

WashU Epigenome Browser

	2024-9				2024-10				2024-11				2024-12				2025-1					2025-2				
	36	37	38	39	40	41	42	43	44	45	46	47	48	49	50	51	52	1	2	3	4	5	6	7	8	
WashU Epigenome Browser	WashU Epigenome Browser																									
Safari 5 navigation display disorder																										
gene set list in hub support url																										
ability to cancel a super-enhancer job																										
Line plot																										
public datahub unload button?																										
operation history																										
Quick capture of the central canvas																										
add time information for session and status																										
bam track (full) view range limit																										
batch GEO id output																										
Coordinate issue in hg19																										
automatically refresh cache																										
general suggestions from UCSF																										
Improve session?																										
Improve gene search function																										
Improve gene set view																										
[System] killing subtleKnife process which (...)																										
"donor" terms in hg19 metadata																										
track background color not applied on 2nd (...)																										
2nd panel lineplot not applied to main panel																										
browser background color																										
session with lineplot TypeError																										
user defined track color																										
function that move track(s) to top or bottom																										
compute pearson correlation in scatterplot																										
hub check button																										
Tracks disappear when using config panel																										

tracks do not relocate and some disappear (...)
add track name to genome snapshot
search auto completion by right clicking (...)
Track configuration on ipad
adjust track order in genome snapshot
Session cannot save more than 1 custom metadata (...)
refSeq gene position didn't keep in session
wrapping for tooltip read information
Include scaffolds on genome snapshot
long responsive time while track missed in (...)
'pack and hide' status not saved in session
refGene track automatically be put to bottom (...)
MCM (metadata colormap) shows blank when (...)
geneplot on methylC track
Json syntax error when trying to jump to (...)
juxtaposition issue on genome align tracks
refresh cache issue
gene set view not working
missing 0% methylated CpG display in methylC (...)
load hub and session issue
Matplot not working with sessions
dbSNP track crashes
Error in SE paper datahub
Gene shading feature inconsistent and needs (...)
Roadmap analysis hub crashes
Potential bug in the json hub name?
zooming in on a genomic region doesn't give (...)
can't zoom out in gene set view in mm9
Please add the option to remove genome comparison (...)
Incorrect SNP coordinate based on SNP ID
Pop-up window for confirming zooming in on (...)
Missing display of 0% methylated Cs in methylC (...)
Change names of custom tracks
Multiple upload of custom tracks

Custom track can not be submitted again after (...) track hub include feature slow loading of big hub json syntax error when loading genome comparison (...) Complete Epigenome Browser WUGB popup link get blocked	
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