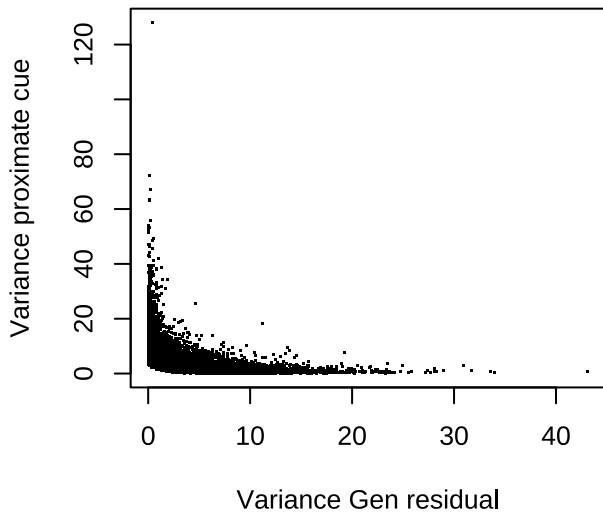
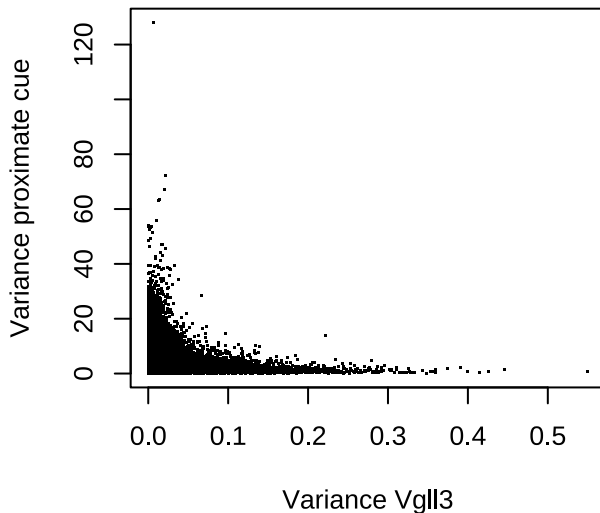


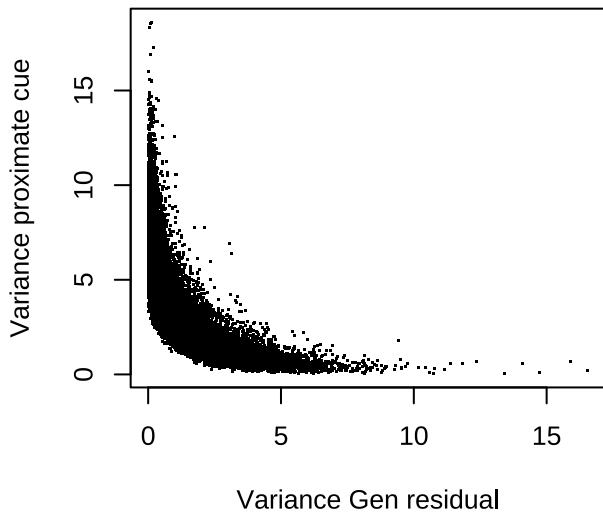
Male



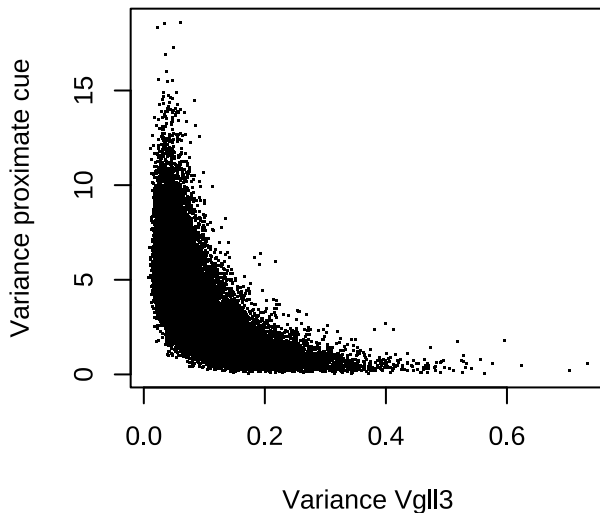
Male

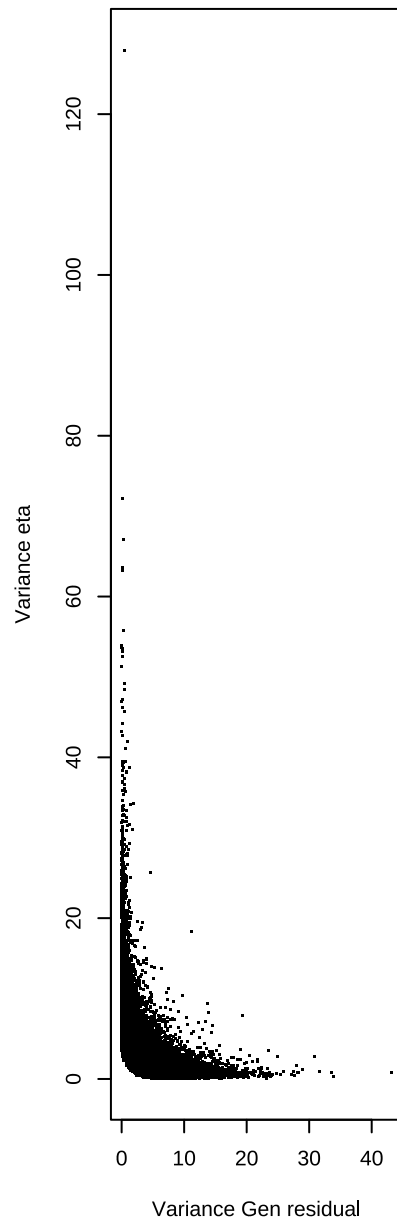
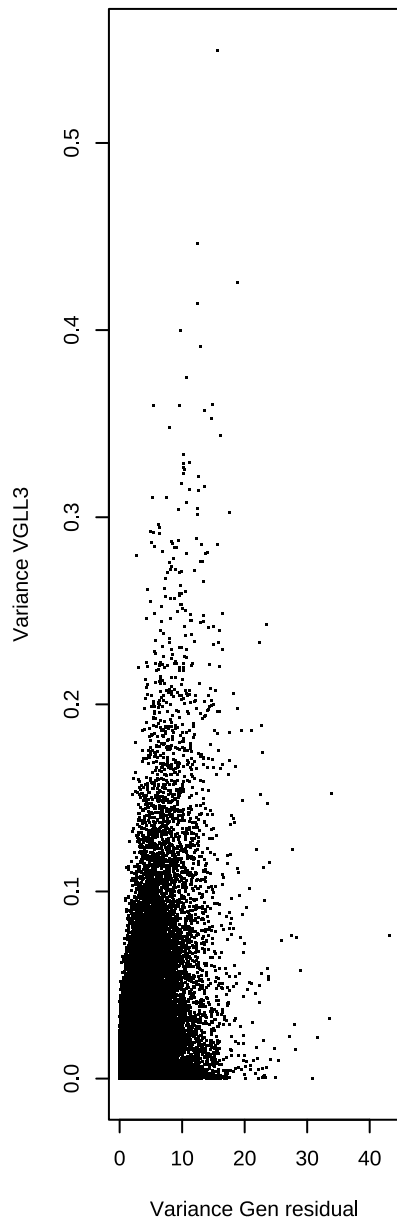
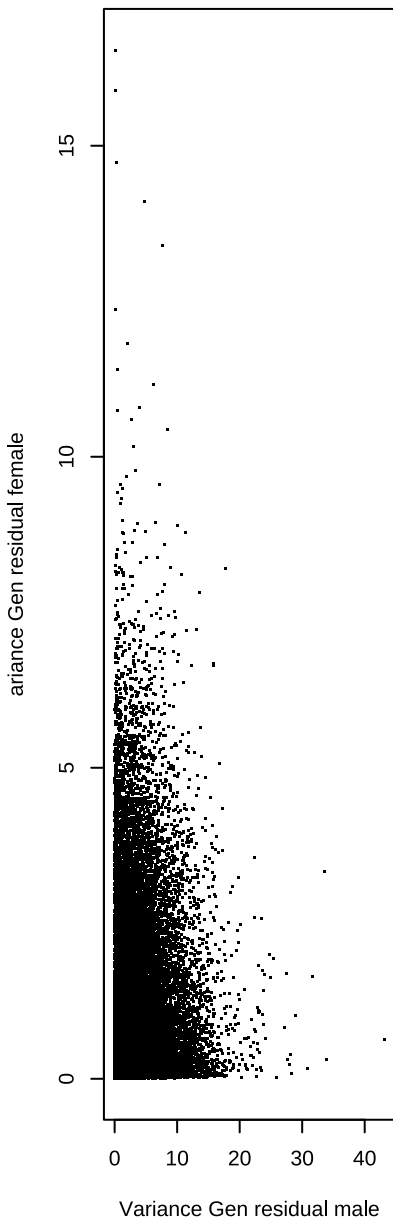


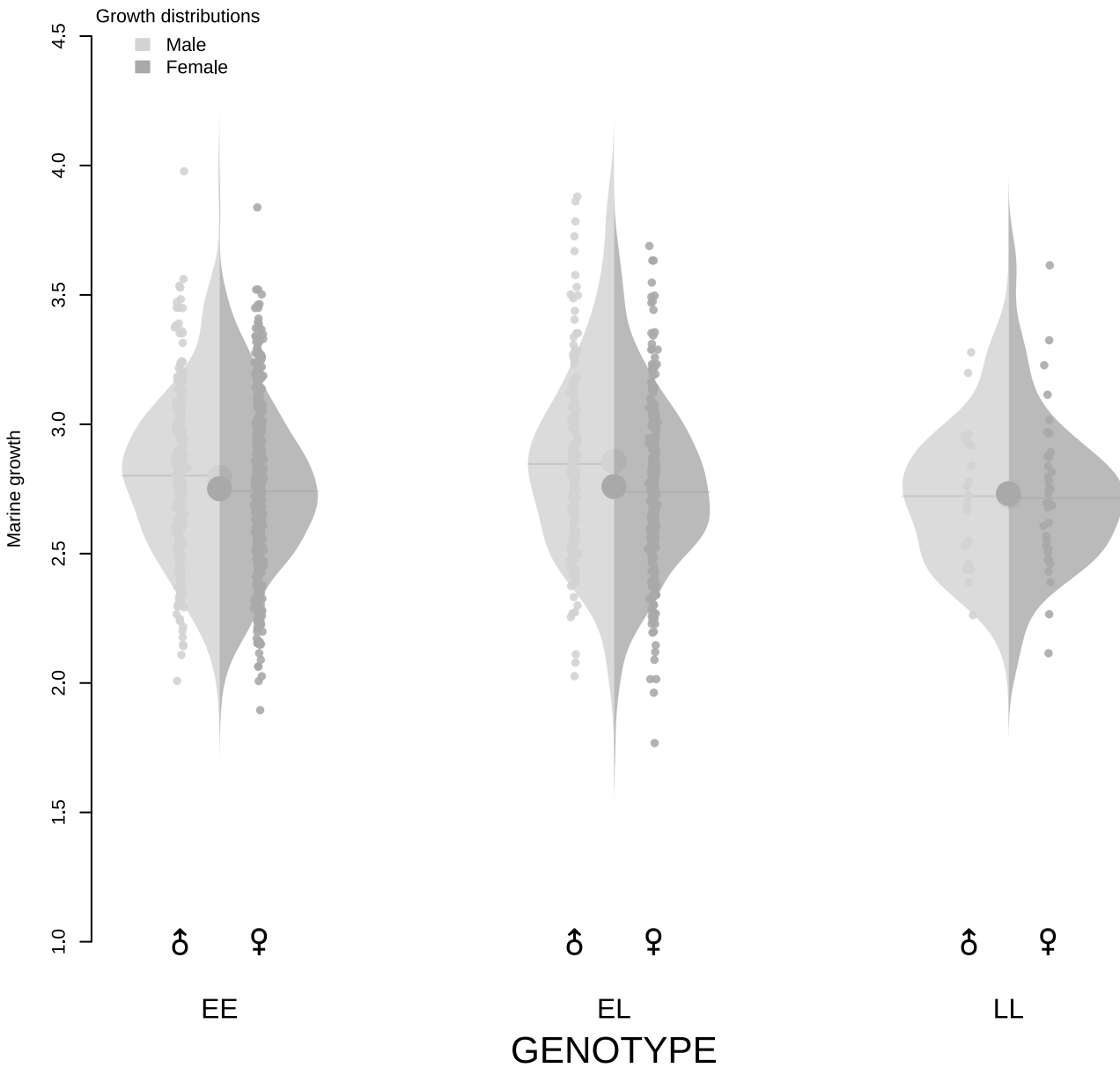
Female

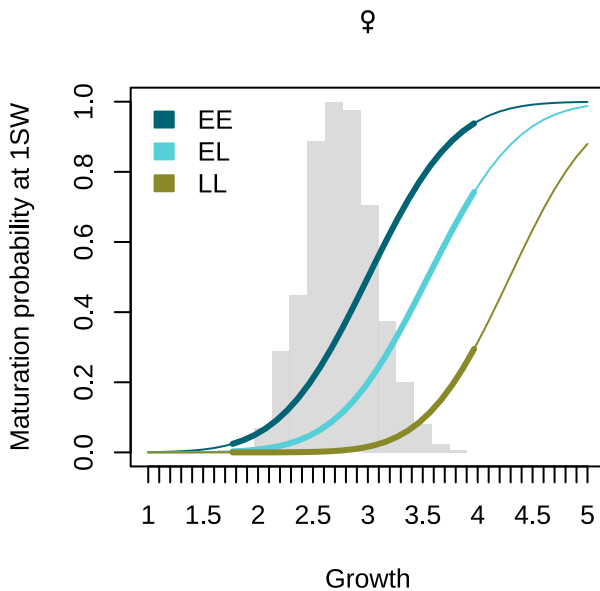
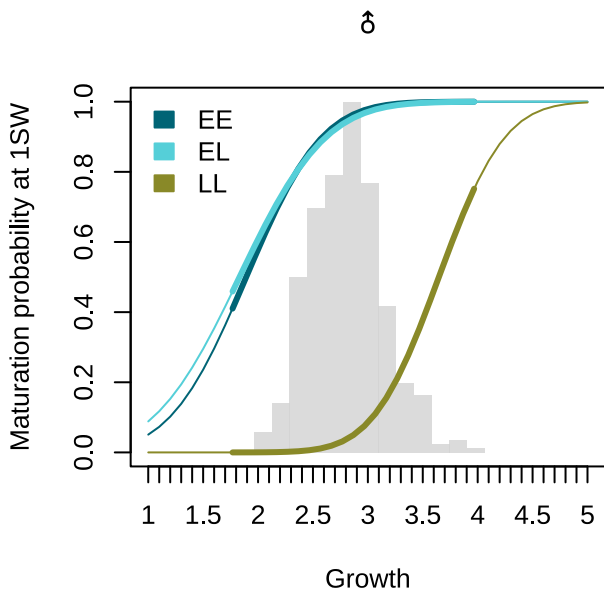
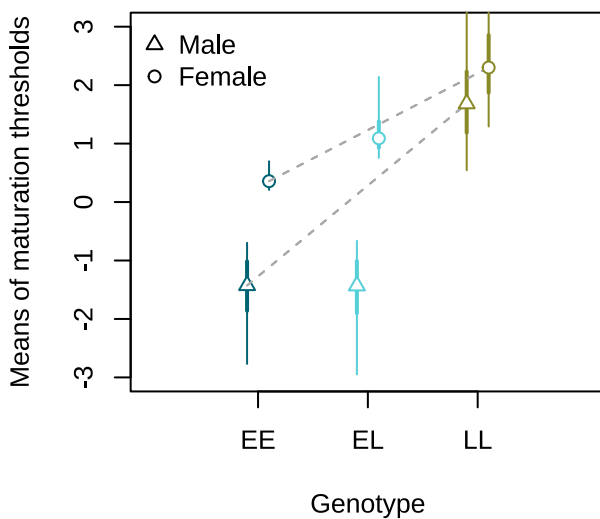
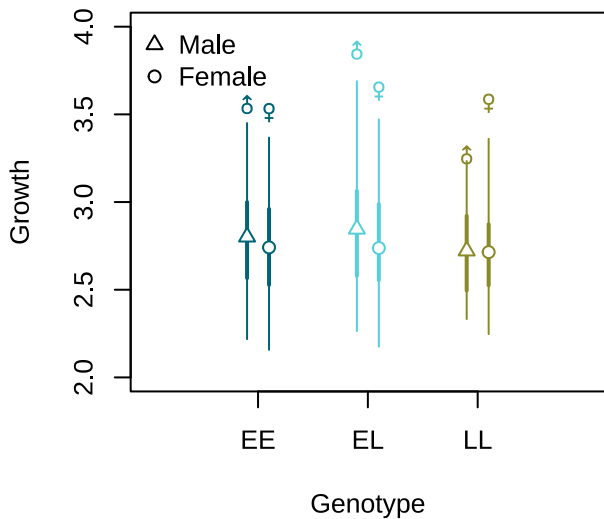


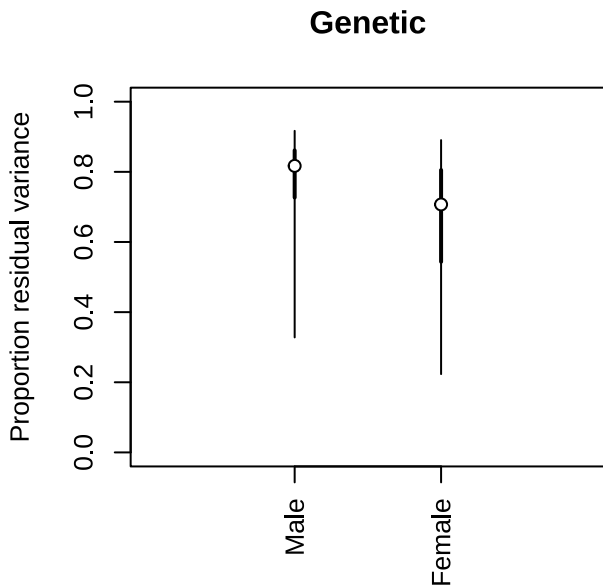
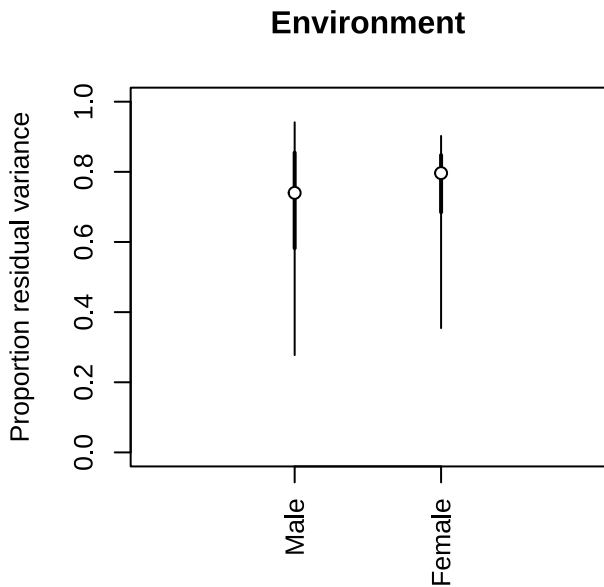
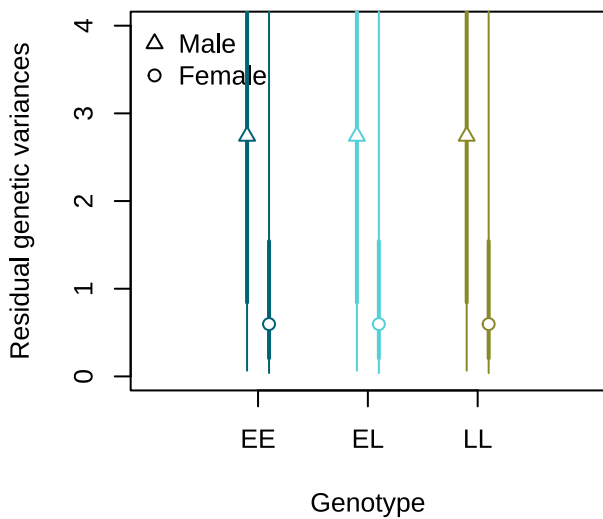
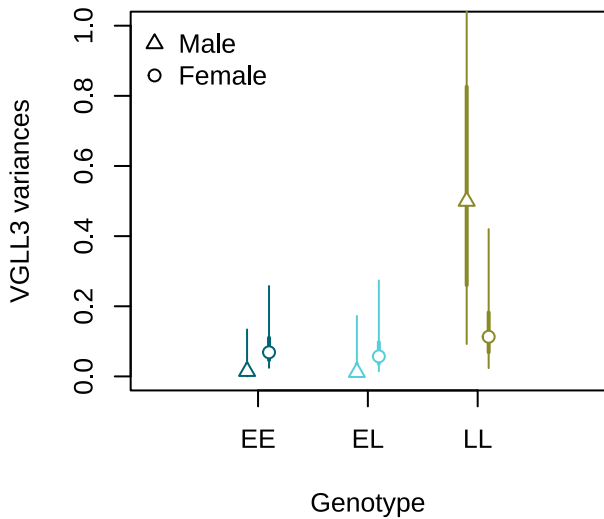
Female



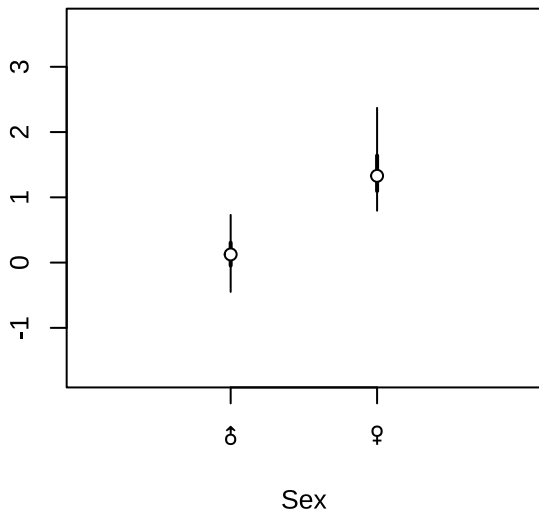




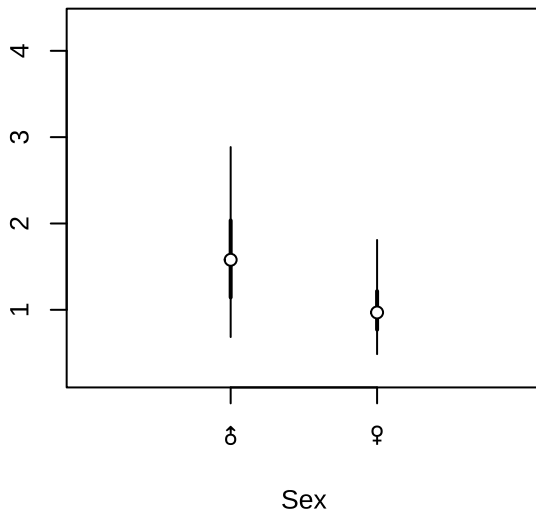




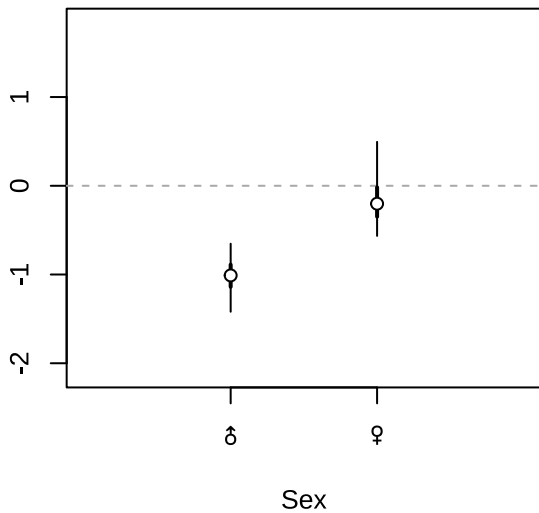
Mean Maturation thresholds



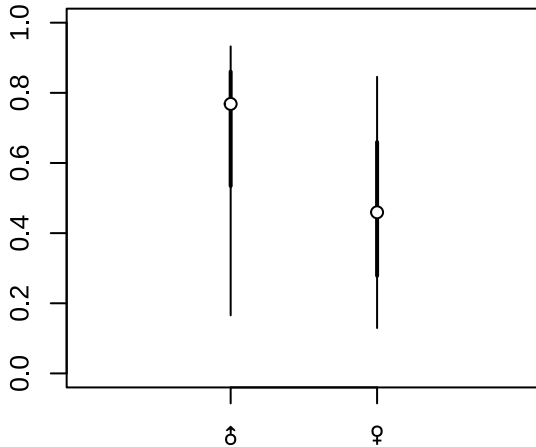
Genotypic value



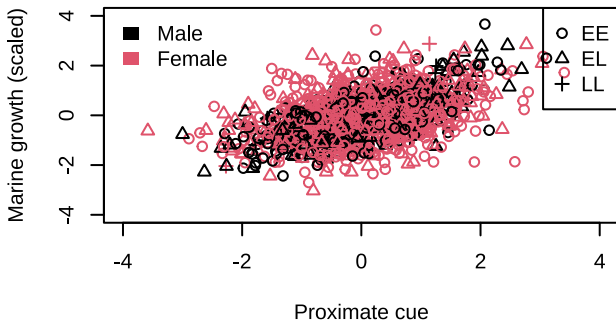
Dominance deviation (scaled)



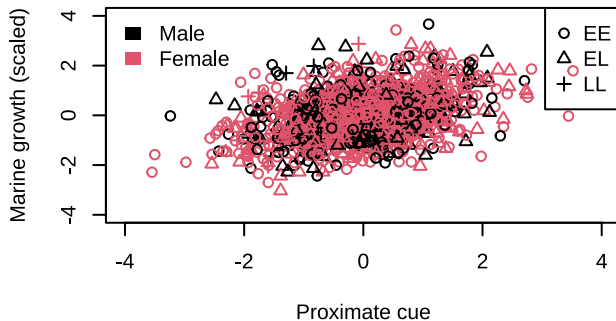
Total variance contribution



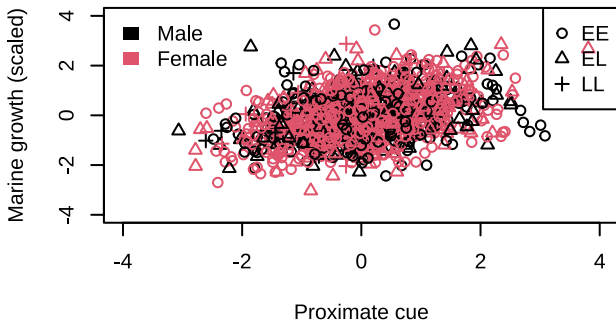
Iteration: 22996



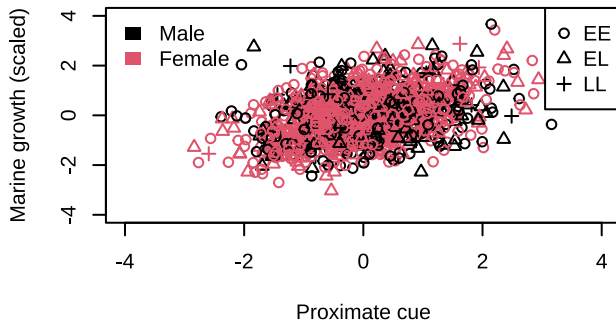
Iteration: 23195



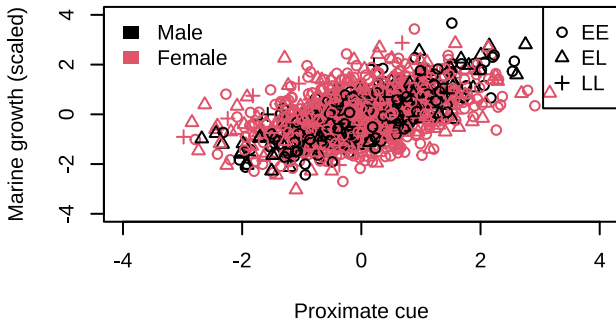
Iteration: 14129



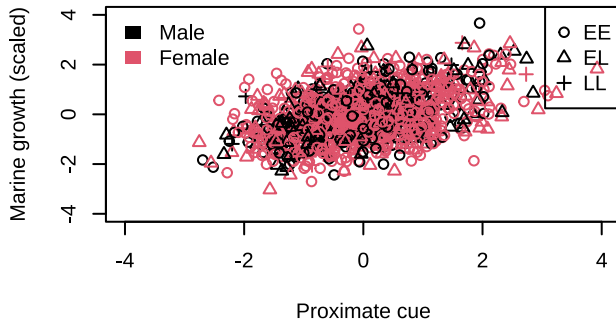
Iteration: 12398

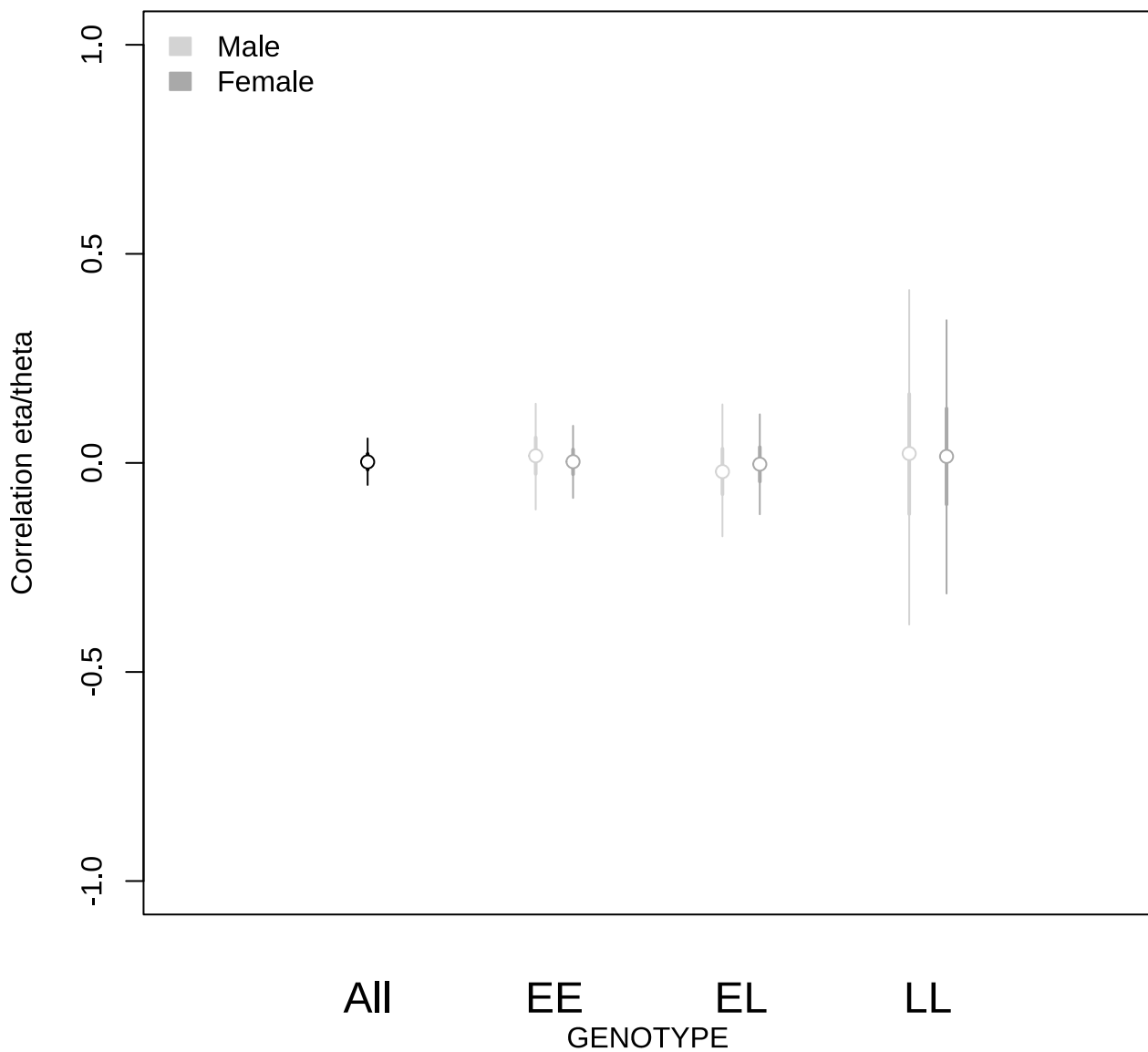


Iteration: 27258

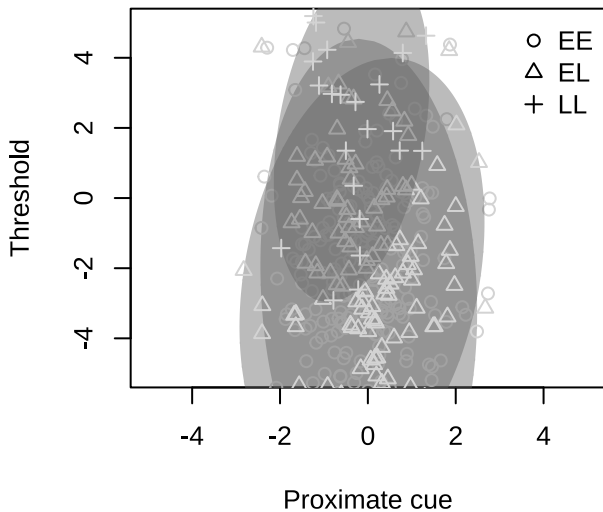


Iteration: 7781

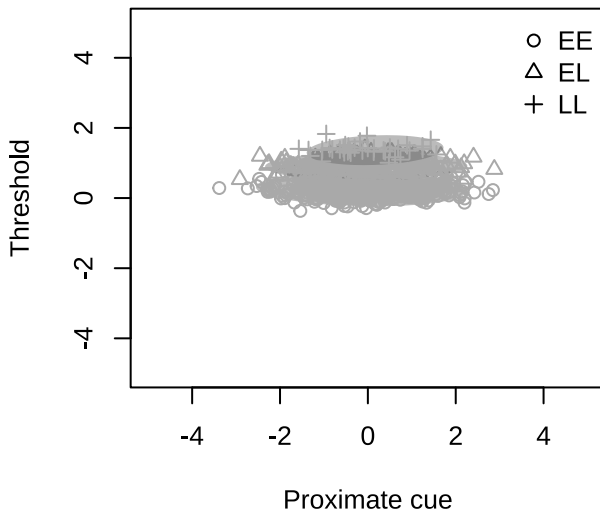




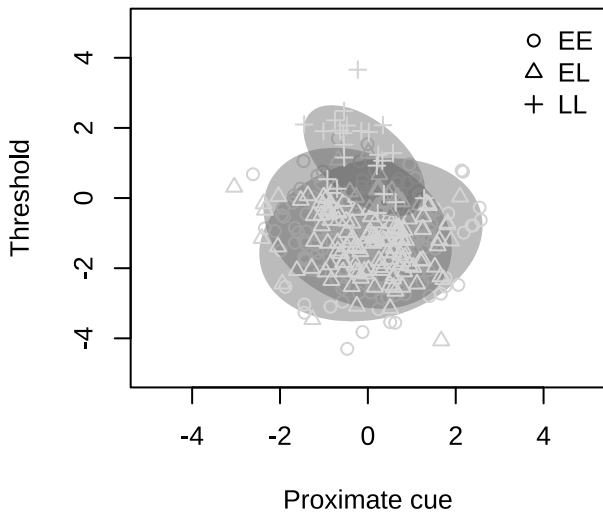
Male (iteration: 23252)



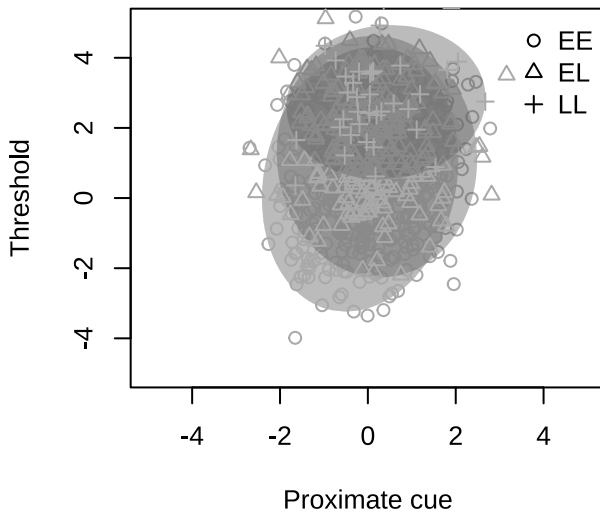
Female (iteration: 23252)

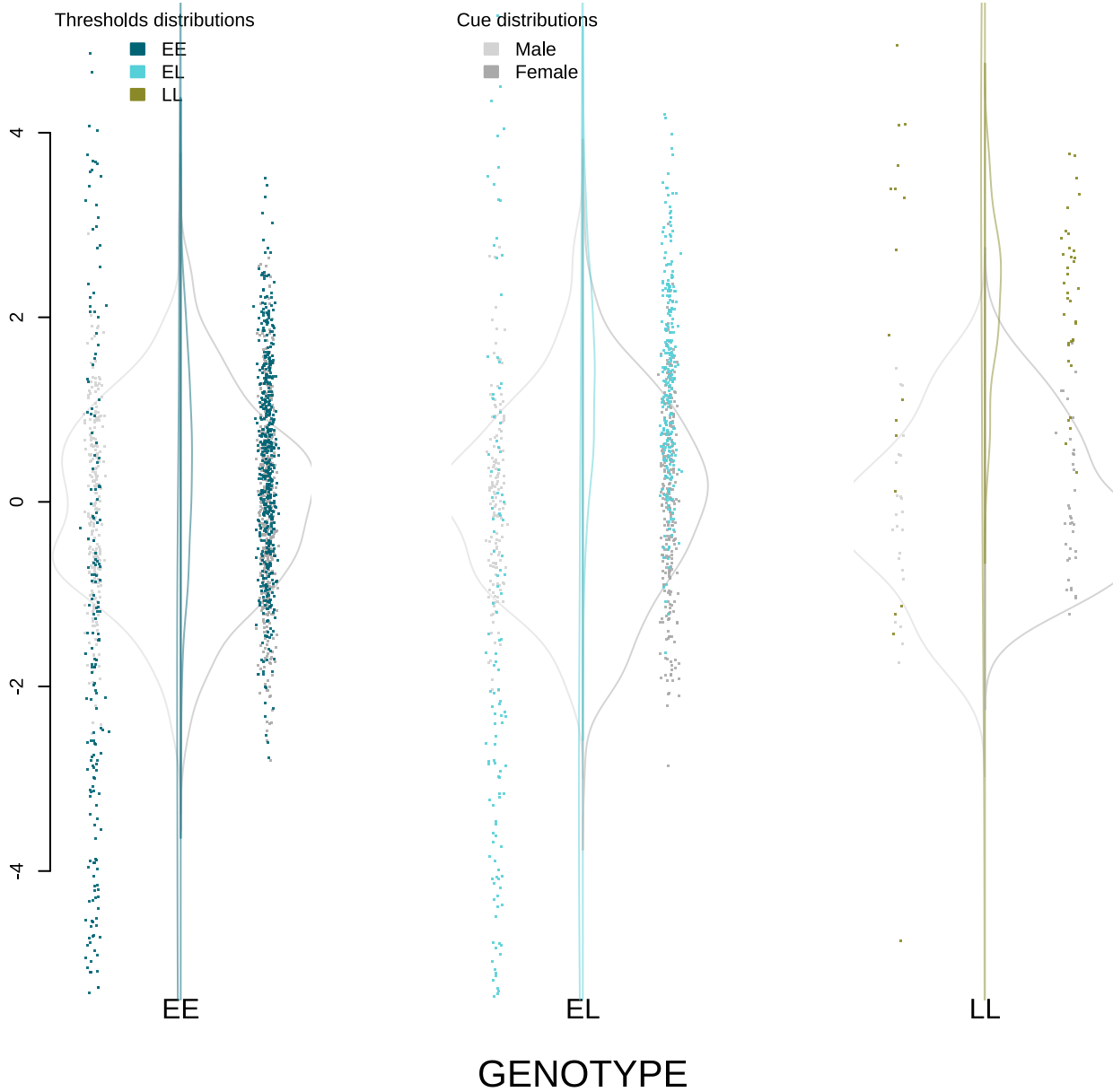


Male (iteration: 1607)



Female (iteration: 1607)





Tresholds (medians)

