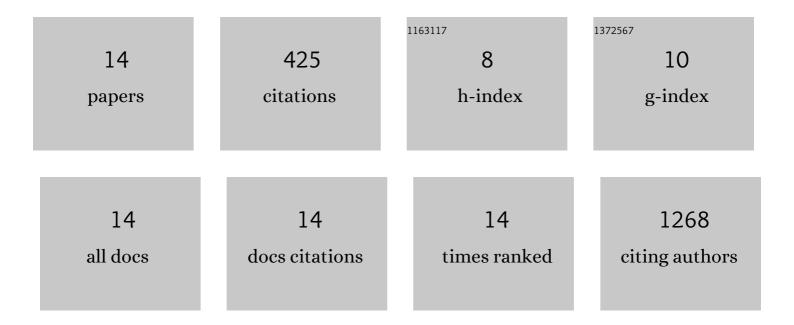
## Victoria Petri

List of Publications by Year in descending order

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VICTORIA DETRI

#	Article	IF	CITATIONS
1	The Disease Portals, disease–gene annotation and the RGD disease ontology at the Rat Genome Database. Database: the Journal of Biological Databases and Curation, 2016, 2016, baw034.	3.0	20
2	Exploring human disease using the Rat Genome Database. DMM Disease Models and Mechanisms, 2016, 9, 1089-1095.	2.4	27
3	Guidelines for the functional annotation of microRNAs using the Gene Ontology. Rna, 2016, 22, 667-676.	3.5	35
4	Comprehensive coverage of cardiovascular disease data in the disease portals at the Rat Genome Database. Physiological Genomics, 2016, 48, 589-600.	2.3	3
5	Disease, Models, Variants and Altered Pathways—Journeying RGD Through the Magnifying Glass. Computational and Structural Biotechnology Journal, 2016, 14, 35-48.	4.1	4
6	Structural libraries of protein models for multiple species to understand evolution of the renin-angiotensin system. General and Comparative Endocrinology, 2015, 215, 106-116.	1.8	10
7	The Rat Genome Database 2015: genomic, phenotypic and environmental variations and disease. Nucleic Acids Research, 2015, 43, D743-D750.	14.5	213
8	PhenoMiner: a quantitative phenotype database for the laboratory rat, Rattus norvegicus . Application in hypertension and renal disease. Database: the Journal of Biological Databases and Curation, 2015, 2015, .	3.0	11
9	The pathway ontology $\hat{a} \in $ updates and applications. Journal of Biomedical Semantics, 2014, 5, 7.	1.6	83
10	PhenoMiner: an interactive tool for physiologists integrating phenotype data using multiple ontologies. FASEB Journal, 2012, 26, 717.1.	0.5	0
11	The Rat Genome Database Pathway Portal. Database: the Journal of Biological Databases and Curation, 2011, 2011, bar010-bar010.	3.0	18
12	Pathway Resources at the Rat Genome Database. , 2011, , 316-336.		1
13	Gene Curation Software at the Rat Genome Database (RGD). Nature Precedings, 2010, , .	0.1	0
14	Physiology Pathway diagrams: new interactive online tools that provide efficient access to genomic and phenomic information through biological pathway analysis FASEB Journal, 2009, 23, 801.4.	0.5	0