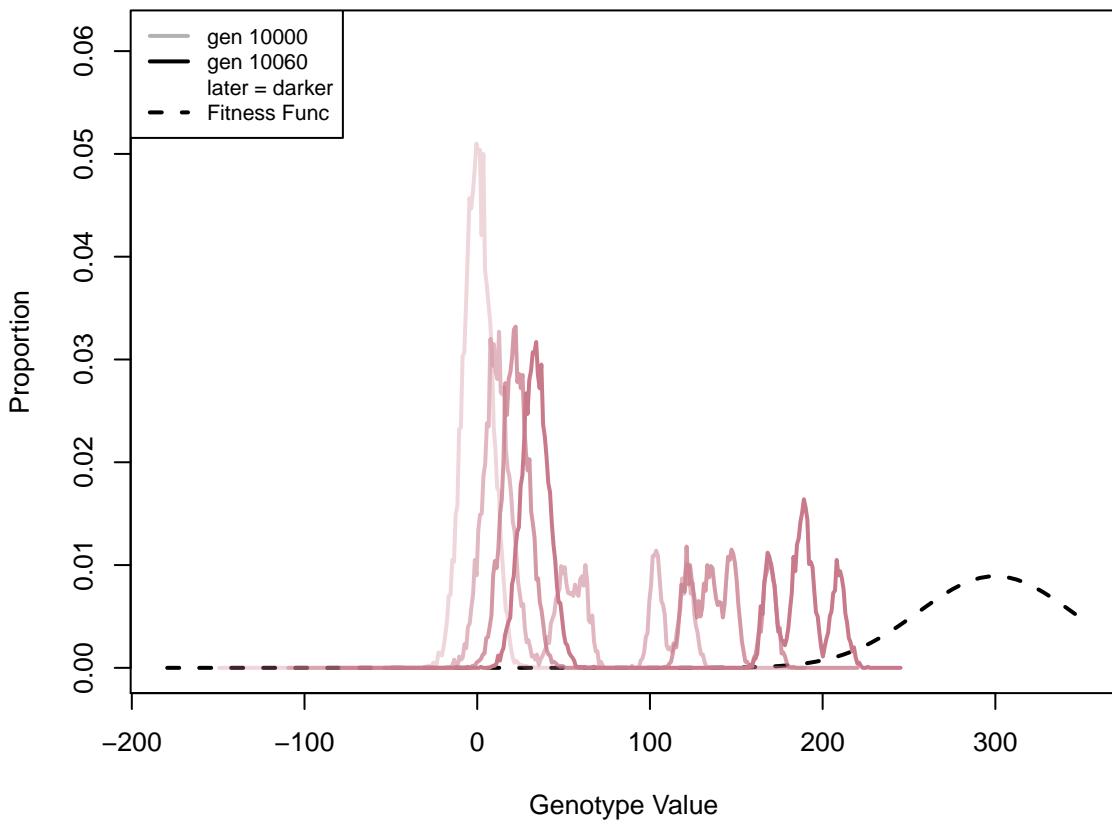
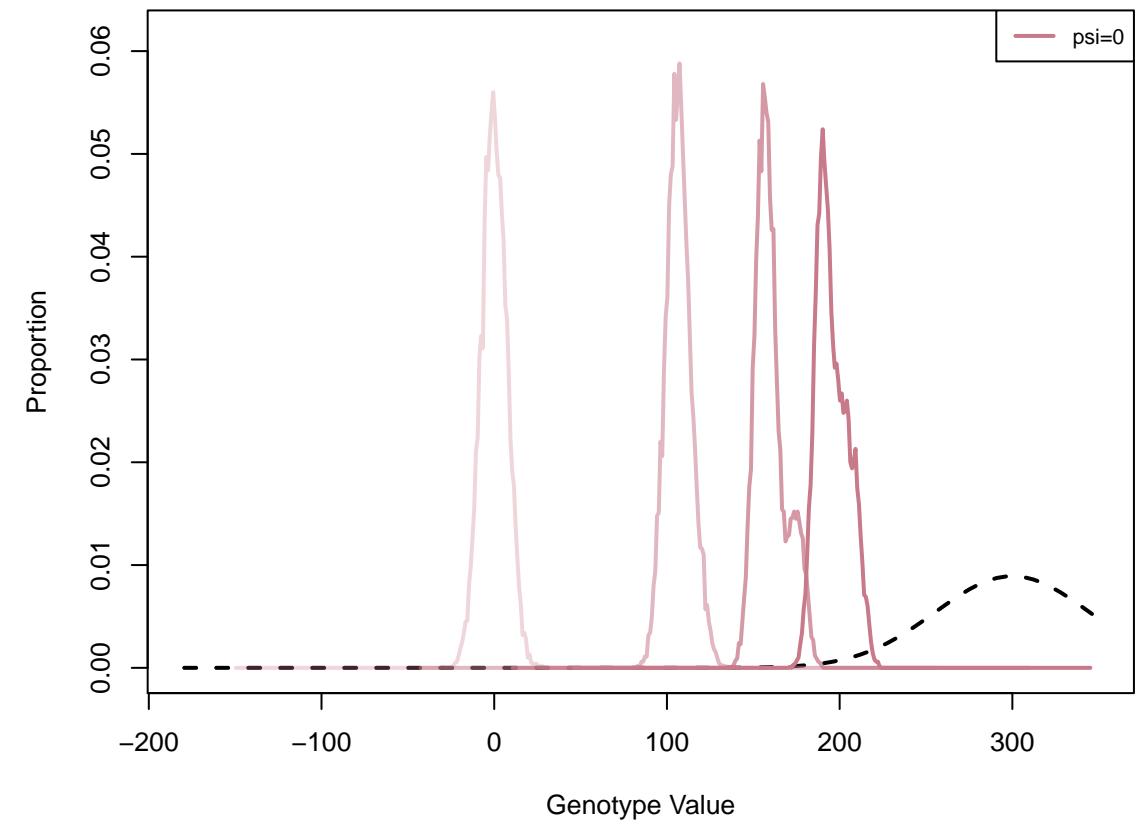


Genotype distributions |  $n = 20$  |  $shift\_size = 300$  by 20 | models: ave, ext, fit, inv

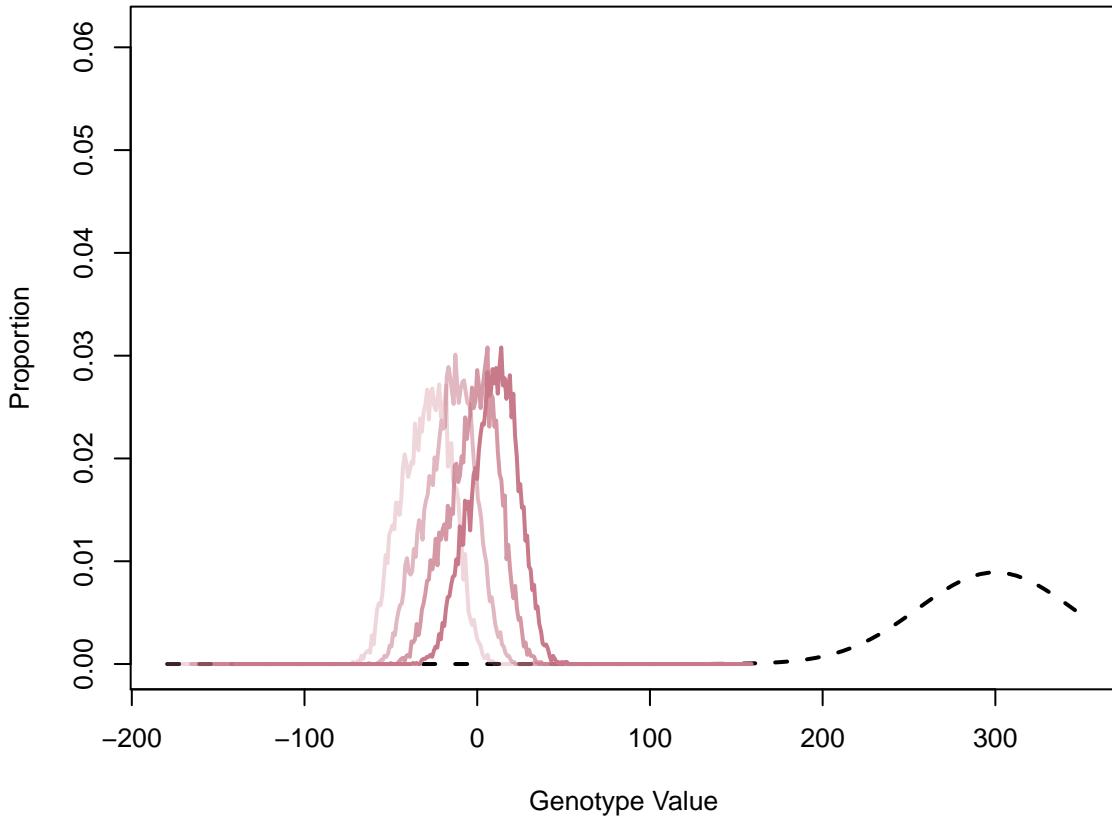
**Model = ave**



**Model = ext**



**Model = fit**



**Model = inv**

