Insyght, a conserved synteny - homolog browser - Issues

#	Tracker	Status	Priority	Subject	Author	Assignee	Updated	Category
3381	Bug	Closed	High	pb login genome privé uncaught exception (example S salivarius)	Thomas Lacroix	Thomas Lacroix	04/30/2014 02:59 PM	
3269	Bug	Closed	High	expand synteny show more genes than expected if those genes overlap	Thomas Lacroix	Thomas Lacroix	04/30/2014 02:56 PM	
3210	Feature	Closed	High	View "Annotation comparator" : add comparison of "Product" et "Note"	Thomas Lacroix	Thomas Lacroix	04/30/2014 02:56 PM	
3209	Bug	Closed	High	when search page -> exclude genomes : ERROR: syntax error at or near ")"	Thomas Lacroix	Thomas Lacroix	04/30/2014 02:59 PM	
3121	Feature	Closed	High	copyright INRA	Thomas Lacroix	Thomas Lacroix	04/30/2014 02:58 PM	
3120	Feature	Closed	High	remove 100 organisms limit	Thomas Lacroix	Thomas Lacroix	04/30/2014 02:56 PM	
3119	Bug	Closed	High	alignment info is not displayed correctly	Thomas Lacroix	Thomas Lacroix	04/30/2014 02:56 PM	
3118	Bug	Closed	High	paper and website: change gene to CDS	Thomas Lacroix	Thomas Lacroix	04/30/2014 02:57 PM	
3116	Feature	Closed	High	in the search tab, rename "clear selected / clear all"	Thomas Lacroix	Thomas Lacroix	04/30/2014 02:57 PM	
3115	Feature	Closed	High	not all genomic location have comma separating the thousands	Thomas Lacroix	Thomas Lacroix	04/30/2014 02:57 PM	
3114	Bug	Rejected	High	no link to outside db for the compared gene annotations	Thomas Lacroix	Thomas Lacroix	04/16/2014 04:20 PM	
3113	Feature	Closed	High	button Add all in the search tab when building the gene set	Thomas Lacroix	Thomas Lacroix	04/30/2014 02:57 PM	
3112	Bug	Closed	High	red stain marker in the proportional view	Thomas Lacroix	Thomas Lacroix	04/30/2014 02:57 PM	
3111	Bug	Closed	High	remove update pipeline	Thomas Lacroix	Thomas Lacroix	04/30/2014 02:58 PM	
3110	Support	Closed	High	public svn realease	Thomas Lacroix	Thomas Lacroix	04/30/2014 02:58 PM	
3109	Bug	Rejected	High	locus tag can have multilpe feature id qualifiers	Thomas Lacroix	Thomas Lacroix	04/16/2014 02:47 PM	
3108	Bug	Closed	High	impossible to sort the homolog table by a reference gene	Thomas Lacroix	Thomas Lacroix	04/30/2014 02:58 PM	
3107	Bug	Rejected	High	back button of the web browser -> error ?	Thomas Lacroix	Thomas Lacroix	04/11/2014 10:16 AM	
3106	Bug	Closed	High	resize web browser -> empty result page	Thomas Lacroix	Thomas Lacroix	04/30/2014 02:59 PM	