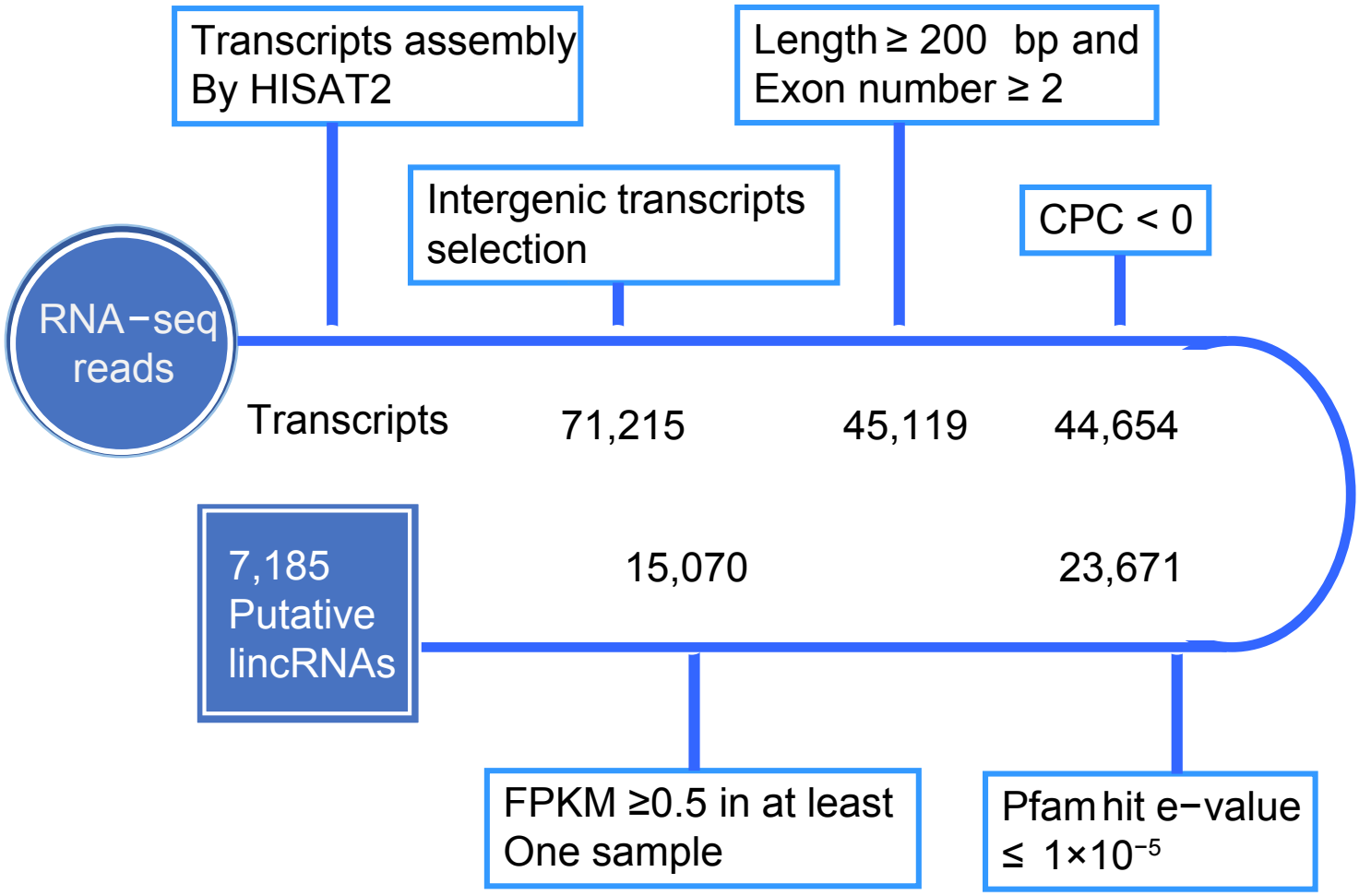
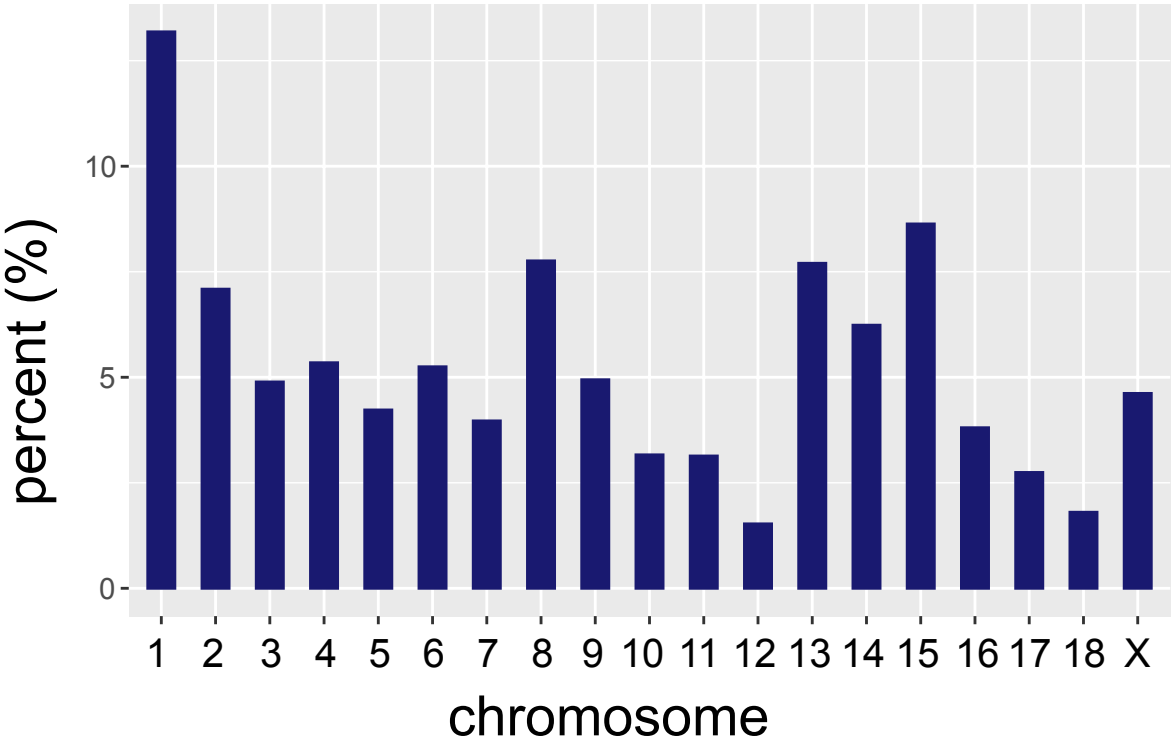
A


Transcripts assembly
By HISAT2
Length ≥ 200 bp and
Exon number ≥ 2
Intergenic transcripts
selection
CPC < 0
RNA−seq
reads
Transcripts
71,215
45,119
44,654
7,185
Putative
lincRNAs
15,070
23,671
FPKM ≥0.5 in at least
One sample
Pfam hit e−value
≤ 1×10−5

B


percent (%)
10
5
0
1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 X
chromosome