

Insyght, a conserved synteny - homolog browser - Issues

#	Tracker	Status	Priority	Subject	Author	Assignee	Updated	Category
8731	Bug	Closed	Normal	Pipeline rev 471 ; check_for_element_types_and_reorganize_elements_into_organisms.pl want to update already updated element type from previous run	Thomas Lacroix	Thomas Lacroix	08/25/2017 12:08 PM	
8730	Bug	Closed	Normal	Pipeline rev 471 ; ObjectElement.pm -> setaccession -> deal with multiple digit in accnum name, do not file prefix and postfix	Thomas Lacroix	Thomas Lacroix	08/25/2017 12:07 PM	
8714	Bug	Closed	Normal	Navigation jumps when moving with >> or << in ortholog table or genomic organisation	Thomas Lacroix	Thomas Lacroix	08/25/2017 10:58 AM	
8713	Bug	Closed	Normal	Compared organisms not sorted correctly for scope Whole organism	Thomas Lacroix	Thomas Lacroix	08/25/2017 10:58 AM	
8712	Bug	Closed	Normal	Synteny symbol not displaying correct number of pairs, it was missing gene mismatch and insertion	Thomas Lacroix	Thomas Lacroix	08/25/2017 10:57 AM	
8711	Bug	Closed	Normal	When transferring q genes from synteny to ortholog table view, s_insertion genes were transeferd as well	Thomas Lacroix	Thomas Lacroix	08/25/2017 10:57 AM	
8710	Bug	Closed	Normal	Error when the database was empty. Redirect user to use the guide to add data or check out http://genome.jouy.inra.fr/Insyght/ to test functionalities	Thomas Lacroix	Thomas Lacroix	08/25/2017 10:57 AM	
8708	Bug	Closed	Normal	Fixed functionality to send email via smtp inra to report uncaught exception	Thomas Lacroix	Thomas Lacroix	08/25/2017 10:55 AM	
8707	Bug	Closed	Normal	Fixed examples taxon ids for release Archaea	Thomas Lacroix	Thomas Lacroix	08/25/2017 10:55 AM	
8705	Bug	Closed	Normal	Sometimes 2 references genomes are selected in the taxonomic browser (2 nodes in yellow) ; impossible to change the group for them...	Thomas Lacroix	Thomas Lacroix	08/25/2017 10:37 AM	
6506	Bug	Closed	Normal	max connection idle : better configuration of the context.xml config file (max total 97 SQL request in parrallel)	Thomas Lacroix	Thomas Lacroix	06/03/2016 03:14 PM	
6501	Bug	Closed	Normal	text above taxonomic browser search tab step 2 was not displaying properly	Thomas Lacroix	Thomas Lacroix	06/03/2016 03:09 PM	
6147	Bug	Closed	Normal	color is not displayed correctly at first within the core / dispensable genome browser of the search page, it appears right after clicking on another node	Thomas Lacroix	Thomas Lacroix	04/06/2016 02:45 PM	
6146	Bug	Closed	Normal	Uncaught exception: java.lang.IndexOutOfBoundsException when viewing resultq from core/ disp gene set directly in ortholog table	Thomas Lacroix	Thomas Lacroix	04/06/2016 02:45 PM	
6145	Bug	Closed	Normal	FATAL: remaining connection slots are reserved for non-replication superuser connections	Thomas Lacroix	Thomas Lacroix	04/06/2016 02:44 PM	
6144	Bug	Closed	Normal	the functionality "browse core / dispensable" genome" in the search tab is not longuer functioning correctly	Thomas Lacroix	Thomas Lacroix	04/06/2016 02:44 PM	
6143	Bug	Closed	Normal	The presence / absence homology filter did not work correctly with multiple organisms in the box : java.lang.Exception: could not parse orga id as integer : presence	Thomas Lacroix	Thomas Lacroix	04/06/2016 02:43 PM	
6140	Bug	Closed	Normal	The compared genomes list in the tab orthologs table and genomic organisation did contains public organisms with no element ; those organisms are now hidden from the result list	Thomas Lacroix	Thomas Lacroix	04/06/2016 02:42 PM	
6139	Bug	Closed	Normal	In the search tab, the pre-made examples were not working properly	Thomas Lacroix	Thomas Lacroix	04/06/2016 02:42 PM	

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6138	Bug	Closed	Normal	some gene have qualifier protein_id with ref to other db (EMBL) enclosed in curly braces, this did pose a problem for the blast2seq link when it was the case ; exemple gene id 41305 metS of Listeria Monocytogenes strain 08-5578 (db origami_test)	Thomas Lacroix	Thomas Lacroix	04/06/2016 02:41 PM	
6137	Bug	Closed	Normal	some genes have multiple qualifier protein_id, display them all in the info panel with correct link to external db	Thomas Lacroix	Thomas Lacroix	04/06/2016 02:41 PM	
6135	Bug	Closed	Normal	some genes have multiple qualifier protein_id (some ref to other db such as tigr, start with REF), this did pose a problem for the blast2seq link when it was the case ; exemple gene id 5727 EF0944 of E faecalis v583 (db origami_test)	Thomas Lacroix	Thomas Lacroix	04/06/2016 02:40 PM	
5819	Bug	Closed	Normal	WEB APP: some special characters were not displaying correctly on some web browser, changed to more widely supported characters	Thomas Lacroix	Thomas Lacroix	02/09/2016 10:30 AM	
3788	Bug	Closed	Normal	transfert gene set from genomic orga to ortho table : some genes are missing sometime ?	Thomas Lacroix		01/14/2016 04:06 PM	
3756	Bug	Closed	Normal	NullPointerException when navigating within a result page while data still loading	Thomas Lacroix	Thomas Lacroix	07/31/2014 03:27 PM	
3409	Bug	Closed	Normal	Ontologies comparator : genes aer not sorted according to bigger element as in other views	Thomas Lacroix	Thomas Lacroix	04/30/2014 02:54 PM	
3381	Bug	Closed	High	pb login genome privé uncaught exception (example S salivarius)	Thomas Lacroix	Thomas Lacroix	04/30/2014 02:59 PM	
3380	Bug	Closed	Normal	exception on logout	Thomas Lacroix	Thomas Lacroix	04/30/2014 02:56 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	
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	
3145	Bug	Closed	Normal	navigation button when more rows displayed	Thomas Lacroix	Thomas Lacroix	04/30/2014 02:54 PM	
3138	Bug	Closed	Low	Export feature homolog table	Thomas Lacroix	Thomas Lacroix	07/31/2014 03:28 PM	
3137	Bug	Closed	Low	Export feature in the search tab	Thomas Lacroix	Thomas Lacroix	07/31/2014 03:28 PM	
3134	Bug	Rejected	Normal	when sorting the homolog table by a reference gene, bug in the result quick navigation ?	Thomas Lacroix	Thomas Lacroix	04/18/2014 04:13 PM	
3133	Bug	Rejected	Normal	contextual menu find genes does not work for browse all homologs of an organism ?	Thomas Lacroix	Thomas Lacroix	04/18/2014 04:02 PM	
3131	Bug	Rejected	Normal	IndexOutOfBoundsException ??	Thomas Lacroix	Thomas Lacroix	04/18/2014 03:48 PM	
3130	Bug	Rejected	Normal	NoSuchElementException ??	Thomas Lacroix	Thomas Lacroix	04/18/2014 03:26 PM	
3129	Bug	Closed	Normal	error if taxon id = 0	Thomas Lacroix	Thomas Lacroix	04/30/2014 02:55 PM	
3127	Bug	Rejected	Normal	taxon_id link	Thomas Lacroix	Thomas Lacroix	04/17/2014 05:42 PM	
3123	Bug	Closed	Normal	export sequence all genes of gene set	Thomas Lacroix	Thomas Lacroix	04/30/2014 02:55 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	
3117	Bug	Closed	Normal	paper : homolog and synteny browser	Thomas Lacroix	Thomas Lacroix	04/17/2014 05:43 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	

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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	
3109	Bug	Rejected	High	locus tag can have multiple 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	