Bioinformatics

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GEMSIRV: a software platform for GEnome-scale metabolic model simulation, reconstruction and visualization

Data and text mining

Boosting automatic event extraction from the literature using domain adaptation and coreference resolution
Y.C.Liao, F.Thompson and S.Aranidou

Statistical interpretation of machine learning-based feature importance scores for biomarker discovery
V.A.Heyn-Th, T.Zelev, L.Wehenkel and P.Guerits

Two effective methods for correcting experimental high-throughput screening data
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