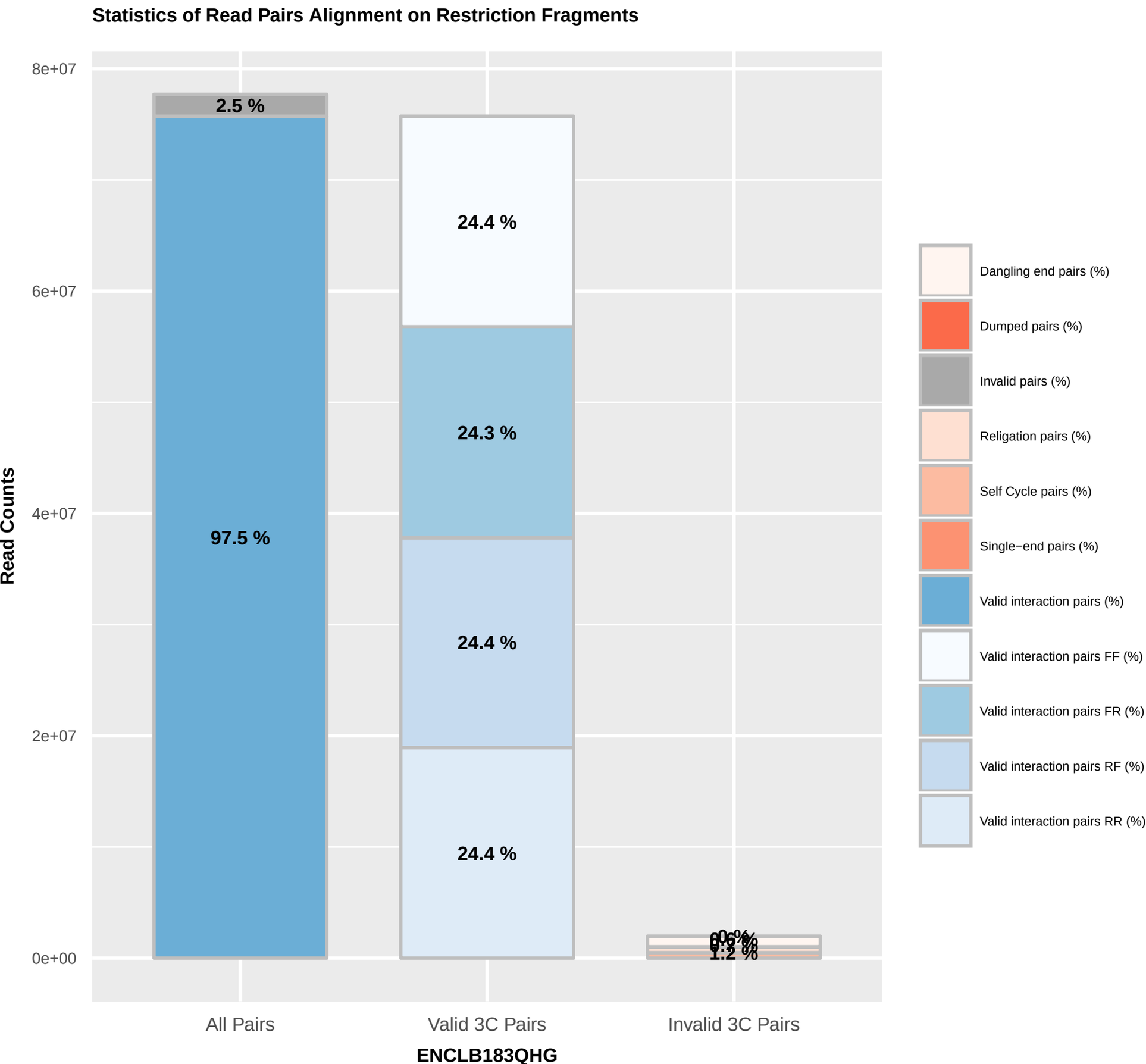
Statistics of Read Pairs Alignment on Restriction Fragments
8e+07
6e+07
4e+07
2e+07
0e+00
Read Counts
2.5 %
97.5 %
24.4 %
24.3 %
24.4 %
24.4 %
1.2 %
All Pairs
Valid 3C Pairs
Invalid 3C Pairs
ENCLB183QHG
Dangling end pairs (%)
Dumped pairs (%)
Invalid pairs (%)
Religation pairs (%)
Self Cycle pairs (%)
Single−end pairs (%)
Valid interaction pairs (%)
Valid interaction pairs FF (%)
Valid interaction pairs FR (%)
Valid interaction pairs RF (%)
Valid interaction pairs RR (%)