

# Suzanne M Noble

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2325364/publications.pdf>

Version: 2024-02-01

18  
papers

3,116  
citations

623188

14  
h-index

839053

18  
g-index

19  
all docs

19  
docs citations

19  
times ranked

3952  
citing authors

#	ARTICLE	IF	CITATIONS
1	Candida albicans oscillating UME6 expression during intestinal colonization primes systemic Th17 protective immunity. Cell Reports, 2022, 39, 110837.	2.9	17
2	Unbelievable but True: Epigenetics and Chromatin in Fungi. Trends in Genetics, 2021, 37, 12-20.	2.9	13
3	Recording of DNA-binding events reveals the importance of a repurposed Candida albicans regulatory network for gut commensalism. Cell Host and Microbe, 2021, 29, 1002-1013.e9.	5.1	19
4	Adaptive immunity induces mutualism between commensal eukaryotes. Nature, 2021, 596, 114-118.	13.7	110
5	Leveraging machine learning essentiality predictions and chemogenomic interactions to identify antifungal targets. Nature Communications, 2021, 12, 6497.	5.8	33
6	A natural histone H2A variant lacking the Bub1 phosphorylation site and regulated depletion of centromeric histone CENP-A foster evolvability in Candida albicans. PLoS Biology, 2019, 17, e3000331.	2.6	16
7	Candida albicans Morphogenesis Programs Control the Balance between Gut Commensalism and Invasive Infection. Cell Host and Microbe, 2019, 25, 432-443.e6.	5.1	154
8	Visualization of <i>Candida albicans</i> in the Murine Gastrointestinal Tract Using Fluorescent In Situ Hybridization. Journal of Visualized Experiments, 2019, , .	0.2	3
9	Genome-wide association and HLA region fine-mapping studies identify susceptibility loci for multiple common infections. Nature Communications, 2017, 8, 599.	5.8	298
10	Candida albicans cell-type switching and functional plasticity in the mammalian host. Nature Reviews Microbiology, 2017, 15, 96-108.	13.6	399
11	Passage through the mammalian gut triggers a phenotypic switch that promotes Candida albicans commensalism. Nature Genetics, 2013, 45, 1088-1091.	9.4	260
12	Candida albicans specializations for iron homeostasis: from commensalism to virulence. Current Opinion in Microbiology, 2013, 16, 708-715.	2.3	85
13	Post-Transcriptional Regulation of the Sef1 Transcription Factor Controls the Virulence of Candida albicans in Its Mammalian Host. PLoS Pathogens, 2012, 8, e1002956.	2.1	74
14	An Iron Homeostasis Regulatory Circuit with Reciprocal Roles in Candida albicans Commensalism and Pathogenesis. Cell Host and Microbe, 2011, 10, 118-135.	5.1	287
15	Systematic screens of a Candida albicans homozygous deletion library decouple morphogenetic switching and pathogenicity. Nature Genetics, 2010, 42, 590-598.	9.4	632
16	Genetics and Molecular Biology in Candida albicans. Methods in Enzymology, 2010, 470, 737-758.	0.4	76
17	Genetics of <i>Candida albicans</i> , a Diploid Human Fungal Pathogen. Annual Review of Genetics, 2007, 41, 193-211.	3.2	110
18	Strains and Strategies for Large-Scale Gene Deletion Studies of the Diploid Human Fungal Pathogen Candida albicans. Eukaryotic Cell, 2005, 4, 298-309.	3.4	530