

Insyght, a conserved synteny - homolog browser - Issues

#	Tracker	Status	Priority	Subject	Author	Assignee	Updated	Category
8732	Amélioration	Closed	Normal	Pipeline rev 471 ; improve the accessibility of the VM by simplifying the pipeline into 1 master script (keeping error handling proper)	Thomas Lacroix	Thomas Lacroix	08/25/2017 12:09 PM	
8729	Amélioration	Closed	Normal	Pipeline rev 471 ; check_db_consistency.pl : test that element lenght = sequence lenght	Thomas Lacroix	Thomas Lacroix	08/25/2017 12:07 PM	
8728	Amélioration	Closed	Normal	Pipeline rev 471 ; check_db_consistency.pl : test that genes.residues = micado.prot.feats.residues	Thomas Lacroix	Thomas Lacroix	08/25/2017 12:06 PM	
8727	Amélioration	Closed	Normal	Pipeline rev 471 ; script download_from_ncbi.pl : add the option -restrict_to_list_taxon_id_and_subnodes	Thomas Lacroix	Thomas Lacroix	08/25/2017 12:05 PM	
8724	Amélioration	Closed	Normal	Pipeline rev 471 ; Task_add_entry (insert of primary data) : Make the script more resilient to errors and re-run, faster (generator and integrator)	Thomas Lacroix	Thomas Lacroix	08/25/2017 11:47 AM	
8723	Amélioration	Closed	Normal	Pipeline rev 471 ; fill_gene_entry.pl : generate data for table gene directly instead of after schema micado is filled w/ data	Thomas Lacroix	Thomas Lacroix	08/25/2017 11:46 AM	
8722	Amélioration	Closed	Normal	Pipeline rev 471 ; fill_gene_entry.pl : infer correct type directly here instead of subsequent script	Thomas Lacroix	Thomas Lacroix	08/25/2017 11:45 AM	
8719	Amélioration	Closed	Normal	Pipeline rev 471 ; improved analyse_stats_db.pl	Thomas Lacroix	Thomas Lacroix	08/25/2017 11:34 AM	
8717	Amélioration	Closed	Normal	Pipeline rev 471 ; fill_gene_entry.pl : add Locus_tag directly in table genes	Thomas Lacroix	Thomas Lacroix	08/25/2017 11:30 AM	
8716	Amélioration	Closed	Normal	Pipeline rev 471 ; Task_add_entry : cleaner option -GENOME_FILES_FORMAT	Thomas Lacroix	Thomas Lacroix	08/25/2017 11:25 AM	
8715	Amélioration	Closed	Normal	Pipeline rev 471 ; Task_add_entry : added option -SKIP_NEIGHBOR_INFO as this info is not useful	Thomas Lacroix	Thomas Lacroix	08/25/2017 11:23 AM	
8709	Amélioration	Closed	Normal	Modified user interface to deal with database changes in schema: removal of elements.description (-> sequences.definition), elements.date_seq (-> sequences.date_seq), elements.version (-> sequences.version), genes.residue (-> prot_feat.proteine)	Thomas Lacroix	Thomas Lacroix	08/25/2017 10:56 AM	
5833	Amélioration	Closed	Normal	Automatic and anonymous bug reporting tool	Thomas Lacroix	Thomas Lacroix	02/09/2016 02:19 PM	