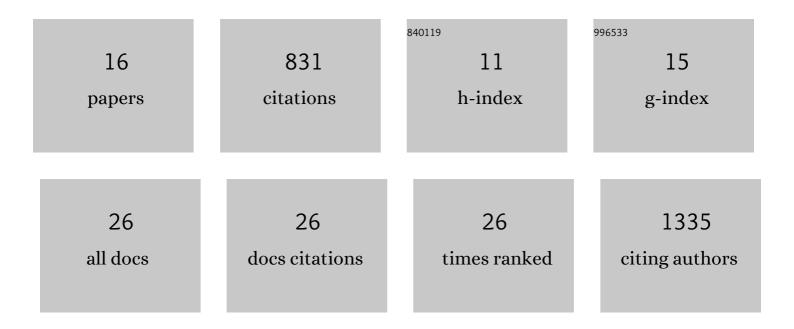
## Sean R Collins

List of Publications by Year in descending order

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SEAN R COLLINS

#	Article	IF	CITATIONS
1	WASP integrates substrate topology and cell polarity to guide neutrophil migration. Journal of Cell Biology, 2022, 221, .	2.3	28
2	Host cells subdivide nutrient niches into discrete biogeographical microhabitats for gut microbes. Cell Host and Microbe, 2022, 30, 836-847.e6.	5.1	29
3	Optimized iLID Membrane Anchors for Local Optogenetic Protein Recruitment. ACS Synthetic Biology, 2021, 10, 1009-1023.	1.9	23
4	SUMO is a pervasive regulator of meiosis. ELife, 2021, 10, .	2.8	50
5	Directional reorientation of migrating neutrophils is limited by suppression of receptor input signaling at the cell rear through myosin II activity. Nature Communications, 2021, 12, 6619.	5.8	27
6	Optogenetic control of receptors reveals distinct roles for actin- and Cdc42-dependent negative signals in chemotactic signal processing. Nature Communications, 2021, 12, 6148.	5.8	14
7	Decoding GEFs of animated cells. Nature Chemical Biology, 2020, 16, 812-813.	3.9	0
8	Efficient Front-Rear Coupling in Neutrophil Chemotaxis by Dynamic Myosin II Localization. Developmental Cell, 2019, 49, 189-205.e6.	3.1	59
9	Identification of phagocytosis regulators using magnetic genome-wide CRISPR screens. Nature Genetics, 2018, 50, 1716-1727.	9.4	135
10	A map of gene expression in neutrophil-like cell lines. BMC Genomics, 2018, 19, 573.	1.2	100
11	Mass Spectrometric Identification of Urinary Biomarkers of Pulmonary Tuberculosis. EBioMedicine, 2018, 31, 157-165.	2.7	46
12	"Rhoâ€ing a Cellular Boat with Rearward Membrane Flow. Developmental Cell, 2018, 46, 1-3.	3.1	12
13	Parallel High-Resolution Imaging of Leukocyte Chemotaxis Under Agarose with Rho-Family GTPase Biosensors. Methods in Molecular Biology, 2018, 1821, 71-85.	0.4	11
14	Locally excitable Cdc42 signals steer cells duringÂchemotaxis. Nature Cell Biology, 2016, 18, 191-201.	4.6	166
15	Using light to shape chemical gradients for parallel and automated analysis of chemotaxis. Molecular Systems Biology, 2015, 11, 804.	3.2	38
16	Quantitative Genetic Interaction Mapping Using the E-MAP Approach. Methods in Enzymology, 2010, 470, 205-231.	0.4	80