

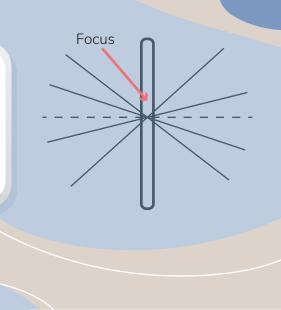
## The challenge

Current technologies for analyzing the spatial organization of cell-surface proteins have limitations:

microscopy and imaging flow cytometry – typically only provide data in one focal plane and have limited multiplexing capacity and throughput.

Imaging-based techniques e.g fluorescence

Immunological activities



**Super-resolution microscopy** – limited in

multiplexing and throughput.

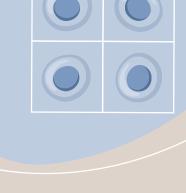
Considered expensive.

Single-cell targeted proteomics using

only provides relative expression levels without any subcellular spatial information.

requires single-cell compartmentalization and

DNA-tagged antibodies e.g CITE-seq -



## Molecular Pixelation (MPX<sup>TM</sup>) is a new technology that overcomes these limitations by capturing 3D spatial maps of surface proteins on thousands of single immune cells simultaneously in a highly

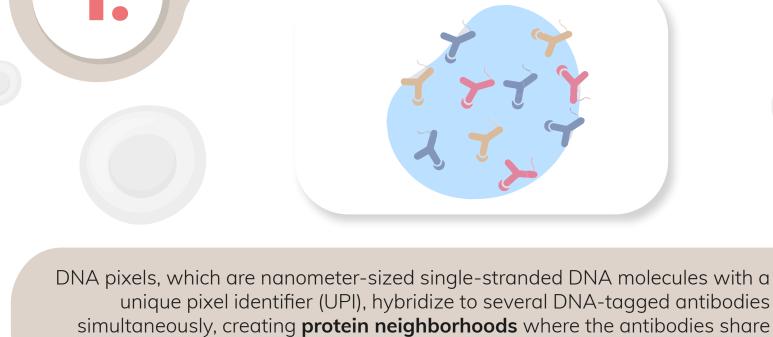
The solution: Molecular Pixelation

How it works:

surface of thousands of PFA-fixed immune cells.

## DNA-tagged antibodies are bound to **proteins** on the

multiplexed manner.



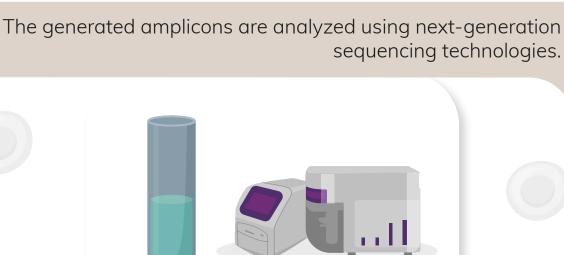
to create a **protein map**.



the same UPI sequence.



A second set of DNA pixels is incorporated, linking the protein neighborhoods



colocalization with unprecedented resolution.

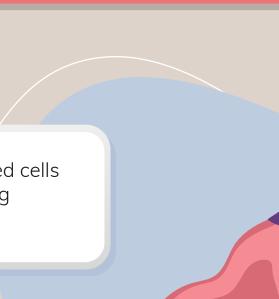
A dedicated software package called Pixelator™ uses sequencing reads to create **graphs** of protein maps. Each graph is a reconstruction of the surface of a cell, quantifying proteins and describing their spatial distribution and protein

with MPX

**Uropods** are the hind part of polarized cells that stabilize and move the cell during migration.

Cytotoxic T cell

Detecting T-cell uropod formation



patient survival.

T-cell uropod

They are essential for cytotoxic T cells to

**infiltrate tumors**, and their formation correlates

to immune checkpoint inhibition efficacy and

**MPX** was used to detect uropod formation in migrating immune cells, providing important insights into T-cell motility, which could help the development of **immune therapies**<sup>1</sup>.

1. Karlsson F, Kallas T, Thiagarajan D et al. Molecular Pixelation: Single cell spatial proteomics by sequencing. bioRxiv doi:10.1101/2023.06.05.543770 (2023).



