

## WashU Epigenome Browser - Bug # 235

<b>Status:</b>	New	<b>Priority:</b>	Normal
<b>Author:</b>	Rebecca Lowdon	<b>Category:</b>	
<b>Created:</b>	10/16/2014	<b>Assignee:</b>	
<b>Updated:</b>	10/16/2014	<b>Due date:</b>	
<b>Subject:</b>	Gene shading feature inconsistent and needs to be made optional		
<b>Description</b>			
<p>I guess there is a new feature where genes you relocate to are shaded? I can see the value in this, but it needs to be made optional!</p> <p>Also it seems to be inconsistent and dependent on gene size:</p> <ol style="list-style-type: none"><li>1. When I relocate to a gene (I've been testing TFAP2C, SFN, KRT5 this morning) it is NOT shaded. Once I zoom out it becomes shaded, but the shorter genes (KRT5 and SFN) are much darker than the longer ones (TFAP2C, CASZ1). Then again, I think it is more related to how much you zoom out.</li><li>2. But only sometimes when I zoom out does it become shaded. The feature seems pretty unstable.</li></ol> <p>My session: <a href="http://epigenomegateway.wustl.edu/browser/?genome=hg19&amp;#38;session=aPGTwll8Yx">http://epigenomegateway.wustl.edu/browser/?genome=hg19&amp;#38;session=aPGTwll8Yx</a> You can delete this session when finished.</p>			

### History

Files			
Screen Shot 2014-10-16 at 9.43.50 AM.png	15 kB	10/16/2014	Rebecca Lowdon
Screen Shot 2014-10-16 at 9.51.28 AM.png	59.8 kB	10/16/2014	Rebecca Lowdon