

# Suzanne Miyamoto

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1759634/publications.pdf>

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9  
papers

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citations

1040056

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1016  
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#	ARTICLE	IF	CITATIONS
1	Multiple Reaction Monitoring for the Quantitation of Serum Protein Glycosylation Profiles: Application to Ovarian Cancer. <i>Journal of Proteome Research</i> , 2018, 17, 222-233.	3.7	37
2	Glycoproteomic Analysis of Malignant Ovarian Cancer Ascites Fluid Identifies Unusual Glycopeptides. <i>Journal of Proteome Research</i> , 2016, 15, 3358-3376.	3.7	28
3	Protein-Specific Differential Glycosylation of Immunoglobulins in Serum of Ovarian Cancer Patients. <i>Journal of Proteome Research</i> , 2016, 15, 1002-1010.	3.7	87
4	Chip-based nLC-TOF-MS is a highly stable technology for large-scale high-throughput analyses. <i>Analytical and Bioanalytical Chemistry</i> , 2013, 405, 4953-4958.	3.7	35
5	Absolute Quantitation of Immunoglobulin G and Its Glycoforms Using Multiple Reaction Monitoring. <i>Analytical Chemistry</i> , 2013, 85, 8585-8593.	6.5	111
6	Developments in the Identification of Glycan Biomarkers for the Detection of Cancer. <i>Molecular and Cellular Proteomics</i> , 2013, 12, 846-855.	3.8	130
7	The glycolyzer: Automated glycan annotation software for high performance mass spectrometry and its application to ovarian cancer glycan biomarker discovery. <i>Proteomics</i> , 2012, 12, 2523-2538.	2.2	44
8	Profile of native N-linked glycan structures from human serum using high performance liquid chromatography on a microfluidic chip and time-of-flight mass spectrometry. <i>Proteomics</i> , 2009, 9, 1939-1951.	2.2	131
9	The development of retrosynthetic glycan libraries to profile and classify the human serum N-linked glycome. <i>Proteomics</i> , 2009, 9, 2986-2994.	2.2	110