

## **ЦИТИРАНИЯ**

### **Цитирания (без автоцитирания)**

**Цитирана: Crous PW, Wingfield MJ, Guarro J, Hernández-Restrepo M, Sutton DA, Acharya K, Barber PA, Boekhout T, Dimitrov RA, Dueñas M, Dutta AK, Gené J, Gouliamova DE et al. Fungal Planet description sheets: 320-370. Persoonia. 2015, 34:167-266. doi: 10.3767/003158515X688433.**

### **Цитиращи:**

1. K. Tanaka, K. Hirayama, H. Yonezawa, G. Sato, A. Toriyabe, H. Kudo, A. Hashimoto, M. Matsumura, Y. Harada, Y. Kurihara, T. Shirouzu, and T. Hosoya. 2015. Revision of the Massarineae (Pleosporales, Dothideomycetes). *Studies in Mycology*. 82: 75–136.

**Цитирана: Yurkov AM, Kachalkin AV, Daniel HM, Groenewald M, Libkind D, de Garcia V, Zalar P, Gouliamova DE, Boekhout T, Begerow D 2015. Two yeast species *Cystobasidium psychroaquaticum* f.a. sp. nov. and *Cystobasidium rietchieii* f.a. sp. nov. isolated from natural environments, and the transfer of *Rhodotorula minuta* clade members to the genus *Cystobasidium*. *Antonie Van Leeuwenhoek*. 107:173–185.**

### **Цитиращи:**

2. Marcin Piątek & Matthias Lutz & Nourou S. Yorou. *Mycol Progress* 2015. A molecular phylogenetic framework for *Anthracoystis* (Ustilaginales), including five new combinations (inter alia for the asexual *Pseudozyma flocculosa*), and description of *Anthracoystis grodzinskae* sp. nov. *Mycol Progress*. 14: 88 DOI 10.1007/s11557-015-1114-3.

**Цитирана: Alexander Hristov, Dilnora Gouliamova, Lilyana Nacheva, Kolishka Tsekova. 2012. Biodegradation of Phenol in the Presence of Heavy Metals by Free and Immobilized Cells of Yeast Association. *Comptes Rendues de l'academie Bulgare des Sciences*. 65(3).**

### **Цитиращи:**

3. Maria Gerginova, Plamena Zlateva, Nadejda Peneva & Zlatka Alexieva. 2014. Influence of phenolic substrates utilised by yeast *Trichosporon cutaneum* on the degradation kinetics. *Biotechnology & Biotechnological Equipment*. 28(1): 33-37.

**Цитирана: Péter G, Dlačhy D, Tornai-Lehoczki J, Gouliamova D, Kurtzman CP. 2011, *Ogataea saltuana* sp. nov., a novel methanol-assimilating yeast species. *Antonie Van Leeuwenhoek*. 100 (3):375-383.**

### **Цитиращи:**

4. Čadež N, Dlačhy D, Raspor P. 2013. *Ogataea kolombanensis* sp. nov., *Ogataea histrianica* sp. nov. and *Ogataea deakii* sp. nov., three novel yeast species from plant sources. *Int J Syst Evol Microbiol*. 63(8): 3115-31223.

5. Hui, Feng-Li; Chen, Liang; Li, Zhi-Hui; et al. 2013. *Metschnikowia henanensis* sp nov., a new anamorphic yeast species isolated from rotten wood in China. *Antonie van Leeuwenhoek*. 103(4): 899-904.

**Цитирана: Petrova PM, Gouliamova DE. 2006. Rapid screening of plasmid-encoded small hsp genes. *Current Microbiology*. 53:422-426.**

**Цитираща:**

6. Tian H, Tan J, Zhang L, Gu X, Xu W, Guo X, Luo Y. 2012. Increase of stress resistance in *Lactococcus lactis* via a novel food-grade vector expressing a shsp gene from *Streptococcus thermophilus*. *Braz J Microbiol*. 43(3):1157-64.

**Цитирана: Stoyancheva G, Gouliamova D, Petrova P. 2009. Molecular study of the useful and the contaminant microflora in fermented dairy products. *Biotech & Biotech Equipment*. 23:551-555.**

**Цитиращи:**

7. Gayathri DA, Devaraja TN. 2011. *Lactobacillus* sp. as Probiotics for Human Health with Special Emphasis on Colorectal Cancer. *Indian J of Science and Technology*. 4(8):1008-1014.

**Цитирана: Richards TA, Vepritskiy AA, Gouliamova DE, Nierzwicki-Bauer S.A. 2005. The molecular diversity of freshwater picoeukaryotes from an oligotrophic lake reveals diverse, distinctive and globally dispersed lineages. *Environmental Microbiol*. 7(9), 1413–1425.**

**Цитиращи:**

8. J. Berglund. 2005. Pelagic microorganisms in the northern Baltic Sea Ecology, diversity and food web dynamics. PhD Thesis Department of Ecology and Environmental Science. Umea University, Sweden.

9. R. Wit and T. Bouvier. 2006. 'Everything is everywhere, but, the environment selects'; what did Baas Becking and Beijerinck really say? *Environ. Microbiol*. 8(4):755-758.

10. T. Weisse. 2006. Biodiversity of freshwater microorganisms-achievements, problems, and perspectives. *Polish Journal of Ecology*. 54(4):633-652.

11. B. Brown, G.V. Wolfe. 2006. Protist genetic diversity in the acidic hydrothermal environments of Lassen Volcanic National Park, USA. *Journal of Eukaryotic Microbiology*. 53(6):420-431.

12. E. Lefèvre, C. Bardot, C. Noël, J. Carrias, E. Viscogliosi, C. Amblard, T. Sime-Ngando. 2007. Unveiling fungal zooflagellates as members of freshwater picoeukaryotes: evidence from a molecular diversity study in a deep meromictic lake. *Environ. Microbiol*. 9 (1):61-71.

13. E. Lara, C. Berney, H. Harms & A. Chatzinotas. 2007. Cultivation-independent analysis reveals a shift in ciliate 18S rRNA gene diversity in a polycyclic aromatic hydrocarbon-polluted soil. *FEMS Microbiology Ecology*. 62(3):365-373.
14. E. Lefèvre, B. Roussel, C. Amblard, and T. Sime-Ngando..2008. The molecular diversity of freshwater picoeukaryotes reveals high occurrence of putative parasitoids in the plankton. *PLoS ONE*. 3(6): e2324.
15. S. Epstein and L. García. Missing” protists: a molecular prospective. 2008. *Biodivers Conserv.* 17:261-276.
16. M. Chen, F. Chen, Y. Yu, J. Ji, and F. Kong. 2008. Genetic Diversity of Eukaryotic Microorganisms in Lake Taihu, a Large Shallow Subtropical Lake in China. *Microb. Ecol.* 56(3):572-583.
17. M. Doherty. 2009. Distribution and diversity of planktonic ciliates: patterns and processes. Open Access Dissertations, University of Massachusetts Amherst, USA.
18. F. Lehembre. 2009. Réponses adaptatives des microorganismes eucaryotes du sol aux pollutions métalliques These de l’universite de Lyon délivrée par l’universite Claude Bernard Lyon, France.
19. Lecroq Béatrice. 2009. New insights into the diversity of deep-sea benthic foraminifera. PhD thesis. University of Geneve, Switzerland.
20. T. Stoeck, A. Behnke, R. Christen, L. Amaral-Zettler, M. J Rodriguez-Mora, A. Chistoserdov, W. Orsi and V. P. Edgcomb. 2009. Massively parallel tag sequencing reveals the complexity of anaerobic marine protistan communities. *BMC Biology*. 7-72. doi:10.1186/1741-7007-7-72.
21. E. Alexander, A. Stock, H. W. Breiner, A. Behnke, J. Bunge, M. M. Yakimov and T. Stoeck. 2009. Microbial eukaryotes in the hypersaline anoxic L'Atalante deep-sea basin. *Environ Microbiol.* 11(2):360-381.
22. R. Logares, J. Bråte, S. Bertilsson, J. L. Clasen, K. Shalchian-Tabrizi and K. Rengefors. 2009. Infrequent marine–freshwater transitions in the microbial world. *Trends in Microbiology*. 17(9):414-422.
23. Terrado, W. F. Vincent, C. Lovejoy. 2009. Mesopelagic protists: diversity and succession in a coastal Arctic ecosystem. *Aquat Microb Ecol.* 56: 25-40.
24. J. S. Park and A. G. B. Simpson. 2010. Characterization of halotolerant Bicosoecida and Placididea (Stramenopila) that are distinct from marine forms, and the phylogenetic pattern of salinity preference in heterotrophic stramenopiles. *Environmental Microbiology*. 12(5):1173–1184.

25. James William Harrison. A new and diverse plastid-bearing microbial eukaryote and its position on the eukaryotic tree of life. 2010. Masters thesis. University of Exeter, UK.
26. D. Klaveness, J. Brate, V. Patil et al. 2011. The 18S and 28S rDNA identity and phylogeny of the common lotic chrysophyte *Hydrurus foetidus*. *European Journal of Phycology* 46(3):283-291.
27. R. Medinger, V. Nolte, R. V. Pandey, S. Jost, Birgit Ottenwalder, C. Schlotterer and J. Boenigk. 2010. Diversity in a hidden world: potential and limitation of next-generation sequencing for surveys of molecular diversity of eukaryotic microorganisms. *Molecular Ecology*. 19(1):32–40.
28. Atteia A, van Lis R, Tielens AG, Martin WF. 2013. Anaerobic energy metabolism in unicellular photosynthetic eukaryotes. *Biochim Biophys Acta*. 1827(2):210-223.
29. J. Jung, H. Kim, S. Ryu et al. 2012. Development of single-nucleotide polymorphism-based phylum-specific PCR amplification technique: Application to the community analysis using ciliates as a reference organism. *Molecules and cells*. 34(4):383-391.
30. C. Wolf. 2007. Protist diversity and biogeography in the Pacific sector of the Southern Ocean. PhD thesis. Jacob University. Bremen, Germany.
31. Simon M, López GP, Moreira D, Jardillier L. 2013. New haptophyte lineages and multiple independent colonizations of freshwater ecosystems. *Environ Microbiol Rep*. 5(2):322-332.
32. S. Charvet. 2013. Diversité et dynamique des communautés de protistes dans le haut Arctique canadien. PhD thesis. Université Laval, Canada.
33. Estelle Silvia Kiliass. 2013. Mapping the Genetic Diversity of Eukaryotic Protists in the Arctic Ocean. PhD thesis. Jacobs University, Bremen, Germany.
- Цитирана: Gouliamova DE, Hennebert GL. 1998. Phylogenetic relationships in the *Saccharomyces cerevisiae* complex of species. *Mycotaxon*. 66: 337-353.**
- Цитираци:**
34. Pulvirenti, HV Nguyen, C Caggia. 2000. *Saccharomyces uvarum*, a proper species within *Saccharomyces sensu stricto*. *FEMS Microbiol. Letters*. 192:191–196.
35. GI Naumov, I Masneuf, ES Naumova, M Aigle et al. 2000. Association of *Saccharomyces bayanus* with some French wines: genetic analysis of yeast populations. *Research in Microbiology*. 151:683–691.
36. JHG von der Schulenburg, JM Hancock. 2001. Extreme length and length variation in the first ribosomal internal transcribed spacer of ladybird beetles (Coleoptera: Coccinellidae). *Mol Biol Evol*. 18:648-660.

37. GI Naumov, HV Nguyen, ES Naumova, A Michel. 2001. Genetic identification of *Saccharomyces bayanus* var. *uvarum*, a cider-fermenting yeast. *Int. J. Food Microbiol.* 65:163–171.
38. M Barros Lopes, JR Bellon, NJ Shirley et al. 2002. Evidence for multiple interspecific hybridization in *Saccharomyces sensu stricto* species. *FEMS Yeast Research.* 1:323–331.
39. M. Teresa Fernández, E. Barrio, Querol A. 2003. Analysis of the genetic variability in the species of the *Saccharomyces sensu stricto* complex. *Yeast.* 20:1213–1226.
40. E. S. Naumova, G. I. Naumov, Masneuf-Pomarède, M. 2005. Molecular genetic study of introgression between *Saccharomyces bayanus* and *S. cerevisiae*. *Yeast.* 22:1099-1115.
41. S. Hosoi-Tanabe, Y. Sako. 2006. Genetic differentiation in the marine dinoflagellates *Alexandrium tamarense* and *Alexandrium catenella* based on DNA–DNA hybridization. *Plankton and Benthos Research.* 1:138-146
42. Naumov GI, Naumova ES, Martynenko NN. et al. 2011. Taxonomy, ecology, and genetics of the yeast *Saccharomyces bayanus*: A new object for science and practice. *Microbiology.* 80(6): 735-742.

**Цитирана: Gouliamova DE, Hennebert GL. 1989. Diversity and affinities among species and strains of *Lipomyces*. 74 (4): 283-291.**

**Цитираци:**

43. C.P. Kurtzman, J Albertyn, E. Basehoar-Powers Multigene phylogenetic analysis of the Lipomycetaceae and the proposed transfer of *Zygozoma* species to *Lipomyces* and *Babjeviaanomala* to *Dipodascopsis*. *FEMS Yeast Research.* 2007. 7:1027-1034.
44. S Gamerschlag, H Mehlhorn, J Heukelbach, H. 2008. Repetitive sequences in the ITS1 region of the ribosomal DNA of *Tunga penetrans* and other flea species (Insecta, Siphonaptera) *Parasitology.* 102:193-199.
45. Kurtzman CP. 2011. In book: *The yeasts a taxonomic study*. Editotors Boekhout T, Fell J., Kurtzman CP. v1, page 560.

**Цитирана: Serdyuk I, Baranov V, Tsalkova T, Gulyamova D, Pavlov M, SpirinA, May R. Structural dynamics of translating ribosomes. Biochimie 74 (4), 1992, 299-306, ISSN: 0300-9084**

**Цитираци:**

46. Marina V. Rodnina, Andreas Savelsbergh, Wolfgang Wintermeyer. 1999. Dynamics of translation on the ribosome: molecular mechanics of translocation. *FEMS microbiology Rev.* 23(3):317 - 333.

47. J Frank, RK Agrawal. 2000. A ratchet-like inter-subunit reorganization of the ribosome during translocation. *Nature*. 406:318-322.
48. AT Gudkov. Structure and functions of the prokaryotic elongation factor G. 2001. *Molecular Biology*. 35:552-558.
49. MM Yusupov, GZ Yusupova, A Baucom, K Lieberman. 2001. Crystal structure of the ribosome at 5.5 Å resolution. *Science*. 29:883-896.
50. S Joseph. 2003. After the ribosome structure: how does translocation work? *RNA*. 9(2):160-164.
51. A. Lescoute and E. Westhof. 2006. SURVEY AND SUMMARY. The interaction networks of structured RNAs. *Nucleic Acids Research*, 2006, Vol. 34, No. 22 6587–6604.
52. Frank J. Gao H. Sengupta J. Gao N. Taylor D. 2007. The process of mRNA–tRNA translocation. *Proc Natl Acad Sci U S A*. 104(50):19671-8.