Adam Byron

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Integrin ligands at a glance. Journal of Cell Science, 2006, 119, 3901-3903.	2.0	1,393
2	Definition of a consensus integrin adhesome and its dynamics during adhesion complex assembly andÂdisassembly. Nature Cell Biology, 2015, 17, 1577-1587.	10.3	442
3	Nuclear FAK Controls Chemokine Transcription, Tregs, and Evasion of Anti-tumor Immunity. Cell, 2015, 163, 160-173.	28.9	304
4	Proteomic Analysis of Integrin-Associated Complexes Identifies RCC2 as a Dual Regulator of Rac1 and Arf6. Science Signaling, 2009, 2, ra51.	3.6	220
5	Glioblastomas acquire myeloid-affiliated transcriptional programs via epigenetic immunoediting to elicit immune evasion. Cell, 2021, 184, 2454-2470.e26.	28.9	165
6	Global Analysis Reveals the Complexity of the Human Glomerular Extracellular Matrix. Journal of the American Society of Nephrology: JASN, 2014, 25, 939-951.	6.1	158
7	Anti-integrin monoclonal antibodies. Journal of Cell Science, 2009, 122, 4009-4011.	2.0	153
8	Defining the phospho-adhesome through the phosphoproteomic analysis of integrin signalling. Nature Communications, 2015, 6, 6265.	12.8	150
9	Defining the extracellular matrix using proteomics. International Journal of Experimental Pathology, 2013, 94, 75-92.	1.3	137
10	A Syndecan-4 Hair Trigger Initiates Wound Healing through Caveolin- and RhoG-Regulated Integrin Endocytosis. Developmental Cell, 2011, 21, 681-693.	7.0	115
11	Giving off mixed signals—Distinct functions of α ₅ 1² ₁ and α _v 1² ₃ integrins in regulating cell behaviour. IUBMB Life, 2009, 61, 731-738.	3.4	96
12	Glomerular Cell Cross-Talk Influences Composition and Assembly of Extracellular Matrix. Journal of the American Society of Nephrology: JASN, 2014, 25, 953-966.	6.1	88
13	A proteomic approach reveals integrin activation state-dependent control of microtubule cortical targeting. Nature Communications, 2015, 6, 6135.	12.8	71
14	Rac1 is deactivated at integrin activation sites via an IQGAP1/filamin-A/RacGAP1 pathway. Journal of Cell Science, 2013, 126, 4121-35.	2.0	68
15	Proteomic analysis of extracellular matrix from the hepatic stellate cell line LX-2 identifies CYR61 and Wnt-5a as novel constituents of fibrotic liver. Journal of Proteome Research, 2012, 11, 4052-4064.	3.7	66
16	IL-33 and ST2 mediate FAK-dependent antitumor immune evasion through transcriptional networks. Science Signaling, 2017, 10, .	3.6	64
17	E-cadherin loss induces targetable autocrine activation of growth factor signalling in lobular breast cancer. Scientific Reports, 2018, 8, 15454.	3.3	55
18	Proteomic analysis of α4β1 integrin adhesion complexes reveals αâ€subunitâ€dependent protein recruitment. Proteomics, 2012, 12, 2107-2114.	2.2	52

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19	Adhesion signalling complexes. Current Biology, 2010, 20, R1063-R1067.	3.9	50
20	Comparative Proteomic Analysis of Supportive and Unsupportive Extracellular Matrix Substrates for Human Embryonic Stem Cell Maintenance. Journal of Biological Chemistry, 2013, 288, 18716-18731.	3.4	50
21	Isolation of Integrinâ€Based Adhesion Complexes. Current Protocols in Cell Biology, 2015, 66, 9.8.1-9.8.15.	2.3	48
22	Nuclear FAK and Runx1 Cooperate to Regulate IGFBP3, Cell-Cycle Progression, and Tumor Growth. Cancer Research, 2017, 77, 5301-5312.	0.9	48
23	Structural basis of Focal Adhesion Kinase activation on lipid membranes. EMBO Journal, 2020, 39, e104743.	7.8	47
24	Proteomic Analysis of Integrin Adhesion ComplexesA presentation from the 6th British Society for Proteome Research (BSPR)–European Bioinformatics Institute (EBI) Meeting "Multiscale Proteomics: From Cells to Organisms―at the Wellcome Trust Conference Centre, Cambridge, UK, 14 to 16 July 2009. The Presentation also complements the <i>Science Signaling</i> Research Article by Humphries <i>et al.</i> published 8 September 2009 Science Signaling, 2011, 4, pt2.	3.6	45
25	Adhesion protein networks reveal functions proximal and distal to cell-matrix contacts. Current Opinion in Cell Biology, 2016, 39, 93-100.	5.4	42
26	mTORC1 activity is supported by spatial association with focal adhesions. Journal of Cell Biology, 2021, 220, .	5.2	41
27	Genetic Background is a Key Determinant of Glomerular Extracellular Matrix Composition and Organization. Journal of the American Society of Nephrology: JASN, 2015, 26, 3021-3034.	6.1	39
28	Basement membrane ligands initiate distinct signalling networks to direct cell shape. Matrix Biology, 2020, 90, 61-78.	3.6	38
29	Microtubule-Dependent Modulation of Adhesion Complex Composition. PLoS ONE, 2014, 9, e115213.	2.5	34
30	Ambra1 spatially regulates Src activity and Src/FAK-mediated cancer cell invasion via trafficking networks. ELife, 2017, 6, .	6.0	32
31	Proteomic analysis of integrinâ€associated complexes from mesenchymal stem cells. Proteomics - Clinical Applications, 2016, 10, 51-57.	1.6	31
32	Identification of novel pathways linking epithelial-to-mesenchymal transition with resistance to HER2-targeted therapy. Oncotarget, 2016, 7, 11539-11552.	1.8	27
33	Analyzing the Anatomy of Integrin Adhesions. Science Signaling, 2011, 4, jc3.	3.6	26
34	Mapping the ligand-binding pocket of integrin α5β1 using a gain-of-function approach. Biochemical Journal, 2009, 424, 179-189.	3.7	24
35	The effect of peptide adsorption on signal linearity and a simple approach to improve reliability of quantification. Journal of Proteomics, 2013, 85, 160-164.	2.4	21

Exploring mechanisms of acquired resistance to HER2 (human epidermal growth factor receptor) Tj ETQq0 0 0 rgBT₃/Qverlock 10 Tf 50 6

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37	Kindlin-1 Promotes Pulmonary Breast Cancer Metastasis. Cancer Research, 2018, 78, 1484-1496.	0.9	17
38	Ligand-induced Epitope Masking. Journal of Biological Chemistry, 2016, 291, 20993-21007.	3.4	16
39	A Synergistic Anticancer FAK and HDAC Inhibitor Combination Discovered by a Novel Chemical–Genetic High-Content Phenotypic Screen. Molecular Cancer Therapeutics, 2020, 19, 637-649.	4.1	16
40	FAK regulates IL-33 expression by controlling chromatin accessibility at c-Jun motifs. Scientific Reports, 2021, 11, 229.	3.3	14
41	The autophagy protein Ambra1 regulates gene expression by supporting novel transcriptional complexes. Journal of Biological Chemistry, 2020, 295, 12045-12057.	3.4	13
42	Characterization of the Phospho-Adhesome by Mass Spectrometry-Based Proteomics. Methods in Molecular Biology, 2017, 1636, 235-251.	0.9	13
43	Characterisation of the Stromal Microenvironment in Lobular Breast Cancer. Cancers, 2022, 14, 904.	3.7	13
44	Trafficking of Adhesion and Growth Factor Receptors and Their Effector Kinases. Annual Review of Cell and Developmental Biology, 2018, 34, 29-58.	9.4	11
45	Proteomic Profiling of Integrin Adhesion Complex Assembly. Methods in Molecular Biology, 2018, 1764, 193-236.	0.9	10
46	Novel roles of PRK1 and PRK2 in cilia and cancer biology. Scientific Reports, 2020, 10, 3902.	3.3	10
47	Integrative analysis of multi-platform reverse-phase protein array data for the pharmacodynamic assessment of response to targeted therapies. Scientific Reports, 2020, 10, 21985.	3.3	9
48	Network Analysis of Integrin Adhesion Complexes. Methods in Molecular Biology, 2021, 2217, 149-179.	0.9	7
49	Reproducibility and Crossplatform Validation of Reverse-Phase Protein Array Data. Advances in Experimental Medicine and Biology, 2019, 1188, 181-201.	1.6	7
50	Clustering and Network Analysis of Reverse Phase Protein Array Data. Methods in Molecular Biology, 2017, 1606, 171-191.	0.9	6
51	Loss of Integrin-Linked Kinase Sensitizes Breast Cancer to SRC Inhibitors. Cancer Research, 2022, 82, 632-647.	0.9	6
52	Utilisation of the budding yeastSaccharomyces cerevisiae for the generation and isolation of non-lethal ricin A chain variants. Yeast, 2005, 22, 1287-1297.	1.7	5
53	Alternative cellular roles for proteins identified using proteomics. Journal of Proteomics, 2012, 75, 4184-4185.	2.4	5
54	Evaluation of Gene Expression Data From Cybrids and Tumours Highlights Elevated NDRG1-Driven Proliferation in Triple-Negative Breast Cancer. Breast Cancer: Basic and Clinical Research, 2020, 14, 117822342093444.	1.1	5

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55	Characterisation of a nucleo-adhesome. Nature Communications, 2022, 13, .	12.8	4
56	Regulation of Cell-Matrix Adhesion Networks: Insights from Proteomics. Biology of Extracellular Matrix, 2020, , 183-208.	0.3	2