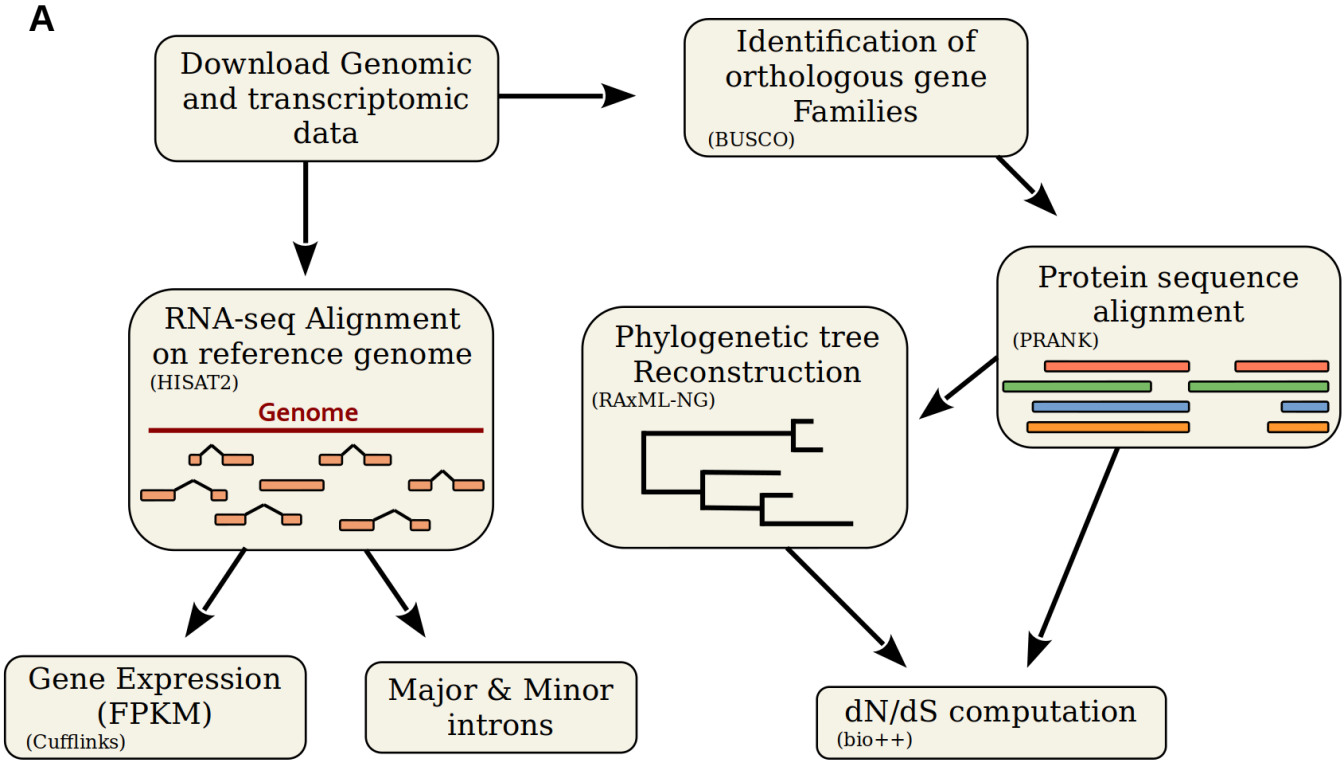
A
Download Genomic and transcriptomic data
Identification of orthologous gene Families
(BUSCO)
RNA-seq Alignment on reference genome
(HISAT2)
Genome
Phylogenetic tree Reconstruction
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(PRANK)
Gene Expression (FPKM)
(Cufflinks)
Major & Minor introns
dN/dS computation
(bio++)