

Investigating the Tumor Immune Microenvironment using Cancer Genomics and Computational Pathology

Vésteinn Þórsson, Institute for Systems Biology
Frontiers of Predictive Oncology and Computing II; Oct 17, 2017; NYC

INSTITUTE FOR SYSTEMS BIOLOGY

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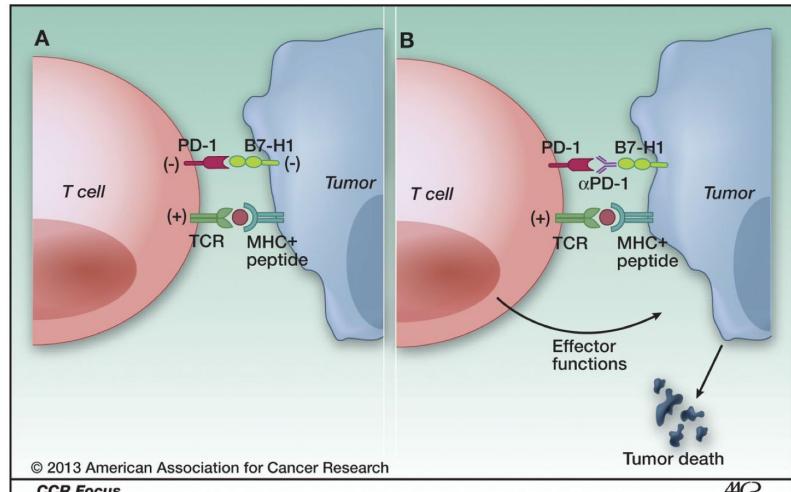
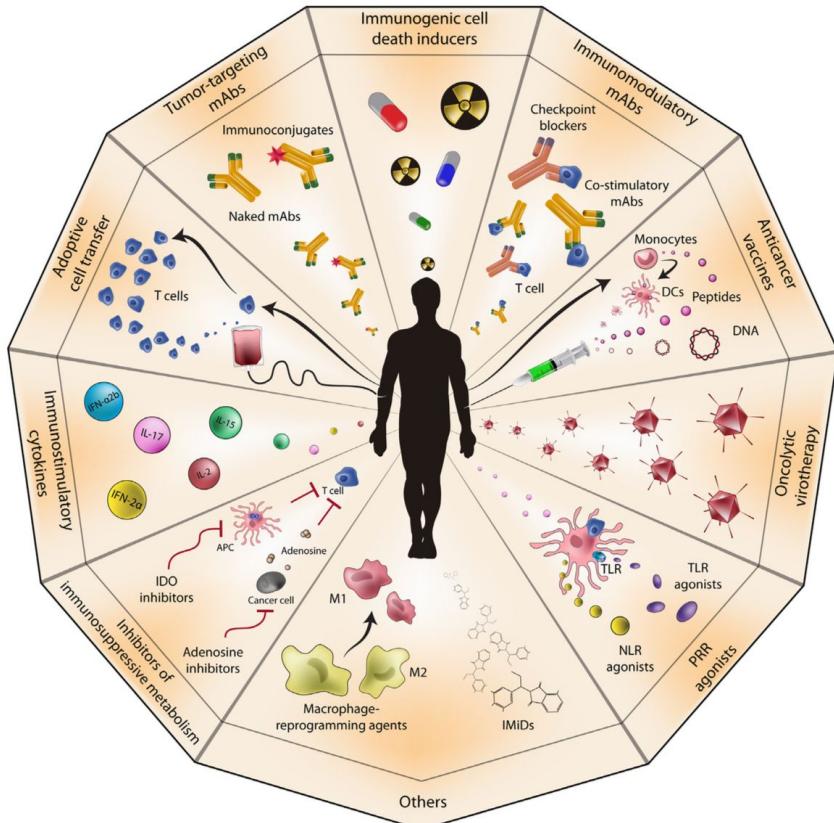
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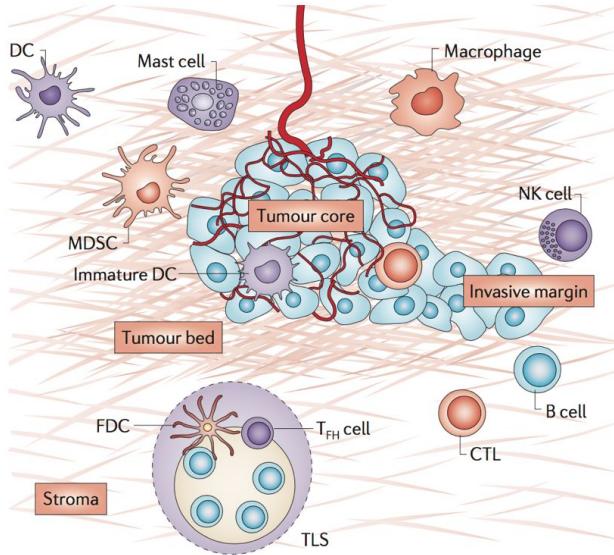
Cancer Immunotherapy

Checkpoint Inhibition



AACR Clinical Cancer Research

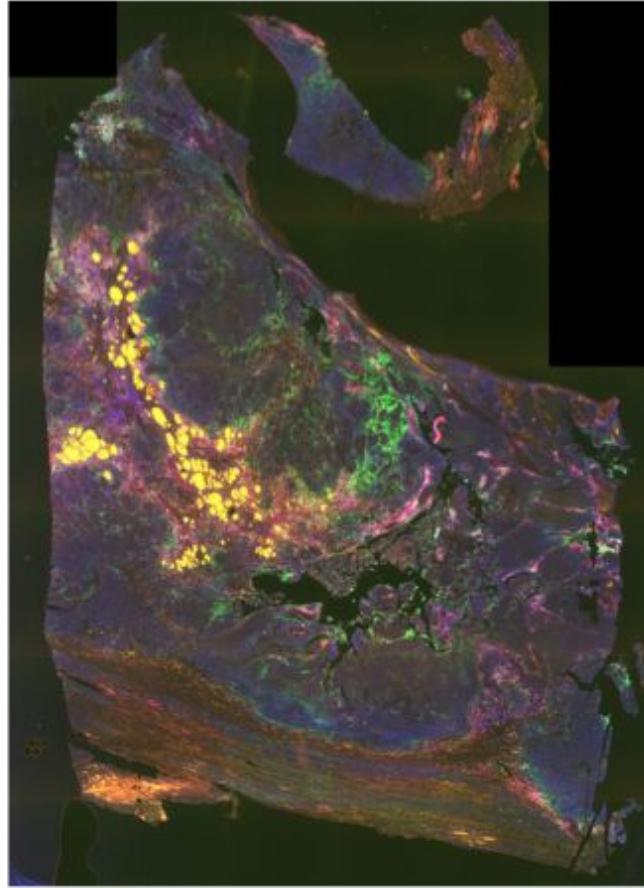
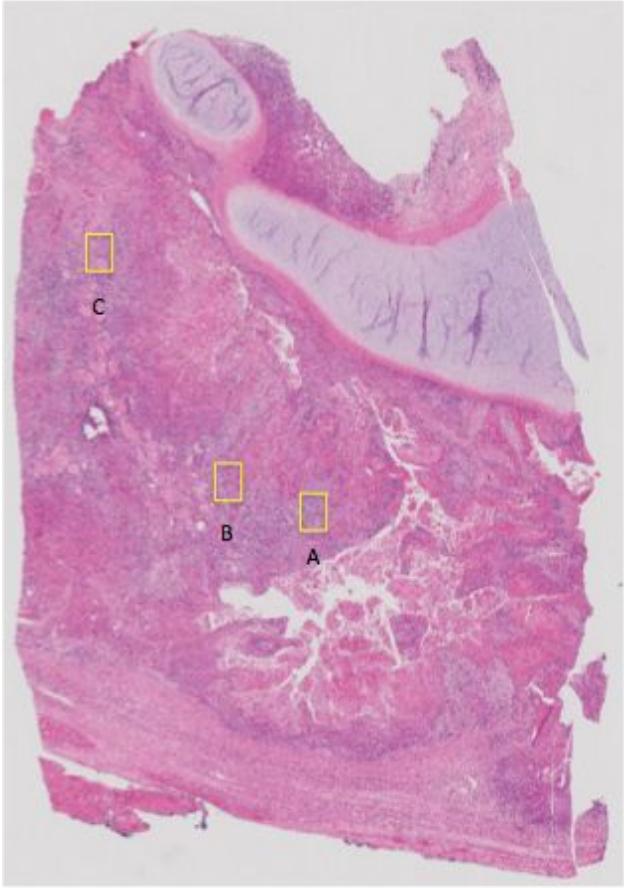
Constitution of Tumor Microenvironment Correlates with Prognosis



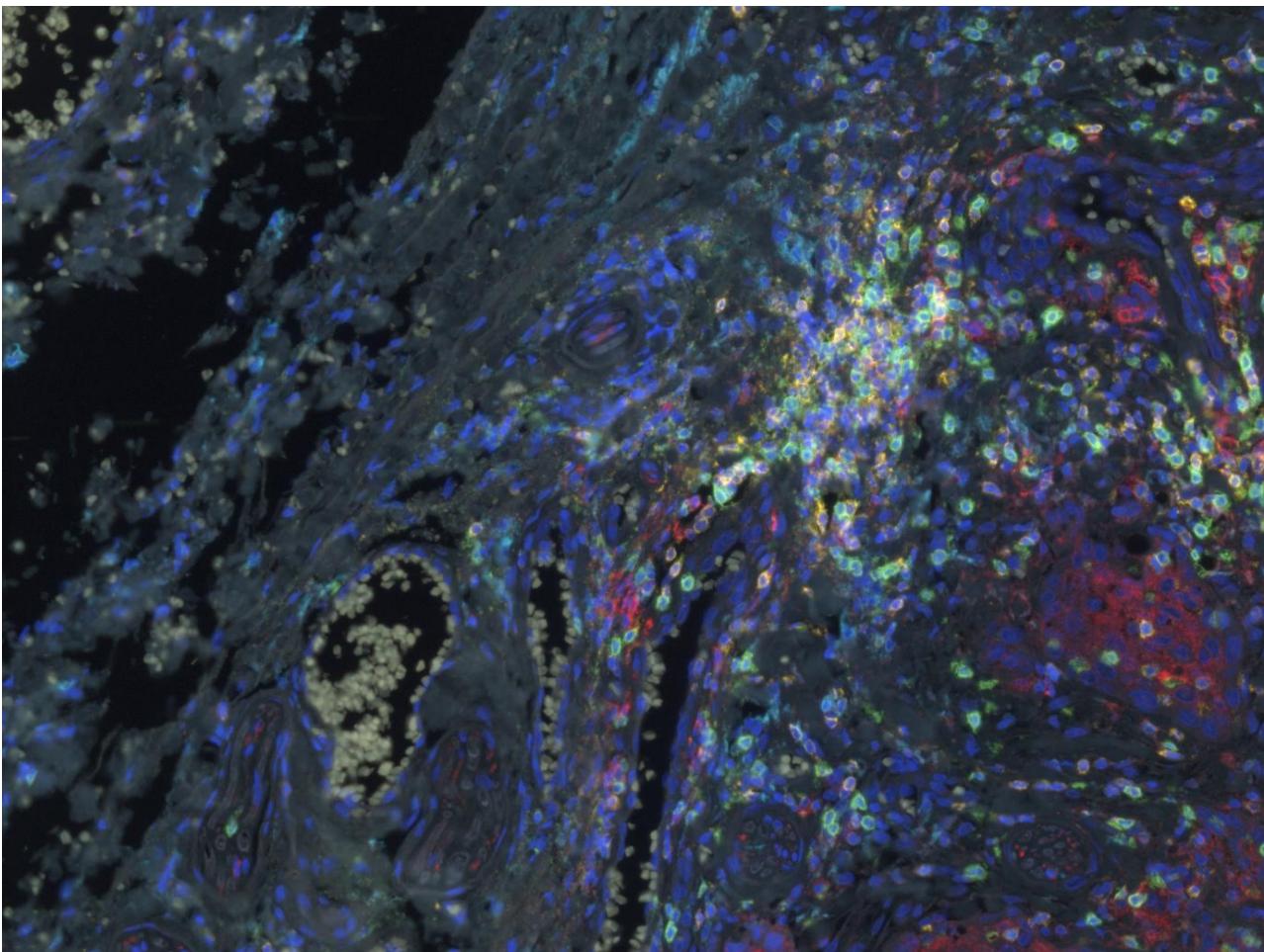
Fridman WH, Pagès F, Sautès-Fridman C, Galon J. *The immune contexture in human tumours: impact on clinical outcome*; Nat Rev Cancer. 2012 Mar 15;12(4):298-306

Galon et al , ASCO 2016 *Validation of the Immunoscore (IM) as a prognostic marker in stage I/II/III colon cancer: Results of a worldwide consortium-based analysis of 1,336 patients.*

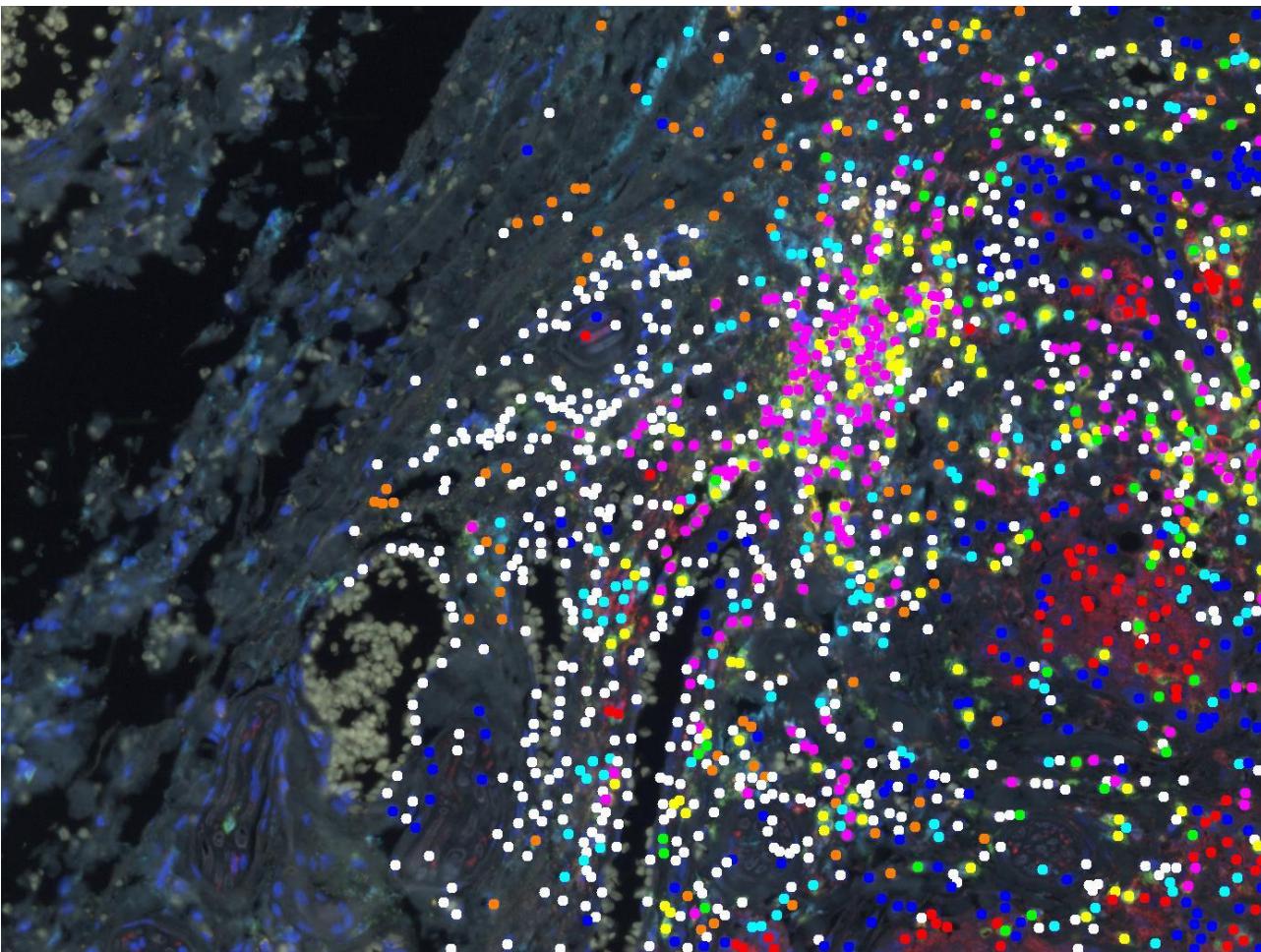
<http://meetinglibrary.asco.org/record/123627/abstract>



Carmen Ballesteros-Merino and Bernard FoxEarl A. Chiles Research Institute



Bernard Fox lab, Earl A. Chiles Research Institute



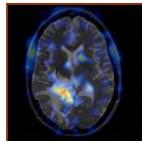
Bernard Fox lab, Earl A. Chiles Research Institute

	Spatial	Molecular	Cellular	Tissue	Organism
H&E	✓	✗	✓	✓	
IHC	✓	✓	✓	✓	
Cancer Genomics	✗	✓	✓	✓	

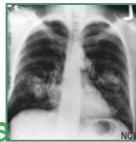
TCGA: Tumor analysis on multiple platforms

33 types of cancer

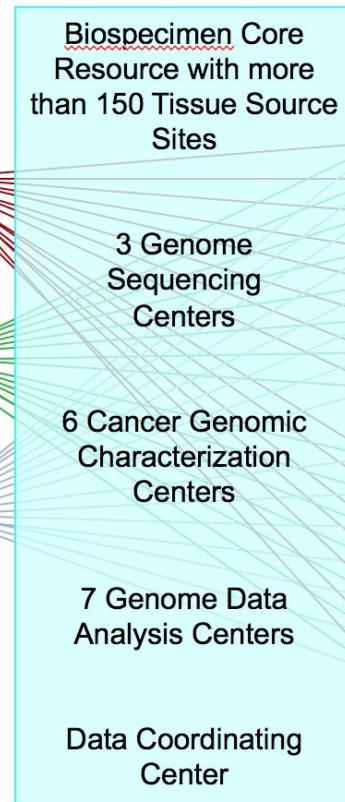
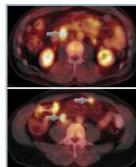
glioblastoma
multiforme
(brain)



squamous carcinoma
(lung)

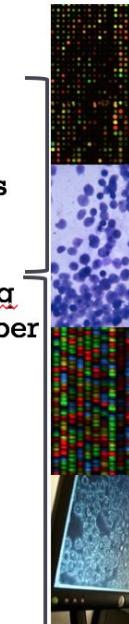


serous
cystadenocarcinoma
(ovarian)

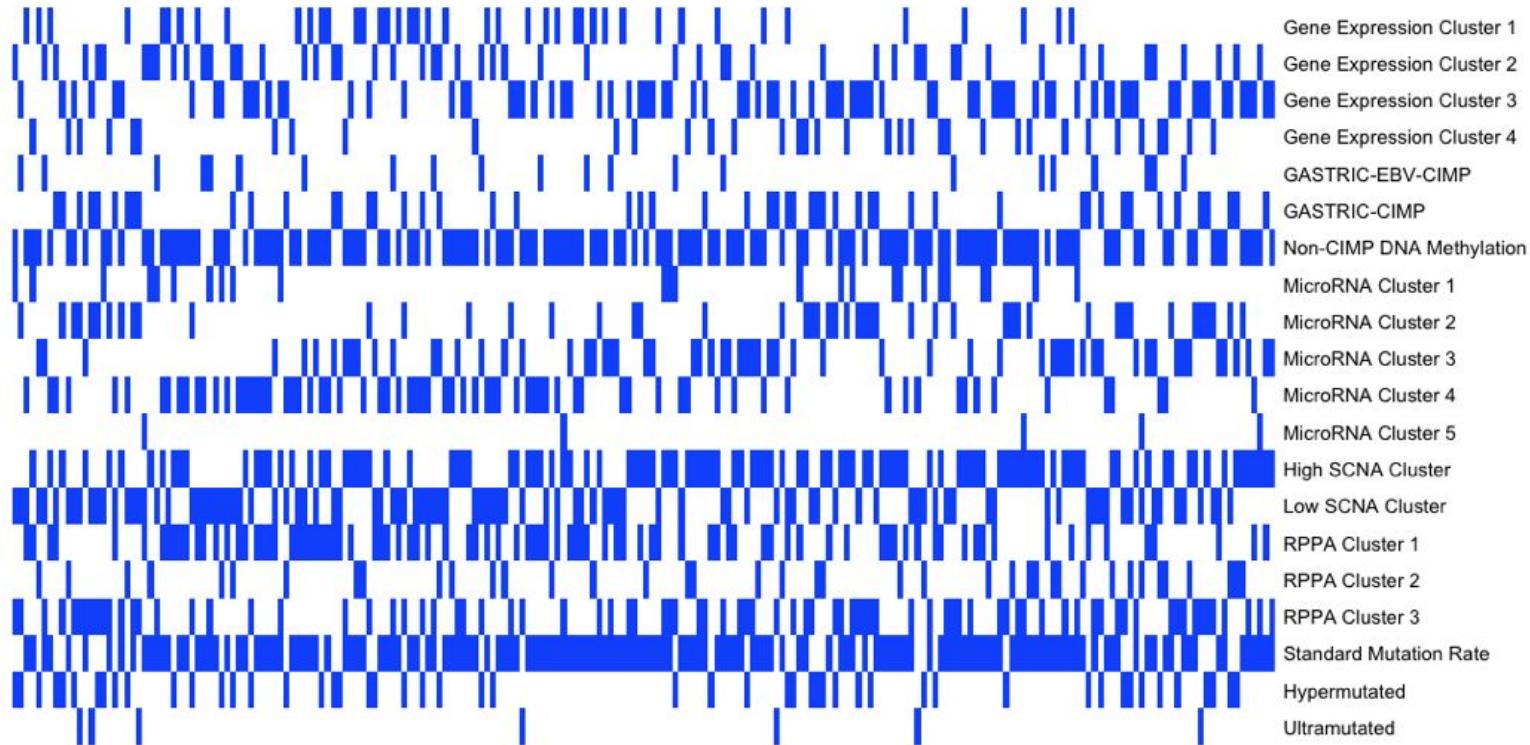


Multiple data types

- Clinical diagnosis
- Treatment history
- Histologic diagnosis
- Pathologic report/images
- Tissue anatomic site
- Surgical history
- Gene Expression/RNAseq
- Chromosomal copy number
- DNA sequence
- DNA mutations
- Methylation patterns
- miRNA expression
- RPPA (protein)
- Loss of heterozygosity

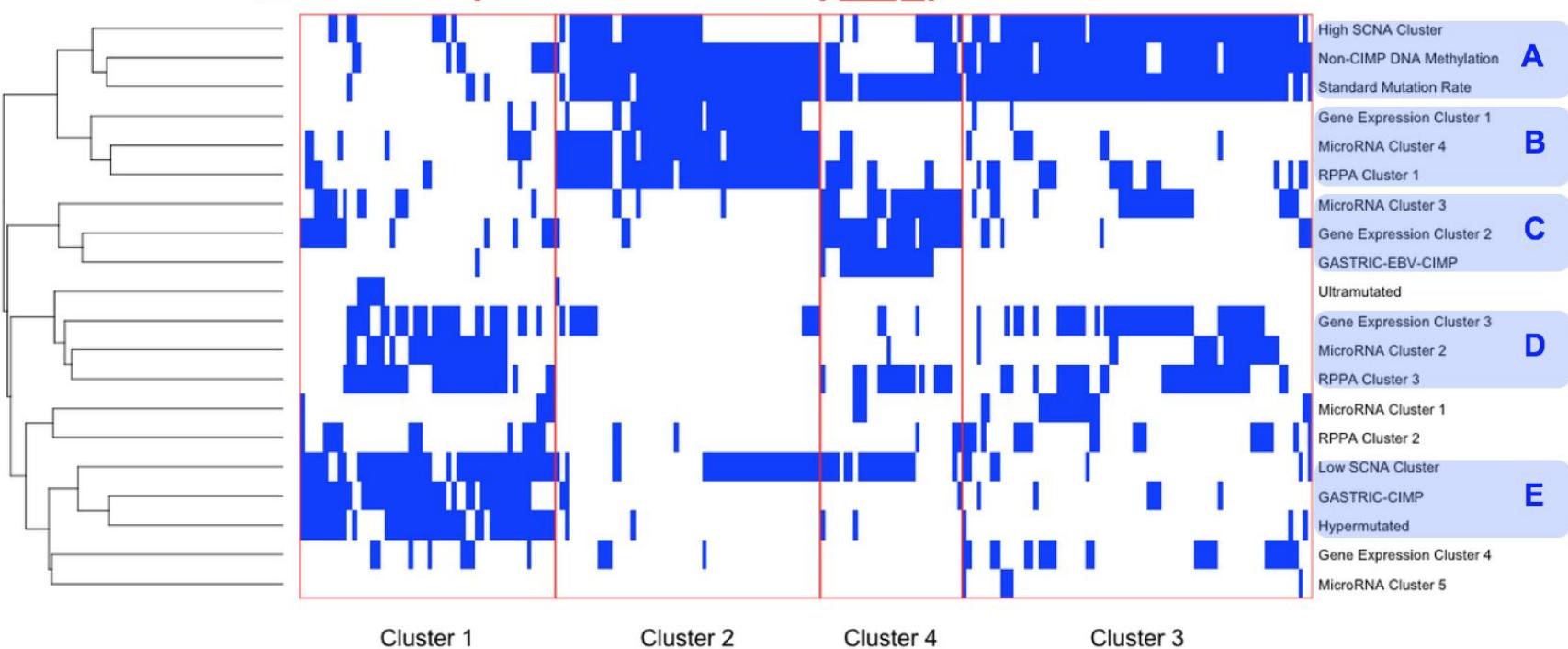


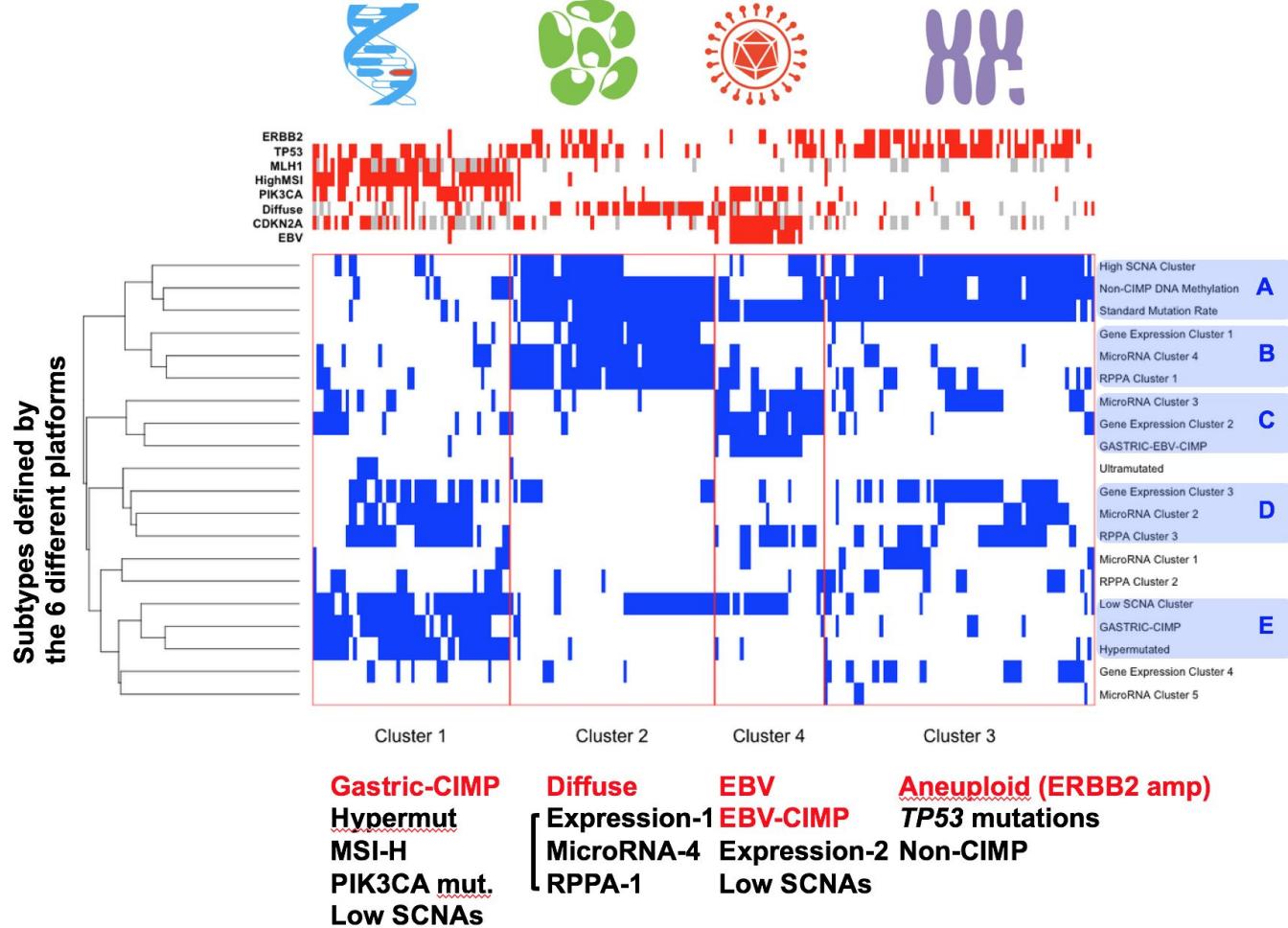
Gastric Cancer - Platform-Specific Clusters

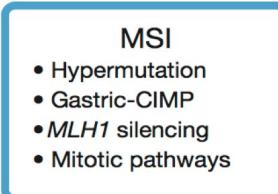
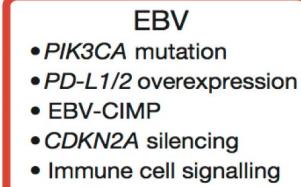
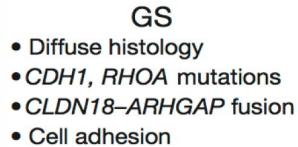
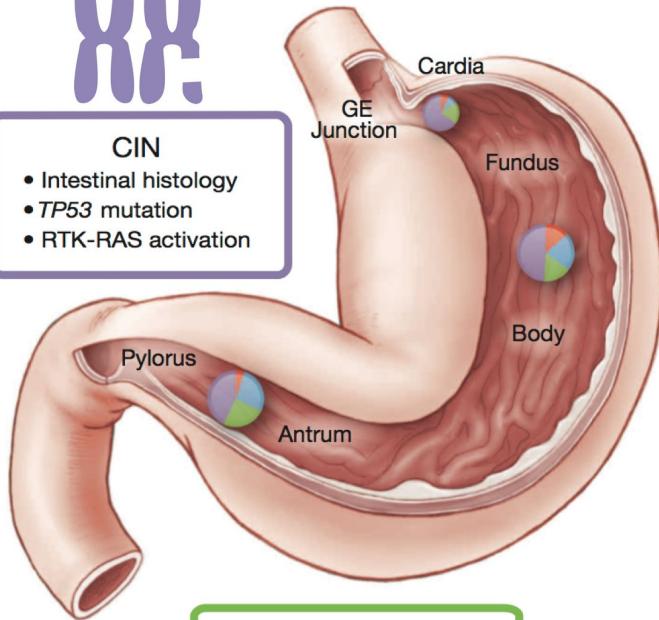
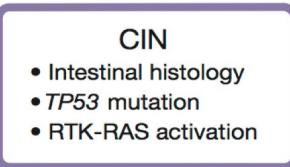


295 Tumor Samples

**Subtypes defined by
the 6 different platforms**

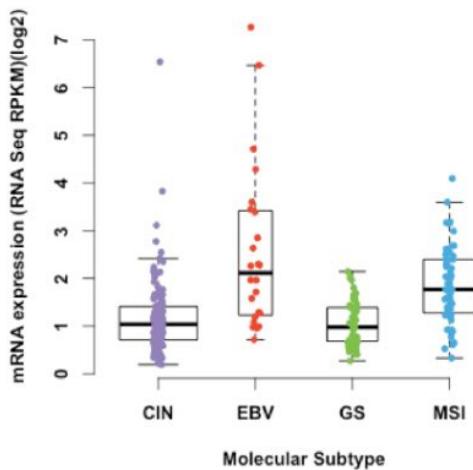






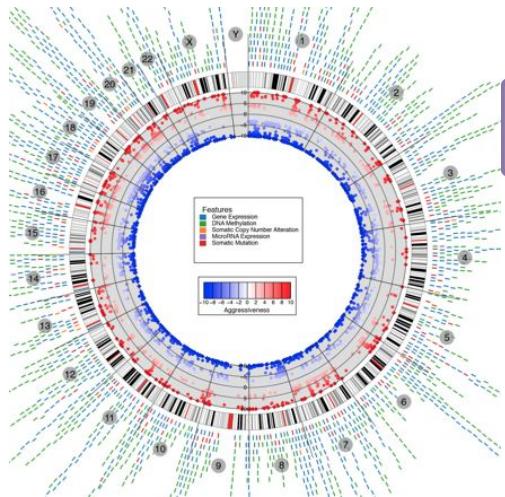
	Spatial	Molecular	Cellular	Tissue
H&E	✓	✗	✓	✓
IHC	✓	✓	✓	✓
Cancer Genomics	✗	✓	✓	✓

PD-L1 / CD274

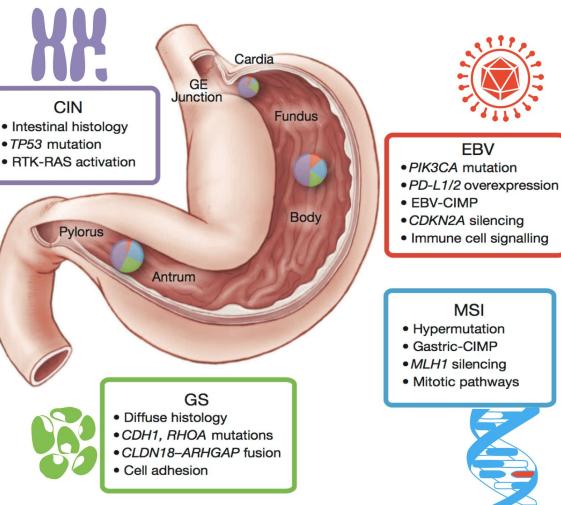


Research on Tumors of the Gastrointestinal Tract by the TCGA

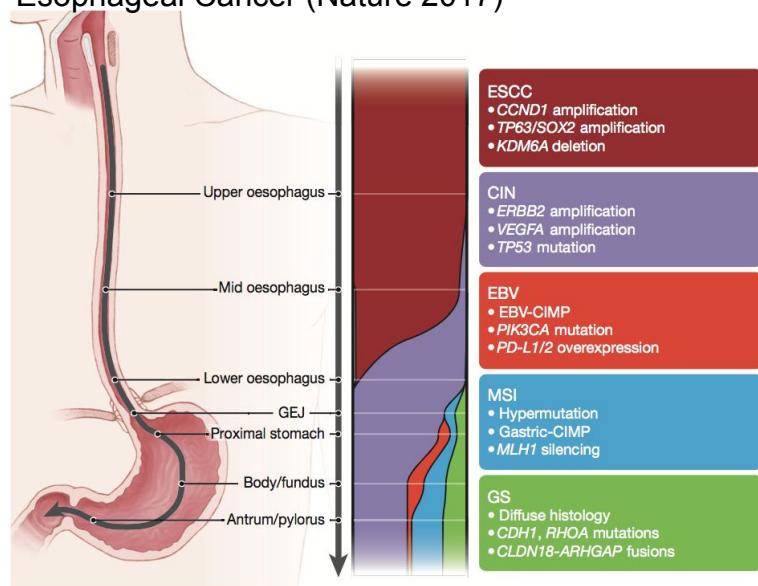
Colorectal Cancer (Nature 2012)



Gastric Cancer (Nature 2014)



Esophageal Cancer (Nature 2017)



TCGA PanCancer Atlas

Cells-Of-Origin

Squamous

Stemness

Gastro

Gyn

Kidney

Pathways / Therapies

MYC

HIPPO

DDR

PI3K

NOTCH

RAS TGFB

Spliceosome

SWI/SNF

Oncogenic Processes

Mutation

Signatures

Germline

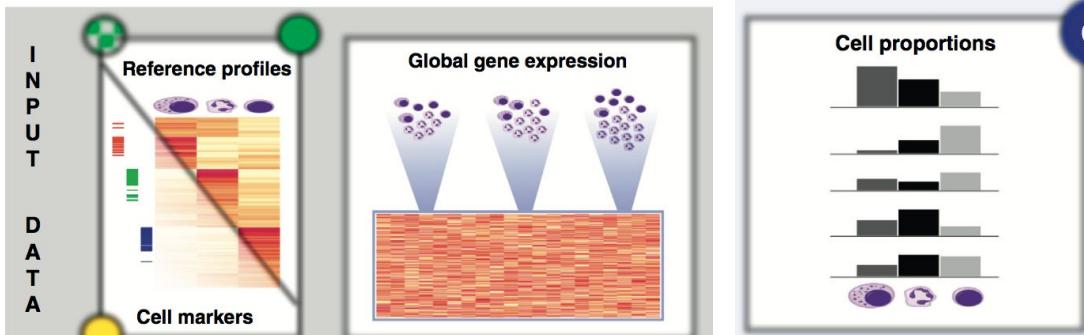
Immune

Drivers /

Essentials

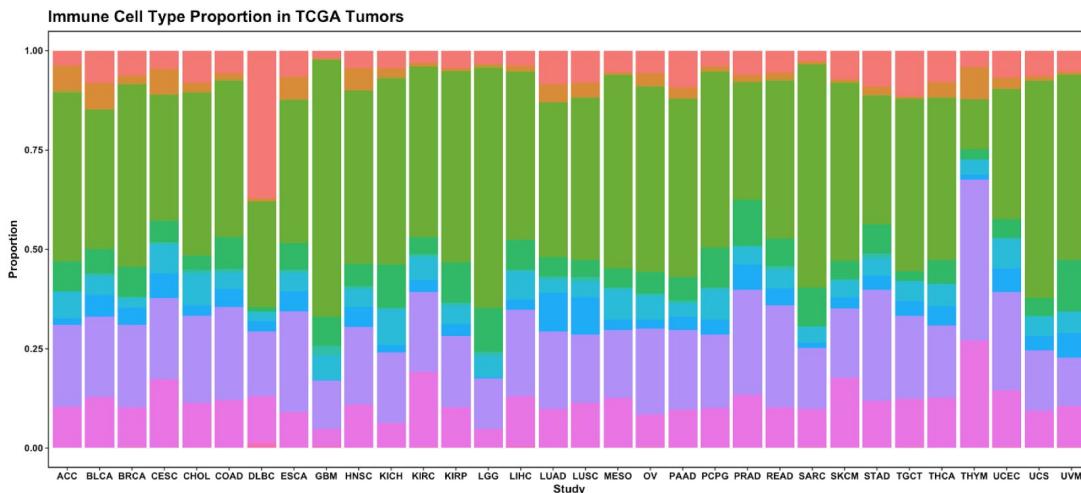
Aneuploidy

Expression deconvolution



Shen-Orr and Gaujoux, *Curr. Op. Immunology* 2013

	Spatial	Molecular	Cellular	Tissue
H&E	✓	✗	✓	✓
IHC	✓	✓	✓	✓
Cancer Genomics	✗	✓	✓	✓



Cell_Type

- B_cells
- Dendritic_cells
- Eosinophils
- Macrophage
- Mast_cells
- Neutrophils
- NK_cells
- Plasma_cells
- T_cells_CD4
- T_cells_CD8
- T_cells_gamma_delta

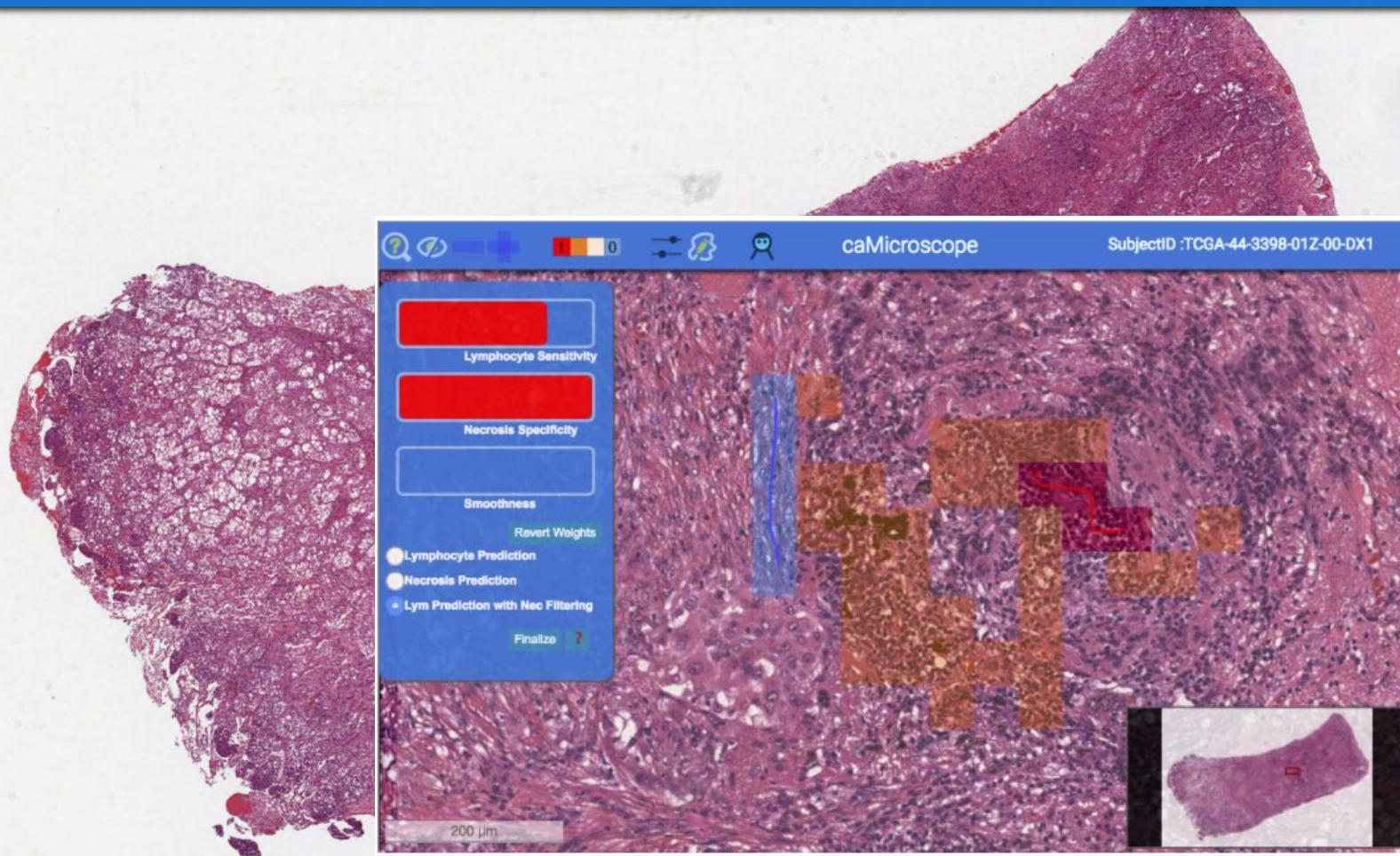
TCGA Immune Response Working Group

1 | 0



caMicroscope

SubjectID :TCGA-44-3398-01Z-00-DX1



	Spatial	Molecular	Cellular	Tissue
H&E	✓	✗	✓	✓
IHC	✓	✓	✓	✓
Cancer Genomics	✗	✓	✓	✓

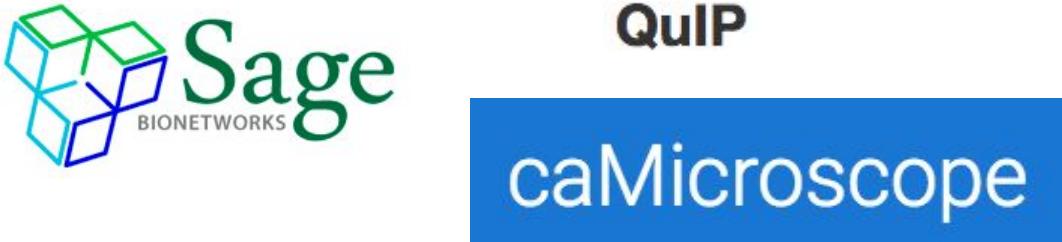
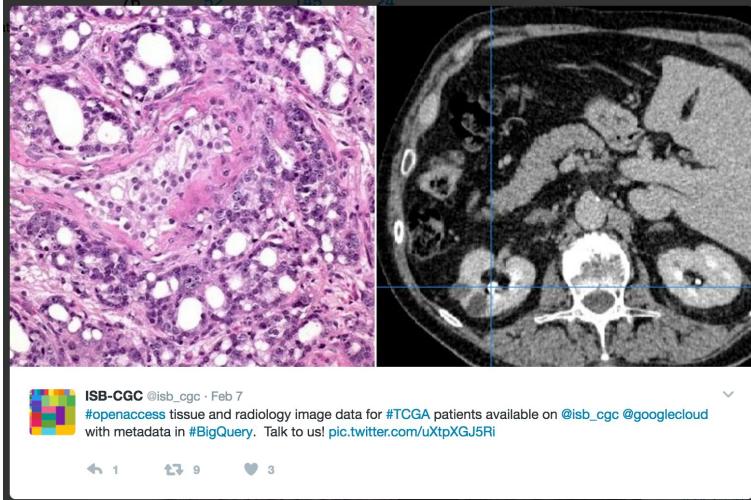
Data Management and Integration

NCI Cloud Resources

Bringing data and computation together to create knowledge that accelerates cancer research and enables precision medicine



Broad Institute Seven Bridges Institute for Systems Biology



Sage BIONETWORKS

QuIP

caMicroscope

CANCER RESEARCH INSTITUTE

iATLAS



Harnessing bioinformatics to speed discovery in cancer immunotherapy.

CANCER RESEARCH INSTITUTE
The Leader in Immunotherapy

Example Query: CTLA-4 Gene Expression



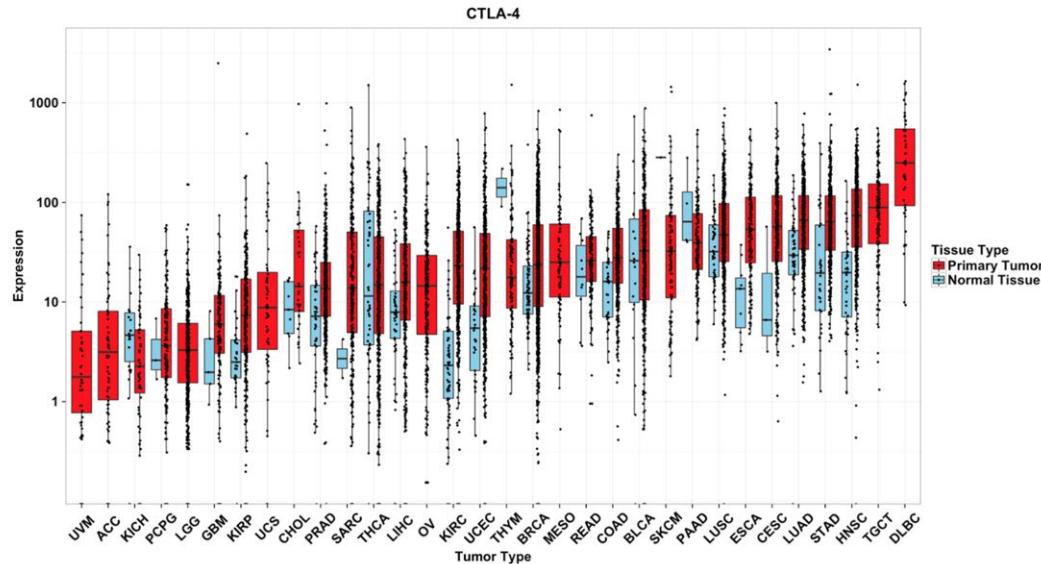
```
select * from [isb-cgc-01-0008:Filtered.EBpp_AdjustPANCAN_RNASeqV2_filtered] where Symbol="CTLA4"
```

Row	ParticipantBarcode	SampleBarcode	AliquotBarcode	SampleTypeLetterCode	SampleType	Study	Symbol	Entrez	normalized_count
1	TCGA-OR-A5JB	TCGA-OR-A5JB-01A	TCGA-OR-A5JB-01A-11R-A29S-07	TP	Primary solid Tumor	ACC	CTLA4	1493	60.1537
2	TCGA-OR-A5LG	TCGA-OR-A5LG-01A	TCGA-OR-A5LG-01A-11R-A29S-07	TP	Primary solid Tumor	ACC	CTLA4	1493	1.365
3	TCGA-4Z-AA7N	TCGA-4Z-AA7N-01A	TCGA-4Z-AA7N-01A-11R-A39I-07	TP	Primary solid Tumor	BLCA	CTLA4	1493	346.102
4	TCGA-DK-A6AV	TCGA-DK-A6AV-01A	TCGA-DK-A6AV-01A-12R-A30C-07	TP	Primary solid Tumor	BLCA	CTLA4	1493	55.1331
5	TCGA-FD-A5BR	TCGA-FD-A5BR-01A	TCGA-FD-A5BR-01A-11R-A26T-07	TP	Primary solid Tumor	BLCA	CTLA4	1493	105.179

Can query
directly from R
using the

bigrquery

library, then do a
plot



Thank you

ISB Team David Gibbs, Sheila M. Reynolds, Michael Miller, Ilya Shmulevich

TCGA Gastroesophageal Cancer Working Group Adam J. Bass (Dana Farber Cancer Institute), Peter W. Laird (Van Andel Research Institute), Ilya Shmulevich (ISB), Francisco Sánchez-Vega (MSKCC), Barbara G. Schneider (Vanderbilt), Toshinori Hinoue (Van Andel),...

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TIL Map Collaboration: Joel Saltz Le Hou, Rajasri R. Gupta, Tahsin Kurc, Vu Nyugen, Dimitri Samaras, Rebecca Batiste, John Van Arnam (Stony Brook), Ashish Sharma, Lee Cooper (Emory), Arvind Rao, Pankaj J. Singh, Alexander Lazar (MDAnderson), Ilya Shmulevich (ISB)

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TCGA Research Network, TCGA Patients and Families

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