

# Gege Xu

## List of Publications by Year in descending order

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Version: 2024-02-01

35  
papers

2,919  
citations

331259

21  
h-index

360668

35  
g-index

37  
all docs

37  
docs citations

37  
times ranked

4562  
citing authors

#	ARTICLE	IF	CITATIONS
1	Serum Glycoprotein Markers in Nonalcoholic Steatohepatitis and Hepatocellular Carcinoma. <i>Journal of Proteome Research</i> , 2022, 21, 1083-1094.	1.8	14
2	Glycanâ€ protein cross-linking mass spectrometry reveals sialic acid-mediated protein networks on cell surfaces. <i>Chemical Science</i> , 2021, 12, 8767-8777.	3.7	14
3	Glycan biomarkers of autoimmunity and bile acid-associated alterations of the human glycome: Primary biliary cirrhosis and primary sclerosing cholangitis-specific glycans. <i>Clinical Immunology</i> , 2021, 230, 108825.	1.4	2
4	The glycoproteomics of hawk and caiman tears. <i>BMC Veterinary Research</i> , 2021, 17, 381.	0.7	0
5	Indole-3-lactic acid associated with Bifidobacterium-dominated microbiota significantly decreases inflammation in intestinal epithelial cells. <i>BMC Microbiology</i> , 2020, 20, 357.	1.3	117
6	A site-specific map of the human plasma glycome and its age and gender-associated alterations. <i>Scientific Reports</i> , 2020, 10, 17505.	1.6	14
7	PB-Net: Automatic peak integration by sequential deep learning for multiple reaction monitoring. <i>Journal of Proteomics</i> , 2020, 223, 103820.	1.2	18
8	Metastasis of cholangiocarcinoma is promoted by extended high-mannose glycans. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 7633-7644.	3.3	63
9	Deep Structural Analysis and Quantitation of O-Linked Glycans on Cell Membrane Reveal High Abundances and Distinct Glycomic Profiles Associated with Cell Type and Stages of Differentiation. <i>Analytical Chemistry</i> , 2020, 92, 3758-3768.	3.2	23
10	Metabolic flux analysis of the neural cell glycolyx reveals differential utilization of monosaccharides. <i>Glycobiology</i> , 2020, 30, 859-871.	1.3	15
11	The DNA repair enzyme MUTYH potentiates cytotoxicity of the alkylating agent MNNG by interacting with abasic sites. <i>Journal of Biological Chemistry</i> , 2020, 295, 3692-3707.	1.6	10
12	Unveiling the metabolic fate of monosaccharides in cell membranes with glycomic and glycoproteomic analyses. <i>Chemical Science</i> , 2019, 10, 6992-7002.	3.7	19
13	Human Milk Proteins and Their Glycosylation Exhibit Quantitative Dynamic Variations during Lactation. <i>Journal of Nutrition</i> , 2019, 149, 1317-1325.	1.3	41
14	Identification of potential sialic acid binding proteins on cell membranes by proximity chemical labeling. <i>Chemical Science</i> , 2019, 10, 6199-6209.	3.7	33
15	Site-Specific Glycosylation Quantitation of 50 Serum Glycoproteins Enhanced by Predictive Glycopeptidomics for Improved Disease Biomarker Discovery. <i>Analytical Chemistry</i> , 2019, 91, 5433-5445.	3.2	41
16	Infection-generated electric field in gut epithelium drives bidirectional migration of macrophages. <i>PLoS Biology</i> , 2019, 17, e3000044.	2.6	28
17	A rapid-throughput adaptable method for determining the monosaccharide composition of polysaccharides. <i>International Journal of Mass Spectrometry</i> , 2019, 438, 22-28.	0.7	36
18	Genetic Ablation of Butyrate Utilization Attenuates Gastrointestinal Salmonella Disease. <i>Cell Host and Microbe</i> , 2018, 23, 266-273.e4.	5.1	48

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19	Biallelic Mutations in FUT8 Cause a Congenital Disorder of Glycosylation with Defective Fucosylation. <i>American Journal of Human Genetics</i> , 2018, 102, 188-195.	2.6	49
20	Mass Spectrometry Approaches to Glycomic and Glycoproteomic Analyses. <i>Chemical Reviews</i> , 2018, 118, 7886-7930.	23.0	277
21	Composition and Variation of Macronutrients, Immune Proteins, and Human Milk Oligosaccharides in Human Milk From Nonprofit and Commercial Milk Banks. <i>Journal of Human Lactation</i> , 2018, 34, 120-129.	0.8	55
22	Recent Advances in the Mass Spectrometry Methods for Glycomics and Cancer. <i>Analytical Chemistry</i> , 2018, 90, 208-224.	3.2	64
23	Revisiting monosaccharide analysis – quantitation of a comprehensive set of monosaccharides using dynamic multiple reaction monitoring. <i>Analyst</i> , The, 2018, 143, 200-207.	1.7	60
24	Membrane glycomics reveal heterogeneity and quantitative distribution of cell surface sialylation. <i>Chemical Science</i> , 2018, 9, 6271-6285.	3.7	42
25	FGF2 Induces Migration of Human Bone Marrow Stromal Cells by Increasing Core Fucosylations on N-Glycans of Integrins. <i>Stem Cell Reports</i> , 2018, 11, 325-333.	2.3	25
26	Intact glycosphingolipidomic analysis of the cell membrane during differentiation yields extensive glycan and lipid changes. <i>Scientific Reports</i> , 2018, 8, 10993.	1.6	16
27	Bifidobacterium grown on human milk oligosaccharides produce tryptophan metabolite Indole-3-lactic acid that significantly decreases inflammation in intestinal cells in vitro. <i>FASEB Journal</i> , 2018, 32, 1b359.	0.2	20
28	System Metaglycomes: Mapping Dynamic Cell Surface N-glycome, O-glycome and Glycolipidome by Mass Spectrometry. <i>FASEB Journal</i> , 2018, 32, 673.11.	0.2	1
29	Enterocyte glycosylation is responsive to changes in extracellular conditions: implications for membrane functions. <i>Glycobiology</i> , 2017, 27, 847-860.	1.3	31
30	Microbiota-activated PPAR- $\beta$ signaling inhibits dysbiotic Enterobacteriaceae expansion. <i>Science</i> , 2017, 357, 570-575.	6.0	796
31	Persistence of Supplemented Bifidobacterium longum subsp. <i>infantis</i> EVC001 in Breastfed Infants. <i>MSphere</i> , 2017, 2, .	1.3	158
32	Absolute Quantitation of Human Milk Oligosaccharides Reveals Phenotypic Variations during Lactation. <i>Journal of Nutrition</i> , 2017, 147, 117-124.	1.3	122
33	Depletion of Butyrate-Producing Clostridia from the Gut Microbiota Drives an Aerobic Luminal Expansion of Salmonella. <i>Cell Host and Microbe</i> , 2016, 19, 443-454.	5.1	600
34	Graphene matrix for signal enhancement in ambient plasma assisted laser desorption ionization mass spectrometry. <i>Talanta</i> , 2013, 114, 54-59.	2.9	17
35	Online Coupling of Capillary Electrophoresis with Direct Analysis in Real Time Mass Spectrometry. <i>Analytical Chemistry</i> , 2013, 85, 170-176.	3.2	49