

Do-Hyung Kim

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57
papers

17,938
citations

31
h-index

59
g-index

59
ext. papers

19,800
ext. citations

7.4
avg, IF

5.72
L-index

#	Paper	IF	Citations
57	Guidelines for the use and interpretation of assays for monitoring autophagy (3rd edition). <i>Autophagy</i> , 2016 , 12, 1-222	10.2	3838
56	Guidelines for the use and interpretation of assays for monitoring autophagy. <i>Autophagy</i> , 2012 , 8, 445-546	10.2	2783
55	mTOR interacts with raptor to form a nutrient-sensitive complex that signals to the cell growth machinery. <i>Cell</i> , 2002 , 110, 163-75	56.2	2322
54	Rictor, a novel binding partner of mTOR, defines a rapamycin-insensitive and raptor-independent pathway that regulates the cytoskeleton. <i>Current Biology</i> , 2004 , 14, 1296-302	6.3	2067
53	mTOR regulation of autophagy. <i>FEBS Letters</i> , 2010 , 584, 1287-95	3.8	1495
52	ULK-Atg13-FIP200 complexes mediate mTOR signaling to the autophagy machinery. <i>Molecular Biology of the Cell</i> , 2009 , 20, 1992-2003	3.5	1434
51	Insulin signalling to mTOR mediated by the Akt/PKB substrate PRAS40. <i>Nature Cell Biology</i> , 2007 , 9, 316-234	2.34	897
50	GbetaL, a positive regulator of the rapamycin-sensitive pathway required for the nutrient-sensitive interaction between raptor and mTOR. <i>Molecular Cell</i> , 2003 , 11, 895-904	17.6	747
49	Local structural elements in the mostly unstructured transcriptional activation domain of human p53. <i>Journal of Biological Chemistry</i> , 2000 , 275, 29426-32	5.4	257
48	Hsp90-Cdc37 chaperone complex regulates Ulk1- and Atg13-mediated mitophagy. <i>Molecular Cell</i> , 2011 , 43, 572-85	17.6	173
47	The ULK1 complex mediates MTORC1 signaling to the autophagy initiation machinery via binding and phosphorylating ATG14. <i>Autophagy</i> , 2016 , 12, 547-64	10.2	163
46	mTORC1 phosphorylates UVRAG to negatively regulate autophagosome and endosome maturation. <i>Molecular Cell</i> , 2015 , 57, 207-18	17.6	162
45	PRR5, a novel component of mTOR complex 2, regulates platelet-derived growth factor receptor beta expression and signaling. <i>Journal of Biological Chemistry</i> , 2007 , 282, 25604-12	5.4	155
44	Epigenetic regulation of autophagy by the methyltransferase G9a. <i>Molecular and Cellular Biology</i> , 2013 , 33, 3983-93	4.8	133
43	Glycolytic flux signals to mTOR through glyceraldehyde-3-phosphate dehydrogenase-mediated regulation of Rheb. <i>Molecular and Cellular Biology</i> , 2009 , 29, 3991-4001	4.8	127
42	ULK1 inhibits the kinase activity of mTORC1 and cell proliferation. <i>Autophagy</i> , 2011 , 7, 1212-21	10.2	122
41	Transactivation ability of p53 transcriptional activation domain is directly related to the binding affinity to TATA-binding protein. <i>Journal of Biological Chemistry</i> , 1995 , 270, 25014-9	5.4	96

40	ULK1 phosphorylates Ser30 of BECN1 in association with ATG14 to stimulate autophagy induction. <i>Autophagy</i> , 2018 , 14, 584-597	10.2	73
39	SH3BP4 is a negative regulator of amino acid-Rag GTPase-mTORC1 signaling. <i>Molecular Cell</i> , 2012 , 46, 833-46	17.6	67
38	Distinct functions of Ulk1 and Ulk2 in the regulation of lipid metabolism in adipocytes. <i>Autophagy</i> , 2013 , 9, 2103-14	10.2	56
37	Cyclic AMP controls mTOR through regulation of the dynamic interaction between Rheb and phosphodiesterase 4D. <i>Molecular and Cellular Biology</i> , 2010 , 30, 5406-20	4.8	50
36	Crystal structure of the Gtr1p(GTP)-Gtr2p(GDP) protein complex reveals large structural rearrangements triggered by GTP-to-GDP conversion. <i>Journal of Biological Chemistry</i> , 2012 , 287, 29648-53	5.4	50
35	mTORC1 Coordinates Protein Synthesis and Immunoproteasome Formation via PRAS40 to Prevent Accumulation of Protein Stress. <i>Molecular Cell</i> , 2016 , 61, 625-639	17.6	46
34	PLD2 forms a functional complex with mTOR/raptor to transduce mitogenic signals. <i>Cellular Signalling</i> , 2006 , 18, 2283-91	4.9	45
33	Asp-99 donates a hydrogen bond not to Tyr-14 but to the steroid directly in the catalytic mechanism of Delta 5-3-ketosteroid isomerase from <i>Pseudomonas putida</i> biotype B. <i>Biochemistry</i> , 2000 , 39, 903-9	3.2	43
32	Contribution of the hydrogen-bond network involving a tyrosine triad in the active site to the structure and function of a highly proficient ketosteroid isomerase from <i>Pseudomonas putida</i> biotype B. <i>Biochemistry</i> , 2000 , 39, 4581-9	3.2	40
31	mRNA 3'UTR shortening is a molecular signature of mTORC1 activation. <i>Nature Communications</i> , 2015 , 6, 7218	17.4	37
30	Hsf1 activation inhibits rapamycin resistance and TOR signaling in yeast revealed by combined proteomic and genetic analysis. <i>PLoS ONE</i> , 2008 , 3, e1598	3.7	37
29	The role of Tyr248 probed by mutant bovine carboxypeptidase A: insight into the catalytic mechanism of carboxypeptidase A. <i>Biochemistry</i> , 2001 , 40, 10197-203	3.2	35
28	Role of catalytic residues in enzymatic mechanisms of homologous ketosteroid isomerases. <i>Biochemistry</i> , 2000 , 39, 13891-6	3.2	34
27	GABARAPs and LC3s have opposite roles in regulating ULK1 for autophagy induction. <i>Autophagy</i> , 2020 , 16, 600-614	10.2	32
26	Quantitative nuclear proteomics identifies mTOR regulation of DNA damage response. <i>Molecular and Cellular Proteomics</i> , 2010 , 9, 403-14	7.6	31
25	¹⁵ N NMR relaxation studies of backbone dynamics in free and steroid-bound Delta 5-3-ketosteroid isomerase from <i>Pseudomonas testosteroni</i> . <i>Biochemistry</i> , 2001 , 40, 3967-73	3.2	30
24	Expression, purification, and identification of a novel self-cleavage site of the Nla C-terminal 27-kDa protease of turnip mosaic potyvirus C5. <i>Virology</i> , 1995 , 213, 517-25	3.6	26
23	Effects of internal cleavages and mutations in the C-terminal region of Nla protease of turnip mosaic potyvirus on the catalytic activity. <i>Virology</i> , 1996 , 226, 183-90	3.6	23

22	Uncoordinated 51-like kinase 2 signaling pathway regulates epithelial-mesenchymal transition in A549 lung cancer cells. <i>FEBS Letters</i> , 2016 , 590, 1365-74	3.8	22
21	Roles of active site aromatic residues in catalysis by ketosteroid isomerase from <i>Pseudomonas putida</i> biotype B. <i>Biochemistry</i> , 1999 , 38, 13810-9	3.2	19
20	Folding mechanism of ketosteroid isomerase from <i>Comamonas testosteroni</i> . <i>Biochemistry</i> , 2001 , 40, 5011-7	3.2	17
19	Unconventional Secretion of Adipocyte Fatty Acid Binding Protein 4 Is Mediated By Autophagic Proteins in a Sirtuin-1-Dependent Manner. <i>Diabetes</i> , 2019 , 68, 1767-1777	0.9	16
18	Roles of dimerization in folding and stability of ketosteroid isomerase from <i>Pseudomonas putida</i> biotype B. <i>Protein Science</i> , 2001 , 10, 741-52	6.3	16
17	dRAGging amino acid-mTORC1 signaling by SH3BP4. <i>Molecules and Cells</i> , 2013 , 35, 1-6	3.5	14
16	Equilibrium and kinetic analysis of folding of ketosteroid isomerase from <i>Comamonas testosteroni</i> . <i>Biochemistry</i> , 2000 , 39, 13084-92	3.2	14
15	Characterization of Nla protease from turnip mosaic potyvirus exhibiting a low-temperature optimum catalytic activity. <i>Virology</i> , 1996 , 221, 245-9	3.6	14
14	-(1-Benzyl-3,5-dimethyl-1-pyrazol-4-yl)benzamides: Antiproliferative Activity and Effects on mTORC1 and Autophagy. <i>ACS Medicinal Chemistry Letters</i> , 2017 , 8, 90-95	4.3	10
13	Maintenance of alpha-helical structures by phenyl rings in the active-site tyrosine triad contributes to catalysis and stability of ketosteroid isomerase from <i>Pseudomonas putida</i> biotype B. <i>Biochemistry</i> , 2001 , 40, 13529-37	3.2	10
12	Effects of mutations in the C-terminal region of Nla protease on cis-cleavage between Nla and Nlb. <i>Virology</i> , 1998 , 241, 94-100	3.6	9
11	Characterization of active-site residues of the Nla protease from tobacco vein mottling virus. <i>Molecules and Cells</i> , 2000 , 10, 505-11	3.5	9
10	GbetaL regulates TNFalpha-induced NF-kappaB signaling by directly inhibiting the activation of IkkappaB kinase. <i>Cellular Signalling</i> , 2008 , 20, 2127-33	4.9	8
9	Molecular cloning, expression, and purification of nuclear inclusion A protease from tobacco vein mottling virus. <i>Molecules and Cells</i> , 2000 , 10, 148-55	3.5	7
8	A Novel Mechanism for NF- κ B-activation via I κ B-aggregation: Implications for Hepatic Mallory-Denk-Body Induced Inflammation. <i>Molecular and Cellular Proteomics</i> , 2020 , 19, 1968-1986	7.6	6
7	Down regulation of Peroxiredoxin-3 in 3T3-L1 adipocytes leads to oxidation of Rictor in the mammalian-target of rapamycin complex 2 (mTORC2). <i>Biochemical and Biophysical Research Communications</i> , 2017 , 493, 1311-1317	3.4	5
6	An expanded role for mTORC1 in autophagy. <i>Molecular and Cellular Oncology</i> , 2016 , 3, e1010958	1.2	5
5	Contribution of conserved amino acids at the dimeric interface to the conformational stability and the structural integrity of the active site in ketosteroid isomerase from <i>Pseudomonas putida</i> biotype B. <i>Journal of Biochemistry</i> , 2003 , 134, 101-10	3.1	5

- 4 Temperature and salt effects on proteolytic function of turnip mosaic potyvirus nuclear inclusion protein a exhibiting a low-temperature optimum activity. *BBA - Proteins and Proteomics*, **2000**, 1480, 29-40 5
- 3 Defective autophagy and increased apoptosis contribute toward the pathogenesis of FKRP-associated muscular dystrophies. *Stem Cell Reports*, **2021**, 16, 2752-2767 8 1
- 2 Potyvirus NIa Protease **2013**, 2427-2432
- 1 Immunoproteasome Inhibition to Target AML with Activated RAS Pathways. *Blood*, **2016**, 128, 577-577 2.2