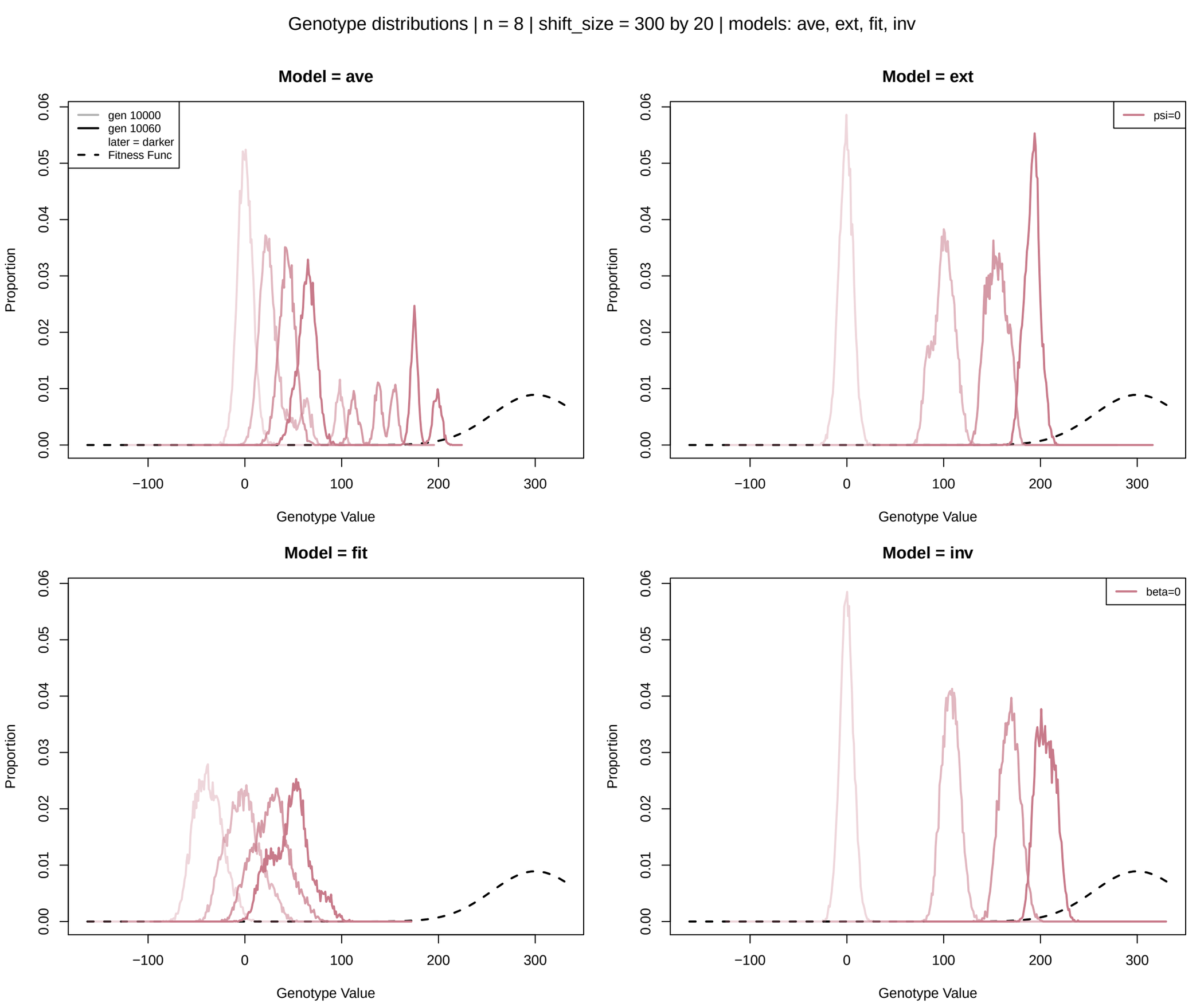
Genotype distributions | n = 8 | shift_size = 300 by 20 | models: ave, ext, fit, inv
Model = ave
gen 10000
gen 10060
later = darker
Fitness Func
Model = ext
psi=0
Model = fit
Model = inv
beta=0
Proportion
Genotype Value
0.00
0.01
0.02
0.03
0.04
0.05
0.06
−100
0
100
200
300