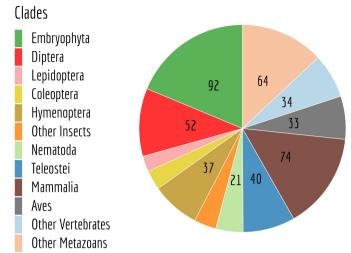
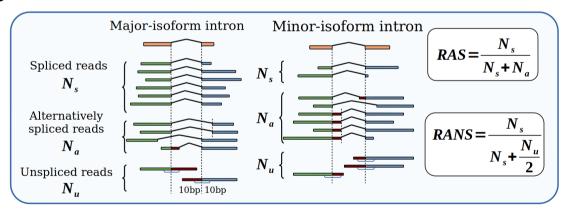


Transcriptomic data for N = 491 species



В



C

Variable	Definition
RAS	Relative abundance of a given spliced isoform compared to other overlapping spliced isoforms
RANS	Relative abundance of a given spliced isoform compared to unspliced transcripts overlaping its splice sites
Major-isoform intron	Intron corresponding to a major isoform (i.e with RAS > 0.5 and RANS > 0.5)
Minor-isoform intron / SV	Splice variant (SV) = intron corresponding to a minor isoform (i.e with RAS \leq 0.5 or RANS \leq 0.5)