WashU Epigenome Browser - Bug # 6

Status:	Closed	**Priority:**	Normal
Author:	Daofeng Li	**Category:**	
Created:	07/19/2013	**Assignee:**	Xin Zhou
Updated:	07/22/2013	**Due date:**	

Subject:	scatter plot shared between genomes and caused Json syntax error

Description

- steps to reproduce
    1. open hg19
    2. add danRer7
    3. in danRer7, submit the default genelist as gene set
    4. in danRer7, choose scatter plot
    5. go back to hg19, choose scatter plot, the plot was already there
    6. if choose any 2 track from hg19, click submit will raise the Json syntax error
    7. see attached figure

- if re-submit gene set list, then it works fine

History

#1 - 07/22/2013 09:33 am - Xin Zhou

- Status changed from New to Resolved

#2 - 07/22/2013 02:09 pm - Xin Zhou

- Status changed from Resolved to Closed

Files

071913-1.png	147.3 kB	07/19/2013	Daofeng Li