

High-Throughput Mass Spectrometry Imaging with Dynamic Sparse Sampling

A deep learning algorithm enhances mass spectrometry imaging by prioritizing molecularly informative tissues, significantly reducing data acquisition time and accelerating molecular insights.

Researchers at Purdue University have developed a method for mass spectrometry imaging that utilizes an artificial intelligence algorithm to prioritize and extract results from the most molecularly informative tissues. This Deep Learning Approach for Dynamic Sampling (DLADS) reduces the number of measurements needed and reconstructs molecular images with high fidelity. While others in mass spectrometry imaging have focused on improving spatial resolution, this approach continues to actively learn based on known/existing data, resulting in less overall experimental time.

Technology Validation: When DLADS was paired with nanospray desorption electrospray ionization (nano-DESI) MSI, the data acquisition time reduced by a factor of 3. DLADS also effectively reconstructed mouse kidney images, after training with mouse uterine images.

Advantages:

- Less experimental time
- Deeper data insights

Applications:

- Molecular imaging
- Artificial intelligence

TRL: 4

Intellectual Property:

Technology ID

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Category

Artificial Intelligence & Machine Learning/Computer Vision & Image Recognition
Artificial Intelligence & Machine Learning/AI-Integrated Imaging Systems & Industrial Vision and Inspection
Biotechnology & Life Sciences/Analytical & Diagnostic Instrumentation

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