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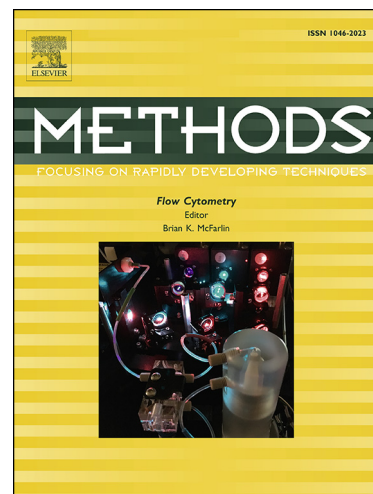
Patanachai Limpikirati, Tianying Liu, Richard W. Vachet

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# Covalent Labeling-Mass Spectrometry with Non-Specific Reagents for Studying Protein Structure and Interactions

Patanachai Limpikirati<sup>‡</sup>, Tianying Liu<sup>‡</sup>, and Richard W. Vachet\*

Department of Chemistry, University of Massachusetts Amherst, Massachusetts 01003, United States

\* Corresponding author

‡ These authors contributed equally to this manuscript.

## Corresponding Author Information

Richard W. Vachet: [rwvachet@chem.umass.edu](mailto:rwvachet@chem.umass.edu)

## Highlights

- Covalent labeling - mass spectrometry (CL-MS) is a valuable tool to study protein structure and protein interactions.
- CL-MS primarily reports on the solvent accessibility of amino acid side chains, and this is used to obtain structural information.
- Non-specific reagents irreversibly modify a wide variety of amino acid side chains, thus enabling broad structural coverage in a single experiment.
- CL-MS techniques are suited for distinguishing protein topologies, conformations, and/or binding sites.

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