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Prediction of S-sulfenylation sites using mRMR feature selection and fuzzy support vector machine algorithm

Zhe Ju , Shi-Yun Wang

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Highlights

- A novel predictor is developed to predict S-sulfenylation sites
- The combined features are used to predict and analyze S-sulfenylation sites
- mRMR feature selection method is adopted to remove the redundant features
- A free online service is available for prediction

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