

WashU Epigenome Browser - Bug # 91

Status:	Closed	Priority:	Normal
Author:	Xin Zhou	Category:	
Created:	11/05/2013	Assignee:	
Updated:	11/18/2013	Due date:	
Subject:	crashed when loading bam track		
Description			
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!