

Figure S1: SNP density map.

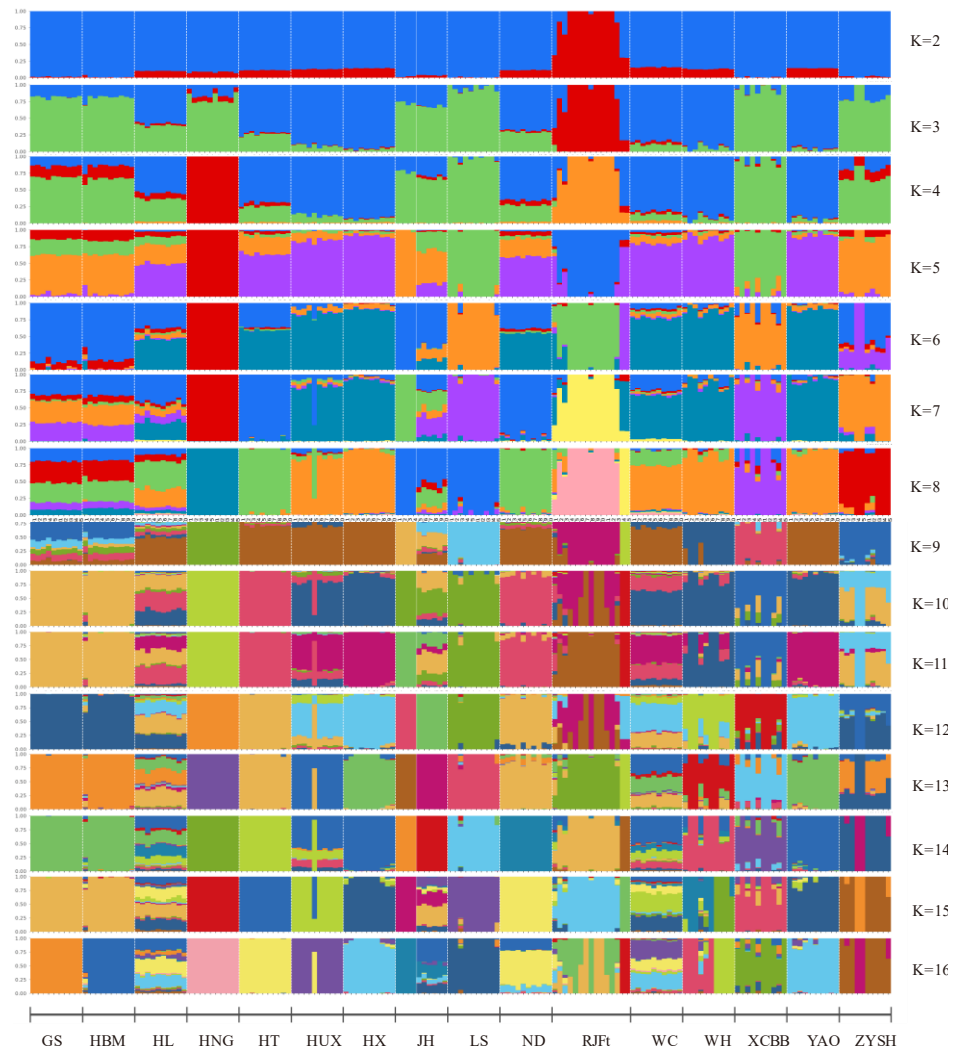


Figure S2. Admixture analysis across 16 chicken populations. Pro-portions of geneticancestry for 16 chicken populations with K=2-16 (K represents the number of inferred ancestral populations).

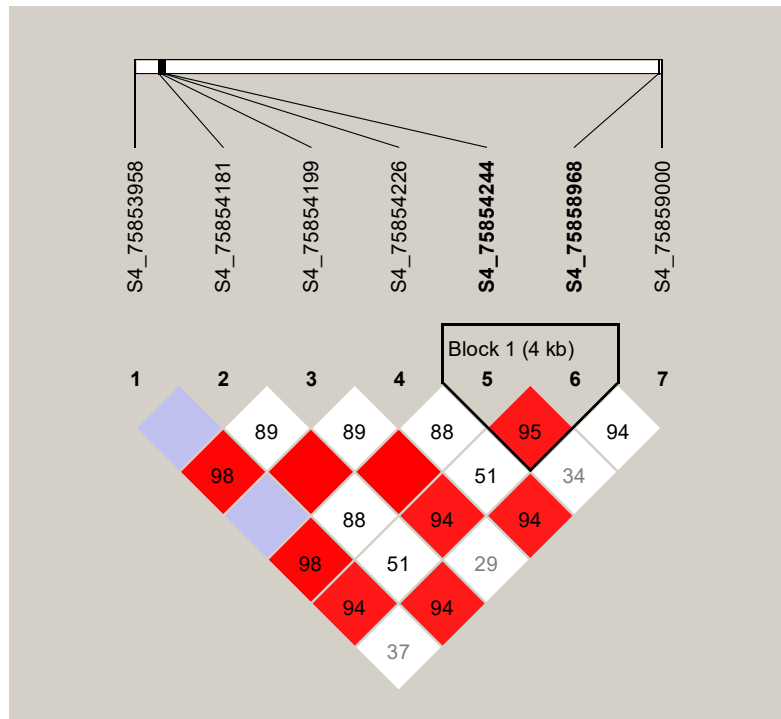


Figure S3: 6 SNPs on *LCORL* gene LD block.