

Advancing drug–target interaction prediction: a comprehensive graph-based approach integrating knowledge graph embedding and ProtBert pretraining

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Caseins from bovine colostrum and milk strongly bind piscidin-1, an antimicrobial peptide from fish

Kütt, Mary-Liis; Stagsted, Jan International journal of biological macromolecules 2014 / p. 364-372 : ill

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A computational approach to study functional monomer–protein molecular interactions to optimize protein molecular imprinting

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EBSI investigation of microstructure and microtexture evolution on additively manufactured TiC-Fe based cermets—Influence of multiple laser scanning

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Effect of agitation on the peptide fibrillization: Alzheimer’s amyloid- b peptide 1-42 but not amylin and insulin fibrils can grow under quiescent conditions

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Formulation of active food packaging by design: Linking composition of the film-forming solution to properties of the chitosan-based film by response surface methodology (RSM) modelling

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Graphene oxide-terminated hyperbranched amino polymer-carboxymethyl cellulose ternary nanocomposite for efficient removal of heavy metals from aqueous solutions

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Heating-induced transition of Potyvirus Potato Virus A coat protein into β -structure

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Isolated Potato Virus A coat protein possesses unusual properties and forms different short virus-like particles

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Solid-state NMR backbone chemical shift assignments of α -synuclein amyloid fibrils at fast MAS regime

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