

# GOATOOLS: A Python library for Gene Ontology analysis

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Citation Report

#	ARTICLE	IF	CITATIONS
1	INTEGRO: an algorithm for data-integration and disease-gene association. , 2018, , .		10
2	GOnet: a tool for interactive Gene Ontology analysis. BMC Bioinformatics, 2018, 19, 470.	1.2	189
3	Identification of cordycepin biosynthesis-related genes through de novo transcriptome assembly and analysis in <i>Cordyceps cicadae</i> . Royal Society Open Science, 2018, 5, 181247.	1.1	21
4	CRlncRC: a machine learning-based method for cancer-related long noncoding RNA identification using integrated features. BMC Medical Genomics, 2018, 11, 120.	0.7	28
5	ZFLNC: a comprehensive and well-annotated database for zebrafish lncRNA. Database: the Journal of Biological Databases and Curation, 2018, 2018, .	1.4	27
6	Population Gene Introgression and High Genome Plasticity for the Zoonotic Pathogen <i>Streptococcus agalactiae</i> . Molecular Biology and Evolution, 2019, 36, 2572-2590.	3.5	36
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8	Pangloss: A Tool for Pan-Genome Analysis of Microbial Eukaryotes. Genes, 2019, 10, 521.	1.0	14
9	Changes in Gene Expression and Metabolite Profiles in <i>Platanus acerifolia</i> Leaves in Response to Feeding Damage Caused by <i>Corythucha ciliata</i> . International Journal of Molecular Sciences, 2019, 20, 3465.	1.8	4
10	Mechanistic insight into the roles of <i>Pseudomonas plecoglossicida</i> <i>clpV</i> gene in host-pathogen interactions with <i>Larimichthys crocea</i> by dual RNA-seq. Fish and Shellfish Immunology, 2019, 93, 344-353.	1.6	32
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14	Genetic interaction networks mediate individual statin drug response in <i>Saccharomyces cerevisiae</i> . Npj Systems Biology and Applications, 2019, 5, 35.	1.4	11
15	A Novel Nitrite-Base Aerobic Denitrifying Bacterium <i>Acinetobacter</i> sp. YTO3 and Its Transcriptome Analysis. Frontiers in Microbiology, 2019, 10, 2580.	1.5	17
16	A glance of the blood stage transcriptome of a Southeast Asian <i>Plasmodium ovale</i> isolate. PLoS Neglected Tropical Diseases, 2019, 13, e0007850.	1.3	5
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18	Deep learning-based model for predicting progression in patients with head and neck squamous cell carcinoma. Cancer Biomarkers, 2019, 27, 19-28.	0.8	15

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20	Effects and transcriptional responses in the hepatopancreas of red claw crayfish <i>Cherax quadricarinatus</i> under cold stress. <i>Journal of Thermal Biology</i> , 2019, 85, 102404.	1.1	40
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