

Qing-Bin Gao, 高庆斌

List of Publications by Year in descending order

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12
papers

402
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933447

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1199594

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g-index

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all docs

12
docs citations

12
times ranked

283
citing authors

#	ARTICLE	IF	CITATIONS
1	Classifying G-protein-coupled receptors to the finest subtype level. <i>Biochemical and Biophysical Research Communications</i> , 2013, 439, 303-308.	2.1	18
2	Prediction of pattern recognition receptor family using pseudo-amino acid composition. <i>Biochemical and Biophysical Research Communications</i> , 2012, 417, 73-77.	2.1	18
3	Improving discrimination of outer membrane proteins by fusing different forms of pseudo amino acid composition. <i>Analytical Biochemistry</i> , 2010, 398, 52-59.	2.4	48
4	Prediction of nuclear receptors with optimal pseudo amino acid composition. <i>Analytical Biochemistry</i> , 2009, 387, 54-59.	2.4	40
5	Improving the Classification of Nuclear Receptors with Feature Selection. <i>Protein and Peptide Letters</i> , 2009, 16, 823-829.	0.9	6
6	EZ-Entry: A clinical data management system. <i>Computers in Biology and Medicine</i> , 2008, 38, 1042-1044.	7.0	11
7	Classification of Amine Type G-Protein Coupled Receptors with Feature Selection. <i>Protein and Peptide Letters</i> , 2008, 15, 834-842.	0.9	10
8	Center-based nearest neighbor classifier. <i>Pattern Recognition</i> , 2007, 40, 346-349.	8.1	90
9	Classification of G-protein coupled receptors at four levels. <i>Protein Engineering, Design and Selection</i> , 2006, 19, 511-516.	2.1	76
10	Feature Subset Selection for Protein Subcellular Localization Prediction. <i>Lecture Notes in Computer Science</i> , 2006, , 433-443.	1.3	1
11	Using Nearest Feature Line and Tunable Nearest Neighbor methods for prediction of protein subcellular locations. <i>Computational Biology and Chemistry</i> , 2005, 29, 388-392.	2.3	16
12	Prediction of protein subcellular location using a combined feature of sequence. <i>FEBS Letters</i> , 2005, 579, 3444-3448.	2.8	68