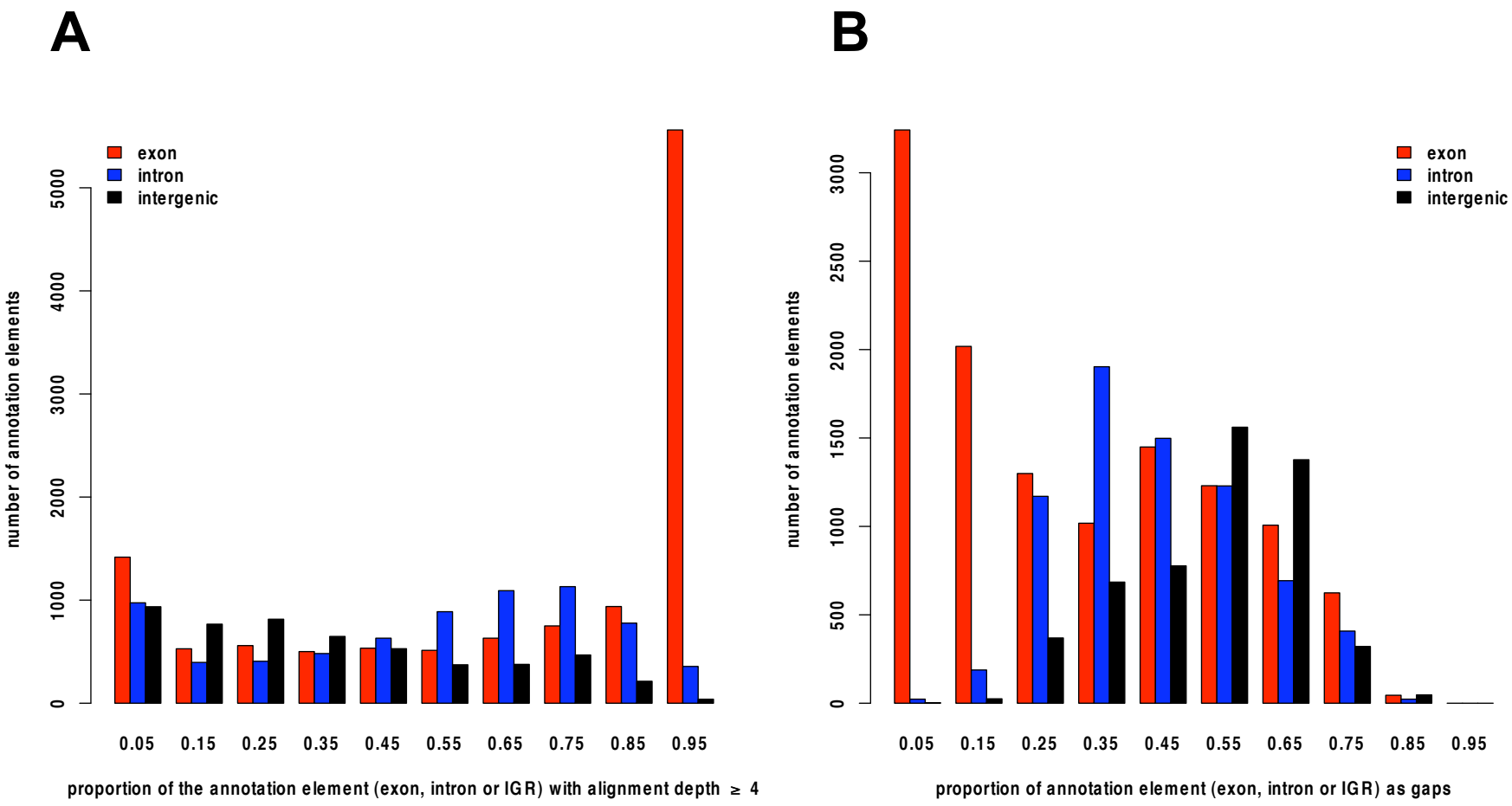
A
exon
intron
intergenic
number of annotation elements
0
1000
2000
3000
4000
5000
0.05
0.15
0.25
0.35
0.45
0.55
0.65
0.75
0.85
0.95
proportion of the annotation element (exon, intron or IGR) with alignment depth ≥ 4
B
exon
intron
intergenic
number of annotation elements
0
500
1000
1500
2000
2500
3000
0.05
0.15
0.25
0.35
0.45
0.55
0.65
0.75
0.85
0.95
proportion of annotation element (exon, intron or IGR) as gaps