

Journalbeiträge

1. Dong Z, Wang K, Dang TKL, Gültas M, Welter M, Wierschin T, Stanke M, Waack S (2014) CRF-based models of protein surfaces improve protein-protein interaction site predictions. BMC BIOINFORMATICS 15: 277, doi: 10.1186/1471-2105-15-277
2. Dönitz J, Wingender E (2014) EndoNet: an information resource about the intercellular signaling network. BMC SYST BIOL 8(1): 49, doi: 10.1186/1752-0509-8-49
3. Gültas M, Düzgün G, Herzog S, Jäger SJ, Meckbach C, Wingender E, Waack S (2014) Quantum coupled mutation finder: predicting functionally or structurally important sites in proteins using quantum Jensen-Shannon divergence and CUDA programming. BMC BIOINFORMATICS 15(1): 96, doi: 10.1186/1471-2105-15-96