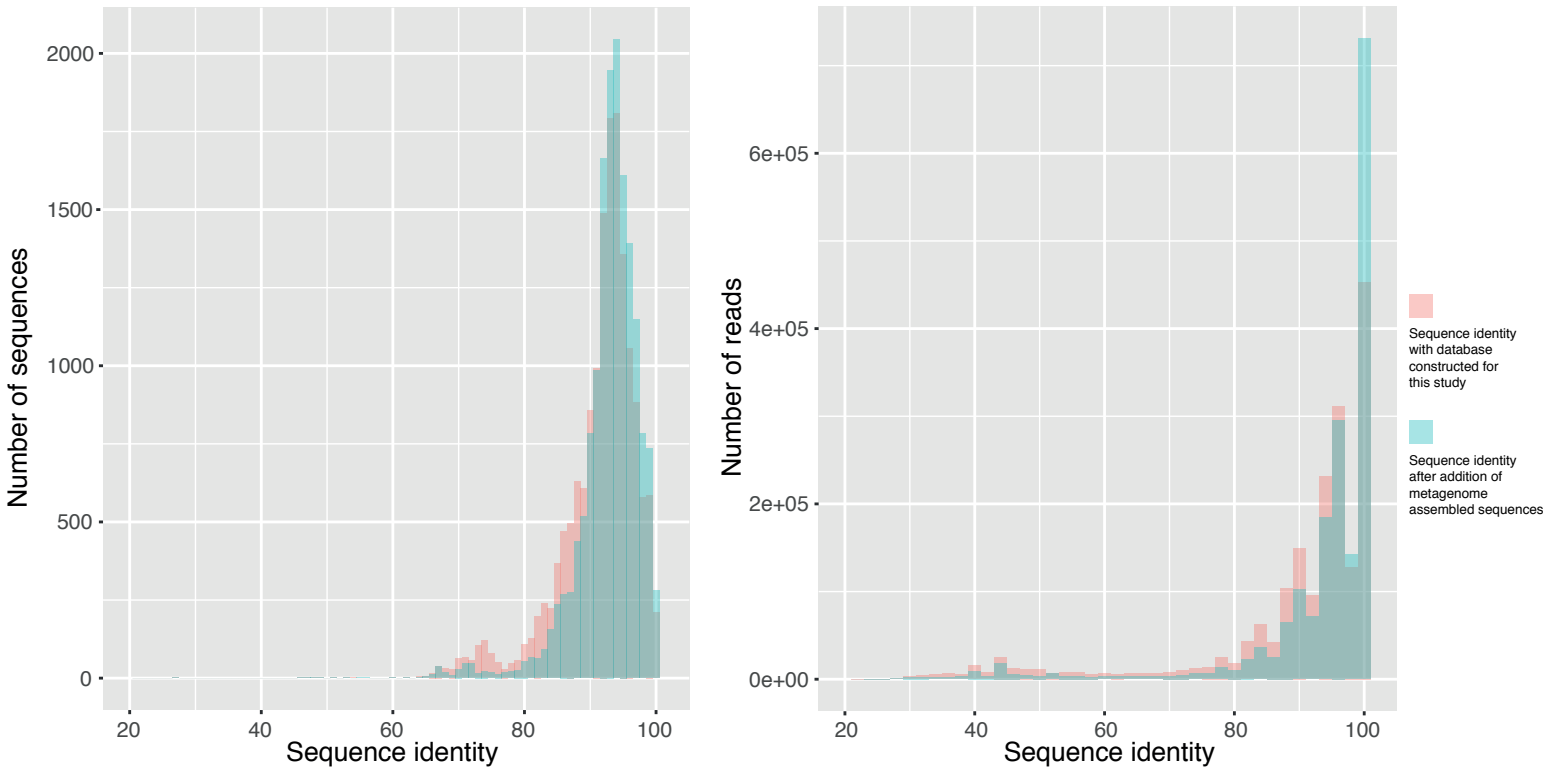
Number of sequences
2000
1500
1000
500
0
20
40
60
80
100
Sequence identity
Number of reads
6e+05
4e+05
2e+05
0e+00
20
40
60
80
100
Sequence identity
Sequence identity with database constructed for this study
Sequence identity after addition of metagenome assembled sequences