

## WashU Epigenome Browser - Feature # 30

<b>Status:</b>	New	<b>Priority:</b>	Low
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<b>Created:</b>	08/05/2013	<b>Assignee:</b>	Xin Zhou
<b>Updated:</b>	08/05/2013	<b>Due date:</b>	
<b>Subject:</b>	bam track (full) view range limit		
<b>Description</b>			
<ul style="list-style-type: none"><li>- in UCSC browser, when you zoom out like 1/4 chromosome range, the repeatmasker tracks will show 'zoom in to &lt;= 10,000,000 bases to view items'</li><li>- I think for bam track in full mode, might we could also adopt this settings to accelerating browsing display?</li></ul>			

### History

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