余光创

数採

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开发二十多个软件包,其中有十个收录于最大的生物信息学软件项目 Bioconductor 中; 以第一作者在 Molecular Biology and Evolution, Methods in Ecology and Evolution 和 Bioinformatcs 等国际期刊发表多篇研究论文。

🎓 教育背景

香港大学	生物信息学	博士	2017
安徽医科大学	生物化学与分子生物学	硕士	2009
华南农业大学	生物技术	学士	2005
Institut Pasteur, Paris	Train the Trainers		2015

1 代表性著作/论文

◆ ESI 热点论文 (1 篇)

學 ESI 高被引论文 (4 篇)

被引次数: 2025, h-index = 15, i10-index = 20

- 1. **G** Yu*, TTY Lam, H Zhu, Y Guan*. Two methods for mapping and visualizing associated data on phylogeny using ggtree. *Molecular Biology and Evolution*. accepted. Impact Factor: 10.217
- 2. **G Yu**. Using meshes for MeSH term enrichment and semantic analyses. *Bioinformatics*. accepted. Impact Factor: 5.481
- 4. **G Yu**, QY He*. ReactomePA: an R/Bioconductor package for reactome pathway analysis and visualization. *Molecular BioSystems*. 2016, 12(2):477-479. Tcitation: 104, Impact Factor: 2.759
- 5. **G** Yu*, LG Wang, QY He*. ChIPseeker: an R/Bioconductor package for ChIP peak annotation, comparision and visualization. *Bioinformatics*. 2015, 31(14):2382-2383. Citation: 162, Impact Factor: 5.481
- 6. **G** Yu*, LG Wang, GR Yan, QY He*. DOSE: an R/Bioconductor package for Disease Ontology Semantic and Enrichment analysis. *Bioinformatics*. 2015, 31(4):608-609. Citation: 94, Impact Factor: 5.481
- 7. **G** Yu, LG Wang, Y Han, QY He*. clusterProfiler: an R package for comparing biological themes among gene clusters. *OMICS: A Journal of Integrative Biology*. 2012, 16(5):284-287. Citation: 733, Impact Factor: 2.370
- 8. **G** Yu[#], F Li[#], Y Qin, X Bo^{*}, Y Wu, S Wang^{*}. GOSemSim: an R package for measuring semantic similarity among GO terms and gene products. *Bioinformatics*. 2010, 26(7):976-978. Citation: 391, Impact Factor: 5.481

▲ 学术任责

Bioinformatics, BMC Bioinformatics, Computational Biology and Chemistry, Current Bioinformatics, Gastroenterology Research and Practice, Molecular BioSystems, Nucleic Acids Research, R Journal 等杂志审稿人