

James B Konopka

List of Publications by Year in descending order

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97
papers

5,984
citations

81900

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74163

75
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103
all docs

103
docs citations

103
times ranked

4715
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|-----|-----------|
| 1 | Plasma Membrane Phosphatidylinositol 4-Phosphate Is Necessary for Virulence of <i>Candida albicans</i> . <i>MBio</i> , 2022, , e0036622. | 4.1 | 0 |
| 2 | Comparison of Experimental Approaches Used to Determine the Structure and Function of the Class D G Protein-Coupled Yeast β -Factor Receptor. <i>Biomolecules</i> , 2022, 12, 761. | 4.0 | 2 |
| 3 | Microdomain Protein Nce102 Is a Local Sensor of Plasma Membrane Sphingolipid Balance. <i>Microbiology Spectrum</i> , 2022, 10, . | 3.0 | 9 |
| 4 | Receptors Pheromone Receptors (Yeast). , 2021, , 236-241. | | 1 |
| 5 | Differential Roles of a Family of Flavodoxin-Like Proteins That Promote Resistance to Quinone-Mediated Oxidative Stress in <i>Candida albicans</i> . <i>Infection and Immunity</i> , 2021, 89, . | 2.2 | 5 |
| 6 | A Conserved Machinery Underlies the Synthesis of a Chitosan Layer in the <i>Candida</i> Chlamyospore Cell Wall. <i>MSphere</i> , 2021, 6, . | 2.9 | 9 |
| 7 | Integrative multi-omics profiling reveals cAMP-independent mechanisms regulating hyphal morphogenesis in <i>Candida albicans</i> . <i>PLoS Pathogens</i> , 2021, 17, e1009861. | 4.7 | 10 |
| 8 | The Sur7 cytoplasmic C terminus regulates morphogenesis and stress responses in <i>Candida albicans</i> . <i>Molecular Microbiology</i> , 2021, 116, 1201-1215. | 2.5 | 6 |
| 9 | N-Acetylglucosamine Regulates Morphogenesis and Virulence Pathways in Fungi. <i>Journal of Fungi</i> (Basel, Switzerland), 2020, 6, 8. | 3.5 | 19 |
| 10 | Plasma Membrane MCC/Eisosome Domains Promote Stress Resistance in Fungi. <i>Microbiology and Molecular Biology Reviews</i> , 2020, 84, . | 6.6 | 30 |
| 11 | Fungal Pathogens: Shape-Shifting Invaders. <i>Trends in Microbiology</i> , 2020, 28, 922-933. | 7.7 | 27 |
| 12 | <i>Candida albicans</i> Agar Invasion Assays. <i>Bio-protocol</i> , 2020, 10, e3730. | 0.4 | 6 |
| 13 | Plasma membrane architecture protects <i>Candida albicans</i> from killing by copper. <i>PLoS Genetics</i> , 2019, 15, e1007911. | 3.5 | 37 |
| 14 | <i>Candida albicans</i> rvs161^{Δ} and rvs167^{Δ} Endocytosis Mutants Are Defective in Invasion into the Oral Cavity. <i>MBio</i> , 2019, 10, . | 4.1 | 6 |
| 15 | Pathogenic Effects of IFIT2 and Interferon- β during Fatal Systemic <i>Candida albicans</i> Infection. <i>MBio</i> , 2018, 9, . | 4.1 | 11 |
| 16 | Genetic Analysis of NDT80 Family Transcription Factors in <i>Candida albicans</i> Using New CRISPR-Cas9 Approaches. <i>MSphere</i> , 2018, 3, . | 2.9 | 39 |
| 17 | Phagocytes from Mice Lacking the Sts Phosphatases Have an Enhanced Antifungal Response to <i>Candida albicans</i> . <i>MBio</i> , 2018, 9, . | 4.1 | 27 |
| 18 | cAMP-independent signal pathways stimulate hyphal morphogenesis in <i>Candida albicans</i> . <i>Molecular Microbiology</i> , 2017, 103, 764-779. | 2.5 | 36 |

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|----|--|-----|-----------|
| 19 | An N-acetylglucosamine transporter required for arbuscular mycorrhizal symbioses in rice and maize. <i>Nature Plants</i> , 2017, 3, 17073. | 9.3 | 72 |
| 20 | N-Acetylglucosamine Metabolism Promotes Survival of <i>Candida albicans</i> in the Phagosome. <i>MSphere</i> , 2017, 2, . | 2.9 | 29 |
| 21 | MCC/Eisosomes Regulate Cell Wall Synthesis and Stress Responses in Fungi. <i>Journal of Fungi (Basel)</i> , 2017, 3, 10784314. <small>Tj ETQq1 1 0.784314 rgBT /Ove</small> | 3.5 | 31 |
| 22 | Modulating Host Signaling Pathways to Promote Resistance to Infection by <i>Candida albicans</i> . <i>Frontiers in Cellular and Infection Microbiology</i> , 2017, 7, 481. | 3.9 | 24 |
| 23 | Regulation of Hyphal Growth and N-Acetylglucosamine Catabolism by Two Transcription Factors in <i>Candida albicans</i> . <i>Genetics</i> , 2017, 206, 299-314. | 2.9 | 33 |
| 24 | Eisosomes promote the ability of Sur7 to regulate plasma membrane organization in <i>Candida albicans</i> . <i>Molecular Biology of the Cell</i> , 2016, 27, 1663-1675. | 2.1 | 32 |
| 25 | Plasma membrane organization promotes virulence of the human fungal pathogen <i>Candida albicans</i> . <i>Journal of Microbiology</i> , 2016, 54, 178-191. | 2.8 | 29 |
| 26 | Raft-Like Membrane Domains in Pathogenic Microorganisms. <i>Current Topics in Membranes</i> , 2015, 75, 233-268. | 0.9 | 46 |
| 27 | Identification of a New Class of Antifungals Targeting the Synthesis of Fungal Sphingolipids. <i>MBio</i> , 2015, 6, e00647. | 4.1 | 124 |
| 28 | Hyphal growth in <i>Candida albicans</i> does not require induction of hyphal-specific gene expression. <i>Molecular Biology of the Cell</i> , 2015, 26, 1174-1187. | 2.1 | 37 |
| 29 | The mitochondrial protein Mcu1 plays important roles in carbon source utilization, filamentation, and virulence in <i>Candida albicans</i> . <i>Fungal Genetics and Biology</i> , 2015, 81, 150-159. | 2.1 | 20 |
| 30 | Protection from Systemic <i>Candida albicans</i> Infection by Inactivation of the Sts Phosphatases. <i>Infection and Immunity</i> , 2015, 83, 637-645. | 2.2 | 35 |
| 31 | N-acetylglucosamine Regulates Virulence Properties in Microbial Pathogens. <i>PLoS Pathogens</i> , 2015, 11, e1004947. | 4.7 | 36 |
| 32 | Flavodoxin-Like Proteins Protect <i>Candida albicans</i> from Oxidative Stress and Promote Virulence. <i>PLoS Pathogens</i> , 2015, 11, e1005147. | 4.7 | 46 |
| 33 | Fungal Membrane Organization: The Eisosome Concept. <i>Annual Review of Microbiology</i> , 2014, 68, 377-393. | 7.3 | 118 |
| 34 | Distinct roles of cell wall biogenesis in yeast morphogenesis as revealed by multivariate analysis of high-dimensional morphometric data. <i>Molecular Biology of the Cell</i> , 2014, 25, 222-233. | 2.1 | 37 |
| 35 | Clathrin- and Arp2/3-Independent Endocytosis in the Fungal Pathogen <i>Candida albicans</i> . <i>MBio</i> , 2013, 4, e00476-13. | 4.1 | 39 |
| 36 | The MARVEL Domain Protein Nce102 Regulates Actin Organization and Invasive Growth of <i>Candida albicans</i> . <i>MBio</i> , 2013, 4, e00723-13. | 4.1 | 34 |

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|----|---|-----|-----------|
| 37 | N-acetylglucosamine (GlcNAc) Triggers a Rapid, Temperature-Responsive Morphogenetic Program in Thermally Dimorphic Fungi. <i>PLoS Genetics</i> , 2013, 9, e1003799. | 3.5 | 58 |
| 38 | A <i>Candida albicans</i> Temperature-Sensitive <i>cdc12-6</i> Mutant Identifies Roles for Septins in Selection of Sites of Germ Tube Formation and Hyphal Morphogenesis. <i>Eukaryotic Cell</i> , 2012, 11, 1210-1218. | 3.4 | 28 |
| 39 | Novel roles for GlcNAc in cell signaling. <i>Communicative and Integrative Biology</i> , 2012, 5, 156-159. | 1.4 | 41 |
| 40 | Sur7 Promotes Plasma Membrane Organization and Is Needed for Resistance to Stressful Conditions and to the Invasive Growth and Virulence of <i>Candida albicans</i> . <i>MBio</i> , 2012, 3, . | 4.1 | 63 |
| 41 | N-Acetylglucosamine Functions in Cell Signaling. <i>Scientifica</i> , 2012, 2012, 1-15. | 1.7 | 138 |
| 42 | The <i>Candida albicans</i> Sur7 Protein Is Needed for Proper Synthesis of the Fibrillar Component of the Cell Wall That Confers Strength. <i>Eukaryotic Cell</i> , 2011, 10, 72-80. | 3.4 | 50 |
| 43 | N-Acetylglucosamine (GlcNAc) Induction of Hyphal Morphogenesis and Transcriptional Responses in <i>Candida albicans</i> Are Not Dependent on Its Metabolism. <i>Journal of Biological Chemistry</i> , 2011, 286, 28671-28680. | 3.4 | 74 |
| 44 | Membrane Compartment Occupied by Can1 (MCC) and Eisosome Subdomains of the Fungal Plasma Membrane. <i>Membranes</i> , 2011, 1, 394-411. | 3.0 | 35 |
| 45 | Identification of GIG1 , a GlcNAc-Induced Gene in <i>Candida albicans</i> Needed for Normal Sensitivity to the Chitin Synthase Inhibitor Nikkomycin Z. <i>Eukaryotic Cell</i> , 2010, 9, 1476-1483. | 3.4 | 43 |
| 46 | Recognition of Yeast by Murine Macrophages Requires Mannan but Not Glucan. <i>Eukaryotic Cell</i> , 2010, 9, 1776-1787. | 3.4 | 82 |
| 47 | A Photostable Green Fluorescent Protein Variant for Analysis of Protein Localization in <i>Candida albicans</i> . <i>Eukaryotic Cell</i> , 2010, 9, 224-226. | 3.4 | 59 |
| 48 | Strategies for Isolating Constitutively Active and Dominant-Negative Pheromone Receptor Mutants in Yeast. <i>Methods in Enzymology</i> , 2010, 485, 329-348. | 1.0 | 3 |
| 49 | The Sur7 protein resides in punctate membrane subdomains and mediates spatial regulation of cell wall synthesis in <i>Candida albicans</i> . <i>Communicative and Integrative Biology</i> , 2009, 2, 76-77. | 1.4 | 24 |
| 50 | BAR Domain Proteins Rvs161 and Rvs167 Contribute to <i>Candida albicans</i> Endocytosis, Morphogenesis, and Virulence. <i>Infection and Immunity</i> , 2009, 77, 4150-4160. | 2.2 | 49 |
| 51 | Identification of Amino Acids at Two Dimer Interface Regions of the $\hat{1}\pm$ -Factor Receptor (Ste2). <i>Biochemistry</i> , 2009, 48, 7132-7139. | 2.5 | 33 |
| 52 | Scanning mutagenesis of regions in the $\hat{A}\hat{Z}\hat{A}\pm$ protein Gpa1 that are predicted to interact with yeast mating pheromone receptors. <i>FEMS Yeast Research</i> , 2008, 8, 71-80. | 2.3 | 3 |
| 53 | The Sur7 Protein Regulates Plasma Membrane Organization and Prevents Intracellular Cell Wall Growth in <i>Candida albicans</i> . <i>Molecular Biology of the Cell</i> , 2008, 19, 5214-5225. | 2.1 | 77 |
| 54 | <i>Saccharomyces cerevisiae</i> Afr1 Protein Is a Protein Phosphatase 1/Glc7-Targeting Subunit That Regulates the Septin Cytoskeleton during Mating. <i>Eukaryotic Cell</i> , 2008, 7, 1246-1255. | 3.4 | 16 |

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| 55 | Identification of an N-Acetylglucosamine Transporter That Mediates Hyphal Induction in <i>Candida albicans</i> . <i>Molecular Biology of the Cell</i> , 2007, 18, 965-975. | 2.1 | 124 |
| 56 | Sterol-Rich Plasma Membrane Domains in Fungi. <i>Eukaryotic Cell</i> , 2007, 6, 755-763. | 3.4 | 139 |
| 57 | Functional expression of mammalian receptors and membrane channels in different cells. <i>Journal of Structural Biology</i> , 2007, 159, 179-193. | 2.8 | 37 |
| 58 | Accessibility of Cysteine Residues Substituted into the Cytoplasmic Regions of the $\hat{I}\pm$ -Factor Receptor Identifies the Intracellular Residues That Are Available for G Protein Interaction. <i>Biochemistry</i> , 2006, 45, 15310-15317. | 2.5 | 20 |
| 59 | DEP-Domain-Mediated Regulation of GPCR Signaling Responses. <i>Cell</i> , 2006, 126, 1079-1093. | 28.9 | 166 |
| 60 | Cell Cycle Dynamics and Quorum Sensing in <i>Candida albicans</i> Chlamydo spores Are Distinct from Budding and Hyphal Growth. <i>Eukaryotic Cell</i> , 2005, 4, 1191-1202. | 3.4 | 55 |
| 61 | Septin Function in Yeast Model Systems and Pathogenic Fungi. <i>Eukaryotic Cell</i> , 2005, 4, 1503-1512. | 3.4 | 104 |
| 62 | Identification of Residues that Contribute to Receptor Activation through the Analysis of Compensatory Mutations in the G Protein-Coupled $\hat{I}\pm$ -Factor Receptor. <i>Biochemistry</i> , 2005, 44, 1278-1287. | 2.5 | 10 |
| 63 | Comparison of Class A and D G Protein-Coupled Receptors: Common Features in Structure and Activation. <i>Biochemistry</i> , 2005, 44, 8959-8975. | 2.5 | 80 |
| 64 | Successful expression of a functional yeast G-protein-coupled receptor (Ste2) in mammalian cells. <i>Biochemical and Biophysical Research Communications</i> , 2005, 329, 281-287. | 2.1 | 20 |
| 65 | Pheromone Receptors (Yeast). , 2004, , 256-261. | | 1 |
| 66 | Lipid Raft Polarization Contributes to Hyphal Growth in <i>Candida albicans</i> . <i>Eukaryotic Cell</i> , 2004, 3, 675-684. | 3.4 | 208 |
| 67 | A Microdomain Formed by the Extracellular Ends of the Transmembrane Domains Promotes Activation of the G Protein-Coupled $\hat{I}\pm$ -Factor Receptor. <i>Molecular and Cellular Biology</i> , 2004, 24, 2041-2051. | 2.3 | 28 |
| 68 | SUMO Modification of Septin-interacting Proteins in <i>Candida albicans</i> . <i>Journal of Biological Chemistry</i> , 2004, 279, 40861-40867. | 3.4 | 26 |
| 69 | Aromatic Residues at the Extracellular Ends of Transmembrane Domains 5 and 6 Promote Ligand Activation of the G Protein-Coupled $\hat{I}\pm$ -Factor Receptor. <i>Biochemistry</i> , 2003, 42, 293-301. | 2.5 | 46 |
| 70 | <i>Candida albicans</i> Septin Mutants Are Defective for Invasive Growth and Virulence. <i>Infection and Immunity</i> , 2003, 71, 4045-4051. | 2.2 | 85 |
| 71 | Septin Function in <i>Candida albicans</i> Morphogenesis. <i>Molecular Biology of the Cell</i> , 2002, 13, 2732-2746. | 2.1 | 166 |
| 72 | lqg1p links spatial and secretion landmarks to polarity and cytokinesis. <i>Journal of Cell Biology</i> , 2002, 159, 601-611. | 5.2 | 50 |

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|----|---|-----|-----------|
| 73 | Functional Assays for Mammalian G-Protein-Coupled Receptors in Yeast. <i>Methods in Enzymology</i> , 2002, 344, 92-111. | 1.0 | 16 |
| 74 | The Cytoplasmic End of Transmembrane Domain 3 Regulates the Activity of the <i>Saccharomyces cerevisiae</i> G-Protein-Coupled $\hat{\pm}$ -Factor Receptor. <i>Genetics</i> , 2002, 160, 429-443. | 2.9 | 30 |
| 75 | Mutational Analysis of the Role of N-Glycosylation in $\hat{\pm}$ -Factor Receptor Function. <i>Biochemistry</i> , 2001, 40, 9685-9694. | 2.5 | 39 |
| 76 | Constitutive activation of the <i>Saccharomyces cerevisiae</i> transcriptional regulator Ste12p by mutations at the amino-terminus. <i>Yeast</i> , 2000, 16, 1365-1375. | 1.7 | 13 |
| 77 | The C Terminus of the <i>Saccharomyces cerevisiae</i> $\hat{\pm}$ -Factor Receptor Contributes to the Formation of Preactivation Complexes with Its Cognate G Protein. <i>Molecular and Cellular Biology</i> , 2000, 20, 5321-5329. | 2.3 | 65 |
| 78 | Interaction between Transmembrane Domains Five and Six of the $\hat{\pm}$ -Factor Receptor. <i>Journal of Biological Chemistry</i> , 2000, 275, 26492-26499. | 3.4 | 41 |
| 79 | Point Mutations Identify a Conserved Region of the <i>Saccharomyces cerevisiae</i> AFR1 Gene That Is Essential for Both the Pheromone Signaling and Morphogenesis Functions. <i>Genetics</i> , 2000, 155, 43-55. | 2.9 | 15 |
| 80 | Visualization of Receptor-mediated Endocytosis in Yeast. <i>Molecular Biology of the Cell</i> , 1999, 10, 799-817. | 2.1 | 72 |
| 81 | Combining mutations in the incoming and outgoing pheromone signal pathways causes a synergistic mating defect in <i>Saccharomyces cerevisiae</i> . , 1999, 15, 765-780. | | 8 |
| 82 | Identification of a Polar Region in Transmembrane Domain 6 That Regulates the Function of the G Protein-Coupled $\hat{\pm}$ -Factor Receptor. <i>Molecular and Cellular Biology</i> , 1998, 18, 7205-7215. | 2.3 | 50 |
| 83 | Dominant-Negative Mutations in the G-Protein-Coupled $\hat{\pm}$ -Factor Receptor Map to the Extracellular Ends of the Transmembrane Segments. <i>Molecular and Cellular Biology</i> , 1998, 18, 5981-5991. | 2.3 | 59 |
| 84 | Afr1p Regulates the <i>Saccharomyces cerevisiae</i> $\hat{\pm}$ -Factor Receptor by a Mechanism That Is Distinct From Receptor Phosphorylation and Endocytosis. <i>Genetics</i> , 1998, 148, 625-635. | 2.9 | 15 |
| 85 | Two New S-Phase-Specific Genes from <i>Saccharomyces cerevisiae</i> . <i>Yeast</i> , 1997, 13, 1029-1042. | 1.7 | 208 |
| 86 | Two New S-Phase-Specific Genes from <i>Saccharomyces cerevisiae</i> . , 1997, 13, 1029. | | 1 |
| 87 | Mutation of Pro-258 in transmembrane domain 6 constitutively activates the G protein-coupled alpha-factor receptor.. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1996, 93, 6764-6769. | 7.1 | 111 |
| 88 | Signal Transduction: Prokaryotic and Simple Eukaryotic Systems. Janet Kurjan , Barry L. Taylor. <i>Quarterly Review of Biology</i> , 1995, 70, 338-339. | 0.1 | 0 |
| 89 | The pheromone signal pathway in <i>Saccharomyces cerevisiae</i> . <i>Antonie Van Leeuwenhoek</i> , 1992, 62, 95-108. | 1.7 | 24 |
| 90 | The pheromone signal pathway in <i>Saccharomyces cerevisiae</i> . , 1992, , 95-108. | | 11 |

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| 91 | <i>S. cerevisiae</i> $\hat{\pm}$ pheromone receptors activate a novel signal transduction pathway for mating partner discrimination. <i>Cell</i> , 1991, 67, 389-402. | 28.9 | 137 |
| 92 | Conjugation in <i>Saccharomyces cerevisiae</i> . <i>Annual Review of Cell Biology</i> , 1988, 4, 429-455. | 26.1 | 263 |
| 93 | The C-terminus of the <i>S. cerevisiae</i> $\hat{\pm}$ -pheromone receptor mediates an adaptive response to pheromone. <i>Cell</i> , 1988, 54, 609-620. | 28.9 | 237 |
| 94 | Expression of a translocated c-abl gene in hybrids of mouse fibroblasts and chronic myelogenous leukaemia cells. <i>Nature</i> , 1986, 319, 331-333. | 27.8 | 26 |
| 95 | Cell lines and clinical isolates derived from Ph1-positive chronic myelogenous leukemia patients express c-abl proteins with a common structural alteration.. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1985, 82, 1810-1814. | 7.1 | 238 |
| 96 | Activation of the abl oncogene in murine and human leukemias. <i>Biochimica Et Biophysica Acta: Reviews on Cancer</i> , 1985, 823, 1-17. | 7.4 | 41 |
| 97 | An alteration of the human c-abl protein in K562 leukemia cells unmask associated tyrosine kinase activity. <i>Cell</i> , 1984, 37, 1035-1042. | 28.9 | 884 |