

## WashU Epigenome Browser - Issues

| #   | Tracker | Status      | Priority | Subject   | Assignee | Updated             |
|-----|---------|-------------|----------|---|----------|---------------------|
| 3   | Bug     | New         | Normal   | Safari 5 navigation display disorder                            |          | 07/18/2013 01:34 pm |
| 46  | Bug     | New         | Normal   | Coordinate issue in hg19  | Xin Zhou | 09/19/2013 10:16 am |
| 77  | Bug     | New         | Normal   | track background color not applied on 2nd panel of line plot    |          | 10/26/2013 11:25 pm |
| 78  | Bug     | New         | Normal   | 2nd panel lineplot not applied to main panel                    |          | 10/26/2013 11:30 pm |
| 79  | Bug     | New         | Normal   | browser background color  |          | 10/26/2013 11:40 pm |
| 80  | Bug     | New         | Normal   | session with lineplot TypeError                                 |          | 10/26/2013 11:45 pm |
| 111 | Bug     | New         | Normal   | Tracks disappear when using config panel                        |          | 11/24/2013 03:25 pm |
| 115 | Bug     | New         | Normal   | tracks do not relocate and some disappear in GSV                | Xin Zhou | 11/23/2013 04:19 pm |
| 146 | Bug     | New         | Normal   | Session cannot save more than 1 custom metadata vocabularies    |          | 01/23/2014 07:37 am |
| 156 | Bug     | New         | Normal   | WUGB popup link get blocked                                     |          | 02/07/2014 11:32 pm |
| 157 | Bug     | New         | Normal   | refSeq gene position didn't keep in session                     |          | 02/14/2014 04:43 pm |
| 163 | Bug     | New         | Normal   | long responsive time while track missed in hub                  |          | 02/18/2014 03:39 pm |
| 168 | Bug     | New         | Normal   | 'pack and hide' status not saved in session                     |          | 03/01/2014 12:59 am |
| 179 | Bug     | New         | Normal   | refGene track automatically be put to bottom when split panel   |          | 03/19/2014 11:39 pm |
| 186 | Bug     | New         | Normal   | MCM (metadata colormap) shows blank when splitting panel        |          | 04/05/2014 09:40 am |
| 190 | Bug     | New         | Normal   | geneplot on methylC track                                       |          | 04/28/2014 12:11 am |
| 193 | Bug     | New         | Normal   | Json syntax error when trying to jump to a new location         | Xin Zhou | 05/14/2014 12:42 pm |
| 194 | Bug     | New         | Normal   | juxtaposition issue on genome align tracks                      |          | 05/25/2014 12:09 am |
| 198 | Bug     | New         | Normal   | refresh cache issue   |          | 06/18/2014 09:51 am |
| 204 | Bug     | In Progress | Normal   | gene set view not working                                       |          | 07/07/2014 03:04 pm |
| 227 | Bug     | New         | Normal   | missing 0% methylated CpG display in methylC track              |          | 10/03/2014 01:02 pm |
| 231 | Bug     | Resolved    | Normal   | load hub and session issue                                      |          | 10/15/2014 11:15 am |
| 232 | Bug     | Resolved    | Normal   | Matplot not working with sessions                               |          | 10/15/2014 02:09 pm |
| 233 | Bug     | Resolved    | Normal   | dbSNP track crashes   |          | 10/15/2014 08:04 pm |
| 235 | Bug     | New         | Normal   | Gene shading feature inconsistent and needs to be made optional |          | 10/16/2014 09:55 am |
| 237 | Bug     | New         | Normal   | Potential bug in the json hub name?                             | Xin Zhou | 10/17/2014 06:08 pm |
| 239 | Bug     | New         | Normal   | zooming in on a genomic region doesn't give the right region    |          | 10/27/2014 01:57 pm |
| 240 | Bug     | Resolved    | Normal   | can't zoom out in gene set view in mm9                          | Xin Zhou | 10/28/2014 06:46 am |
| 243 | Bug     | New         | Normal   | Incorrect SNP coordinate based on SNP ID                        | Xin Zhou | 11/07/2014 10:43 am |
| 245 | Bug     | Resolved    | Normal   | Missing display of 0% methylated Cs in methylC track            |          | 11/24/2014 01:40 pm |
| 248 | Bug     | New         | Normal   | Custom track can not be submitted again after removal           |          | 12/17/2014 04:21 pm |
| 250 | Bug     | New         | Normal   | slow loading of big hub   |          | 12/19/2014 03:45 pm |

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| 252 | Bug | New | Normal | json syntax error when loading genome comparison tracks |  | 03/10/2015 04:21 pm |
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