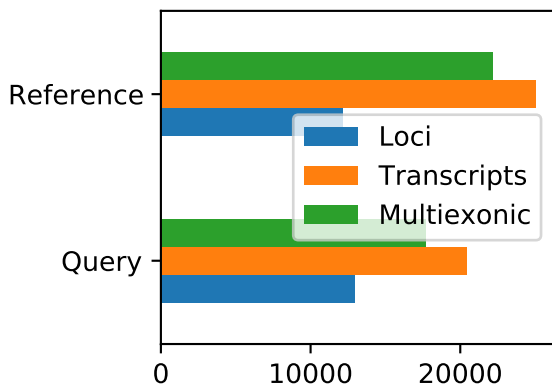
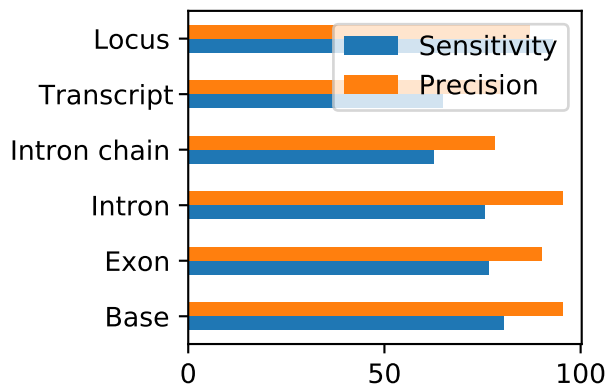


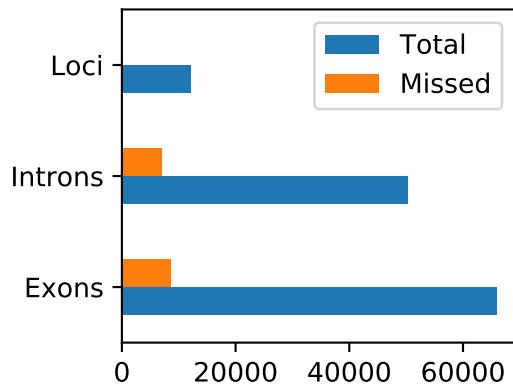
Totals



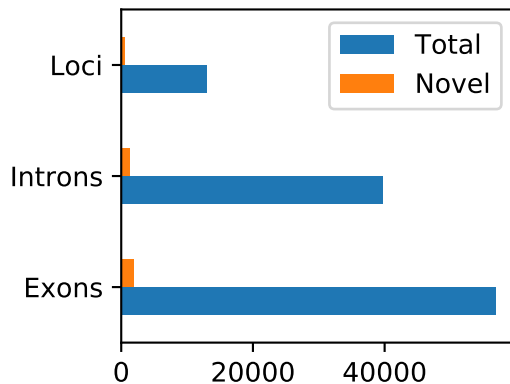
Performance



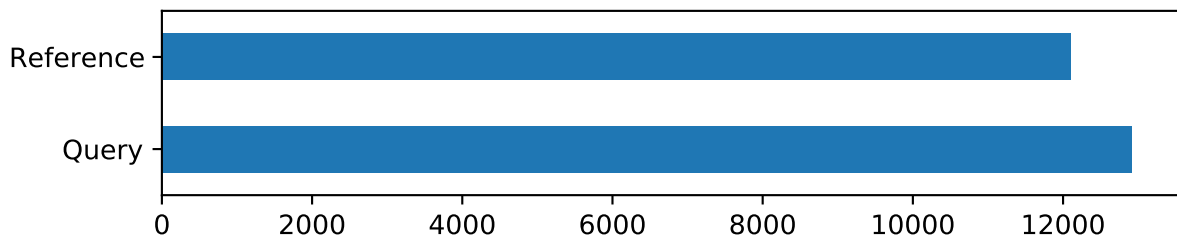
Missed



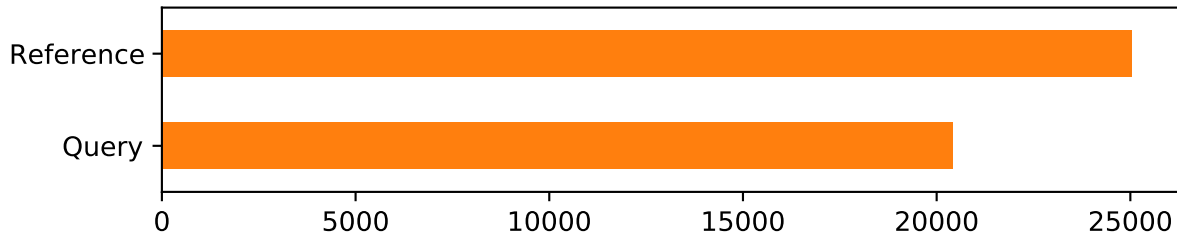
Novel



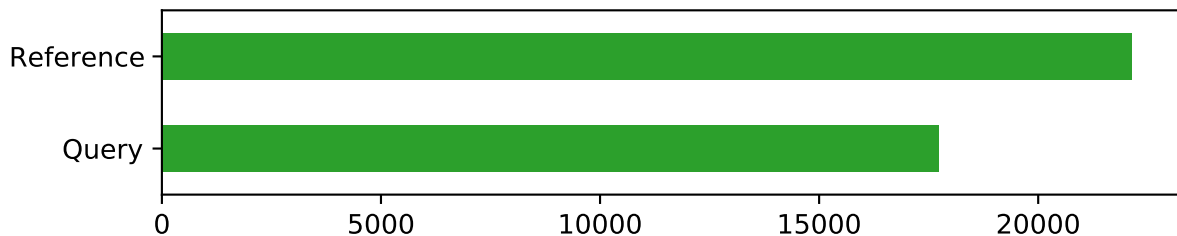
Loci



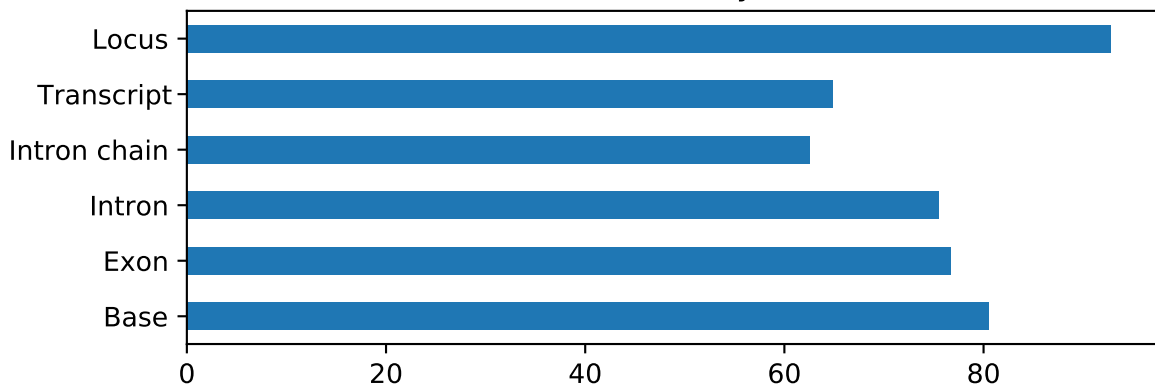
Transcripts



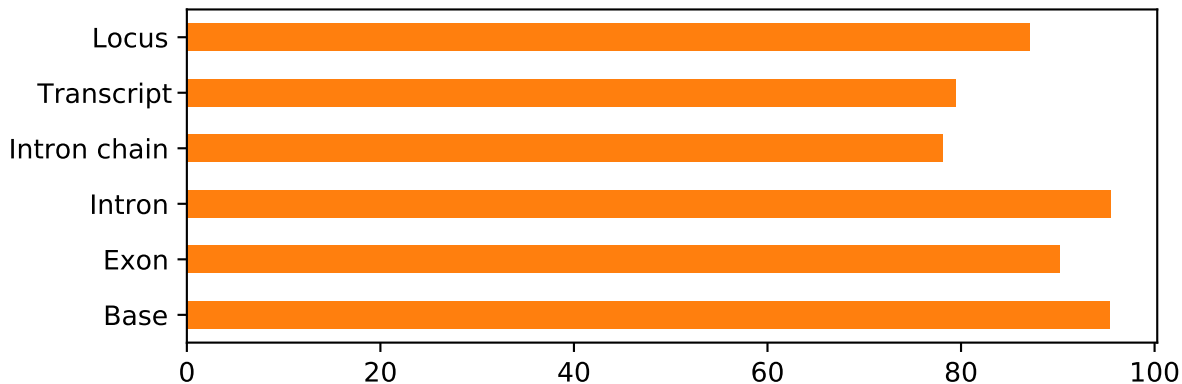
Multiexonic



Sensitivity



Precision



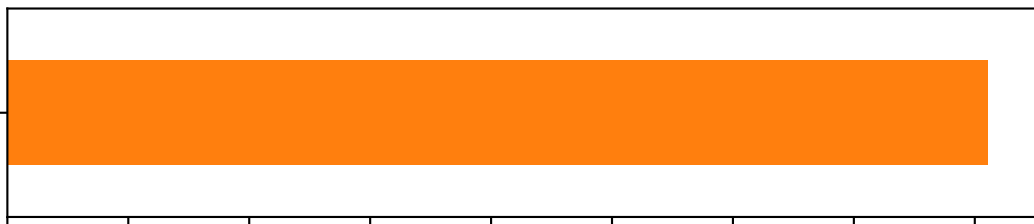
Intron chains

Matching



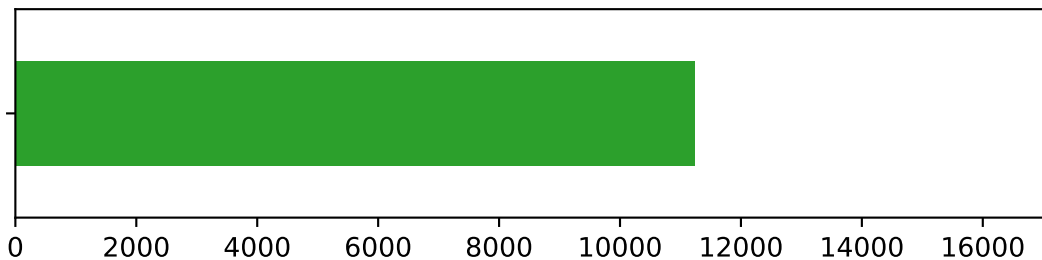
Transcripts

Matching

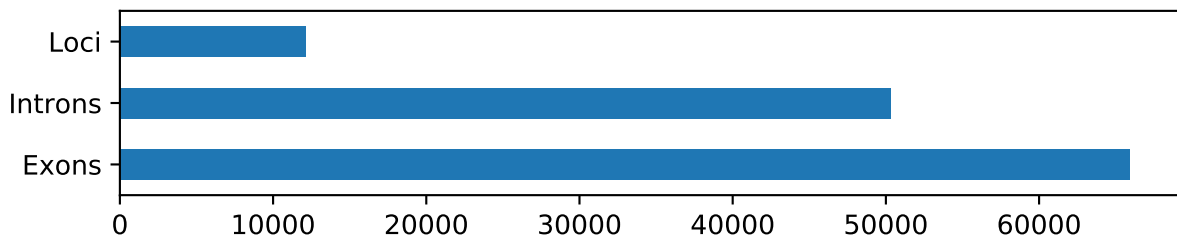


Loci

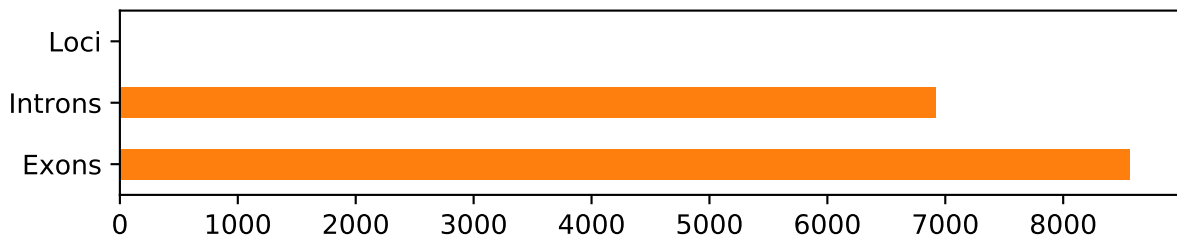
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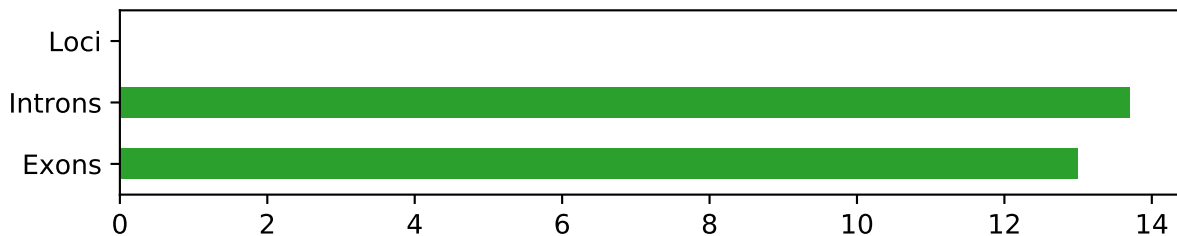
Total



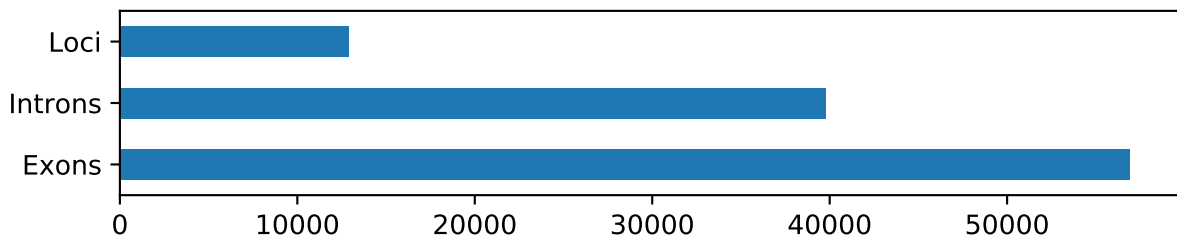
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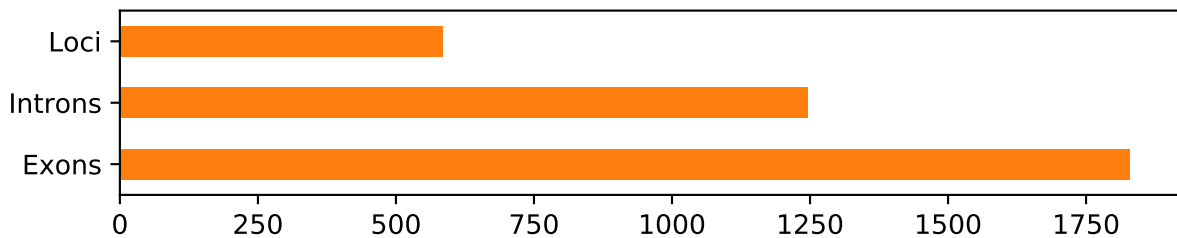
Percent missed



Total



Novel



Percent novel

