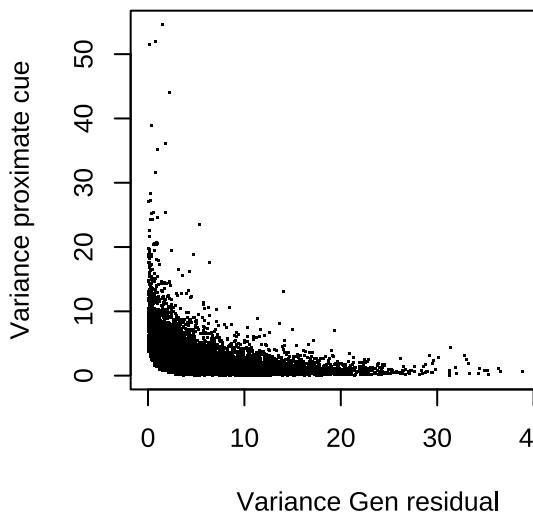
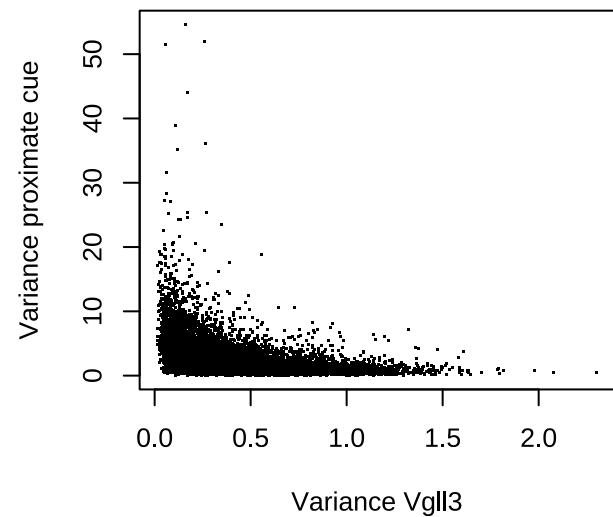
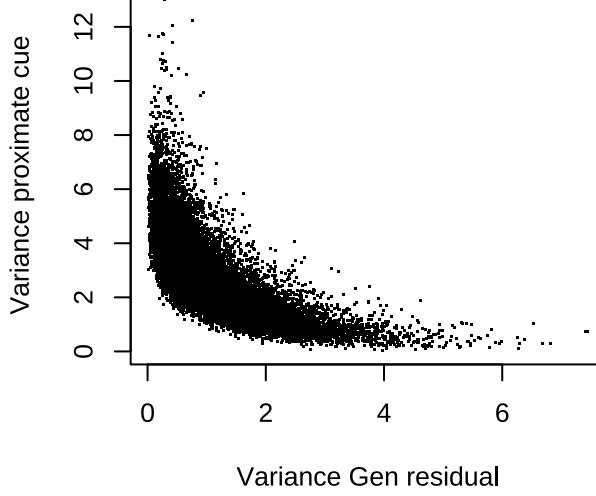
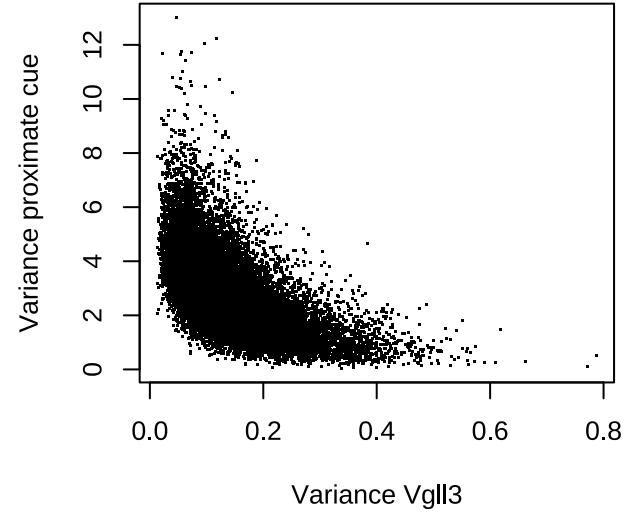
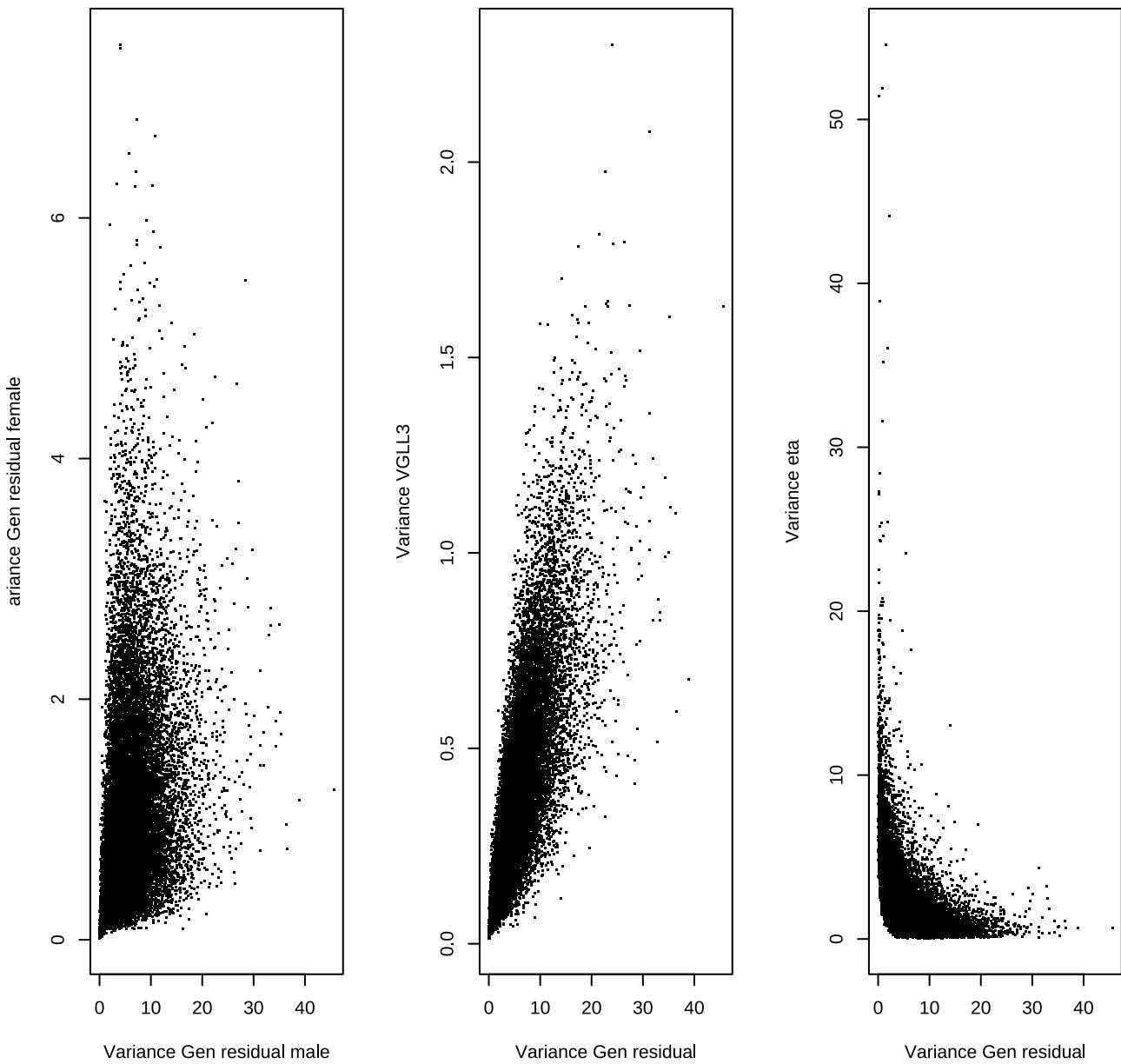
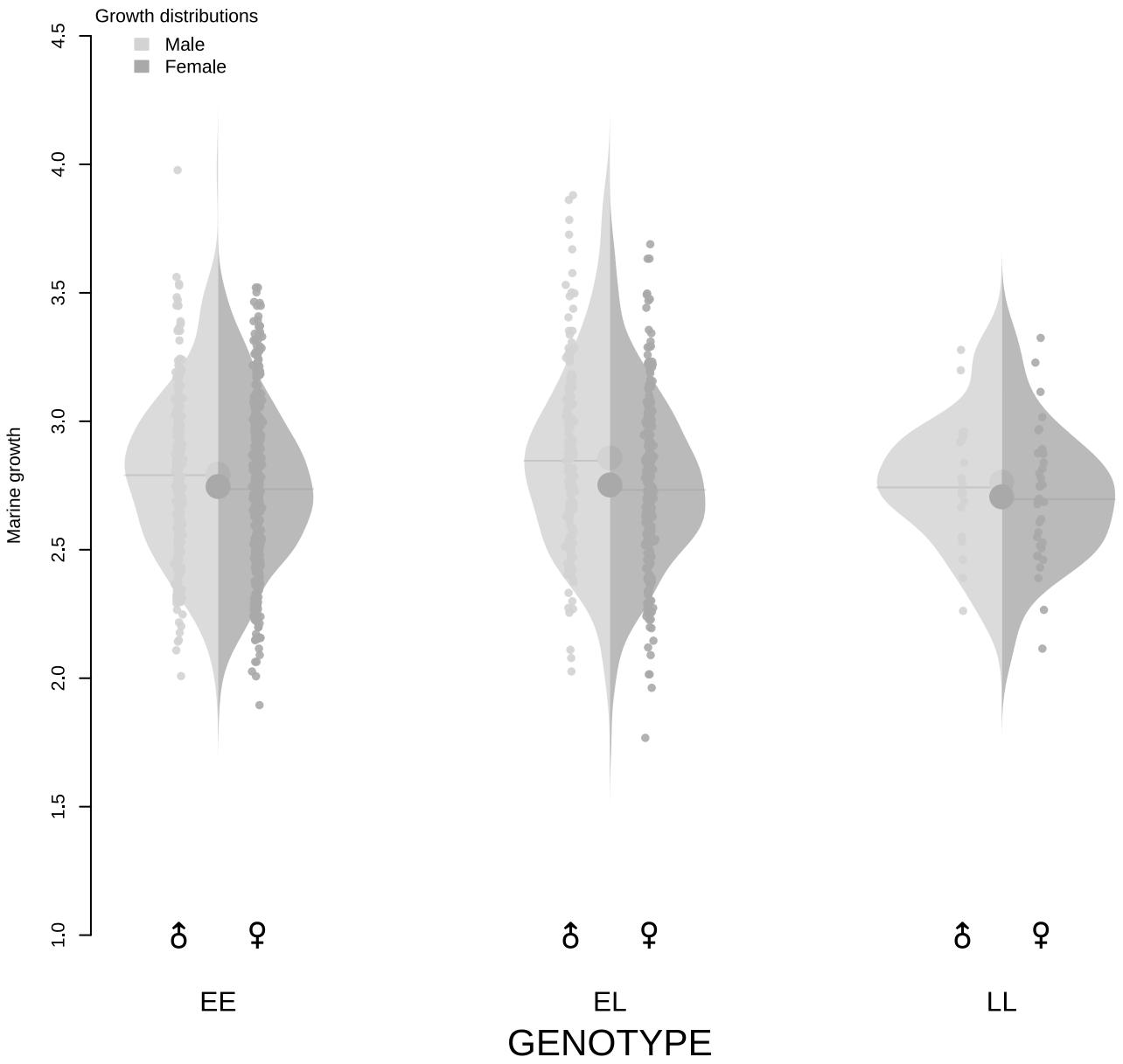
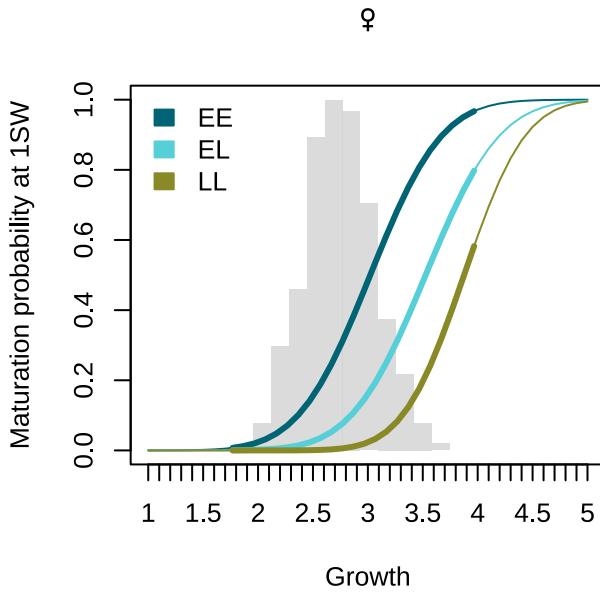
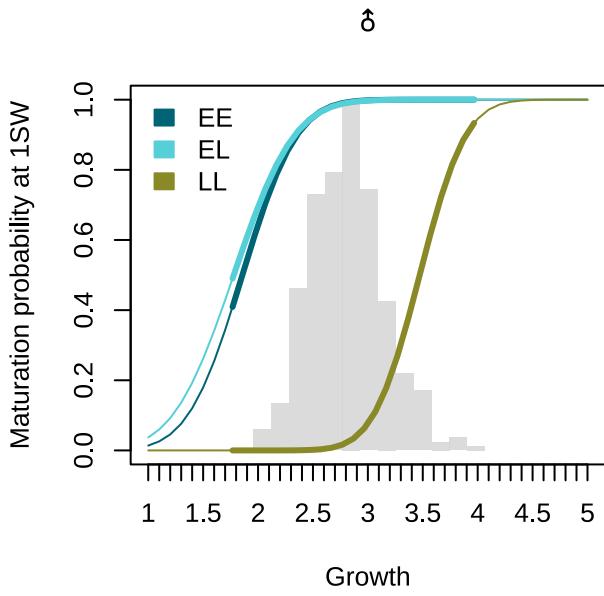
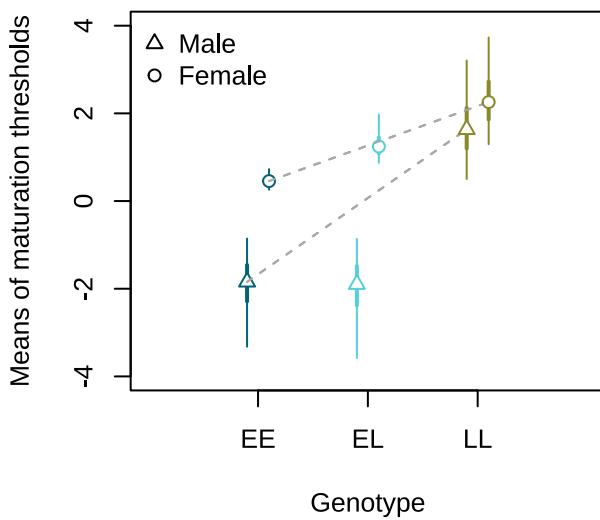
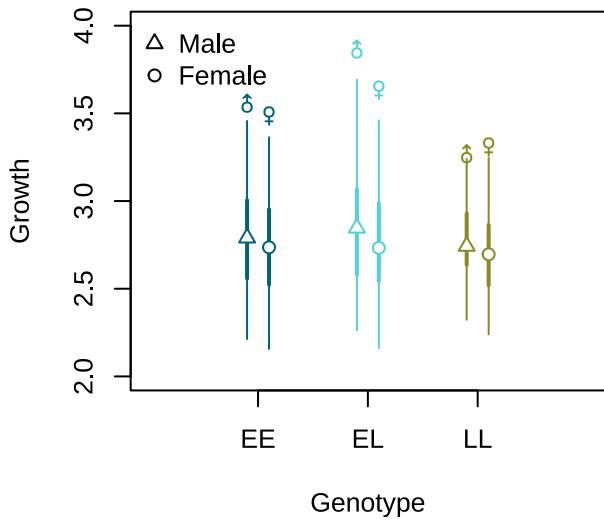
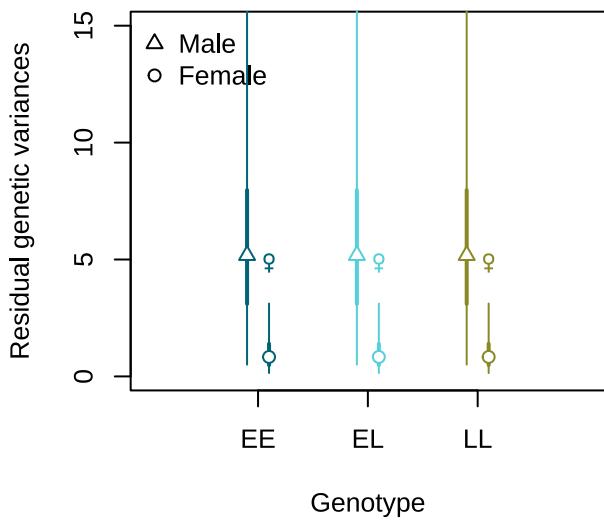
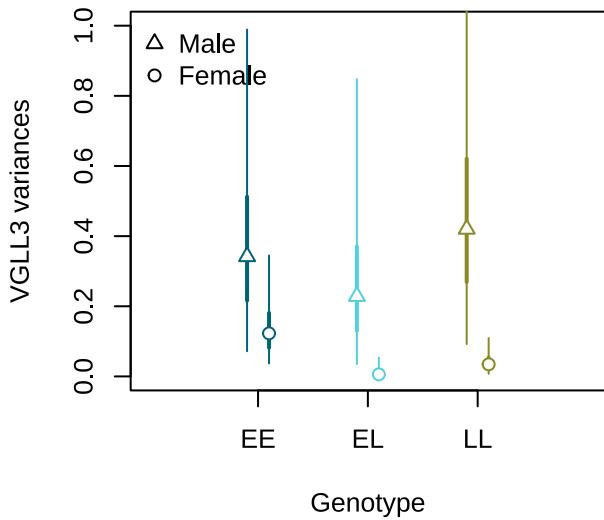


Male**Male****Female****Female**

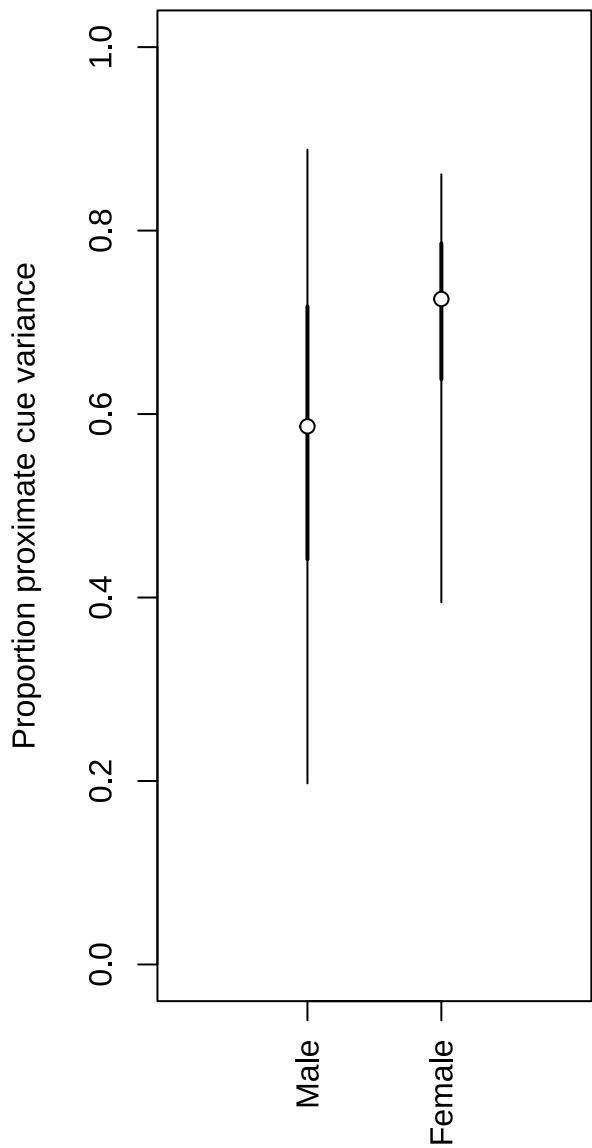




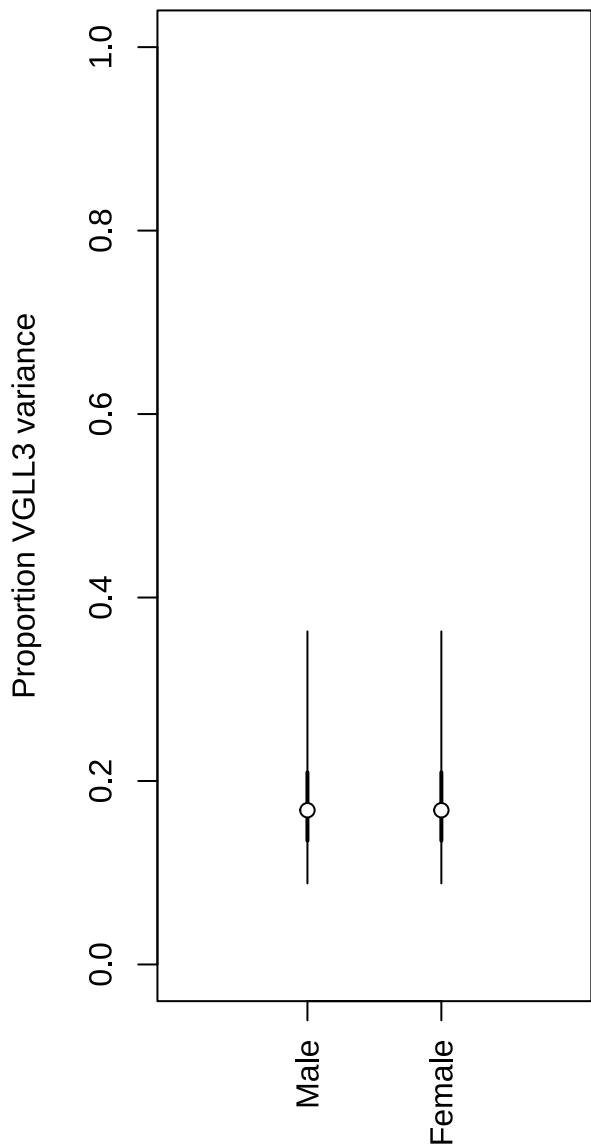




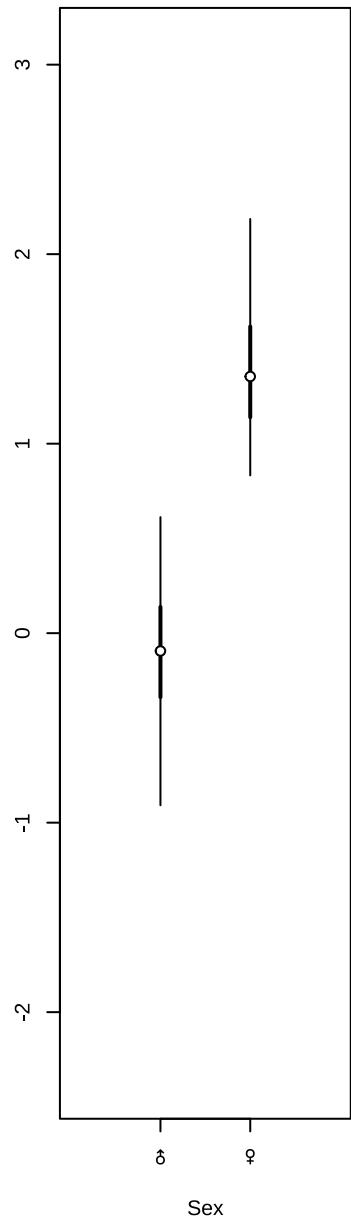
Environment



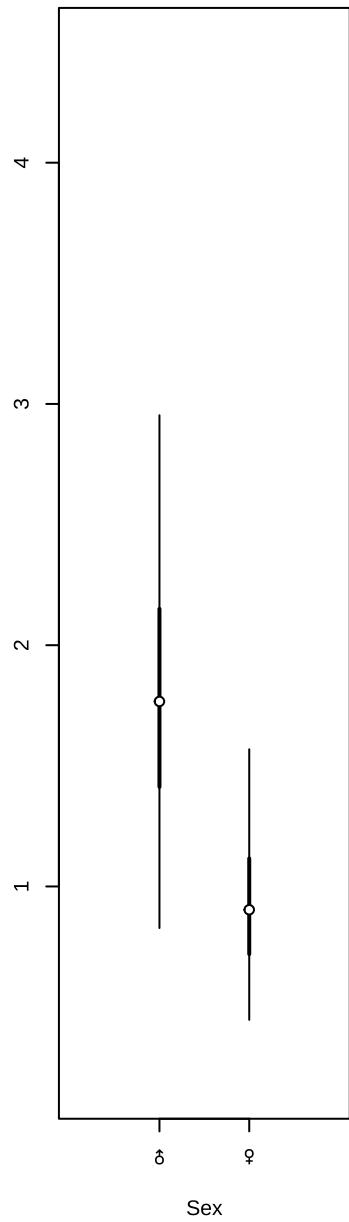
Genetic



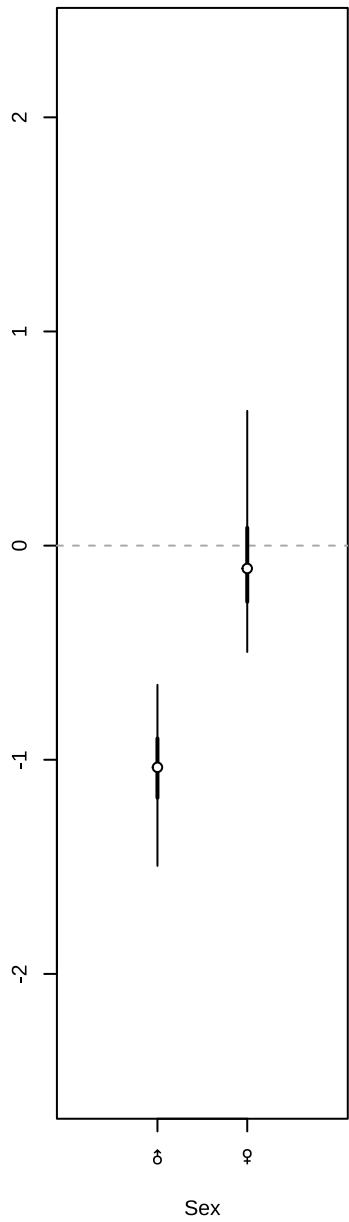
Mean Maturation thresholds

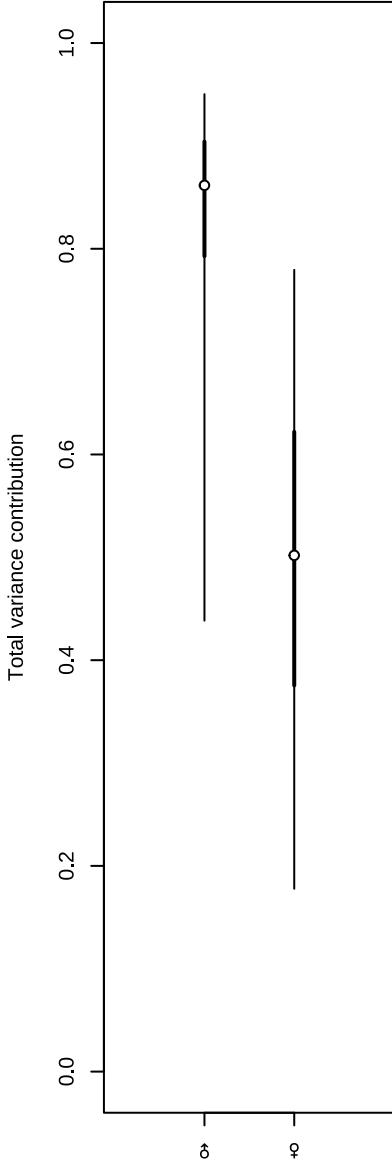


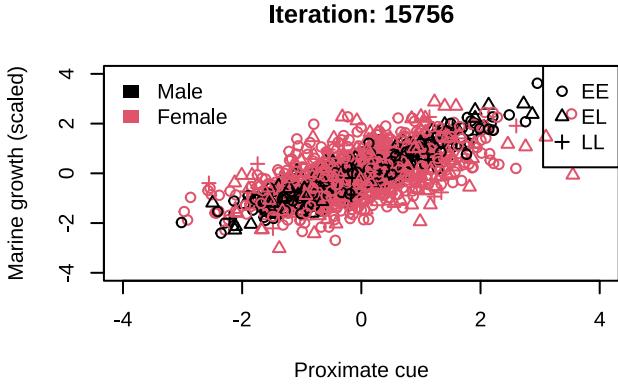
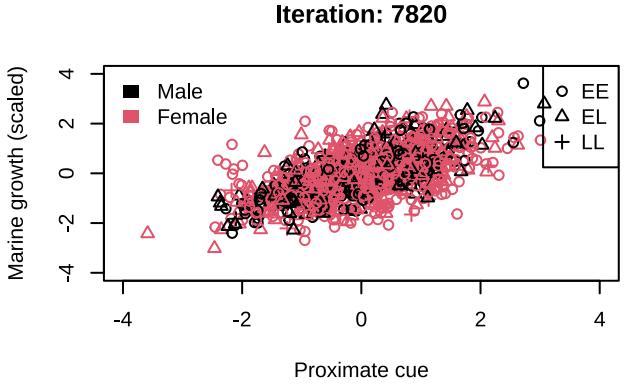
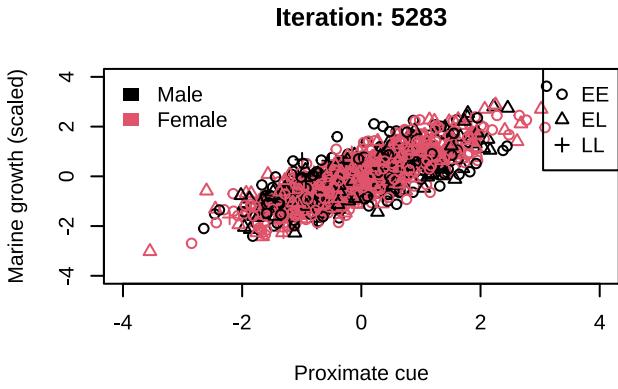
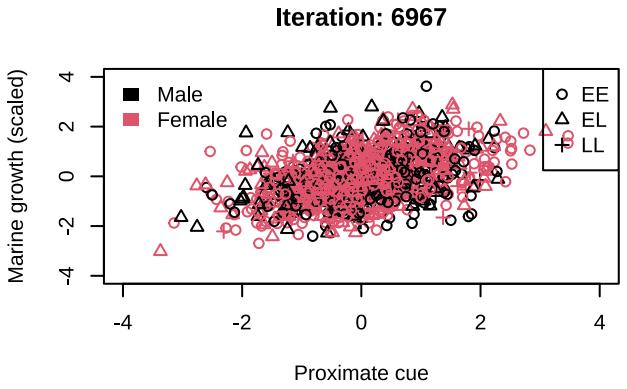
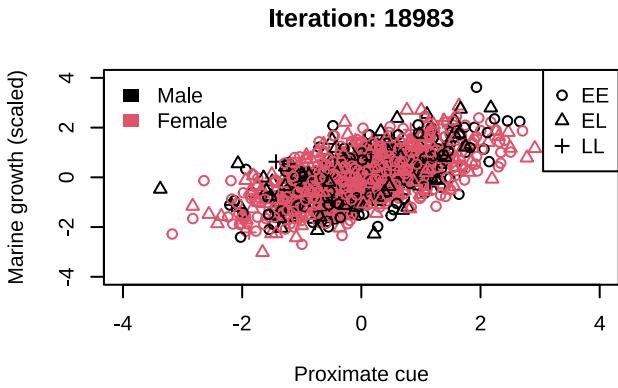
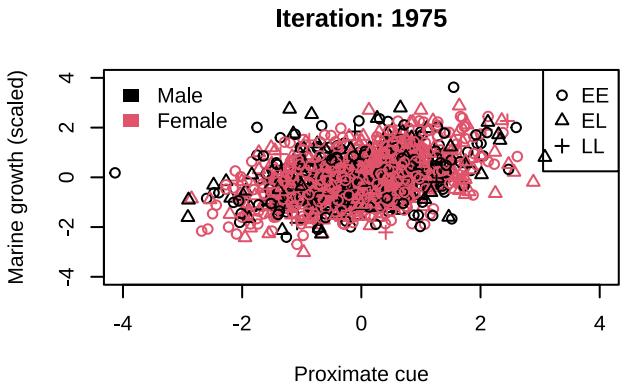
Genotypic value

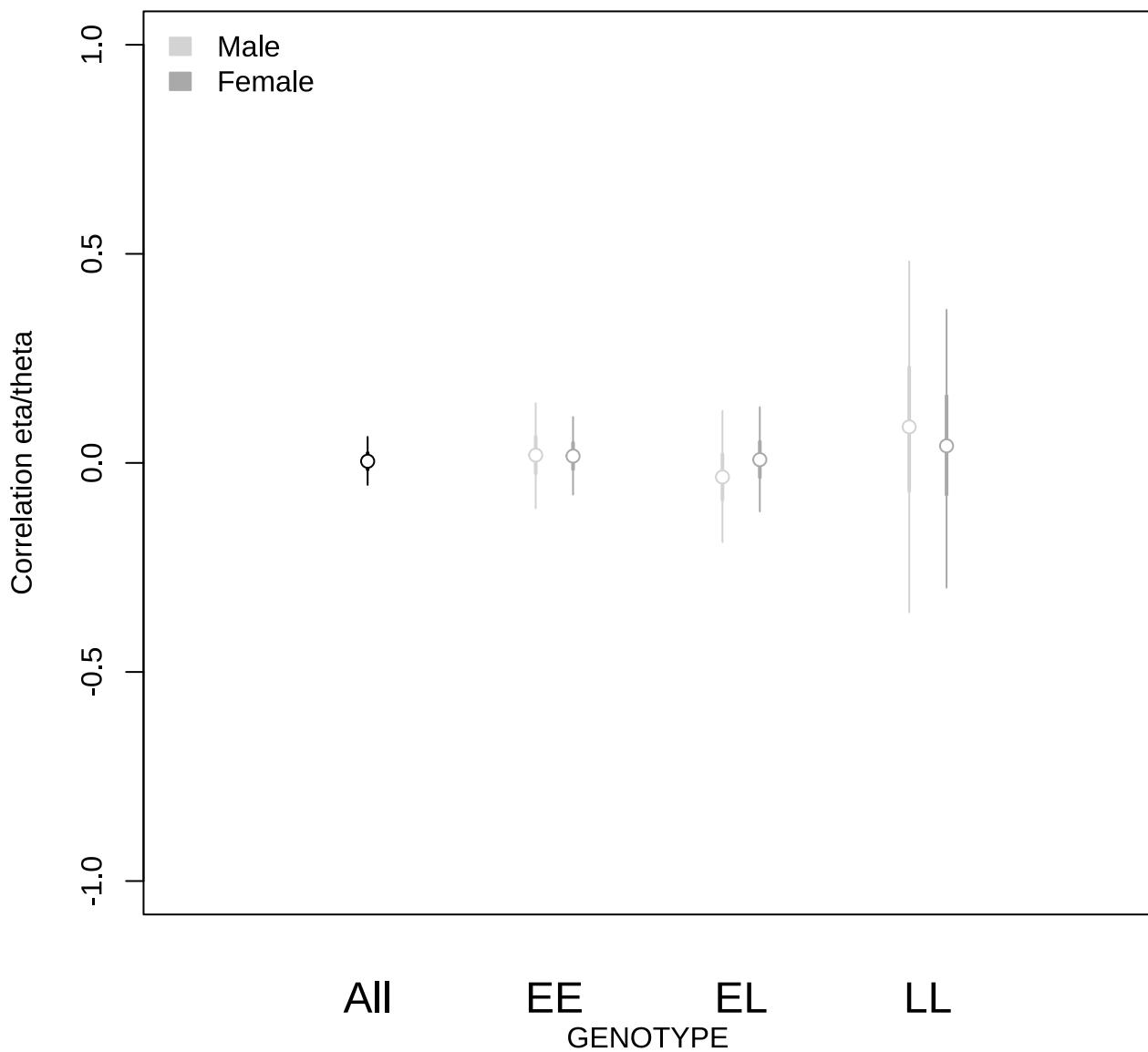


Dominance deviation (scaled)

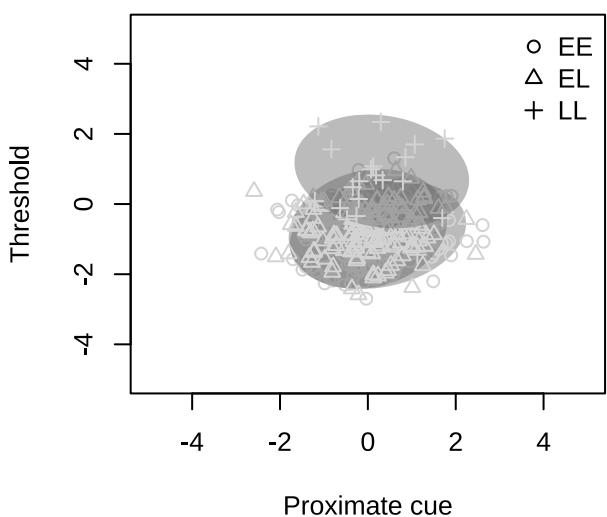




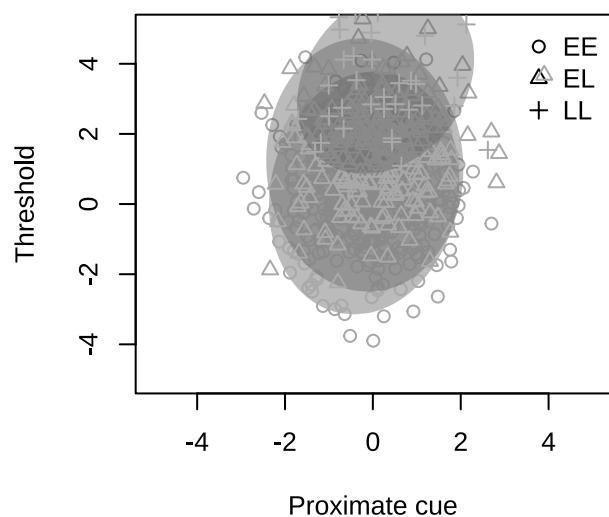




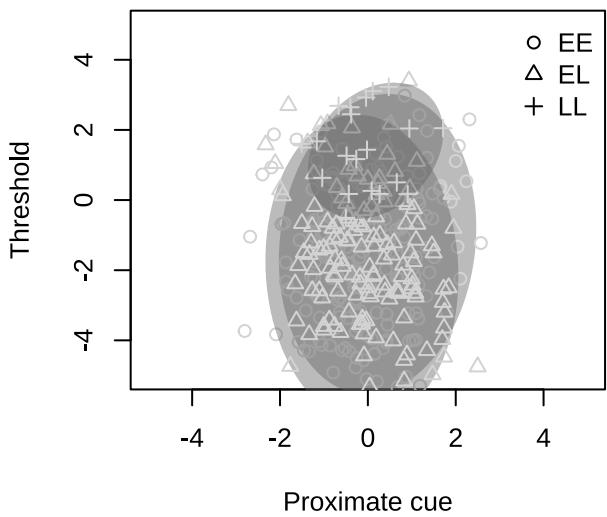
Male (iteration: 15333)



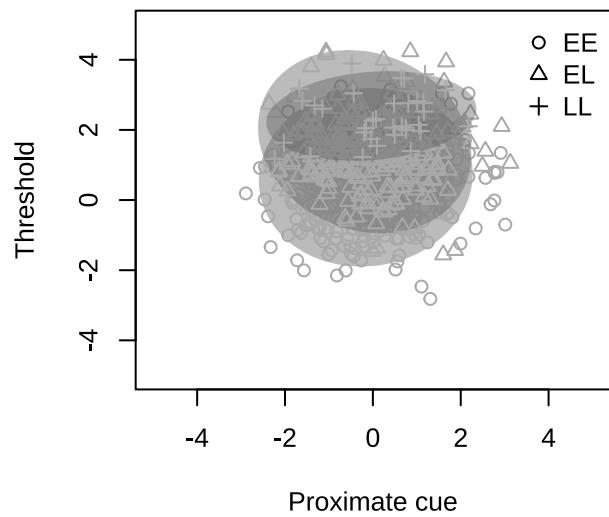
Female (iteration: 15333)



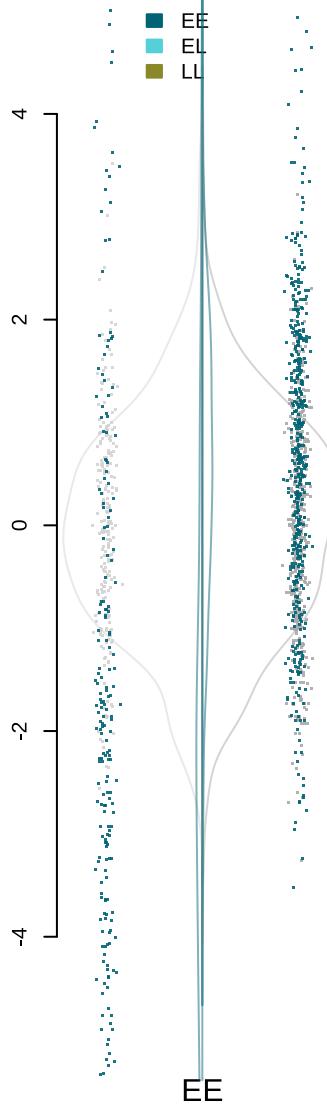
Male (iteration: 2727)



Female (iteration: 2727)

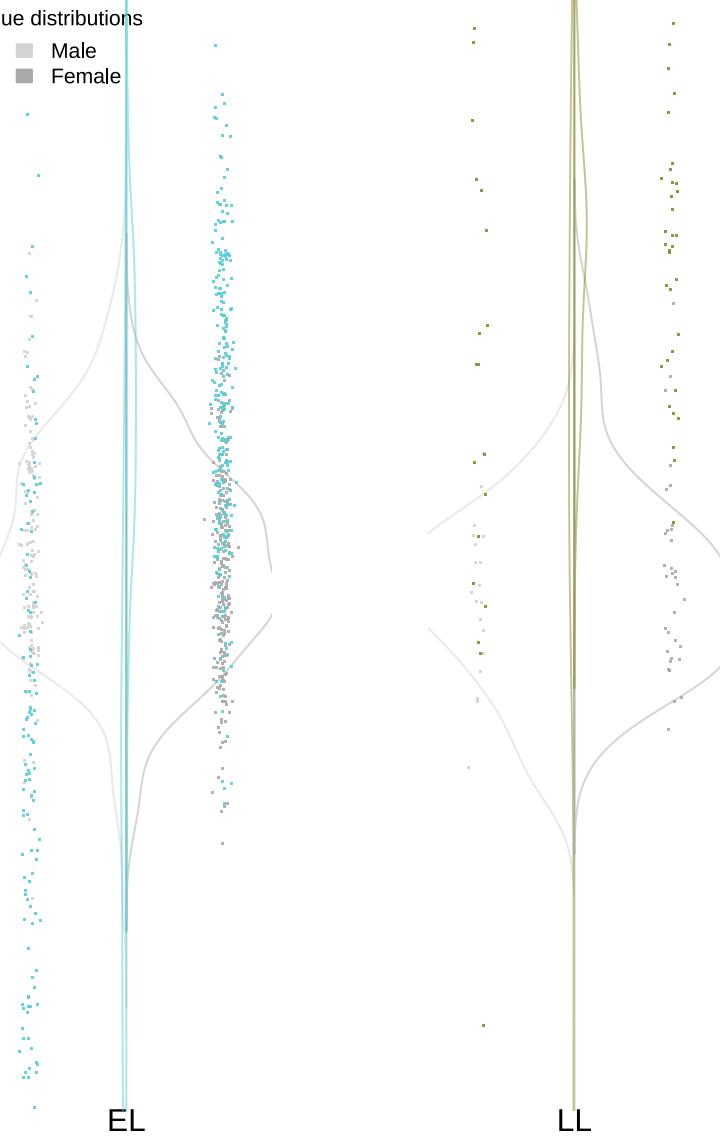


Thresholds distributions

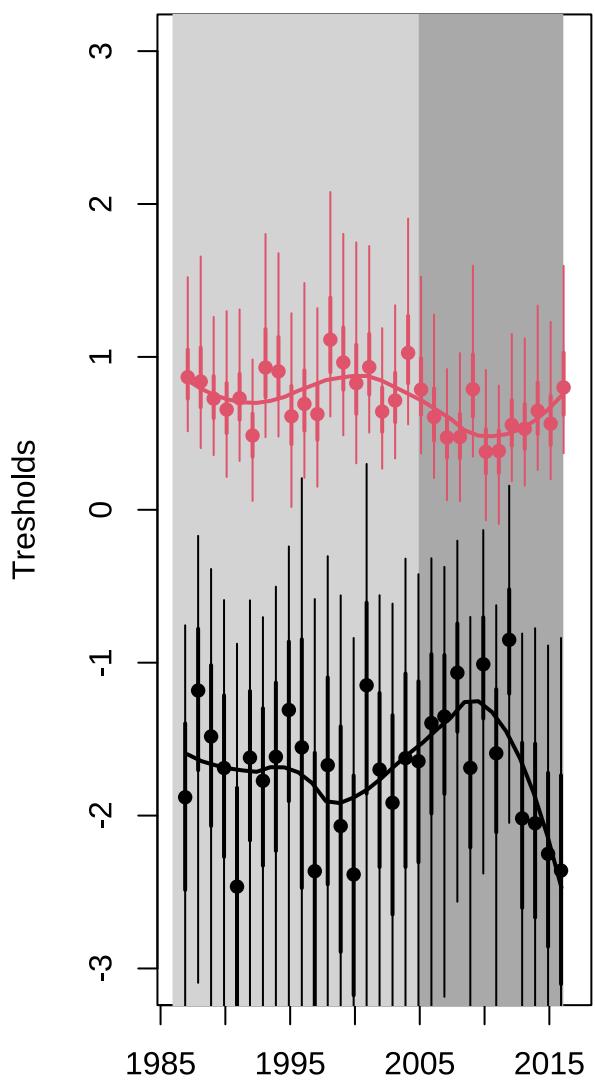
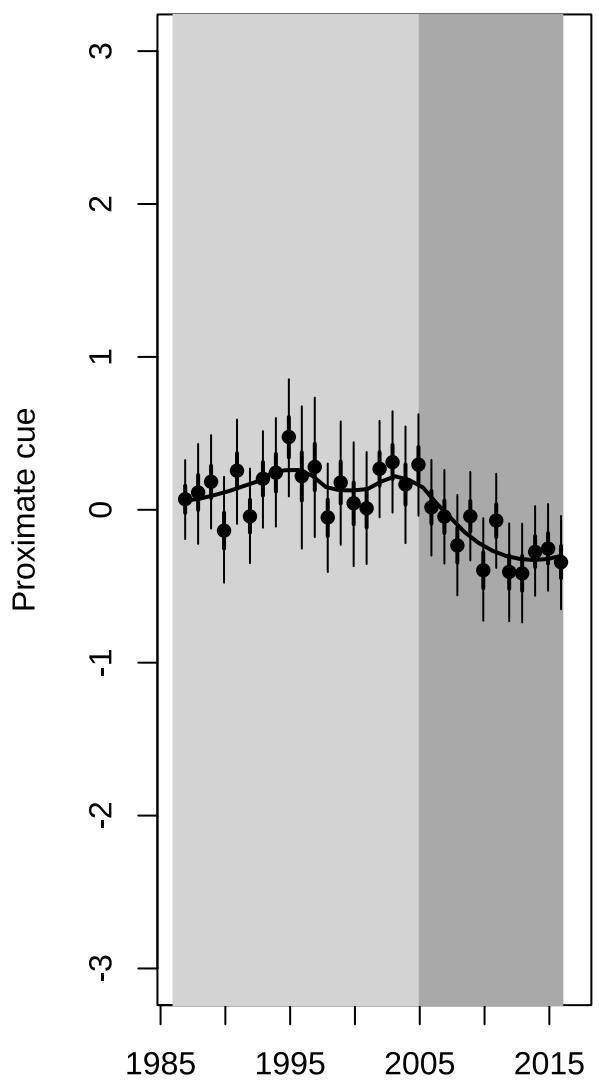


Cue distributions

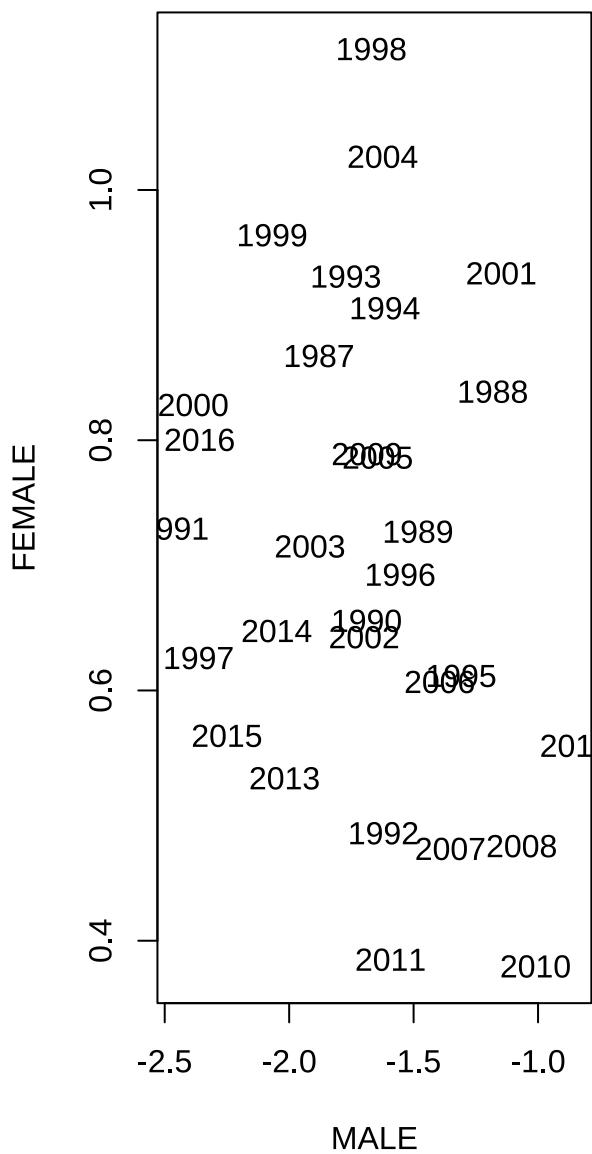
Male
Female

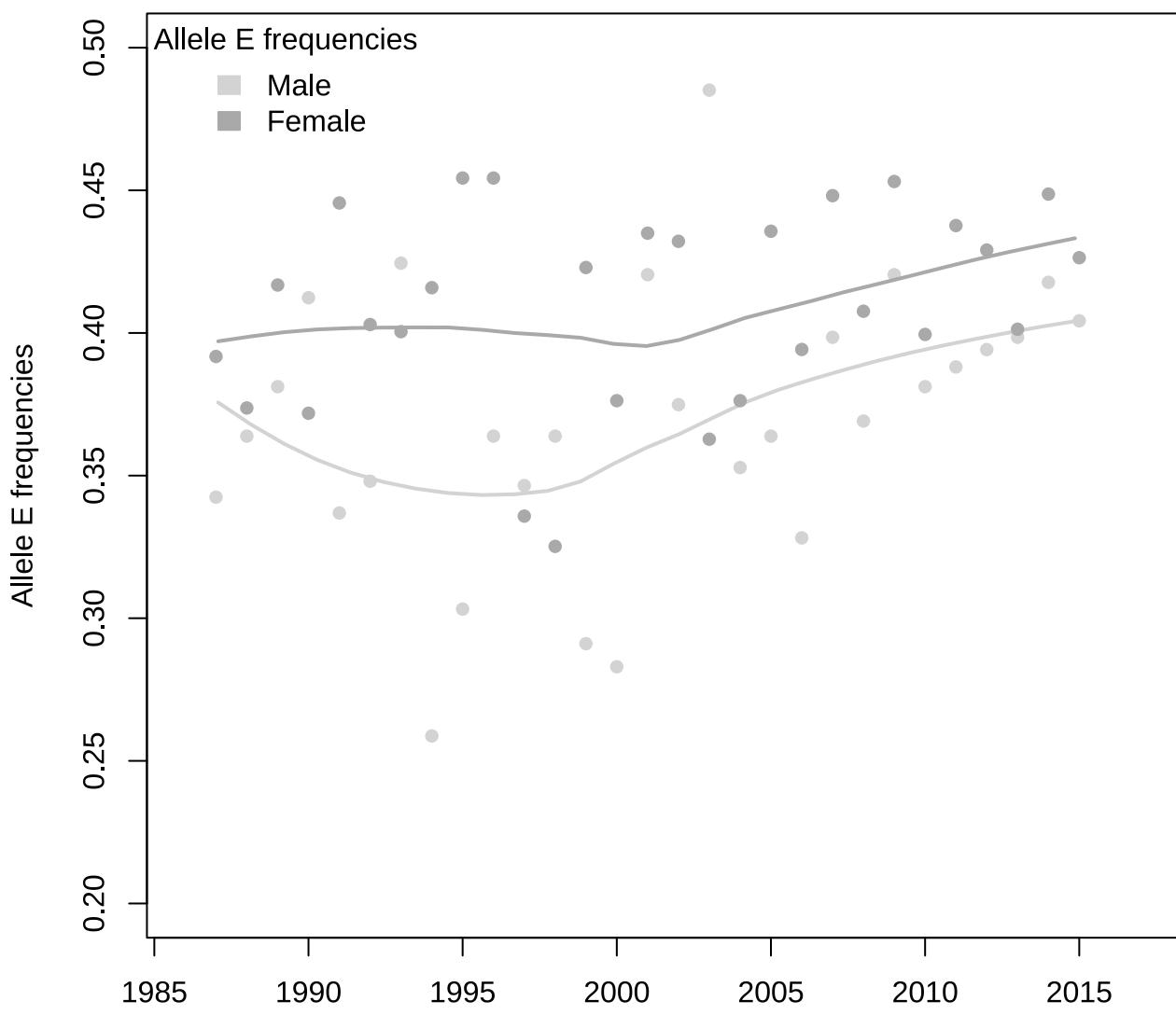


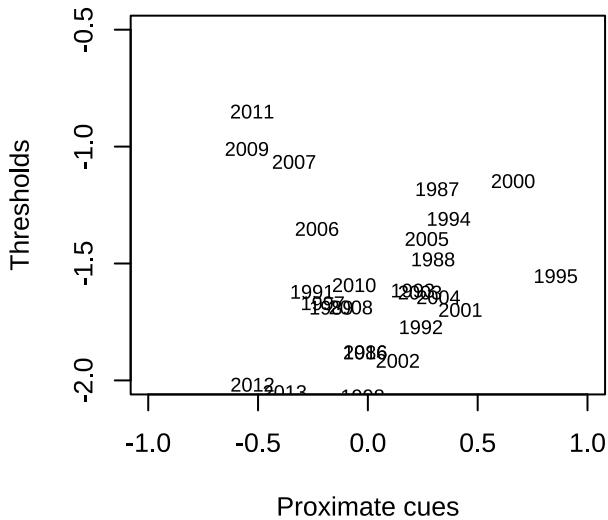
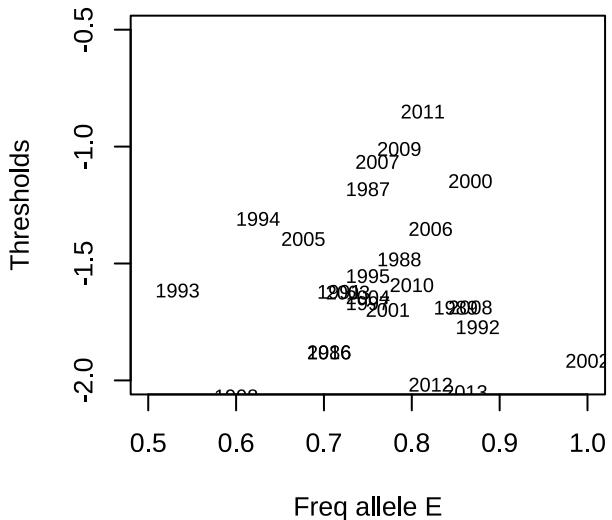
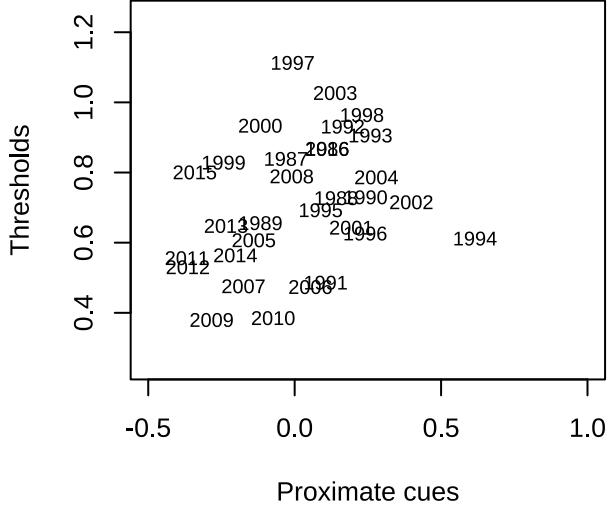
GENOTYPE



Thresholds (medians)





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