

## WashU Epigenome Browser - Issues

#	Tracker	Status	Priority	Subject	Assignee	Updated
244	Feature	New	Normal	Pop-up window for confirming zooming in on a region	Xin Zhou	11/09/2014 11:39 pm
243	Bug	New	Normal	Incorrect SNP coordinate based on SNP ID	Xin Zhou	11/07/2014 10:43 am
240	Bug	Resolved	Normal	can't zoom out in gene set view in mm9	Xin Zhou	10/28/2014 06:46 am
237	Bug	New	Normal	Potential bug in the json hub name?	Xin Zhou	10/17/2014 06:08 pm
193	Bug	New	Normal	Json syntax error when trying to jump to a new location	Xin Zhou	05/14/2014 12:42 pm
161	Feature	New	Normal	Include scaffolds on genome snapshot	Xin Zhou	02/13/2014 03:32 pm
115	Bug	New	Normal	tracks do not relocate and some disappear in GSV	Xin Zhou	11/23/2013 04:19 pm
46	Bug	New	Normal	Coordinate issue in hg19	Xin Zhou	09/19/2013 10:16 am
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
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
13	Feature	New	Normal	Line plot	Xin Zhou	09/22/2013 02:50 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