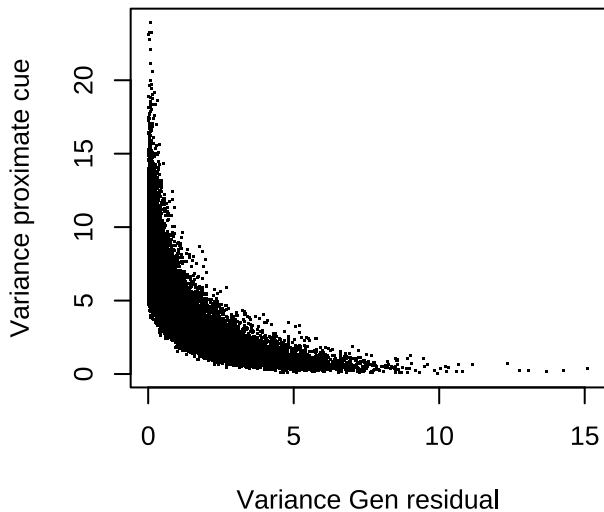
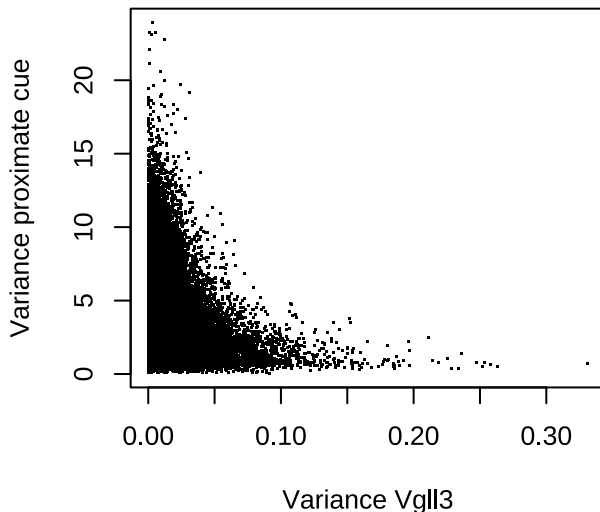


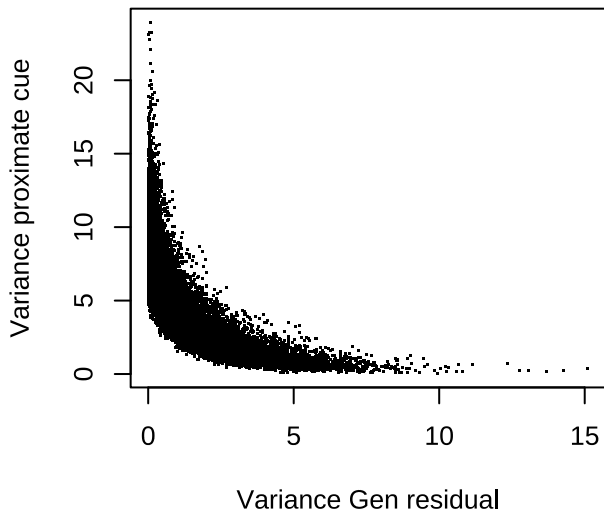
Male



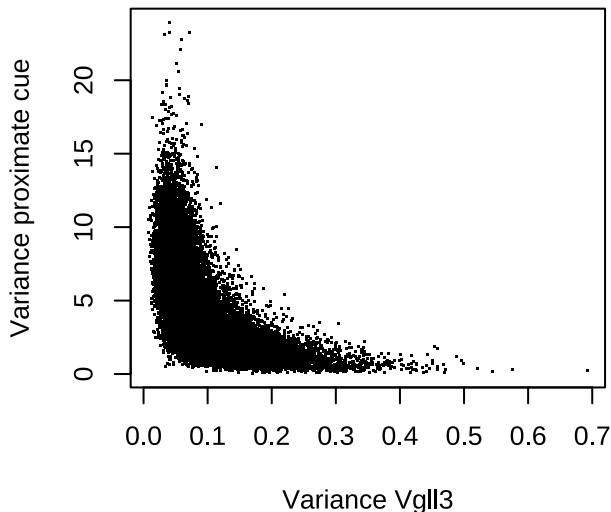
Male

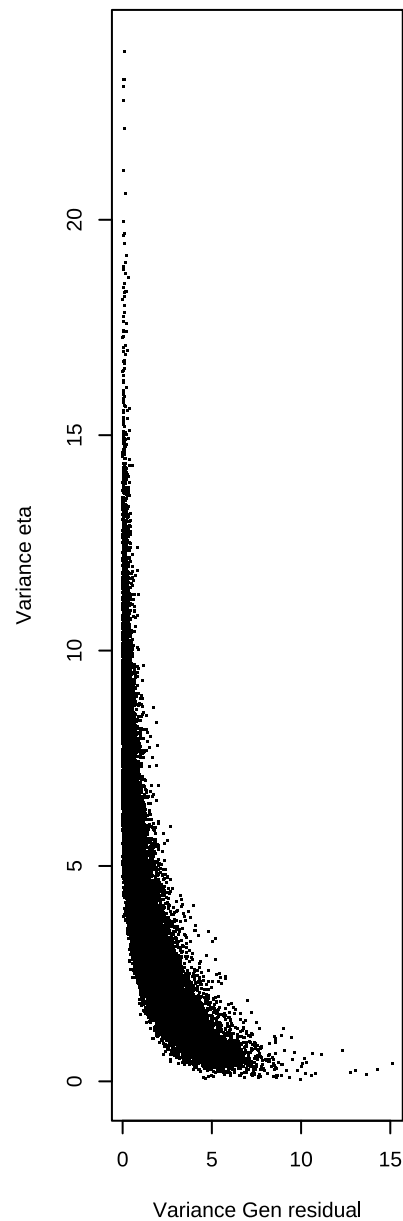
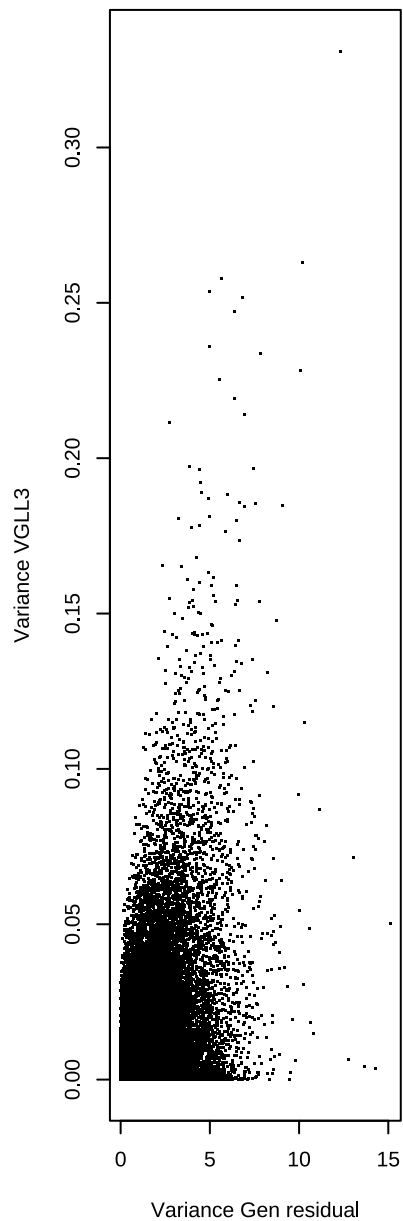
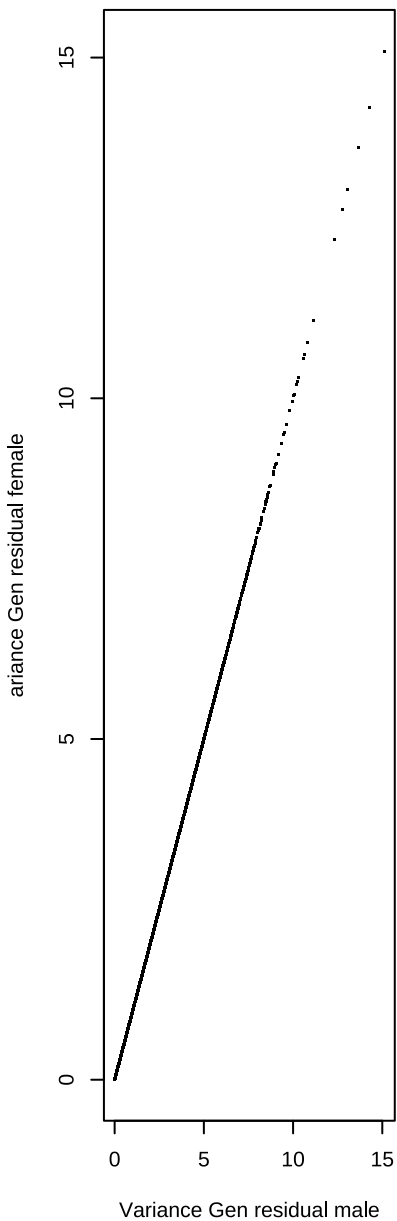


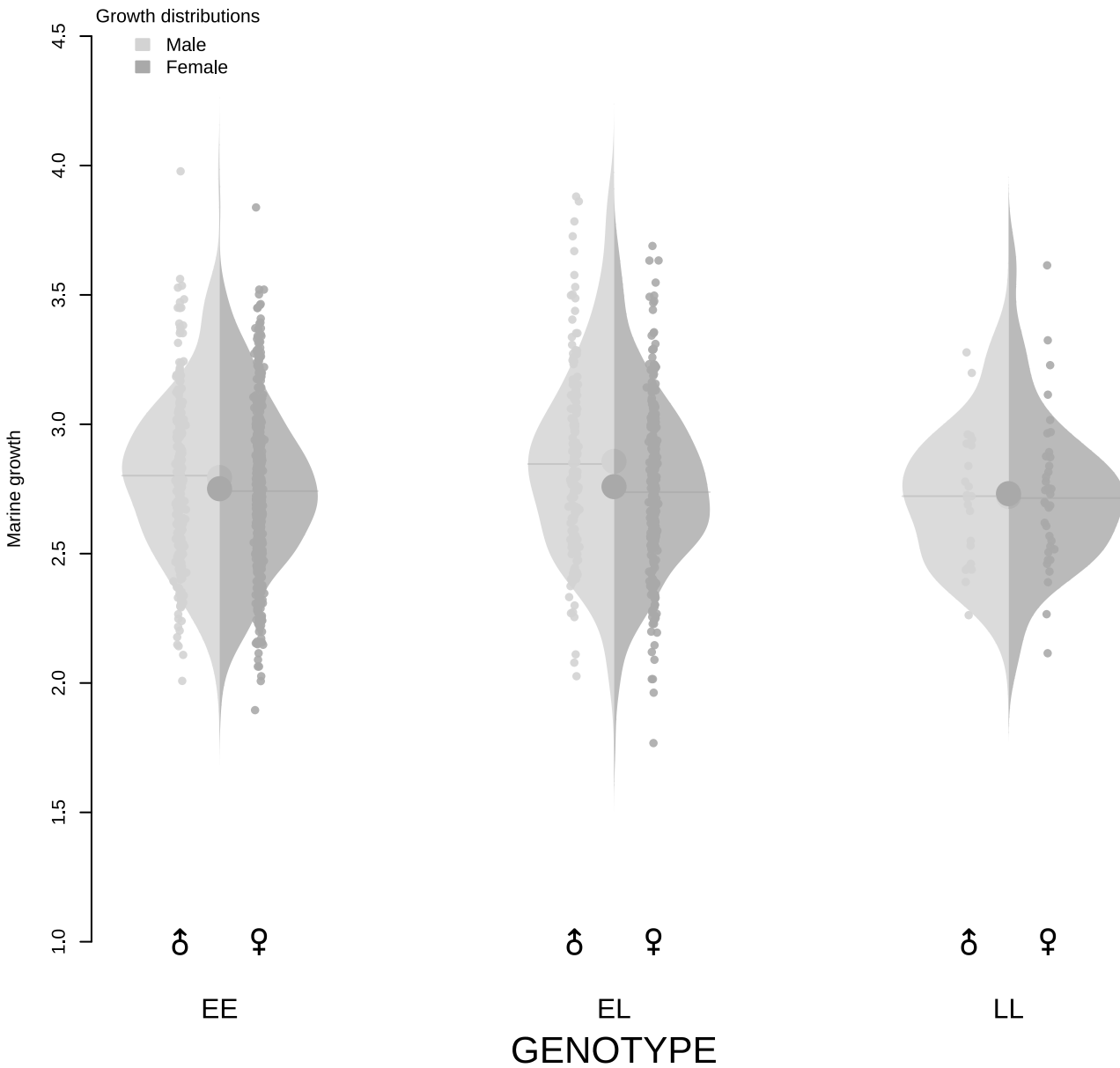
Female

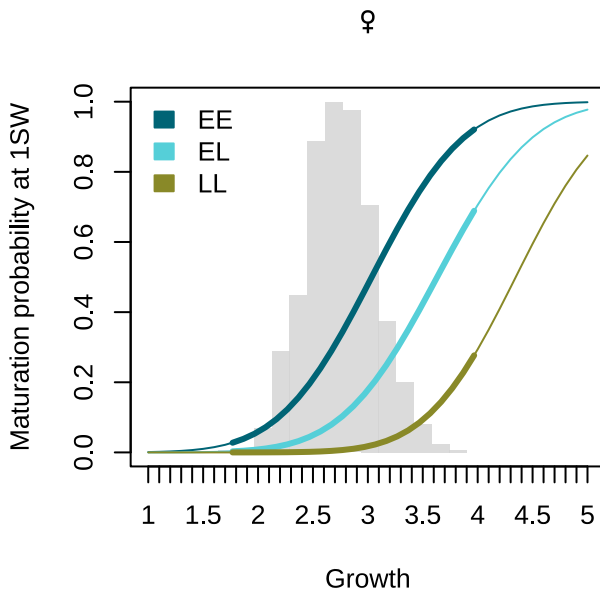
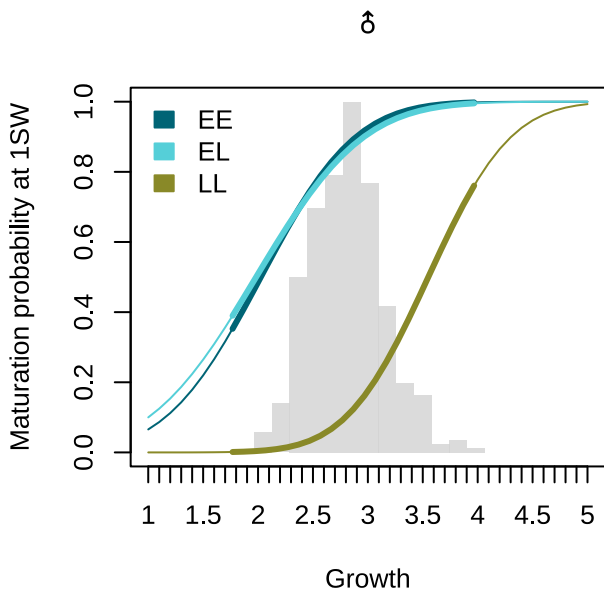
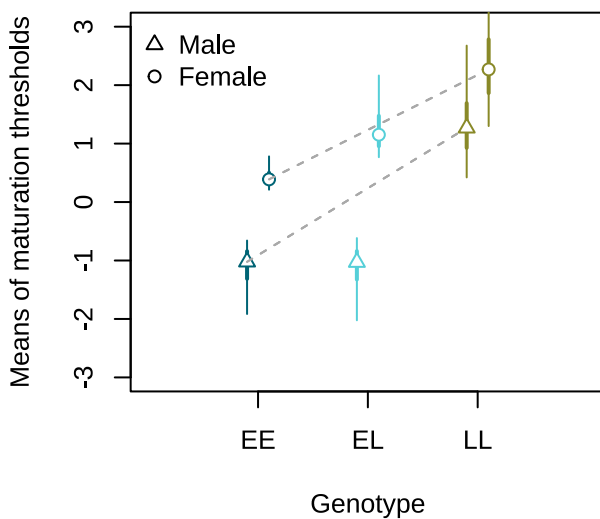
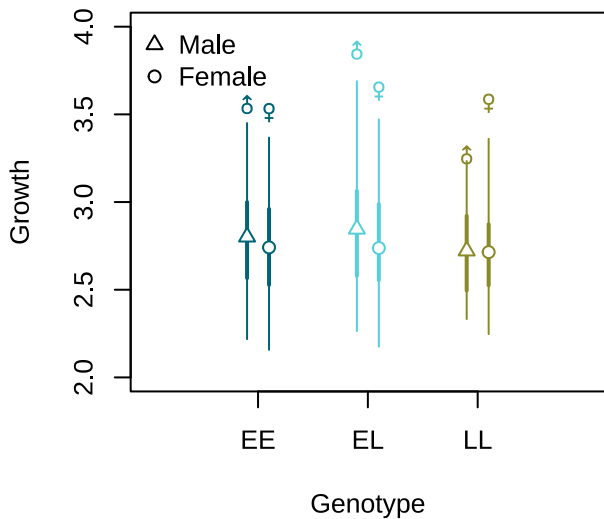


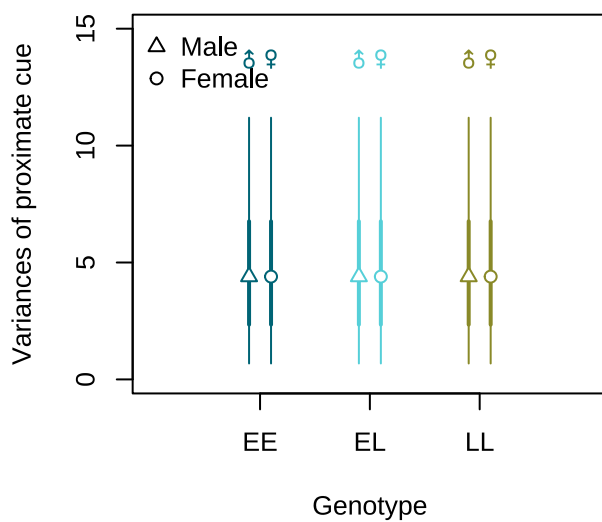
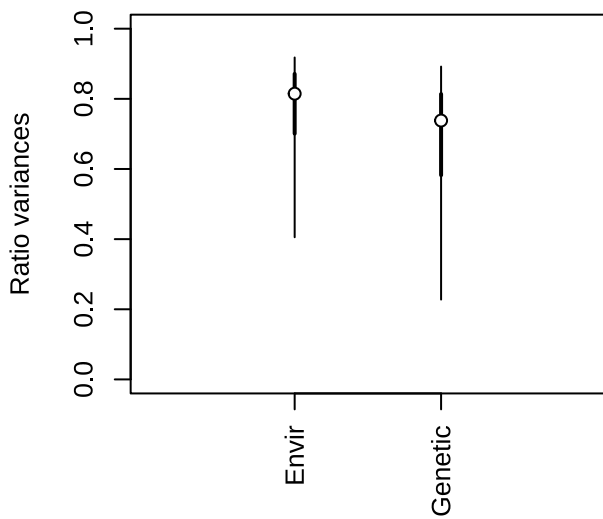
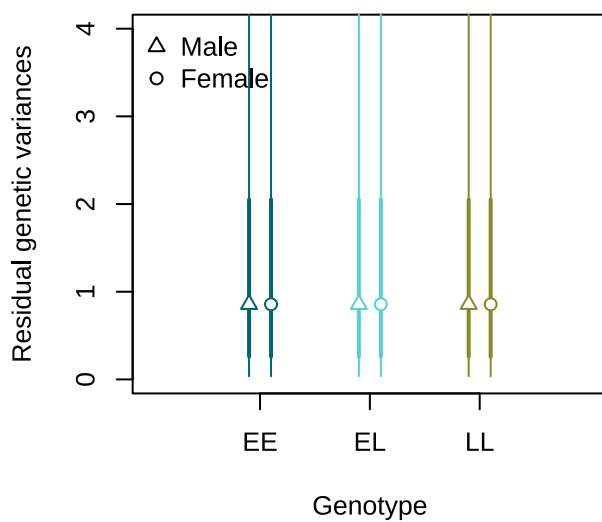
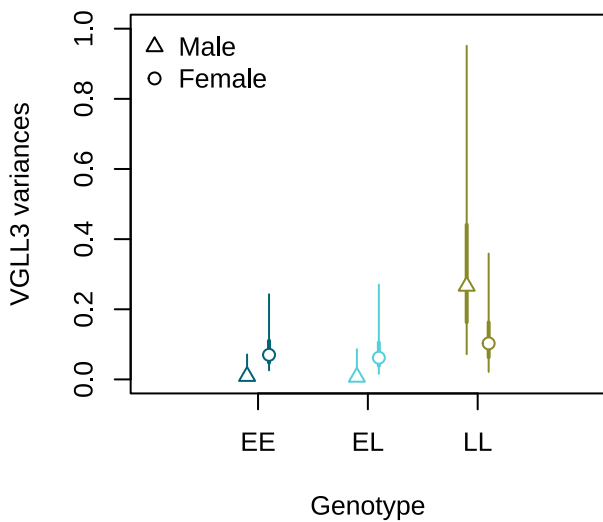
Female



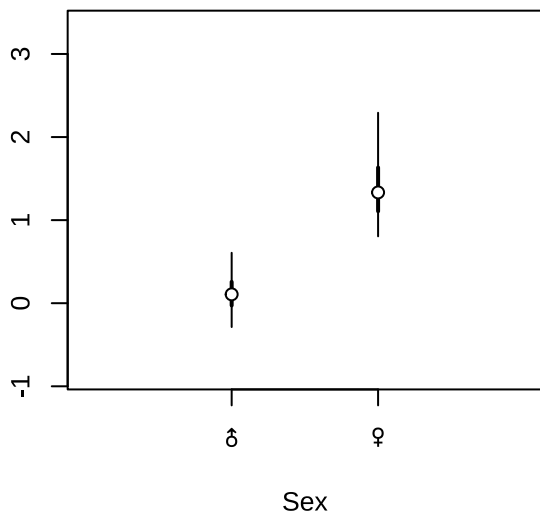




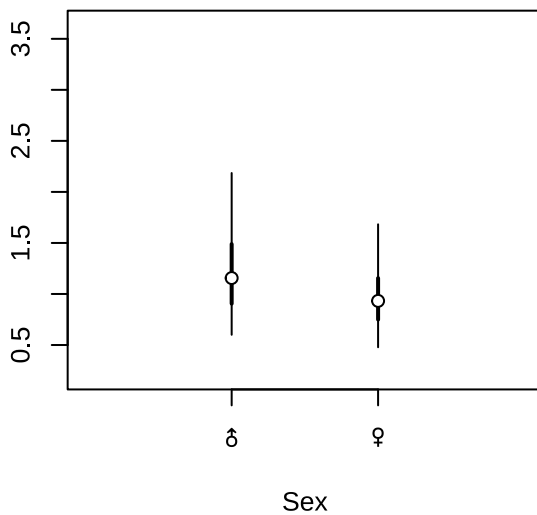




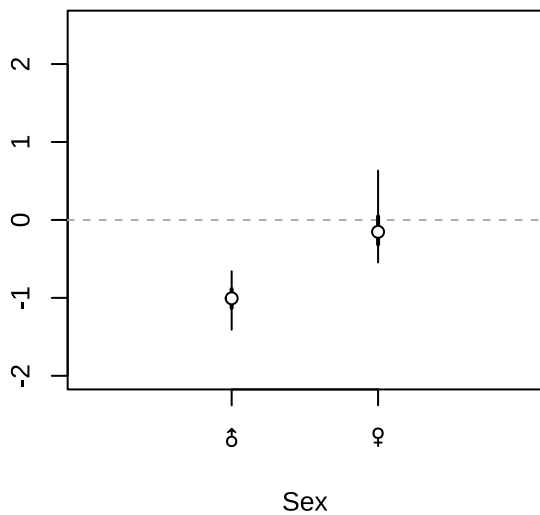
Mean Maturation thresholds



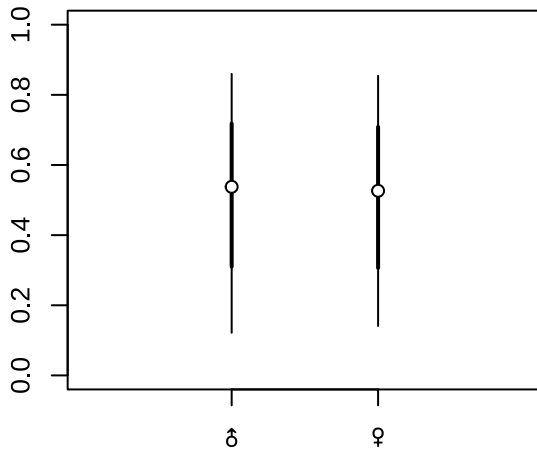
Genotypic value



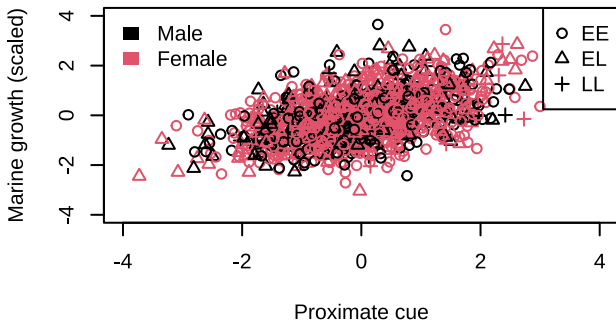
Dominance deviation (scaled)



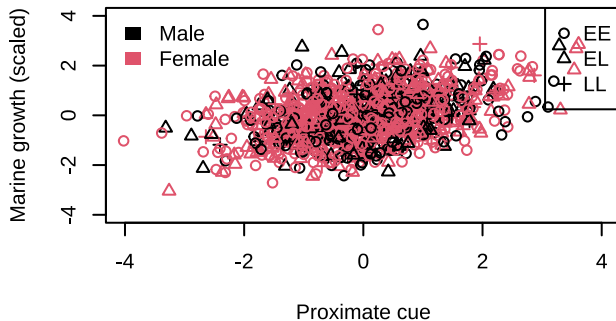
Total variance contribution



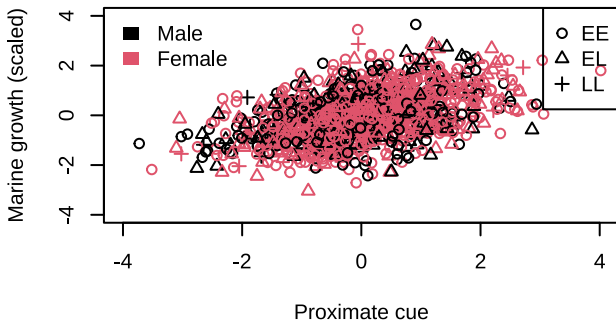
Iteration: 17374



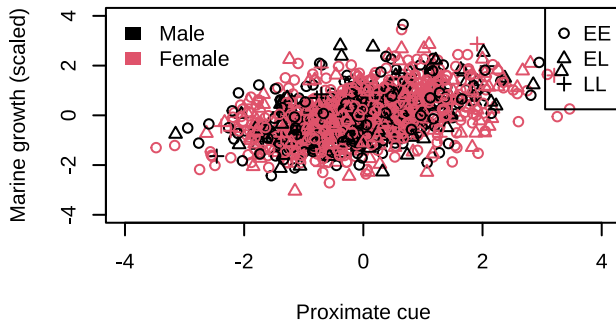
Iteration: 25375



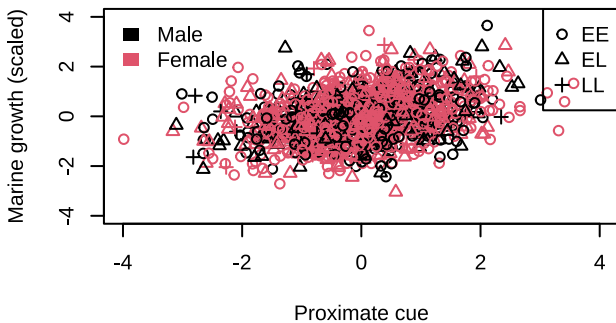
Iteration: 7928



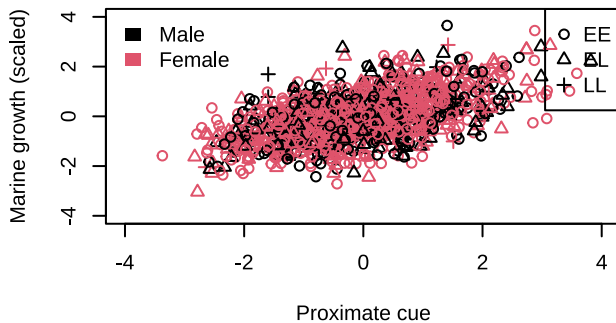
Iteration: 4338

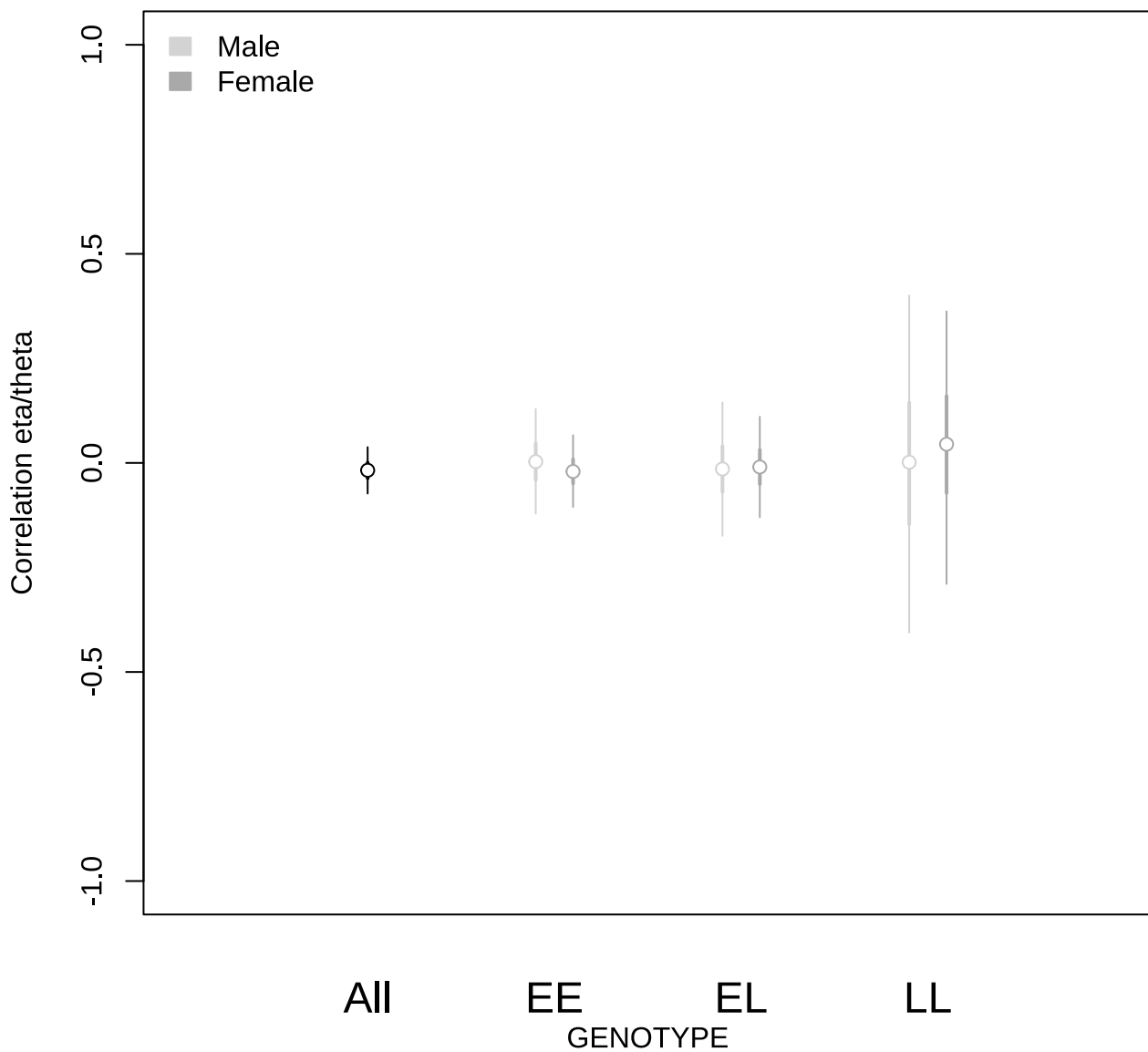


Iteration: 289

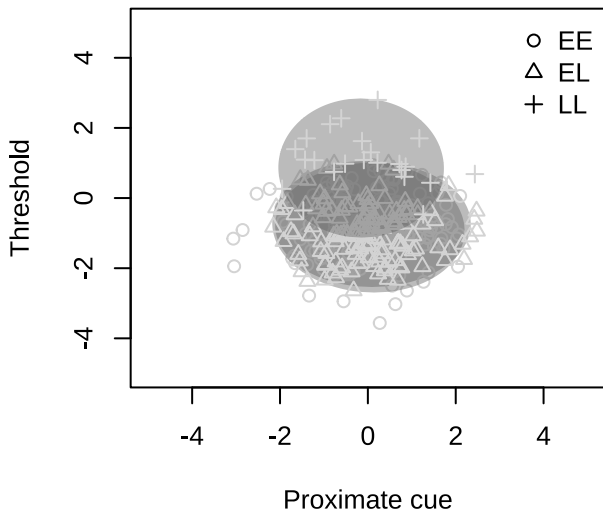


Iteration: 21996

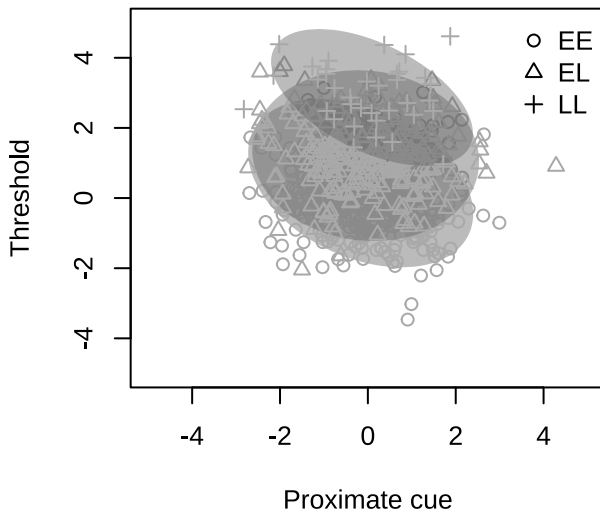




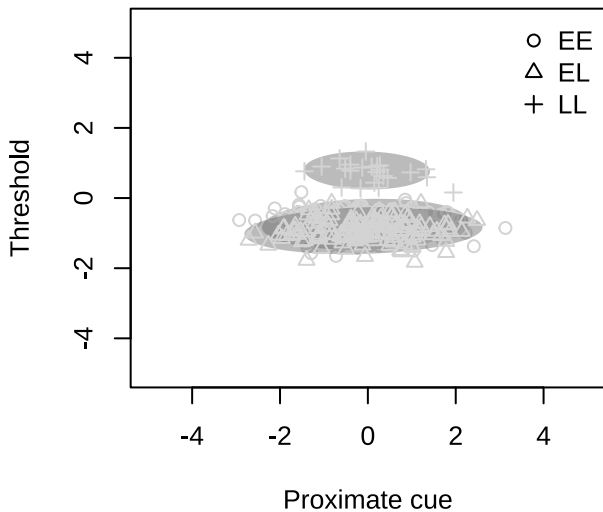
Male (iteration: 17561)



Female (iteration: 17561)



Male (iteration: 13797)



Female (iteration: 13797)

