SITC 2019 Gaylord National Hotel Nov 6 10

Gaylord National Hotel & Convention Center Nov. 6-10

NATIONAL HARBOR, MARYLAND





There are no relevant disclosures to the presentation.





Spatial transcriptomics

Itai Yanai

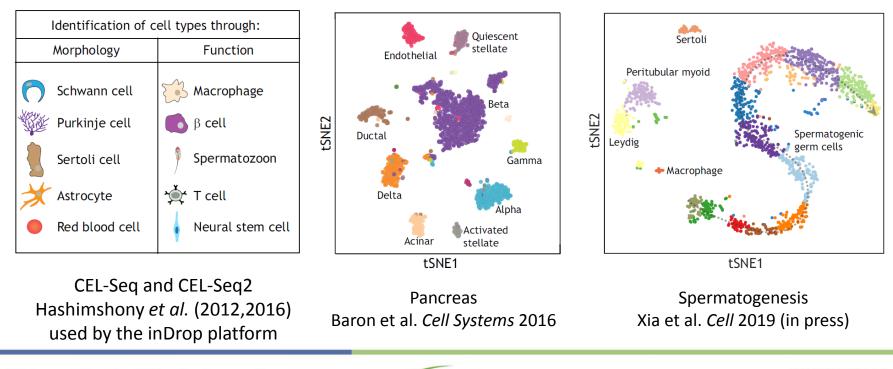
Institute for Computational Medicine

NYU School of Medicine

NYU Langone Health



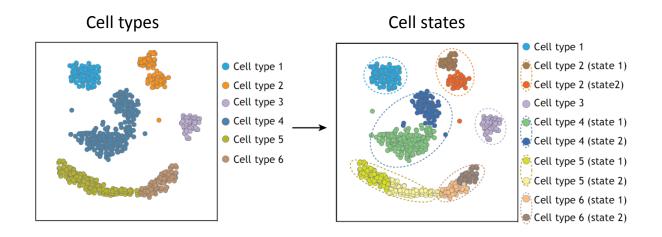
Historical and scRNA-Seq approaches to identifying cell types



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Distinguishing cell types and cell states

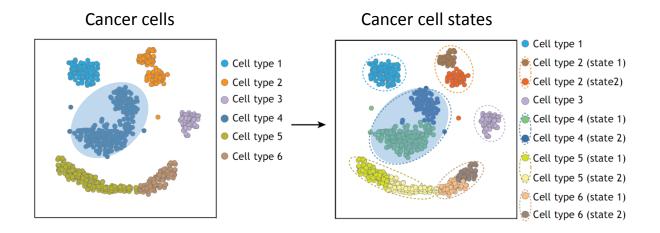


Xia & Yanai Development 2019





Distinguishing cancer cell states

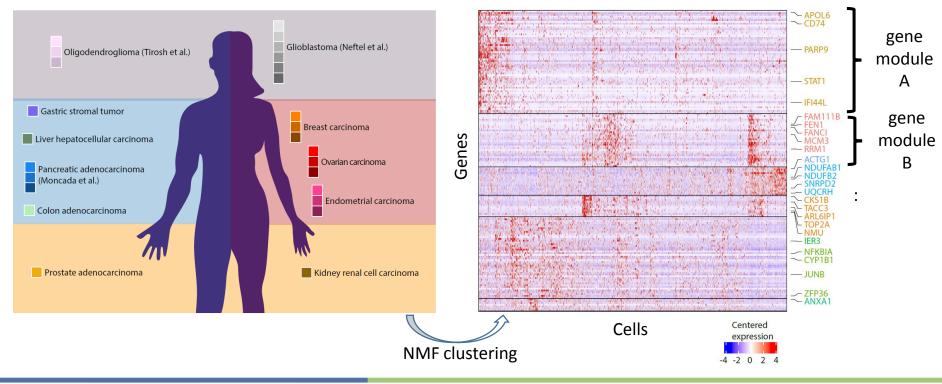


Xia & Yanai Development 2019



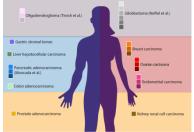


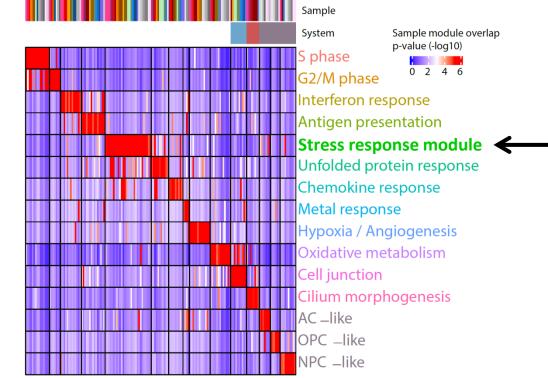
We studied 24 tumors of different cancer types and used scRNA-Seq to detect gene modules





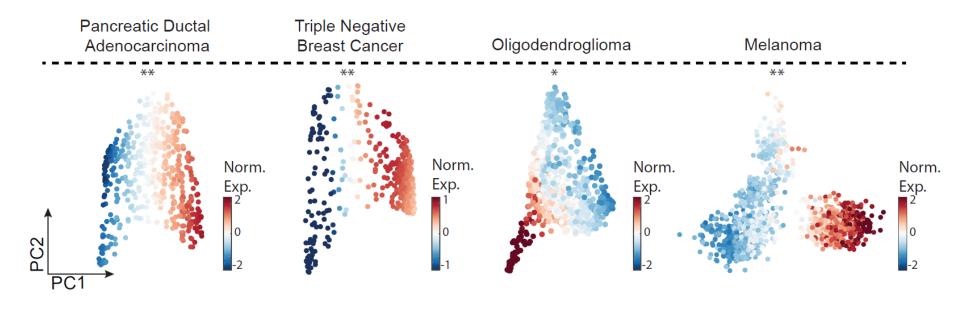
Universal and specific cancer gene modules





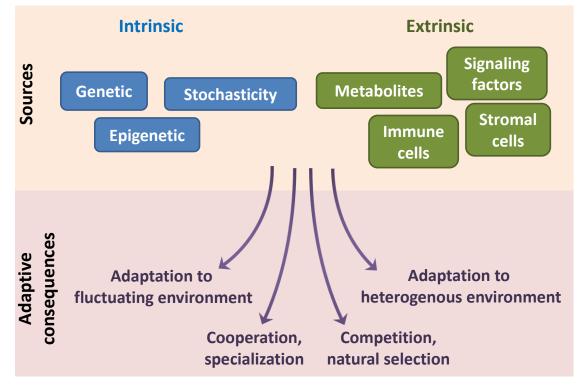


A state expressing the stress module across cancer types





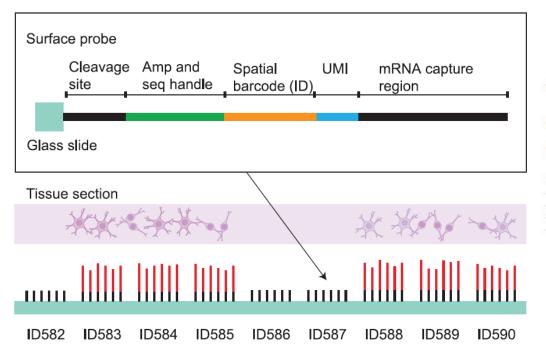
What are the roles of cancer cell states in tumorigenesis?



Barkley & Yanai. Trends in Cancer 2019



Spatial transcriptomics



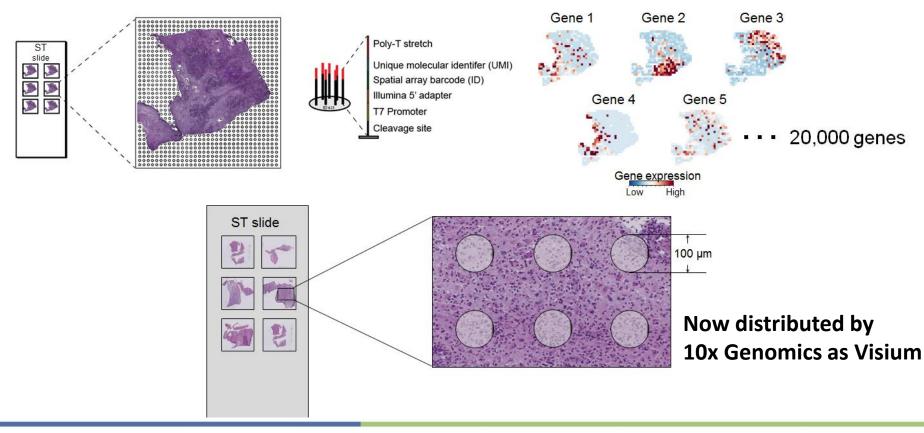
Visualization and analysis of gene expression in tissue sections by spatial transcriptomics

Patrik L. Ståhl,^{1,2*} Fredrik Salmén,^{2*} Sanja Vickovic,²⁺ Anna Lundmark,^{2,3+} José Fernández Navarro,^{1,2} Jens Magnusson,¹ Stefania Giacomello,² Michaela Asp,² Jakub O. Westholm,⁴ Mikael Huss,⁴ Annelie Mollbrink,² Sten Linnarsson,⁵ Simone Codeluppi,^{5,6} Åke Borg,⁷ Fredrik Pontén,⁸ Paul Igor Costea,² Pelin Sahlén,² Jan Mulder,⁹ Olaf Bergmann,¹ Joakim Lundeberg,²⁺ Jonas Frisén¹

Stahl et al. Science 2016



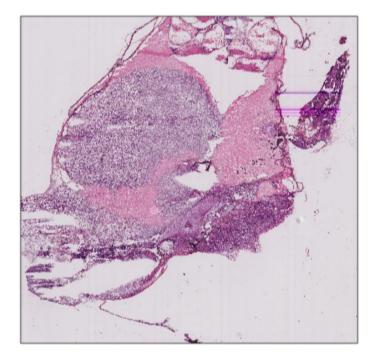
Spatial transcriptomics

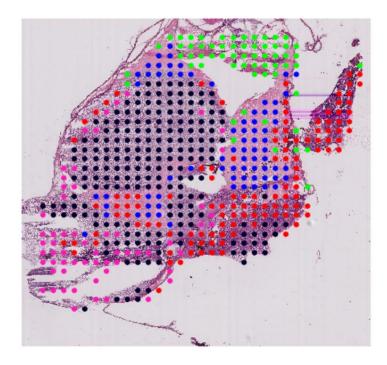


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Spatial transcriptomics on a melanoma tumor

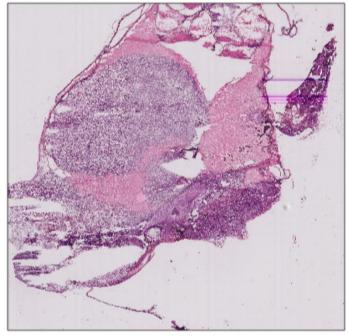


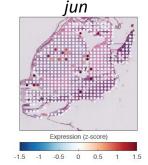


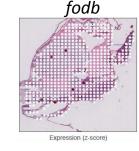




Spatial transcriptomics on a melanoma tumor reveals expression of the stress module in the cancer region

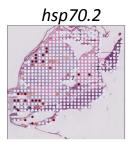


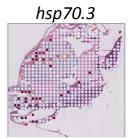






hsp70.1



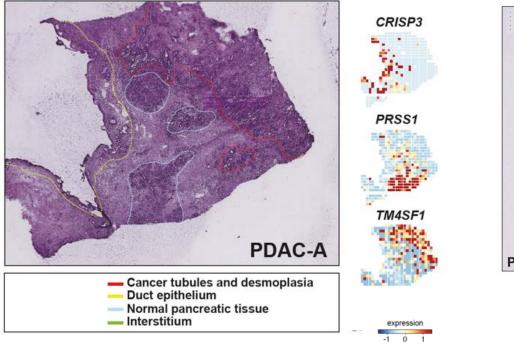


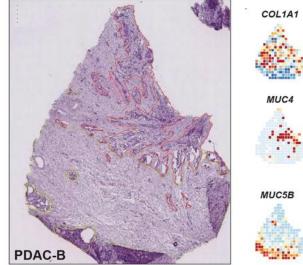






Spatial transcriptomics on pancreatic ductal adenocarcinoma (PDAC) tumor samples

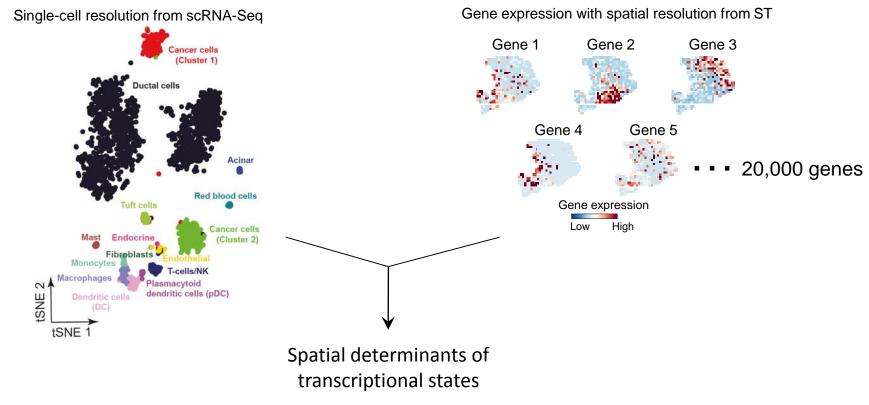




Diane M. Simeone and Cristina H. Hajdu



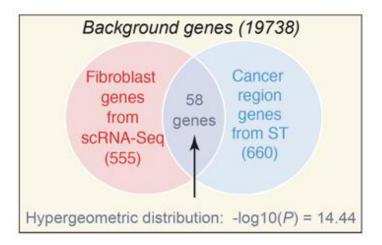
Integration of single-cell and spatial resolution





019

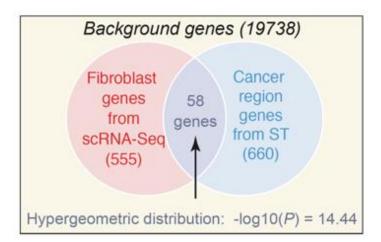
Multimodal Intersection Analysis (MIA) for the tissue-wide inference of cell type localization

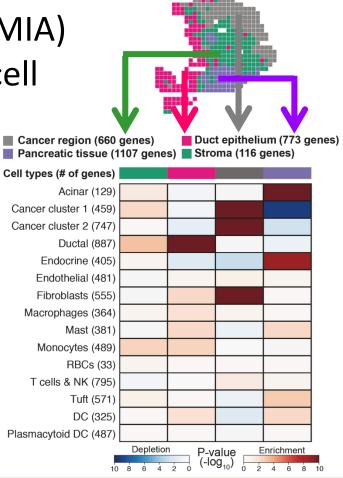






Multimodal Intersection Analysis (MIA) for the tissue-wide inference of cell type localization

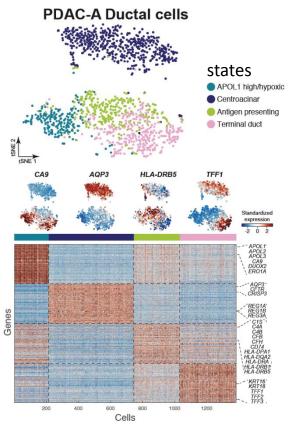


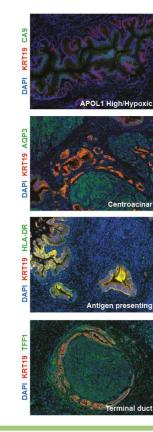


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Identifying states among the ductal cells

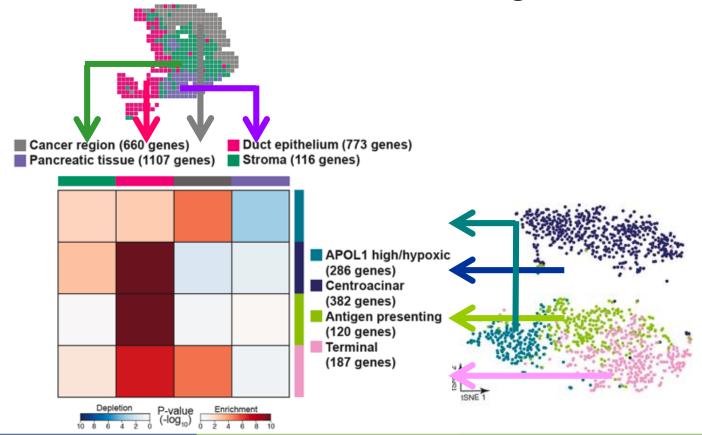








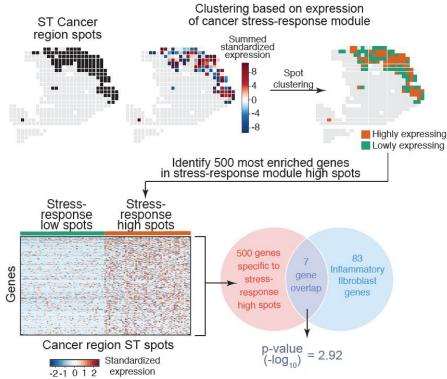
Ductal states across PDAC tissue regions





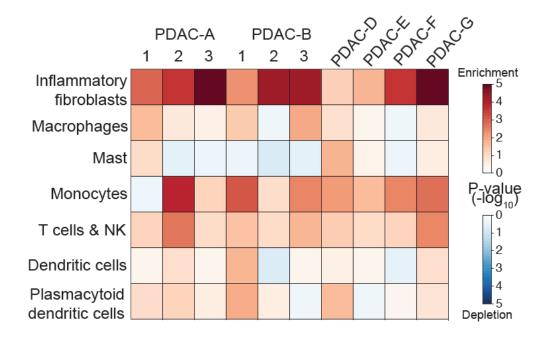


Querying for associations between cell states and cell types of the TME





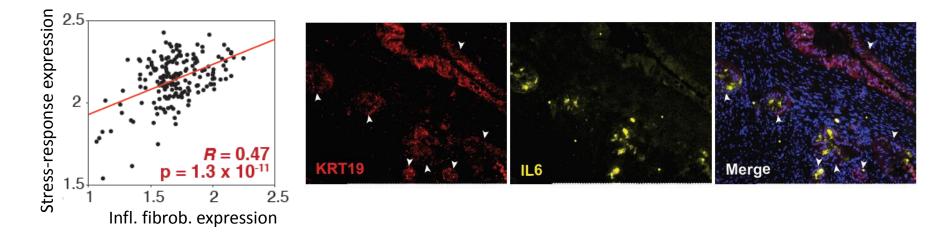
Inflammatory fibroblasts are consistently associated with cancer cells expressing the stress module







The stress module is correlated with inflammatory fibroblasts in the TCGA dataset and in IFs



TCGA data

Immunofluoresence



Summary

PDAC-A Cancer region (660 genes) Duct epithelium (773 genes) Cancer region (660 genes) Duct epithelium (773 genes) Pancreatic tissue (1107 genes) Stroma (116 genes) Pancreatic tissue (1107 genes) Stroma (116 genes) Cell types (# of genes) Multimodal intersection Acinar (129) Cancer cluster 1 (459) Cancer cluster 2 (747) analysis (MIA) maps Ductal (887 Endocrine (405) PDAC-A APOL1 high/hypoxic scRNA-Seg states onto Endothelial (481) (286 genes) Fibroblasts (555) Centroacinar Macrophages (364) (382 genes) ST tissue regions Mast (381 Antigen presenting Monocytes (489) (120 genes) Cluster assignments RBCs (33 Terminal Cancer region T cells & NK (795) (187 genes) Pancreatic tissue Tuft (571 DC (325) Duct epithelium Depletion P-value Enrichment Plasmacytoid DC (487) Stroma etion P-value Enricht 2 4 6 6 20PUL and a start op of the second ACC ACC M PDAC-A PDAC-B 2 3 2 3 1 1 MIA highlights a potential Inflammatory Fibroblasts PDAC TCGA Data functional relationship Macrophages Enrichmer Hypoxia module Ox. phos. module Stress module between inflammatory Mas 2.4 Cancer module expression fibroblasts and a pan-Monocytes P-value (-log₁₀) cancer stress state in T cells & NK R = 0.4734R = 0.16055R = -0.00088209PDAC cancer cells Dendritic cells $p = 1.3 \times 10^{-1}$ p = 0.029927p = 0.99054Depletion 1.5 2 2.5 1.5 2.5 1.5 2 2.5 Plasmacytoid dendritic cells Inflammatory fibroblast signature expression B2 B3 D E F A1 A2 A3 B1 G ST: Cancer stress module 'high' spots

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Acknowledging our wonderful lab and collaborators! NYU Langone Health

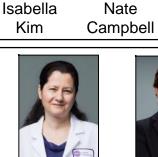
MSK





Dr. Richard White

Alumni: Dr. Tamar Hashimshony Dr. Vlad Grishkevich Dr. Noa Sher Dr. Alison Cole Dr. Harel Zalts Dr. Michal Levin Martin Feder Leon Anavy Sally Khair Natalia Mostov Naftalie Senderovich Shay Ben-Elazar David Silver Avital Polsky



Dr. Cristina H. Hajdu



Dr. Diane M.

Simeone

Mohita Malay Tagore



Miranda Hunter









Dr. Gustavo Franca Dr. Maayan Baron



Bo Xia







Reuben Moncada





Dr. Maayan Pour Dr. Florian Wagner Dr. Gal Avital Felicia Kuperwaser









Itai.Yanai@nyulangone.org

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biomedical challenges.



