

UCSC Genome Browser on D. melanogaster Aug. 2014 (BDGP Release 6 + ISO1 MT/dm6) Assembly

move

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zoom in

1.5x

3x

10x

base

zoom out

1.5x

3x

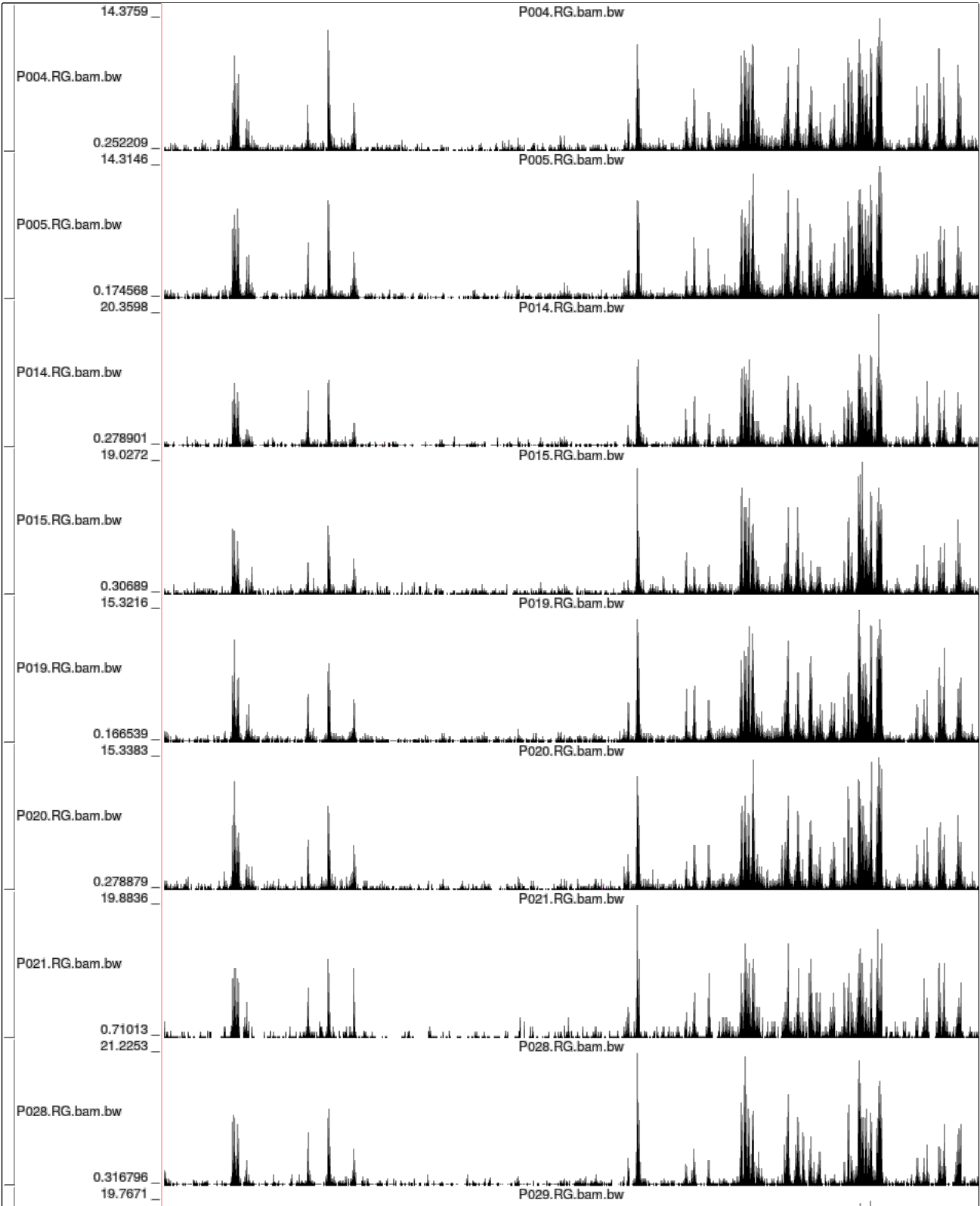
10x

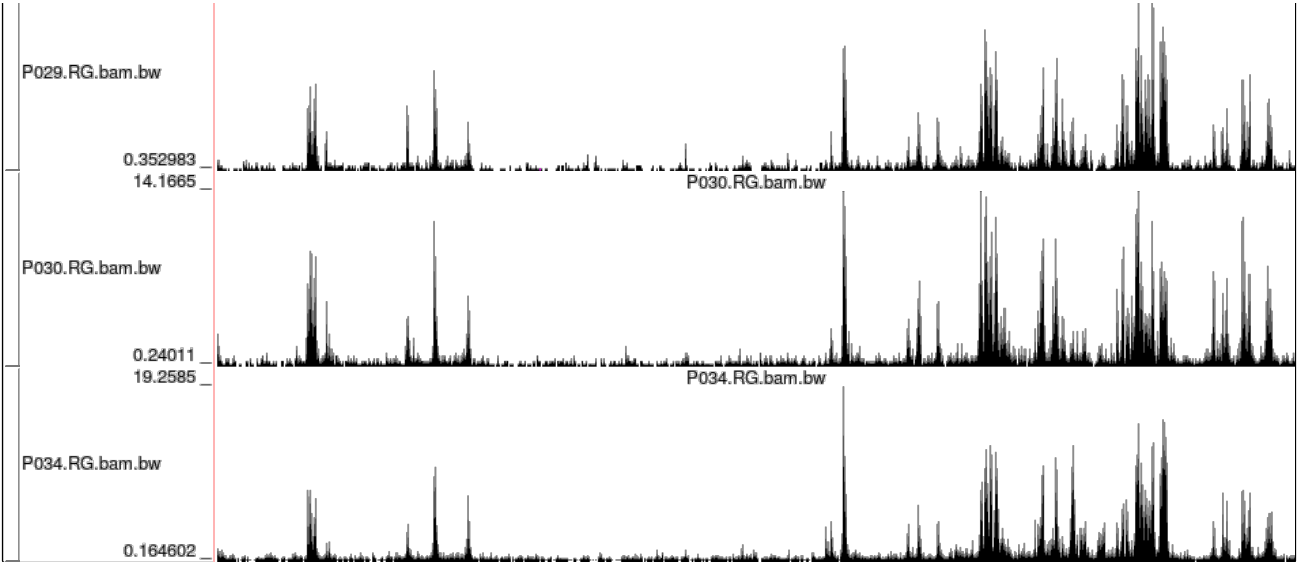
100x

chr2L:1,988,549-2,232,756244,208 bp.

enter position, gene symbol or search terms

go





move start

< 2.0 >

Click on a feature for details. Click+shift+drag to zoom in. Click side bars for track options.
Drag side bars or labels up or down to reorder tracks. Drag tracks left or right to new position.
Press "?" for keyboard shortcuts.

move end

< 2.0 >

track search default tracks default order hide all manage custom tracks track hubs configure multi-region reverse resize refresh

collapse all

Use drop-down controls below and press refresh to alter tracks displayed.
Tracks with lots of items will automatically be displayed in more compact modes.

expand all

Custom Tracks

P004.RG.bam.bw

full

P021.RG.bam.bw

full

P005.RG.bam.bw

full

P028.RG.bam.bw

full

P014.RG.bam.bw

full

P029.RG.bam.bw

full

P015.RG.bam.bw

full

P030.RG.bam.bw

full

P019.RG.bam.bw

full

P034.RG.bam.bw

full

P020.RG.bam.bw

full

refresh

Mapping and Sequencing

Base Position

hide

RefSeq Acc

hide

Chromosome Band

hide

Restr Enzymes

hide

Assembly

hide

Short Match

hide

Gap

hide

GC Percent

hide

INSDC

hide

refresh

Genes and Gene Predictions

Updated NCBI RefSeq

hide

Pfam in RefSeq

hide

Other RefSeq

hide

Updated UniProt

hide

AUGUSTUS

hide

CRISPR...

hide

Ensembl Genes

hide

Genscan Genes

hide

refresh

mRNA and EST

D. melanogaster mRNAs

hide

Spliced ESTs

hide

D. melanogaster ESTs

hide

Other mRNAs

hide

refresh

Expression and Regulation

CpG Islands...

hide

ORegAnno

hide

refresh

Comparative Genomics

Conservation

hide

Insects Chain/Net

hide

Cons 124 Insects

hide

refresh

Variation and Repeats

Interrupted Rpts

hide

Simple Repeats

hide

Microsatellite

hide

UniProt Variants

hide

WM + SDust

hide

RepeatMasker

hide

refresh