

Amparo Querol

List of Publications by Citations

Source: <https://exaly.com/author-pdf/5496252/amparo-querol-publications-by-citations.pdf>

Version: 2024-04-28

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

224
papers

9,568
citations

57
h-index

87
g-index

233
ext. papers

11,343
ext. citations

4.6
avg, IF

6.17
L-index

#	Paper	IF	Citations
224	Identification of yeasts by RFLP analysis of the 5.8S rRNA gene and the two ribosomal internal transcribed spacers. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 1999 , 49 Pt 1, 329-37	2.2	633
223	Molecular monitoring of wine fermentations conducted by active dry yeast strains. <i>Applied and Environmental Microbiology</i> , 1992 , 58, 2948-53	4.8	371
222	A Comparative Study of Different Methods of Yeast Strain Characterization. <i>Systematic and Applied Microbiology</i> , 1992 , 15, 439-446	4.2	286
221	Rapid identification of wine yeast species based on RFLP analysis of the ribosomal internal transcribed spacer (ITS) region. <i>Archives of Microbiology</i> , 1998 , 169, 387-92	3	238
220	Natural hybrids from <i>Saccharomyces cerevisiae</i> , <i>Saccharomyces bayanus</i> and <i>Saccharomyces kudriavzevii</i> in wine fermentations. <i>FEMS Yeast Research</i> , 2006 , 6, 1221-34	3.1	188
219	Molecular characterization of a chromosomal rearrangement involved in the adaptive evolution of yeast strains. <i>Genome Research</i> , 2002 , 12, 1533-9	9.7	186
218	Role of yeasts in table olive production. <i>International Journal of Food Microbiology</i> , 2008 , 128, 189-96	5.8	185
217	Temperature adaptation markedly determines evolution within the genus <i>Saccharomyces</i> . <i>Applied and Environmental Microbiology</i> , 2011 , 77, 2292-302	4.8	155
216	Fermentative stress adaptation of hybrids within the <i>Saccharomyces sensu stricto</i> complex. <i>International Journal of Food Microbiology</i> , 2008 , 122, 188-95	5.8	152
215	Molecular characterization of new natural hybrids of <i>Saccharomyces cerevisiae</i> and <i>S. kudriavzevii</i> in brewing. <i>Applied and Environmental Microbiology</i> , 2008 , 74, 2314-20	4.8	132
214	Adaptive evolution of wine yeast. <i>International Journal of Food Microbiology</i> , 2003 , 86, 3-10	5.8	122
213	Physiological characterization of spoilage strains of <i>Zygosaccharomyces bailii</i> and <i>Zygosaccharomyces rouxii</i> isolated from high sugar environments. <i>International Journal of Food Microbiology</i> , 2007 , 114, 234-42	5.8	120
212	Effects of temperature, pH and sugar concentration on the growth parameters of <i>Saccharomyces cerevisiae</i> , <i>S. kudriavzevii</i> and their interspecific hybrid. <i>International Journal of Food Microbiology</i> , 2009 , 131, 120-7	5.8	118
211	Identification of yeasts isolated from wine-related environments and capable of producing 4-ethylphenol. <i>Food Microbiology</i> , 2003 , 20, 567-574	6	111
210	Scientific Opinion on the update of the list of QPS-recommended biological agents intentionally added to food or feed as notified to EFSA (2017-2019). <i>EFSA Journal</i> , 2020 , 18, e05966	2.3	106
209	Yeasts in table olive processing: desirable or spoilage microorganisms?. <i>International Journal of Food Microbiology</i> , 2012 , 160, 42-9	5.8	102
208	The complex and dynamic genomes of industrial yeasts. <i>FEMS Microbiology Letters</i> , 2009 , 293, 1-10	2.9	100

207	Qualified presumption of safety (QPS): a generic risk assessment approach for biological agents notified to the European Food Safety Authority (EFSA). <i>Trends in Food Science and Technology</i> , 2010 , 21, 425-435	15.3	99
206	Yeast population dynamics during the fermentation and biological aging of sherry wines. <i>Applied and Environmental Microbiology</i> , 2001 , 67, 2056-61	4.8	99
205	RFLP analysis of the ribosomal internal transcribed spacers and the 5.8S rRNA gene region of the genus <i>Saccharomyces</i> : a fast method for species identification and the differentiation of flor yeasts. <i>Antonie Van Leeuwenhoek</i> , 2000 , 78, 87-97	2.1	98
204	The role of indigenous yeasts in traditional Irish cider fermentations. <i>Journal of Applied Microbiology</i> , 2004 , 97, 647-55	4.7	97
203	The prevalence and control of spoilage yeasts in foods and beverages. <i>Trends in Food Science and Technology</i> , 1999 , 10, 356-365	15.3	97
202	Enological characterization of natural hybrids from <i>Saccharomyces cerevisiae</i> and <i>S. kudriavzevii</i> . <i>International Journal of Food Microbiology</i> , 2007 , 116, 11-8	5.8	96
201	Genetically different wine yeasts isolated from Austrian vine-growing regions influence wine aroma differently and contain putative hybrids between <i>Saccharomyces cerevisiae</i> and <i>Saccharomyces kudriavzevii</i> . <i>FEMS Yeast Research</i> , 2007 , 7, 953-65	3.1	94
200	Dry Yeast Strain For Use in Fermentation of Alicante Wines: Selection and DNA Patterns. <i>Journal of Food Science</i> , 1992 , 57, 183-185	3.4	88
199	Study of the authenticity of commercial wine yeast strains by molecular techniques. <i>International Journal of Food Microbiology</i> , 2001 , 70, 1-10	5.8	87
198	Diversity of <i>Saccharomyces</i> strains in wine fermentations: analysis for two consecutive years. <i>Letters in Applied Microbiology</i> , 1998 , 26, 452-5	2.9	86
197	Scientific Opinion on the update of the list of QPS-recommended biological agents intentionally added to food or feed as notified to EFSA. <i>EFSA Journal</i> , 2017 , 15, e04664	2.3	83
196	Mitotic recombination and genetic changes in <i>Saccharomyces cerevisiae</i> during wine fermentation. <i>Applied and Environmental Microbiology</i> , 2000 , 66, 2057-61	4.8	83
195	Fungemia with <i>Saccharomyces cerevisiae</i> in two newborns, only one of whom had been treated with ultra-levura. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2000 , 19, 468-70	5.3	82
194	Population dynamics of natural <i>Saccharomyces</i> strains during wine fermentation. <i>International Journal of Food Microbiology</i> , 1994 , 21, 315-23	5.8	82
193	Differences in the glucose and fructose consumption profiles in diverse <i>Saccharomyces</i> wine species and their hybrids during grape juice fermentation. <i>International Journal of Food Microbiology</i> , 2009 , 134, 237-43	5.8	81
192	Rapid identification and enumeration of <i>Saccharomyces cerevisiae</i> cells in wine by real-time PCR. <i>Applied and Environmental Microbiology</i> , 2005 , 71, 6823-30	4.8	81
191	Use of molecular methods for the identification of yeast associated with table olives. <i>Food Microbiology</i> , 2006 , 23, 791-6	6	80
190	Analysis of the stress resistance of commercial wine yeast strains. <i>Archives of Microbiology</i> , 2001 , 175, 450-7	3	80

189	Saccharomyces cerevisiae wine yeast populations in a cold region in Argentinean Patagonia. A study at different fermentation scales. <i>Journal of Applied Microbiology</i> , 2002 , 93, 608-15	4.7	79
188	Chimeric genomes of natural hybrids of Saccharomyces cerevisiae and Saccharomyces kudriavzevii. <i>Applied and Environmental Microbiology</i> , 2009 , 75, 2534-44	4.8	76
187	Rapid characterization of four species of the Saccharomyces sensu stricto complex according to mitochondrial DNA patterns. <i>International Journal of Systematic Bacteriology</i> , 1994 , 44, 708-14		76
186	Identification of species of the genus Candida by analysis of the 5.8S rRNA gene and the two ribosomal internal transcribed spacers. <i>Antonie Van Leeuwenhoek</i> , 2004 , 85, 175-85	2.1	70
185	Selection and molecular characterization of wine yeasts isolated from the El Penedès area (Spain). <i>Food Microbiology</i> , 2000 , 17, 553-562	6	70
184	Screening of non-Saccharomyces wine yeasts for the production of beta-D-xylosidase activity. <i>International Journal of Food Microbiology</i> , 1999 , 46, 105-12	5.8	70
183	Exploring the yeast biodiversity of green table olive industrial fermentations for technological applications. <i>International Journal of Food Microbiology</i> , 2011 , 147, 89-96	5.8	68
182	Phylogeny of the genus Kluyveromyces inferred from the mitochondrial cytochrome-c oxidase II gene. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2000 , 50 Pt 1, 405-416	2.2	68
181	Mitochondrial import of subunit Va of cytochrome c oxidase characterized with yeast mutants. <i>Journal of Biological Chemistry</i> , 1995 , 270, 3788-95	5.4	67
180	A simplified procedure to analyse mitochondrial DNA from industrial yeasts. <i>International Journal of Food Microbiology</i> , 2001 , 68, 75-81	5.8	64
179	The application of molecular techniques in wine microbiology. <i>Trends in Food Science and Technology</i> , 1996 , 7, 73-78	15.3	64
178	Production of aroma compounds by cryotolerant Saccharomyces species and hybrids at low and moderate fermentation temperatures. <i>Journal of Applied Microbiology</i> , 2013 , 114, 1405-14	4.7	63
177	Comparative genomics among Saccharomyces cerevisiae Saccharomyces kudriavzevii natural hybrid strains isolated from wine and beer reveals different origins. <i>BMC Genomics</i> , 2012 , 13, 407	4.5	63
176	Molecular characterization of Colletotrichum strains derived from strawberry. <i>Mycological Research</i> , 1999 , 103, 385-394		63
175	Metabolomic comparison of Saccharomyces cerevisiae and the cryotolerant species S. bayanus var. uvarum and S. kudriavzevii during wine fermentation at low temperature. <i>PLoS ONE</i> , 2013 , 8, e60135	3.7	62
174	Characterization of Wine Yeast Strains of the Saccharomyces Genus on the Basis of Molecular Markers: Relationships Between Genetic Distance and Geographic or Ecological Origin. <i>Systematic and Applied Microbiology</i> , 1996 , 19, 122-132	4.2	62
173	Molecular profiling of yeasts isolated during spontaneous fermentations of Austrian wines. <i>FEMS Yeast Research</i> , 2008 , 8, 1063-75	3.1	60
172	Modulation of the glycerol and ethanol syntheses in the yeast Saccharomyces kudriavzevii differs from that exhibited by Saccharomyces cerevisiae and their hybrid. <i>Food Microbiology</i> , 2010 , 27, 628-37	6	59

171	Molecular typing of the yeast species <i>Dekkera bruxellensis</i> and <i>Pichia guilliermondii</i> recovered from wine related sources. <i>International Journal of Food Microbiology</i> , 2006 , 106, 79-84	5.8	58
170	Correlation between acetaldehyde and ethanol resistance and expression of HSP genes in yeast strains isolated during the biological aging of sherry wines. <i>Archives of Microbiology</i> , 2002 , 177, 304-12	3	57
169	Phylogenetic Relationships Among <i>Colletotrichum</i> Pathogens of Strawberry and Design of PCR Primers for their Identification. <i>Journal of Phytopathology</i> , 2003 , 151, 135-143	1.8	57
168	Mycotoxins and mycotoxigenic moulds in nuts and sunflower seeds for human consumption. <i>Mycopathologia</i> , 1991 , 115, 121-7	2.9	57
167	Multiple Approaches Detect the Presence of Fungi in Human Breastmilk Samples from Healthy Mothers. <i>Scientific Reports</i> , 2017 , 7, 13016	4.9	55
166	Natural hybrids of <i>S. cerevisiae</i> x <i>S. kudriavzevii</i> share alleles with European wild populations of <i>Saccharomyces kudriavzevii</i> . <i>FEMS Yeast Research</i> , 2010 , 10, 412-21	3.1	55
165	Sour rot-damaged grapes are sources of wine spoilage yeasts. <i>FEMS Yeast Research</i> , 2008 , 8, 1008-17	3.1	55
164	<i>Saccharomyces kudriavzevii</i> and <i>Saccharomyces uvarum</i> differ from <i>Saccharomyces cerevisiae</i> during the production of aroma-active higher alcohols and acetate esters using their amino acidic precursors. <i>International Journal of Food Microbiology</i> , 2015 , 205, 41-6	5.8	53
163	Susceptibility and resistance to ethanol in <i>Saccharomyces</i> strains isolated from wild and fermentative environments. <i>Yeast</i> , 2010 , 27, 1005-15	3.4	52
162	Aroma improving in microvinification processes by the use of a recombinant wine yeast strain expressing the <i>Aspergillus nidulans</i> xlnA gene. <i>International Journal of Food Microbiology</i> , 1999 , 47, 171-8	5.8	52
161	Update of the list of QPS-recommended biological agents intentionally added to food or feed as notified to EFSA 12: suitability of taxonomic units notified to EFSA until March 2020. <i>EFSA Journal</i> , 2020 , 18, e06174	2.3	51
160	Food and probiotic strains from the <i>Saccharomyces cerevisiae</i> species as a possible origin of human systemic infections. <i>International Journal of Food Microbiology</i> , 2006 , 110, 286-90	5.8	50
159	A comparison of clinical and food <i>Saccharomyces cerevisiae</i> isolates on the basis of potential virulence factors. <i>Antonie Van Leeuwenhoek</i> , 2006 , 90, 221-31	2.1	50
158	Quantifying the individual effects of ethanol and temperature on the fitness advantage of <i>Saccharomyces cerevisiae</i> . <i>Food Microbiology</i> , 2011 , 28, 1155-61	6	49
157	Microbiological and Enological Parameters during Fermentation of Musts from Poor and Normal Grape-Harvests in the Region of Alicante (Spain). <i>Journal of Food Science</i> , 1990 , 55, 1603-1606	3.4	49
156	On the origins and industrial applications of <i>Saccharomyces cerevisiae</i> [<i>Saccharomyces kudriavzevii</i> hybrids. <i>Yeast</i> , 2018 , 35, 51-69	3.4	46
155	Evaluation of different genetic procedures for the generation of artificial hybrids in <i>Saccharomyces</i> genus for winemaking. <i>International Journal of Food Microbiology</i> , 2012 , 156, 102-11	5.8	46
154	Effect of Temperature on the Prevalence of Non Species against a Wine Strain in Wine Fermentation: Competition, Physiological Fitness, and Influence in Final Wine Composition. <i>Frontiers in Microbiology</i> , 2017 , 8, 150	5.7	45

153	Enhanced enzymatic activity of glycerol-3-phosphate dehydrogenase from the cryophilic <i>Saccharomyces kudriavzevii</i> . <i>PLoS ONE</i> , 2014 , 9, e87290	3.7	45
152	Genetic and phenotypic diversity of autochthonous cider yeasts in a cellar from Asturias. <i>Food Microbiology</i> , 2010 , 27, 503-8	6	44
151	Analysis of the genetic variability in the species of the <i>Saccharomyces sensu stricto</i> complex. <i>Yeast</i> , 2003 , 20, 1213-26	3.4	44
150	Molecular monitoring of spoilage yeasts during the production of candied fruit nougats to determine food contamination sources. <i>International Journal of Food Microbiology</i> , 2005 , 101, 293-302	5.8	44
149	Lipid composition of wine strains of <i>Saccharomyces kudriavzevii</i> and <i>Saccharomyces cerevisiae</i> grown at low temperature. <i>International Journal of Food Microbiology</i> , 2012 , 155, 191-8	5.8	43
148	Opportunistic Strains of <i>Saccharomyces cerevisiae</i> : A Potential Risk Sold in Food Products. <i>Frontiers in Microbiology</i> , 2015 , 6, 1522	5.7	43
147	On the complexity of the <i>Saccharomyces bayanus</i> taxon: hybridization and potential hybrid speciation. <i>PLoS ONE</i> , 2014 , 9, e93729	3.7	42
146	Update of the list of QPS-recommended biological agents intentionally added to food or feed as notified to EFSA 7: suitability of taxonomic units notified to EFSA until September 2017. <i>EFSA Journal</i> , 2018 , 16, e05131	2.3	41
145	Patagonian wines: the selection of an indigenous yeast starter. <i>Journal of Industrial Microbiology and Biotechnology</i> , 2007 , 34, 539-46	4.2	41
144	Identification of species in the genus <i>Pichia</i> by restriction of the internal transcribed spacers (ITS1 and ITS2) and the 5.8S ribosomal DNA gene. <i>Antonie Van Leeuwenhoek</i> , 2006 , 90, 171-81	2.1	41
143	A new PCR-based method for monitoring inoculated wine fermentations. <i>International Journal of Food Microbiology</i> , 2003 , 81, 63-71	5.8	40
142	The molecular characterization of new types of <i>Saccharomyces cerevisiae</i> β. <i>kudriavzevii</i> hybrid yeasts unveils a high genetic diversity. <i>Yeast</i> , 2012 , 29, 81-91	3.4	38
141	Inter- and intraspecific chromosome pattern variation in the yeast genus <i>Kluyveromyces</i> . <i>Yeast</i> , 1998 , 14, 1341-54	3.4	38
140	Alternative yeasts for winemaking: <i>Saccharomyces non-cerevisiae</i> and its hybrids. <i>Critical Reviews in Food Science and Nutrition</i> , 2018 , 58, 1780-1790	11.5	37
139	Monoterpene alcohols release and bioconversion by <i>Saccharomyces</i> species and hybrids. <i>International Journal of Food Microbiology</i> , 2011 , 145, 92-7	5.8	36
138	Molecular identification of yeasts associated with traditional Egyptian dairy products. <i>Journal of Food Science</i> , 2009 , 74, M341-6	3.4	36
137	A rapid and simple method for the preparation of yeast mitochondrial DNA. <i>Nucleic Acids Research</i> , 1990 , 18, 1657	20.1	36
136	Update of the list of QPS-recommended biological agents intentionally added to food or feed as notified to EFSA 5: suitability of taxonomic units notified to EFSA until September 2016. <i>EFSA Journal</i> , 2017 , 15, e04663	2.3	33

135	Transcriptomics of cryophilic <i>Saccharomyces kudriavzevii</i> reveals the key role of gene translation efficiency in cold stress adaptations. <i>BMC Genomics</i> , 2014 , 15, 432	4.5	33
134	Patagonian wines: implantation of an indigenous strain of <i>Saccharomyces cerevisiae</i> in fermentations conducted in traditional and modern cellars. <i>Journal of Industrial Microbiology and Biotechnology</i> , 2007 , 34, 139-49	4.2	33
133	Molecular identification and characterization of wine yeasts isolated from Tenerife (Canary Island, Spain). <i>Journal of Applied Microbiology</i> , 2007 , 102, 1018-25	4.7	33
132	Authentication and identification of <i>Saccharomyces cerevisiae</i> 'flor' yeast races involved in sherry ageing. <i>Antonie Van Leeuwenhoek</i> , 2004 , 85, 151-8	2.1	33
131	Dynamics of the yeast flora in artisanal country style and industrial dry cured sausage (yeast in fermented sausage). <i>Food Control</i> , 2013 , 29, 143-148	6.2	32
130	Ethanol Cellular Defense Induce Unfolded Protein Response in Yeast. <i>Frontiers in Microbiology</i> , 2016 , 7, 189	5.7	32
129	Nitrogen sources preferences of non- <i>Saccharomyces</i> yeasts to sustain growth and fermentation under winemaking conditions. <i>Food Microbiology</i> , 2020 , 85, 103287	6	32
128	Ecological interactions among <i>Saccharomyces cerevisiae</i> strains: insight into the dominance phenomenon. <i>Scientific Reports</i> , 2017 , 7, 43603	4.9	31
127	Molecular characterization of clinical <i>Saccharomyces cerevisiae</i> isolates and their association with non-clinical strains. <i>Systematic and Applied Microbiology</i> , 2004 , 27, 427-35	4.2	31
126	The qualified presumption of safety assessment and its role in EFSA risk evaluations: 15 years past. <i>FEMS Microbiology Letters</i> , 2019 , 366,	2.9	31
125	Potential benefits of the application of yeast starters in table olive processing. <i>Frontiers in Microbiology</i> , 2012 , 3,	5.7	30
124	Mitochondrial introgression suggests extensive ancestral hybridization events among <i>Saccharomyces</i> species. <i>Molecular Phylogenetics and Evolution</i> , 2017 , 108, 49-60	4.1	29
123	Effect of aromatic precursor addition to wine fermentations carried out with different <i>Saccharomyces</i> species and their hybrids. <i>International Journal of Food Microbiology</i> , 2011 , 147, 33-44	5.8	29
122	Application of a substrate inhibition model to estimate the effect of fructose concentration on the growth of diverse <i>Saccharomyces cerevisiae</i> strains. <i>Journal of Industrial Microbiology and Biotechnology</i> , 2009 , 36, 663-9	4.2	29
121	Phylogenetic reconstruction of the yeast genus <i>Kluyveromyces</i> : restriction map analysis of the 5.8S rRNA gene and the two ribosomal internal transcribed spacers. <i>Systematic and Applied Microbiology</i> , 1998 , 21, 266-73	4.2	29
120	Update of the list of QPS-recommended biological agents intentionally added to food or feed as notified to EFSA 8: suitability of taxonomic units notified to EFSA until March 2018. <i>EFSA Journal</i> , 2018 , 16, e05315	2.3	29
119	An analysis of inter- and intraspecific genetic variabilities in the <i>Kluyveromyces marxianus</i> group of yeast species for the reconsideration of the <i>K. lactis</i> taxon. <i>Yeast</i> , 2002 , 19, 257-68	3.4	28
118	In vivo virulence of commercial <i>Saccharomyces cerevisiae</i> strains with pathogenicity-associated phenotypical traits. <i>International Journal of Food Microbiology</i> , 2011 , 144, 393-9	5.8	27

117	Sequence-based identification of species belonging to the genus <i>Debaryomyces</i> . <i>FEMS Yeast Research</i> , 2005 , 5, 1157-65	3.1	27
116	Dominance of wine <i>Saccharomyces cerevisiae</i> strains over <i>S. kudriavzevii</i> in industrial fermentation competitions is related to an acceleration of nutrient uptake and utilization. <i>Environmental Microbiology</i> , 2019 , 21, 1627-1644	5.2	26
115	Update of the list of QPS-recommended biological agents intentionally added to food or feed as notified to EFSA 9: suitability of taxonomic units notified to EFSA until September 2018. <i>EFSA Journal</i> , 2019 , 17, e05555	2.3	26
114	Characterisation of Four Species of the Genus <i>Kluyveromyces</i> by Mitochondrial DNA Restriction Analysis. <i>Systematic and Applied Microbiology</i> , 1997 , 20, 397-408	4.2	26
113	Stabilization process in <i>Saccharomyces</i> intra and interspecific hybrids in fermentative conditions. <i>International Microbiology</i> , 2014 , 17, 213-24	3	26
112	Update of the list of QPS-recommended biological agents intentionally added to food or feed as notified to EFSA 10: Suitability of taxonomic units notified to EFSA until March 2019. <i>EFSA Journal</i> , 2019 , 17, e05753	2.3	25
111	Probabilistic model for the spoilage wine yeast <i>Dekkera bruxellensis</i> as a function of pH, ethanol and free SO ₂ using time as a dummy variable. <i>International Journal of Food Microbiology</i> , 2014 , 170, 83-90	5.8	25
110	Exclusion of <i>Saccharomyces kudriavzevii</i> from a wine model system mediated by <i>Saccharomyces cerevisiae</i> . <i>Yeast</i> , 2011 , 28, 423-35	3.4	25
109	Molecular characterisation of <i>Hanseniaspora</i> species. <i>Antonie Van Leeuwenhoek</i> , 2001 , 80, 85-92	2.1	25
108	Genetic improvement of non-GMO wine yeasts: Strategies, advantages and safety. <i>Trends in Food Science and Technology</i> , 2015 , 45, 1-11	15.3	24
107	Differences in Enzymatic Properties of the <i>Saccharomyces kudriavzevii</i> and <i>Saccharomyces uvarum</i> Alcohol Acetyltransferases and Their Impact on Aroma-Active Compounds Production. <i>Frontiers in Microbiology</i> , 2016 , 7, 897	5.7	24
106	New Trends in the Uses of Yeasts in Oenology. <i>Advances in Food and Nutrition Research</i> , 2018 , 85, 177-210		23
105	Combined use of killer biotype and mtDNA-RFLP patterns in a Patagonian wine <i>Saccharomyces cerevisiae</i> diversity study. <i>Antonie Van Leeuwenhoek</i> , 2006 , 89, 147-56	2.1	23
104	Pathogenic potential of <i>Saccharomyces</i> strains isolated from dietary supplements. <i>PLoS ONE</i> , 2014 , 9, e98094	3.7	23
103	Statement on the update of the list of QPS-recommended biological agents intentionally added to food or feed as notified to EFSA 1: Suitability of taxonomic units notified to EFSA until October 2014. <i>EFSA Journal</i> , 2014 , 12, 3938	2.3	22
102	Molecular analysis of the genes involved in aroma synthesis in the species <i>S. cerevisiae</i> , <i>S. kudriavzevii</i> and <i>S. bayanus</i> var. <i>uvarum</i> in winemaking conditions. <i>PLoS ONE</i> , 2014 , 9, e97626	3.7	21
101	A comparative study of the wine fermentation performance of <i>Saccharomyces paradoxus</i> under different nitrogen concentrations and glucose/fructose ratios. <i>Journal of Applied Microbiology</i> , 2010 , 108, 73-80	4.7	21
100	PCR-RFLP analysis of the IGS region of rDNA: a useful tool for the practical discrimination between species of the genus <i>Debaryomyces</i> . <i>Antonie Van Leeuwenhoek</i> , 2006 , 90, 211-9	2.1	21

99	Yeast microflora isolated from brazilian cassava roots: taxonomical classification based on molecular identification. <i>Current Microbiology</i> , 2010 , 60, 287-93	2.4	20
98	A molecular genetic study of natural strains of <i>Saccharomyces</i> isolated from Asturian cider fermentations. <i>Journal of Applied Microbiology</i> , 2007 , 103, 778-86	4.7	20
97	Spoilage yeasts from Patagonian cellars: characterization and potential biocontrol based on killer interactions. <i>World Journal of Microbiology and Biotechnology</i> , 2008 , 24, 945-953	4.4	20
96	Update of the list of QPS-recommended biological agents intentionally added to food or feed as notified to EFSA 11: suitability of taxonomic units notified to EFSA until September 2019. <i>EFSA Journal</i> , 2020 , 18, e05965	2.3	20
95	Improving the Cryotolerance of Wine Yeast by Interspecific Hybridization in the Genus. <i>Frontiers in Microbiology</i> , 2018 , 9, 3232	5.7	19
94	Molecular and enological characterization of a natural <i>Saccharomyces uvarum</i> and <i>Saccharomyces cerevisiae</i> hybrid. <i>International Journal of Food Microbiology</i> , 2015 , 204, 101-10	5.8	19
93	Genome-wide gene expression of a natural hybrid between <i>Saccharomyces cerevisiae</i> and <i>S. kudriavzevii</i> under enological conditions. <i>International Journal of Food Microbiology</i> , 2012 , 157, 340-5	5.8	19
92	Alternative Glycerol Balance Strategies among <i>Saccharomyces</i> Species in Response to Winemaking Stress. <i>Frontiers in Microbiology</i> , 2016 , 7, 435	5.7	19
91	A time course metabolism comparison among <i>Saccharomyces cerevisiae</i> , <i>S. uvarum</i> and <i>S. kudriavzevii</i> species in wine fermentation. <i>Food Microbiology</i> , 2020 , 90, 103484	6	18
90	Genomic stability of <i>Saccharomyces cerevisiae</i> baker's yeasts. <i>Systematic and Applied Microbiology</i> , 1999 , 22, 329-40	4.2	18
89	RAPD Analysis of <i>Colletotrichum</i> Species Isolated from Strawberry and the Design of Specific Primers for the Identification of <i>C. fragariae</i> . <i>Journal of Phytopathology</i> , 2002 , 150, 680-686	1.8	17
88	Identification of <i>Colletotrichum</i> species responsible for anthracnose of strawberry based on the internal transcribed spacers of the ribosomal region. <i>FEMS Microbiology Letters</i> , 2000 , 189, 97-101	2.9	17
87	Membrane fluidification by ethanol stress activates unfolded protein response in yeasts. <i>Microbial Biotechnology</i> , 2018 , 11, 465-475	6.3	16
86	Update of the list of QPS-recommended biological agents intentionally added to food or feed as notified to EFSA 6: suitability of taxonomic units notified to EFSA until March 2017. <i>EFSA Journal</i> , 2017 , 15, e04884	2.3	16
85	The Use of Mixed Populations of and to Reduce Ethanol Content in Wine: Limited Aeration, Inoculum Proportions, and Sequential Inoculation. <i>Frontiers in Microbiology</i> , 2017 , 8, 2087	5.7	16
84	Clinical <i>Saccharomyces cerevisiae</i> isolates cannot cross the epithelial barrier in vitro. <i>International Journal of Food Microbiology</i> , 2012 , 157, 59-64	5.8	16
83	Comparative genomic analysis of <i>Saccharomyces cerevisiae</i> yeasts isolated from fermentations of traditional beverages unveils different adaptive strategies. <i>International Journal of Food Microbiology</i> , 2014 , 171, 129-35	5.8	15
82	Identification of <i>Colletotrichum</i> species responsible for anthracnose of strawberry based on the internal transcribed spacers of the ribosomal region.. <i>FEMS Microbiology Letters</i> , 2000 , 189, 97-101	2.9	15

81	Saccharomyces cerevisiae & Saccharomyces uvarum hybrids generated under different conditions share similar winemaking features. <i>Yeast</i> , 2018 , 35, 157-171	3.4	15
80	iTRAQ-based proteome profiling of Saccharomyces cerevisiae and cryotolerant species Saccharomyces uvarum and Saccharomyces kudriavzevii during low-temperature wine fermentation. <i>Journal of Proteomics</i> , 2016 , 146, 70-9	3.9	14
79	Physiological and molecular characterisation of Saccharomyces cerevisiae cachañ strains isolated from different geographic regions in Brazil. <i>World Journal of Microbiology and Biotechnology</i> , 2010 , 26, 579-587	4.4	14
78	Molecular characterisation of the species of the genus Zygosaccharomyces. <i>Systematic and Applied Microbiology</i> , 2003 , 26, 404-11	4.2	14
77	Saccharomyces uvarum is responsible for the traditional fermentation of apple chicha in Patagonia. <i>FEMS Yeast Research</i> , 2017 , 17,	3.1	14
76	Update of the list of QPS-recommended biological agents intentionally added to food or feed as notified to EFSA 13: suitability of taxonomic units notified to EFSA until September 2020. <i>EFSA Journal</i> , 2021 , 19, e06377	2.3	14
75	Enological characterization of Spanish Saccharomyces kudriavzevii strains, one of the closest relatives to parental strains of winemaking and brewing Saccharomyces cerevisiae & S. kudriavzevii hybrids. <i>Food Microbiology</i> , 2016 , 53, 31-40	6	13
74	Statement on the update of the list of QPS-recommended biological agents intentionally added to food or feed as notified to EFSA 3: Suitability of taxonomic units notified to EFSA until September 2015. <i>EFSA Journal</i> , 2015 , 13, 4331	2.3	13
73	Cell-Wall Degrading Enzymes in the Release of Grape Aroma Precursors. <i>Food Science and Technology International</i> , 2001 , 7, 83-87	2.6	13
72	Evaluation of the use of phase-specific gene promoters for the expression of enological enzymes in an industrial wine yeast strain. <i>Biotechnology Letters</i> , 1996 , 18, 887-892	3	13
71	Physiological and genomic characterisation of Saccharomyces cerevisiae hybrids with improved fermentation performance and mannoprotein release capacity. <i>International Journal of Food Microbiology</i> , 2015 , 205, 30-40	5.8	12
70	Characterisation of the broad substrate specificity 2-keto acid decarboxylase Aro10p of Saccharomyces kudriavzevii and its implication in aroma development. <i>Microbial Cell Factories</i> , 2016 , 15, 51	6.4	12
69	Transcriptomics in human blood incubation reveals the importance of oxidative stress response in Saccharomyces cerevisiae clinical strains. <i>BMC Genomics</i> , 2012 , 13, 419	4.5	12
68	Update of the list of QPS-recommended biological agents intentionally added to food or feed as notified to EFSA 14: suitability of taxonomic units notified to EFSA until March 2021. <i>EFSA Journal</i> , 2021 , 19, e06689	2.3	12
67	Molecular profiling of beer wort fermentation diversity across natural Saccharomyces eubayanus isolates. <i>Microbial Biotechnology</i> , 2020 , 13, 1012-1025	6.3	11
66	Genetic Study of Natural Introgression Supports Delimitation of Biological Species in the Saccharomyces Sensu Stricto Complex. <i>Systematic and Applied Microbiology</i> , 1997 , 20, 595-601	4.2	11
65	Molecular evolution in yeast of biotechnological interest. <i>International Microbiology</i> , 2003 , 6, 201-5	3	11
64	Fermentative behaviour and competition capacity of cryotolerant Saccharomyces species in different nitrogen conditions. <i>International Journal of Food Microbiology</i> , 2019 , 291, 111-120	5.8	11

63	and Synthetic Wine Fermentation Performance Dissected by Predictive Modeling. <i>Frontiers in Microbiology</i> , 2018 , 9, 88	5.7	10
62	Ethanol Effects Involve Non-canonical Unfolded Protein Response Activation in Yeast Cells. <i>Frontiers in Microbiology</i> , 2017 , 8, 383	5.7	9
61	Genomic and transcriptomic analysis of aroma synthesis in two hybrids between <i>Saccharomyces cerevisiae</i> and <i>S. kudriavzevii</i> in winemaking conditions. <i>Microbial Cell Factories</i> , 2015 , 14, 128	6.4	9
60	Retrospective analysis of clinical yeast isolates in a hospital in the centre of Portugal: spectrum and revision of the identification procedures. <i>Medical Mycology</i> , 2009 , 47, 836-44	3.9	9
59	Differential proteomic analysis by SWATH-MS unravels the most dominant mechanisms underlying yeast adaptation to non-optimal temperatures under anaerobic conditions. <i>Scientific Reports</i> , 2020 , 10, 22329	4.9	9
58	Interspecific hybridisation among diverse <i>Saccharomyces</i> species: A combined biotechnological solution for low-temperature and nitrogen-limited wine fermentations. <i>International Journal of Food Microbiology</i> , 2019 , 310, 108331	5.8	8
57	Differential Contribution of the Parental Genomes to a \square Hybrid, Inferred by Phenomic, Genomic, and Transcriptomic Analyses, at Different Industrial Stress Conditions. <i>Frontiers in Bioengineering and Biotechnology</i> , 2020 , 8, 129	5.8	8
56	Increased mannoprotein content in wines produced by <i>Saccharomyces kudriavzevii</i> / <i>Saccharomyces cerevisiae</i> hybrids. <i>International Journal of Food Microbiology</i> , 2016 , 237, 35-38	5.8	8
55	Possible use of Biolog methodology for monitoring yeast presence in alcoholic fermentation for wine-making. <i>Journal of Applied Microbiology</i> , 2010 , 108, 1199-206	4.7	8
54	<i>Candida cabralensis</i> sp. nov., a yeast species isolated from traditional Spanish blue-veined Cabrales cheese. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2010 , 60, 2671-2674	2.2	8
53	Heterologous production in <i>saccharomyces cerevisiae</i> of different <i>aspergillus nidulans</i> xylanases of potential interest in oenology. <i>Journal of the Science of Food and Agriculture</i> , 1998 , 78, 315-320	4.3	8
52	Molecular Methods to Identify and Characterize Yeasts in Foods and Beverages 2006 , 55-82		8
51	Relationship between molecular and enological features of Patagonian wine yeasts: relevance in selection protocols. <i>World Journal of Microbiology and Biotechnology</i> , 2006 , 22, 827-833	4.4	8
50	Genome of <i>Saccharomyces cerevisiae</i> and Related Yeasts 2009 , 361-378		8
49	Expression of heterologous transporters in <i>Saccharomyces kudriavzevii</i> : A strategy for improving yeast salt tolerance and fermentation performance. <i>International Journal of Food Microbiology</i> , 2018 , 268, 27-34	5.8	7
48	RNAseq-based transcriptome comparison of <i>Saccharomyces cerevisiae</i> strains isolated from diverse fermentative environments. <i>International Journal of Food Microbiology</i> , 2017 , 257, 262-270	5.8	7
47	Comparative genomic analysis reveals a critical role of de novo nucleotide biosynthesis for <i>Saccharomyces cerevisiae</i> virulence. <i>PLoS ONE</i> , 2015 , 10, e0122382	3.7	7
46	Differences in activation of MAP kinases and variability in the polyglutamine tract of Slt2 in clinical and non-clinical isolates of <i>Saccharomyces cerevisiae</i> . <i>Yeast</i> , 2010 , 27, 549-61	3.4	7

45	Saccharomyces uvarum isolated from patagonian ciders shows excellent fermentative performance for low temperature cidemaking. <i>Food Research International</i> , 2019 , 126, 108656	7	6
44	Mixed growth curve data do not suffice to fully characterize the dynamics of mixed cultures. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020 , 117, 811-813	11.5	6
43	Aroma production and fermentation performance of <i>S. cerevisiae</i> × <i>S. kudriavzevii</i> natural hybrids under cold oenological conditions. <i>International Journal of Food Microbiology</i> , 2019 , 297, 51-59	5.8	5
42	The effect of two antifungal commercial formulations on the metabolism of a commercial <i>Saccharomyces cerevisiae</i> strain and their repercussion on fermentation evolution and phenylalanine catabolism. <i>Food Microbiology</i> , 2020 , 92, 103554	6	5
41	show low levels of traversal across the human blood brain barrier. <i>F1000Research</i> , 2017 , 6, 944	3.6	5
40	A comparison of the performance of natural hybrids <i>Saccharomyces cerevisiae</i> × <i>Saccharomyces kudriavzevii</i> at low temperatures reveals the crucial role of their <i>S. kudriavzevii</i> genomic contribution. <i>International Journal of Food Microbiology</i> , 2018 , 274, 12-19	5.8	5
39	Genetic and molecular study of the inability of the yeast <i>Kluyveromyces lactis</i> var. <i>drosophilae</i> to ferment lactose. <i>Microbiology</i> , 2006 , 75, 248-252	1.4	5
38	Phylogenetic relationships among wine yeast strains based on electrophoretic whole-cell protein patterns. <i>International Journal of Food Microbiology</i> , 1993 , 18, 115-25	5.8	5
37	Wine, Beer and Cider: Unravelling the Aroma Profile 2014 , 261-297		5
36	Genome structure reveals the diversity of mating mechanisms in x hybrids, and the genomic instability that promotes phenotypic diversity. <i>Microbial Genomics</i> , 2020 , 6,	4.4	5
35	A Multiphase Multiobjective Dynamic Genome-Scale Model Shows Different Redox Balancing among Yeast Species of the Genus in Fermentation. <i>MSystems</i> , 2021 , 6, e0026021	7.6	5
34	Metabolome segregation of four strains of <i>Saccharomyces cerevisiae</i> , <i>Saccharomyces uvarum</i> and <i>Saccharomyces kudriavzevii</i> conducted under low temperature oenological conditions. <i>Environmental Microbiology</i> , 2020 , 22, 3700-3721	5.2	4
33	<i>Candida patagonica</i> sp. nov., a new species of yeast from cellar surfaces. <i>Antonie Van Leeuwenhoek</i> , 2007 , 92, 77-81	2.1	4
32	Update of the list of QPS-recommended biological agents intentionally added to food or feed as notified to EFSA 15: suitability of taxonomic units notified to EFSA until September 2021.. <i>EFSA Journal</i> , 2022 , 20, e07045	2.3	4
31	<i>Saccharomyces cerevisiae</i> show low levels of traversal across the human blood brain barrier in vitro. <i>F1000Research</i> , 2017 , 6, 944	3.6	4
30	The effects of strains carrying alcoholic fermentation on the fermentative and varietal aroma profiles of young and aged Tempranillo wines. <i>Food Chemistry: X</i> , 2021 , 9, 100116	4.7	4
29	Phenotypic and genomic differences among <i>S. cerevisiae</i> strains in nitrogen requirements during wine fermentations. <i>Food Microbiology</i> , 2021 , 96, 103685	6	4
28	Screening of <i>Saccharomyces</i> strains for the capacity to produce desirable fermentative compounds under the influence of different nitrogen sources in synthetic wine fermentations. <i>Food Microbiology</i> , 2021 , 97, 103763	6	4

27	Transcriptomic analysis of <i>Saccharomyces cerevisiae</i> x <i>Saccharomyces kudriavzevii</i> hybrids during low temperature winemaking. <i>F1000Research</i> , 2017 , 6, 679	3.6	3
26	Differential proteomic analysis by SWATH-MS unravels the most dominant mechanisms underlying yeast adaptation to non-optimal temperatures under anaerobic conditions		3
25	Inheritance of winemaking stress factors tolerance in <i>Saccharomyces uvarum</i> /S. <i>eubayanus</i> hybrids. <i>International Journal of Food Microbiology</i> , 2020 , 320, 108500	5.8	3
24	Lipid Composition Analysis Reveals Mechanisms of Ethanol Tolerance in the Model Yeast. <i>Applied and Environmental Microbiology</i> , 2021 , 87, e0044021	4.8	3
23	Thermo-adaptive evolution to generate improved <i>Saccharomyces cerevisiae</i> strains for cocoa pulp fermentations. <i>International Journal of Food Microbiology</i> , 2021 , 342, 109077	5.8	3
22	Mitochondrial DNA polymorphism of the yeast <i>Saccharomyces bayanus</i> var. <i>uvarum</i> . <i>Microbiology</i> , 2010 , 79, 520-527	1.4	2
21	Four new <i>Candida cretensis</i> strains isolated from Spanish fermented sausages (chorizo): taxonomic and phylogenetic implications. <i>FEMS Yeast Research</i> , 2008 , 8, 485-91	3.1	2
20	Modulation of aroma and chemical composition of Albariño semi-synthetic wines by non-wine <i>Saccharomyces</i> yeasts and bottle aging.. <i>Food Microbiology</i> , 2022 , 104, 103981	6	2
19	Transcriptomic analysis of x hybrids during low temperature winemaking. <i>F1000Research</i> , 2017 , 6, 679	3.6	2
18	The qualified presumption of safety assessment and its role in EFSA risk evaluations: 15 years past. <i>FEMS Microbiology Letters</i> , 2019 , 366, i17-i23	2.9	2
17	Genomic instability in an interspecific hybrid of the genus : a matter of adaptability. <i>Microbial Genomics</i> , 2020 , 6,	4.4	2
16	Metabolic differences between a wild and a wine strain of <i>Saccharomyces cerevisiae</i> during fermentation unveiled by multi-omic analysis. <i>Environmental Microbiology</i> , 2021 , 23, 3059-3076	5.2	2
15	Molecular Identification and Characterization of Wine Yeasts 2011 , 111-141		1
14	The diverse effects of yeast on the aroma of non-sulfite added white wines throughout aging. <i>LWT - Food Science and Technology</i> , 2022 , 158, 113111	5.4	1
13	Mitochondrial introgression suggests extensive ancestral hybridization events among <i>Saccharomyces</i> species		1
12	Human-associated migration of Holarctic <i>Saccharomyces uvarum</i> strains to Patagonia. <i>Fungal Ecology</i> , 2020 , 48, 100990	4.1	1
11	Temperature Shapes Ecological Dynamics in Mixed Culture Fermentations Driven by Two Species of the Genus. <i>Frontiers in Bioengineering and Biotechnology</i> , 2020 , 8, 915	5.8	1
10	Impact of Nitrogen Addition on Wine Fermentation by Strains with Different Nitrogen Requirements. <i>Journal of Agricultural and Food Chemistry</i> , 2021 , 69, 6022-6031	5.7	1

9	Virulence related traits in yeast species associated with food; <i>Debaryomyces hansenii</i> , <i>Kluyveromyces marxianus</i> , and <i>Wickerhamomyces anomalus</i> . <i>Food Control</i> , 2021 , 124, 107901	6.2	1
8	Effect of transient thermal shocks on alcoholic fermentation performance. <i>International Journal of Food Microbiology</i> , 2020 , 312, 108362	5.8	1
7	Inter- and intraspecific chromosome pattern variation in the yeast genus <i>Kluyveromyces</i> 1998 , 14, 1341		1
6	Genetics of Yeasts 2008 , 167-179		1
5	Effect of non-wine <i>Saccharomyces</i> yeasts and bottle aging on the release and generation of aromas in semi-synthetic Tempranillo wines.. <i>International Journal of Food Microbiology</i> , 2022 , 365, 109554	5.8	0
4	Use of non-conventional yeasts to increase total acidity in the Cava base wines. <i>LWT - Food Science and Technology</i> , 2022 , 158, 113183	5.4	0
3	Convergent adaptation of <i>Saccharomyces uvarum</i> to sulfite, an antimicrobial preservative widely used in human-driven fermentations. <i>PLoS Genetics</i> , 2021 , 17, e1009872	6	0
2	Stt1 transporter mediating the uptake of glycerol is not a weak point of <i>Saccharomyces kudriavzevii</i> 's low osmotolerance. <i>Letters in Applied Microbiology</i> , 2019 , 68, 81-86	2.9	0
1	Genome-wide effect of non-optimal temperatures under anaerobic conditions on gene expression in <i>Saccharomyces cerevisiae</i> .. <i>Genomics</i> , 2022 , 110386	4.3	0