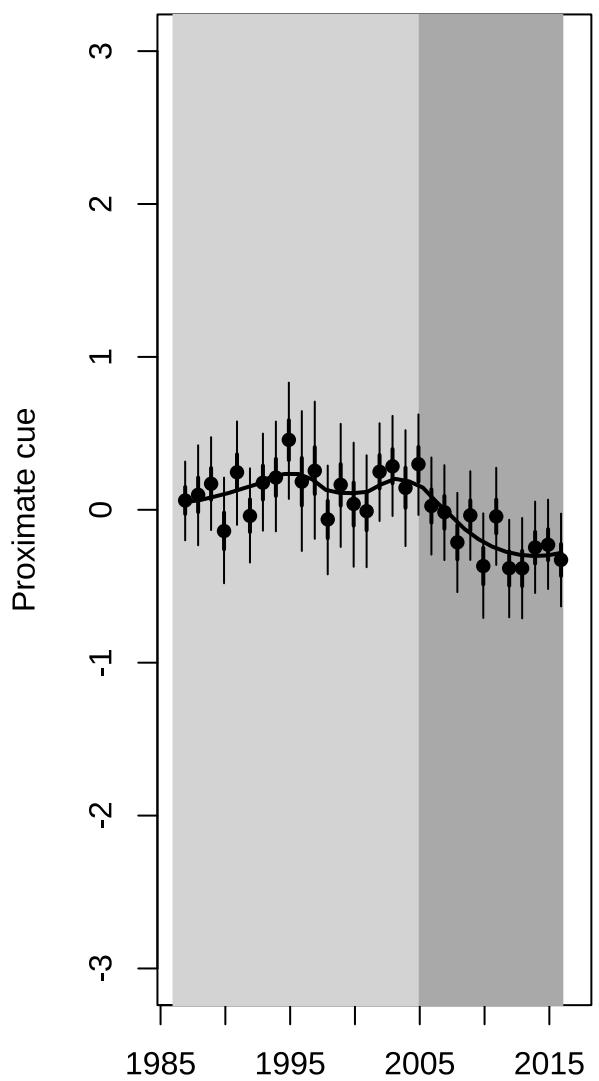
Thresholds distributions



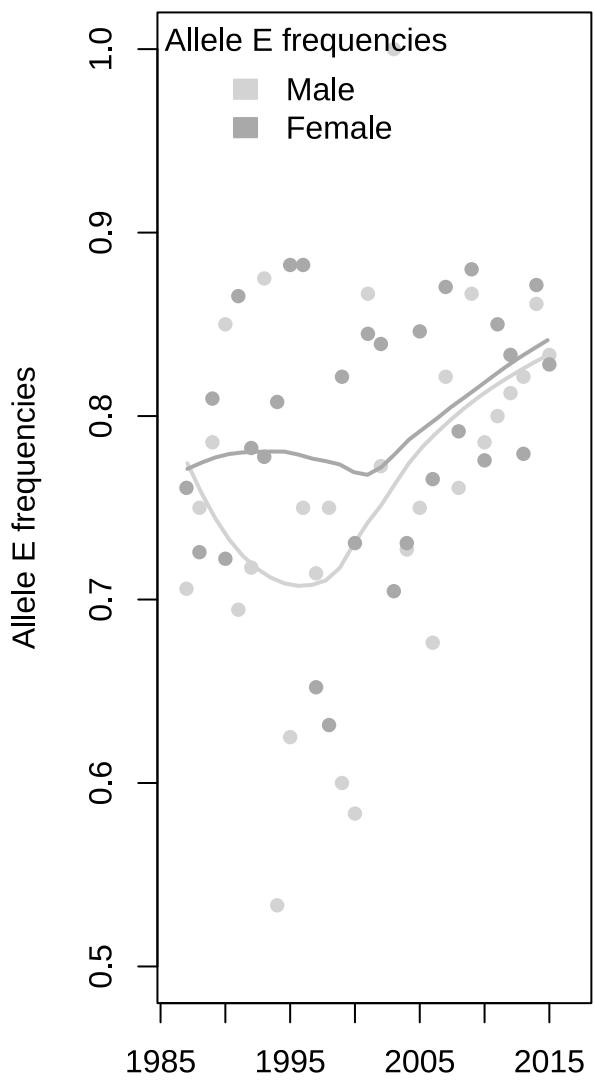
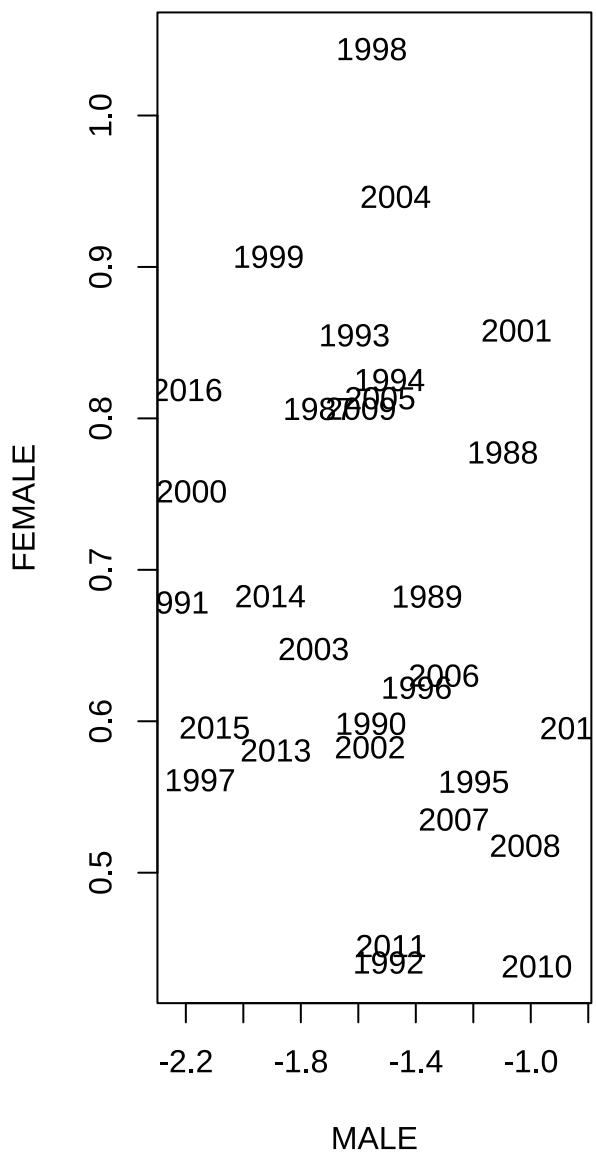
Cue distributions

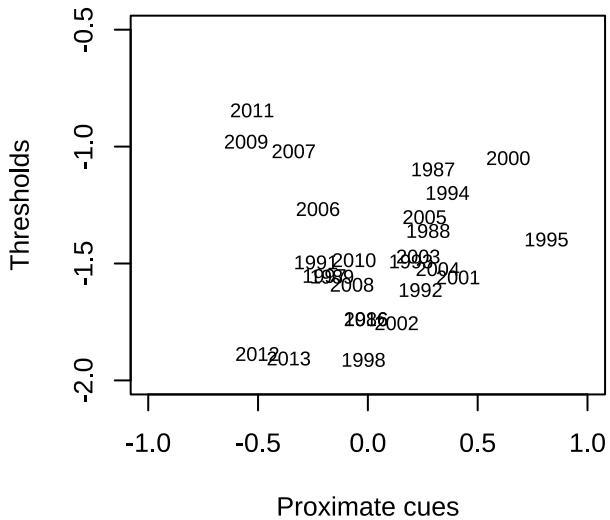
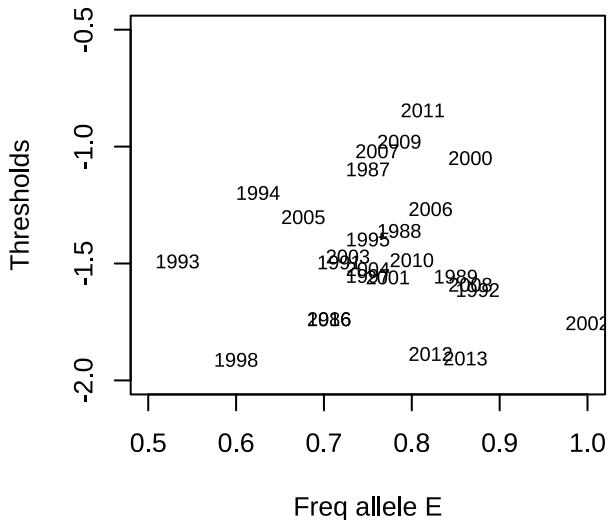
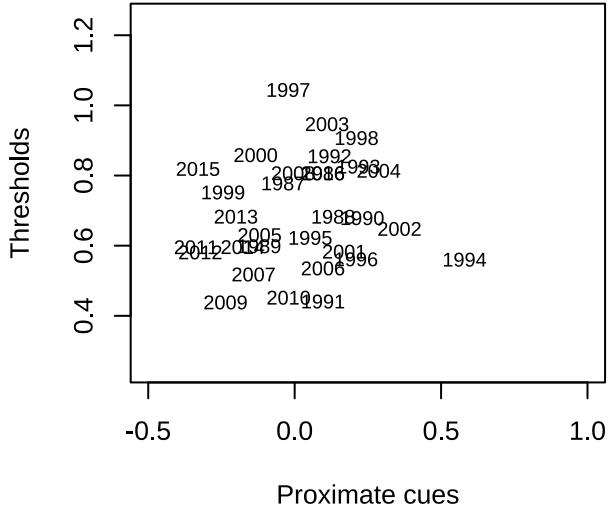
Male
Female

GENOTYPE



Thresholds (medians)



Male**Male****Female****Female**