

<p>导师个人信息</p>	<p>简介：孙海汐 超级细胞所 博导</p> <p>研究方向：生物信息学，目前重点关注多组学技术、单细胞与空间转录组测序相关的生物信息学分析。</p> 
<p>科研项目列表</p>	<p>(1) 造血干细胞分子特征及扩增机理的单细胞水平研究，参与，国家任务 - 国家自然科学基金课题，2020-01-01/ 2023-12-31</p> <p>(2) 单细胞发育大数据，参与，研究所自选 - 研究所自主部署，2018-04-02/ 2021-03-31</p> <p>(3) CD 设计合成，参与，研究所自选 - 研究所自主部署，2018-03-19/ 2021-06-30</p> <p>(4) 噬菌体抗菌新应用，参与，研究所自选 - 研究所自主部署，2017-02-22/ 2021-12-31</p>
<p>培养成果介绍</p>	<p>已培养硕士研究生 1 人，目前在读硕士研究生 3 人、博士研究生 4 人；学生培养期间已发表 SCI 文章 14 篇，获批软著 10 个；其中 5 名学生获各类奖项 20 次。</p>
<p>出版信息</p>	<p>(1) Derivation of Intermediate Pluripotent Stem Cells Amenable to Primordial Germ Cell Specification, Cell Stem Cell, 2021, 第 1 作者</p> <p>(2) Downregulated miR-451a as a feature of the plasma cfRNA landscape reveals regulatory networks of IL-6/IL-6R-associated cytokine storms in COVID-19 patients, Cellular & Molecular Immunology, 2021, 通讯作者</p> <p>(3) Cross-species single-cell transcriptomic analysis reveals pre-gastrulation developmental differences among pigs, monkeys, and humans, Cell Discovery, 2021, 第 2 作者</p> <p>(4) Cell competition constitutes a barrier for interspecies chimerism, nature, 2021, 第 9 作者</p> <p>(5) Extraembryonic Endoderm (XEN) Cells Capable of Contributing to Embryonic Chimeras Established from Pig Embryos, Stem Cell Reports, 2021, 合作组作者</p> <p>(6) Transcriptome Analyses of β-Thalassemia -28(A>G) Mutation Using Isogenic Cell Models Generated by CRISPR/Cas9 and Asymmetric Single-Stranded Oligodeoxynucleotides (assODNs), Frontiers in Genetics, 2020, 第 2 作者</p> <p>(7) Low-Concentration Essential Amino Acids in PZM-3 Improve the Developmental Competence of Porcine Embryos Produced by</p>

	<p>Handmade Cloning, Cellular Reprogramming, 2020, 合作组作者</p> <p>(8) Restoration of β-globin expression with optimally designed lentiviral vector for β-thalassemia treatment in Chinese patients, Human Gene Therapy, 2020, 合作组作者</p> <p>(9) African Arowana Genome Provides Insights on Ancient Teleost Evolution, iScience, 2020, 第 10 作者</p> <p>(10) Single-Cell Sequencing of Peripheral Mononuclear Cells Reveals Distinct Immune Response Landscapes of COVID-19 and Influenza Patients, Immunity, 2020, 第 6 作者</p> <p>(11) Prophage Hunter: an integrative hunting tool for active prophages, Nucleic Acids Research, 2019, 第 1 作者</p> <p>(12) Characterization and allergic role of IL-33-induced neutrophil polarization, Cellular & Molecular Immunology, 2018, 第 1 作者</p> <p>(13) IL-23-induced macrophage polarization and its pathological roles in mice with imiquimod-induced psoriasis, Protein & Cell, 2018, 第 2 作者</p> <p>(14) Bioinformatics Approaches to Studying Plant Long Noncoding RNAs (lncRNAs): Identification and Functional Interpretation of lncRNAs from RNA-Seq Data Sets, Springer, 2019-01-01, 第 1 作者</p>
<p>专利成果 与奖项</p>	<p>专利成果:</p> <p>(1) 检测慢病毒插入位点的测序文库构建方法和慢病毒插入位点检测方法, CN201911376706.7, 2019, 第 5 作者</p> <p>(2) 从细菌全基因组序列中挖掘温和型噬菌体的方法、装置和存储介质, CN201880098544.2, 2018, 第 1 作者</p> <p>(3) 用于动物胚胎解离的试剂盒和方法, CN201810557944.7, 2018, 第 2 作者</p> <p>(4) 连通式层析槽, CN200720093882.6, 2007, 第 2 作者</p> <p>(5) Sequencing library construction method for detecting lentivirus insertion sites, and lentivirus insertion site detection method, CN:201911376706:A, 第 3 作者</p> <p>奖项:</p> <p>(1) 2020 年深圳华大生命科学研究院提名奖, 2020</p> <p>(2) 深圳市海外高层次人才, 2018</p> <p>(3) 盐田区梧桐人才, 2018</p>