## Thomas A Peterson

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3622669/publications.pdf

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1162889 1372474 10 423 8 10 citations h-index g-index papers 12 12 12 852 docs citations times ranked citing authors all docs

| #  | Article  | IF  | CITATIONS |
|----|--|-----|-----------|
| 1  | Quantifying Variation in Treatment Utilization for Type 2 Diabetes Across Five Major University of California Health Systems. Diabetes Care, 2021, 44, 908-914.  | 4.3 | 9         |
| 2  | Discovery of Depression-Associated Factors From a Nationwide Population-Based Survey:<br>Epidemiological Study Using Machine Learning and Network Analysis. Journal of Medical Internet<br>Research, 2021, 23, e27344. | 2.1 | 11        |
| 3  | Protected Health Information filter (Philter): accurately and securely de-identifying free-text clinical notes. Npj Digital Medicine, 2020, 3, 57.   | 5.7 | 38        |
| 4  | Explanatory Model of Dry Eye Disease Using Health and Nutrition Examinations: Machine Learning and Network-Based Factor Analysis From a National Survey. JMIR Medical Informatics, 2020, 8, e16153.                    | 1.3 | 8         |
| 5  | Empirical Null Estimation Using Zero-inflated Discrete Mixture Distributions and Its Application to Protein Domain Data. Biometrics, 2018, 74, 458-471.  | 0.8 | 9         |
| 6  | Oncodomains: A protein domain-centric framework for analyzing rare variants in tumor samples. PLoS Computational Biology, 2017, 13, e1005428.  | 1.5 | 25        |
| 7  | Regulatory Single-Nucleotide Variant Predictor Increases Predictive Performance of Functional Regulatory Variants. Human Mutation, 2016, 37, 1137-1143.  | 1.1 | 13        |
| 8  | Towards Precision Medicine: Advances in Computational Approaches for the Analysis of Human Variants. Journal of Molecular Biology, 2013, 425, 4047-4063.   | 2.0 | 122       |
| 9  | Crosstalk between the Circadian Clock and Innate Immunity in Arabidopsis. PLoS Pathogens, 2013, 9, e1003370.   | 2.1 | 164       |
| 10 | Incorporating molecular and functional context into the analysis and prioritization of human variants associated with cancer. Journal of the American Medical Informatics Association: JAMIA, 2012, 19, 275-283.       | 2.2 | 23        |