{2} RNA-Seqlyze Tickets by Milestone   (5 matches)

2012-05-25 12:00:00 | Solution Conceptualized   (1 match)

Ticket	Priority	Summary	Component	Type	Resolution
Description					
#3	high	Software framework choice	Software	task	

It has to be decided which software frameworks should be used (and which not).

    -> This is almost done, what is still missing, is the corresponding documentation on the project wiki.

2012-06-15 18:00:00 | Core Features Implemented   (1 match)

Ticket	Priority	Summary	Component	Type	Resolution
Description					
#7	high	Implement an initial version of the scoring algorithm	Software	task	

The scoring algorithm is going to be the core feature of the software.

It is expected that there will be a lot of "difficult cases" -- i.e. contradicting data -- that will make it hard to write a robust algorithm that delivers reasonable results for a variety of input data.

The calculation of the final score will be done in a python function and the source code must be documented thoroughly, such that any person with sufficient expertise in bioinformatics will be able to understand what is going within reasonable time.

2012-06-22 18:00:00 | Version 1.0   (2 matches)

Ticket	Priority	Summary	Component	Type	Resolution
Description					
#5	high	The operon data tracks need a serious checkup	Software	defect	

The data integrity needs to be verified.

Specifically:

- there should not be any single gene operons
- when linking multiple (di-cistronic) operons the score should be adjusted

#6	low	The SRS is still incomplete	Presentation	defect	

Fill in the missing parts of the SRS (TBD)

2012-07-13 12:00:00 | Version 1.1   (1 match)

Ticket	Priority	Summary	Component	Type	Resolution
Description					
#8	low	Add References to SRS	Report	task	

References to the project homepage as well as the Trac page should be added to the SRS in a "References" section in chapter one once these sites become publicly available.