

BED

Browser Extensible Data format introduced by UCSC genome browser

Widely used to annotate genomes with intervals of interest

Simple, tab-delimited text file

Not very extensible, so used for very simple features (e.g. enhancers)

FYI gene models typically stored in GFF or GTF format, which is much more complex.

Chromosome	Start		Stop	Feature name	“Score”	Strand
I	15062083		15063836	WBGene00004512	255	+
I	15064301		15064453	WBGene00004567	255	+
I	15064838		15068346	WBGene00004622	255	+
I	15069280		15071033	WBGene00004513	255	+
MtDNA	898	1593	WBGene00014454	255	+	
MtDNA	10403	11354	WBGene00014472	255	+	
V	17115903		17116021	WBGene00077465	255	+
V	17117863		17117981	WBGene00077466	255	+
V	17118837		17118935	WBGene00077467	255	+