

Judul:

Prediksi interaksi protein berdasarkan barisan asam amino pada human immunodeficiency virus menggunakan pseudo substitution matrix representation feature dikombinasikan dengan rotation forest ensemble classifier = Sequence base prediction of protein-protein interactions using pseudo substitution matrix representation feature and ensemble rotation forest classifier in HIV (human immunodeficiency virus)

Pengarang/Penulis:

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Subjek:

Protein-protein interactions--Mathematical models

Nomor Panggil:

T49440

Penerbitan:

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