

GUANGCHUANG YU

Professor and dean of the Department of Bioinformatics, School of Basic Medicine, Southern Medical University. My research group has been focusing on the development and application of bioinformatics methods and tools, and has developed a number of widely recognized bioinformatics tools in phylogeny, omics data analysis, and data visualization to help biologists explore and understand data.

教授、博导，现任南方医科大学基础医学院生物信息学系系主任。

课题组一直专注于生物信息学方法工具开发及应用，在系统发育、组学数据分析和数据可视化上开发了多款广受同行认可的生物信息学工具，帮助生物学家探索和理解数据。



RESEARCH EXPERIENCE

- | | |
|----------------------|--|
| present

2018 | <ul style="list-style-type: none">● Professor
School of Basic Medical Sciences  Southern Medical University<ul style="list-style-type: none">• 2021-present, Director, Department of Bioinformatics• 2019-2021, Associate director, Department of Bioinformatics |
| 2018

2017 | <ul style="list-style-type: none">● Post-doctoral Fellow
State Key Laboratory of Infectious Diseases  The University of Hong Kong |
| 2013

2009 | <ul style="list-style-type: none">● Bioinformatician
Key Laboratory of Functional Protein Research of Guangdong Higher Education Institutes  |

EDUCATION

- | | |
|-------------------|--|
| 2017

2013 | <ul style="list-style-type: none">● PhD., Bioinformatics
The University of Hong Kong  |
| | <ul style="list-style-type: none">• Thesis: Big-data computational methods for studying molecular evolution: from accurate genome assembly to phylogenetic analysis and data integration |
| 2009

2006 | <ul style="list-style-type: none">● M.S., Biochemistry and Molecular Biology
Anhui Medical University  |
| | <ul style="list-style-type: none">• Thesis: miRNA靶基因计算分析新方法及其应用研究 |
| 2005

2001 | <ul style="list-style-type: none">● B.S., Biotechnology
South China Agricultural University  |
| 2019 | <ul style="list-style-type: none">● Antigenic cartography and antibody landscapes course
University of Cambridge  |

CONTACT

- ✉ gcyu1@smu.edu.cn
- 🐦 [guangchuangyu](#)
- /github.com/YuLab-SMU
- 🔗 <https://yulab-smu.top/>
- wechat: YuLabSMU (公众号)
- 📞 (86) 13826432521

Citation = 27852

H-index = 28

I10-index = 42



Last updated on 2023-03-30.

- 2018 ● 南方医科大学高层次人才“不忘初心 牢记使命”学习教育
延安干部培训学院 📍 Yan'an, CN
- 2015 ● Train the Trainers
Institut Pasteur 📍 Paris, FR
- 2009 | 2007 ● Visiting student
Beijing Institute of Radiation Medicine 📍 Beijing, CN

sertificate

- 2019 ● 高等学校教师资格
广东省教育厅
- 2015 ● Certificate of Teaching and Learning in Higher Education
Centre for Enhancement of Teaching and Learning 📍 The University of Hong Kong

trophy SCHOLARSHIPS & AWARDS

- 2023 | 2021 ● 连续3年入选“中国高被引学者”
Elsevier
- 2022 | 2021 ● 连续2年入选“终身科学影响力”排行榜
全球前2%顶尖科学家榜单
- 2022 | 2020 ● 连续3年入选“年度科学影响力”排行榜
全球前2%顶尖科学家榜单
- 2022 ● 2020学年教学优秀一等奖
南方医科大学
- 2017 | 2013 ● Postgraduate studentships
The University of Hong Kong
- 2012 ● Best paper of the year 2010
Journal of Integrated OMICS



GRANTS

- 2026
|
2023
- 基于噬菌体-细菌-人类宿主互作的多囊卵巢综合征患者肠道微生态解析与干预 (32270677)
主持, CNY ￥540,000, 国家自然科学基金, 面上项目
- 2025
|
2022
- 应用型生物信息学专项人才培养计划 (特色班)
主持, CNY ￥300,000, 广东省高校教学质量与教学改革工程建设项目
- 2022
|
2020
- 大样本多组学数据联合分析系统研究与构建
主持, CNY ￥600,000, 省部级项目
- 2020
|
2018
- 南方医科大学第三层次人才引进科研项目
主持, CNY ￥1,500,000
- 病毒性新发传染病的研究与防治 (2017YFE0190800)
参与, CNY ￥990,000, 中华人民共和国科学技术部, 国家重点研发计划
- 2013
- hnRNP蛋白在调控人肺癌细胞周期胞质分裂中的作用研究 (21613414)
主持, CNY ￥100,000, 中央高校基本科研业务费专项资金
- 2011
- HeLa细胞周期动态蛋白质组学分析 (21611303)
主持, CNY ￥20,000, 中央高校基本科研业务费专项资金

BOOKS

- 2023
- **R实战：系统发育树的数据集成操作及可视化**
ISBN: 9787121451829
 - 余光创 (著), 李林, 罗晓 (译)
 - 电子工业出版社
- 2022
- **Data Integration, Manipulation and Visualization of Phylogenetic Trees**
ISBN: 9781032233574, DOI: 10.1201/9781003279242
 - G Yu
 - Chapman and Hall/CRC
 - **Biomedical Data Visualization: Methods and Applications**
ISBN: 9782889761906, DOI: 10.3389/978-2-88976-190-6
 - G Yu, TTY Lam, CL Xiao, M Zhou (eds.)
 - Lausanne: Frontiers Media SA

2020

- **Gene Ontology Semantic Similarity Analysis Using GOSemSim**

ISBN: 9781071603000, DOI: 10.1007/978-1-0716-0301-7_11

- G Yu
- In: Kidder B. (eds) *Stem Cell Transcriptional Networks, Methods in Molecular Biology*, 2117:207-215.
- Humana, New York, NY

TEACHING EXPERIENCE

present
|
2022

- **生物信息学**

School of Basic Medical Sciences

 Southern Medical University

- Graduate level class

- **毒理学原理与方法**

School of Public Health

 Southern Medical University

- Graduate level class

- **生物信息学导论**

School of Basic Medical Sciences

 Southern Medical University

- Undergraduate level class

present
|
2021

- **基础医学研究与医学发展**

School of Basic Medical Sciences

 Southern Medical University

- Undergraduate level class

present
|
2019

- **基因组学**

School of Basic Medical Sciences

 Southern Medical University

- Undergraduate level class

2021
|
2019

- **基础医学新进展讲座**

School of Basic Medical Sciences

 Southern Medical University

- Undergraduate level class

2019

- **生物信息学科研训练**

School of Basic Medical Sciences

 Southern Medical University

- Undergraduate level class

- **BMB209: Workshops in Applied Bioinformatics**

Department of Biochemistry and Molecular Biology

 University of Southern Denmark

- PhD level class, invited lecture

2012
|
2011

● 功能蛋白质研究

College of Life Science and Technology

· Graduate level class

Jinan University

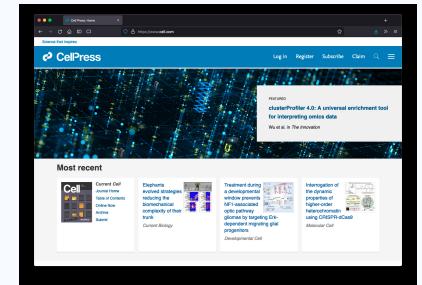
2023

■ PUBLICATIONS

- **MicrobiotaProcess: A comprehensive R package for deep mining microbiome**
The Innovation. 2023, 4(2):100388.
 - S Xu, L Zhan, W Tang, Q Wang, Z Dai, L Zhou, T Feng, M Chen, T Wu, E Hu, G Yu*
- **Editorial: Computational and Systematic Analysis of Multi-omics Data for Drug Discovery and Development**
Frontiers in Medicine. 2023, 10:1146896.
 - S Guo*, D Zhang, H Wang, Q An, G Yu, J Han, C Jiang, J Huang
- **Ggtree: A serialized data object for visualization of a phylogenetic tree and annotation data**
iMeta, 2022, 1(4):e56.
 - S Xu, L Li, X Luo, M Chen, W Tang, L Zhan, Z Dai, TT. Lam, Y Guan, G Yu*
- **Exploring epigenomic datasets by ChIPseeker**
Current Protocols in Bioinformatics, 2022, 2(10):e585.
 - Q Wang#, M Li#, T Wu, L Zhan, L Li, M Chen, W Xie, Z Xie, E Hu, S Xu, G Yu*
- **Characterization of toxin-antitoxin systems from public sequencing data: a case study in *Pseudomonas aeruginosa***
Frontiers in Microbiology. 2022, 13:951774.
 - Z Dai, T Wu, S Xu, L Zhou, W Tang, E Hu, L Zhan M Chen, G Yu*
- **Stemness Analysis Uncovers That The Peroxisome Proliferator-Activated Receptor Signaling Pathway Can Mediate Fatty Acid Homeostasis In Sorafenib-Resistant Hepatocellular Carcinoma Cells**
Frontiers in Oncology. 2022, 12:912694.
 - T Feng#, T Wu#, Y Zhang#, L Zhou#, S Liu, L Li, M Li, E Hu, Q Wang, X Fu, L Zhan, Z Xie, W Xie, X Huang*, X Shang*, G Yu*
- **ggmsa: a visual exploration tool for multiple sequence alignment and associated data**
Briefings in Bioinformatics. 2022, 23(4):bbac222.
 - L Zhou#, T Feng#, S Xu, F Gao, TT Lam, Q Wang, T Wu, H Huang, L Zhan, L Li, Y Guan, Z Dai*, G Yu*

2021

- **CBNplot: Bayesian network plots for enrichment analysis**
Bioinformatics. 2022, 38(10):2959-2960.
 - N Sato, Y Tamada, G Yu, Y Okuno*
- **Editorial: Biomedical Data Visualization: Methods and Applications**
Frontiers in Genetics. 2022, 13:890775.
 - T Wu, CL Xiao, TT Lam, G Yu*
- **Altered gut microbiota as an auxiliary diagnostic indicator for patients with fracture-related infection**
Frontiers in Microbiology. 2022, 13:723791.
 - X Zhao#, W Tang#, H Wan#, Z Lan H Qin, Q Lin, Y Hu, G Yu*, N Jiang*, B Yu*
- **Use ggbreak to effectively utilize plotting space to deal with large datasets and outliers**
Frontiers in Genetics. 2021, 12:774846.
 - S Xu#, M Chen#, T Feng, L Zhan, L Zhou, G Yu*
- **clusterProfiler 4.0: A universal enrichment tool for interpreting omics data**
The Innovation. 2021, 2(3):100141.
 - T Wu#, E Hu#, S Xu, M Chen, P Guo, Z Dai, T Feng, L Zhou, W Tang, L Zhan, X Fu, S Liu, X Bo*, G Yu*
- **ggtreeExtra: Compact visualization of richly annotated phylogenetic data**
Molecular Biology and Evolution. 2021, 38(9):4039-4042.
 - S Xu, Z Dai, P Guo, X Fu, S Liu, L Zhou, W Tang, T Feng, M Chen, L Zhan, T Wu, E Hu, Y Jiang*, X Bo*, G Yu*
- **ggVennDiagram: An intuitive, easy-to-use, and highly customizable R package to generate Venn diagram**
Frontiers in Genetics. 2021, 12:706907.
 - CH Gao, G Yu, P Cai*
- **IOBR: Multi-Omics Immuno-Oncology Biological Research to Decode Tumor Microenvironment and Signatures**
Frontiers in Immunology. 2021, 12:687975.
 - D Zeng#, Z Ye#, R Shen, G Yu, J Wu, Y Xiong, R Zhou, W Qiu, N Huang, L Sun, X Li, J Bin, Y Liao, M Shi, W Liao*
- **nCov2019: an R package for studying the COVID-19 coronavirus pandemic**
PeerJ. 2021, 9:e11421.
 - T Wu, E Hu, X Ge*, G Yu**



clusterProfiler (Wu et al. 2021),
the featured article in 2021-08-24's website banner of Cell Press

- **TreeSummarizedExperiment: a S4 class for data with hierarchical structure**
F1000Research. 2021, 9:1246.
· R Huang, C Soneson, FGM Ernst, KC Rue-Albrecht, G Yu, SC Hicks, MD Robinson*
- **Identification of prognostic Stromal–Immune Score–Based Genes in Hepatocellular Carcinoma microenvironment**
Frontiers in Genetics. 2021, 12: 625236.
· S Liu, G Yu*, L Liu*, X Zou, L Zhou, E Hu, Y Song
- 2020
- **Gut microbiota and metabolite alterations associated with reduced bone mineral density or bone metabolic indexes in postmenopausal osteoporosis**
Aging (Albany NY). 2020, 12(9):8583-8604.
· J He#, S Xu#, B Zhang, C Xiao, Z Chen, F Si, J Fu, X Lin, G Zheng*, G Yu*, J Chen*
- **Using ggtree to Visualize Data on Tree-Like Structure**
Current Protocols in Bioinformatics, 69(1):e96
· G Yu*
- **treeio: an R package for phylogenetic tree input and output with richly annotated and associated data**
Molecular Biology and Evolution. 2020, 37(2):599-603.
· LG Wang, TTY Lam, S Xu, Z Dai, L Zhou, T Feng, P Guo, CW Dunn, BR Jones, T Bradley, H Zhu, Y Guan, Y Jiang, G Yu*
· Impact Factor = 16.24
· 🏆 ESI Highly Cited Paper
- **RIdogram: drawing SVG graphics to visualize and map genome-wide data on the idiograms**
PeerJ Computer Science. 2020, 6:e251.
· Z Hao, D Lv, Y Ge, J Shi, D Weijers, G Yu*, J Chen*
· 🏆 ESI Highly Cited Paper
- 2018
- **Two methods for mapping and visualizing associated data on phylogeny using ggtree**
Molecular Biology and Evolution. 2018, 35(12):3041-3043.
· G Yu*, TTY Lam, H Zhu, Y Guan*
· Impact Factor = 14.797
- **Using meshes for MeSH term enrichment and semantic analyses**
Bioinformatics. 2018, 34(21):3766–3767.
· G Yu*

2017

- **ggtree: an R package for visualization and annotation of phylogenetic trees with their covariates and other associated data**

Methods in Ecology and Evolution. 2017, 8(1):28-36

- G Yu, DK Smith, H Zhu, Y Guan, TTY Lam *
- 🏆 ESI Highly Cited Paper

2016

- **ReactomePA: an R/Bioconductor package for reactome pathway analysis and visualization**

Molecular BioSystems. 2016, 12(2):477-479

- G Yu, QY He *
- 🏆 ESI Highly Cited Paper

- **Integrated Translomics with Proteomics to Identify Novel Iron-Transporting Proteins in *Streptococcus pneumoniae***

Frontiers in Microbiology. 2016, 7:78

- XY Yang, K He, G Du, X Wu, G Yu, Y Pan, G Zhang *, X Sun *, QY He *

- **Genetic characterization of highly pathogenic H5 influenza viruses from poultry in Taiwan, 2015**

Infection, Genetics and Evolution. 2016, 38:96-100

- PY Huang, CCD Lee, CH Yip, CL Cheung, G Yu, TTY Lam, DK Smith, H Zhu *, Y Guan *

2015

- **ChIPseeker: an R/Bioconductor package for ChIP peak annotation, comparision and visualization**

Bioinformatics. 2015, 31(14):2382-2383

- G Yu *, LG Wang, QY He *
- 🏆 ESI Highly Cited Paper

- **DOSE: an R/Bioconductor package for Disease Ontology Semantic and Enrichment analysis**

Bioinformatics. 2015, 31(4):608-609

- G Yu *, LG Wang, GR Yan, QY He *
- 🏆 ESI Highly Cited Paper

- **ACK1 Promotes Gastric Cancer Epithelial-Mesenchymal Transition and Metastasis through AKT-POU2F1-ECD Signaling**

Journal of Pathology. 2015, 236(2):175-185

- SH Xu, JZ Huang, ML Xu, G Yu, XF Yin, D Chen *, GR Yan *

- **Proteomic analysis on the copper resistance of *Streptococcus pneumoniae***

Metallomics. 2015, 7(3):448-454

- Z Guo, J Han, XY Yang, K Cao, K He, G Du, G Zeng, L Zhang, G Yu, Z Sun, QY He *, X Sun *



The *ggtree* (Yu et al. 2017) paper was selected as a feature article to celebrate the 10th anniversary of the launch of *Methods in Ecology and Evolution*.

- **Proteomic analysis on the antibacterial activity of a Ru(II) complex against *Streptococcus pneumonia***
Journal of Proteomics. 2015, 115:107-116
 · XY Yang, L Zhang, J Liu, N Li, G Yu, K Cao, J Han, G Zeng, Y Pan, X Sun*, QY He*
- 2014
 ● **Proteomic analysis of putative heme-binding proteins in *Streptococcus pyogenes***
Metalomics. 2014, 6(8):1451-1459
 · N Wang#, J Zhang#, L Zhang, XY Yang, N Li, G Yu, J Han, K Cao, Z Guo, X Sun*, QY He*
- 2013
 ● **Putative cobalt- and nickel-binding proteins and motifs in *Streptococcus pneumonia***
Metalomics. 2013, 5(7):928-935
 · X Sun#, G Yu#, Q Xu, N Li, C Xiao, X Yin, K Cao, J Han, QY He*
- **A novel andrographolide derivative AL-1 exerts its cytotoxicity on K562 cells through a ROS-dependent mechanism**
Proteomics. 2013, 13(1):169-178
 · YY Zhu#, G Yu#, Y Zhang, Z Xu, YQ Huang, GR Yan*, QY He*
- **A DNA-Based Semantic Fusion Model for Remote Sensing Data**
PLoS One. 2013, 8(10):e77090
 · H Sun*, J Weng, G Yu, RH Massawe
- **Quantitative Proteomics Characterization on the Anti-tumor Effects of Isodexyelephantopin against Nasopharyngeal Carcinoma**
Proteomics. 2013, 13(21):3222-3232
 · GR Yan#, Z Tan#, Y Wang, ML Xu, G Yu, Y Li*, QY He*
- **Design, SAR, angiogenic activities evaluation and pro-angiogenic mechanism of new marine cyclopeptide analogs**
Current Medicinal Chemistry. 2013, 20(9):1183-1194
 · J Li, X Lu, Q Wu, G Yu, Z Xu, L Qiu, Z Pei, Y Lin, J Pang*
- **Proteomic investigation of the interactome of FMNL1 in hematopoietic cells unveils a role in calcium-dependent membrane plasticity**
Journal of Proteomics. 2013, 78:72-82
 · Y Han, G Yu, H Sarioglu, A Caballero-Martinez, F Schlott, M Ueffing, H Haase, C Peschel, AM Krackhardt*
- 2012
 ● **LXtoo: an integrated live Linux distribution for the bioinformatics community**
BMC Research Notes. 2012, 5(1):360
 · G Yu#, LG Wang#, XH Meng, QY He*



clusterProfiler (Yu et al. 2012) 被中国科学技术信息研究所列为“2011-2021我国高被引论文中被引次数最高

- **clusterProfiler: an R package for comparing biological themes among gene clusters**
OMICS: A Journal of Integrative Biology. 2012, 16(5):284-287
· G Yu, LG Wang, Y Han, QY He^{*}
· 🏆 ESI Highly Cited Paper
- **Cellular MicroRNA let-7c Inhibits M1 Protein Expression of the H1N1 Influenza A Virus in Infected Human Lung Epithelial Cells**
Journal of Cellular and Molecular Medicine. 2012, 16(10):2539-2546
· Y Ma[#], J Yang[#], X Fan[#], H Zhao, W Hu, Z Li, G Yu, X Ding, J Wang, X Bo, X Zheng, Z Zhou^{*}, S Wang^{*}
- **Genistein-induced mitotic arrest of gastric cancer cells by downregulating KIF2oA, a proteomics study**
Proteomics. 2012, 12(14):2391-2399
· GR Yan[#], FY Zhou[#], BL Dang, Y Zhang, G Yu, X Liu, QY He^{*}
- 2011 ● **Functional similarity analysis of human virus-encoded miRNAs**
Journal of Clinical Bioinformatics. 2011, 1(1):15
· G Yu, QY He^{*}
- **A new method for measuring functional similarity of microRNAs**
Journal of Integrated OMICS. 2011, 1(1):49-54
· G Yu[#], CL Xiao[#], X Bo, CH Lu, Y Qin, S Zhan, QY He^{*}
- **Phosphoproteome profile of human lung cancer cell line A549**
Molecular BioSystems. 2011, 7(2):472-479
· G Yu[#], CL Xiao[#], CH Lu[#], HT Jia, F Ge, W Wang, XF Yin, HL Jia, JX He^{*}, QY He^{*}
- **Proteomic Analysis of Membrane Proteins from *Streptococcus pneumoniae* with Multiple Separation Methods plus High Accuracy Mass Spectrometry**
OMICS: A Journal of Integrative Biology. 2011, 15(10):683-694
· X Sun[#], XY Yang[#], XF Yin, G Yu, CL Xiao, X He, QY He^{*}
- 2010 ● **GOSemSim: an R package for measuring semantic similarity among GO terms and gene products**
Bioinformatics. 2010, 26(7):976-978
· G Yu[#], F Li[#], Y Qin, X Bo^{*}, Y Wu, S Wang^{*}
- 2009 ● **结合基因组分析预测炎症相关miRNA及其靶标**
军事医学科学院院刊. 2009, 33(3):245-247
· 余光创, 王超, 许尹, 伯晓晨, 王升启^{*}, 秦宜德^{*}

2007

- 依赖于5'端非编码区高级结构的真核生物mRNA翻译调控
中国生物化学与分子生物学报. 2007, 23(11):881-887
· 余光创, 秦宜德, 伯晓晨, 王升启*

CONFERENCE PROCEEDINGS

2021

- clusterProfiler 4.0: A universal enrichment tool for interpreting omics data
The 7th National Conference on Computational Biology and Bioinformatics

2020

- Data integration and visualization of phylogenetic trees
BioC Asia, Oct 2020  Virtual Conference

2019

- R语言在生物信息学数据可视化中的应用
第八届南方医院博士论坛 - 研而有信之修“生”养“信”, Oct 2019  Southern Medical University
- Data integration and visualization of phylogenetic trees
Managing Zoonotic Infections in the era of One Health, Oct 2019  Guangzhou Medical University

2017

- ggtree for tree visualization & annotation
Phylogenetic tree visualization workshop/hackathon, May 2017  The Field Museum of Natural History
- Computational Methods in Ecology and Evolution
The Second International Workshop on Environmental Microbiomes, Apr 2017  Sun Yat-sen University

2016

- R语言与生物信息学
潮博沙龙  Guangzhou, CN
- ggtree for visualization and annotation of phylogenetic trees
The 9th China-R Conference, May 2016  Renmin University of China

PROFESSIONAL AFFILIATIONS

- present | 2022
- **iMeta**
Associate Editor
 - **The Innovation**
Youth Editor
 - **南方医科大学基础医学院第二届学术分委员会**
委员
- present | 2021
- **中国生物工程学会**
计算生物学与生物信息学专业委员会委员
- present | 2020
- **广东省精准医学应用学会**
微生态医学分会常务委员 📍 Guangdong, CN
 - **广东省医学会**
医学遗传学分会青年委员会副主任委员 📍 Guangdong, CN
- present | 2019
- **中国计算机学会**
会员
- present | 2017
- **香港大学校友会**
会员
- present | 2014
- **广东潮博智库**
专家 📍 Guangdong, CN
- 2016
- **International Society for Influenza and other Respiratory Virus Disease**
Member
- 2015
- **International Network for Data Analysis**
Steering Committee 📍 Institut Pasteur