

Supplemental Figure 2

A

Accuracy on simulated rna-seq data

Use nf-unsplicedcds to get empirical AUC

B

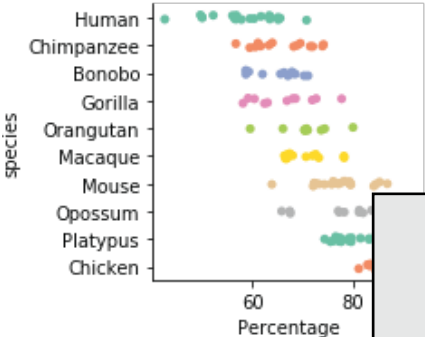
Show distribution of  
number of trans  
lated frames per read

C

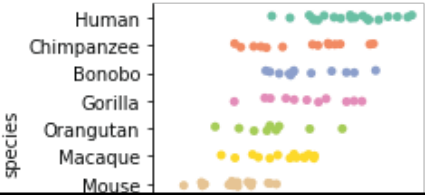
Vignette of extracting transcription  
factors only

D

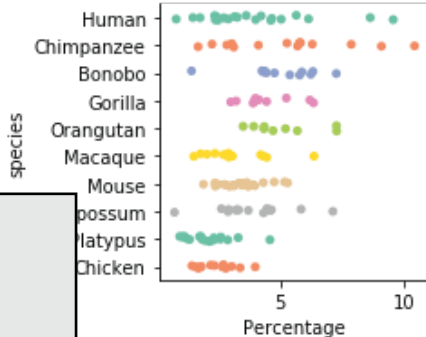
Non-coding



Coding

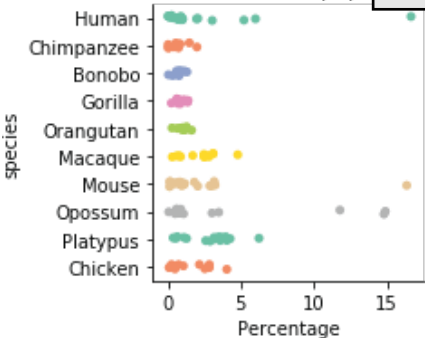


Stop codon in all translation frames



Rerun using uniprot data as reference

All translations shorter than pept



Low complexity peptide

