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## IN SILICO DETERMINATION OF METHYLATED RESIDUES IN MAMMALIAN PROTEINS



LAP Lambert Academic Publishing Mai 2012, 2012. Taschenbuch. Book Condition: Neu. 220x150x5 mm. This item is printed on demand - Print on Demand Neuware - Bioinformatics is a helpful tool to experimental studies. It is less time consuming and may provide a platform for experimental work. Post translational modification is investigated in silico, with the help of the bioinformatics tool MAPRes. This tool mines association pattern around a modified residue. In this work association rules were mined around methylated Lys/Arg...

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- · Authored by Anila Saeed
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