COMPLEX CELLULAR PHENOTYPE ANALYSIS



Cuithbert, Carmen, Claresta, Joey Oscar, Lit-Hsin, Guorui



Spatial profiling of cells and tissues

- Spatial multi-omics profiling of human tissues
- High-throughput imagebased phenotypic profiling



Bioimage databases and portals

- Visualization and sharing of cellular and tissue images
- HistoPath Analytics (HPA)
 Platform and ImmunoAtlas



Digital medicine for cancers

- Biomarkers for patient stratification and targeted intervention
- Pathogenicity of mutations



- Tumor-associated antigen prediction
- Xenobiotic-induced toxicity prediction









We have updated cX2 to handle large hyperplexed











Spatial profiling of cells and tissues

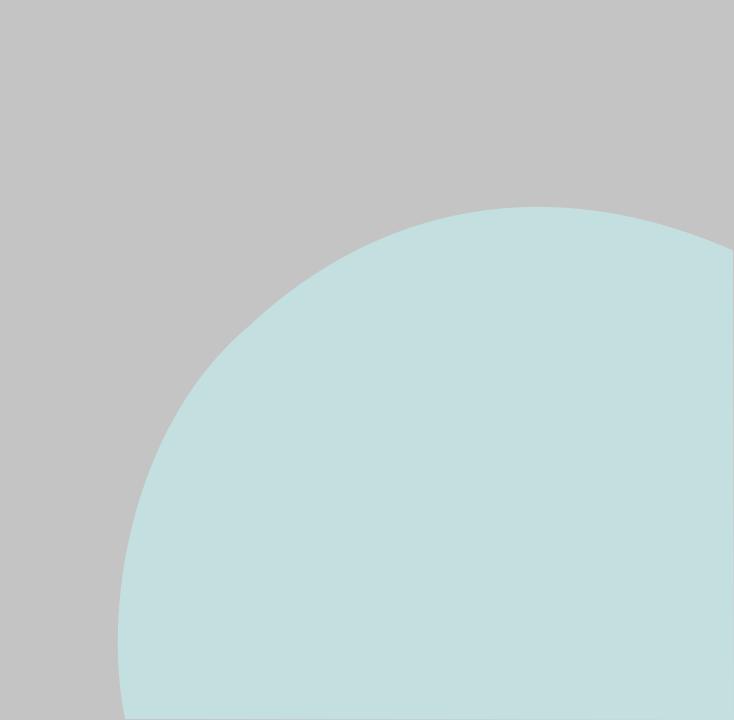
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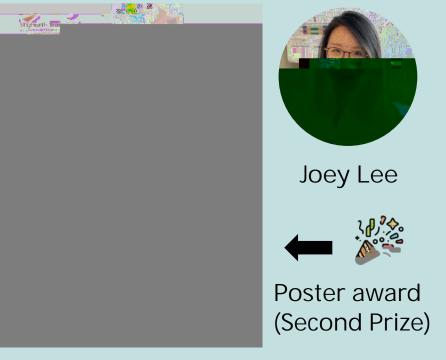


SingHealth Immunopathology Workshop (8th – 9th May 2023)













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Biomarkers for Liver Cancer







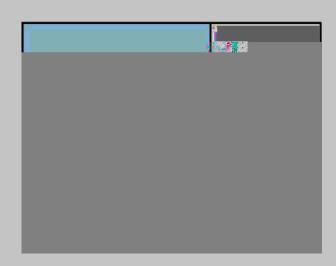
Spatial metabolomics landscape of Liver Cancer









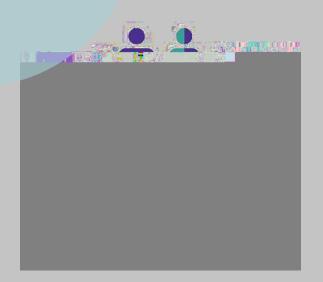




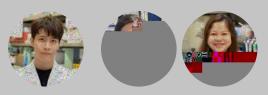
The 5th Singapore Liver Cancer Consortium Scientific Symposium

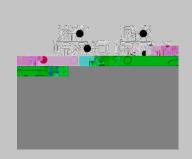
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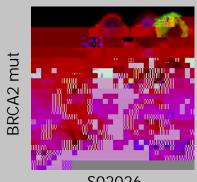


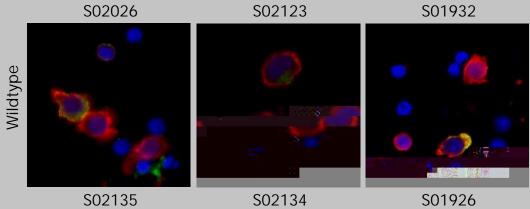
profiling to predict pathogenic mutations

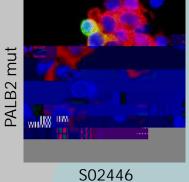


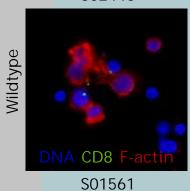


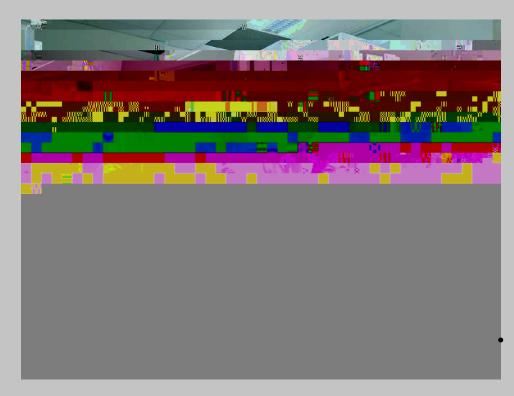














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High-throughput assays and Al models for tumor-associated antigen classification





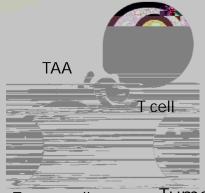












IAF-ICP funded: \$3,897,195 (1 Sept 2022 to 31 Aug 2025)

Tumor cell

Tumor-associated antigens (TAAs)

Viral

Differentiation

Cancergermline

Overexpressed

Mutated

DNA

Protein

Tumor cell

HPV EBV MART-1 CEA

NY-ESO-1 MAGE-A3 **hTRRT EGFR**

BCR-ABL p53



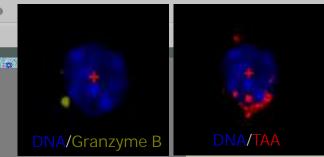
to predict tumor-associated antigens



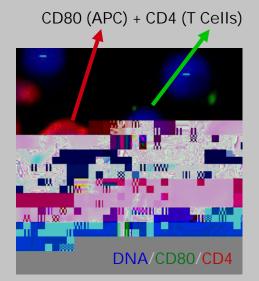


We have developed the assays to image immune cells under different activated conditions

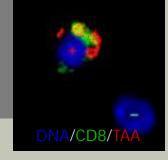
Mouse expressing TAAs

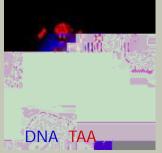


Granzyme B + TAA











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