

WashU Epigenome Browser - Issues

#	Tracker	Status	Priority	Subject	Assignee	Updated
252	Bug	New	Normal	json syntax error when loading genome comparison tracks		03/10/2015 04:21 pm
250	Bug	New	Normal	slow loading of big hub		12/19/2014 03:45 pm
249	Feature	New	Normal	track hub include feature		12/18/2014 05:24 pm
231	Bug	Resolved	Normal	load hub and session issue		10/15/2014 11:15 am
204	Bug	In Progress	Normal	gene set view not working		07/07/2014 03:04 pm
198	Bug	New	Normal	refresh cache issue		06/18/2014 09:51 am
194	Bug	New	Normal	juxtaposition issue on genome align tracks		05/25/2014 12:09 am
190	Bug	New	Normal	geneplot on methylC track		04/28/2014 12:11 am
179	Bug	New	Normal	refGene track automatically be put to bottom when split panel		03/19/2014 11:39 pm
168	Bug	New	Normal	'pack and hide' status not saved in session		03/01/2014 12:59 am
163	Bug	New	Normal	long responsive time while track missed in hub		02/18/2014 03:39 pm
157	Bug	New	Normal	refSeq gene position didn't keep in session		02/14/2014 04:43 pm
135	Feature	New	Normal	adjust track order in genome snapshot		01/05/2014 11:11 pm
134	Feature	New	Normal	Track configuration on ipad		01/01/2014 10:09 pm
118	Feature	New	Normal	search auto completion by right clicking from gene tracks		11/29/2013 11:11 pm
117	Feature	New	Normal	add track name to genome snapshot		11/29/2013 11:10 pm
110	Feature	New	Normal	hub check button		11/20/2013 03:13 pm
99	Feature	New	Normal	function that move track(s) to top or bottom		11/07/2013 05:36 pm
97	Feature	New	Normal	user defined track color		11/07/2013 05:33 pm
80	Bug	New	Normal	session with lineplot TypeError		10/26/2013 11:45 pm
79	Bug	New	Normal	browser background color		10/26/2013 11:40 pm
78	Bug	New	Normal	2nd panel lineplot not applied to main panel		10/26/2013 11:30 pm
77	Bug	New	Normal	track background color not applied on 2nd panel of line plot		10/26/2013 11:25 pm
55	Feature	New	Normal	[System] killing subtleKnife process which had been stucked longer than 10min		09/26/2013 09:14 am
52	Feature	New	Normal	Improve gene set view		09/21/2013 07:14 pm
51	Feature	New	Normal	Improve gene search function		09/21/2013 07:14 pm
50	Feature	New	Normal	Improve session?		09/21/2013 07:13 pm
49	Feature	New	Normal	general suggestions from UCSF		09/21/2013 07:13 pm
48	Feature	New	Normal	automatically refresh cache		09/18/2013 10:59 pm
33	Feature	New	Low	batch GEO id output		08/20/2013 10:57 pm
30	Feature	New	Low	bam track (full) view range limit	Xin Zhou	08/05/2013 04:50 pm
28	Feature	New	Low	add time information for session and status	Xin Zhou	08/03/2013 11:43 pm

25	Feature	New	Normal	Quick capture of the central canvas	Daofeng Li	08/02/2013 07:49 am
21	Feature	New	Low	operation history	Xin Zhou	07/26/2013 02:47 pm
17	Feature	New	Low	public datahub unload button?	Xin Zhou	07/25/2013 09:42 pm
11	Feature	New	Low	ability to cancel a super-enhancer job	Xin Zhou	07/22/2013 05:40 pm
10	Feature	New	Low	gene set list in hub support url	Xin Zhou	07/22/2013 05:35 pm
3	Bug	New	Normal	Safari 5 navigation display disorder		07/18/2013 01:34 pm