

WashU Epigenome Browser - Bug # 73

Status:	Closed	Priority:	Normal
Author:	Daofeng Li	Category:	
Created:	10/23/2013	Assignee:	
Updated:	11/18/2013	Due date:	
Subject:	bam track in json hub set to show cause syntax error		
Description	<ul style="list-style-type: none">- Use bam track in a json hub, the hub works when all bam tracks set to 'hide' mode- while any of the bam track set to 'show', a syntax error happen- could use this for example: http://budweiser/example/H1Es_cmp_Brian_4/hub.json		

History

#1 - 11/18/2013 12:03 pm - Xin Zhou

- Status changed from New to Resolved

Fixed in v27.3 on test browser

In the datahub, if a bam track is set to "show", it will be shown as a read density track when the hub is loaded.

#2 - 11/18/2013 10:01 pm - Daofeng Li

- Status changed from Resolved to Closed

fixed