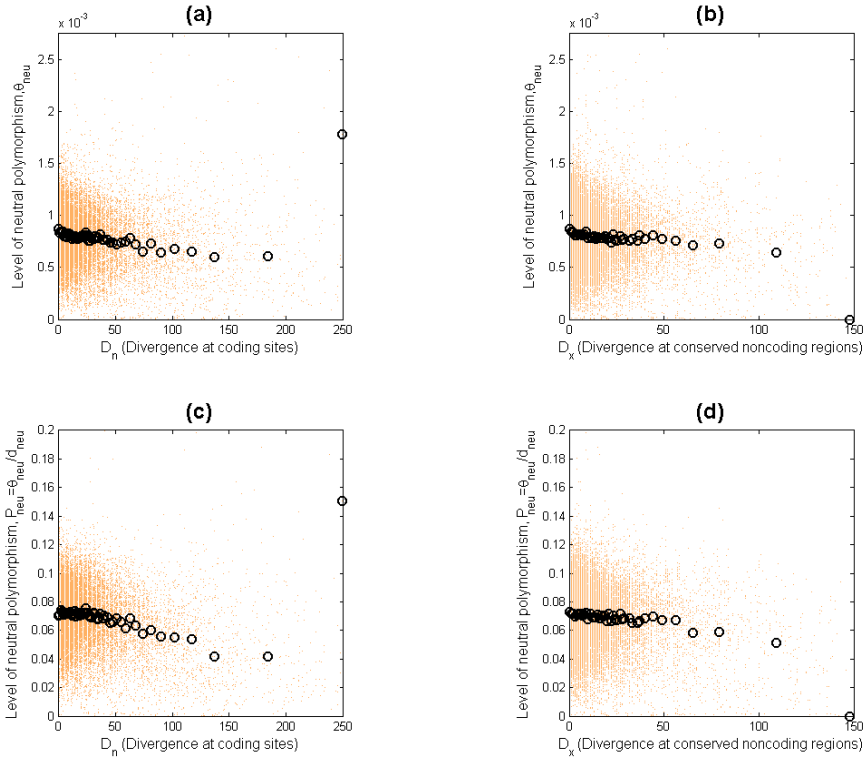
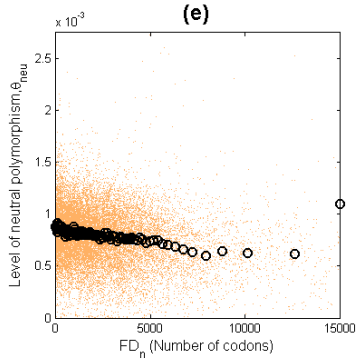
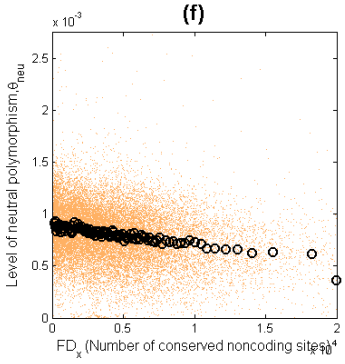
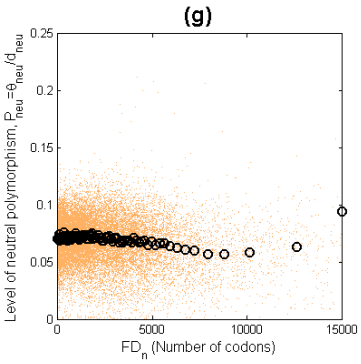
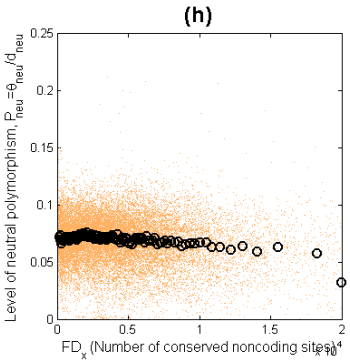
Figure S10. Results derived from the sliding windows of 600 kb. Correlations between functional density [i.e. the number of codons (FD_n) or the number of conserved noncoding sites (FD_x)], and divergence [i.e. the divergence at coding sites (D_n) or the divergence at conserved noncoding region (D_x)] and neutral polymorphism [i.e. the level of neutral polymorphism (θ_{neu}) and the level of normalized neutral polymorphism ($P_{neu}=\theta_{neu}/d_{neu}$)] are given. The results are based on the Watson data.




(e)
x 10^-3
Level of neutral polymorphism, θneu
FDn (Number of codons)


(f)
x 10^-3
Level of neutral polymorphism, θneu
FDx (Number of conserved noncoding sites)
x 10^4


(g)
Level of neutral polymorphism, Pneu =θneu/dneu
FDn (Number of codons)


(h)
Level of neutral polymorphism, Pneu =θneu/dneu
FDx (Number of conserved noncoding sites)
x 10^4