

WashU Epigenome Browser - Issues

#	Tracker	Status	Priority	Subject	Assignee	Updated
10	Feature	New	Low	gene set list in hub support url	Xin Zhou	07/22/2013 05:35 pm
11	Feature	New	Low	ability to cancel a super-enhancer job	Xin Zhou	07/22/2013 05:40 pm
13	Feature	New	Normal	Line plot	Xin Zhou	09/22/2013 02:50 pm
17	Feature	New	Low	public datahub unload button?	Xin Zhou	07/25/2013 09:42 pm
21	Feature	New	Low	operation history	Xin Zhou	07/26/2013 02:47 pm
25	Feature	New	Normal	Quick capture of the central canvas	Daofeng Li	08/02/2013 07:49 am
28	Feature	New	Low	add time information for session and status	Xin Zhou	08/03/2013 11:43 pm
30	Feature	New	Low	bam track (full) view range limit	Xin Zhou	08/05/2013 04:50 pm
33	Feature	New	Low	batch GEO id output		08/20/2013 10:57 pm
48	Feature	New	Normal	automatically refresh cache		09/18/2013 10:59 pm
49	Feature	New	Normal	general suggestions from UCSF		09/21/2013 07:13 pm
50	Feature	New	Normal	Improve session?		09/21/2013 07:13 pm
51	Feature	New	Normal	Improve gene search function		09/21/2013 07:14 pm
52	Feature	New	Normal	Improve gene set view		09/21/2013 07:14 pm
55	Feature	New	Normal	[System] killing subtleKnife process which had been stuck longer than 10min		09/26/2013 09:14 am
97	Feature	New	Normal	user defined track color		11/07/2013 05:33 pm
99	Feature	New	Normal	function that move track(s) to top or bottom		11/07/2013 05:36 pm
108	Feature	New	Normal	compute pearson correlation in scatterplot		11/19/2013 08:06 am
110	Feature	New	Normal	hub check button		11/20/2013 03:13 pm
117	Feature	New	Normal	add track name to genome snapshot		11/29/2013 11:10 pm
118	Feature	New	Normal	search auto completion by right clicking from gene tracks		11/29/2013 11:11 pm
134	Feature	New	Normal	Track configuration on ipad		01/01/2014 10:09 pm
135	Feature	New	Normal	adjust track order in genome snapshot		01/05/2014 11:11 pm
158	Feature	New	Low	wrapping for tooltip read information		02/13/2014 05:59 pm
161	Feature	New	Normal	Include scaffolds on genome snapshot	Xin Zhou	02/13/2014 03:32 pm
242	Feature	New	Normal	Please add the option to remove genome comparison track		11/07/2014 09:37 am
244	Feature	New	Normal	Pop-up window for confirming zooming in on a region	Xin Zhou	11/09/2014 11:39 pm
246	Feature	New	Low	Change names of custom tracks		12/17/2014 04:14 pm
247	Feature	New	Low	Multiple upload of custom tracks		12/17/2014 04:16 pm
249	Feature	New	Normal	track hub include feature		12/18/2014 05:24 pm