

Sox2

Sox2

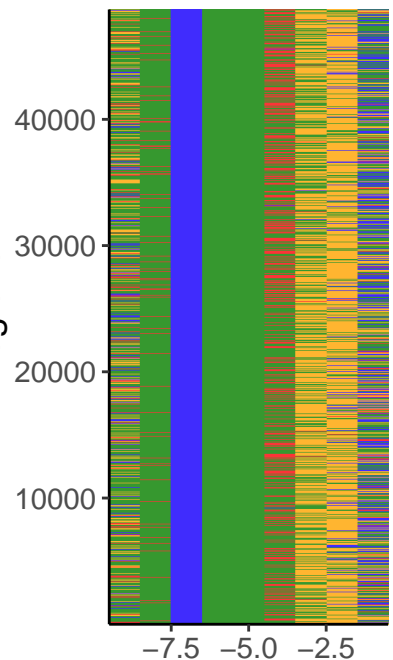
Sox2

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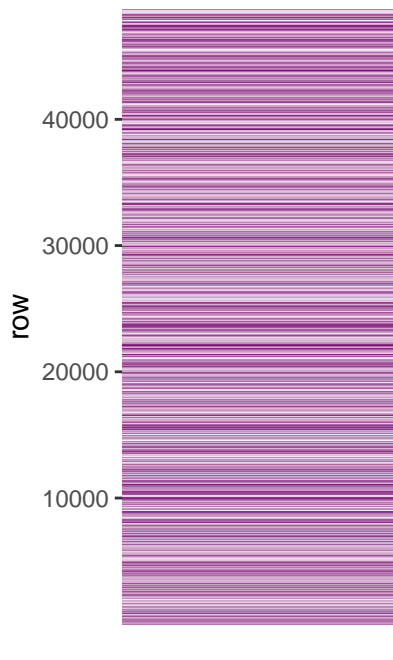
Sox2

Regions



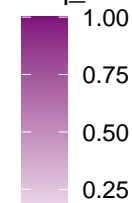
Motif distance (bp)

Context ordered

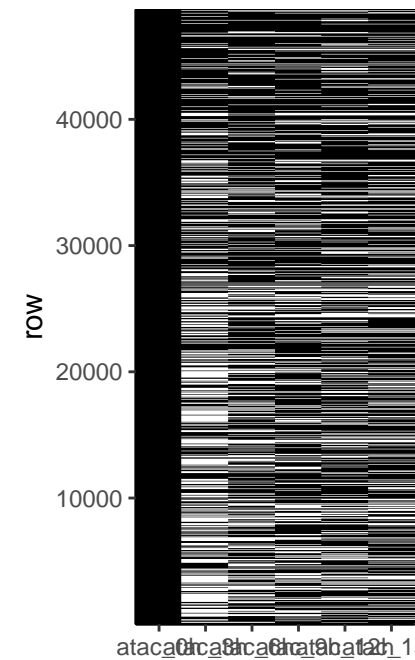


x

seq_match_quantile

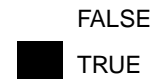


row

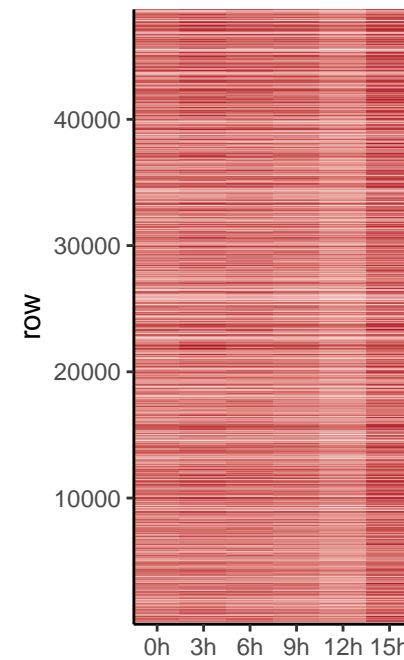


timepoint

overlap

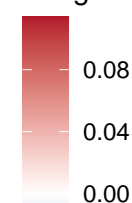


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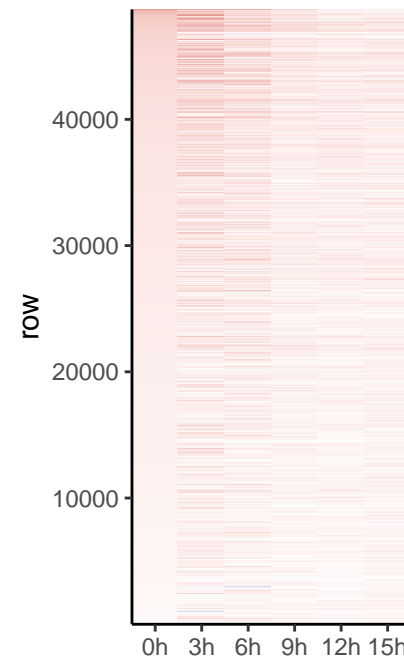


timepoint

Marg. score

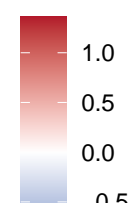


row

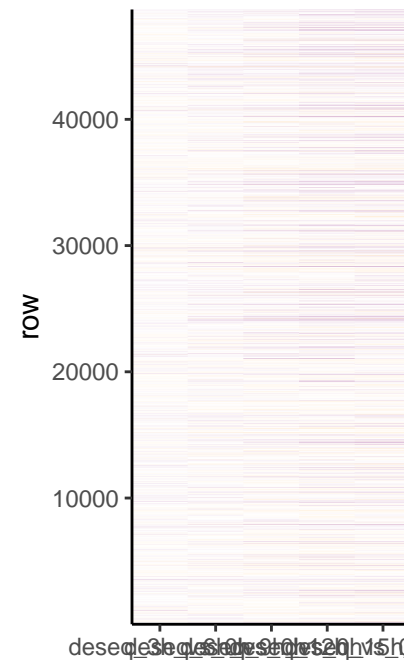


timepoint

Perturb score

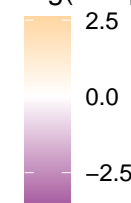


row



timepoint

log(timepoint/WT)



Nucleotide A C G T