

Professor Xinghua (Victor) Pan, PhD, MD

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(Profile in English and Chinese 英文和中文簡曆)

CURRENT RESEARCH INTERESTS IN BRIEF 當前研究方向

- (1) 單細胞組學技術創新和應用
Single cell technology innovation and application
- (2) 基因組醫學
Genome medicine
- (3) 幹細胞與衰老和再生醫學
Stem cell and aging and regenerative medicine
- (4) 腫瘤異質性與精準醫學
Cancer heterogeneity and precision medicine
- (5) 組學生物信息學
Bioinformatics for omics data
- (6) 中醫藥組學
Chinese medicine omics

CURRENT POSITION AND WORKING EXPERIENCES:

Xinghua Pan, PhD, MD, Professor, Precision Regenerative Medicine Research Centre (PRMRC), Medical Science Division (MSD), and secondarily affiliated to State Key Laboratory of Quality Research in Chinese Medicine (SKL-QRCM), Macau University of Science and Technology (MUST), Macao 999078, China. He is a Supervisor for PhD candidate in Biomedical Science program in School of Medicine, and a Supervisor for Master candidate in Pharmacy in School of Pharmacy, MSD, MUST.

Prior to this from 2017-2024, Dr. Pan served as the Chairman and a Professor of Department Biochemistry and Molecular Biology, Southern Medical University (SMU), in Guangzhou, China, wherein he was the founding Director of Guangdong Provincial Key Laboratory of Single Cell Technology and Application, currently Vice Chairman of its Academic Committee, and an adjunct Professor there. In SMU he was initially appointed as the Director of the Guangdong Provincial Key Laboratory of Biochip in SMU. He was a Principal Investigator of an open fund grant of Shenzhen Bay Laboratory, and invited as an adjunct Professor at Hangzhou Cancer Hospital, Sichuan University, Guangdong Pharmaceutical University, and Jiujiang University. He has been an Adjunct Professor of Nanfang Hospital, Zhujiang Hospital, Guangdong Provincial People's Hospital, and Dongguan Maternal and Child Health Hospital. He has also been an adjunct PI of the Key Laboratory of Infectious Disease Prevention and Control of the Ministry of Education in South China, and of the Key Laboratory of Mental Health Research of the Ministry of Education, in SMU.

From 2004 to 2017, he entered as an Associate Research Scientist serving at Department of Genetics of Yale University School of Medicine, USA, then promoted to be a Research Scientist (PI) by Yale School of Medicine, while as a member of Yale Center Excellence in Genomics, Cancer Center, and Stem Cell Center.

From 2000 to 2004, he was employed as a Research Scientist and Enzymology Director at the Genomics Section of Molecular Staging Inc. (later acquired by Qiagen) in USA, meanwhile working as a Visiting Researcher at Yale University.

From 1994 year end to 1997, he was employed as an Associate Professor of Cell Biology and Medical Genetics at the Department of Biology in the Navy Medical University, Shanghai, China.

EDUCATIONS AND TRAINING EXPERIENCES

Dr. Pan got his Bachelor of medicine (MD equivalent) and MS of medicine at Southern Medical University (1980-1988, Guangzhou), PhD in genetics at Fudan University (1989-1993, Shanghai) with supervisors Dr. CC Tan (an Academician of CAS and NAS USA) and Dr. Zhencheng Geng. He got 2 terms of postdoc training: the 1st was in molecular oncology at the State Key Laboratory of Molecular Oncology of the Cancer Hospital of Chinese Academy of Medical Sciences and Peking Union Medical College (CAMS & PUMC, 1993-1994, Beijing) with Dr. Min Wu (an Academician of CAS), and the 2nd was in genomics at Yale University School of Medicine (1997-1999, USA) with Dr. Sherman M. Weissman (an Academician of NAS USA, and Yale Sterling Professor) on the Molecular Oncology and Development Program of Boyer Center for Molecular Medicine at Yale.

During this period, Dr. Pan Pan also got short time training, particularly on *Drosophila* cytogenetics in Peking University Department of Biology; on molecular oncology in Melbourne University and Ludwig Cancer Institute, Australia, supported by International Union Against Cancer (UICC) Scholarship; and on artificial yeast chromosome in Cold Spring Harbor Laboratory (CSHL, New York) supported by CSHL Scholarship.

In return, Dr. Pan since then has personally trained over 60 postdoctors, visiting scholars, and PhD, MS and BS candidates, and given lectures (biochemistry and molecular biology, medical genetics, cell biology) for near 10,000 undergraduate BS and postgraduate (and MD equivalent) students.

RESEARCH INTERESTS AND ACHIEVEMENTS:

Technological Explorations:

On the background of medicine, genetics and oncology, Dr. Pan started his technological exploration on single cell sequencing in early 2000s as a member of international ENCODE consortium. He has developed a dozen of novel technologies

for single cell analysis, including genomics (whole genome amplification WPA, medium-throughput scCNV-seq, and contributed to development REPLiG), epigenomics (2 methods for DNA methylome sequencing: scCGI-seq and msRRBS, and a method for closed chromatin profiling), transcriptomics (PMA, SMA and MUST-seq), telomere length measurement (SCT-pqPCR, USC-STELA), Single cell dual-multiplexing sequencing (NAMUL-seq) , and paved single-cell multiomics (the 1st approach for single cell co-analysis of transcriptome and genome). Currently Dr. Pan continues the exploration of the methods msCNVS and msRRBS for single cell detection of genomic and epigenomic variation/alternation in clinics for embryos (PGT, NIPT) and cancers (cancer liquid biopsy, circulating cancer cells/CTC/MRD), and scaling up the throughput for research applications.

Basic Biomedical Researches:

Meanwhile, Dr. Pan mostly applies the cutting edge technologies in single cell omics, spatial omics, multiomics, bioinformatics and machine learning, combining with other advanced cellular and molecular approaches, animal experiments, large clinical samples and public data to dissect the special mechanisms as well as the common intrinsic law for different physiological systems and disorders. Recent years his studies cover solid tumors (particularly liver fibrosis and liver cancer, and thyroid cancer, colorectal cancer, lung cancer, osteosarcoma), as well as leukemia (MDS/AML and ALL) for heterogeneity and evolution, stem/initiation cells and plasticity, cellular microenvironment, biomarker and precision medicine (diagnosis, treatment, drug response) models; stem cells and progenitor cells (particularly hematopoietic stem progenitor cells/HSC, mesenchymal stem cells/MSC), organoid, development, aging and regeneration; neuron and brain diseases; infection diseases and inflammatory bowel disease/IBD; and traditional Chinese medicine. He always enthusiastically seeks a collaboration with clinicians and scientists in variants of disciplines.

Summary of Publications and Innovations:

With a series of discoveries and innovations, Dr. Pan has contributed 160 publications cited in Google scholar, including in these journals: PNAS x7, Adv Science x4, Nucleic Acids Res x3, Nat Comm x3, Cell Mol Life Sci x3, Stem Cell Reports x3, Cancer Res x2, Cell Discovery x1, as well as coauthor in Cell x1 and Nature x4 papers, with a total impact factor >1000, and citations >8500. An ebook, Introduction to Single Cell Omics based on an associated research topic, led by Dr. Pan and published by Frontiers Medias in 2019, has been hit for more than 340 thousands, and downloaded for 50 thousands of time. He has contributed 15 invention patents issued in USA and China, and couples of PCTs, with a few more patents pending. On some of his research achievements Dr. Pan was interviewed by Genetic Engineering & Biotechnology News (GEN, 2015), PNAS Club (2013) and Technology Networks (2018).

The studies conducted by Dr. Pan and his team were supported by Natural Science Foundation of China (NSFC youth fund) and China Postdoctoral Foundation (1990s in China); National Institute of Health (NIH), Natural Science Foundation (NSF) and State of Connecticut (early in USA); and NSFC, China Minister of Science and Technology (MOST) , China Minister of Health/Education, Guangdong NSF, and Shenzhen Municipal foundation (recent years in China).

SCHOLASTIC SERVICES AND HONORS

Journal Reviewer and Editor:

Dr. Pan is a funding Associate Editor of Precision Clinical Medicine (by Oxford University Press) and Monocytomics (MCM), and currently the Executive Editor of the Special Issue on "Single Cell and Spatial Omics" of the Chinese Journal of Biochemistry and Molecular Biology, and an Editorial Board Member of the "Medical Molecular Cell Genetics Fundamentals" Core Textbook of the Ministry of Education 101 Plan.

He has been servicing as a reviewer or editorial board member for more than 30 internationally prestigious journals including Nature series such as Molecular Psychiatry, Nature Communications, Nature Protocols, Scientific Data and Science Reports; BMC series such as Molecular Cancer, BMC Genomics, BMC Biotechnology, BMC Medical Genetics; Frontiers series such as Frontiers in Cell and Developmental Biology, Frontiers in Genetics, Frontiers in Bioengineering and Biotechnology; journal on genomics and bioinformatics such as Genome Biology, Genome Medicine, Aging Cell, Protein and Cell, Giga Science, Gene, Genomics Proteomics and Bioinformatics, Journal of Genetics and Genomics, Current Bioinformatics, Computational and Structural Biotechnology; journals on cancer such as Cancer Communication, Cancer Letters, JNCC (J National Cancer Center), BBA Reviews on Cancer, Oncology Reports, Cancers; and others journals such as Science in China Life Sciences, Science Bulletin, Bone Research, Biology of Reproduction, Molecular Ecology Resource, Clinical and Translation Medicine, Intl J Mol Sci, Zoological Research, Hereditas (Beijing, in Chinese), Acta Academiae Medicinae Sinicae (in Chinese), Chinese Journal of Cell Biology (in Chinese), etc.

Grant Reviewer:

Dr. Pan also served or takes the role as a committee member or an invited reviewer for a number of research foundations such as the Medical Research Council (MRC, UK), Foundation against Cancer Belgium (i.e. Stichting tegen Kanker / Fondation contre le Cancer), NSFC, China MOST funds, Guangdong NSF, Chinese Academy of Sciences (CAS), and a few other local research foundations such as foundations of Zhejiang Province, Fujian Province, Guangxi Province, municipal foundations of Shenzhen, Guangzhou, Foshan, Dongguang, Nanjing, Zhongqin, Chengdu, Xi'an, Jiujiang, etc.

Academic degree and professional title reviewer:

Dr. Pan is an invited expert for doctoral thesis review by the Degree and Graduate Education Development Center of the Ministry of Education, and he has been an committee member of invited reviewer for PhD degree defenses or professorship evaluation for dozens of universities such as Shanghai Jiaotong University, Tsinghua University, Sun Yat-sen University, South China University of Technology, South China University of Technology, Guangdong Medical University, and Guangzhou Medical University.

Conference organization and speeches:

Dr. Pan leads or co-organizes some national and international academic conferences, particularly recently as the initiator and a co-Chairman of the 1st to currently the 3rd International Conference on Single cell and Spatial Omics (TICSSO) held in 2022 to 2025 (Hangzhou/Guangzhou, Shenzhen, and Shanghai to come), and the Chairman of the Forum of Single Cell Elite (FOSCE) for the session number 1 to 10 held in 2021 to 2023. In addition, he has given invited speech on over 100 prestigious academic meetings in the past 20 years.

Scholastic organization leadership:

Dr. Pan currently is a Vice Chairman of Guangdong Biochemical and Molecular Biology Society of Chinese Society of Biochemistry and Molecular Biology (CSBMB), a member of CSBMB Basic Medical Professional Committee, a member of the Biomedical Professional Committee of the All-China Federation of Overseas Chinese, an Executive Committee member of the Guangdong Medical Genetics Society, an Executive Committee member of the Genetics and Reproduction Committee of the Cross Strait Medical and Health Exchange Association. He is also a member of the Academic Committee of the Key Laboratory of Precision Medicine in Sichuan Province, and the Chief Advisory of the Academic Committee (former Committee Chairman) of the Jiangxi Provincial Key Laboratory of Systemic Biomedical Sciences.

He was elected as the President, then the Chairman of Board of Governor, and the Chairman of Board consecutively of the 26th to 28th committee consecutively from 2018 to 2020, and again the Chairman of Board of the 32nd committee in 2024 of Chinese Association of Science and Technology in USA. He was invited as a member of the International Advisory Committee for Advances in NGS in India.

He was invited as an Expert of Hangzhou Cancer Hospital, an Researcher of Jinan University (Guangzhou), a Guest Professor of Sichuan University, and of Guangdong Pharmaceutical University. He was granted as Nanjing Leading Talent, Guangzhou High Level Talent, and Ganpo Talent- Distinguished Professor of Jiangxi Province. He once was granted UICC (International Union Against Cancer) Scholarship, CSHL (Cold Spring Harbor Laboratory) Scholarship, PolyGenomics Fellowship, international science organization VEBLEO Fellow, and CastUSA Single Cell Genomics Pioneer Award.

SELECTED RECENT PUBLICATIONS 部分主要近期著作

(*通訊作者, IF 為最高影響因子)

1. Su H, Zhou X, Lin G, Luo C, Meng W, Lv C, Chen Y, Wen Z, Li X, Wu Y, Xiao C, Yang J, Lu J, Luo X, Hong X, Chen Y, Tam PKH*, Li C*, Sun H*, **Pan X***. Deciphering the Oncogenic Landscape of Hepatocytes through Integrated Single-Nucleus and Bulk RNA-Seq of Hepatocellular Carcinoma. *Adv Sci* (Weinh), 12, 2412944, doi:10.1002/advs.202412944 (2025). **(IF 15.1)**
2. Huang Y, Wang Q, Zhou W, Jiang Y, He K, Huang W, Feng Y, Wu H, Liu L, Pan Y, Huang Y, Chen Z, Li W, Huang Y, Lin G, Zhang Y, Ren Y, Xu K, Yu Y, Peng Y, **Pan X***, Pan S*, Hu H*, Hu Y*. Prenatal p25-activated Cdk5 induces pituitary tumorigenesis through MCM2 phosphorylation-mediated cell proliferation. *Neoplasia* 57, 101054, doi:10.1016/j.neo.2024.101054 (2024). **(IF 6.3)**
3. Gao J, Wu Y, Yu J, Qiu Y, Yi T, Luo C, Zhang J, Lu G, Li X, Xiong F, Wu X*, **Pan X***. Impact of genomic and epigenomic alterations of multigene on a multicancer pedigree. *Cancer Medicine* 13(13):e7394. doi: 10.1002/cam4.7394 (2024). **(IF 4.7)**
4. Chen F, Zhang K, Wang M, He Z, Yu B, Wang X, **Pan X**, Luo Y, Xu S, Lau JTY, Han C, Shi Y, Sun YE, Li S, Hu YP. VEGF-FGF Signaling Activates Quiescent CD63(+) Liver Stem Cells to Proliferate and Differentiate. *Adv Sci* (Weinh) 11, e2308711, doi:10.1002/advs.202308711 (2024). **(IF 14.3)**
5. Mai L, Wen Z, Zhang Y, Gao Y, Lin G, Lian Z, Yang X, Zhou J, Lin X, Luo C, Peng W, Chen C, Peng J, Liu D, Marjani SL, Tao Q, Cui Y, Zhang J, Wu X, Weissman SM, **Pan X***. Shortcut barcoding and early pooling for scalable multiplex single-cell reduced-representation CpG methylation sequencing at single nucleotide resolution. *Nucleic Acids Res* 51, e108, doi:10.1093/nar/gkad892 (2023). **(IF 19.1)**
6. Wu F, Wu F, Zhou Q, Liu X, Fei J, Zhang D, Wang W, Tao Y, Lin Y, Lin Q, **Pan X**, Sun K, Xie F, Bai L. A CCL2(+)DPP4(+) subset of mesenchymal stem cells expedites aberrant formation of creeping fat in humans. *Nat Commun* 14, 5830, doi:10.1038/s41467-023-41418-z (2023). **(IF 24.9)**
7. Bai X, Guo ZQ, Zhang YP, Fan ZZ, Liu LJ, Liu L, Long LL, Ma SC, Wang J, Fang Y, Tang XR, Zeng YJ, **Pan X***, Wu DH*, Dong ZY*. CDK4/6 inhibition triggers ICAM1-driven immune response and sensitizes LKB1 mutant lung cancer to immunotherapy. *Nat Commun* 14, 1247, doi:10.1038/s41467-023-36892-4 (2023). **(IF 24.9)**
8. Long LL, Ma SC, Guo ZQ, Zhang YP, Fan Z, Liu LJ, Liu L, Han DD, Leng MX, Wang J, Guo XJ, Tan JL, Cai XT, Lin Y, **Pan X**, Wu DH, Bai X, Dong ZY. PARP Inhibition Induces Synthetic Lethality and Adaptive Immunity in

LKB1-Mutant Lung Cancer. **Cancer Res** 83, 568-581,
doi:10.1158/0008-5472.CAN-22-1740 (2023). (IF 13.8)

9. Trubetskoy V, Pardiñas AF.....**Pan X**,Ripke S, Walters JTR, O'Donovan MC; Schizophrenia Working Group of the Psychiatric Genomics Consortium. Mapping genomic loci implicates genes and synaptic biology in schizophrenia. **Nature** 604, 502-508, doi:10.1038/s41586-022-04434-5 (2022).(IF 83.4)

10. Zhang Y, Xu S, Wen Z, Gao J, Li S, Weissman SM, **Pan X***. Sample-multiplexing approaches for single-cell sequencing. **Cell Mol Life Sci** 79, 466, doi:10.1007/s00018-022-04482-0 (2022).(IF 9.2)

11. Zhong C, Liu M, **Pan, X*** & Zhu H*. Tumorigenicity risk of iPSCs in vivo: nip it in the bud. **Precis Clin Med** 5, pbac004, doi:10.1093/pccmedi/pbac004 (2022).(IF 5.3, 潘為創刊副主編)

12. Qu R, He K, Yang Y, Fan T, Sun B, Khan AU, Huang W*, Ouyang J*, **Pan X***, Dai J*. The role of serum amyloid A1 in the adipogenic differentiation of human adipose-derived stem cells basing on single-cell RNA sequencing analysis. **Stem Cell Res Ther** 13, 187, doi:10.1186/s13287-022-02873-5 (2022).(IF 8.1)

13. Zhang H*, Wang L, Qiu Y, Gong F, Nong B and **Pan X***. Discovery of 194 Unreported Conopeptides and Identification of a New Protein Disulfide Isomerase in *Conus characteristicus* Using Integrated Transcriptomic and Proteomic Analysis. **Front Mar Sci** 9:792908. doi: 10.3389/fmars.2022.792908. (2022).(JCR Q1)

14. Zhang H* , Liang A and **Pan X***. Preparation and Functional Identification of a Novel Conotoxin QcMNCL-XIII0.1 from *Conus quercinus*. **Toxins** (Basel) 14(2):99. doi.org/10.3390/toxins14020099 (2022).(JCR Q1)

15. Lu Y, Liu M, Yang J, Weissman SM, **Pan X***, Katz SG*, Wang S*. Spatial transcriptome profiling by MERFISH reveals fetal liver hematopoietic stem cell niche architecture. **Cell Discov** 7, 47, doi:10.1038/s41421-021-00266-1 (2021).(IF 38.9)

16. Mai L, Qiu Y, Lian Z, Chen C, Wang L, Yin Y, Wang S, Yang X, Li Y, Peng W, Luo C, **Pan X***. MustSeq, an alternative approach for multiplexible strand-specific 3' end sequencing of mRNA transcriptome confers high efficiency and practicality. **RNA Biol** 18, 232-243, doi:10.1080/15476286.2021.1974208 (2021).(IF 5.48)

17. He K*, Chen X, Qiu YB, Liu Z, Wang WZ, Woodman N, Maldonado JE, **Pan X***. Mitogenome and phylogenetic analyses support rapid diversification among species groups of small-eared shrews genus *Cryptotis* (Mammalia: Eulipotyphla: Soricidae). **Zool Res** 42, 739-745, doi:10.24272/j.issn.2095-8137.2021.199 (2021).(JCR Q1)

18. Luo C, Peng W, Kang J, Chen C, Peng J, Wang Y, Tang Q, Xie H, Li Y, **Pan X***. Glutamine Regulates Cell Growth and Casein Synthesis through the CYTHs/ARFGAP1-Arf1-mTORC1 Pathway in Bovine Mammary Epithelial Cells

(cover story). **J Agric Food Chem** 69, 6810-6819, doi:10.1021/acs.jafc.1c02223 (2021).(JCR Q1)

19. Wang H, Gong P, Chen T, Gao S, Wu Z, Wang X, Li J, Marjani SL, Costa J, Weissman SM, Qi F*, **Pan X***, Liu L*. Colorectal Cancer Stem Cell States Uncovered by Simultaneous Single-Cell Analysis of Transcriptome and Telomeres. **Adv Sci** (Weinh) 8, 2004320, doi:10.1002/advs.202004320 (2021).(IF 15.1)

20. Zhou Y, Yang D, Yang Q, Lv X, Huang W, Zhou Z, Wang Y, Zhang Z, Yuan T, Ding X, Tang L, Zhang J, Yin J, Huang Y, Yu W, Wang Y, Zhou C, Su Y, He A, Sun Y, Shen Z, Qian B, Meng W, Fei J, Yao Y*, **Pan X***, Chen P*, Hu H*. Single-cell RNA landscape of intratumoral heterogeneity and immunosuppressive microenvironment in advanced osteosarcoma. **Nat Commun** 11, 6322, doi:10.1038/s41467-020-20059-6 (2020).(IF 24.9) 高被引論文。

21. Ma X, Guo J, Liu K, Chen L, Liu D, Dong S, Xia J, Long Q, Yue Y, Zhao P, Hu F, Xiao Z, **Pan X**, Xiao K, Cheng Z, Ke Z, Chen ZS, Zou C. Identification of a distinct luminal subgroup diagnosing and stratifying early stage prostate cancer by tissue-based single-cell RNA sequencing. **Mol Cancer** 19, 147, doi:10.1186/s12943-020-01264-9 (2020).(IF 41.4)

22. ENCODE Project Consortium (**Pan X** as a member); Moore JE, Purcaro MJ, Pratt HE, Epstein CB, et al. Expanded encyclopaedias of DNA elements in the human and mouse genomes. **Nature** 583, 699-710, doi:10.1038/s41586-020-2493-4 (2020).(IF 83.4)

23. ENCODE Project Consortium (**Pan X** as a member); Snyder MP, Gingeras TR, Moore JE, Weng Z, et al. Perspectives on ENCODE. **Nature** 583:693-698. doi: 10.1038/s41586-020-2449-8 (2020).(IF 83.4)

24. Huang P, Zhao Y, Zhong J, Zhang X, Liu Q, Qiu X, Chen S, Yan H, Hillyer C, Mohandas N, **Pan X***, Xu X*. Putative regulators for the continuum of erythroid differentiation revealed by single-cell transcriptome of human BM and UCB cells. **Proc Natl Acad Sci U S A** 117, 12868-12876, doi:10.1073/pnas.1915085117 (2020).(IF 12.8)

25. Situ B, Ye X, Zhao Q, Mai L, Huang Y, Wang S, Chen J, Li B, He B, Zhang Y, Zou J, Tang BZ, **Pan X**, Zheng L. Identification and Single-Cell Analysis of Viable Circulating Tumor Cells by a Mitochondrion-Specific AIE Bioprobe. **Adv Sci** (Weinh) 7, 1902760, doi:10.1002/advs.201902760 (2020).(IF 15.1)

26. Cen B, Wei Y, Huang W, Teng M, He S, Li J, Wang W, He G, Bai X, Liu X, Yuan Y, **Pan X***, Ji A*. An Efficient Bivalent Cyclic RGD-PIK3CB siRNA Conjugate for Specific Targeted Therapy against Glioblastoma In Vitro and In Vivo. **Mol Ther Nucleic Acids** 13, 220-232, doi:10.1016/j.omtn.2018.09.002 (2018).(IF 8.1)

27. Cen B, Liao W, Wang Z, Gao L, Wei Y, Huang W, He S, Wang W, Liu X, **Pan X***, Ji A*. Gelofusine Attenuates Tubulointerstitial Injury Induced by

cRGD-Conjugated siRNA by Regulating the TLR3 Signaling Pathway. **Mol Ther Nucleic Acids** 11, 300-311, doi:10.1016/j.omtn.2018.03.006 (2018). **(IF 8.1)**

28. Zhang J*, Spath SS, Marjani SL, Zhang W & **Pan X***. Characterization of cancer genomic heterogeneity by next-generation sequencing advances precision medicine in cancer treatment. **Precis Clin Med** 1, 29-48, doi:10.1093/pccmedi/pby007 (2018). **(IF 5.3)**(潘為創刊副主編)

29. Han L, Wu HJ, Zhu H, Kim KY, Marjani SL, Riester M, Euskirchen G, Zi X, Yang J, Han J, Snyder M, Park IH, Irizarry R, Weissman SM, Michor F*, Fan R*, **Pan X***. Bisulfite-independent analysis of CpG island methylation enables genome-scale stratification of single cells. **Nucleic Acids Res** 45(10):e77, doi:10.1093/nar/gkx026 (2017). **(IF 19.1)**

30. Yang J, Tanaka Y, Seay M, Li Z, Jin J, Garmire LX, Zhu X, Taylor A, Li W, Euskirchen G, Halene S, Kluger Y, Snyder MP, Park IH, **Pan X***, Weissman SM*. Single cell transcriptomics reveals unanticipated features of early hematopoietic precursors. **Nucleic Acids Res** 45, 1281-1296, doi:10.1093/nar/gkw1214 (2017). **(IF 19.1)**

31. Wu H, Zhang XY, Hu Z, Hou Q, Zhang H, Li Y, Li S, Yue J, Jiang Z, Weissman SM, **Pan X***, Ju BG*, Wu S*. Evolution and heterogeneity of non-hereditary colorectal cancer revealed by single-cell exome sequencing. **Oncogene** 36, 2857-2867, doi:10.1038/onc.2016.438 (2017). **(IF 9.9)**

32. Zhu W, Zhang XY, Marjani SL, Zhang J, Zhang W, Wu S*, **Pan X***. Next-generation molecular diagnosis: single-cell sequencing from bench to bedside. **Cell Mol Life Sci** 74, 869-880, doi:10.1007/s00018-016-2368-x (2017). **(IF 9.2)**

33. Zhang X, Marjani SL, Hu Z, Weissman SM, **Pan X***, Wu S*. Single-Cell Sequencing for Precise Cancer Research: Progress and Prospects. **Cancer Res** 76, 1305-1312, doi:10.1158/0008-5472.CAN-15-1907 (2016). **(IF 12.5)**

34. Cheng J, Roden CA, Pan W, Zhu S, Baccei A, **Pan X**, Jiang T, Kluger Y, Weissman SM, Guo S, Flavell RA, Ding Y, Lu J. A Molecular Chipper technology for CRISPR sgRNA library generation and functional mapping of noncoding regions. **Nat Commun** 7, 11178, doi:10.1038/ncomms11178 (2016). **(IF 24.9)**

35. Hysolli E, Tanaka Y, Su J, Kim KY, Zhong T, Janknecht R, Zhou XL, Geng L, Qiu C, **Pan X**, Jung YW, Cheng J, Lu J, Zhong M, Weissman SM, Park IH. Regulation of the DNA Methylation Landscape in Human Somatic Cell Reprogramming by the miR-29 Family. **Stem Cell Reports** 7, 43-54, doi:10.1016/j.stemcr.2016.05.014 (2016). **(IF 7.3)**

36. Gagliani N, Amezcuca Vesely MC, Iseppon A, Brockmann L, Xu H, Palm NW, de Zoete MR, Licona-Limón P, Paiva RS, Ching T, Weaver C, Zi X, **Pan X**, Fan R, Garmire LX, Cotton MJ, Drier Y, Bernstein B, Geginat J, Stockinger B, Esplugues E, Huber S, Flavell RA. Th17 cells transdifferentiate into regulatory T cells during

resolution of inflammation. **Nature** 523, 221-225, doi:10.1038/nature14452 (2015).(IF 83.4)

37. Tanaka Y, Hysolli E, Su J, Xiang Y, Kim KY, Zhong M, Li Y, Heydari K, Euskirchen G, Snyder MP, **Pan X**, Weissman SM, Park IH. Transcriptome Signature and Regulation in Human Somatic Cell Reprogramming. **Stem Cell Reports** 4, 1125-1139, doi:10.1016/j.stemcr.2015.04.009 (2015).(IF 7.3)

38. Kim KY, Hysolli E, Tanaka Y, Wang B, Jung YW, **Pan X**, Weissman SM, Park IH. X Chromosome of female cells shows dynamic changes in status during human somatic cell reprogramming. **Stem Cell Reports** 2, 896-909, doi:10.1016/j.stemcr.2014.04.003 (2014).(IF 7.3)

39. Liu N, Liu L* & **Pan X***. Single-cell analysis of the transcriptome and its application in the characterization of stem cells and early embryos. **Cell Mol Life Sci** 71, 2707-2715, doi:10.1007/s00018-014-1601-8 (2014).(IF 9.2)

40. Guo S, Zi X, Schulz VP, Cheng J, Zhong M, Koochaki SH, Megyola CM, **Pan X**, Heydari K, Weissman SM, Gallagher PG, Krause DS, Fan R, Lu J. Nonstochastic reprogramming from a privileged somatic cell state. **Cell** 156, 649-662, doi:10.1016/j.cell.2014.01.020 (2014).(IF 66.8)

41. Han L, Zi X, Garmire LX, Wu Y, Weissman SM, **Pan X***, Fan R*. Co-detection and sequencing of genes and transcripts from the same single cells facilitated by a microfluidics platform. **Sci Rep** 4, 6485, doi:10.1038/srep06485 (2014).(IF 5)

42. **Pan X**. Single Cell Analysis: From Technology to Biology and Medicine. **Single Cell Biol** 3, doi:10.4172/2168-9431.1000106 (2014).(潘為創刊主編)

43. Tanaka Y, Kim KY, Zhong M, **Pan X**, Weissman SM, Park IH. Transcriptional regulation in pluripotent stem cells by methyl CpG-binding protein 2 (MeCP2). **Hum Mol Genet** 23, 1045-1055, doi:10.1093/hmg/ddt500 (2014).(IF 9.3)

44. Dan J, Liu Y, Liu N, Chiourea M, Okuka M, Wu T, Ye X, Mou C, Wang L, Wang L, Yin Y, Yuan J, Zuo B, Wang F, Li Z, **Pan X**, Yin Z, Chen L, Keefe DL, Gagos S, Xiao A, Liu L. Rif1 maintains telomere length homeostasis of ESCs by mediating heterochromatin silencing. **Developmental Cell** 29, 7-19, doi:10.1016/j.devcel.2014.03.004 (2014).(IF 13.4)

45. Wang F, **Pan X***, Kalmbach K, Seth-Smith ML, Ye X, Antunes DM, Yin Y, Liu L*, Keefe DL*, Weissman SM*. Robust measurement of telomere length in single cells. **Proc Natl Acad Sci U S A** 110, E1906-1912, doi:10.1073/pnas.1306639110 (2013).(IF 12.8) Interviewed by **PNAS club**: <http://firstlook.pnas.org/new-ruler-for-telomere-length/>, New ruler for telomere length, May 10, 2013 by Sarah CP Williams; featured in "This week in PNAS - In this Issue": <http://www.pnas.org/content/110/21/8315.full>, May 21, 2013 by C.R., and featured as "Telomere length measurement in single cells" by Hannah Stower published in **Nature Reviews Genetics**, 2013. 14, 444, doi:10.1038/nrg3529, June 18, 2013.

46. **Pan X**, Durrett RE, Zhu H, Tanaka Y, Li Y, Zi X, Marjani SL, Euskirchen G, Ma C, Lamotte RH, Park IH, Snyder MP, Mason CE, Weissman SM. Two methods for full-length RNA sequencing for low quantities of cells and single cells. **Proc Natl Acad Sci U S A** 110, 594-599, doi:10.1073/pnas.1217322109 (2013). **(IF 12.8)**

47. Zhang Y, Schulz VP, Reed BD, Wang Z, **Pan X**, Mariani J, Euskirchen G, Snyder MP, Vaccarino FM, Ivanova N, Weissman SM, Szekely AM. Functional genomic screen of human stem cell differentiation reveals pathways involved in neurodevelopment and neurodegeneration. **Proc Natl Acad Sci U S A** 110, 12361-12366, doi:10.1073/pnas.1309725110 (2013). **(IF 12.8)**

48. Wang Y, Yang F, Zhang HX, Zi XY, **Pan XH**, Chen F, Luo WD, Li JX, Zhu HY, Hu YP. Cuprous oxide nanoparticles inhibit the growth and metastasis of melanoma by targeting mitochondria. **Cell Death Dis** 4, e783, doi:10.1038/cddis.2013.314 (2013). **(IF 9.7)**

49. Wang F, Yin Y, Ye X, Liu K, Zhu H, Wang L, Chiourea M, Okuka M, Ji G, Dan J, Zuo B, Li M, Zhang Q, Liu N, Chen L, **Pan X**, Gagos S, Keefe DL, Liu L. Molecular insights into the heterogeneity of telomere reprogramming in induced pluripotent stem cells. **Cell Res** 22, 757-768, doi:10.1038/cr.2011.201 (2012). **(IF 46.3)**

50. Liu Z, Hu Z, **Pan X**, Li M, Togun TA, Tuck D, Pelizzola M, Huang J, Ye X, Yin Y, Liu M, Li C, Chen Z, Wang F, Zhou L, Chen L, Keefe DL, Liu L. Germline competency of parthenogenetic embryonic stem cells from immature oocytes of adult mouse ovary. **Hum Mol Genet** 20, 1339-1352, doi:10.1093/hmg/ddr016 (2011). **(IF 9.3)**

51. Yasukochi Y, Maruyama O, Mahajan MC, Padden C, Euskirchen GM, Schulz V, Hirakawa H, Kuhara S, **Pan XH**, Newburger PE, Snyder M, Weissman SM. X chromosome-wide analyses of genomic DNA methylation states and gene expression in male and female neutrophils. **Proc Natl Acad Sci U S A** 107, 3704-3709, doi:10.1073/pnas.0914812107 (2010). **(IF 12.8)**

52. **Pan X**, Urban AE, Palejev D, Schulz V, Grubert F, Hu Y, Snyder M, Weissman SM. A procedure for highly specific, sensitive, and unbiased whole-genome amplification. **Proc Natl Acad Sci U S A** 105, 15499-15504, doi:10.1073/pnas.0808028105 (2008). **(IF 12.8)**

53. Xu Y, He Z, Zhu H, Chen X, Li J, Zhang H, **Pan X***, Hu Y*. Murine fertilized ovum, blastomere and morula cells lacking SP phenotype. **Sci China C Life Sci** 50, 762-765, doi:10.1007/s11427-007-0097-y (2007). **(IF 10.4)**

54. **Pan X** & Weissman SM. An approach for global scanning of single nucleotide variations. **Proc Natl Acad Sci U S A** 99, 9346-9351, doi:10.1073/pnas.132218699 (2002). **(IF 12.8)**

Books and Chapters 部分書及章節:

55. **Pan X**, Wu S, Weissman S. eds. Introduction to Single Cell Omics. Lausanne (Switzerland): **Frontiers Media**. ISSN 1664-8714, ISBN 978-2-88945-920-9, DOI 10.3389/978-2-88945-920-9. August 1, 2019 (Invited Editors).

56. **Pan X**, Urban AE and Weissman SM. Chapter 20: Enriching DNA sequencing with nucleotide variation by thymidine glycosylase combined with suppression PCR. In: PCR Technology: Current Innovations (3rd Edition), edited by Tania Nolan and Stephen A. Bustin, **CRC Press**. June 17, 2013. pp. 285-297 (total 475 pages). ISBN 9781439848050.

57. **Pan X**, Weissman SM. Chapter 16: Global analysis of DNA allelic variation (GADAV) by specific enrichment of mismatches and selective amplification of heterohybrids. In: PCR Technology: Current Innovations (2nd ed), edited by Thomas Weissensteiner, Hugh G. Griffin and Annette Griffin. **CRC Press**. November 13, 2003. pp. 163-174 (total 475 pages). ISBN 9780849311840.

58. **潘星華**, 李亞子。第 57 章, 單細胞測序技術 (第 1127-1149 頁)。《表觀遺傳學》(於文強、徐國良主編)。科學出版社, ISBN: 9787030737892。2023 年 3 月 1 日。

59. **潘星華**, 殷志新。第三章 遺傳信息的複製、轉錄與翻譯。《醫學分子生物學》第三版(胡維新、劉靜主編)。科學出版社, ISBN: 9787030666680。2021-01 (國家精品課程教材, 國家精品資源共享課配套教材)。

60. 趙紅珊, 楊玉霞, **潘星華**, 李傳洲。第十七章, 人類基因組和染色體。《醫學分子細胞遺傳基礎》(喬傑、高國全、左伋主編), 第 413-428 頁。北京大學醫學出版社, ISBN: 978-7-5659-3196-3, 2024 年 7 月(教育部 101 計劃核心教材)。

61. 易霞, **潘星華**。第二十二章, 基因表達調控。《醫學分子細胞遺傳基礎》(喬傑、高國全、左伋主編), 第 524-546 頁。北京大學醫學出版社, ISBN: 978-7-5659-3196-3, 2024 年 7 月(教育部 101 計劃核心教材)。

SELECTED RECENT INNOVATION PATENTS 部分近期發明專利

1. Weissman SM, **Pan X**. Methods for preparing cDNA from low quantities of cells. 2018-07-10, USPTO, US10017761 B2. (美國專利)

2. **Pan X**, Weissman SM. Methods for closed chromatin mapping and DNA methylation analysis for single cells, 2019-11-19, USPTO, US10480021 B2. (美國專利)

3. Keefe D, Weissman SM, Liu L, Wang F, **Pan X**. A method for a single cell analysis of telomere length. 2018-10-09, USPTO, US10093970 B2. (美國專利)

4. Weissman SM, Lasken R, **Pan X**. Methods for reducing the complexity of DNA sequences, 2001-09-11, US6,287,825; 2022-04-16, US6,372,434; 2022-04-12, US6,346,399 (3 patents in a series). (美國專利)

5. Weissman SM, **Pan X**. Methods for identifying genes associated with diseases or specific phenotypes. 2005-08-02, US6924104 B2. (美國專利)

6. Marchese FP, **Pan X**, Harold J. Trehalose-containing topical drying composition and method of using same. 2011-11-29, US 8067037B2. PCT Filing date 6/24/2008. (美國專利)

7. Wang J, Zhang J, Huang Z, and **Pan X**. A method for accurately identification of molecular interactions and their polarity and directionality. China Patent authorization number # ZL201910571327.7; Authorization date: June 19, 2020. (中國專利) 汪佳宏.....潘星華。一種準確識別分子相互作用及其極性和方向性方法。授權號#ZL201910571327.7; 授權日: 2020年6月19日。

8. Wang J, Zhang J, Huang Z, and **Pan X**. A method for screening gene keywords from PubMed literature. China Patent Authorization Number # ZL201910571336.6; Authorization date: June 16, 2020. (中國專利) 汪佳宏.....潘星華。一種從 PubMed 文獻篩選基因關鍵字的方法。授權號#ZL201910571336.6, 授權日: 2020年6月16日。

9. Huang Z, Zhang J, Wang J, Lin X, and **Pan X**. A method for annotating cell identity based on single-cell transcriptome clustering results. China patent authorization number # ZL201910242519.0; Authorization date: February 28, 2020. (中國專利) 黃仲曦.....潘星華。一種基於單細胞轉錄組聚類結果注釋細胞身份的方法, 專利授權號#ZL201910242519.0; 授權日: 2020年2月28日。

10. **Pan X**, Xu S, Zhang Y, Zhang J, Lin X, Wang J. Unmarked multisample mixed single cell sequencing technology based on sample genotype data splitting. China Patent Authorization Number #ZL202211555116.2, Authorization date: May 14, 2024. (中國專利) 潘星華等。基於樣品基因型的數據拆分的無標記多樣品混合單細胞測序技術。專利號# ZL202211555116.2, 授權日 2024年5月14日。

11. Huang Z, Shen W, Wen Y, Zhang J, and **Pan X**. A tumor suppressor gene and its application. Invention patents. China Patent Authorization Number # ZL201911152615.5; Authorization date: January 10, 2023. (中國專利) 黃仲曦...潘星華。一種以抑癌基因及其應用。專利號#ZL201911152615.5, 授權日 2022年1月10日。

12. **Pan X**, Lin G, Huang Z, Zhang J. A unique fragment sequence capture method based on single cell sequencing data. Application date: Oct 30, 2020. China authorization# ZL 202011200039.X. Authorization announcement# CN 112309500 B, and announcement date: Aug 30, 2024. (中國專利) 潘星華等。一種基於單細胞測序數據唯一片段序列捕獲方法。專利號#ZL 202011200039.X, 授權日 2024年8月30日。

13. **Pan X**, Mai L, Wang L, Qiu Y, Yin Y, Wang S. A new method, primer set, and reagent kit for high-throughput RNA sequencing and their application. China patent authorization # ZL2020 10248230.5; Authorization date: April 15, 2022. (中國

專利) 潘星華等。一種新的 RNA 高通量測序的方法、引物組和試劑盒及其應用。專利號#ZL202010248230.5, 授權日: 2022 年 4 月 15 日。

14. **Pan X**, Mai L, Lian Z, Zhang Y, Lin X, Li S, Yang X, Peng J. A set of barcode connectors and a medium throughput multiplex single-cell representative DNA methylation library construction and sequencing method. China patent application number CN2021103368157, application date: March 25, 2021; Patent acceptance number # 202110336815.7; Publication number CN115125624A, publication date: September 30, 2022. PCT application acceptance on January 21, 2022, PCT/CN2022/073322. (中國專利, 國際專利 PCT) 一組條形碼接頭以及中高通量多重單細胞代表性 DNA 甲基化建庫和測序方法。

15. **Pan X**, Mai L, Lian Z. Method for medium-throughput multi-single-cell representative DNA methylation library construction and sequencing. USPTO submission: EFS ID #48631744, application #18372695, mailed date: Nov 15th, 2023; Confirmation Number: #2711. This applicant is a CIP of PCT/CN2022/073322. (美國專利)

16. **Pan X**, Lin G, Chen C, Dong Z. A method for constructing a medium throughput single-cell copy number library and its application. China patent application number CN2021101331285, application date 2021-02-01; Patent acceptance number 202110133128.5; Publication number CN114836838A, publication date: August 2, 2022. PCT application accepted on January 21, 2022, PCT/CN2022/073321. (中國專利, 國際專利 PCT) 一種中高通量單細胞拷貝數文庫構建的方法及其應用。

17. **Pan X**, Lin G, Caiming C, Dong Z. Method for traceable medium-throughput single-cell copy number sequencing. USPTO submission: EFS ID #48631744, application #18372695, mailed date: Nov 15th, 2023; Confirmation Number: #2711. US Patent App. 18/228,664. This applicant is a CIP of PCT/CN2022/073321. (美國專利)

工作和學習簡歷 WORKING AND TRAINING EXPERIENCES

潘星華, 理學博士, 醫學碩士, 醫學學士; 澳門科技大學醫學部精准再生醫學研究中心教授, 兼澳門科技大學中藥質量控制國家重點實驗室教授, 生物醫學科學博士生導師、藥學碩士生導師。

2017-2024 年擔任南方醫科大學教授, 基礎醫學院學術委員會委員、生物化學與分子生物學教研室(即基因工程研究所)主任、廣東省單細胞技術與應用重點實驗室創始人和主任(現學術委員會副主任); 初入職該校時擔任廣東省生物芯片重點實驗室主任; 後申報省珠江學者學科和省特支計劃學科獲批並擔任負責人。先後擔任深圳灣實驗室開放基金 PI、南方醫院、珠江醫院、廣東省人民醫院、東莞婦幼保健院雙聘教授, 也兼為精神健康研究教育部重點實驗室和華南傳染病防治教育部重點實驗室 PI。2004-2017 年為美國耶魯大學(Yale University)醫學院遺傳系副研究員、研究員和項目 PI, 並兼為耶魯卓越基因組中心、癌症

中心和幹細胞中心成員。2000-2004 年在美國分子平臺公司（Molecular Staging Inc., 獲 Qiagen 並購）基因組學部擔任研究員和酶學主任, 兼耶魯大學客座研究員。1994 年底到 1997 間在上海海軍醫科大學生物學教研室擔任醫學遺傳學和細胞生物學副教授。

先後在南方醫科大學臨床醫學和基礎醫學院（1980-1988）、復旦大學遺傳與遺傳工程系（1989-1993）、中國醫學科學院暨北京協和醫學院（CAMS & PUMC）腫瘤醫院分子腫瘤學國家重點實驗室（1993-1994）和美國耶魯大學遺傳學系暨分子醫學波義中心（Boyer Center）分子腫瘤學與發育項目（1997-1999）獲得了學士、碩士、博士學位和 2 期博士後培訓, 先後師從談家楨、吳旻和 Sherman M. Weissman 教授等中國和美國國家科學院院士三位院士和庚鎮城教授。期間在北京大學生物系（學習果蠅染色體技術）、澳大利亞墨爾本大學及路德維希癌症研究所（獲得日內瓦國際抗癌聯盟/UICC Scholarship 支持學習分子腫瘤學）和美國紐約冷泉港實驗室（受 CSHL Scholarship 支持學習人工酵母染色體技術）短期進修。

潘博士先後培養博士後/訪問學者、博士、碩士 40 多名, 本科生和醫學生 10 多位, 近年曾為近一萬名本科生和研究生授課。

研究興趣和貢獻 RESEARCH INTERESTS AND CONTRIBUTIONS

潘星華課題組的主要研究方向主要是單細胞組學技術創新和應用、基因組醫學、幹細胞與衰老和再生醫學、腫瘤異質性與精準醫學、組學生物信息學及中醫藥組學等。

作為單細胞組學技術創新的早期探索者之一, 近二十餘年來專攻單細胞組學核心技術創新及其生物醫藥多個領域的應用研究, 先後創新單細胞技術 10 餘項, 包括單細胞基因組（參與開發 REPLIg 技術, 創建全基因組擴增技術 WPA 和拷貝數變異測序技術 msCNVS）、轉錄組（2 種技術 PMA, SMA 及 MUST-seq）和表觀組多種技術（包括 DNA 甲基化 2 種測序技術 scCGI-seq 和 msRRBS, 及染色質圖譜技術）、單細胞雙重多樣品測序方案 (NAMUL-seq), 首創單細胞多維組學技術(首報單細胞轉錄組和基因組共分析技術)及單細胞端粒長度檢測技術 (SCT-pqPCR, USC-STELA) 等。當前正在進一步探索 msCNVS 和 msRRBS 在腫瘤（液體活檢, 循環癌細胞 CTC/殘留病/MRD）和產前及植入前胚胎遺傳病（PGT, NIPT）的基因組突變和表觀基因組檢測等, 並進行高通量升級為基礎科研服務。

同時, 主要應用單細胞組學、空間組學、多維組學、生物信息學和機器學習等前沿技術手段, 結合其他先進的細胞和分子方法、動物實驗、大型臨床樣本和公共數據, 探索不同生理系統和疾病的特殊機制及精準防治, 和他們的共同內在規律。近年來, 他研究多種實體瘤（尤其是肝纖維化和肝癌, 以及甲狀腺癌、結直腸癌、肺癌、骨肉瘤）以及白血病 MDS/AML 和 ALL 的異質性和進化、幹細胞/起始細胞和可塑性、細胞微環境、生物標志物和精準醫學（診斷、治療、藥物反應）模型; 多種幹細胞和祖細胞（特別是造血幹祖細胞/HSPC、間充質幹細胞/MSC）、類器官、發育、衰老和再生; 神經元和腦病、HBV 及炎症性腸病/IBD 等疾病組學; 以及中醫藥組學。潘星華熱情尋求與不同學科的臨床醫生和科學家的合作。

先後獲中美發明專利授權 15 項，及發明專利多項在審查中，獲得廣泛應用或成功轉化產品或試劑盒 5 項。迄今學術著作獲 Google Scholar 收錄 160 餘篇，總影響因子超 1000，總被引用 8500 多次。主要論文包括期刊 PNAS x7、Adv Science x4、Nucleic Acids Res x3、Nat Comm x3、Cell Mol Life Sci x3、Stem Cell Reports x3、Cancer Res x2、Cell Discov x1、Oncogene x1 及 Cell x1 和 Nature x4 篇，其中作為人類基因組 DNA 元素百科全書計劃/ENCODE 聯盟成員參與 Nature x3 篇；主編研究專題/單一著作 Introduction to Single Cell Omics 2019 年電子版獲得全球點閱 34 萬多次,下載 5 萬多次。

基於其部分單細胞組學研究成果，潘星華獲得 GEN (Genetic Engineering & Biotechnology News, 2015) 獨家報導、美國國家科學院進展雜誌俱樂部的採訪 (PNAS Club, 2013) 及 Technology Networks 的專門採訪和報導 (2018)。

研究課題獲得中國國家自然科學基金、科技部重大專項基金、衛健委、教育部、廣東省重大基礎培育項目、廣東省基礎與應用基礎研究重點項目和面上項目、廣東省珠江領軍人才創新團隊、廣東省重點實驗室基金和澳門科技大學研究基金等資助，早前獲得美國國立衛生研究院、自然科學基金會及康州幹細胞基金會等多項基金和中國博士後科學基金及國自然青年基金支持。

學術兼職和榮譽 SCHOLASTIC SERVICES AND HONORS

潘星華擔任廣東省生物化學與分子生物學學會副理事長，廣東省醫學遺傳學會常務理事，中國生物化學與分子生物學會基礎醫學專委會委員，全國僑聯特聘專家和生物醫藥專委會委員，海峽兩岸醫藥衛生交流協會遺傳與生殖專委會常委，中國抗癌協會生物標志專委會委員，江西省系統生物醫學重點實驗室學術委員會主任(現顧問)，精準醫學四川省重點實驗室學術委員；曾任印度 Advances in NGS 國際顧問委員會委員；獲選相繼擔任全美中國旅美科技協會第 26-28 屆委員會會長、董事會主席和理事會主席 (第 32 屆再任理事會主席)。

獲邀擔任 Nature Protocols、Nature Communications、Nucleic Acids Res、Genome Biology、Genome Research、Genome Medicine、Aging Cell、Protein and Cell、Genomics Proteomics and Bioinformatics、Computational and Structural Biotechnology、Science Bulletin 等 30 多種 SCI 雜誌特邀評審專家或編輯及數種新興專業雜誌顧問。是 Monocytomics (MCM) 創刊副主編，及 Precision Clinical Medicine 雜誌 (牛津大學出版社/OUP 出版) 創刊副主編，也曾擔任歐洲出版社雜誌 Frontiers 的《Single Cell Omics》研究專題執行主編。目前還擔任中國生物化學與分子生物學學報《單細胞與空間組學》主題專刊執行主編，教育部 101 計劃核心教材《醫學分子細胞遺傳基礎》編委。

在國際國內專業大會發表主題報告或專題報告超 100 次，主持組織和參與組織國際學術會議多次，包括主導發起和擔任大會共同主席的首屆至迄今第三屆國際單細胞和空間組學大會 (TICSSO, 2022-2025, 杭州/廣州、深圳、上海)，擔任主席組織了國際壹細胞菁英論壇第一到十期 (FOSCE, 2021-2023)，參與組織第二至十屆華西精準醫學國際學術論壇等。

是英國醫學研究理事會 (MRC)、比利時癌症研究基金會和中國國家自然科技基金委、科技部、中國科學院及佛山、成都、杭州、西安等多個省市科研基金的特邀評審/論證專家，教育部學位與研究生教育發展中心博士論文評審專家

（答辯前盲審，答辯後抽檢），是廣東省、浙江省、廣西省及深圳、東莞、九江等地市科技專家委員會委員；曾為清華大學深圳研究院、中山大學、上海交通大學、華南理工大學、廣東醫科大學、廣州醫科大學等博士學位論文答辯、論文評審或教授職稱評審服務。

曾獲聘兼任杭州市引進智力專家及杭州市腫瘤醫院特邀研究員，暨南大學研究員，四川大學和廣東藥科大學等客座教授等。獲得廣州市高層次人才、南京市領軍人才、江西省贛鄱英才-高端柔性特聘教授。獲得國際科學組織 VEBLEO 會士、PolyGenomics 多基因組學學術獎及 CastUSA 單細胞基因組學先鋒獎。

MUST scholar: <https://scholar.must.edu.mo/scholar/106820>

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Being interviewed by scientific magazines:

<https://www.genengnews.com/insights/single-and-loving-it/?q=fluidigm>

<https://www.technologynetworks.com/genomics/articles/studying-the-genome-one-cell-at-a-time-308054>

<http://blog.pnas.org/2013/05/new-ruler-for-telomere-length/>

An the leading Editor of a research topic and its corresponding eBook

<https://www.frontiersin.org/research-topics/4004/single-cell-genomics-technology-and-application>

Precision Clinical Medicine editorial board in OUP Oxford Press:

https://academic.oup.com/pcm/pages/Editorial_Board

Monocytomics editorial board:

https://www.accscience.com/journal/MCM/about/editorial_board

The international conference 2023 TICSSO-1 and 2024 TICSSO-2:

<https://www.prnewswire.com/news-releases/the-first-international-conference-on-single-cell-and-spatial-omics-is-held-in-guangzhou-china-the-golden-age-of-single-cell-analysis-301819021.html>;

<http://www.fimmu.com/info/1139/17590.htm>;

<https://www.ticssso.org.cn/>

<https://www.prnewswire.com/news-releases/the-second-international-conference-on-single-cell-and-spatial-omics-ticssso-2--showcasing-the-state-of-the-art-bioscience-achievements-30211962.html>

<https://www.tmcnet.com/submit/2024/04/09/9991594.htm>

<https://baijiahao.baidu.com/s?id=1795218461823037932&wfr=spider&for=pc>

<https://zhuanlan.zhihu.com/p/664535376>