IQC Validator

NCATS Informatics

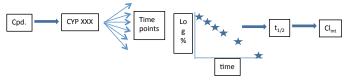
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Overview

- Desktop client
- Proposed scoring function to evaluate data fitting

Features

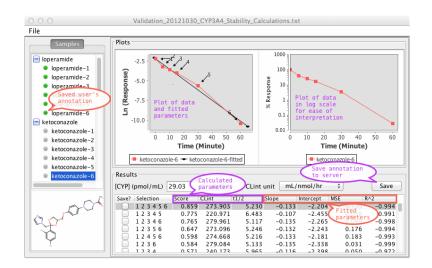
 A simple desktop client to help validate the CYP isozyme data generation workflow



- Minimal user interaction
 - Precalculate results for all possible combinations
 - Rank results based on scoring function
- A web-based protocol for data management (i.e., data upload, download, and annotation)
- Available at http://tripod.nih.gov/ws/iqc/iqc.jnlp

Desktop client (cont'd)

Screenshot



Proposed scoring function

- Developed based on discussion with Scott Obach and Ed Kerns
- Let the raw score be defined as follows

$$Score_{raw} = NR^2 \exp(-\sigma_e) \sum_{i} 2^{-i}, \qquad (1)$$

where N is the number of data points used to build the linear model, $\sigma_e \geq 0$ and $R \in [-1,1]$ are the estimated mean square error and Pearson's correlation of the model, respectively.

- Weighted contribution of T_0 , T_5 , T_{10} , T_{15} , T_{30} , T_{60} as $1, \frac{1}{2}, \frac{1}{4}, \frac{1}{8}, \frac{1}{16}, \frac{1}{32}$, respectively.
- The best possible raw score (i.e., $\sigma_e=0$, |R|=1, and N=6) is

Score_{best} =
$$6 \sum_{i=0}^{5} 2^{-i} = 11.8125$$

with i corresponds to the index of $\{T_0, T_5, T_{10}, T_{15}, T_{30}, T_{60}\}$.



Scoring function (cont'd)

Normalized score is the ratio of raw and best

$$Score = \frac{Score_{raw}}{Score_{best}}$$

where Score \in [0, 1].

- Initial observations
 - Good fit when score ≥ 0.9
 - In good agreement with with manual evaluation (based on limited annotations from Ed Kerns)
- Require additional validation
 - Collect additional manual annotations
 - Evaluation metric (e.g., Spearman's rank correlation)