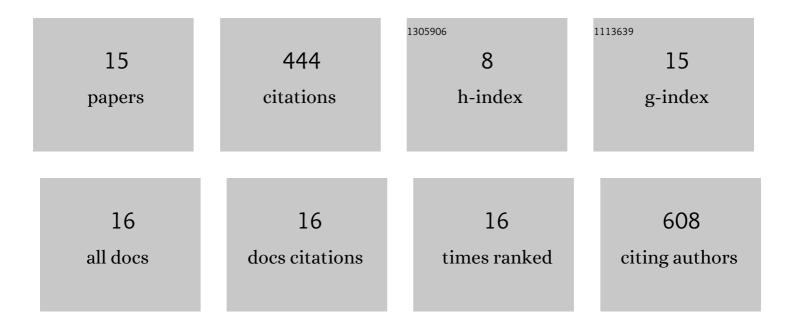
Varsha D Badal

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1092167/publications.pdf Version: 2024-02-01



ΝΑΡΩΗΛ Π ΒΛΠΛΙ

#	Article	IF	CITATIONS
1	Confidence, accuracy judgments and feedback in schizophrenia and bipolar disorder: a time series network analysis. Psychological Medicine, 2023, 53, 4200-4209.	2.7	2
2	Dynamics of Loneliness Among Older Adults During the COVID-19 Pandemic: Pilot Study of Ecological Momentary Assessment With Network Analysis. Frontiers in Digital Health, 2022, 4, 814179.	1.5	2
3	Natural language processing of medical records: new understanding of suicide ideation by dementia subtypes. International Psychogeriatrics, 2022, 34, 319-321.	0.6	1
4	Automated Analysis of Drawing Process to Estimate Global Cognition in Older Adults: Preliminary International Validation on the US and Japan Data Sets. JMIR Formative Research, 2022, 6, e37014.	0.7	9
5	Prediction of Loneliness in Older Adults Using Natural Language Processing: Exploring Sex Differences in Speech. American Journal of Geriatric Psychiatry, 2021, 29, 853-866.	0.6	27
6	Text mining for modeling of protein complexes enhanced by machine learning. Bioinformatics, 2021, 37, 497-505.	1.8	2
7	Computational methods for integrative evaluation of confidence, accuracy, and reaction time in facial affect recognition in schizophrenia. Schizophrenia Research: Cognition, 2021, 25, 100196.	0.7	8
8	Do Words Matter? Detecting Social Isolation and Loneliness in Older Adults Using Natural Language Processing. Frontiers in Psychiatry, 2021, 12, 728732.	1.3	12
9	Dynamic contextual influences on social motivation and behavior in schizophrenia: a case-control network analysis. NPJ Schizophrenia, 2021, 7, 62.	2.0	11
10	The Gut Microbiome, Aging, and Longevity: A Systematic Review. Nutrients, 2020, 12, 3759.	1.7	207
11	Blind prediction of homo―and heteroâ€protein complexes: The CASP13â€CAPRI experiment. Proteins: Structure, Function and Bioinformatics, 2019, 87, 1200-1221.	1.5	99
12	Modeling CAPRI targets 110â€120 by templateâ€based and free docking using contact potential and combined scoring function. Proteins: Structure, Function and Bioinformatics, 2018, 86, 302-310.	1.5	19
13	Natural language processing in text mining for structural modeling of protein complexes. BMC Bioinformatics, 2018, 19, 84.	1.2	27
14	Genome-Wide Structural Modeling of Protein-Protein Interactions. Lecture Notes in Computer Science, 2016, , 95-105.	1.0	1
15	Text Mining for Protein Docking. PLoS Computational Biology, 2015, 11, e1004630.	1.5	17