

Akoya 5-plex Signature Panels

START WITH ASKING THE RIGHT QUESTIONS



IMMUNO-CONTEXTURE PANEL

Is the tumor “hot” or “cold”?



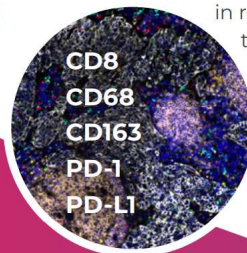
IMMUNE PROFILE PANEL

Where are the immune cells in the TME?



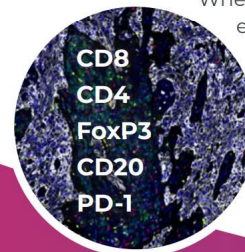
ACTIVATED TIL STATUS PANEL

Are the tumor cells proliferating or lymphocytes activated?



M1/M2 POLARIZATION PANEL

Where are the TAMs in relation to the tumor margin?



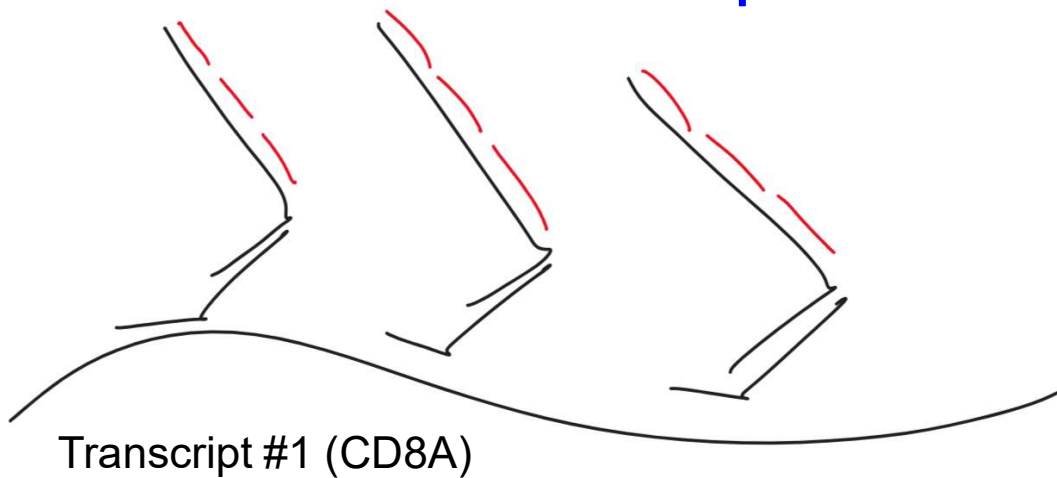
EXHAUSTION PANEL

Where are the exhausted and regulatory T cells?

	A	B	C	D	E	F	G	H	I	J	K	L	M
1	Akoya Signature Panel	CD8	CD68	PanCK	CD3	CD20	PD-1	PD-L1	CD4	CD163	FoxP3	Ki-67	GrzB
2	Immune Profile Human Protein Panel	x	x	x	x	x							
3	Immuno-Contexture Human Protein Panel	x	x	x				x			x		
4	M1/M2 Polarization Panel	x	x				x	x		x			
5	Activated TIL Status Human Protein Panel	x		x	x							x	x
6	Exhaustion Panel	x				x	x		x		x		

PhenoCode™ Signature
5-Plex Base Panels

Demo #2: Multiplexed RNA-FISH (w/ SABER)

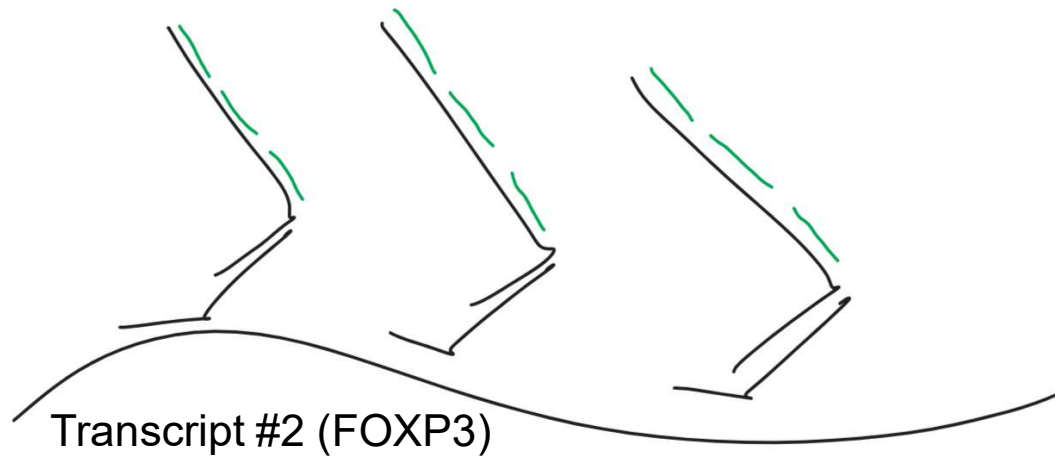


[...]

Transcript #3 (CD4)

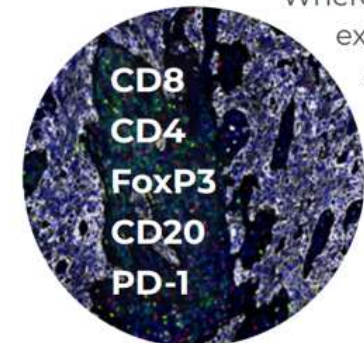
Transcript #4 (CD20)

Transcript #5 (PD-1)



EXHAUSTION PANEL

Where are the exhausted and regulatory T cells?



Demo #2: Multiplexed RNA-FISH (w/ SABER)

PaintSHOP workflow

1. Go to <https://paintshop.io> in a browser, click [Launch PaintSHOP](#)
2. Query database for homology sequences
 1. Click on the [RNA Probe Design](#) tab
 2. Select the [hg38 newBalance \(isoform-flattened\)](#) probe collection
 3. Enter [CD8A](#), [MS4A1](#), [PDCCD1](#), [CD4](#), [FOXP3](#) and click [Submit](#)
 4. Use [Optimize Set](#) → [Trim](#) feature to set max [30](#) probes per target
3. Append a SABER 42mer bridge to the 3' end (unique bridge per target)
 1. Click on the [Appending](#) tab
 2. Under [3' Appending Scheme](#), select [Primer/Bridge/Universal](#) (unique / target), select the [Kishi et al. 2019 Bridges](#) and click [Append](#).
4. [Download](#) your sequences