

# Kathryn M Ferguson

## List of Publications by Year in descending order

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Version: 2024-02-01

21  
papers

4,597  
citations

516710

16  
h-index

839539

18  
g-index

22  
all docs

22  
docs citations

22  
times ranked

5791  
citing authors

#	ARTICLE	IF	CITATIONS
1	Glioblastoma mutations alter EGFR dimer structure to prevent ligand bias. <i>Nature</i> , 2022, 602, 518-522.	27.8	36
2	Insulin and epidermal growth factor receptor family members share parallel activation mechanisms. <i>Protein Science</i> , 2020, 29, 1331-1344.	7.6	31
3	The Mechanism of Ligand-Induced Activation of the Tie Family of Receptor Tyrosine Kinases. <i>FASEB Journal</i> , 2019, 33, 809.10.	0.5	0
4	Molecular Basis for Necitumumab Inhibition of EGFR Variants Associated with Acquired Cetuximab Resistance. <i>Molecular Cancer Therapeutics</i> , 2018, 17, 521-531.	4.1	45
5	Structural Basis for MARK1 Kinase Autoinhibition by Its KA1 Domain. <i>Structure</i> , 2018, 26, 1137-1143.e3.	3.3	15
6	Dimerization of Tie2 mediated by its membrane-proximal FNIII domains. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, 4382-4387.	7.1	29
7	Molecular determinants of KA1 domain-mediated autoinhibition and phospholipid activation of MARK1 kinase. <i>Biochemical Journal</i> , 2017, 474, 385-398.	3.7	21
8	EGFR Ligands Differentially Stabilize Receptor Dimers to Specify Signaling Kinetics. <i>Cell</i> , 2017, 171, 683-695.e18.	28.9	276
9	Intramolecular autoinhibition of checkpoint kinase 1 is mediated by conserved basic motifs of the C-terminal kinase-associated 1 domain. <i>Journal of Biological Chemistry</i> , 2017, 292, 19024-19033.	3.4	15
10	Comparison of <i>Saccharomyces cerevisiae</i> F-BAR Domain Structures Reveals a Conserved Inositol Phosphate Binding Site. <i>Structure</i> , 2015, 23, 352-363.	3.3	40
11	Complex Relationship between Ligand Binding and Dimerization in the Epidermal Growth Factor Receptor. <i>Cell Reports</i> , 2014, 9, 1306-1317.	6.4	78
12	The EGFR Family: Not So Prototypical Receptor Tyrosine Kinases. <i>Cold Spring Harbor Perspectives in Biology</i> , 2014, 6, a020768-a020768.	5.5	345
13	Discoidin Discoveries. <i>Structure</i> , 2012, 20, 568-570.	3.3	0
14	Structural aspects of extracellular EGFR signaling. <i>FASEB Journal</i> , 2009, 23, 198.3.	0.5	0
15	Structure-Based View of Epidermal Growth Factor Receptor Regulation. <i>Annual Review of Biophysics</i> , 2008, 37, 353-373.	10.0	306
16	Structural basis for inhibition of the epidermal growth factor receptor by cetuximab. <i>Cancer Cell</i> , 2005, 7, 301-311.	16.8	949
17	Epidermal Growth Factor Receptor Dimerization and Activation Require Ligand-Induced Conformational Changes in the Dimer Interface. <i>Molecular and Cellular Biology</i> , 2005, 25, 7734-7742.	2.3	247
18	EGF Activates Its Receptor by Removing Interactions that Autoinhibit Ectodomain Dimerization. <i>Molecular Cell</i> , 2003, 11, 507-517.	9.7	675

#	ARTICLE	IF	CITATIONS
19	An Open-and-Shut Case? Recent Insights into the Activation of EGF/ErbB Receptors. <i>Molecular Cell</i> , 2003, 12, 541-552.	9.7	774
20	Signal-dependent membrane targeting by pleckstrin homology (PH) domains. <i>Biochemical Journal</i> , 2000, 350, 1-18.	3.7	656
21	Scratching the surface with the PH domain. <i>Nature Structural and Molecular Biology</i> , 1995, 2, 715-718.	8.2	59