DEVELOPMENT SCIENCES: Innovation & Technology

TITLE

Automating Quality Assessment of Mass Spectrometry Peaks Using Machine Learning

SUMMARY

Targeted mass spectrometry data analysis requires manual inspection of chromatographic peaks from large datasets to ensure quality of biomarker data. A QC framework and toolset has been developed that leverages feature engineering and machine learning to automate this process.

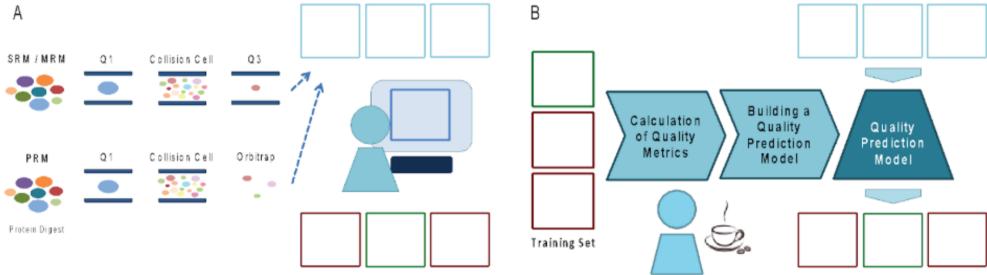
IMPACT

Improves throughput and reproducibility of MS biomarker data analysis

Facilitates faster and more robust biomarker assay development and implementation

Manual MS Peak QC is time- and labor-intensive 0.1 Collision Cell Q 3

ML-assisted QC improves the efficiency of data processing



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