

## Insyght, a conserved synteny - homolog browser - Feature #8725

**adapt to database schema rev 2: additional links and info from ncbi, add columns for convenience when querying the database (avoid cross tables querying)**

08/25/2017 11:53 AM - Thomas Lacroix

<b>Status:</b>	Closed	<b>Start date:</b>	08/25/2017
<b>Priority:</b>	Normal	<b>Due date:</b>	
<b>Assignee:</b>	Thomas Lacroix	<b>% Done:</b>	100%
<b>Category:</b>		<b>Estimated time:</b>	0.00 hour
<b>Target version:</b>			
<b>Description</b>			

### History

#### #1 - 08/25/2017 12:00 PM - Thomas Lacroix

- Subject changed from Pipeline rev 471 ; adapt to database schema rev 2 : modify pipeline to deal with additional info in table elements: NCBI\_internal\_id, NCBI\_sourcedb, NCBI\_tech, NCBI\_geneticcode, NCBI\_topology, NCBI\_completeness, NCBI\_status, NCBI\_comment ; also warn if ge to adapt to database schema rev 2: additional links and info from ncbi, add columns for convenience when querying the database (avoid cross tables querying)

-> additional info in table organisms: NCBI\_assemblyaccession\_IT, NCBI\_assemblyname\_IT, NCBI\_lastupdateddate\_IT, NCBI\_seqreleasedate\_IT, NCBI\_isolate\_IT, NCBI\_speciestaxid\_IT, NCBI\_biosampleid\_IT, NCBI\_biosampleaccn\_IT, NCBI\_list\_bioprojectid\_IT, NCBI\_list\_bioprojectaccn\_IT, NCBI\_assemblyclass\_IT, NCBI\_assemblystatus\_IT

-> additional info in table elements: NCBI\_internal\_id, NCBI\_sourcedb, NCBI\_tech, NCBI\_geneticcode, NCBI\_topology, NCBI\_completeness, NCBI\_status, NCBI\_comment ; also warn if genome status = suppressed