

空间多组学临床转化医学创新研究策略

李振亚 博士

经典的多组学研究策略

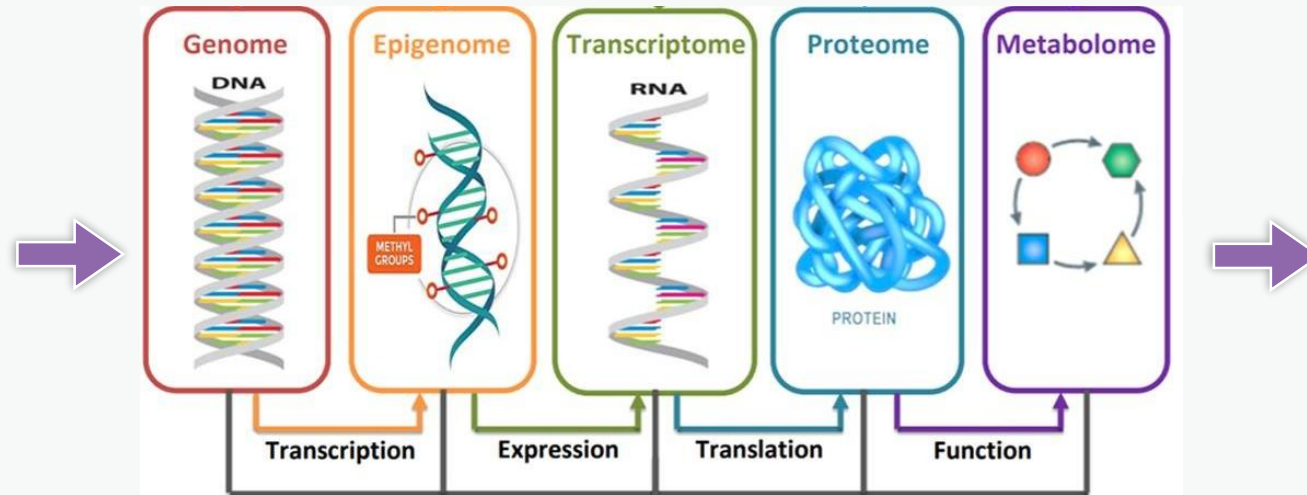
病人组织样本

外周血

外泌体

细胞模型

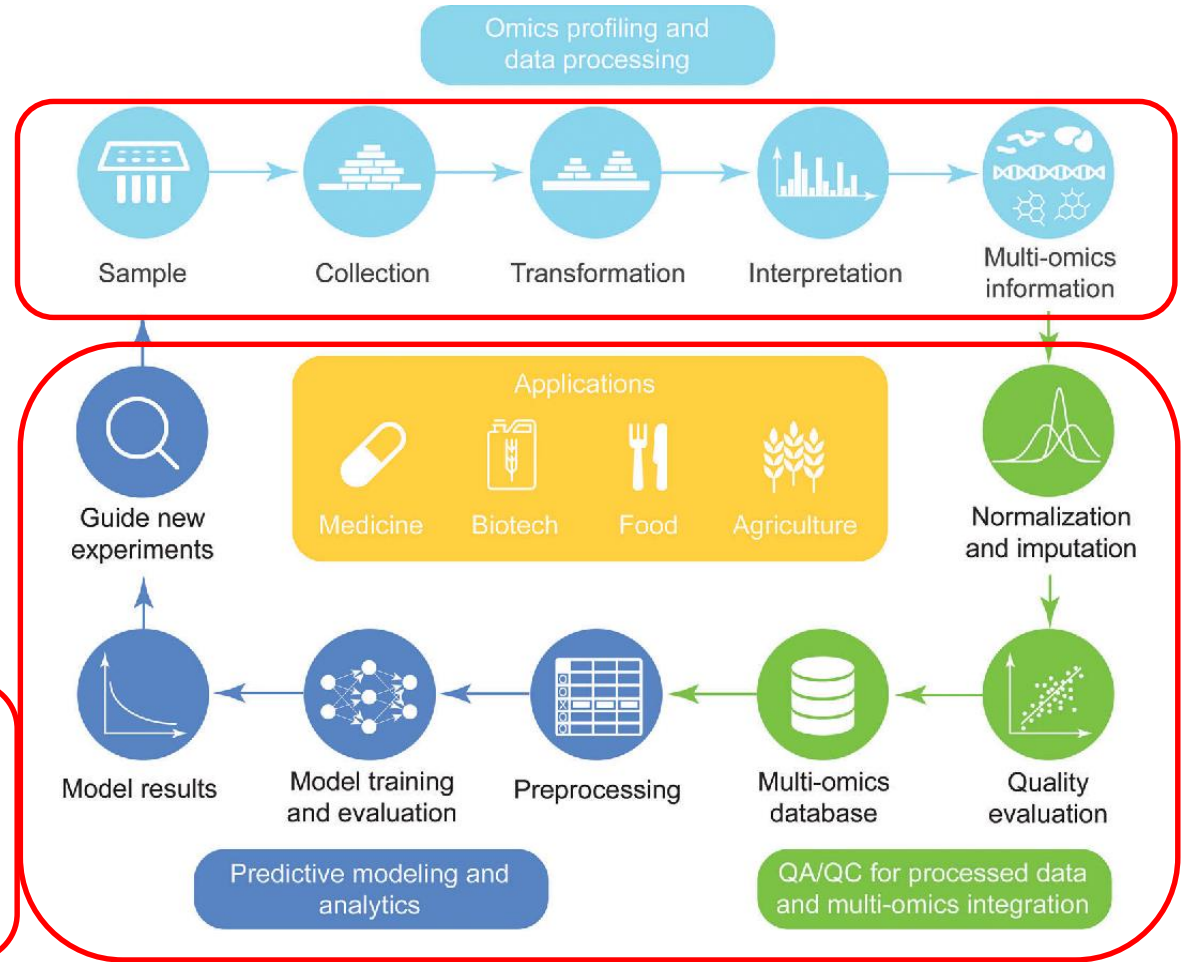
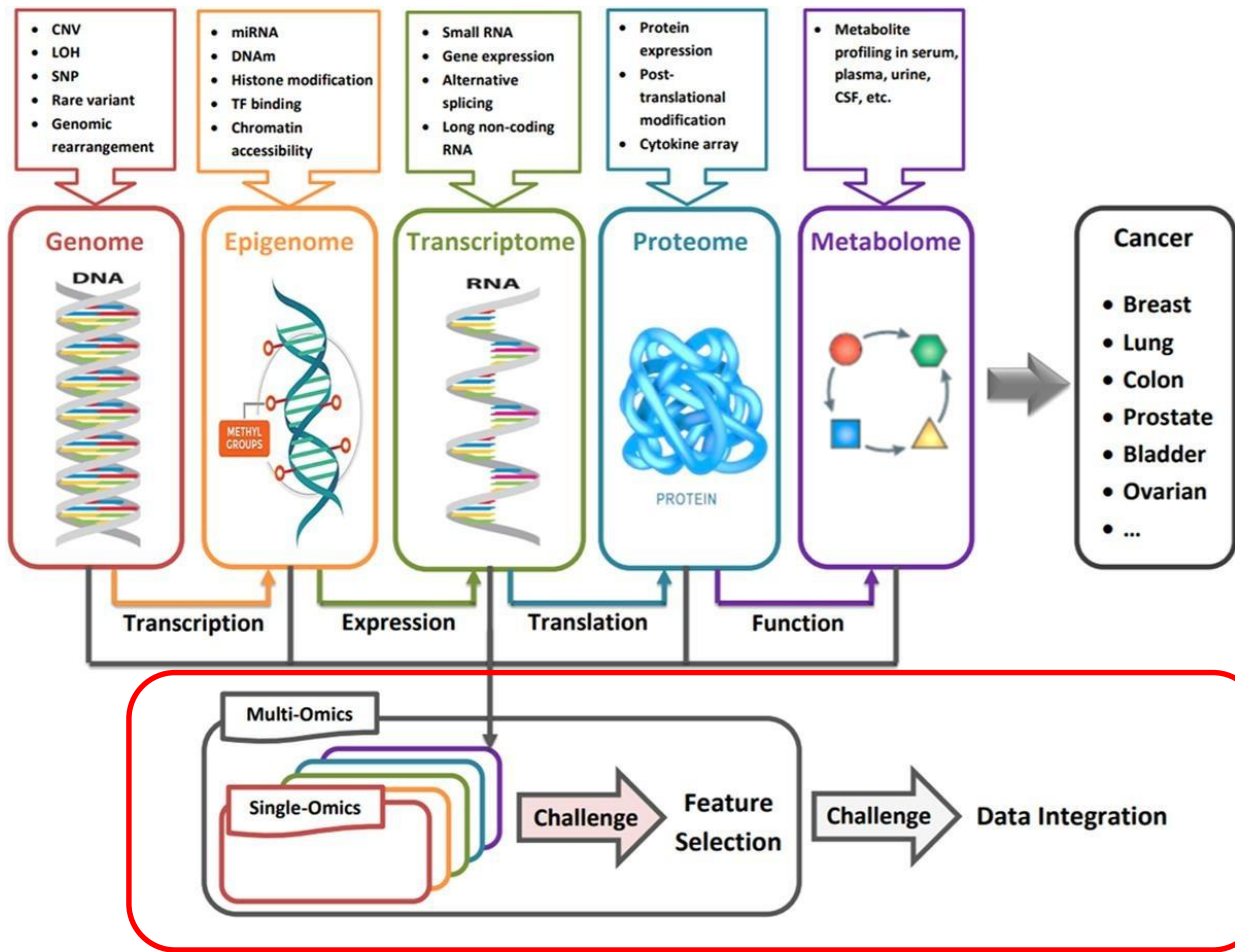
动物模型



整合型多组学分析

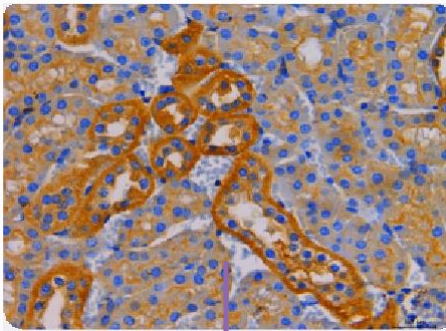


多组学交互研究



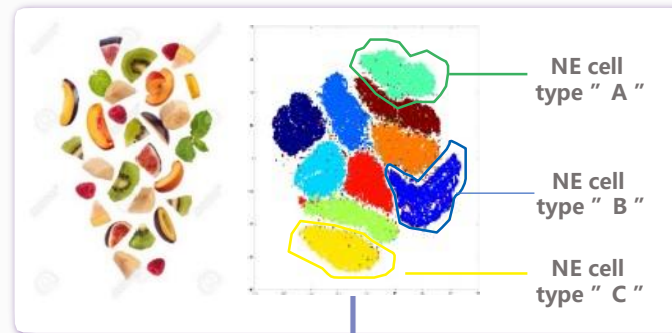
空间生物学的革命正在发生

探究微环境中组织结构、细胞亚型分布、特殊基因表达的重要性



TCGA 和bulk组织
NGS 分析
2006 年至今

组织形态及功能
传统方式



单细胞测序, 细胞分群
2012 年至今

组学结果如何与组织原位空间
信息对应起来进行验证?



数字化空间分析(DSP)技术的出现

nature biotechnology

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Letter | [Published: 11 May 2020](#)

Multiplex digital spatial profiling of proteins and RNA in fixed tissue

[Christopher R. Merritt](#) , [Giang T. Ong](#), [Sarah E. Church](#), [Kristi Barker](#), [Patrick Danaher](#), [Gary Geiss](#), [Margaret Hoang](#), [Jaemyeong Jung](#), [Yan Liang](#), [Jill McKay-Fleisch](#), [Karen Nguyen](#), [Zach Norgaard](#), [Kristina Sorg](#), [Isaac Sprague](#), [Charles Warren](#), [Sarah Warren](#), [Philippa J. Webster](#), [Zoey Zhou](#), [Daniel R. Zollinger](#), [Dwayne L. Dunaway](#), [Gordon B. Mills](#) & [Joseph M. Beechem](#) 

Nature Biotechnology **38**, 586–599 (2020) | [Cite this article](#)

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Abstract

Digital Spatial Profiling (DSP) is a method for highly multiplex spatial profiling of proteins or RNAs suitable for use on formalin-fixed, paraffin-embedded (FFPE) samples. The approach relies on (1) multiplexed readout of proteins or RNAs using oligonucleotide tags; (2) oligonucleotide tags attached to affinity reagents (antibodies or RNA probes) through a photocleavable (PC) linker; and (3) photocleaving light projected onto the tissue sample to release PC oligonucleotides in any spatial pattern across a region of interest (ROI) covering 1 to ~5,000 cells. DSP is capable of single-cell sensitivity within an ROI using the antibody readout, with RNA detection feasible down to ~600 individual mRNA transcripts. We show spatial profiling of up to 44 proteins and 96 genes (928 RNA probes) in lymphoid, colorectal tumor and autoimmune tissues by using the nCounter system and 1,412 genes (4,998 RNA probes) by using next-generation sequencing (NGS). DSP may be used to profile not only proteins and RNAs in biobanked samples but also immune markers in patient samples, with potential prognostic and predictive potential for clinical decision-making.



高维度分析与空间表达信息的解决方案

Gordon B. Mills



Dr. Joseph Beechem



Digital Spatial Profiler (DSP) 是一个全新的技术平台, Dr. Mills 是该项技术的共同发明人之一。它将传统的免疫荧光技术与数字可视化编码技术相结合, 因此不仅能获得清晰的空间信息, 而且可以做到多重蛋白/RNA 靶标的检测。目前在单个样本中的检测上限能够达到 150 种蛋白 / 18000 种 RNA, 很好的将空间信息及其他多组学数据结合起来, 极大的帮助科研工作者来解读 肿瘤异质性及其复杂的微环境。



Implementation of a Multiplex and Quantitative Proteomics Platform for Assessing Protein Lysates Using DNA-Barcoded Antibodies[†]

Jinho Lee¹, Gary K. Geiss¹, Gokhan Demirkan¹, Christopher P. Vellano¹, Brian Filanoski¹, Yiling Lu¹, Zheslin Ju¹, Shuangxing Yui¹, Huihang Guo¹, Lisa Y. Bogatzki¹, Warren Carter¹, Rhonda K. Meredith¹, Savitri Krishnamurthy¹, Zhiyong Dong¹, Joseph M. Beechem¹, and Gordon B. Mills¹



Multiplex digital spatial profiling of proteins and RNA in fixed tissue

Christopher B. Moritt¹, Giang T. Ong¹, Sarah E. Church¹, Kristi Barker¹, Patrick Danaher¹, Gary Geiss¹, Margaret Hoang¹, Jaemyeong Jung¹, Yan Liang¹, Jill McKay-Fleisch¹, Karen Nguyen¹, Zach Norgaard¹, Kristina Sorg¹, Isaac Sprague¹, Charles Warren¹, Sarah Warren¹, Philippe J. Webster¹, Zoey Zhou¹, Daniel R. Zollinger¹, Dwayne L. Dunaway¹, Gordon B. Mills¹ and Joseph M. Beechem^{1,2}

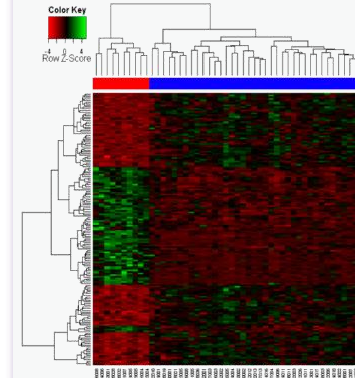


解决现有技术存在问题, 可进行RNA - 蛋白共分析

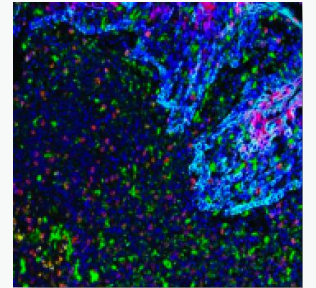
高维度表达谱分析

OR

低维度图像分析



- + 高维度 -
- 空间性 +
- + 定量性 -
- + 准确性 -



GEP:
Multi-plex PCR (bulk),
RNAseq (bulk),
Single-cell RNAseq,
PEP:
Flow Cytometry,
Mass cytometry,
DSP

IHC,
Immunofluorescence,
FISH

- ▶ 相对荧光强度定义的表达量, 各个荧光基团的激发强弱差异
- ▶ 定量存在偏倚
- ▶ 自荧光

- ▶ 荧光信号交叠
- ▶ 多重靶标检测能力限制在30-40重 (实操)



高维度分析与空间表达信息的解决方案

Gordon B. Mills



Dr. Joseph Beechem



Digital Spatial Profiler (DSP) 是一个全新的技术平台, Dr. Mills 是该项技术的共同发明人之一。它将传统的免疫荧光技术与数字可视化编码技术相结合, 因此不仅能获得清晰的空间信息, 而且可以做到多重蛋白/RNA靶标的检测。目前在单个样本中的检测上限能够达到 150 种蛋白 / 18000 种 RNA, 很好的将空间信息及其他多组学数据结合起来, 极大的帮助科研工作者来解读 肿瘤异质性及其复杂的微环境。

适用于FFPE样本和新鲜样本



样本制备简便容易



兼顾形态学与原位信息对应关系



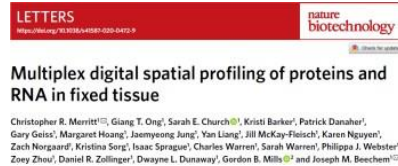
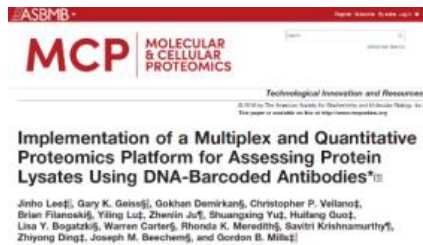
适用于大样本空间组分析 (TMA)



样本覆盖区域大, 不同组织区域或结构异质性信息更为全面丰富



可同时分析蛋白和RNA, 可实现原位全转录组高通量分析



蛋白组和空间组为核心的多组学研究策略

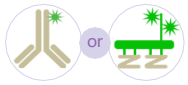


DSP技术原理及流程

- 石蜡样本/冰冻切片/穿刺样本等多样本、珍贵样本策略
- 前处理简单, 无需优化, 视野区域广阔, 无需盲选
- 靶向检测, 对中低表达靶点友好
- 自动化程度高, 结果可重复性强

特定标记荧光抗体(或探针)与所选待测指标共孵育

形态学标记荧光抗体或RNA探针



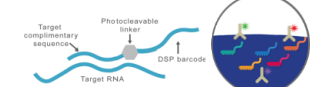
蛋白检测

待测靶点抗体 + 特定标记荧光抗体(或探针) 共孵育

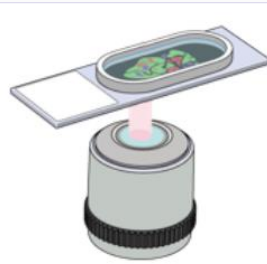
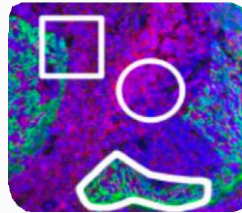


RNA 检测

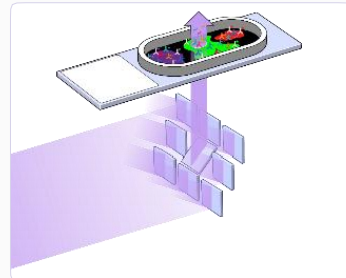
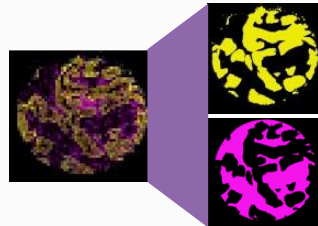
探针孵育 + 特定标记荧光抗体(或探针) 共孵育



形态学标记荧光成像 ROI (感兴趣区域) 圈选

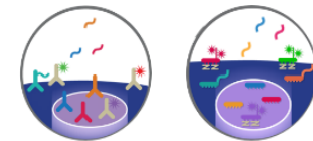


区域划分 (Segment, 可选) 紫外光解 DSP barcode

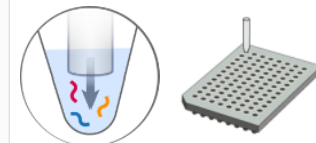


逐个 ROI 收集 DSP barcode 到 96 孔收集板

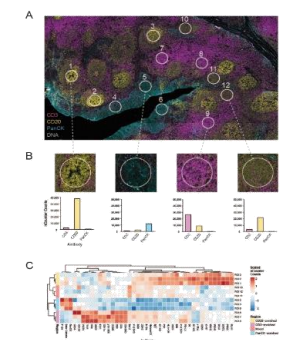
紫外光照射, linker断裂, DSP barcode解离至上层缓冲液



毛细管将解离的DSP barcode收集至96孔板



计数、测序和数据分析



DSP技术的应用场景

运用 DSP 作为探索性工具

CLINICAL CANCER RESEARCH

High-Plex Predictive Marker Discovery for Melanoma Immunotherapy-Treated Patients Using Digital Spatial Profiling

CLINICAL CANCER RESEARCH

Biomarkers Associated with Beneficial PD-1 Checkpoint Blockade in Non-Small Cell Lung Cancer (NSCLC) Identified Using High-Plex Digital Spatial Profiling

nature cancer

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ESMO

High multiplex analysis of the immune microenvironment in bone marrow trephine samples using GeoMx™ digital spatial profiling

cancers

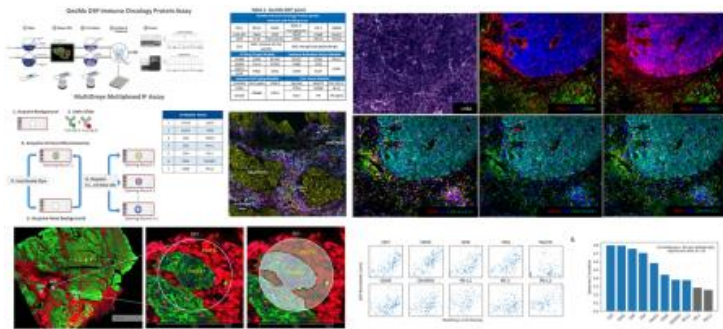
High-Plex and High-Throughput Digital Spatial Profiling of Non-Small-Cell Lung Cancer (NSCLC)

James Monkman, Tomraj Tabiri, Majid Ebrahimi Warkiani, Connor O'Leary, Rabal Ludwa, Derek Richard, Ken O'Byrne, and Aratha Kalasinghe

IOTECH

High multiplex analysis of the immune microenvironment in bone marrow trephine samples using GeoMx™ digital spatial profiling

DSP 与单细胞空间组联合分析



运用 DSP 与高通量多组学数据进行整合

Journal for Immunotherapy of Cancer

Distinct tumor microenvironments of lytic and blastic bone metastases in prostate cancer patients

Qian L. He, Meredith D. Prosser, Christine M. Seignin, E. Din Smith, Susan M. Edgermon, Adrie Van Balkhoven, M. Scott Lucia, and Philip Ouyang

Immunologic alterations in the pancreatic cancer microenvironment of patients treated with neoadjuvant chemotherapy and radiotherapy

Matthew R. Farnen, Layal Sayegh, Michael Brandon Ware, Hsiao-Rong Chen, Jingling Gong, Yan Liang, Alyssa Kravitskaya, Shichik K. Mutha, Mohammad Zaidi, Juan M. Samir, David Gaddy, Patrick Patel, Bassem El-Rayes, Walid Ibrahim, and Gregory B. Leshem

Journal for Immunotherapy of Cancer

Inter- and intra-tumor heterogeneity of metastatic prostate cancer determined by digital spatial gene expression profiling

Lauren Brady, Michelle Kriner, Isaac Coleman, Cole Morrissey, Martine Roudier, Lawrence D. True, Roman Gulati, Stephen R. Plymate, Zoey Zhou, Brian Biditt, Rhonda Meredith, Gary Geiss, Margaret Hoang, Joseph Beechem, and Peter S. Nelson

High dose-rate brachytherapy of localized prostate cancer converts tumors from cold to hot

Simon P. Kaam, Heiwoo Hahn, Thi Nguyen, Minyu Wang, Nicholas Van Kesteren, Catherine Mitchell, Franco Carraia, David J Byrne, Sue Haupt, Georgina Flynn, Phillip K Darcy, Shaheen Sandhu, Piers Blombery, Ygal Haupt, Scott G Williams, Paul J Neeson

运用 DSP 与单细胞测序交叉验证

nature medicine

Opposing immune and genetic mechanisms shape oncogenic programs in synovial sarcoma

Livnat Jerby-Aron, Cyril Nefzi, Marri E. Shon, Hannah R. Weisman, Nathan D. Mathewson, Matthew J. McBride, Brian Hays, Benjamin Izan, Angela Volker, Gayle Boulay, Luisa Ghent, Alyssa R. Richman, Liliane C. Breyer

Onco-fetal Reprogramming of Endothelial Cells Drives Immunosuppressive Macrophages in Hepatocellular Carcinoma

Ankur Sharma, Justine Jia Wen Seow

Cell

Spatially organized multicellular immune hubs in human colorectal cancer

scRNA-seq and spatial profiling of CRC patients

Tertiary lymphoid structures improve immunotherapy and survival in melanoma

Wito Cabrita, Martin Lasser, Adriana Serrao, Marco Donati, Mathilde Stangor, Shreshth Mittal, Ivo Johansson, Bengt Philip, Kristina Herberich, John Wilton Christensen, Alben van Schrick, Katharina Langner, Sarah Warner, Karim Alnaitan, Mikael Olsson, Kristian Preker, Christian Ingemar, Karoline Sultmann, Erik Schindler, Henrik Schmalz, Lars Bastholt, Ana Carmona, Jennifer A. Wargo, Inga Marie Svane, and Ola Jonsson



高维度分析与空间表达信息的解决方案

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Letter | [Published: 11 May 2020](#)

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Abstract

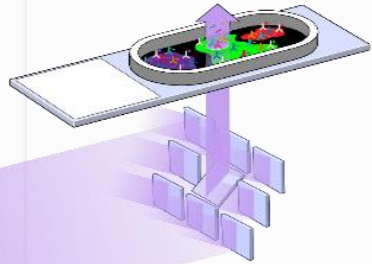
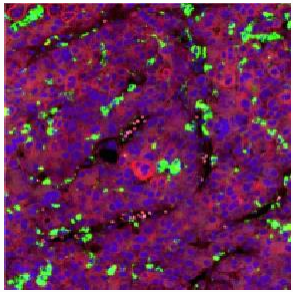
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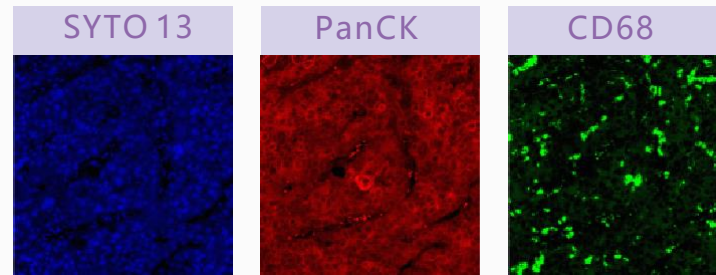
特殊光学技术助力灵活的ROI 选择和Masking 覆盖

细胞亚群纯化策略

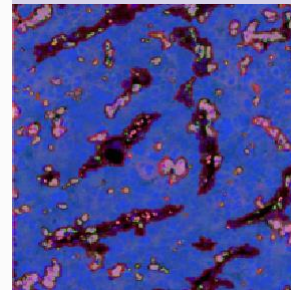
Full Channel Overlay



Single Channel (Mask)

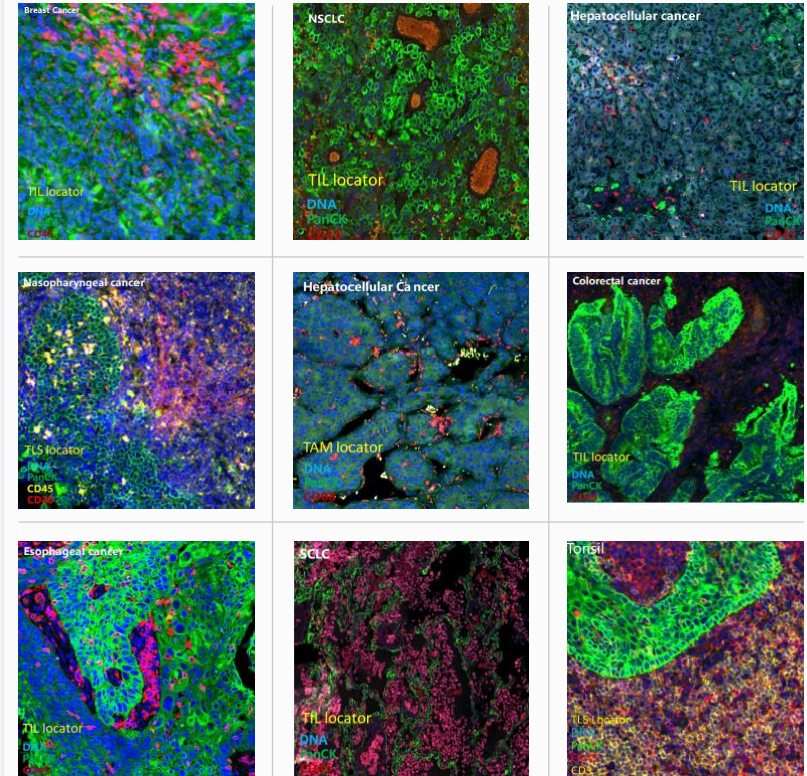


Segmentation



细胞亚群及特殊组织结构定位

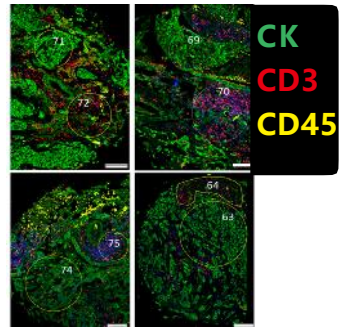
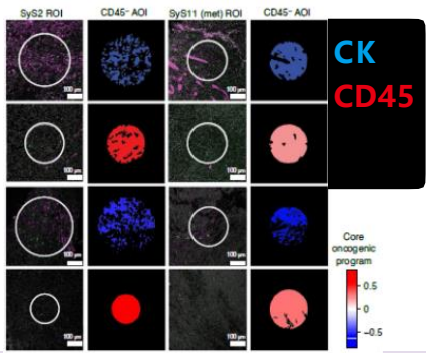
(TLS/TIL/TAM)



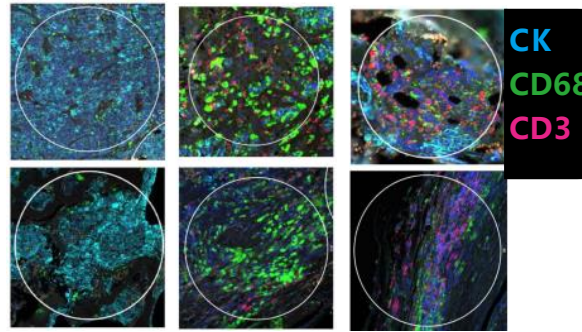
DSP染色解决方案

- 合适的亚组织结构或细胞群标志物，有助于ROI的选择和后续分析。

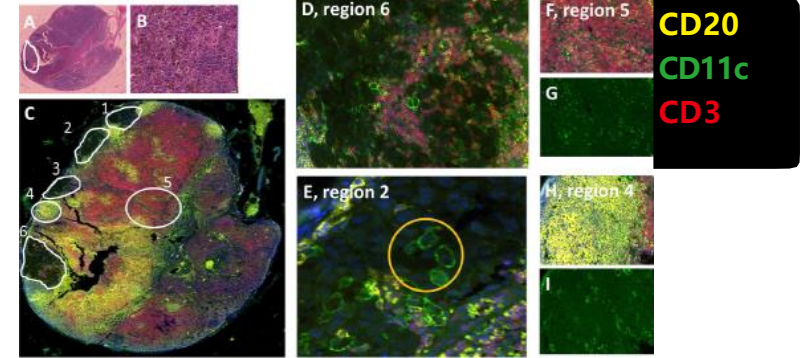
肿瘤及肿瘤浸润免疫细胞(白细胞, T细胞)



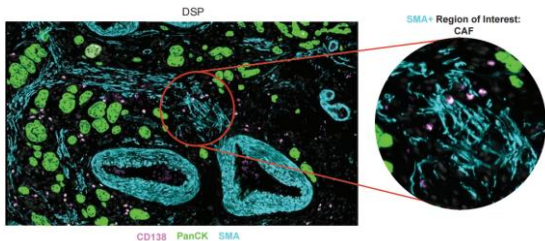
肿瘤及肿瘤浸润免疫细胞(T细胞,巨噬细胞)



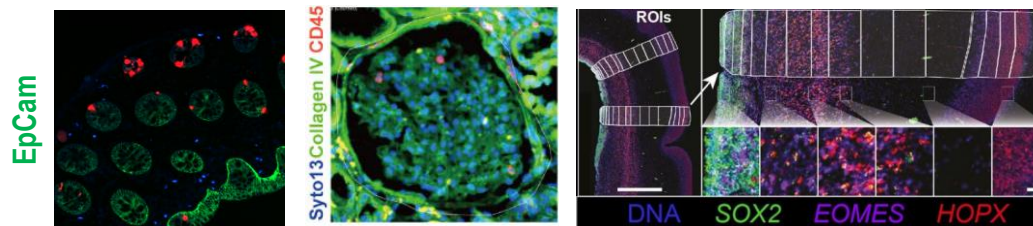
肿瘤浸润免疫细胞(T细胞, B细胞, 髓系细胞)



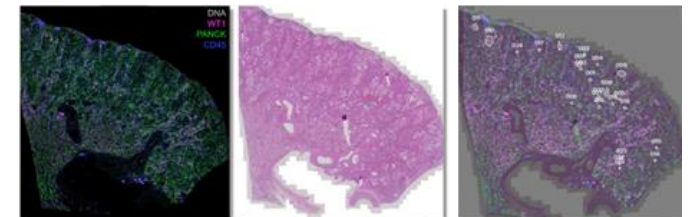
肿瘤相关基质细胞(成纤维类)



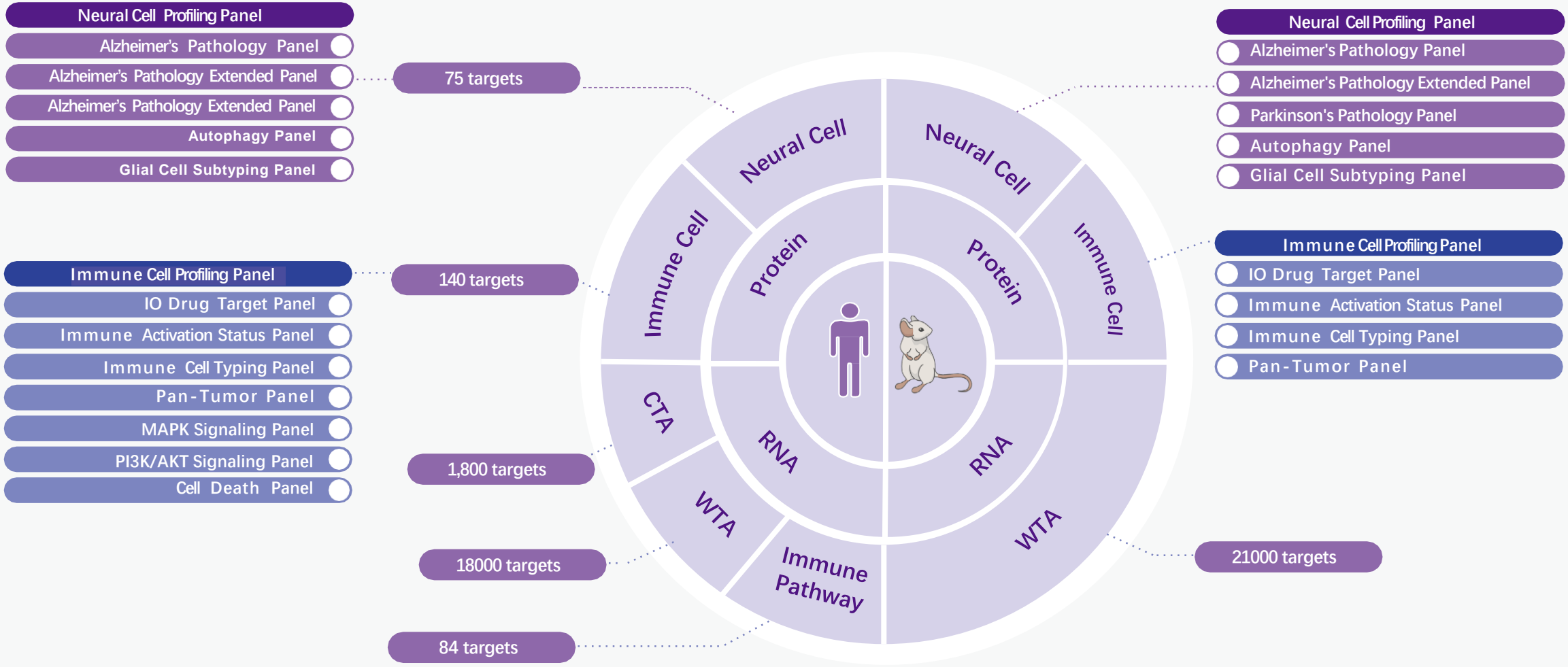
其他特殊标志物染色



通过HE确定待分析结构



DSP检测panel 设置 (蛋白+RNA)



DSP在肿瘤微环境相关研究中的切入点

运用 DSP 作为探索性工具



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High-Plex Predictive Marker Discovery for Melanoma Immunotherapy-Treated Patients Using Digital Spatial Profiling



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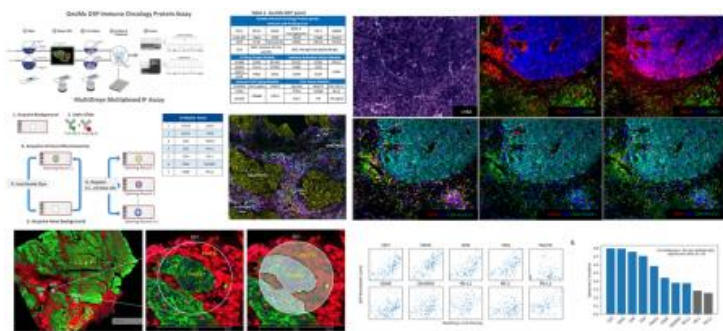


ESMO
High multiplex analysis of the immune microenvironment in bone marrow trephine samples using GeoMx™ digital spatial profiling

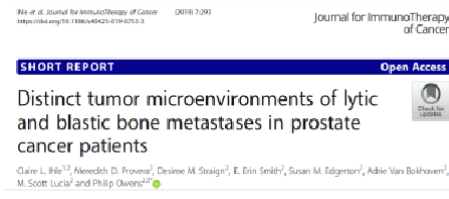


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DSP 与单细胞空间组联合分析



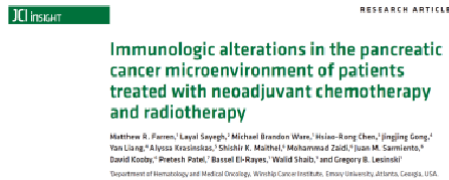
运用 DSP 与高通量多组学数据进行整合



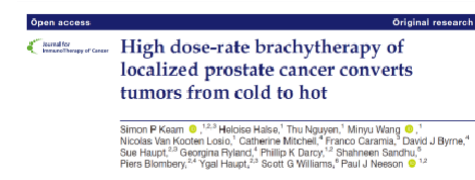
Journal for Immunotherapy of Cancer
Distinct tumor microenvironments of lytic and blastic bone metastases in prostate cancer patients



ARTICLE
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JCI INSIGHT
Immunologic alterations in the pancreatic cancer microenvironment of patients treated with neoadjuvant chemotherapy and radiotherapy

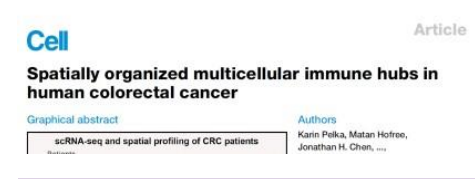


Open access
High dose-rate brachytherapy of localized prostate cancer converts tumors from cold to hot

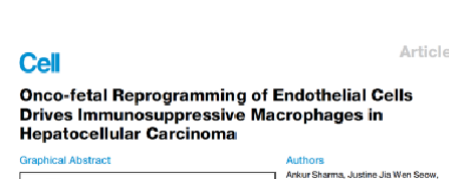
运用 DSP 与单细胞测序交叉验证



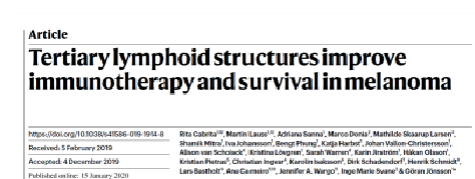
nature medicine
Opposing immune and genetic mechanisms shape oncogenic programs in synovial sarcoma



Cell
Spatially organized multicellular immune hubs in human colorectal cancer



Cell
Onco-fetal Reprogramming of Endothelial Cells Drives Immunosuppressive Macrophages in Hepatocellular Carcinoma



Article
Tertiary lymphoid structures improve immunotherapy and survival in melanoma

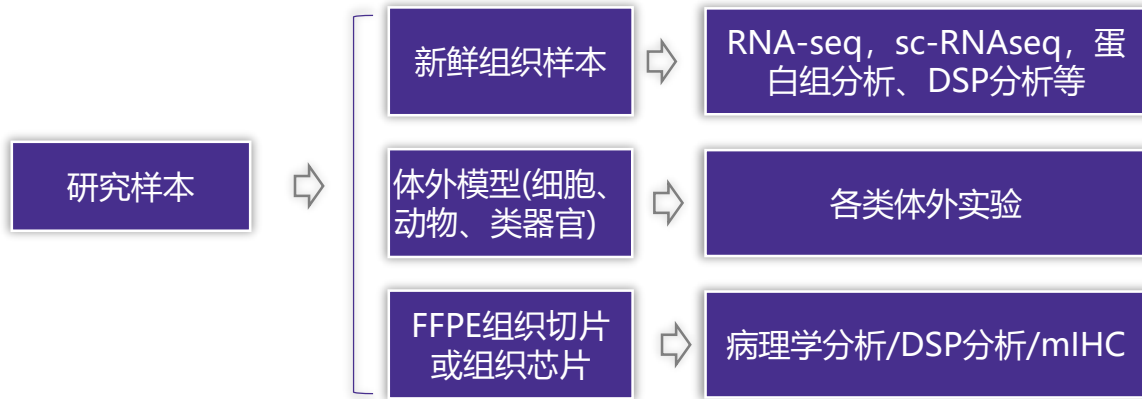


本次系列讲座探讨的核心内容

- 如何合理设计研究方案并最大化利用样本，开展空间多组学研究？
- 如何对多组学技术平台获得的数据进行集成分析，并有效进行验证？



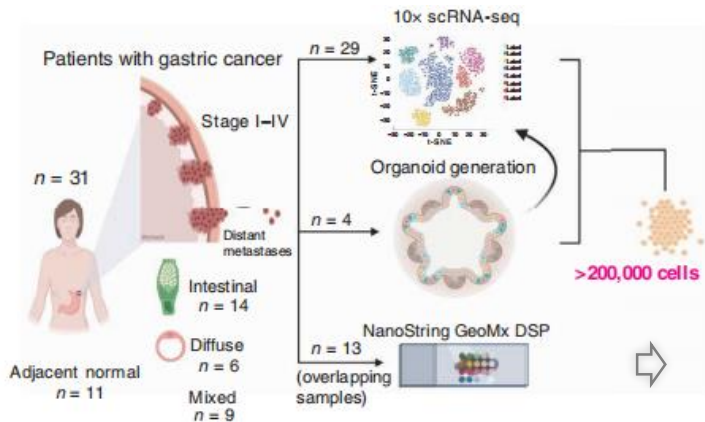
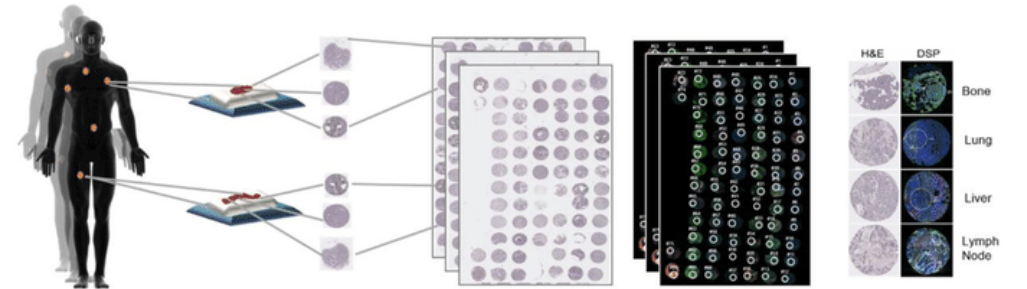
DSP空间组学的样本规划



普通组织切片

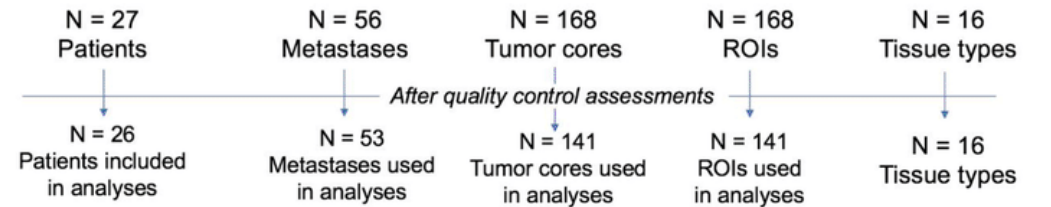
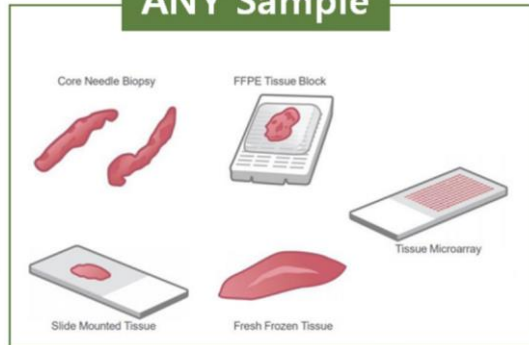


组织芯片(TMA)



满足DSP分析的样品类型

ANY Sample



DSP空间组学的样本规划

样本收集 (穿刺/手术)

样本收集(组织芯片)

瘤内异质性

Clinical Cancer Res 2020 (软骨肉瘤)

肿瘤核心/病灶异质性

Nat Comm 2020 (前列腺癌---常规转录组/基因组联用)

原发转移灶异质性

Journal of Immunotherapy of Cancer 2022 (肉瘤)

肿瘤免疫微环境

Clinical Cancer Res 2020 (肺癌生物标志物挖掘)

外周基质微环境

Journal of Immunotherapy of Cancer 2020 (前列腺癌骨转移)

外周基质微环境

Cell Rep 2021 (神经母细胞瘤相关基质微环境)

特定细胞亚群

Cancer Cell 2021 (胰腺癌)

特定细胞亚群

Clinical Cancer Res 2019 (黑素瘤-巨噬细胞/免疫基质)

特定病理学形态学结构

Nature 2019 (黑素瘤相关三级淋巴结构)

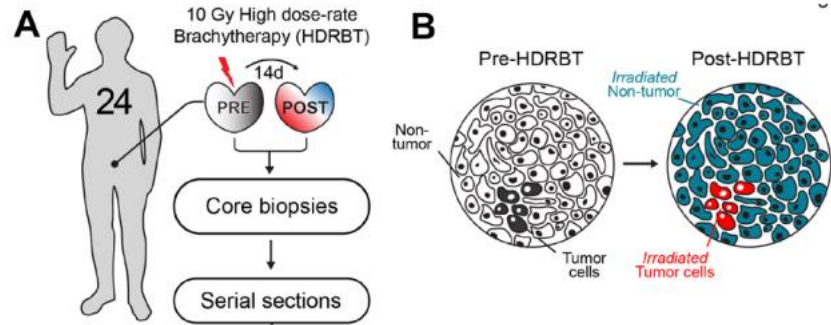
特定病理学特征/形态学结构

Nature 2019 (黑素瘤相关三级淋巴结构)

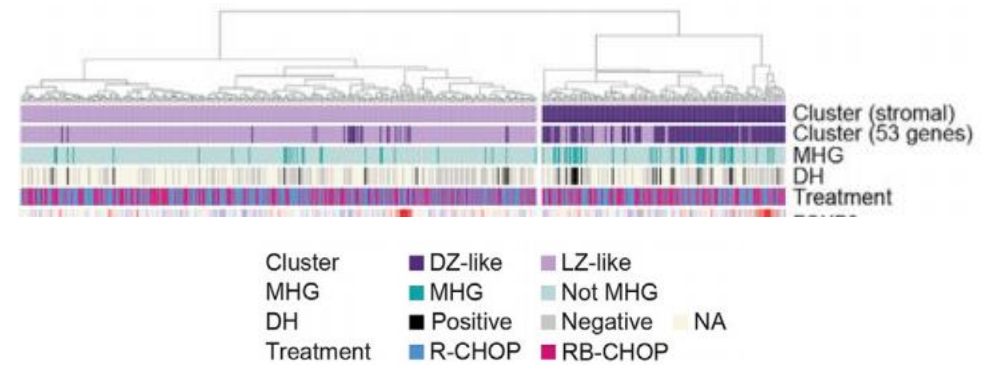


DSP空间组学的样本分组

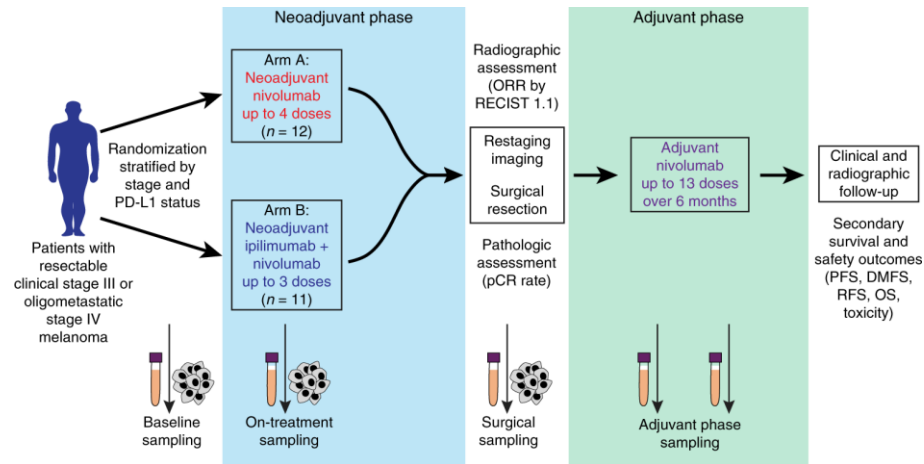
样本分组(治疗前与治疗后)



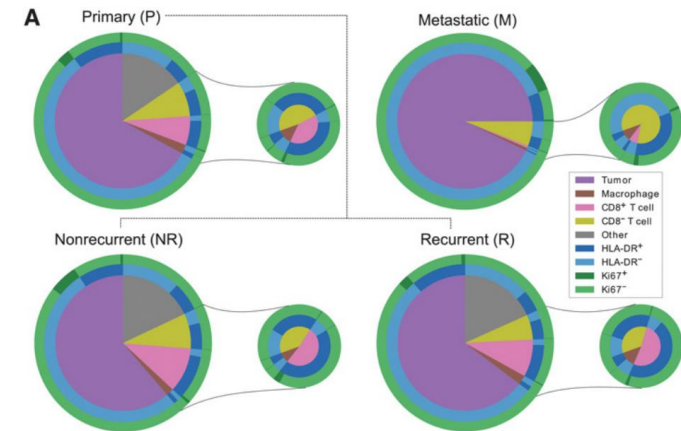
样本分组(疾病亚型与治疗分组)



样本分组(治疗进程取样)



样本分组(疾病发展过程)



DSP空间组学的样本分组

样本分组(病理分型和分子分型)

Table 1 Summary of patient and tumor characteristics

| Patient | Tissue | Diagnosis | ER(%)/ PR(%)/ HER2(+) | PAM50 Subtype | TIS Status | Treatment at the time of sample collection |
|---------|-------------|------------------|-----------------------------|---------------|------------|--|
| 1 | Breast | IDC, DCIS | 2/0/3 | Luminal A | High | Gemcitabine and trastuzumab |
| 1 | Breast | Tumor emboli | N/A | Luminal A | High | Nab-paclitaxel and trastuzumab |
| 1 | Lung | Metastatic | 0/0/3 | HER2 enriched | High | Nab-paclitaxel and trastuzumab |
| 2 | Breast | ICD, DCIS | 0/0/3 | HER2 enriched | Low | None. Initial diagnosis |
| 2 | Breast | Residual IDC | 0/0/N/A | HER2 enriched | High | TDM1, pertuzumab, doxorubicin and cyclophosphamide* |
| 2 | Soft tissue | Local recurrence | 0/0/3 | HER2 enriched | Low | Trastuzumab and pertuzumab |
| 3 | Brain | Metastatic | 15/0/3 | HER2 enriched | Low | None. Discontinued due to toxicities |
| 4 | Breast | IDC | 0/0/3 | Basal like | High | None. Initial diagnosis |
| 4 | Breast | Local recurrence | N/A | Luminal A | High | Doxorubicin, cyclophosphamide, paclitaxel and trastuzumab |
| 5 | Brain | Metastatic | 0/0/3 | Basal like | Low | TDM1 and anastrozole |
| 6 | Breast | IDC, DCIS | 80/70/2** | Luminal A | High | None. Initial diagnosis |
| 7 | Breast | IDC, DCIS | 0/0/3 | HER2 enriched | High | None. Initial diagnosis |
| 7 | Breast | Local recurrence | N/A | Basal like | High | None. Completed neoadjuvant and adjuvant therapy |
| 8 | Breast | IDC, DCIS | 10/15/1 | Luminal A | Low | None. Initial diagnosis |
| 8 | Soft tissue | Local recurrence | 0/0/3 | Basal like | Low | Nab-paclitaxel and atezolizumab (previously had triple negative disease) |

DCIS: ductal carcinoma in situ; IDC: ductal carcinoma in situ; ER: estrogen receptor; HER2: human epidermal growth factor 2; IDC: invasive ductal carcinoma; IDC: invasive ductal carcinoma; PR: progesterone receptor; TDM1: ado-trastuzumab emtansine; TIS: tumor inflammation score

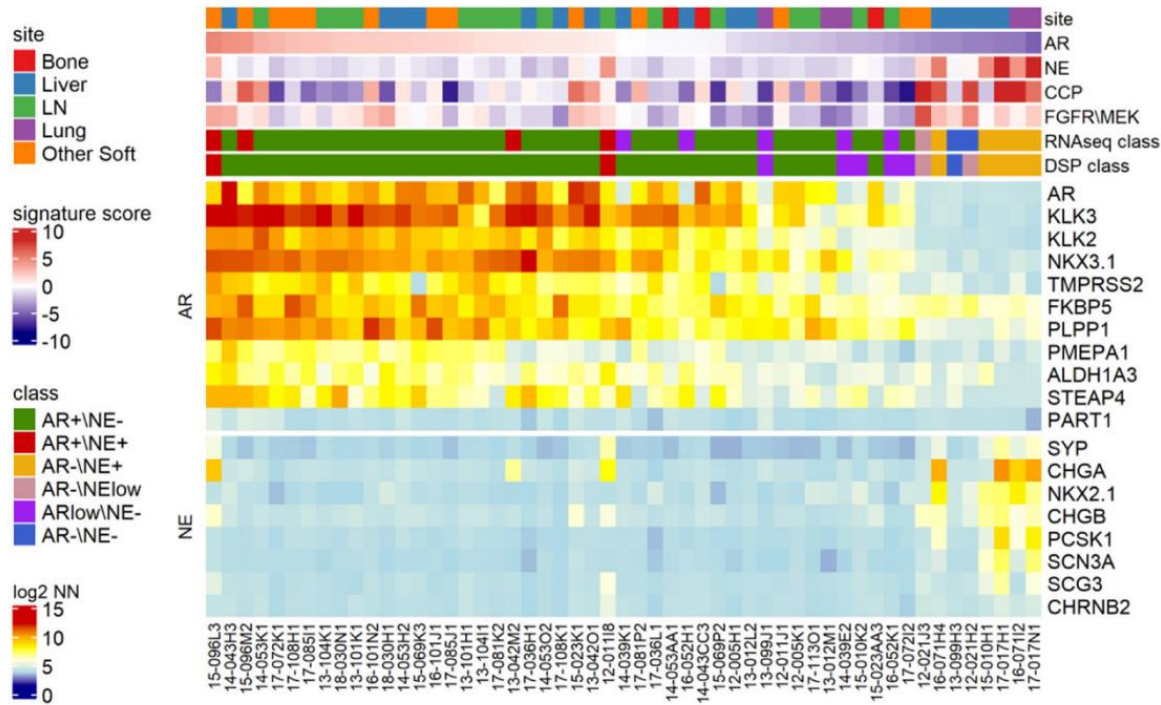
* Neoadjuvant therapy in the ISPY2 clinical trial

** HER2/CEP17 ratio: 2.13 (amplified by FISH)



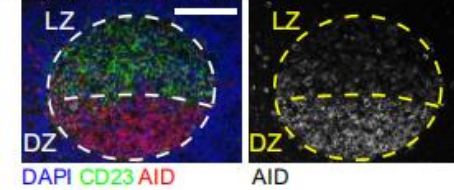
DSP空间组学的样本分组

样本分组(特征基因集分型)

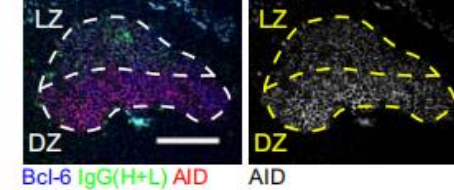


样本分组(病理组织结构结合特征基因集分型)

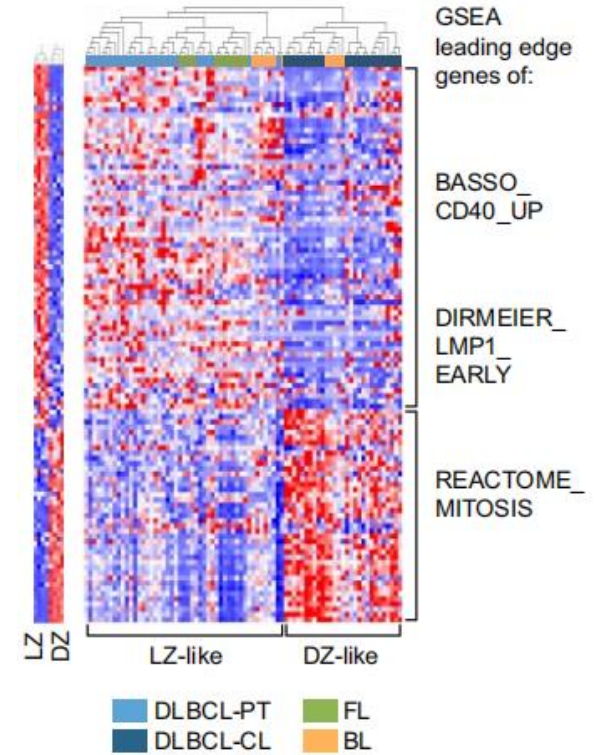
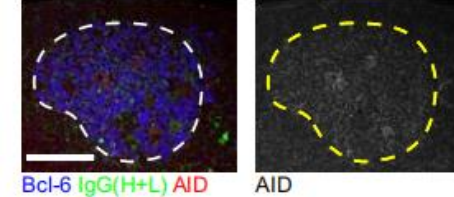
Human lymph node



Mouse spleen (wild type)

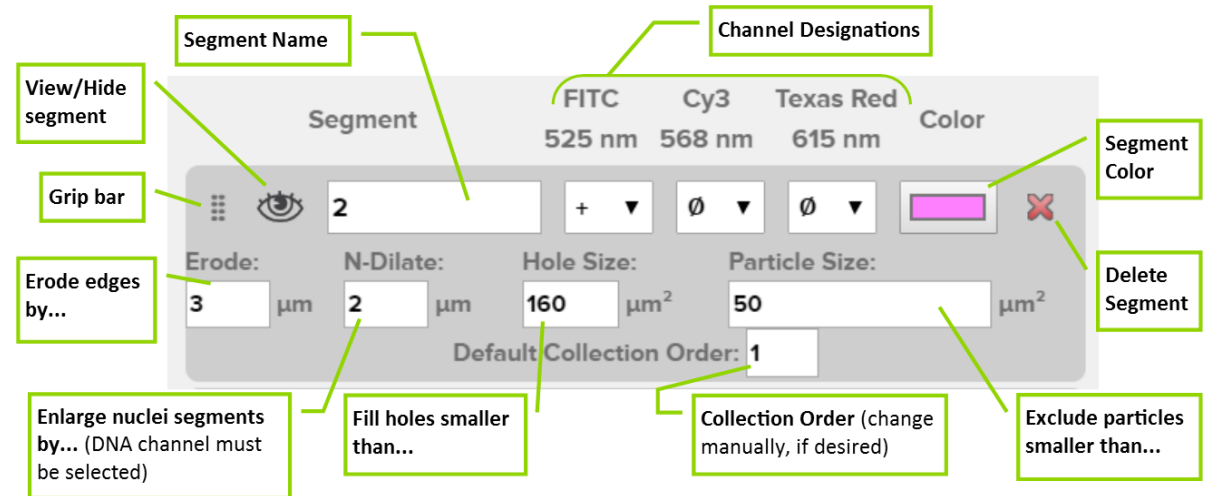
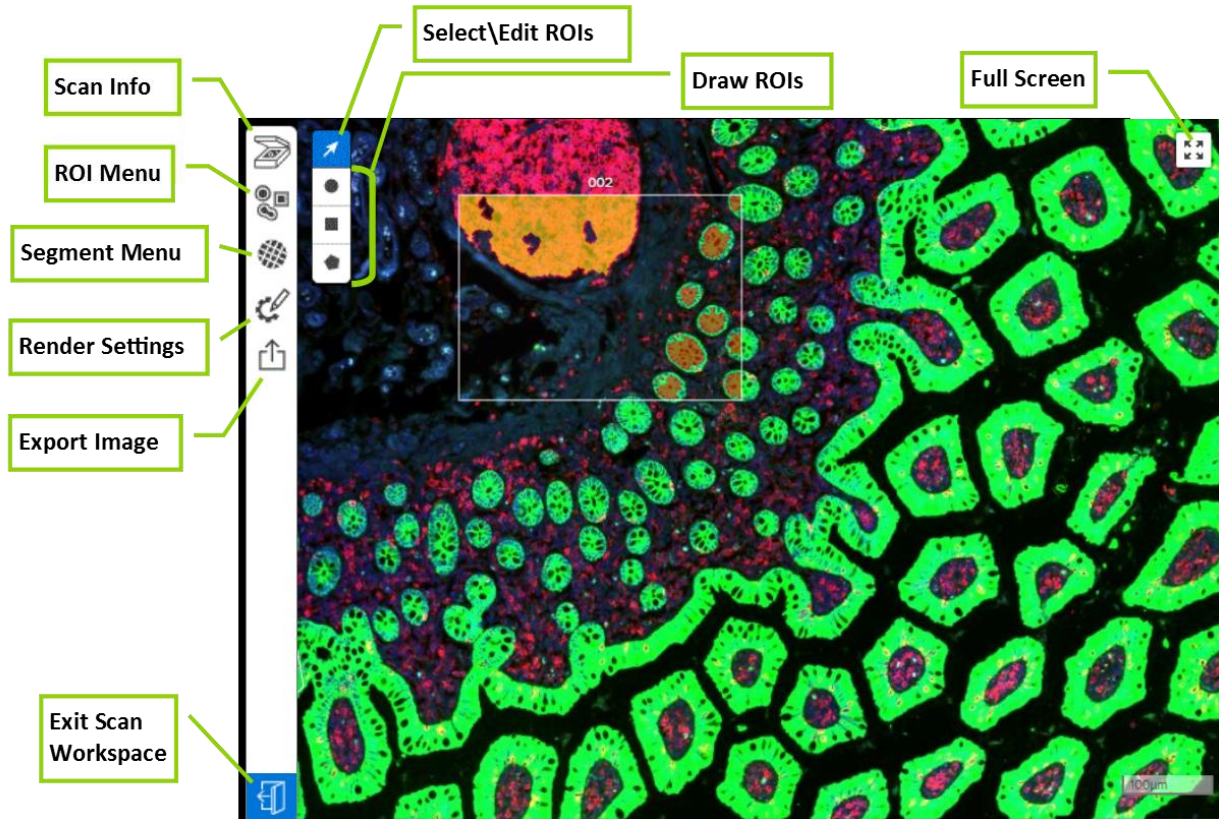


Mouse lymph node (*Aicda*⁺)



DSP空间组学的兴趣区域(ROI)选择

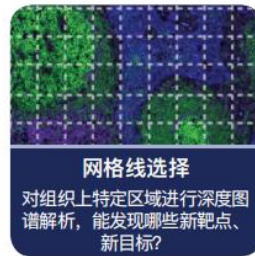
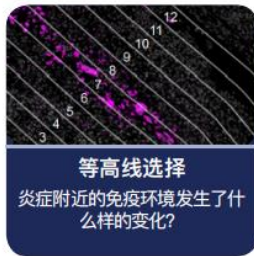
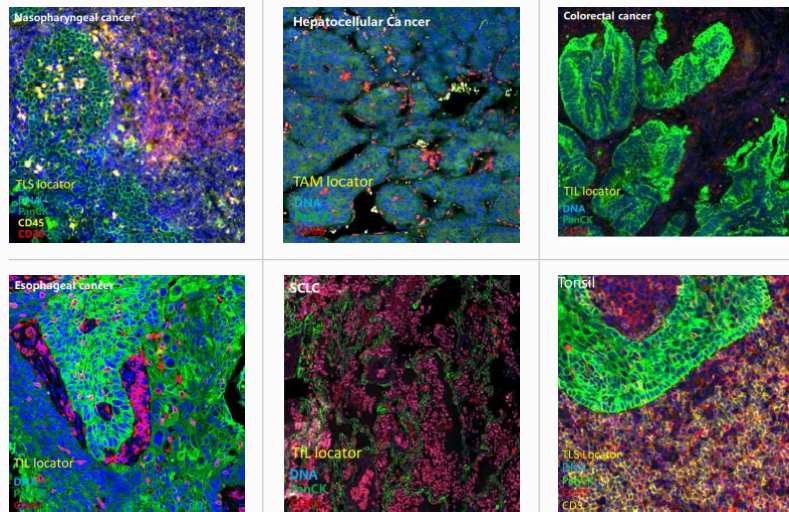
ROI的选择是DSP技术最大化应用的核心环节



DSP空间组学的兴趣区域(ROI)选择

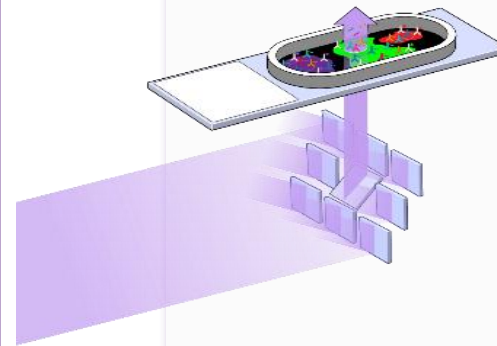
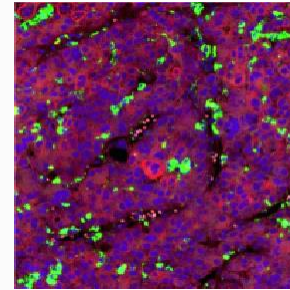
ROI的选择是DSP技术最大化应用的核心环节

特定组织结构区域选取策略

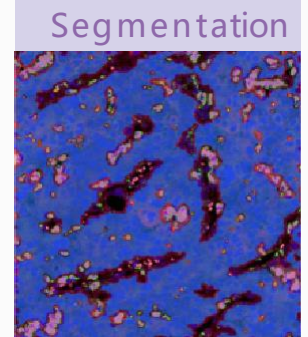
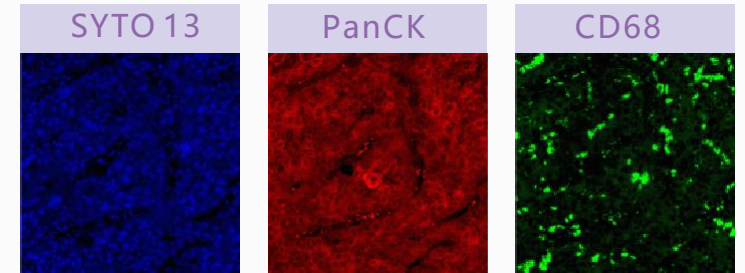


特定细胞群选取策略

Full Channel Overlay



Single Channel (Mask)



DSP空间组学的兴趣区域(ROI)选择

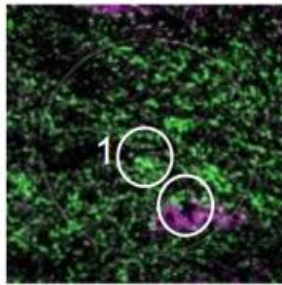
ROI的选择是DSP技术最大化应用的核心环节

Five Unique Fluorescent Profiling Modalities

Geometric



CD3 PanCK DNA

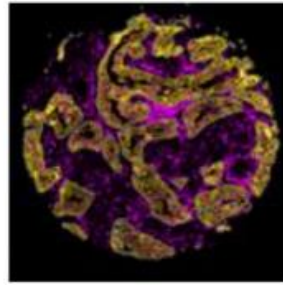


Assess heterogeneity
(e.g. tumor margin vs. core, cortical layers)

Segmentation

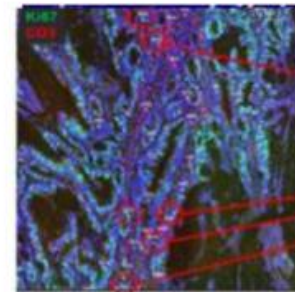


PanCK DNA



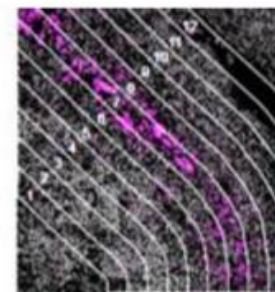
Evaluate biological compartments
(e.g. Tumor-TME)?

Cell Type/Rare cell



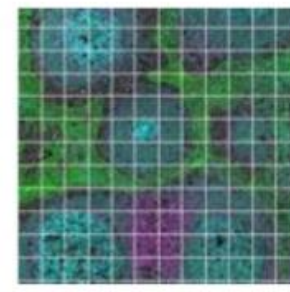
Profile cell populations
(e.g. CD3+ cells, GFAP+ cells)

Contour



Determine proximal response
(e.g. CART, AD plaque)

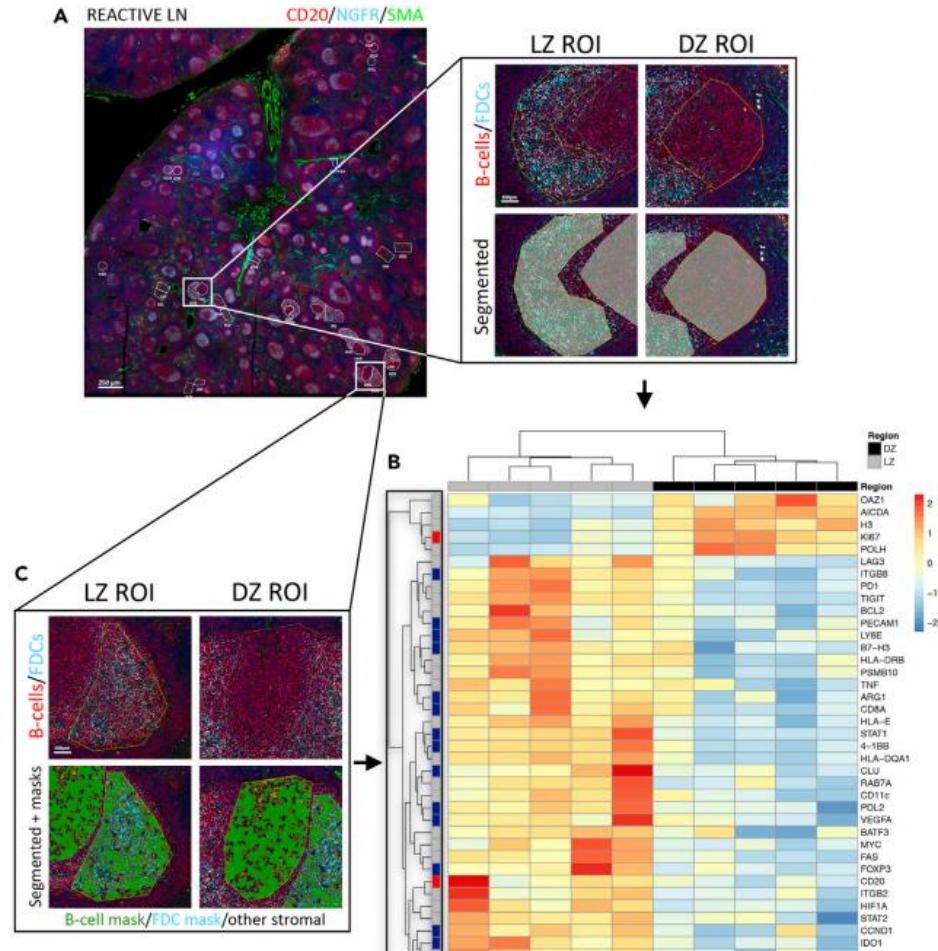
Gridded



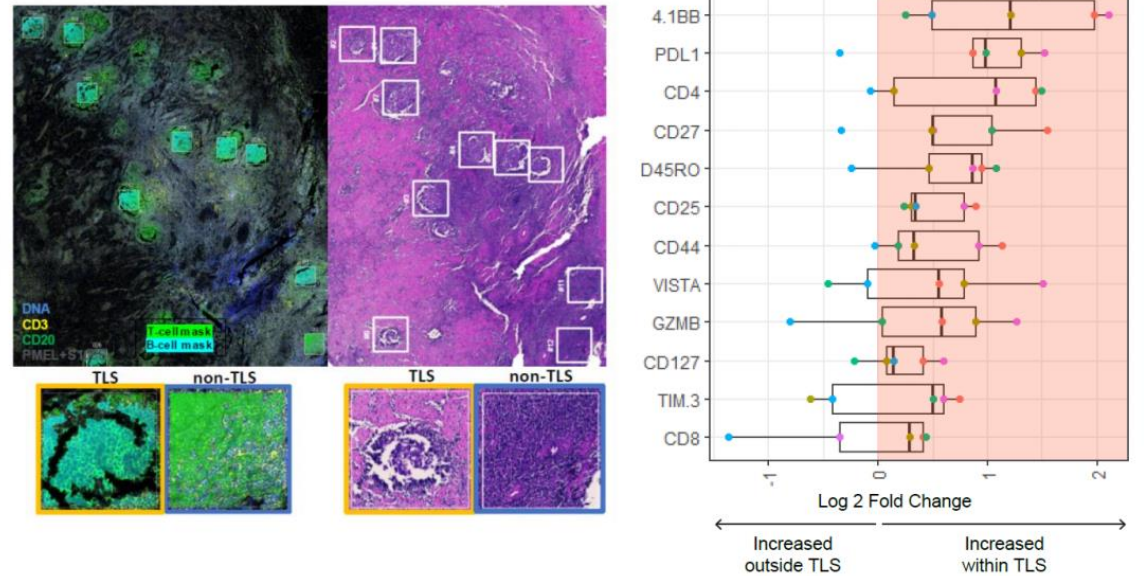
Perform deep mapping
(e.g. 96 ROI)



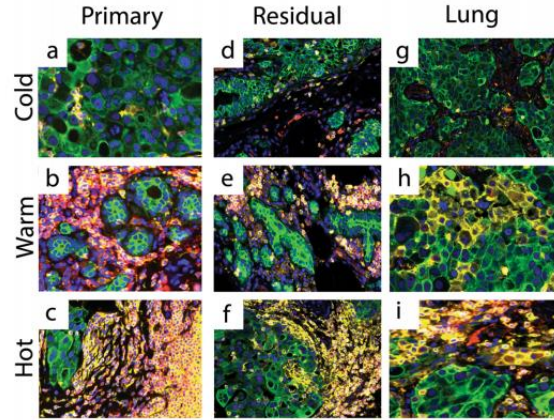
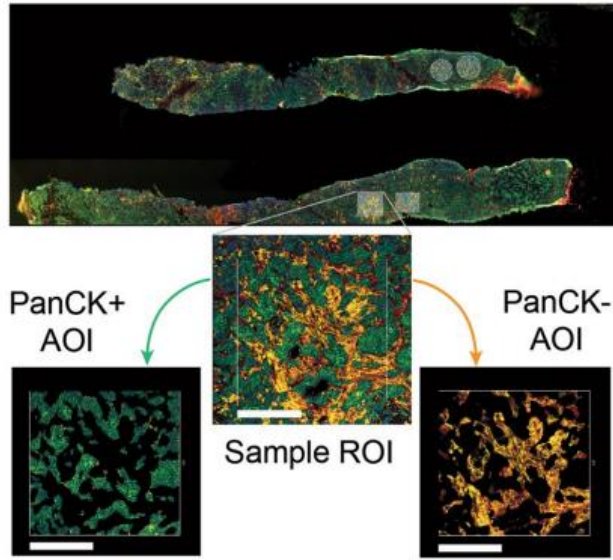
DSP空间组学的兴趣区域(ROI)选择



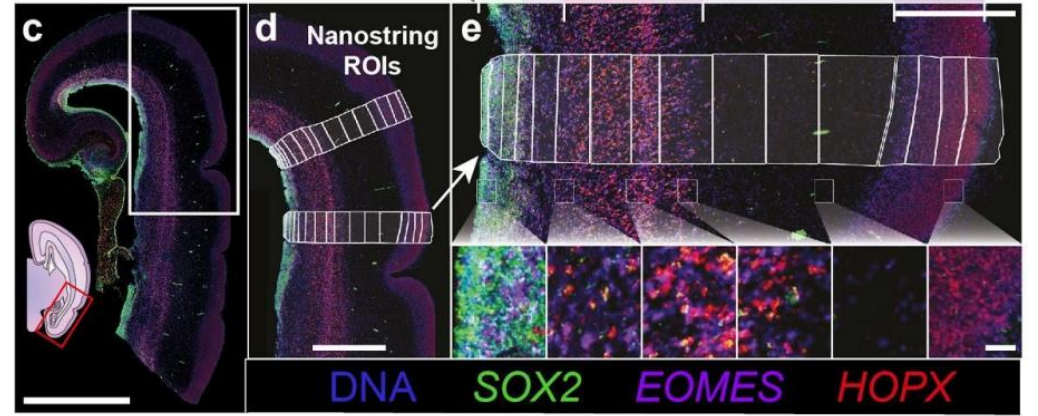
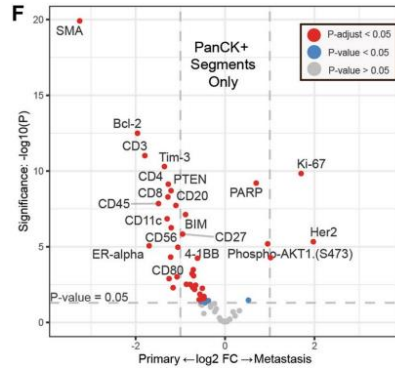
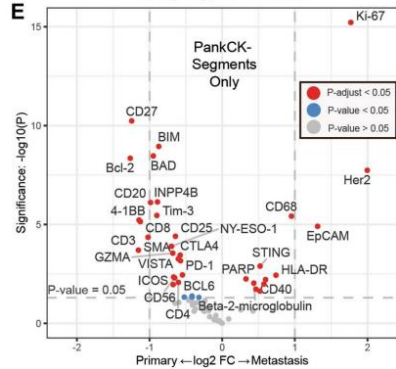
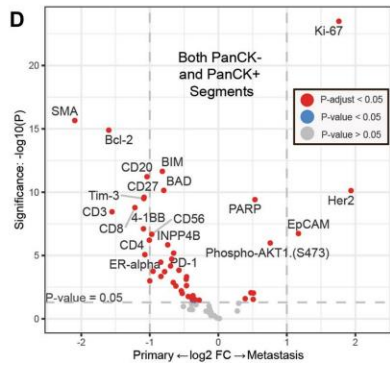
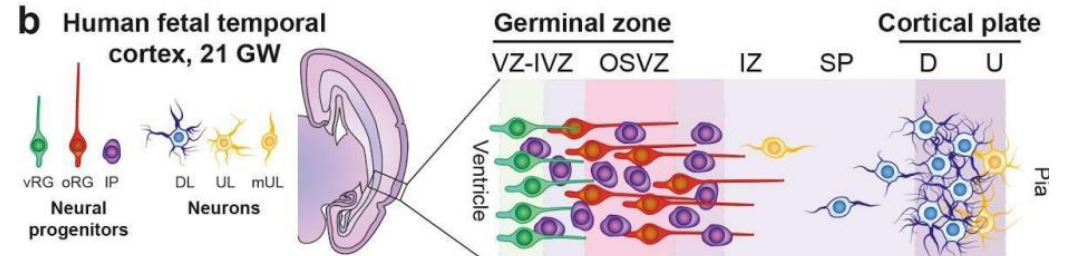
Spatially-distinct changes in immune cell activation



DSP空间组学的兴趣区域(ROI)选择

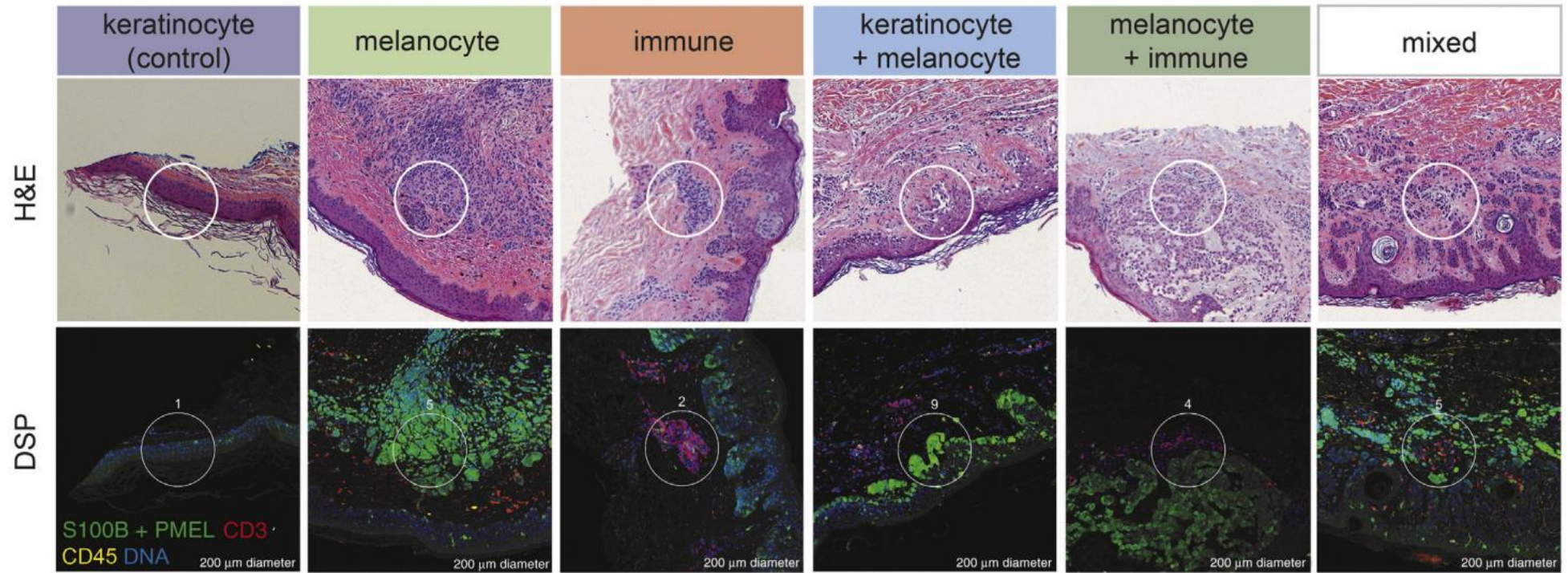


| Timepoint | Segment | PAM50 | ROI.Type | Tissue | TIS.Status |
|-----------|-----------|------------|----------|--------|------------|
| Primary 1 | PanCK neg | Basal like | Cold | Brain | High |
| Primary 2 | PanCK pos | HER2e | Warm | Breast | Low |
| Met | | Luminal A | Hot | Lung | |



DSP空间组学的兴趣区域(ROI)选择

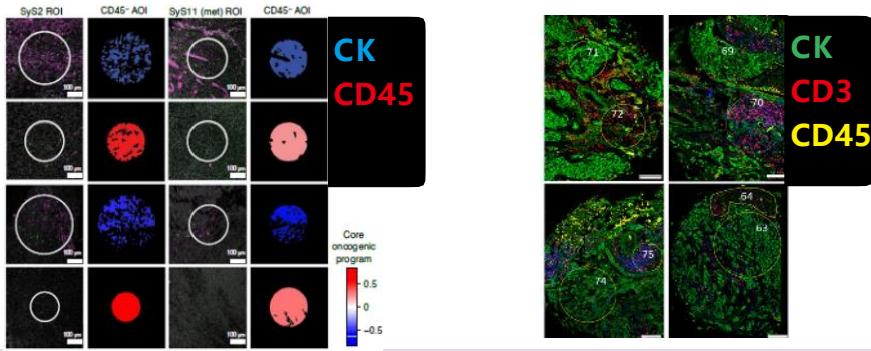
圈选ROI时，HE/IHC图片与DSP扫描的荧光图像互为参考，所选区域会更加准确。



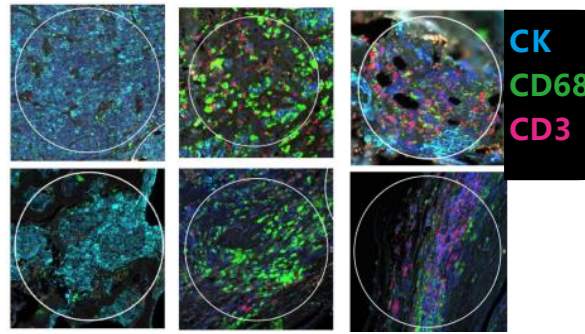
DSP空间组学分析相关细胞标志物选择

合适的组织结构或细胞群标志物，有助于ROI的选择和后续分析。

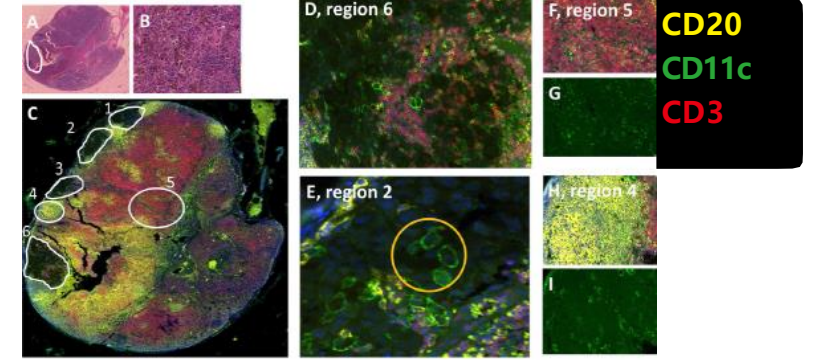
肿瘤及肿瘤浸润免疫细胞(白细胞, T细胞)



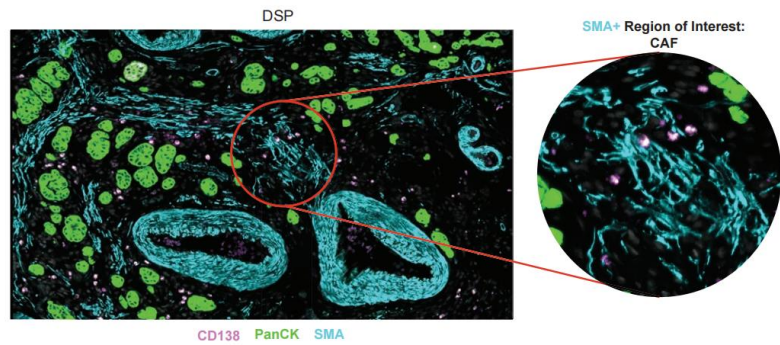
肿瘤及肿瘤浸润免疫细胞(T细胞,巨噬细胞)



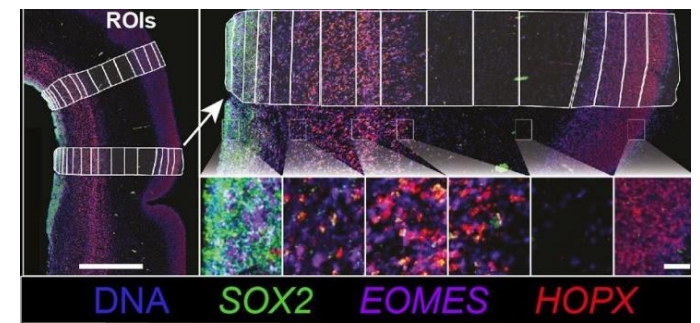
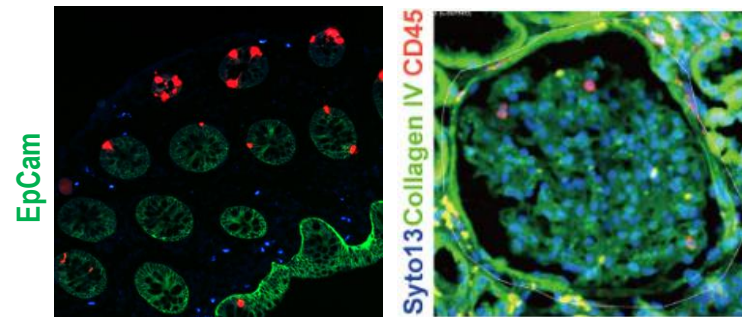
肿瘤浸润免疫细胞(T细胞, B细胞, 髓系细胞)



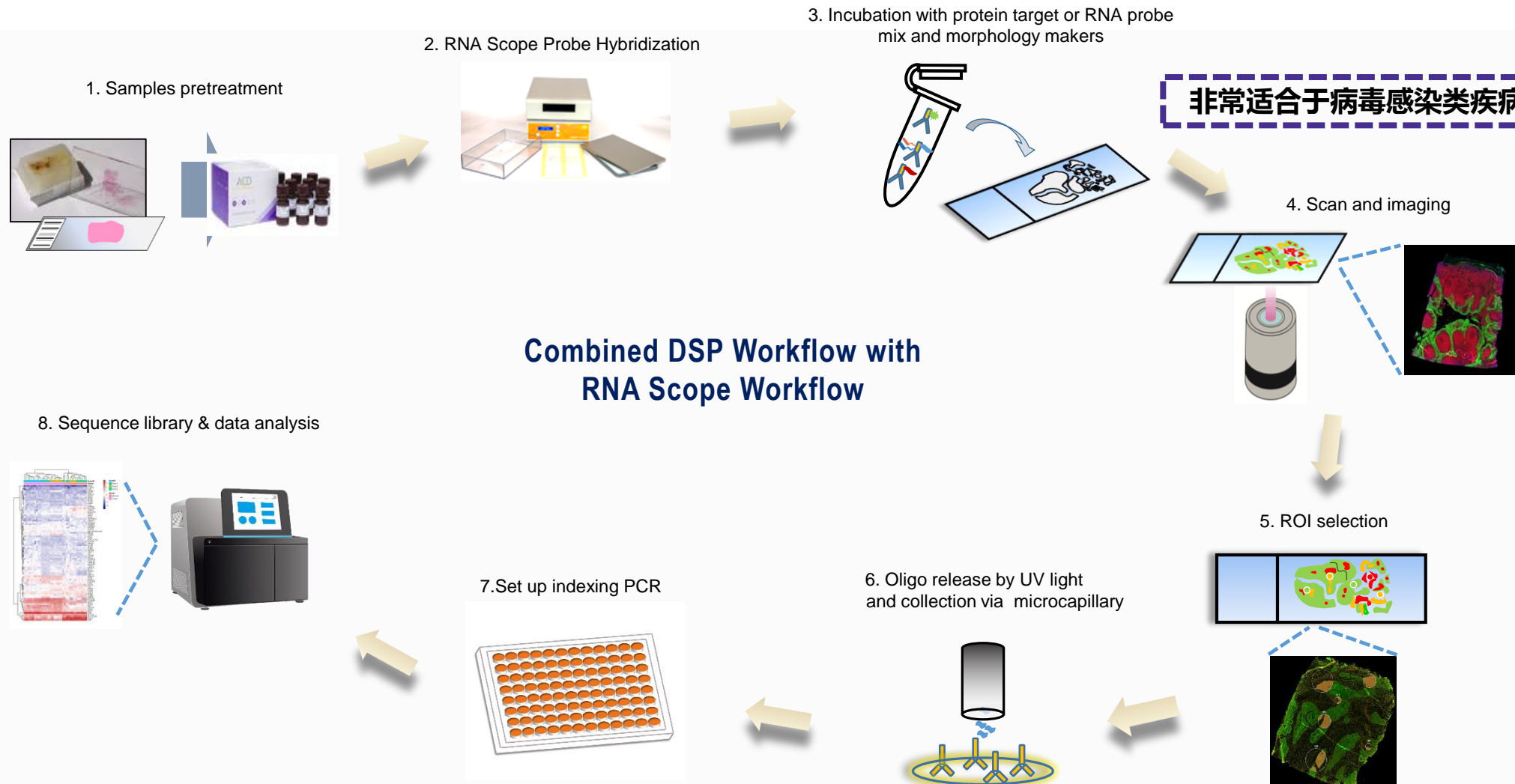
肿瘤相关基质细胞(成纤维类)



其他特殊标志物染色

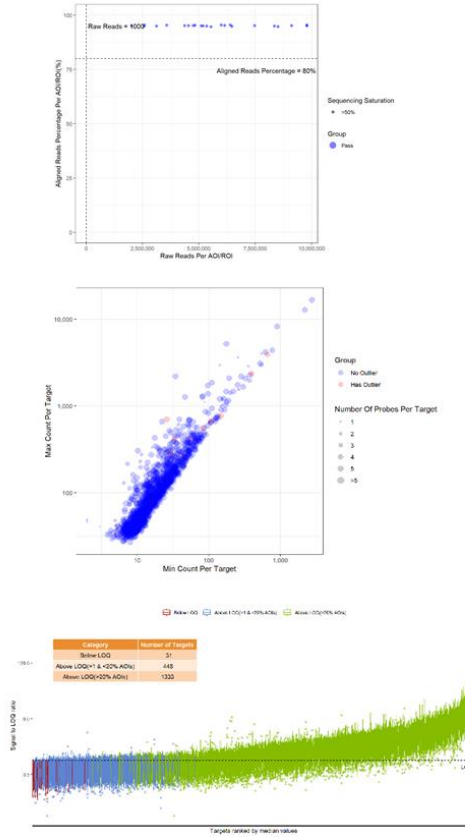
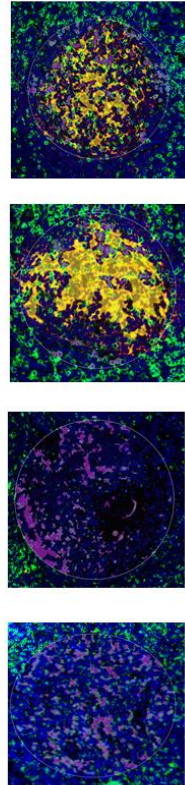
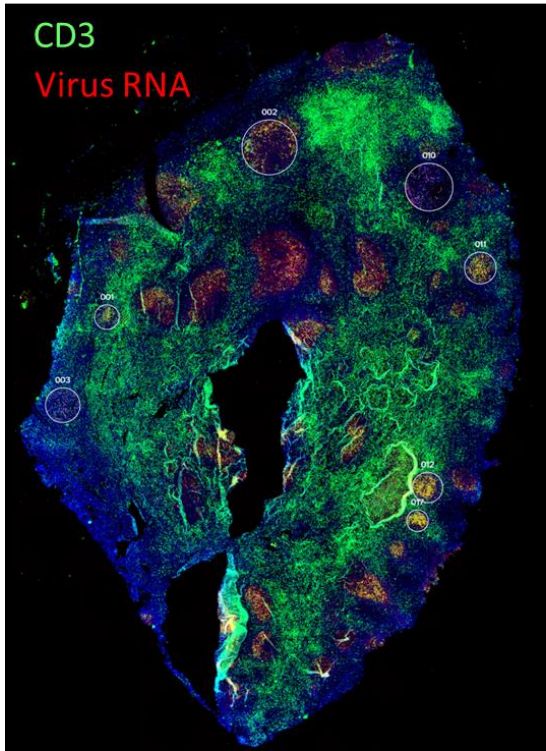


结合RNA原位杂交的DSP空间分析策略



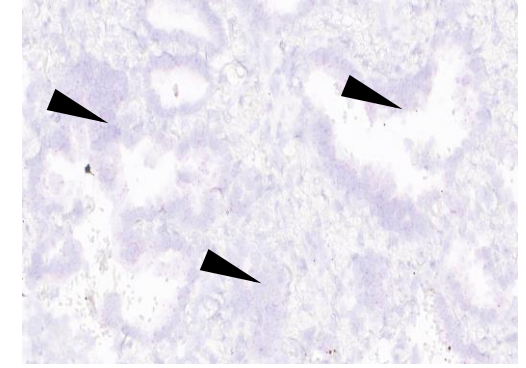
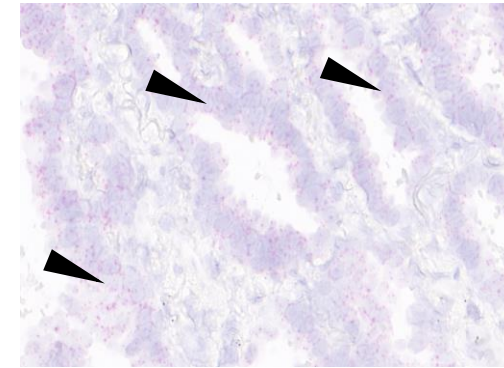
结合RNA原位杂交的DSP空间分析策略

RNAScope

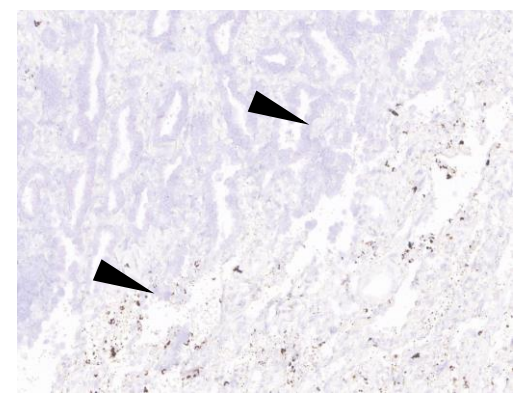
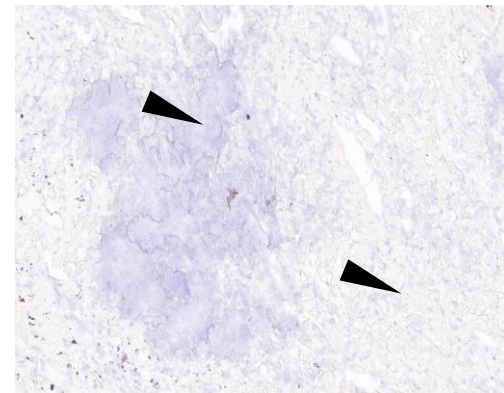


Base Scope

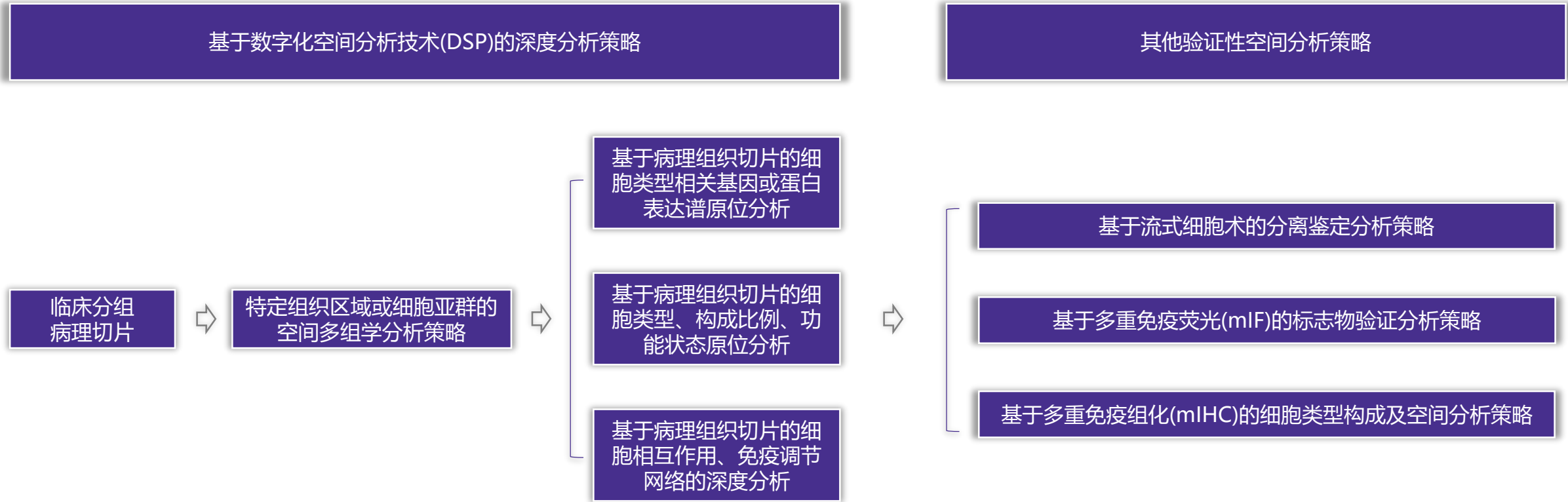
Amplified HER2



Non-Amplified regions

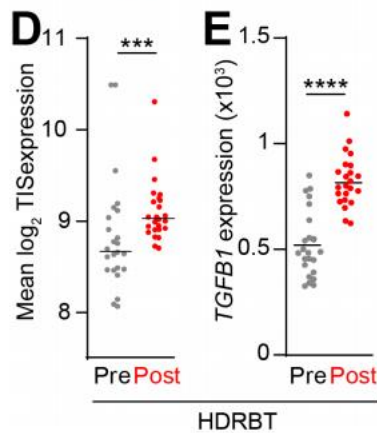
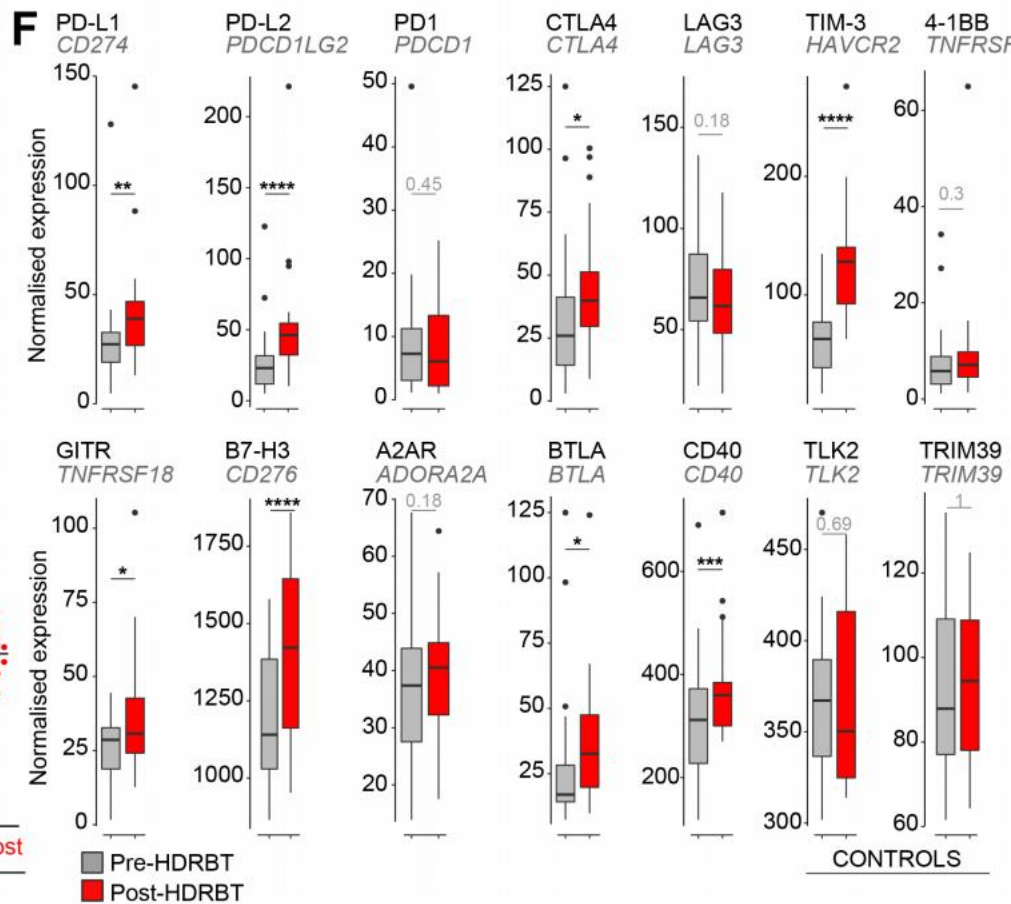
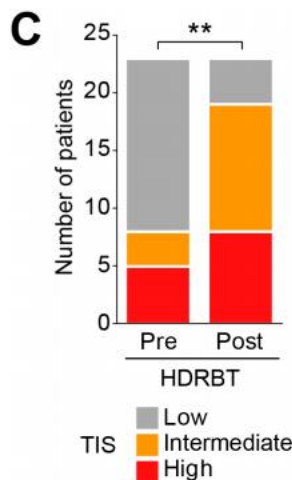
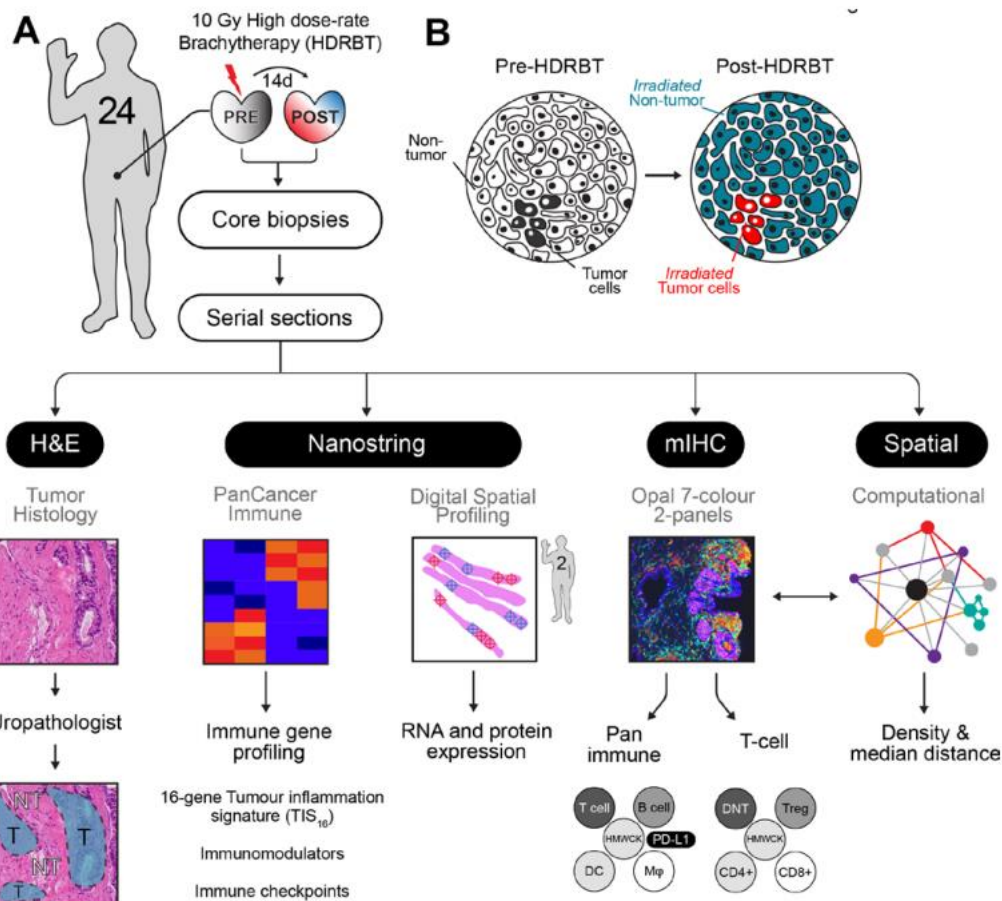


基于临床病理切片的常见空间分析策略组合



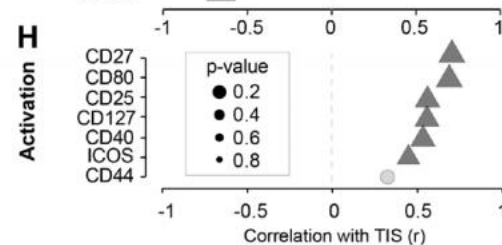
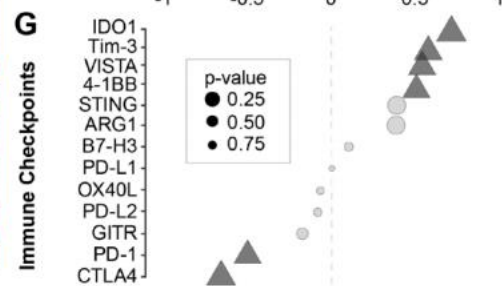
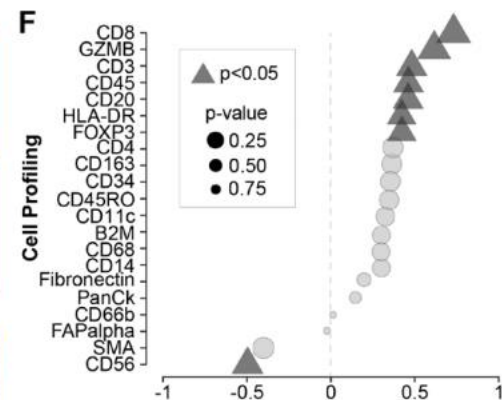
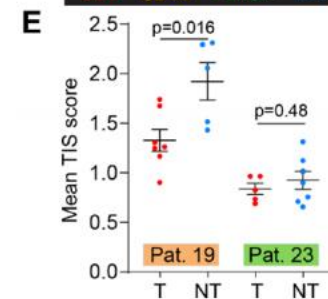
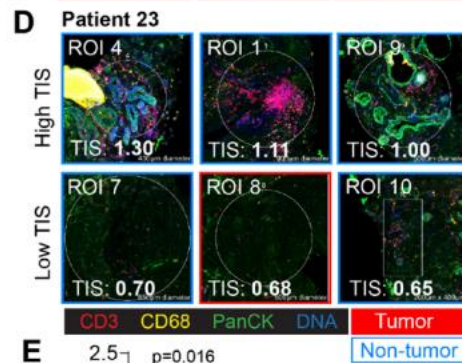
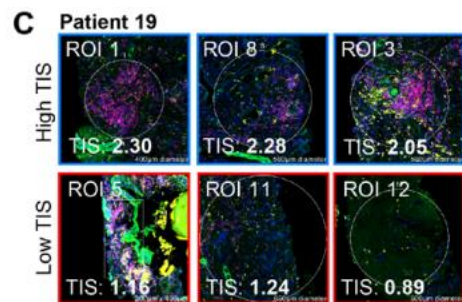
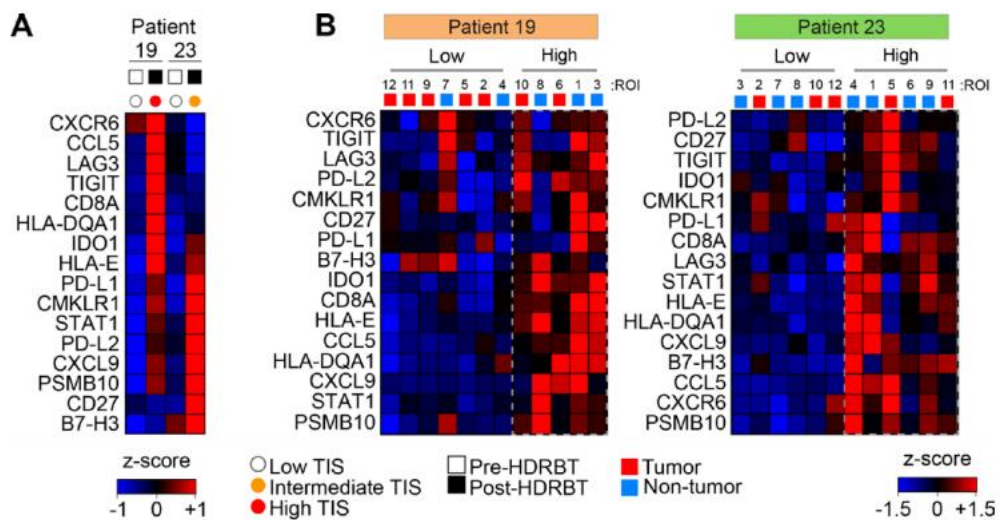
基于临床病理切片的常见空间分析策略组合

通过DSP技术对自定义的肿瘤免疫特征基因或蛋白集群进行聚类 and 分型



基于临床病理切片的常见空间分析策略组合

通过DSP技术对不同患者的不同ROI区域的肿瘤免疫特征基因或蛋白集群进行聚类 and 分型

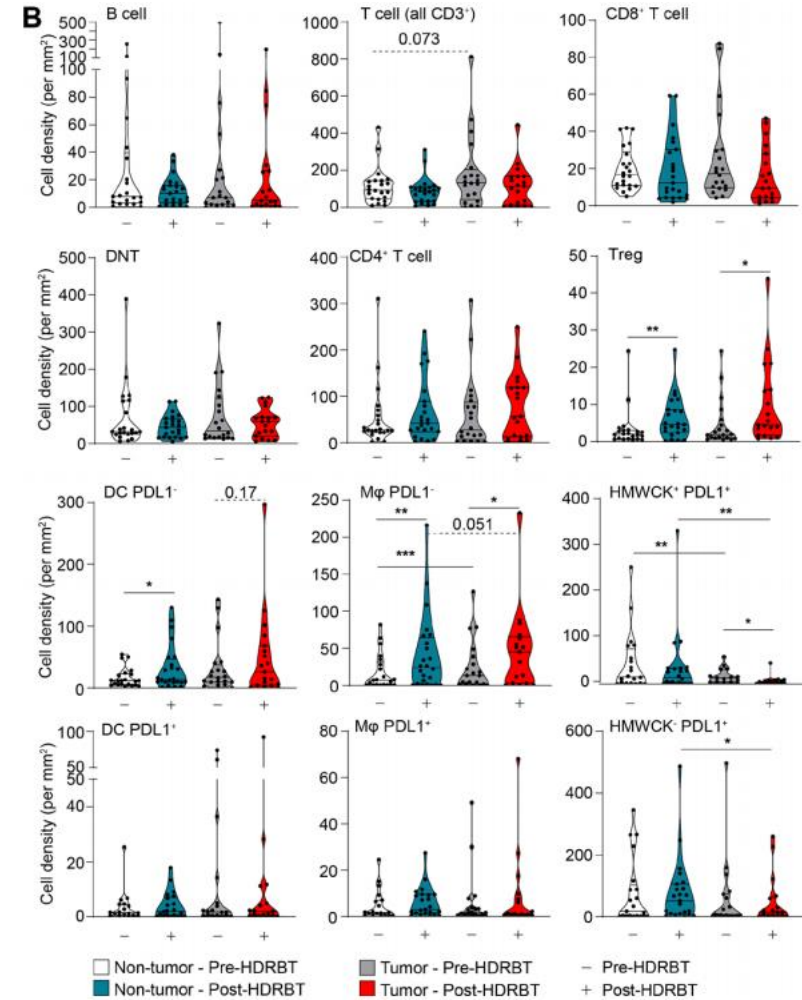
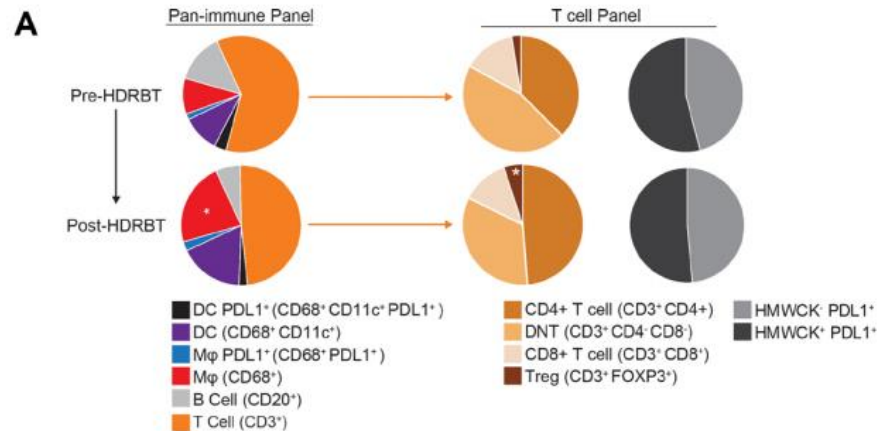


基于临床病理切片的常见空间分析策略组合

通过mIHC技术对治疗前后的各类免疫细胞类型分布、密度进行分析

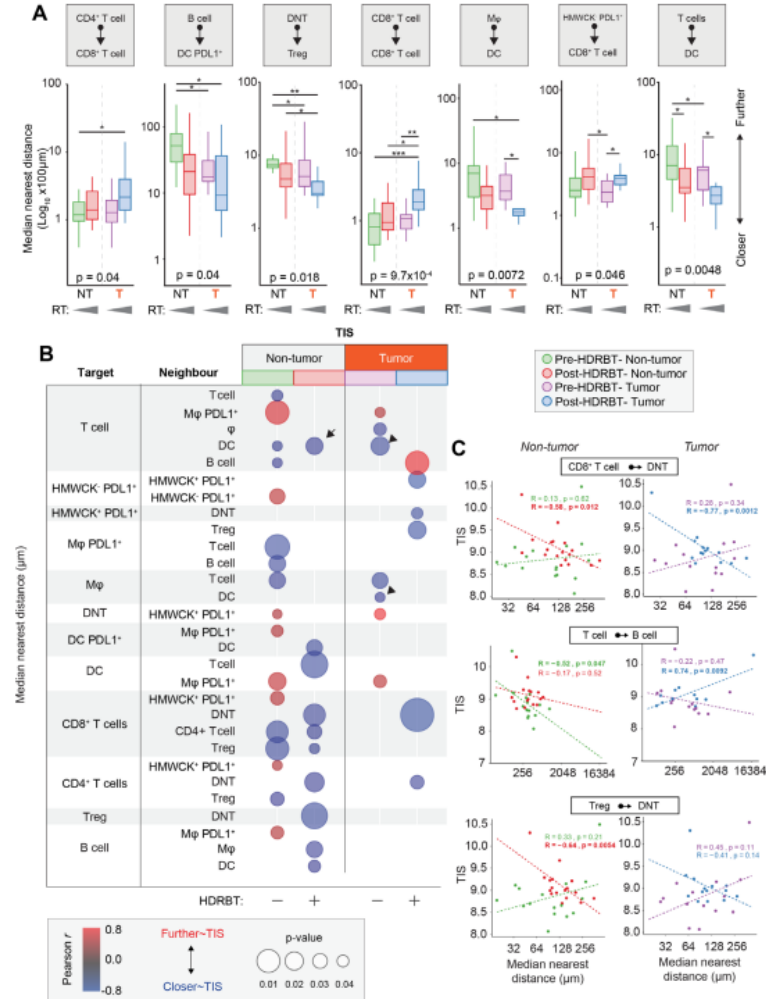
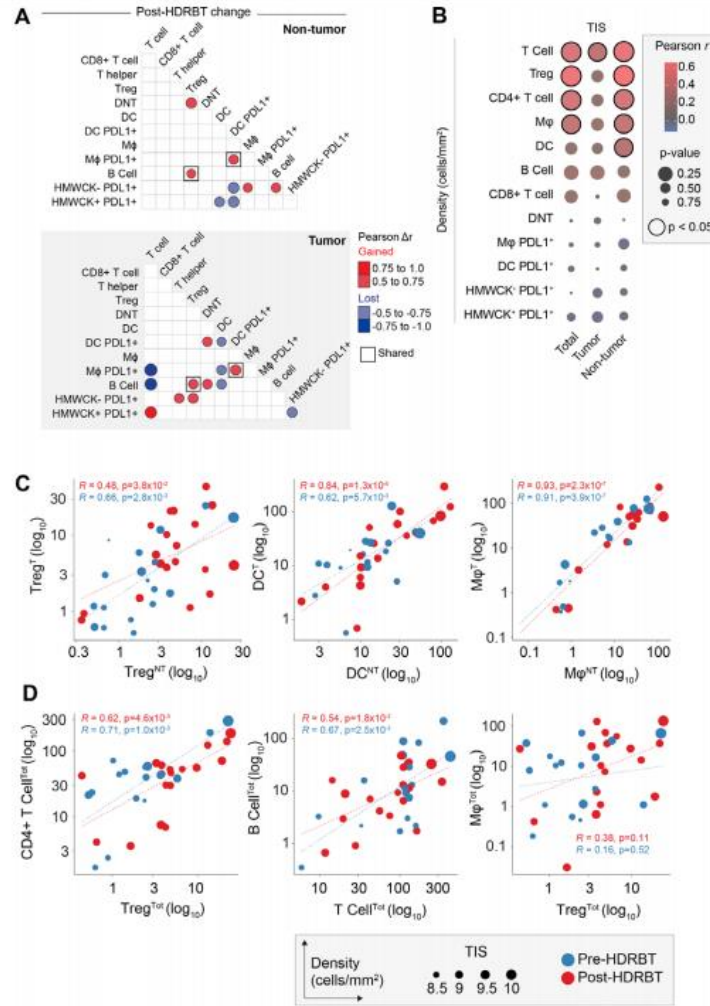
| Table 2 Immune subsets assessed by multiplex IHC | | | |
|--|------------|----------------------------------|--------------------------------------|
| Marker expression | Panel | Cell subset | Abbreviation |
| CD3 ⁺ CD4 ⁺ CD8 ⁻ | T cell | T-helper | T-helper |
| CD3 ⁺ CD8 ⁺ CD4 ⁻ | T cell | CD8 ⁺ T cells | CD8 ⁺ T cells |
| CD3 ⁺ CD4 ⁺ FOXP3 ⁺ | T cell | Regulatory T cell | Treg |
| CD3 ⁺ CD4 ⁻ CD8 ⁻ | T cell | Double negative T cell | DNT |
| CD3 ⁺ | Pan immune | Pan T cell | T cell |
| CD20 ⁺ | Pan immune | B cell | B cell |
| CD68 ⁺ | Pan immune | Macrophage | φ |
| CD68 ⁺ CD11c ⁺ | Pan immune | Dendritic cell | DC |
| CD68 ⁺ PDL1 ⁺ | Pan immune | PDL1 ⁺ macrophage | φ ^{PDL1+} |
| CD68 ⁺ CD11c ⁺ PDL1 ⁺ | Pan immune | PDL1 ⁺ dendritic cell | DC ^{PDL1+} |
| HMWCK ⁺ PDL1 ⁺ | Pan immune | PDL1 ⁺ basal cells | HMWCK ⁺ PDL1 ⁺ |
| HMWCK ⁻ PDL1 ⁺ | Pan immune | PDL1 ⁺ nonbasal cells | HMWCK ⁻ PDL1 ⁺ |

IHC, immunohistochemistry.

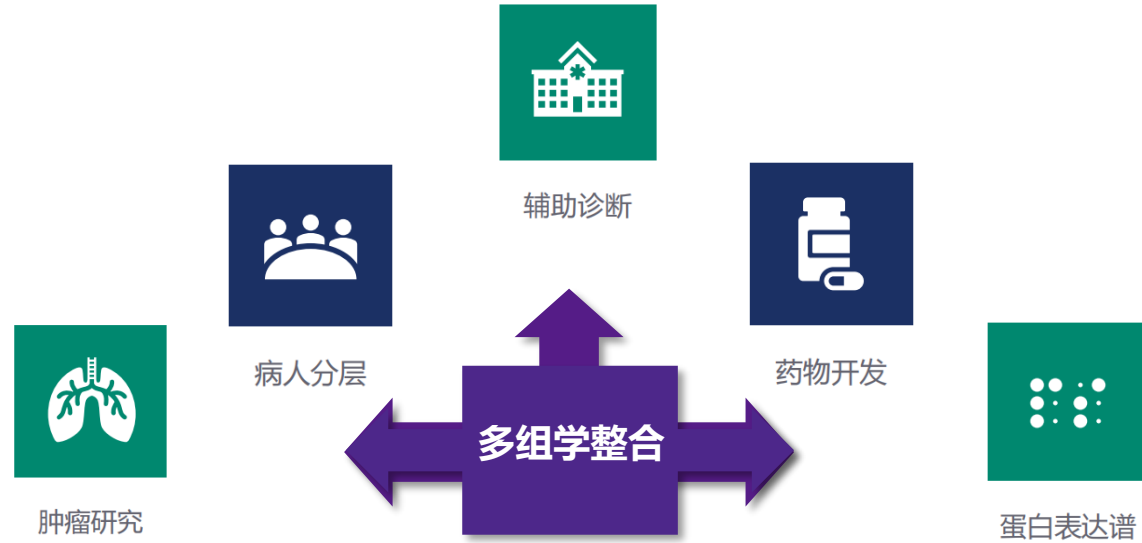


基于临床病理切片的常见空间分析策略组合

通过DSP技术分析获得**肿瘤免疫特征基因信息**与mIHC技术分析获得的**细胞密度与空间分布信息**，可以进行交互分析，获得两者与疾病治疗有关的相关性数据



非因生物多组学研究策略助力转化医学研究



药物筛选平台

- 肿瘤细胞系(80+)
- PDC平台
- 分子克隆平台
- 基因敲除平台

空间多组学平台

- DSP空间组技术
- 空间单细胞技术

靶向蛋白组平台

- RPPA靶向蛋白组技术 (400+药物靶点)
- TargetScan信号通路分析平台

单细胞转录组平台

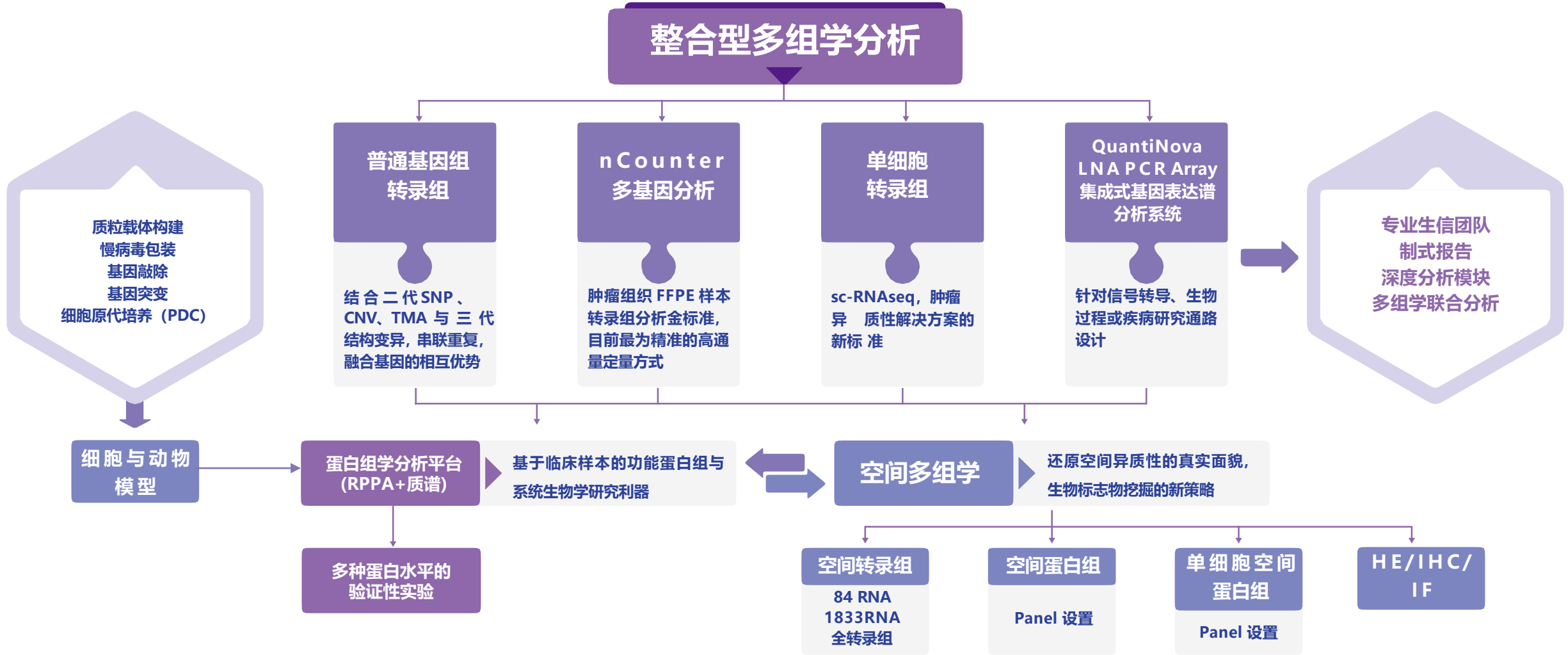
- GexScope技术

靶向转录组平台

- nCounter技术
- PCR-Array技术



非因生物多组学研究策略



非因生物简介——致力于转化医学

转化医学研究综合解决方案

基础研究

- ◆ **RPPA 蛋白组学**
国内唯一技术服务商；
350+ 经严格验证的抗体资源库；
1700+ 学术论文, 8+ 平均影响因子
- ◆ **DSP空间多组学**
DSP技术合作开发方；
国内首个官方认证技术服务商；
80+项目, 800+样本；
100+学术论文, 10+平均影响因子
- ◆ **单细胞空间蛋白组学**
国内领先的空间单细胞蛋白组学服务商；
350+经严格验证的抗体资源库；
40+学术论文, 5+平均影响因子

转化医学

- ◆ **RPPA蛋白组学**
临床蛋白生物标志转化的完美解决方案；
高样本通量（单次上样多达1000+）；
多标志物平行分析（同时分析10-500个潜在生物标志物）；
高度可定制化（靶点根据需求开发）；
高准确度（基于高度定量的免疫学反应原理）；
广泛适用于组织和体液来源的样本
- ◆ **空间多组学**
基于微环境的生物标志物转化解决方案

临床应用

- ◆ **RPPA蛋白组学**
作为新型的LDT药物伴随诊断技术，已经在2021年开始实现临床应用；

临床适应症开发涵盖乳腺癌，结直肠癌，肺癌，胰腺癌等领域；

非因生物致力打造以DSP技术为核心竞争力的临床伴随诊断产品。



非因生物搭建的符合国际标准的精准医疗研究中心



Mills癌症个体化诊疗研究中心



RPPA蛋白组学平台



细胞学平台



分子生物学平台



nCounter及空间生物学平台



空间单细胞表型平台



病理学平台

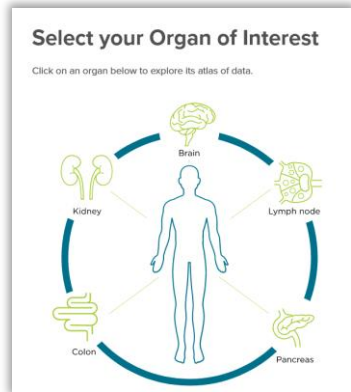
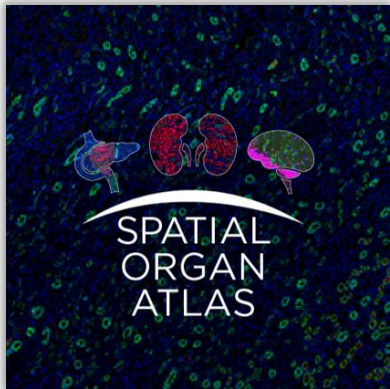
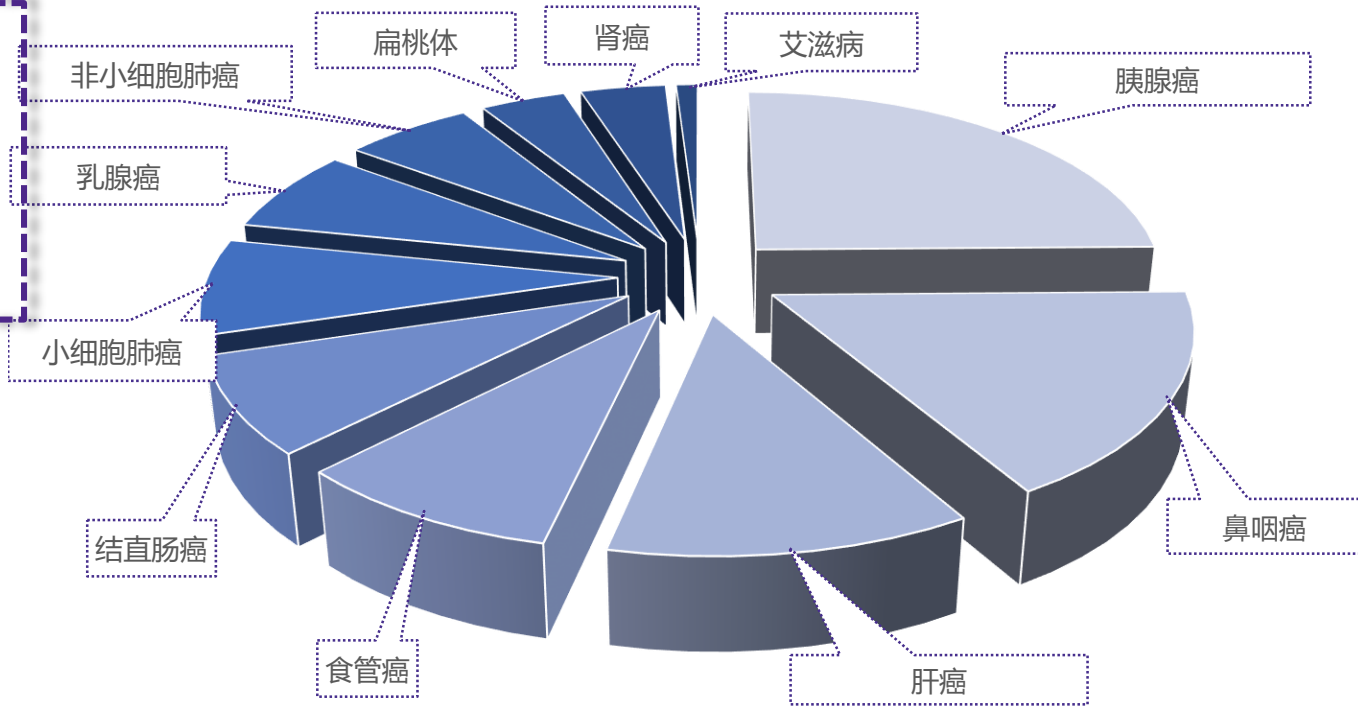


单细胞转录组平台



非因生物建设的国内最大的空间组学研究平台

- ✓ nanoString全球8大DSP空间组Premier access site
- ✓ 全球第一批5家获得nanoString官方认证DSP空间组学服务商之一 (国内唯一)
- ✓ Organbook空间组学国际研究合作方



非因生物助力创新科学研究



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TECHNOLOGICAL ADVANCES OPEN ACCESS

Spatially-resolved proteomics and transcriptomics: An emerging digital spatial profiling approach for tumor microenvironment

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Biotechnology Journal

MINI-REVIEW

Spatial transcriptomics and proteomics technologies for deconvoluting the tumor microenvironment

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<https://doi.org/10.1186/s12943-022-01526-8>

Molecular Cancer

REVIEW Open Access

Proteomics technologies for cancer liquid biopsies

Zhiyong Ding^{1*}, Nan Wang¹, Ning Ji^{2,3} and Zhe-Sheng Chen^{2*}

Abstract

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Journal of Ethnopharmacology

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Ameliorative effect of *Gastrodia elata* Blume extracts on depression in zebrafish and cellular models through modulating reticulum 4 receptors and apoptosis

Rongchun Wang^{b,c,e,1}, Qingyu Ren^{a,b,c,e,1}, Daili Gao^{b,c,e,1}, Yam Nath Paudel^d, Xia Li^e, Lizhen Wang^{b,c}, Pengyu Zhang^{a,b,c}, Baokun Wang^{b,c}, Xueliang Shang^{a,c}, Meng Jin^{b,c,e,*}

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EVI1 overexpression promotes ovarian cancer progression by regulating estrogen signaling

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REVIEW

Recent progress on targeting leukemia stem cells

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frontiers in Bioengineering and Biotechnology

BRIEF RESEARCH REPORT
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Tumor Microenvironment Profiles Reveal Distinct Therapy-Oriented Proteogenomic Characteristics in Colorectal Cancer

Nan Wang¹, Rongshui Wang¹, Xia Li¹, Zhenhao Song¹, Lingbo Xia¹, Jue Wang², Li Zhang³, Aiwu Wu^{4*} and Zhiyong Ding^{1*}

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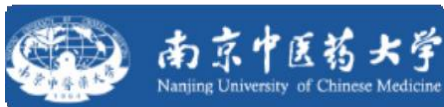
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