

# Yvetta Gbelska

## List of Publications by Year in descending order

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26  
papers

255  
citations

1040056

9  
h-index

940533

16  
g-index

27  
all docs

27  
docs citations

27  
times ranked

337  
citing authors

#	ARTICLE	IF	CITATIONS
1	The UPC2 gene in <i>Kluyveromyces lactis</i> stress adaptation. <i>Folia Microbiologica</i> , 2022, , 1.	2.3	1
2	UPC2 gene deletion modifies sterol homeostasis and susceptibility to metabolic inhibitors in <i>Kluyveromyces lactis</i> . <i>Yeast</i> , 2020, 37, 647-657.	1.7	3
3	The Absence of PDR16 Gene Restricts the Overexpression of CaSNQ2 Gene in the Presence of Fluconazole in <i>Candida albicans</i> . <i>Mycopathologia</i> , 2020, 185, 455-465.	3.1	9
4	Stb5p is involved in <i>Kluyveromyces lactis</i> response to 4-nitroquinoline-N-oxide stress. <i>Folia Microbiologica</i> , 2019, 64, 579-586.	2.3	1
5	Erg6 gene is essential for stress adaptation in <i>Kluyveromyces lactis</i> . <i>FEMS Microbiology Letters</i> , 2018, 365, .	1.8	7
6	Differences in the arrangement of the Pdr5p multidrug transporter binding pocket of <i>Saccharomyces cerevisiae</i> and <i>Kluyveromyces lactis</i> . <i>FEMS Yeast Research</i> , 2017, 17, .	2.3	1
7	Measurement of Energy-dependent Rhodamine 6G Efflux in Yeast Species. <i>Bio-protocol</i> , 2017, 7, e2428.	0.4	7
8	Sterol Analysis in <i>Kluyveromyces lactis</i> . <i>Bio-protocol</i> , 2017, 7, e2527.	0.4	0
9	<i>ERG6</i> gene deletion modifies <i>Kluyveromyces lactis</i> susceptibility to various growth inhibitors. <i>Yeast</i> , 2016, 33, 621-632.	1.7	23
10	Insight into the <i>Kluyveromyces lactis</i> Pdr1p regulon. <i>Canadian Journal of Microbiology</i> , 2016, 62, 918-931.	1.7	0
11	The major facilitator superfamily transporter Knq1p modulates boron homeostasis in <i>Kluyveromyces lactis</i> . <i>Folia Microbiologica</i> , 2016, 61, 101-107.	2.3	0
12	Deletion of the <i>PDR16</i> gene influences the plasma membrane properties of the yeast <i>Kluyveromyces lactis</i> . <i>Canadian Journal of Microbiology</i> , 2015, 61, 273-279.	1.7	7
13	Stress response and expression of fluconazole resistance associated genes in the pathogenic yeast <i>Candida glabrata</i> deleted in the <i>CgPDR16</i> gene. <i>Microbiological Research</i> , 2015, 174, 17-23.	5.3	10
14	Isolation and functional analysis of the <i>KIPDR16</i> gene. <i>FEMS Yeast Research</i> , 2014, 14, 337-345.	2.3	10
15	Gain of function mutation in the <i>KIPDR1</i> gene encoding multidrug resistance regulator in <i>Kluyveromyces lactis</i> . <i>Yeast</i> , 2013, 30, 71-80.	1.7	1
16	Mutation of the <i>CgPDR16</i> gene attenuates azole tolerance and biofilm production in pathogenic <i>Candida glabrata</i> . <i>Yeast</i> , 2013, 30, 403-414.	1.7	15
17	Cytosolic proteome of <i>Kluyveromyces lactis</i> affected by the multidrug resistance regulating transcription factor KIPdr1p. <i>Journal of Proteomics</i> , 2012, 75, 5316-5326.	2.4	4
18	Interplay among regulators of multidrug resistance in <i>Kluyveromyces lactis</i> . <i>General Physiology and Biophysics</i> , 2011, 30, 77-81.	0.9	1

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19	Functional analysis of the <i>Kluyveromyces lactis</i> PDR1 gene. FEMS Yeast Research, 2009, 9, 321-327.	2.3	10
20	Evolution of gene families: the multidrug resistance transporter genes in five related yeast species. FEMS Yeast Research, 2006, 6, 345-355.	2.3	85
21	-mediated expression in. FEMS Yeast Research, 2005, 5, 323-329.	2.3	10
22	KNQ1, a <i>Kluyveromyces lactis</i> gene encoding a drug efflux permease. Current Genetics, 2004, 45, 1-8.	1.7	15
23	Identification and functional analysis of a <i>Kluyveromyces lactis</i> homologue of the SPT4 gene of <i>Saccharomyces cerevisiae</i> . Current Genetics, 1998, 34, 375-378.	1.7	2
24	Cloning and characterization of KICOX18, a gene required for activity of cytochrome oxidase in <i>Kluyveromyces lactis</i> . Current Genetics, 1997, 32, 267-272.	1.7	18
25	Isolation and molecular analysis of the gene for cytochrome c 1 from <i>Kluyveromyces lactis</i> . Current Genetics, 1996, 30, 145-150.	1.7	13
26	The properties of the multicopy suppressor of the <i>ogdl</i> mutation in yeast. Journal of Basic Microbiology, 1995, 35, 229-232.	3.3	2