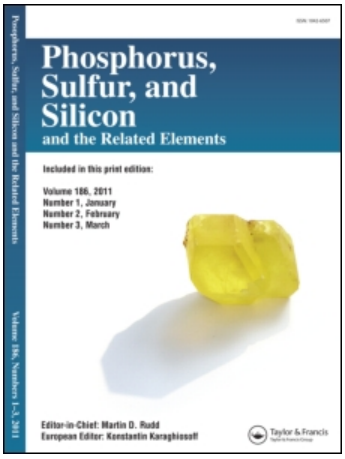

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A Novel Phosphoryl Derivative Method for Peptide Sequencing by ESI-MS/MS

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A NOVEL PHOSPHORYL DERIVATIVE METHOD FOR PEPTIDE SEQUENCING BY ESI-MS/MS

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In the past decade, mass spectrometry has become a powerful tool on peptide sequencing. Different derivative methods for peptide have been used to simplify the mass spectra analysis.[1,2]

Here, we report a novel derivative method for peptide sequencing. A homo-peptide $(Phe)_6$ was derived by N-Di-methyloxyphosphoryl group at the N-terminal. The sodium adduct was selected and analyzed by ESI-MS/MS which presented very simple fragments.

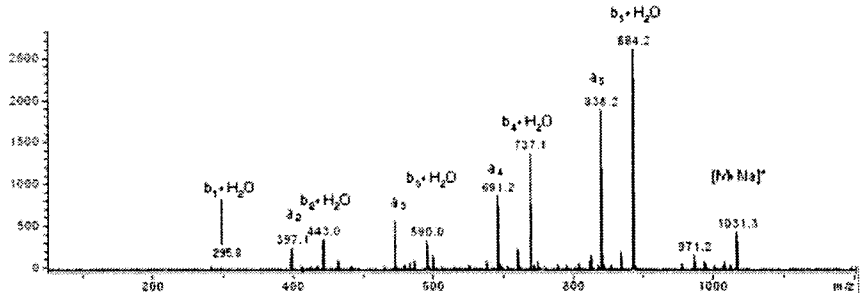

FIGURE 1 Tandem mass spectrometry of DMP-Phe_6.

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