

## Documents

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- 1) Bastami, Z., Sheikhpour, R., Razzaghi, P., Ramazani, A., Gharaghani, S.

**Proteochemometrics modeling for prediction of the interactions between caspase isoforms and substrates**  
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- 2) Li, Z., Miao, Q., Yan, F., Meng, Y., Zhou, P.

**Machine learning in quantitative protein-peptide affinity prediction: Implications for the design of peptide-based drugs**  
(2019) *Current Drug Metabolism*, 20 (3), pp. 170-176.

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