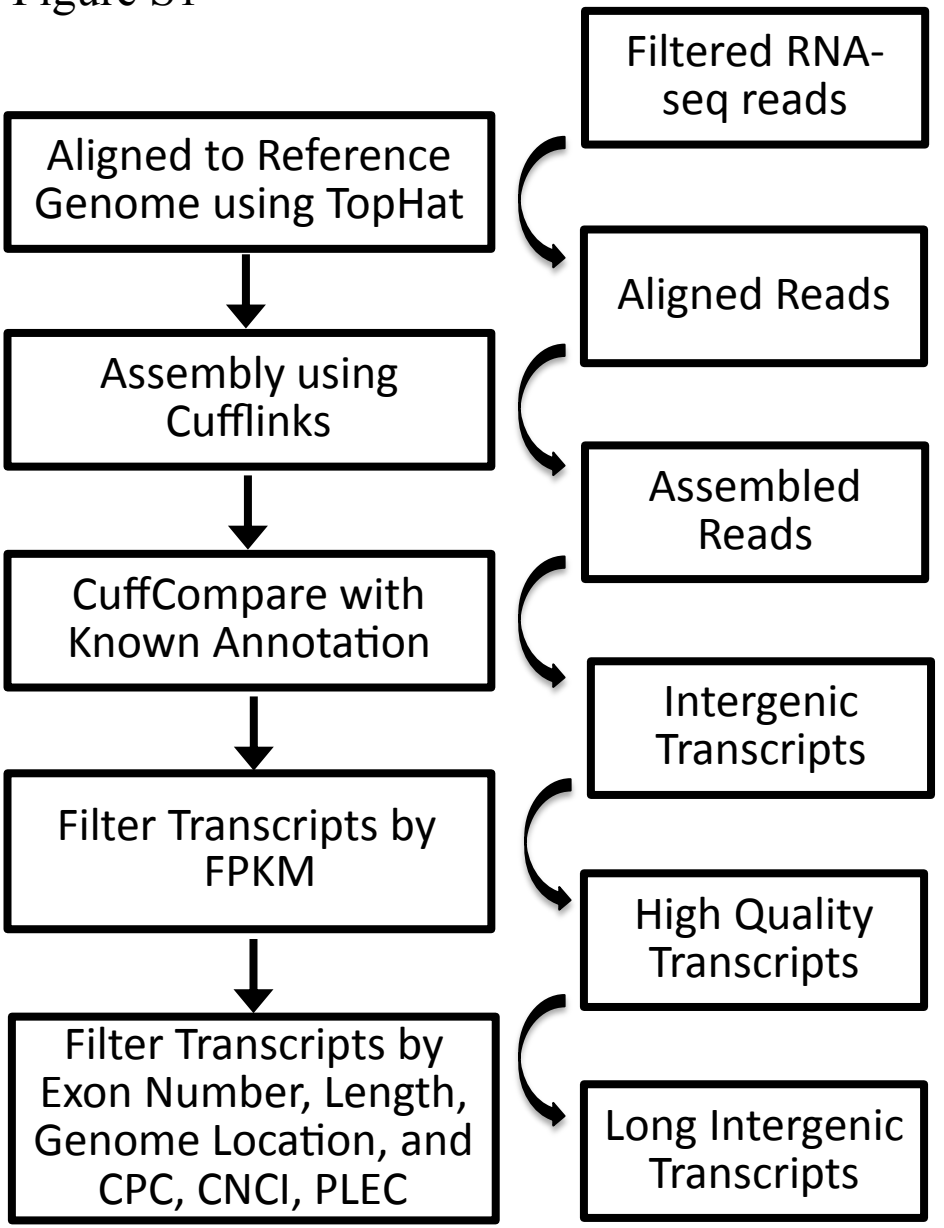
Figure S1


Filtered RNA-seq reads
Aligned to Reference Genome using TopHat
Aligned Reads
Assembly using Cufflinks
Assembled Reads
CuffCompare with Known Annotation
Intergenic Transcripts
Filter Transcripts by FPKM
High Quality Transcripts
Filter Transcripts by Exon Number, Length, Genome Location, and CPC, CNCI, PLEC
Long Intergenic Transcripts