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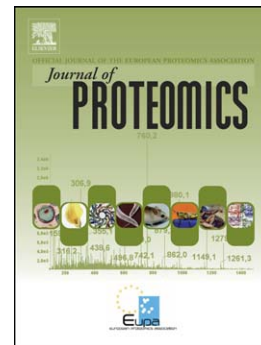
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REVIEW ARTICLE (revised)

In silico prediction and characterization of protein post-translational modifications.

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