A
Count
Haplotype SNP Count
B
Count
Frequency in Strain
C
High
Low
D. azt
D. fal
D. inn
PR
GA
HU/SR
CH
PC2 (26.7%)
PC1 (52.1%)
D
Between viral haplotype SNPs
Between viral haplotype SNPs and other SNPs
Between other SNPs
r2
Distance between SNPs (bp)