

Tiziana Lodi

List of Publications by Year in descending order

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67
papers

1,930
citations

279798

23
h-index

289244

40
g-index

67
all docs

67
docs citations

67
times ranked

2799
citing authors

#	ARTICLE	IF	CITATIONS
1	Drug repositioning as a therapeutic strategy for neurodegenerations associated with OPA1 mutations. <i>Human Molecular Genetics</i> , 2021, 29, 3631-3645.	2.9	13
2	Efficient clofilium tosylate-mediated rescue of POLG-related disease phenotypes in zebrafish. <i>Cell Death and Disease</i> , 2021, 12, 100.	6.3	13
3	The Power of Yeast in Modelling Human Nuclear Mutations Associated with Mitochondrial Diseases. <i>Genes</i> , 2021, 12, 300.	2.4	15
4	A Yeast-Based Screening Unravels Potential Therapeutic Molecules for Mitochondrial Diseases Associated with Dominant ANT1 Mutations. <i>International Journal of Molecular Sciences</i> , 2021, 22, 4461.	4.1	10
5	A Yeast-Based Repurposing Approach for the Treatment of Mitochondrial DNA Depletion Syndromes Led to the Identification of Molecules Able to Modulate the dNTP Pool. <i>International Journal of Molecular Sciences</i> , 2021, 22, 12223.	4.1	6
6	Mechanistic insights on the mode of action of an antiproliferative thiosemicarbazone-nickel complex revealed by an integrated chemogenomic profiling study. <i>Scientific Reports</i> , 2020, 10, 10524.	3.3	17
7	Modeling of pathogenic variants of mitochondrial DNA polymerase: insight into the replication defects and implication for human disease. <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2020, 1864, 129608.	2.4	3
8	Amino and carboxy-terminal extensions of yeast mitochondrial DNA polymerase assemble both the polymerization and exonuclease active sites. <i>Mitochondrion</i> , 2019, 49, 166-177.	3.4	5
9	Sabotage at the Powerhouse? Unraveling the Molecular Target of 2-Isopropylbenzaldehyde Thiosemicarbazone, a Specific Inhibitor of Aflatoxin Biosynthesis and Sclerotia Development in <i>Aspergillus flavus</i> , Using Yeast as a Model System. <i>Molecules</i> , 2019, 24, 2971.	3.8	4
10	Yeast expression of mammalian Onzin and fungal FCR1 suggests ancestral functions of PLAC8 proteins in mitochondrial metabolism and DNA repair. <i>Scientific Reports</i> , 2019, 9, 6629.	3.3	17
11	Deciphering OPA1 mutations pathogenicity by combined analysis of human, mouse and yeast cell models. <i>Biochimica Et Biophysica Acta - Molecular Basis of Disease</i> , 2018, 1864, 3496-3514.	3.8	36
12	Dominance of yeast aac2 R96H and aac2 R252G mutations, equivalent to pathological mutations in ant1, is due to gain of function. <i>Biochemical and Biophysical Research Communications</i> , 2017, 493, 909-913.	2.1	8
13	Structural modification of cuminaldehyde thiosemicarbazone increases inhibition specificity toward aflatoxin biosynthesis and sclerotia development in <i>Aspergillus flavus</i> . <i>Applied Microbiology and Biotechnology</i> , 2017, 101, 6683-6696.	3.6	17
14	Defective mitochondrial rRNA methyltransferase MRM2 causes MELAS-like clinical syndrome. <i>Human Molecular Genetics</i> , 2017, 26, 4257-4266.	2.9	63
15	Combined use of <i>Saccharomyces cerevisiae</i> , <i>Caenorhabditis elegans</i> and patient fibroblasts leads to the identification of clofilium tosylate as a potential therapeutic chemical against POLG-related diseases. <i>Human Molecular Genetics</i> , 2016, 25, 715-727.	2.9	18
16	Recurrent De Novo Dominant Mutations in SLC25A4 Cause Severe Early-Onset Mitochondrial Disease and Loss of Mitochondrial DNA Copy Number. <i>American Journal of Human Genetics</i> , 2016, 99, 860-876.	6.2	93
17	DNA polymerase γ^3 and disease: what we have learned from yeast. <i>Frontiers in Genetics</i> , 2015, 6, 106.	2.3	23
18	In vitro evaluation of the activity of thiosemicarbazone derivatives against mycotoxigenic fungi affecting cereals. <i>International Journal of Food Microbiology</i> , 2015, 200, 104-111.	4.7	39

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19	Polymorphisms in DNA polymerase β affect the mtDNA stability and the NRTI-induced mitochondrial toxicity in <i>Saccharomyces cerevisiae</i> . <i>Mitochondrion</i> , 2015, 20, 52-63.	3.4	16
20	Biallelic Mutations of Methionyl-tRNA Synthetase Cause a Specific Type of Pulmonary Alveolar Proteinosis Prevalent on Runion Island. <i>American Journal of Human Genetics</i> , 2015, 96, 826-831.	6.2	94
21	Validation of a MGM1/OPA1 chimeric gene for functional analysis in yeast of mutations associated with dominant optic atrophy. <i>Mitochondrion</i> , 2015, 25, 38-48.	3.4	16
22	Mitochondrial thiol oxidase Erv1: both shuttle cysteine residues are required for its function with distinct roles. <i>Biochemical Journal</i> , 2014, 460, 199-210.	3.7	16
23	<i>FLO11</i> expression and lipid biosynthesis are required for air-liquid biofilm formation in a <i>Saccharomyces cerevisiae</i> flor strain. <i>FEMS Yeast Research</i> , 2012, 12, 864-866.	2.3	25
24	Overexpression of DNA Polymerase Zeta Reduces the Mitochondrial Mutability Caused by Pathological Mutations in DNA Polymerase Gamma in Yeast. <i>PLoS ONE</i> , 2012, 7, e34322.	2.5	20
25	Insights into Physiological and Genetic Mupirocin Susceptibility in Bifidobacteria. <i>Applied and Environmental Microbiology</i> , 2011, 77, 3141-3146.	3.1	37
26	Predicting the contribution of novel POLG mutations to human disease through analysis in yeast model. <i>Mitochondrion</i> , 2011, 11, 182-190.	3.4	23
27	Construction and validation of a yeast model system for studying in vivo the susceptibility to nucleoside analogues of DNA polymerase gamma allelic variants. <i>Mitochondrion</i> , 2010, 10, 183-187.	3.4	17
28	A variable neurodegenerative phenotype with polymerase γ mutation. <i>Journal of Neurology, Neurosurgery and Psychiatry</i> , 2009, 80, 1181-1182.	1.9	18
29	Oxygen is required to restore flor strain viability and lipid biosynthesis under fermentative conditions. <i>FEMS Yeast Research</i> , 2009, 9, 217-225.	2.3	21
30	Construction and characterization of centromeric, episomal and GFP-containing vectors for <i>Saccharomyces cerevisiae</i> prototrophic strains. <i>Journal of Biotechnology</i> , 2009, 143, 247-254.	3.8	10
31	The Mitochondrial Disulfide Relay System Protein GFER Is Mutated in Autosomal-Recessive Myopathy with Cataract and Combined Respiratory-Chain Deficiency. <i>American Journal of Human Genetics</i> , 2009, 84, 594-604.	6.2	121
32	Characterization of <i>KLGUT2</i> , a gene of the glycerol-3-phosphate shuttle, in <i>Kluyveromyces lactis</i> . <i>FEMS Yeast Research</i> , 2008, 8, 697-705.	2.3	16
33	Behaviour of <i>Saccharomyces cerevisiae</i> wine strains during adaptation to unfavourable conditions of fermentation on synthetic medium: Cell lipid composition, membrane integrity, viability and fermentative activity. <i>International Journal of Food Microbiology</i> , 2008, 121, 84-91.	4.7	91
34	Deletion of the Glucose-6-Phosphate Dehydrogenase Gene Kl ZWF1 Affects both Fermentative and Respiratory Metabolism in <i>Kluyveromyces lactis</i> . <i>Eukaryotic Cell</i> , 2007, 6, 19-27.	3.4	26
35	A Single Nucleotide Polymorphism in the DNA Polymerase Gamma Gene of <i>Saccharomyces cerevisiae</i> Laboratory Strains Is Responsible for Increased Mitochondrial DNA Mutability. <i>Genetics</i> , 2007, 177, 1227-1231.	2.9	25
36	Induction and characterization of morphologic mutants in a natural <i>Saccharomyces cerevisiae</i> strain. <i>Canadian Journal of Microbiology</i> , 2007, 53, 223-230.	1.7	3

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37	KNQ1, a <i>Kluyveromyces lactis</i> gene encoding a transmembrane protein, may be involved in iron homeostasis. <i>FEMS Yeast Research</i> , 2007, 7, 715-721.	2.3	6
38	Evolution of the carboxylate Jen transporters in fungi. <i>FEMS Yeast Research</i> , 2007, 7, 646-656.	2.3	22
39	Mutation D104G in ANT1 gene: Complementation study in <i>Saccharomyces cerevisiae</i> as a model system. <i>Biochemical and Biophysical Research Communications</i> , 2006, 341, 810-815.	2.1	17
40	KIADH3, a gene encoding a mitochondrial alcohol dehydrogenase, affects respiratory metabolism and cytochrome content in <i>Kluyveromyces lactis</i> . <i>FEMS Yeast Research</i> , 2006, 6, 1184-1192.	2.3	10
41	Galactose transport in <i>Kluyveromyces lactis</i> : major role of the glucose permease Hgt1. <i>FEMS Yeast Research</i> , 2006, 6, 1235-1242.	2.3	48
42	Genetic and chemical rescue of the <i>Saccharomyces cerevisiae</i> phenotype induced by mitochondrial DNA polymerase mutations associated with progressive external ophthalmoplegia in humans. <i>Human Molecular Genetics</i> , 2006, 15, 2846-2855.	2.9	80
43	Secretion of Human Serum Albumin by <i>Kluyveromyces lactis</i> Overexpressing KIPDI1 and KIERO1. <i>Applied and Environmental Microbiology</i> , 2005, 71, 4359-4363.	3.1	43
44	Lactose-induced cell death of β -galactosidase mutants in. <i>FEMS Yeast Research</i> , 2005, 5, 727-734.	2.3	17
45	Complete loss-of-function of the heart/muscle-specific adenine nucleotide translocator is associated with mitochondrial myopathy and cardiomyopathy. <i>Human Molecular Genetics</i> , 2005, 14, 3079-3088.	2.9	165
46	The Deletion of the Succinate Dehydrogenase Gene KISDH1 in <i>Kluyveromyces lactis</i> Does Not Lead to Respiratory Deficiency. <i>Eukaryotic Cell</i> , 2004, 3, 589-597.	3.4	23
47	Mutations in AAC2, equivalent to human adPEO-associated ANT1 mutations, lead to defective oxidative phosphorylation in <i>Saccharomyces cerevisiae</i> and affect mitochondrial DNA stability. <i>Human Molecular Genetics</i> , 2004, 13, 923-934.	2.9	71
48	Carboxylic acids permeases in yeast: two genes in <i>Kluyveromyces lactis</i> . <i>Gene</i> , 2004, 339, 111-119.	2.2	33
49	MIG1-dependent and MIG1-independent regulation of GAL gene expression in <i>Saccharomyces cerevisiae</i> : role of Imp2p. <i>Yeast</i> , 2003, 20, 1085-1096.	1.7	9
50	LYS2 gene and its mutation in <i>Kluyveromyces lactis</i> . <i>Yeast</i> , 2003, 20, 1171-1175.	1.7	4
51	Co-ordinate regulation of lactate metabolism genes in yeast: the role of the lactate permease gene JEN1. <i>Molecular Genetics and Genomics</i> , 2002, 266, 838-847.	2.1	39
52	Three Target Genes for the Transcriptional Activator Cat8p of <i>Kluyveromyces lactis</i> : Acetyl Coenzyme A Synthetase Genes KIACS1 and KIACS2 and Lactate Permease Gene KIJEN1. <i>Journal of Bacteriology</i> , 2001, 183, 5257-5261.	2.2	24
53	Cloning and characterization of the lactate-specific inducible gene KICYB2, encoding the cytochrome b2 of <i>Kluyveromyces lactis</i> . <i>Yeast</i> , 2000, 16, 657-665.	1.7	16
54	Cloning and characterization of the lactate-specific inducible gene KICYB2, encoding the cytochrome b2 of <i>Kluyveromyces lactis</i> . <i>Yeast</i> , 2000, 16, 657-665.	1.7	0

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55	A <i>Klaac</i> null mutant of <i>Kluyveromyces lactis</i> is complemented by a single copy of the <i>Saccharomyces cerevisiae</i> AAC1 gene. <i>Current Genetics</i> , 1999, 36, 29-36.	1.7	12
56	Regulation of the <i>Saccharomyces cerevisiae</i> DLD1 gene encoding the mitochondrial protein D-lactate ferricytochrome c oxidoreductase by HAP1 and HAP2/3/4/5. <i>Molecular Genetics and Genomics</i> , 1999, 262, 623-632.	2.4	30
57	Transcriptional regulation of the KIDL gene, encoding the mitochondrial enzyme D-lactate ferricytochrome c oxidoreductase in <i>Kluyveromyces lactis</i> : effect of <i>Klhap2</i> and <i>fog</i> mutations. <i>Current Genetics</i> , 1998, 34, 12-20.	1.7	12
58	Expression of a lipocalin in <i>Pichia pastoris</i> : secretion, purification and binding activity of a recombinant mouse major urinary protein. <i>FEBS Letters</i> , 1997, 401, 73-77.	2.8	47
59	FOG1 and FOG2 genes, required for the transcriptional activation of glucose-repressible genes of <i>Kluyveromyces lactis</i> , are homologous to GAL83 and SNF1 of <i>Saccharomyces cerevisiae</i> . <i>Current Genetics</i> , 1996, 29, 316-326.	1.7	13
60	FOG1 and FOG2 genes, required for the transcriptional activation of glucose-repressible genes of <i>Kluyveromyces lactis</i> , are homologous to GAL83 and SNF1 of <i>Saccharomyces cerevisiae</i> . <i>Current Genetics</i> , 1996, 29, 316-326.	1.7	35
61	Characterization of a promoter mutation in the CYP3 gene of <i>Saccharomyces cerevisiae</i> which cancels regulation by Cyp1p (<i>Hap1p</i>) without affecting its binding site. <i>Molecular Genetics and Genomics</i> , 1996, 253, 103-110.	2.4	2
62	Genes coding for mitochondrial proteins are more strongly biased in <i>Kluyveromyces lactis</i> than in <i>Saccharomyces cerevisiae</i> . <i>Current Genetics</i> , 1994, 26, 91-93.	1.7	6
63	Carbon catabolite repression in <i>Kluyveromyces lactis</i> : isolation and characterization of the KINLD gene encoding the mitochondrial enzyme D-lactate ferricytochrome c oxidoreductase. <i>Molecular Genetics and Genomics</i> , 1994, 244, 622-629.	2.4	39
64	Isolation of the DLD gene of <i>Saccharomyces cerevisiae</i> encoding the mitochondrial enzyme D-lactate ferricytochrome c oxidoreductase. <i>Molecular Genetics and Genomics</i> , 1993, 238, 315-324.	2.4	70
65	IMP2, a nuclear gene controlling the mitochondrial dependence of galactose, maltose and raffinose utilization in <i>Saccharomyces cerevisiae</i> . <i>Yeast</i> , 1992, 8, 83-93.	1.7	26
66	Antimycin A- and hydroxamate-insensitive respiration in yeasts. <i>Antonie Van Leeuwenhoek</i> , 1985, 51, 57-64.	1.7	11
67	Respiratory pathways in <i>Hansenula saturnus</i> . <i>Antonie Van Leeuwenhoek</i> , 1983, 49, 537-549.	1.7	5