

Insyght, a conserved synteny - homolog browser - Issues

#	Tracker	Status	Priority	Subject	Author	Assignee	Updated	Category
3442	Feature	New	Normal	display information if homology is 1-1, 1-n or n-n	Thomas Lacroix	Thomas Lacroix	01/14/2016 04:10 PM	
3428	Bug	New	Low	Annot compa view do not take private genomes into account	Thomas Lacroix	Thomas Lacroix	04/18/2014 05:31 PM	
3139	Bug	New	Low	expand collapse partial genomic region	Thomas Lacroix	Thomas Lacroix	12/03/2013 05:47 PM	
3135	Feature	New	Low	center display on a reference gene in the middle of the screen	Thomas Lacroix	Thomas Lacroix	12/03/2013 05:45 PM	
3132	Bug	New	Low	off shoot expand error	Thomas Lacroix	Thomas Lacroix	04/17/2014 03:21 PM	
3122	Feature	New	Normal	change the result quick nav to be a cell widget that can handle a large quantity of data	Thomas Lacroix	Thomas Lacroix	07/31/2014 03:07 PM	
8720	Feature	In Progress	Normal	Pipeline rev 471 ; avoid non essential redondant data, decrease total database size by around 2 fold	Thomas Lacroix	Thomas Lacroix	08/28/2017 10:33 AM	
3794	Bug	Feedback	Normal	find gene and go back to ortholog view	Thomas Lacroix	Thomas Lacroix	01/14/2016 04:10 PM	
3787	Bug	Feedback	Normal	double click on empty cell in ortholog browser triggers error	Thomas Lacroix	Thomas Lacroix	01/14/2016 04:10 PM	