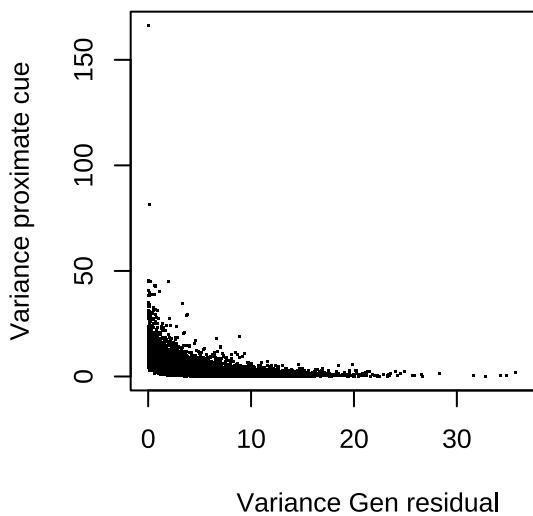
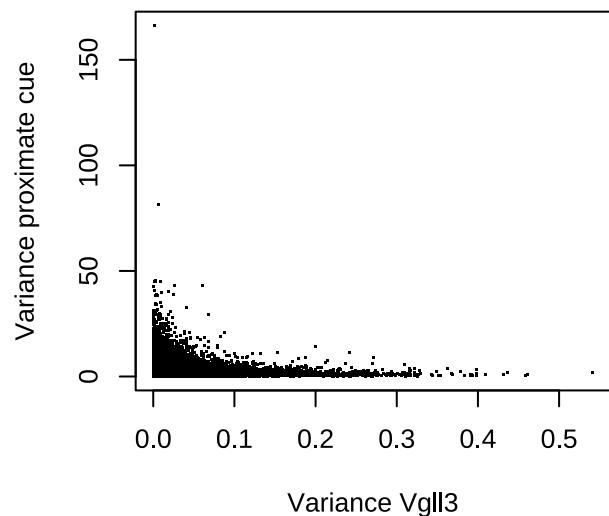
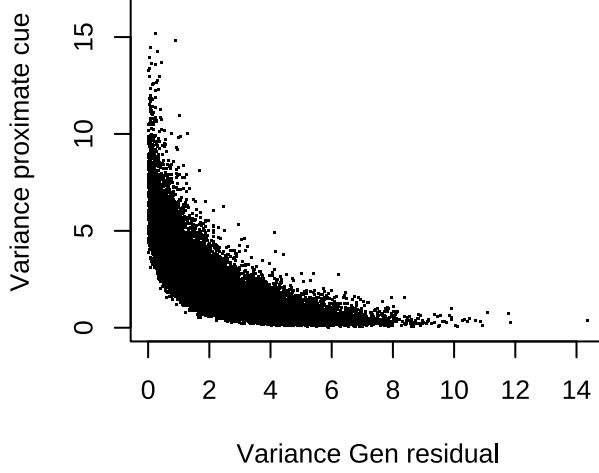
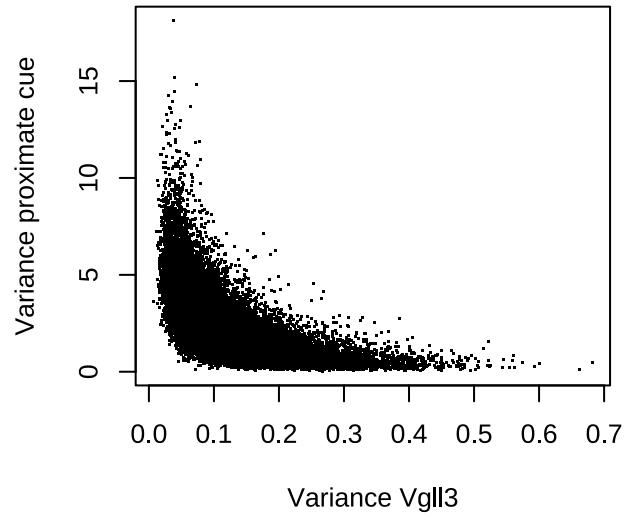
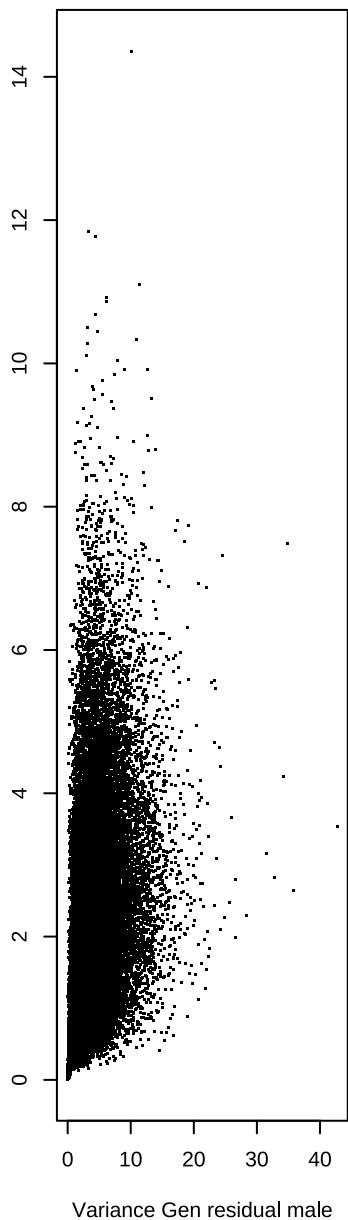
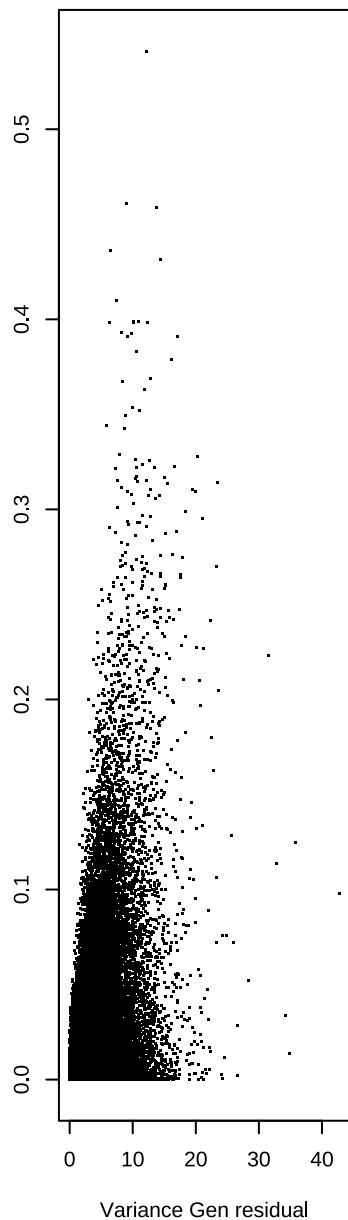


**Male****Male****Female****Female**

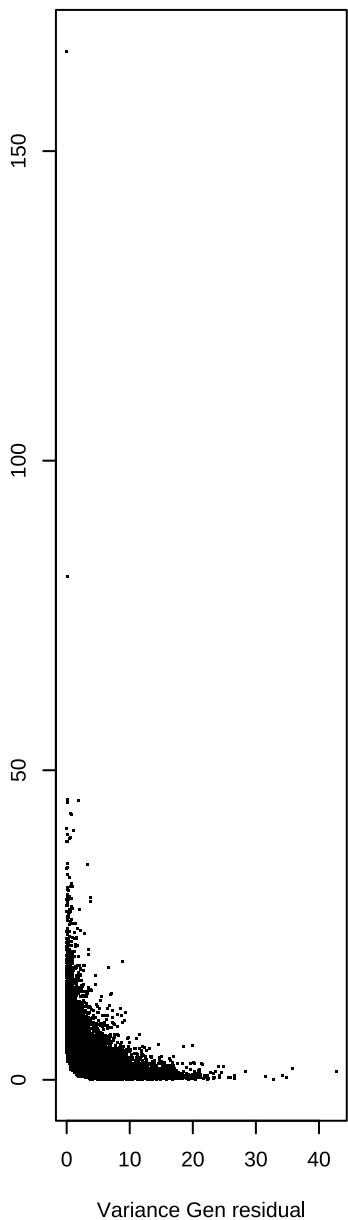
Variance Gen residual female

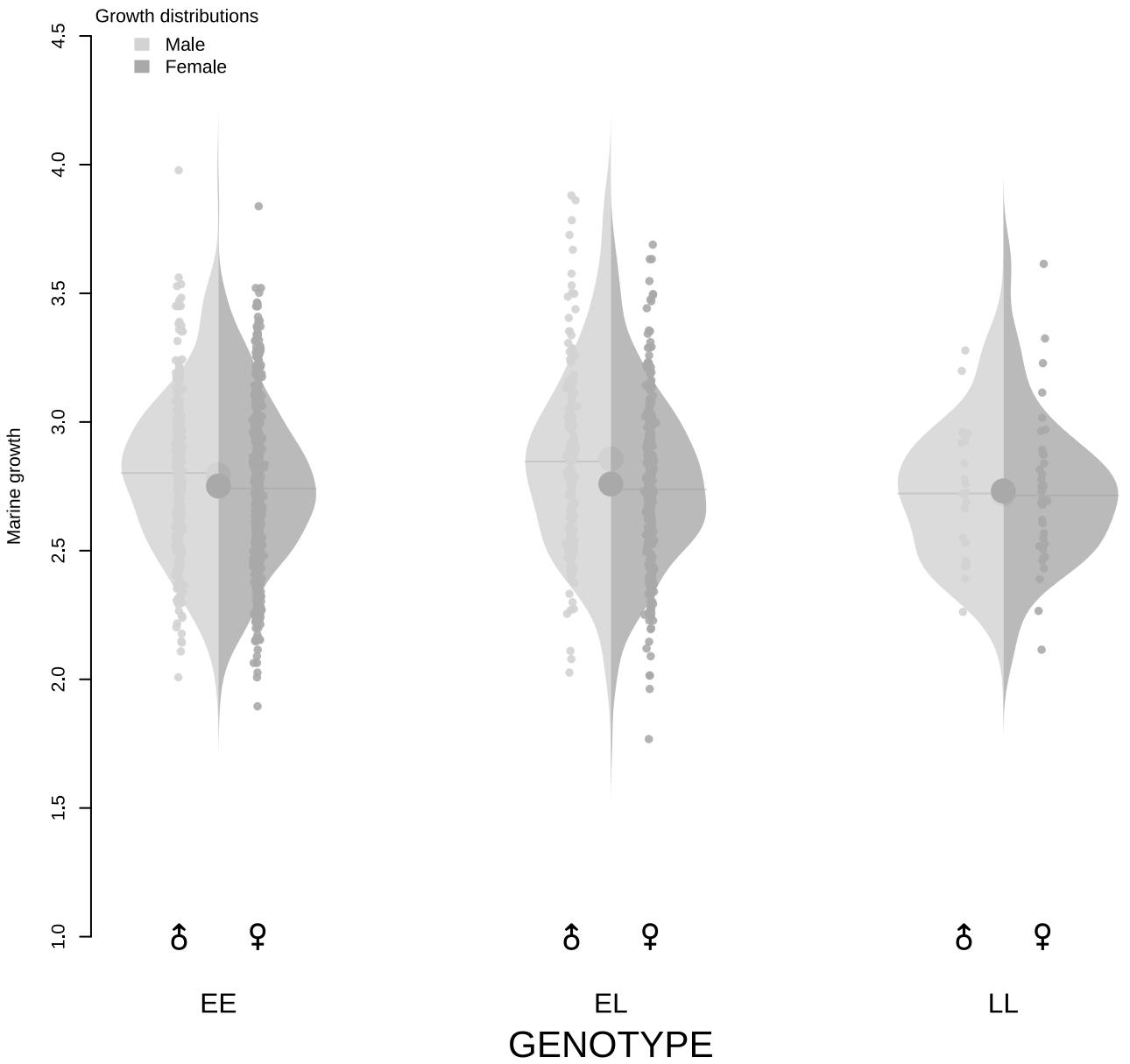


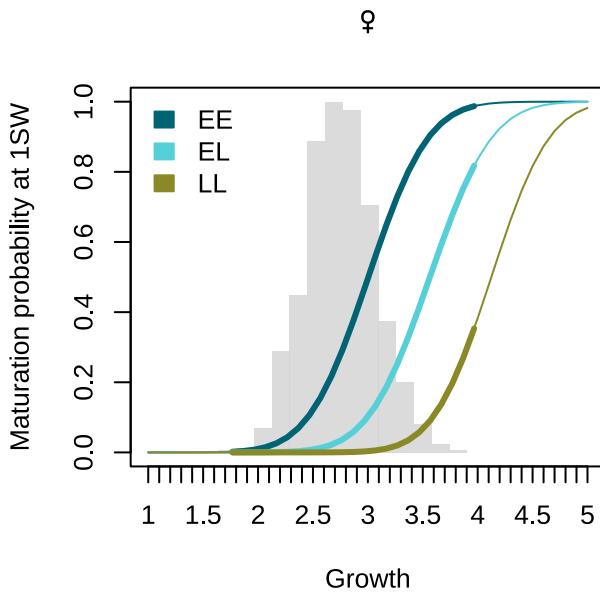
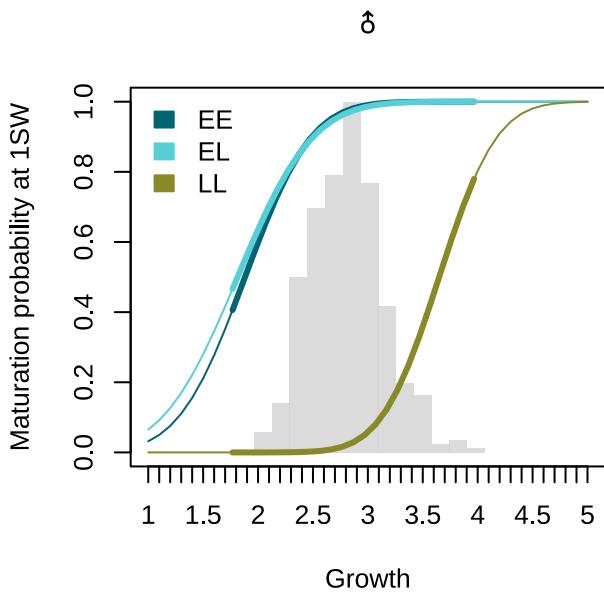
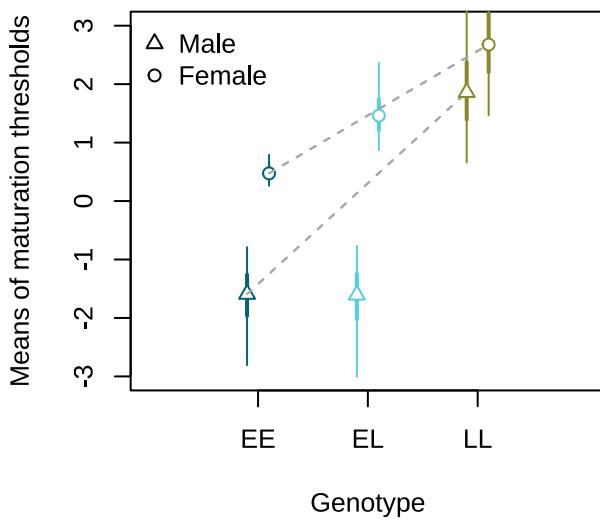
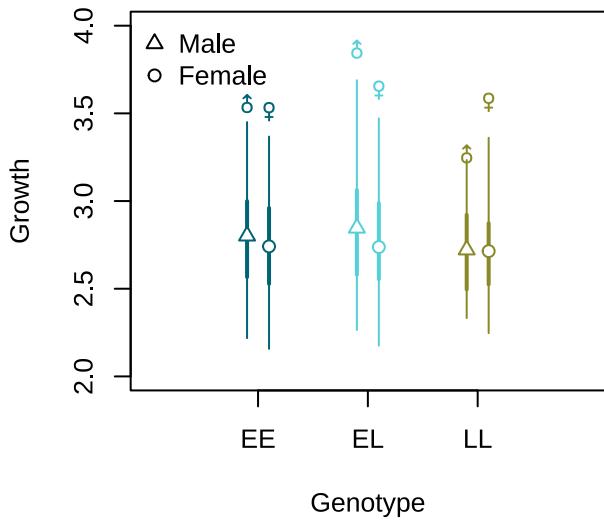
Variance VGLL3

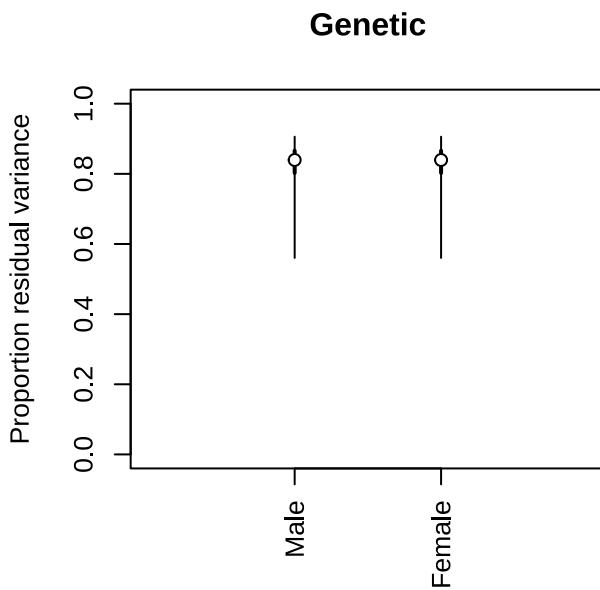
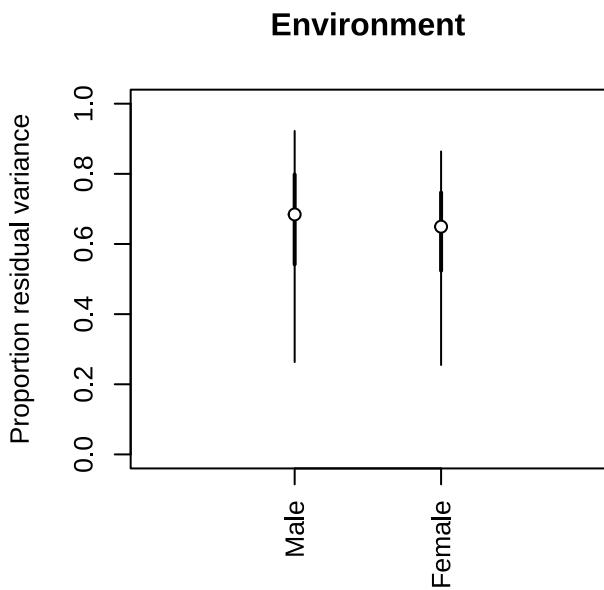
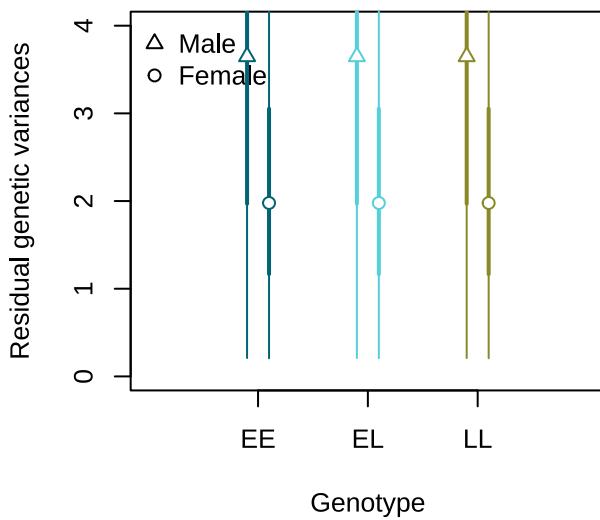
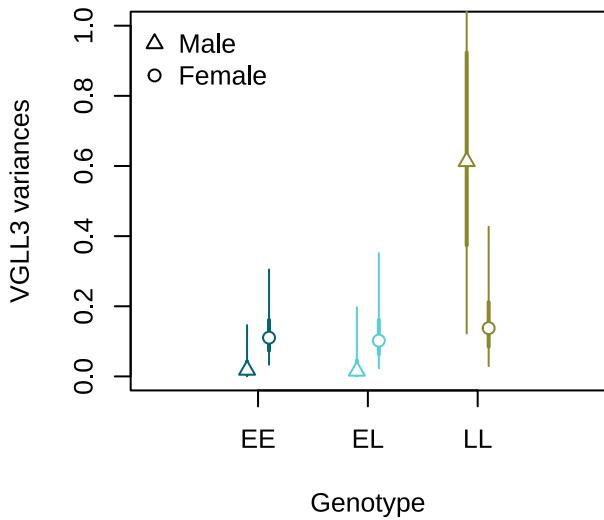


Variance eta

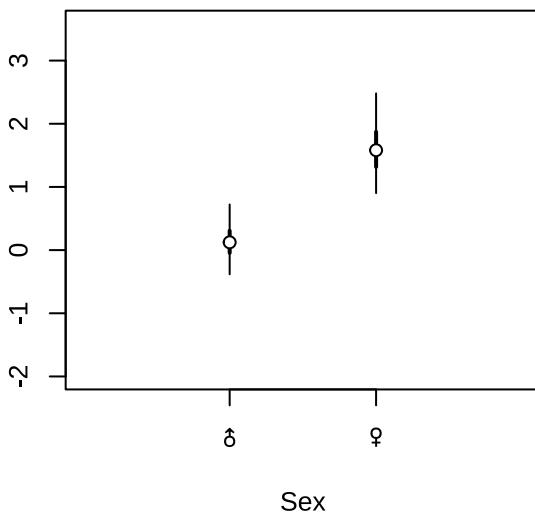




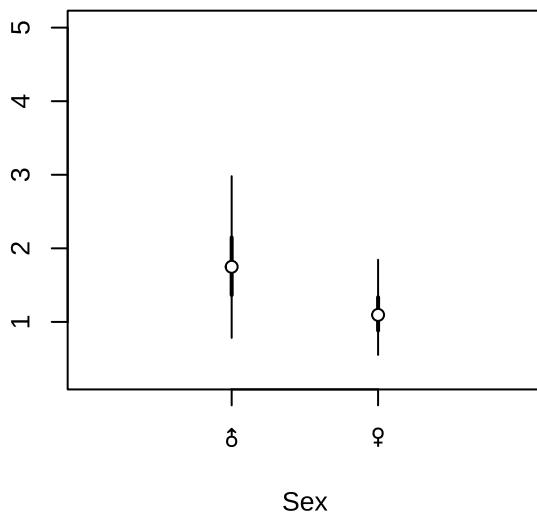




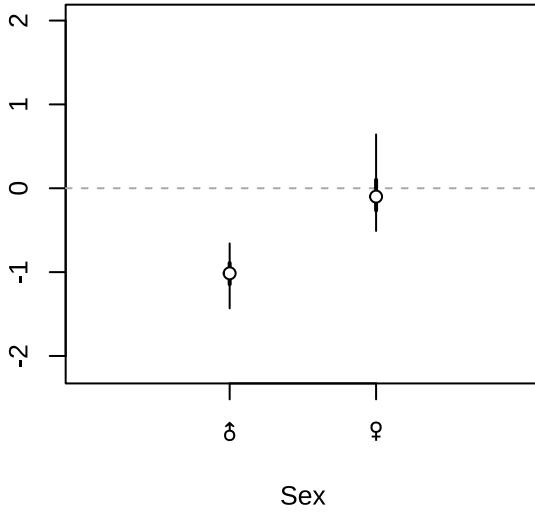
Mean Maturation thresholds



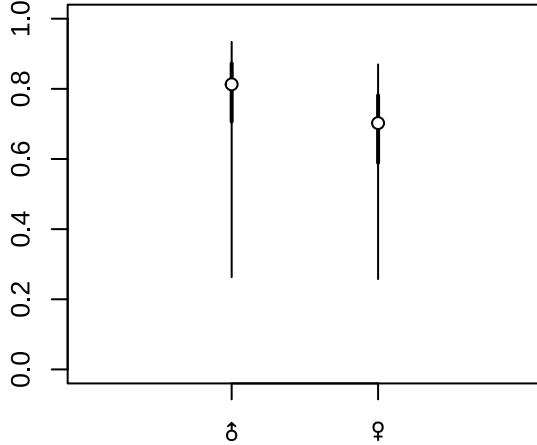
Genotypic value

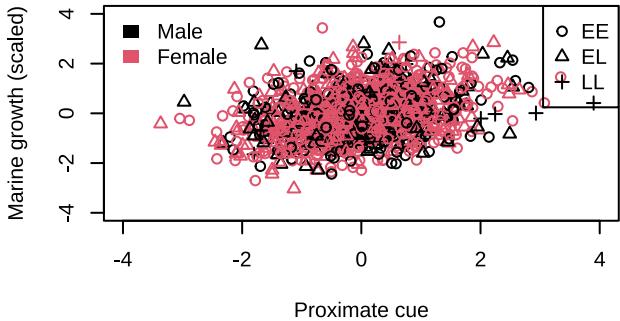
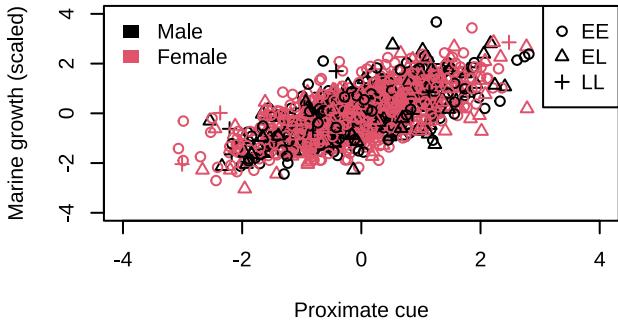
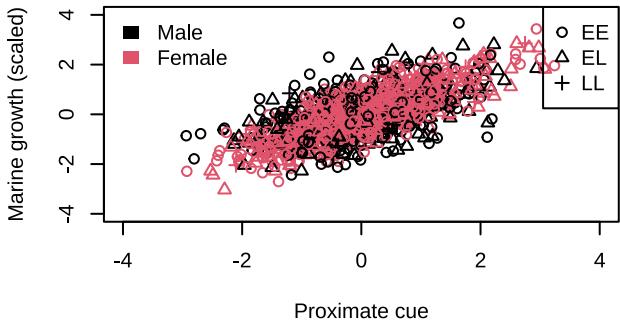
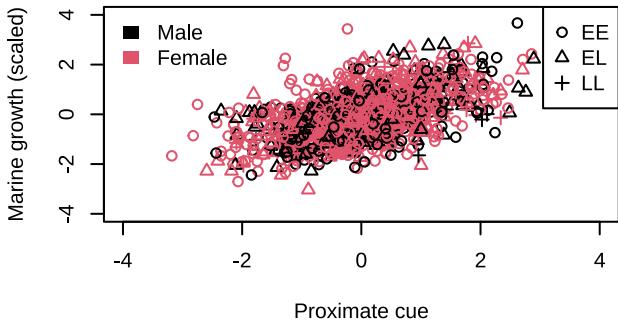
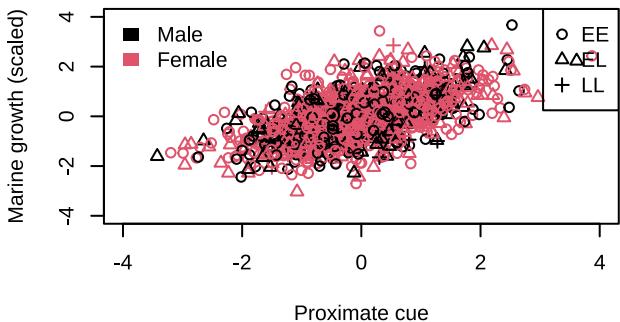
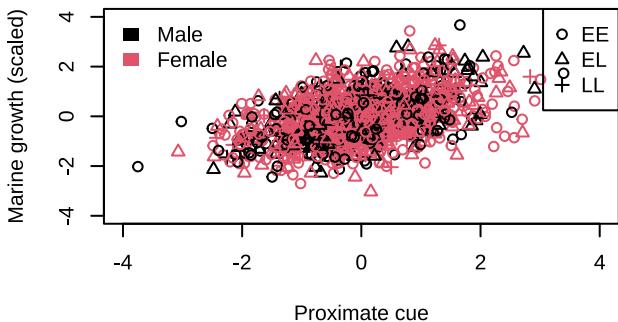


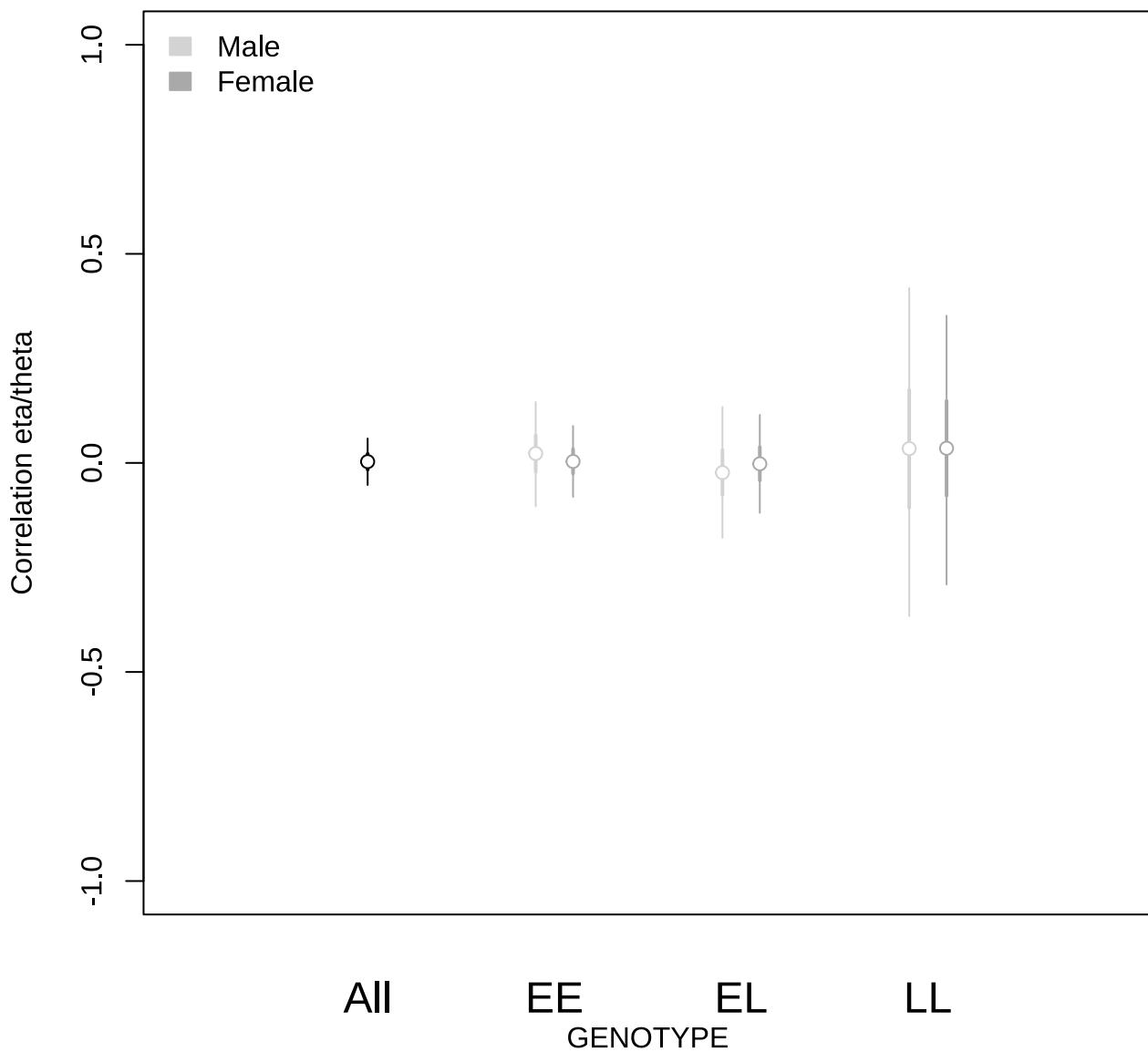
Dominance deviation (scaled)



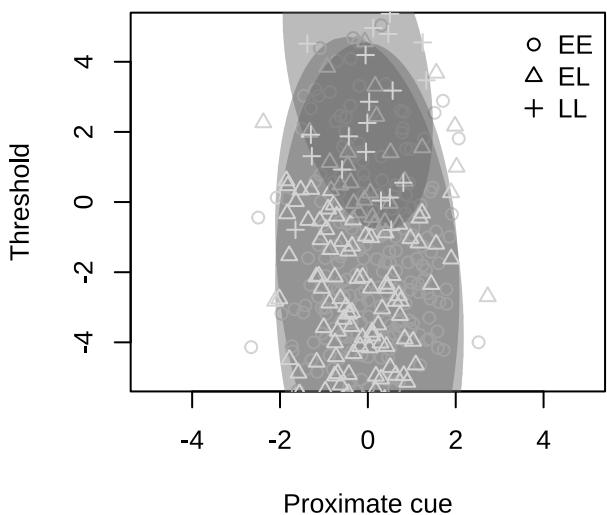
Total variance contribution



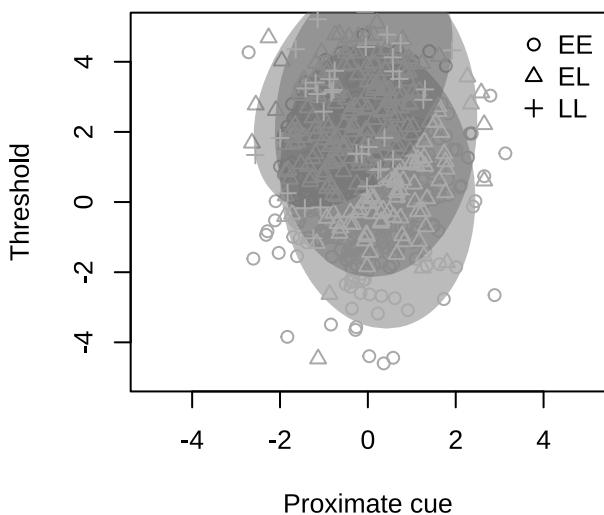
**Iteration: 23616****Iteration: 17872****Iteration: 4482****Iteration: 7613****Iteration: 16565****Iteration: 9501**



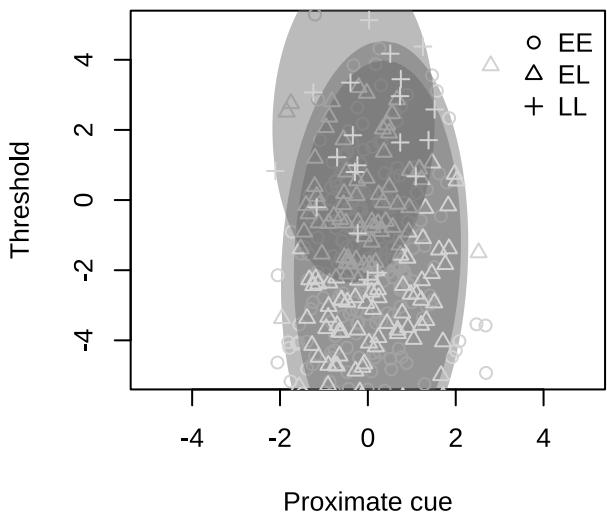
**Male (iteration: 17957)**



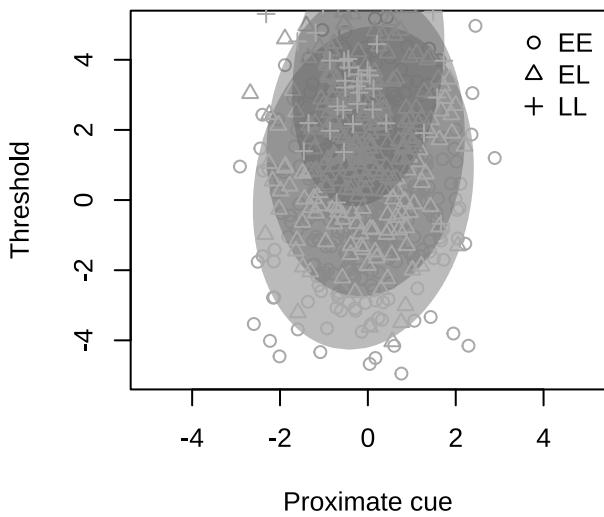
**Female (iteration: 17957)**



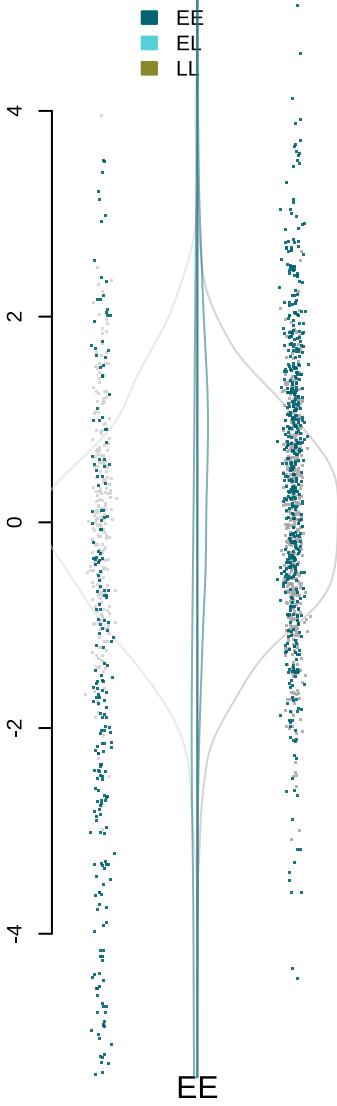
**Male (iteration: 3165)**



**Female (iteration: 3165)**

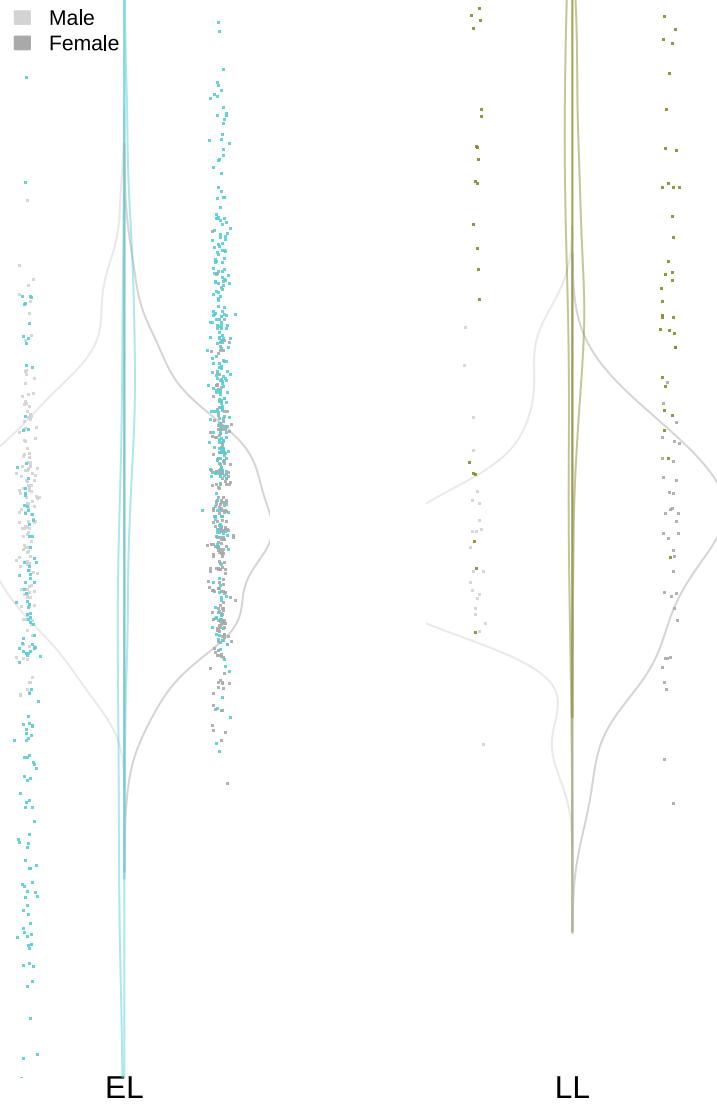


Thresholds distributions

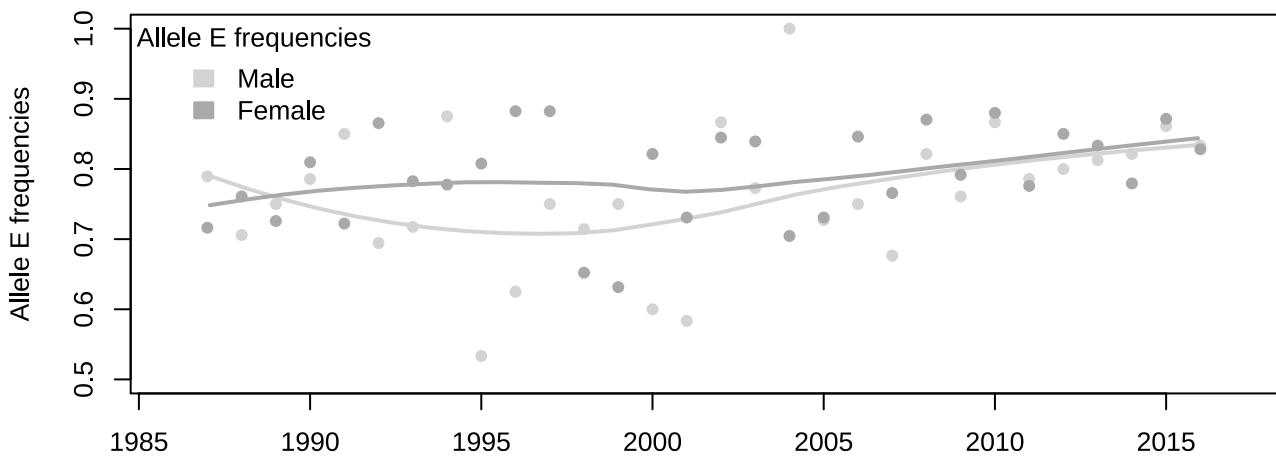
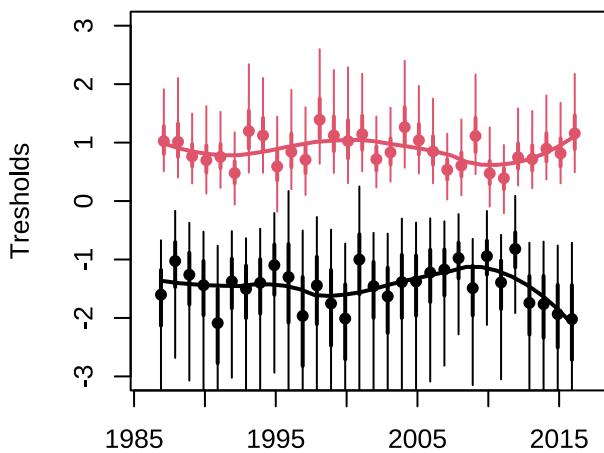
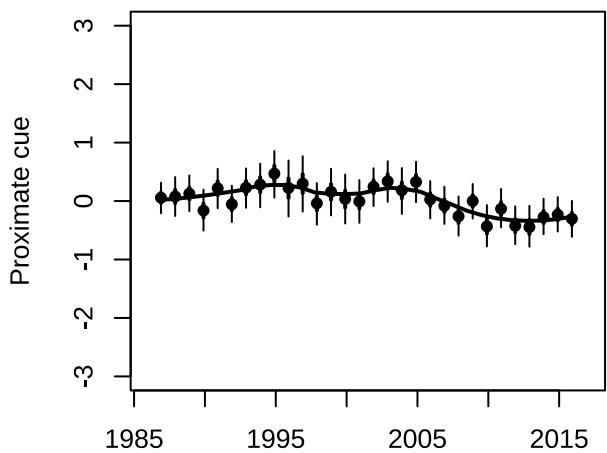


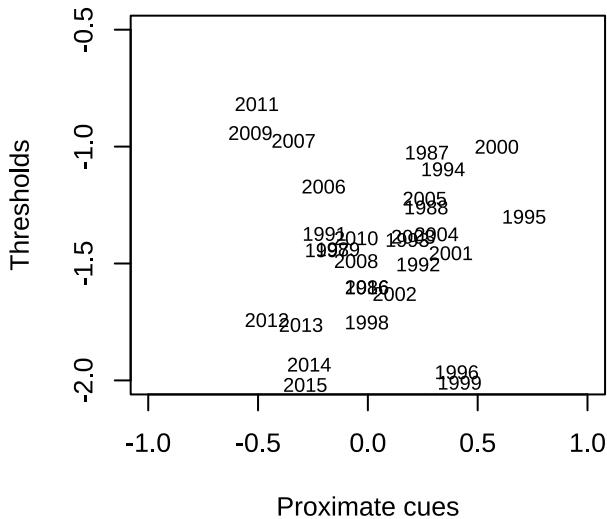
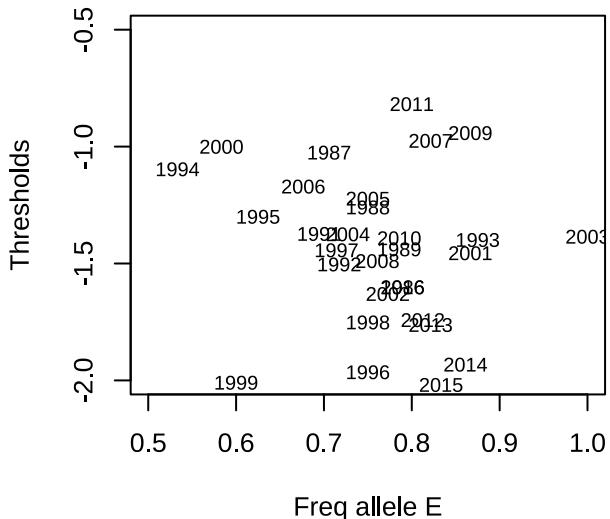
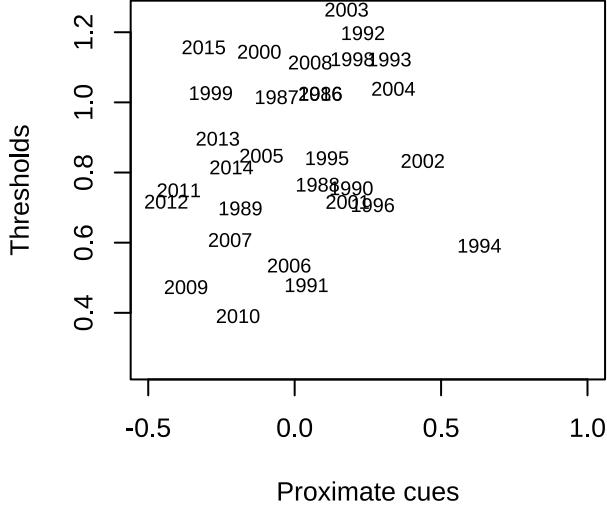
Cue distributions

Male  
Female



GENOTYPE



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