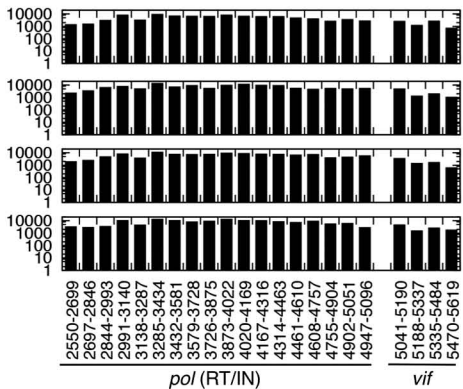


A

Sample ID#

17
82
84
180

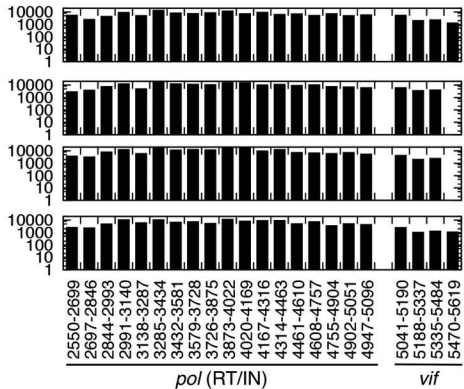
Coverage depth (fold)



Sample ID#

8
47
73
83

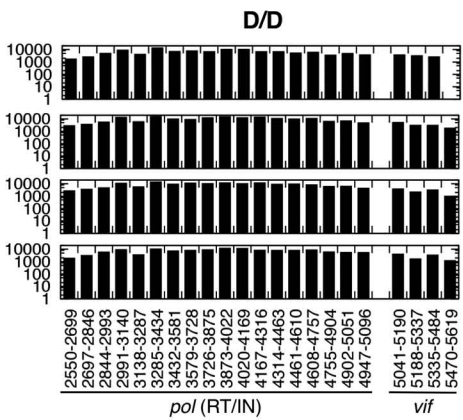
Coverage depth (fold)

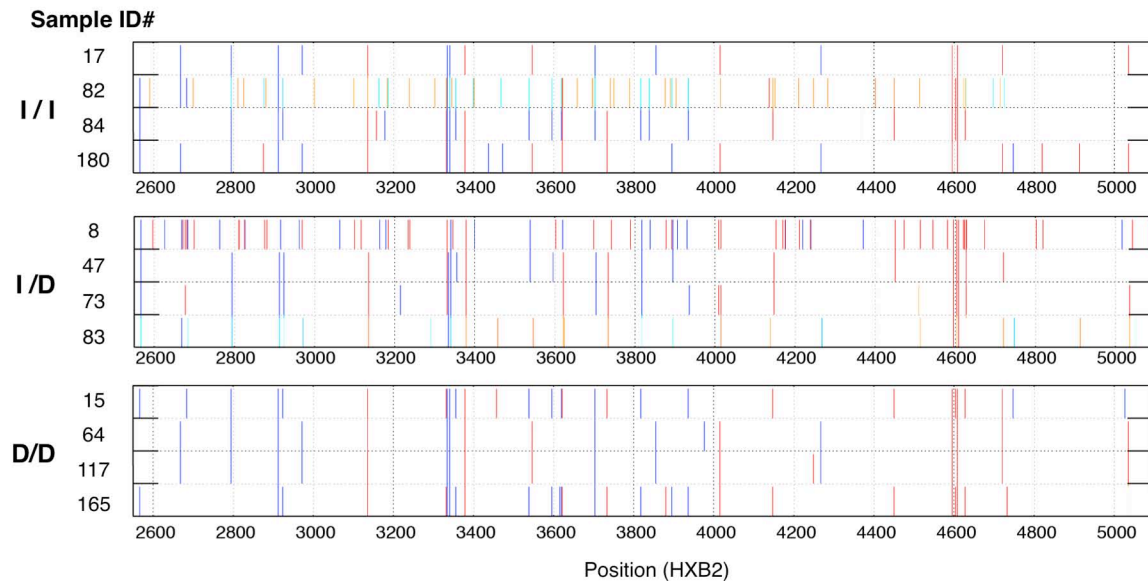
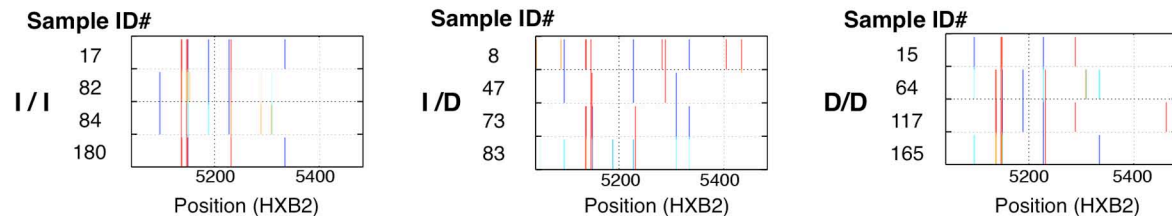


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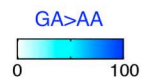
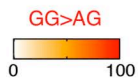
15
64
117
165

Coverage depth (fold)

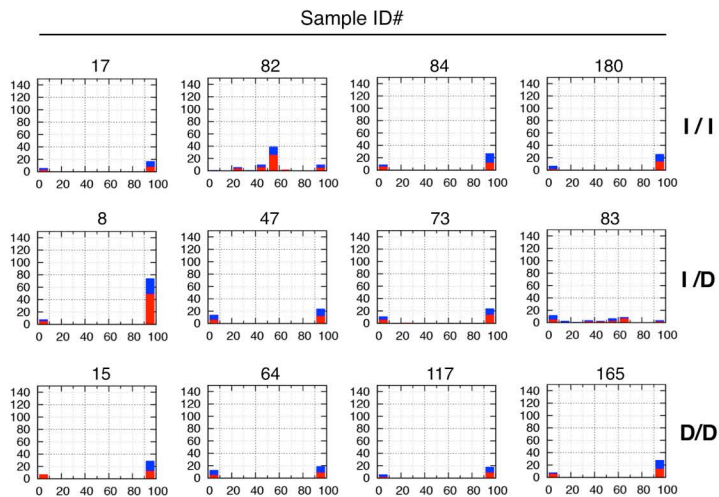
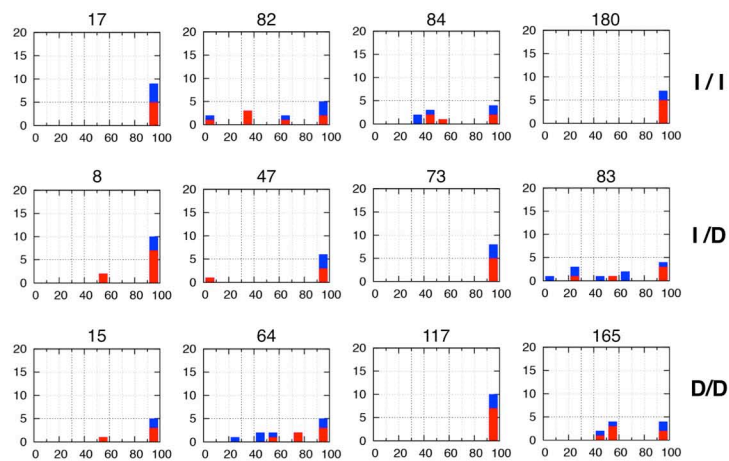


B*pol**vif*

Frequency
of hypermutations (%)



C

pol*vif*
GG>AG
GA>AA

 Number of dinucleotide sequences

 Frequency of hypermutations (%)