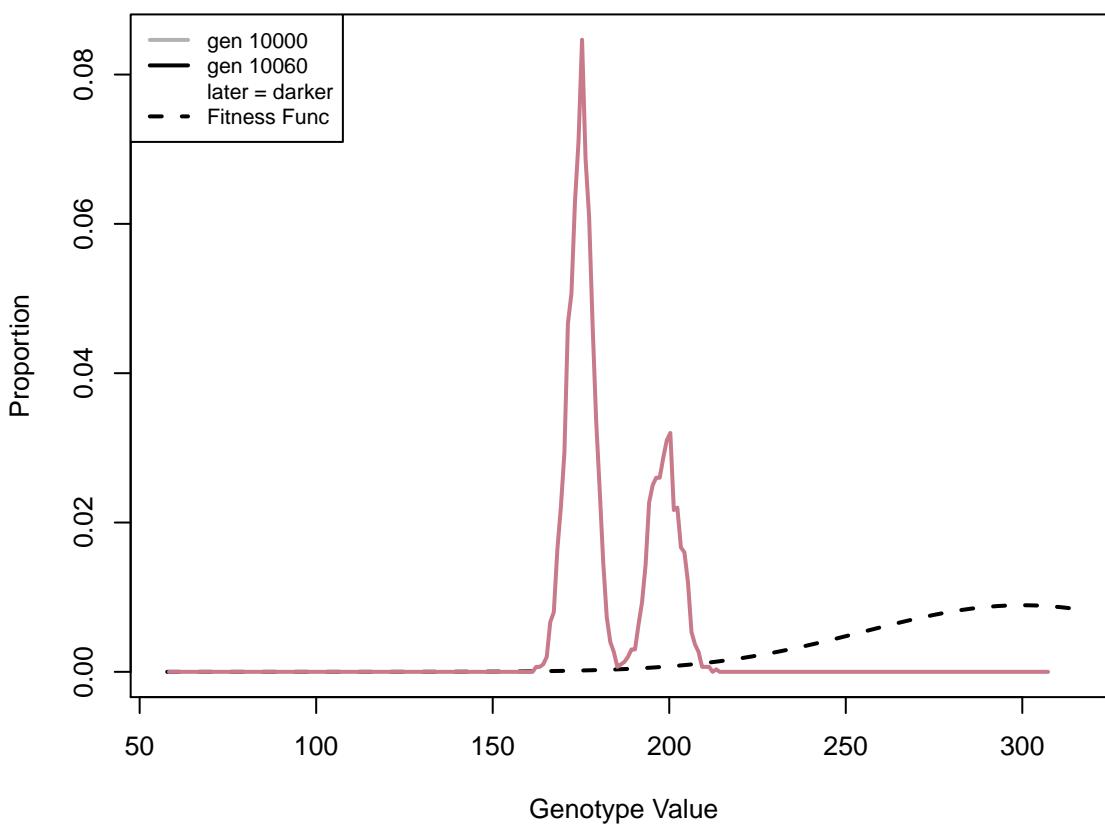
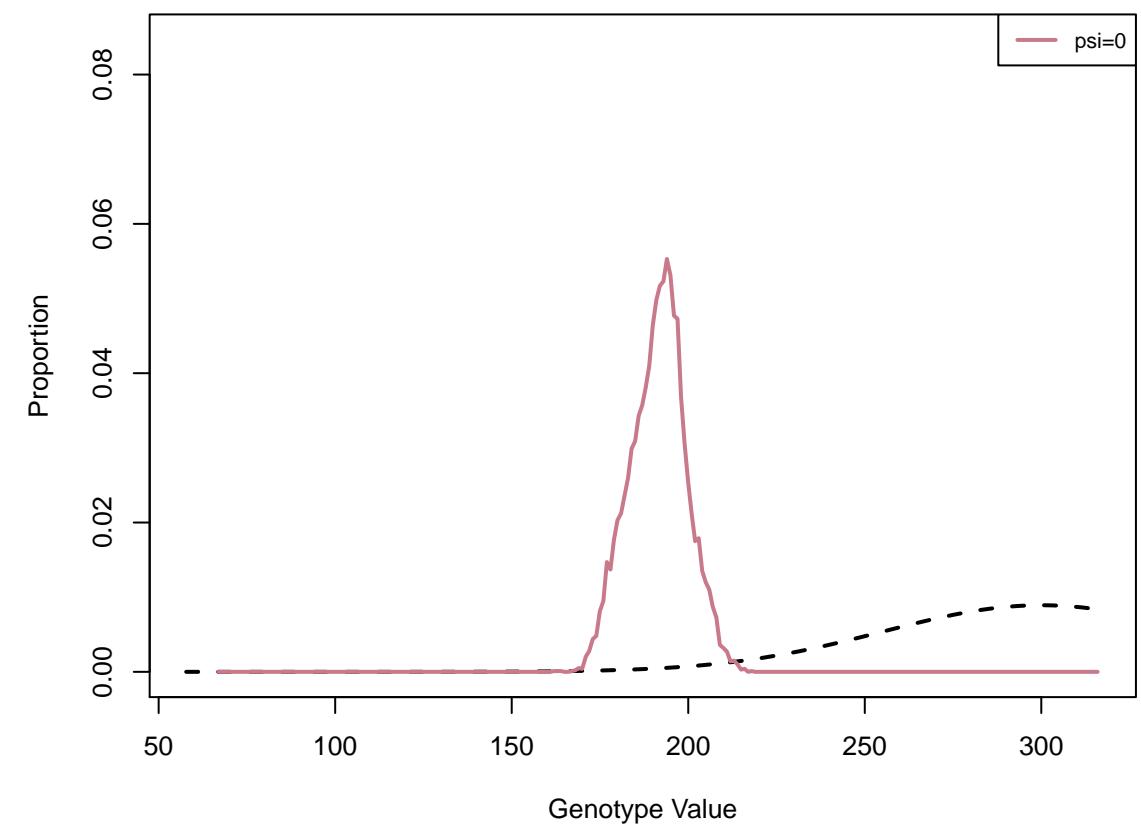


Genotype distributions | $n = 8$ | $\text{shift_size} = 300$ by NA | models: ave, ext, fit, inv

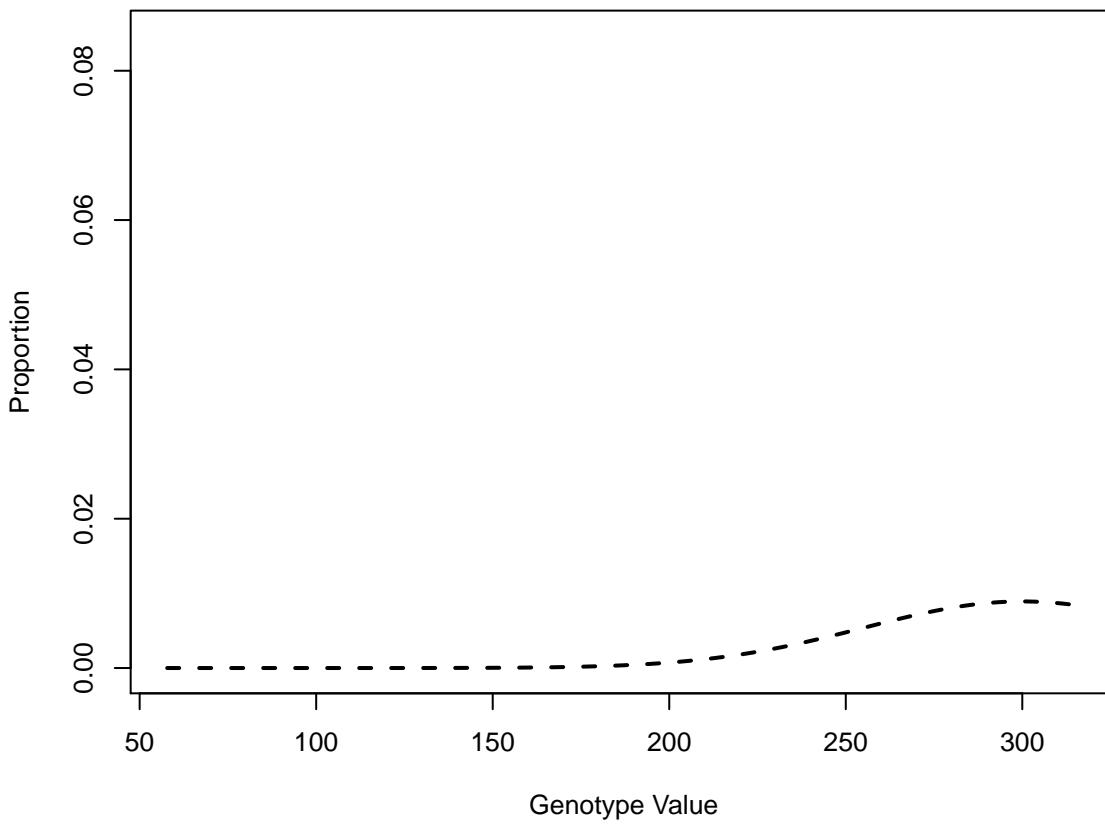
Model = ave



Model = ext



Model = fit



Model = inv

