WashU Epigenome Browser - Issues

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

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