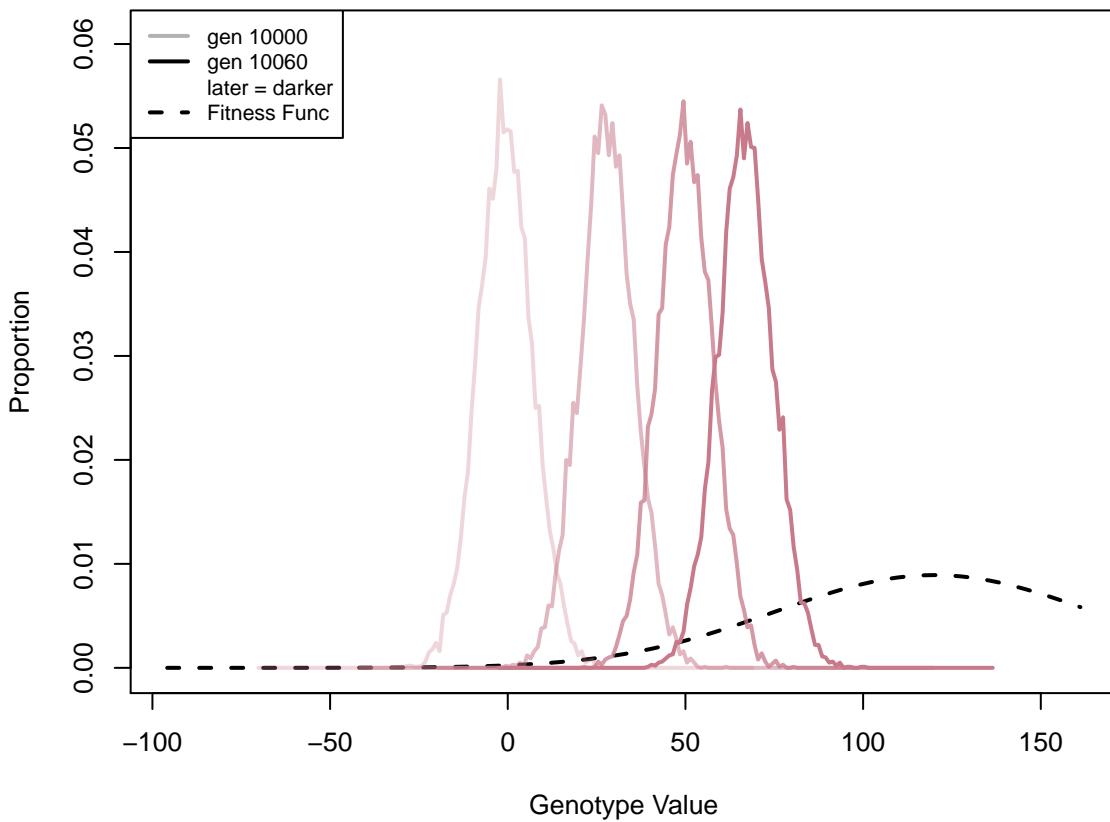
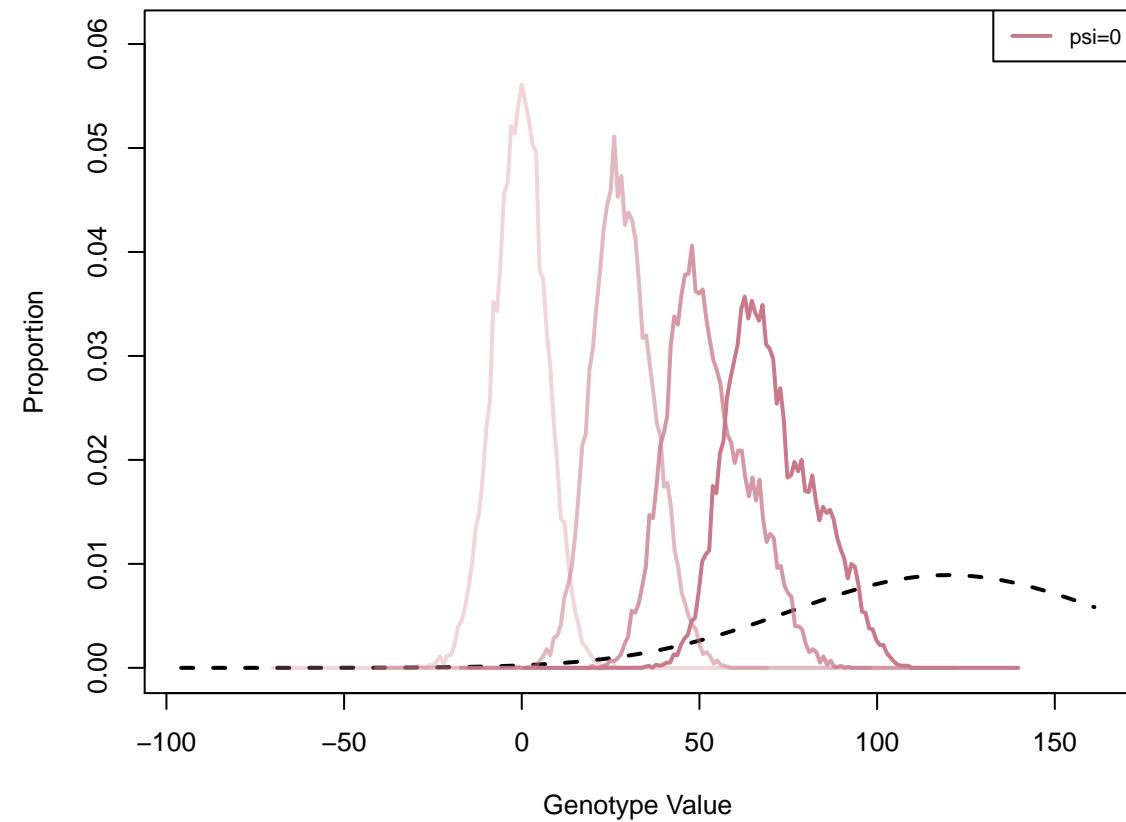


Genotype distributions | $n = 2$ | shift_size = 120 by 20 | models: ave, ext, fit, inv

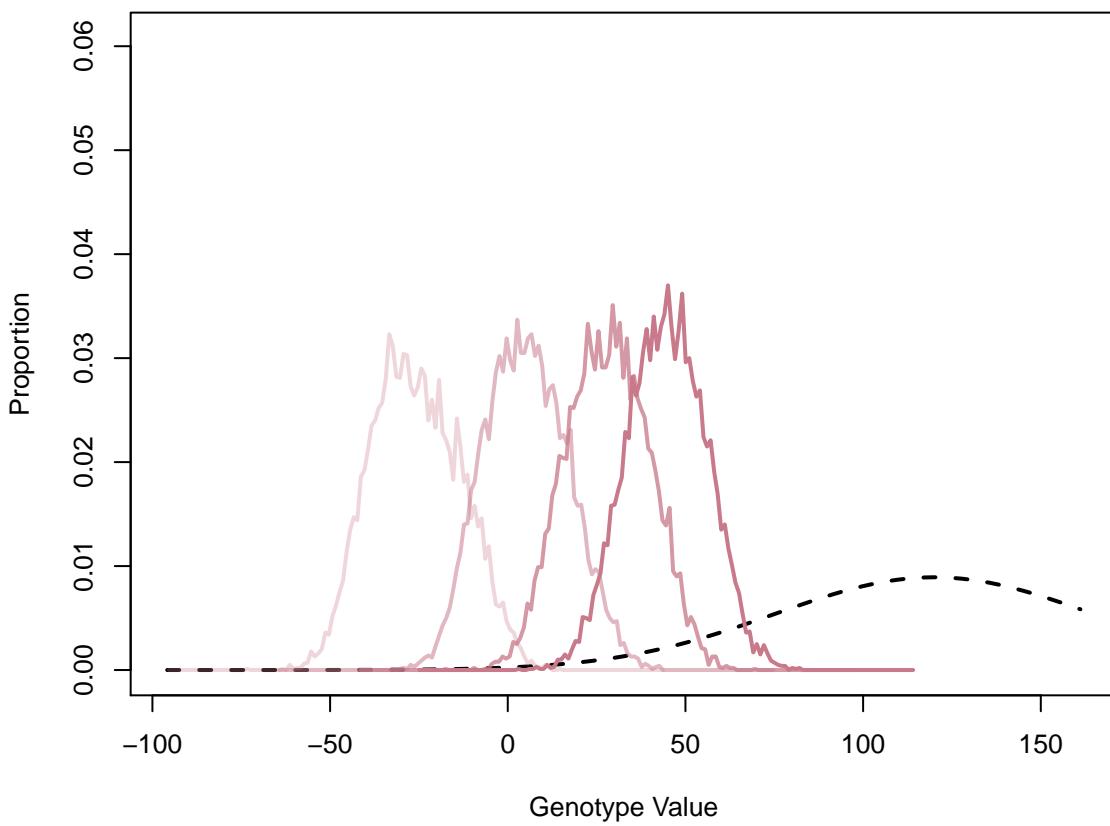
Model = ave



Model = ext



Model = fit



Model = inv

