

WashU Epigenome Browser - Bug # 233

Status:	Resolved	Priority:	Normal
Author:	Rebecca Lowdon	Category:	
Created:	10/15/2014	Assignee:	
Updated:	10/15/2014	Due date:	
Subject:	dbSNP track crashes		
Description			
Loaded the dbSNP track on a wide view range. The browser thinks the track loaded, but no data are shown. (image 1)			
Change to "density" mode and there is signal (image 2). But when I change back to "full mode" still no data appear.			

History

#1 - 10/15/2014 04:55 pm - Rebecca Lowdon

When I zoom in on a region, data appear in "full" mode. Maybe create a warning if there is too much data in a view range to show them all?

#2 - 10/15/2014 08:04 pm - Xin Zhou

- Status changed from New to Resolved

Fixed on both test and public server. If you open dbSNP track when view range is greater than 20kb, the track is automatically shown in "density" mode.
But try not to turn this track right into "full" mode since it may halt the browser.

Files

Screen Shot 2014-10-15 at 4.50.08 PM.png	107.9 kB	10/15/2014	Rebecca Lowdon
Screen Shot 2014-10-15 at 4.51.01 PM.png	25.6 kB	10/15/2014	Rebecca Lowdon