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Bi-PSSM: position specific scoring matrix based intelligent
Computational model for identification of mycobacterial membrane
proteins

Muslim Khan , Maqsood Hayat , Sher Afzal Khan , Saeed Ahmad ,
Nadeem Iqbal

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Highlights

- Computational model Bi-PSSM is developed for prediction of mycobacterial membrane proteins.
- Three feature extraction schemes are used.
- Various classification algorithms are utilized
- SVM with combination with PSSM achieved quite promising results.

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