

Mehdi Mollapour

List of Publications by Year in descending order

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Version: 2024-02-01

87
papers

6,058
citations

66343

42
h-index

71685

76
g-index

102
all docs

102
docs citations

102
times ranked

6640
citing authors

#	ARTICLE	IF	CITATIONS
1	Seventh BHD international symposium: recent scientific and clinical advancement. <i>Oncotarget</i> , 2022, 13, 173-181.	1.8	4
2	Therapeutic potential of CDK4/6 inhibitors in renal cell carcinoma. <i>Nature Reviews Urology</i> , 2022, 19, 305-320.	3.8	9
3	TRAP1 Chaperones the Metabolic Switch in Cancer. <i>Biomolecules</i> , 2022, 12, 786.	4.0	14
4	Emerging Link between Tsc1 and FNIP Co-Chaperones of Hsp90 and Cancer. <i>Biomolecules</i> , 2022, 12, 928.	4.0	2
5	A specialized Hsp90 co-chaperone network regulates steroid hormone receptor response to ligand. <i>Cell Reports</i> , 2022, 40, 111039.	6.4	15
6	The Role of Heat Shock Protein-90 in the Pathogenesis of Birt-Hogg-Dubé and Tuberous Sclerosis Complex Syndromes. <i>Urologic Oncology: Seminars and Original Investigations</i> , 2021, 39, 322-326.	1.6	6
7	MMPs, tyrosine kinase signaling and extracellular matrix proteolysis in kidney cancer. <i>Urologic Oncology: Seminars and Original Investigations</i> , 2021, 39, 316-321.	1.6	9
8	First Virtual International Congress on Cellular and Organismal Stress Responses, November 5-6, 2020. <i>Cell Stress and Chaperones</i> , 2021, 26, 289-295.	2.9	0
9	Comprehensive genomic profiling of metastatic collecting duct carcinoma, renal medullary carcinoma, and clear cell renal cell carcinoma. <i>Urologic Oncology: Seminars and Original Investigations</i> , 2021, 39, 367.e1-367.e5.	1.6	11
10	Clinically Advanced Pheochromocytomas and Paragangliomas: A Comprehensive Genomic Profiling Study. <i>Cancers</i> , 2021, 13, 3312.	3.7	9
11	The tumor suppressor folliculin inhibits lactate dehydrogenase A and regulates the Warburg effect. <i>Nature Structural and Molecular Biology</i> , 2021, 28, 662-670.	8.2	19
12	Hsp90 chaperone code and the tumor suppressor VHL cooperatively regulate the mitotic checkpoint. <i>Cell Stress and Chaperones</i> , 2021, 26, 965-971.	2.9	9
13	Comprehensive genomic profiling of histologic subtypes of urethral carcinomas. <i>Urologic Oncology: Seminars and Original Investigations</i> , 2021, 39, 731.e1-731.e15.	1.6	7
14	Decrypting the chaperone code. <i>Journal of Biological Chemistry</i> , 2021, 296, 100293.	3.4	12
15	Fumarate hydratase as a therapeutic target in renal cancer. <i>Expert Opinion on Therapeutic Targets</i> , 2020, 24, 923-936.	3.4	12
16	Post-translational modifications of Hsp90 and translating the chaperone code. <i>Journal of Biological Chemistry</i> , 2020, 295, 11099-11117.	3.4	115
17	Structure and function of the co-chaperone protein phosphatase 5 in cancer. <i>Cell Stress and Chaperones</i> , 2020, 25, 383-394.	2.9	28
18	Chemical Perturbation of Oncogenic Protein Folding: from the Prediction of Locally Unstable Structures to the Design of Disruptors of Hsp90-Client Interactions. <i>Chemistry - A European Journal</i> , 2020, 26, 9459-9465.	3.3	39

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19	Structural and functional regulation of lactate dehydrogenase-A in cancer. <i>Future Medicinal Chemistry</i> , 2020, 12, 439-455.	2.3	33
20	Design of Disruptors of the Hsp90â€Cdc37 Interface. <i>Molecules</i> , 2020, 25, 360.	3.8	14
21	Clinically advanced renal cell carcinoma (RCC) and renal sarcoma (RSC) in young patients: A comprehensive genomic profiling (CGP) study.. <i>Journal of Clinical Oncology</i> , 2020, 38, 5066-5066.	1.6	0
22	Co-chaperones TIMP2 and AHA1 Competitively Regulate Extracellular HSP90:Client MMP2 Activity and Matrix Proteolysis. <i>Cell Reports</i> , 2019, 28, 1894-1906.e6.	6.4	50
23	Post-translational Regulation of FNIP1 Creates a Rheostat for the Molecular Chaperone Hsp90. <i>Cell Reports</i> , 2019, 26, 1344-1356.e5.	6.4	38
24	Renal cell carcinoma and brain metastasis: Questioning the dogma of role for cytoreductive nephrectomy. <i>Urologic Oncology: Seminars and Original Investigations</i> , 2019, 37, 182.e9-182.e15.	1.6	10
25	The multiple facets of the Hsp90 machine. <i>Nature Structural and Molecular Biology</i> , 2019, 26, 92-95.	8.2	9
26	Mutation of the co-chaperone Tsc1 in bladder cancer diminishes Hsp90 acetylation and reduces drug sensitivity and selectivity. <i>Oncotarget</i> , 2019, 10, 5824-5834.	1.8	18
27	Extracellular Phosphorylation of TIMP-2 by Secreted c-Src Tyrosine Kinase Controls MMP-2 Activity. <i>IScience</i> , 2018, 1, 87-96.	4.1	29
28	Detecting Posttranslational Modifications of Hsp90. <i>Methods in Molecular Biology</i> , 2018, 1709, 209-219.	0.9	13
29	Detection and Analysis of Extracellular Hsp90 (eHsp90). <i>Methods in Molecular Biology</i> , 2018, 1709, 321-329.	0.9	9
30	The mTOR Independent Function of Tsc1 and FNIPs. <i>Trends in Biochemical Sciences</i> , 2018, 43, 935-937.	7.5	14
31	<i>PBRM1</i> mutation and immunotherapy efficacy: A comprehensive genomic profiling (CGP) assessment.. <i>Journal of Clinical Oncology</i> , 2018, 36, 12091-12091.	1.6	4
32	Sporadic renal angiomyolipoma in a patient with Birt-Hogg-DubÃ©: chaperones in pathogenesis. <i>Oncotarget</i> , 2018, 9, 22220-22229.	1.8	11
33	Carcinomas of the renal medulla: A comprehensive genomic profiling (CGP) study.. <i>Journal of Clinical Oncology</i> , 2018, 36, 640-640.	1.6	0
34	Carcinomas of the renal medulla: A comprehensive genomic profiling (CGP) study.. <i>Journal of Clinical Oncology</i> , 2018, 36, e16586-e16586.	1.6	0
35	Phosphorylation and Ubiquitination Regulate Protein Phosphatase 5 Activity and Its Prosurvival Role in Kidney Cancer. <i>Cell Reports</i> , 2017, 21, 1883-1895.	6.4	40
36	Tumor suppressor Tsc1 is a new Hsp90 coâ€chaperone that facilitates folding of kinase and nonâ€kinase clients. <i>EMBO Journal</i> , 2017, 36, 3650-3665.	7.8	64

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37	Impact of Posttranslational Modifications on the Anticancer Activity of Hsp90 Inhibitors. <i>Advances in Cancer Research</i> , 2016, 129, 31-50.	5.0	36
38	Structural and functional basis of protein phosphatase 5 substrate specificity. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 9009-9014.	7.1	66
39	The FNIP co-chaperones decelerate the Hsp90 chaperone cycle and enhance drug binding. <i>Nature Communications</i> , 2016, 7, 12037.	12.8	56
40	Chromophobe Renal Cell Carcinoma is the Most Common Nonclear Renal Cell Carcinoma in Young Women: Results from the SEER Database. <i>Journal of Urology</i> , 2016, 195, 847-851.	0.4	14
41	Mps1 Mediated Phosphorylation of Hsp90 Confers Renal Cell Carcinoma Sensitivity and Selectivity to Hsp90 Inhibitors. <i>Cell Reports</i> , 2016, 14, 872-884.	6.4	60
42	Targeting Hsp90 in Non-Cancerous Maladies. <i>Current Topics in Medicinal Chemistry</i> , 2016, 16, 2792-2804.	2.1	12
43	The dynamic interactome of human Aha1 upon Y223 phosphorylation. <i>Data in Brief</i> , 2015, 5, 752-755.	1.0	10
44	c-Abl Mediated Tyrosine Phosphorylation of Aha1 Activates Its Co-chaperone Function in Cancer Cells. <i>Cell Reports</i> , 2015, 12, 1006-1018.	6.4	54
45	Targeting Hsp90 in urothelial carcinoma. <i>Oncotarget</i> , 2015, 6, 8454-8473.	1.8	31
46	Asymmetric Hsp90 α N Domain SUMOylation Recruits Aha1 and ATP-Competitive Inhibitors. <i>Molecular Cell</i> , 2014, 53, 317-329.	9.7	101
47	The HSP90 Inhibitor Ganetespib Synergizes with the MET Kinase Inhibitor Crizotinib in both Crizotinib-Sensitive and -Resistant MET-Driven Tumor Models. <i>Cancer Research</i> , 2013, 73, 7022-7033.	0.9	49
48	Contributions of co-chaperones and post-translational modifications towards Hsp90 drug sensitivity. <i>Future Medicinal Chemistry</i> , 2013, 5, 1059-1071.	2.3	54
49	Post-translational modification and conformational state of Heat Shock Protein 90 differentially affect binding of chemically diverse small molecule inhibitors. <i>Oncotarget</i> , 2013, 4, 1065-1074.	1.8	58
50	Combined inhibition of Wee1 and Hsp90 activates intrinsic apoptosis in cancer cells. <i>Cell Cycle</i> , 2012, 11, 3649-3655.	2.6	23
51	Charged linker sequence modulates eukaryotic heat shock protein 90 (Hsp90) chaperone activity. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 2937-2942.	7.1	107
52	Tumor-Intrinsic and Tumor-Extrinsic Factors Impacting Hsp90- Targeted Therapy. <i>Current Molecular Medicine</i> , 2012, 12, 1125-1141.	1.3	47
53	Dynamic Tyrosine Phosphorylation Modulates Cycling of the HSP90-P50CDC37-AHA1 Chaperone Machine. <i>Molecular Cell</i> , 2012, 47, 434-443.	9.7	113
54	Activity of the yeast zinc-finger transcription factor War1 is lost with alanine mutation of two putative phosphorylation sites in the activation domain. <i>Yeast</i> , 2012, 29, 39-44.	1.7	9

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55	Post-translational modifications of Hsp90 and their contributions to chaperone regulation. <i>Biochimica Et Biophysica Acta - Molecular Cell Research</i> , 2012, 1823, 648-655.	4.1	248
56	Threonine 22 Phosphorylation Attenuates Hsp90 Interaction with Cochaperones and Affects Its Chaperone Activity. <i>Molecular Cell</i> , 2011, 41, 672-681.	9.7	146
57	A systematic protocol for the characterization of Hsp90 modulators. <i>Bioorganic and Medicinal Chemistry</i> , 2011, 19, 684-692.	3.0	80
58	Detecting HSP90 Phosphorylation. <i>Methods in Molecular Biology</i> , 2011, 787, 67-74.	0.9	7
59	Casein kinase 2 phosphorylation of Hsp90 threonine 22 modulates chaperone function and drug sensitivity. <i>Oncotarget</i> , 2011, 2, 407-417.	1.8	53
60	Targeting the dynamic HSP90 complex in cancer. <i>Nature Reviews Cancer</i> , 2010, 10, 537-549.	28.4	1,306
61	Hsp90 phosphorylation, Wee1 and the cell cycle. <i>Cell Cycle</i> , 2010, 9, 2310-2316.	2.6	74
62	Swe1/Wee1-Dependent Tyrosine Phosphorylation of Hsp90 Regulates Distinct Facets of Chaperone Function. <i>Molecular Cell</i> , 2010, 37, 333-343.	9.7	165
63	Fumarate Hydratase Deficiency in Renal Cancer Induces Glycolytic Addiction and Hypoxia-Inducible Transcription Factor β Stabilization by Glucose-Dependent Generation of Reactive Oxygen Species. <i>Molecular and Cellular Biology</i> , 2009, 29, 4080-4090.	2.3	212
64	The complex dance of the molecular chaperone Hsp90. <i>Trends in Biochemical Sciences</i> , 2009, 34, 223-226.	7.5	40
65	The Hsp90/Cdc37p chaperone system is a determinant of molybdate resistance in <i>Saccharomyces cerevisiae</i> . <i>Yeast</i> , 2009, 26, 339-347.	1.7	9
66	Hsp90 charged-linker truncation reverses the functional consequences of weakened hydrophobic contacts in the N domain. <i>Nature Structural and Molecular Biology</i> , 2009, 16, 1141-1147.	8.2	78
67	Visualizing the twists and turns of a molecular chaperone. <i>Nature Structural and Molecular Biology</i> , 2009, 16, 235-236.	8.2	13
68	Novel stress responses facilitate <i>Saccharomyces cerevisiae</i> growth in the presence of the monocarboxylate preservatives. <i>Yeast</i> , 2008, 25, 169-177.	1.7	60
69	Hsp90-Dependent Activation of Protein Kinases Is Regulated by Chaperone-Targeted Dephosphorylation of Cdc37. <i>Molecular Cell</i> , 2008, 31, 886-895.	9.7	184
70	Hog1 Mitogen-Activated Protein Kinase Phosphorylation Targets the Yeast Fps1 Aquaglyceroporin for Endocytosis, Thereby Rendering Cells Resistant to Acetic Acid. <i>Molecular and Cellular Biology</i> , 2007, 27, 6446-6456.	2.3	225
71	In the Yeast Heat Shock Response, Hsf1-Directed Induction of Hsp90 Facilitates the Activation of the Slt2 (Mpk1) Mitogen-Activated Protein Kinase Required for Cell Integrity. <i>Eukaryotic Cell</i> , 2007, 6, 744-752.	3.4	49
72	Expressed as the sole Hsp90 of yeast, the β and γ isoforms of human Hsp90 differ with regard to their capacities for activation of certain client proteins, whereas only Hsp90 β generates sensitivity to the Hsp90 inhibitor radicicol. <i>FEBS Journal</i> , 2007, 274, 4453-4463.	4.7	76

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73	Hog1p mitogen-activated protein kinase determines acetic acid resistance in <i>Saccharomyces cerevisiae</i> . <i>FEMS Yeast Research</i> , 2006, 6, 1274-1280.	2.3	84
74	Expressed in the Yeast <i>Saccharomyces cerevisiae</i> , Human ERK5 Is a Client of the Hsp90 Chaperone That Complements Loss of the Slt2p (Mpk1p) Cell Integrity Stress-Activated Protein Kinase. <i>Eukaryotic Cell</i> , 2006, 5, 1914-1924.	3.4	60
75	Qri2/Nse4, a component of the essential Smc5/6 DNA repair complex. <i>Molecular Microbiology</i> , 2005, 55, 1735-1750.	2.5	43
76	Overexpressed Sod1p acts either to reduce or to increase the lifespans and stress resistance of yeast, depending on whether it is Cu ²⁺ -deficient or an active Cu,Zn-superoxide dismutase. <i>Aging Cell</i> , 2005, 4, 41-52.	6.7	43
77	Screening the yeast deletant mutant collection for hypersensitivity and hyper-resistance to sorbate, a weak organic acid food preservative. <i>Yeast</i> , 2004, 21, 927-946.	1.7	73
78	Mnsod overexpression extends the yeast chronological (G0) life span but acts independently of Sir2p histone deacetylase to shorten the replicative life span of dividing cells. <i>Free Radical Biology and Medicine</i> , 2003, 34, 1599-1606.	2.9	89
79	Weak organic acid stress inhibits aromatic amino acid uptake by yeast, causing a strong influence of amino acid auxotrophies on the phenotypes of membrane transporter mutants. <i>FEBS Journal</i> , 2003, 270, 3189-3195.	0.2	110
80	Sensitivity to Hsp90-targeting drugs can arise with mutation to the Hsp90 chaperone, cochaperones and plasma membrane ATP binding cassette transporters of yeast. <i>FEBS Journal</i> , 2003, 270, 4689-4695.	0.2	52
81	Moderately lipophilic carboxylate compounds are the selective inducers of the <i>Saccharomyces cerevisiae</i> Pdr12p ATP-binding cassette transporter. <i>Yeast</i> , 2003, 20, 575-585.	1.7	63
82	Yeast is selectively hypersensitized to heat shock protein 90 (Hsp90)-targeting drugs with heterologous expression of the human Hsp90 α , a property that can be exploited in screens for new Hsp90 chaperone inhibitors. <i>Gene</i> , 2003, 302, 165-170.	2.2	51
83	War1p, a Novel Transcription Factor Controlling Weak Acid Stress Response in Yeast. <i>Molecular and Cellular Biology</i> , 2003, 23, 1775-1785.	2.3	129
84	Activation of the ATPase Activity of Hsp90 by the Stress-Regulated Cochaperone Aha1. <i>Molecular Cell</i> , 2002, 10, 1307-1318.	9.7	487
85	The shortened replicative life span of prohibitin mutants of yeast appears to be due to defective mitochondrial segregation in old mother cells. <i>Aging Cell</i> , 2002, 1, 149-157.	6.7	69
86	The ZbYME2 gene from the food spoilage yeast <i>Zygosaccharomyces bailii</i> confers not only YME2 functions in <i>Saccharomyces cerevisiae</i> , but also the capacity for catabolism of sorbate and benzoate, two major weak organic acid preservatives. <i>Molecular Microbiology</i> , 2001, 42, 919-930.	2.5	54
87	Targeted gene deletion in <i>Zygosaccharomyces bailii</i> . <i>Yeast</i> , 2001, 18, 173-186.	1.7	47