## Neil Arvin Bretaña

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	dbPTM 3.0: an informative resource for investigating substrate site specificity and functional association of protein post-translational modifications. Nucleic Acids Research, 2013, 41, D295-D305.	14.5	179
2	Exploiting maximal dependence decomposition to identify conserved motifs from a group of aligned signal sequences. Bioinformatics, 2011, 27, 1780-1787.	4.1	100
3	PlantPhos: using maximal dependence decomposition to identify plant phosphorylation sites with substrate site specificity. BMC Bioinformatics, 2011, 12, 261.	2.6	63
4	A prospective study of hepatitis <scp>C</scp> incidence in Australian prisoners. Addiction, 2014, 109, 1695-1706.	3.3	58
5	A two-layered machine learning method to identify protein O-GlcNAcylation sites with O-GlcNAc transferase substrate motifs. BMC Bioinformatics, 2015, 16, S10.	2.6	41
6	Transmission of Hepatitis C Virus among Prisoners, Australia, 2005–2012. Emerging Infectious Diseases, 2015, 21, 765-774.	4.3	37
7	Identifying Protein Phosphorylation Sites with Kinase Substrate Specificity on Human Viruses. PLoS ONE, 2012, 7, e40694.	2.5	36
8	Carboxylator: incorporating solvent-accessible surface area for identifying protein carboxylation sites. Journal of Computer-Aided Molecular Design, 2011, 25, 987-995.	2.9	26