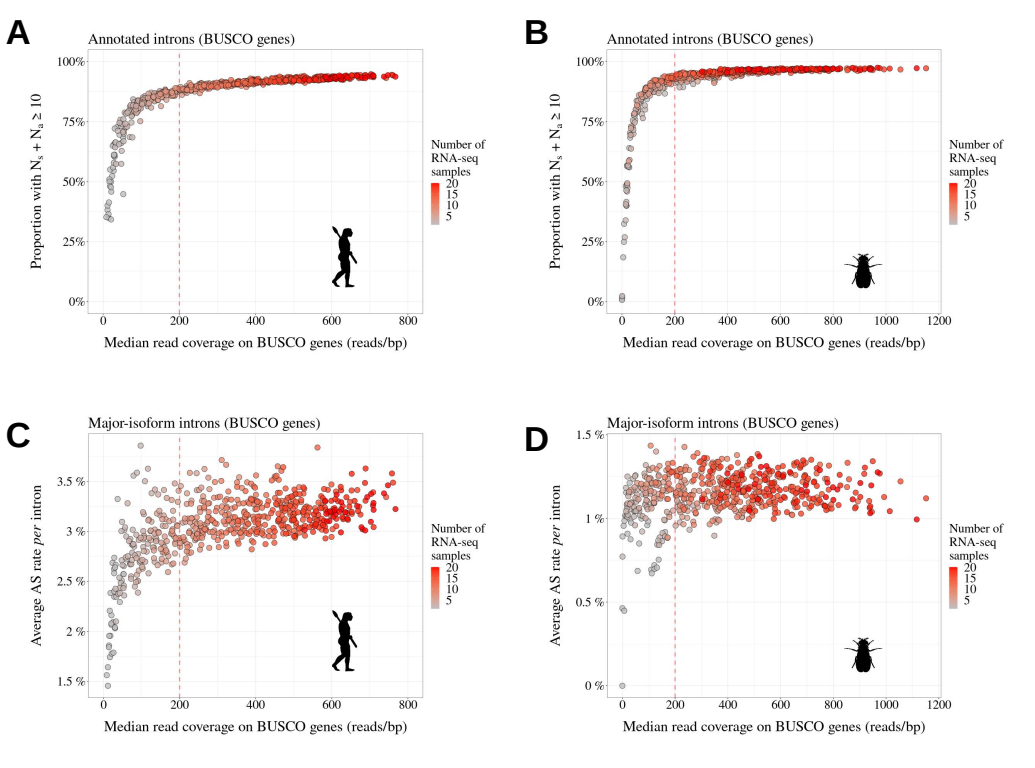
A
Annotated introns (BUSCO genes)
Proportion with $N_s + N_a \geq 10$
100%
75%
50%
25%
0%
0
200
400
600
800
Median read coverage on BUSCO genes (reads/bp)
Number of RNA-seq samples
20
15
10
5
B
Annotated introns (BUSCO genes)
Proportion with $N_s + N_a \geq 10$
100%
75%
50%
25%
0%
0
200
400
600
800
1000
1200
Median read coverage on BUSCO genes (reads/bp)
Number of RNA-seq samples
20
15
10
5
C
Major-isoform introns (BUSCO genes)
Average *AS* rate *per* intron
3.5 %
3 %
2.5 %
2 %
1.5 %
0
200
400
600
800
Median read coverage on BUSCO genes (reads/bp)
Number of RNA-seq samples
20
15
10
5
D
Major-isoform introns (BUSCO genes)
Average *AS* rate *per* intron
1.5 %
1 %
0.5 %
0 %
0
200
400
600
800
1000
1200
Median read coverage on BUSCO genes (reads/bp)
Number of RNA-seq samples
20
15
10
5