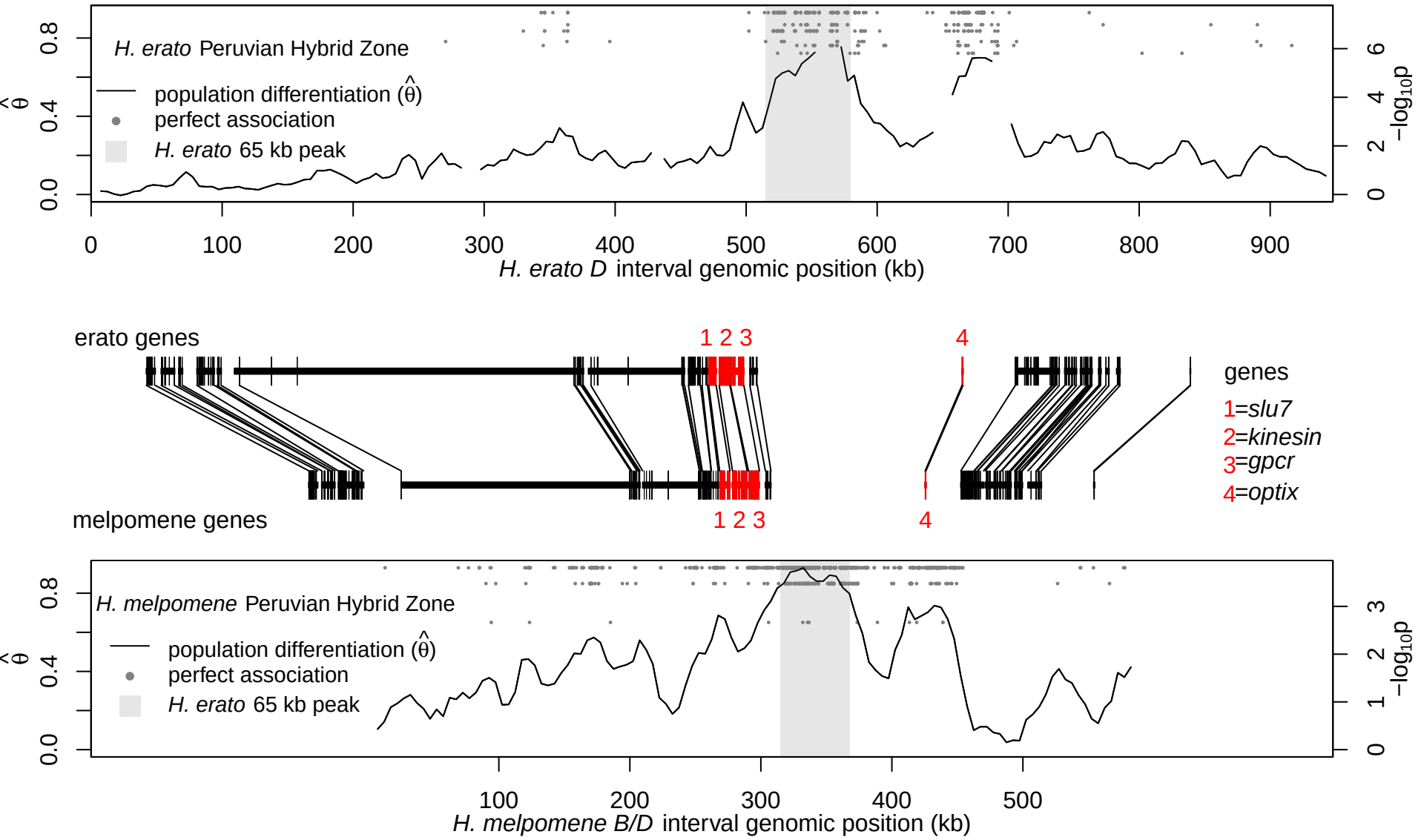
H. erato Peruvian Hybrid Zone
population differentiation ($\hat{\theta}$)
perfect association
H. erato 65 kb peak
$\hat{\theta}$
0.0
0.4
0.8
$-\log_{10}p$
0
2
4
6
0
100
200
300
400
500
600
700
800
900
H. erato D interval genomic position (kb)
erato genes
1 2 3
4
genes
1=slu7
2=kinesin
3=gpcr
4=optix
melpomene genes
1 2 3
4
H. melpomene Peruvian Hybrid Zone
population differentiation ($\hat{\theta}$)
perfect association
H. erato 65 kb peak
$\hat{\theta}$
0.0
0.4
0.8
$-\log_{10}p$
0
1
2
3
100
200
300
400
500
H. melpomene B/D interval genomic position (kb)