

## WashU Epigenome Browser - Bug # 245

<b>Status:</b>	Resolved	<b>Priority:</b>	Normal
<b>Author:</b>	Junchen Gu	<b>Category:</b>	
<b>Created:</b>	11/14/2014	<b>Assignee:</b>	
<b>Updated:</b>	11/24/2014	<b>Due date:</b>	
<b>Subject:</b>	Missing display of 0% methylated Cs in methylC track		
<b>Description</b>			
<p>Currently, when a CpG has a 0% methylation, with read coverage, there is no grey bar to indicate 0% methylation.</p> <p>E.g. go to this hub: <a href="http://epigenomegateway.wustl.edu/browser/?genome=mm9&amp;#38;datahub_jsonfile=http://remc.wustl.edu/jgu/mouse/8w/tracks/8w_ouse_hub.json">http://epigenomegateway.wustl.edu/browser/?genome=mm9&amp;#38;datahub_jsonfile=http://remc.wustl.edu/jgu/mouse/8w/tracks/8w_ouse_hub.json</a></p> <p>For the first C in chr4 66488665 64488667, there is an entry in bedGraph file</p> <pre>chr4 66488665 66488666 0.0000</pre> <p>But there is no display on the browser. See attached image.</p>			

### History

#1 - 11/24/2014 01:40 pm - Xin Zhou

- Status changed from New to Resolved

Fixed in v39.3.5

Please check

### Files

methylCtrack no 0 meth display.png	43.9 kB	11/14/2014	Junchen Gu
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