

## WashU Epigenome Browser - Bug # 91

<b>Status:</b>	Closed	<b>Priority:</b>	Normal
<b>Author:</b>	Xin Zhou	<b>Category:</b>	
<b>Created:</b>	11/05/2013	<b>Assignee:</b>	
<b>Updated:</b>	11/18/2013	<b>Due date:</b>	
<b>Subject:</b>	crashed when loading bam track		
<b>Description</b>	try loading a bam track from submission panel, crashes		

### History

#### #1 - 11/05/2013 07:29 pm - Xin Zhou

- Status changed from New to Resolved

Fixed in v26.2 on PUBLIC browser, still there on the test for you to see it

the error was because the code was trying to access .qtc.bedcolor which is annotation track color that does not exist for bam track

#### #2 - 11/18/2013 09:58 pm - Daofeng Li

- Status changed from Resolved to Closed

- % Done changed from 0 to 100

- tested loading custom bam file, works! Closed!