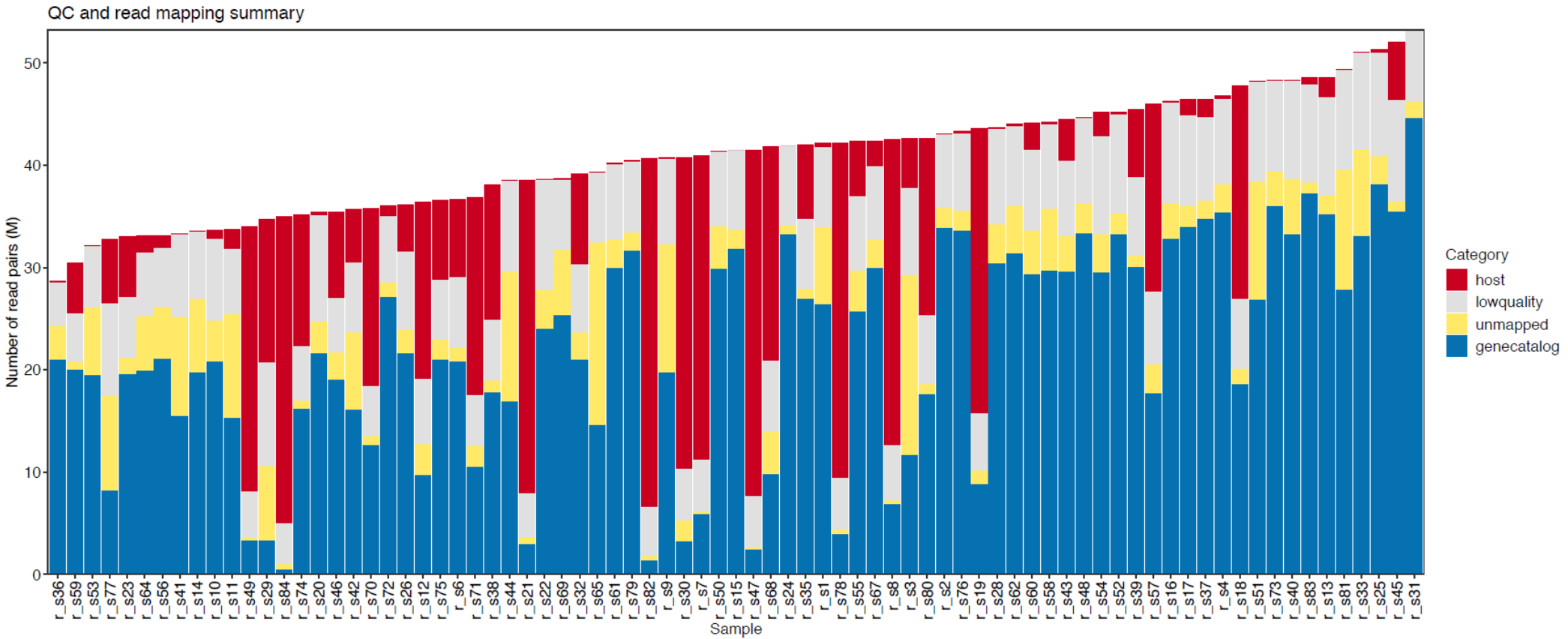

QC and read mapping summary
Number of read pairs (M)
0
10
20
30
40
50
Sample
r_s36
r_s59
r_s53
r_s77
r_s23
r_s64
r_s56
r_s41
r_s14
r_s10
r_s11
r_s49
r_s29
r_s84
r_s74
r_s20
r_s46
r_s42
r_s70
r_s72
r_s26
r_s12
r_s75
r_s6
r_s71
r_s38
r_s44
r_s21
r_s22
r_s69
r_s32
r_s65
r_s61
r_s79
r_s82
r_s9
r_s30
r_s7
r_s50
r_s15
r_s47
r_s68
r_s24
r_s35
r_s1
r_s78
r_s55
r_s67
r_s8
r_s3
r_s80
r_s2
r_s76
r_s19
r_s28
r_s62
r_s60
r_s58
r_s43
r_s48
r_s54
r_s52
r_s39
r_s57
r_s16
r_s17
r_s37
r_s4
r_s18
r_s51
r_s73
r_s40
r_s83
r_s13
r_s81
r_s33
r_s25
r_s45
r_s31
Category
host
lowquality
unmapped
genecatalog