WashU Epigenome Browser - Bug # 137

Status:	Rejected	**Priority:**	Normal
Author:	Rebecca Lowdon	**Category:**	
Created:	01/15/2014	**Assignee:**	Xin Zhou
Updated:	01/16/2014	**Due date:**	
Subject:	cutsom BED track not displayed		

Description

Custom bed tracks are not displayed properly when added to a session with a data hub.

First loaded data hub: http://cgs.wustl.edu/~dli/placenta.json

Then added two bed tracks: http://cgs.wustl.edu/~rlowdon/placentaDMRs/MUU.bed.gz and

http://cgs.wustl.edu/~rlowdon/placentaDMRs/UMM.bed.gz

Genome snapshot of these two tracks shows the data are there and detected by the browser (see image). But when I zoom in to a region, nothing is displayed in the track. For example, I selected chr4 p14 from the genome snapshot, which looks to have many annotations there, but no data appear on the browser.

History

#1 - 01/16/2014 07:48 am - Xin Zhou

- Status changed from New to Rejected

Sorry but for a bed track to be displayed, it must have at least 5 fields:

4th - name, if no name, use dot or leave empty

5th - unique integer id

Though a bed track only needs its first 3 fields to be shown as density track as in genome snapshot.

Files

genomesnapshot.png	177.3 kB	01/15/2014	Rebecca Lowdon
nodata.png	153.4 kB	01/15/2014	Rebecca Lowdon