

AllerTOP v.2â€™a server for in silico prediction of allerg

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Citation Report

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1	In silico tools for exploring potential human allergy to proteins. <i>Drug Discovery Today: Disease Models</i> , 2015, 17-18, 3-11.	1.2	13
2	Accurate Classification of Biological Data Using Ensembles. , 2015, , .		0
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5	Bioactive hydrolysates from bovine blood globulins: Generation, characterisation, and in silico prediction of toxicity and allergenicity. <i>Journal of Functional Foods</i> , 2016, 24, 142-155.	1.6	39
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9	In silico approaches towards the exploration of rice bran proteins-derived angiotensin-I-converting enzyme inhibitory peptides. <i>International Journal of Food Properties</i> , 2017, , 1-14.	1.3	7
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14	Immunoinformatics Approach for Epitope-Based Peptide Vaccine Design and Active Site Prediction against Polyprotein of Emerging Oropouche Virus. <i>Journal of Immunology Research</i> , 2018, 2018, 1-22.	0.9	102
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17	Immunoinformatics-aided design of a potential multi-epitope peptide vaccine against <i>Leishmania infantum</i> . <i>International Journal of Biological Macromolecules</i> , 2018, 120, 1127-1139.	3.6	63
18	In Silico Analysis of Peptide Potential Biological Functions. <i>Russian Journal of Bioorganic Chemistry</i> , 2018, 44, 367-385.	0.3	11

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