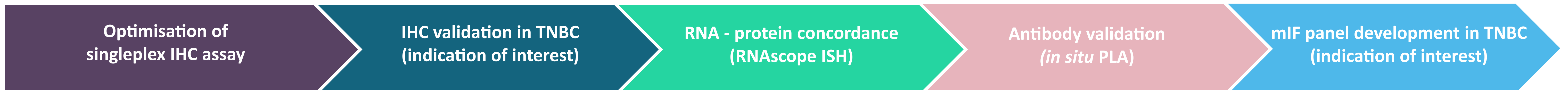


## Background and purpose

Understanding immune-checkpoint ligand distribution within the tumour microenvironment is essential for robust biomarker analysis in triple-negative breast cancer (TNBC), where PD-L1 and PD-L2 expression is highly heterogeneous, and in some cases very low. Reliable detection of these heterogeneous markers underpins actionable immunohistochemistry (IHC) and/or multiplex immunofluorescence (mIF) staining and the downstream quantitative image analysis. However, accurate detection of some of these immune-checkpoint ligands, such as PD-L2, can be technically challenging due to variable antibody specificity and contradictory staining patterns, evidenced by comparing multiple antibodies against the same target.

To overcome this challenge, we designed and conducted a structured, optimisation programme to evaluate commercial PD-L2 antibodies using singleplex IHC, orthogonal validation with RNAScope™ *in situ* hybridization (ISH) and *in situ* proximity-ligation assay (*isPLA*), to enable design and optimisation of a mIF panel to robustly localise PD-L1 and PD-L2 populations in TNBC. This approach enabled the development of a mIF panel capable of delivering reliable, spatially resolved PD-L2 measurements, alongside other key tumour microenvironment markers in a cohort of TNBC samples.

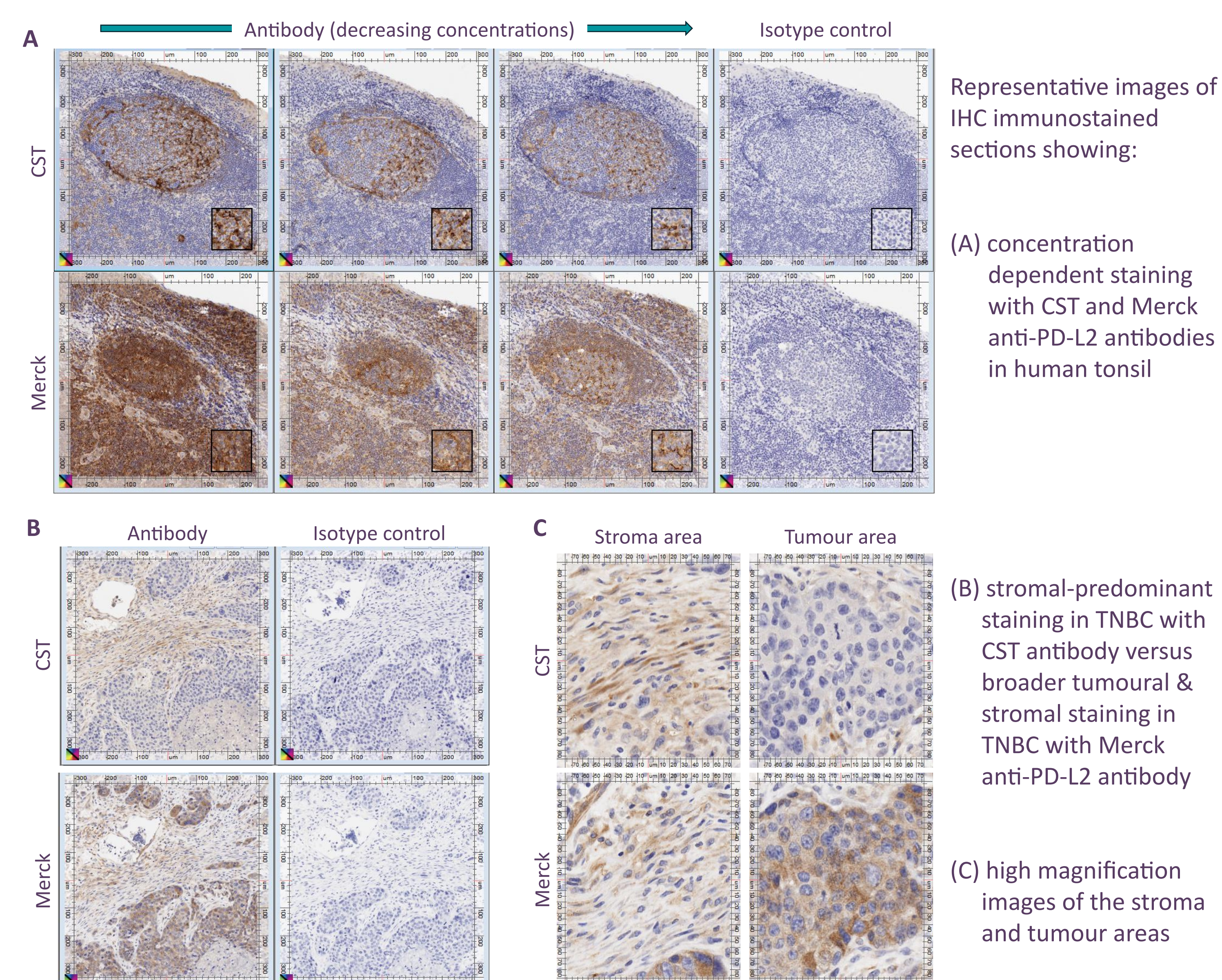
## Study design overview



## Results

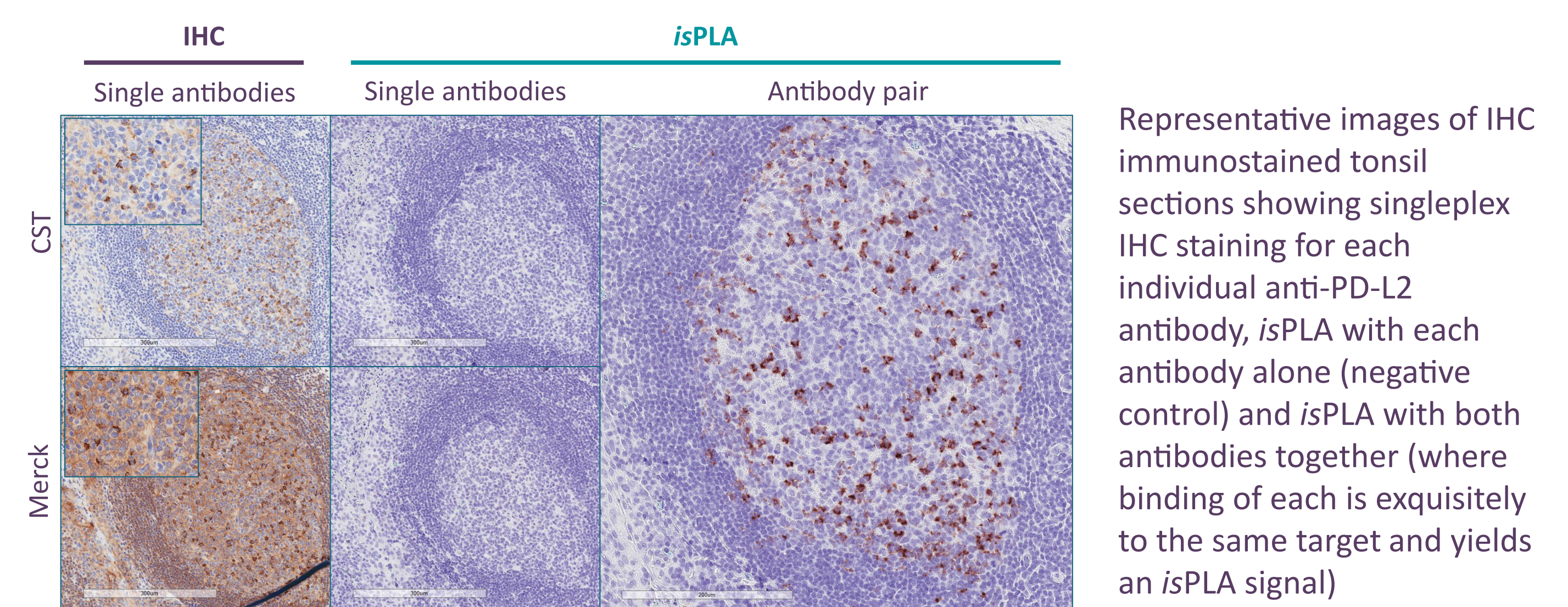
### Optimisation and comparison of PD-L2 antibodies

IHC assay optimisation was performed using a known positive tissue (tonsil) and a cohort of indication specific (TNBC) samples. The immunostaining showed concentration-dependent binding of each antibody to tonsil sections. However, discrepancies between the patterns of immunostaining were evident for the different antibodies in both tonsil and TNBC samples.



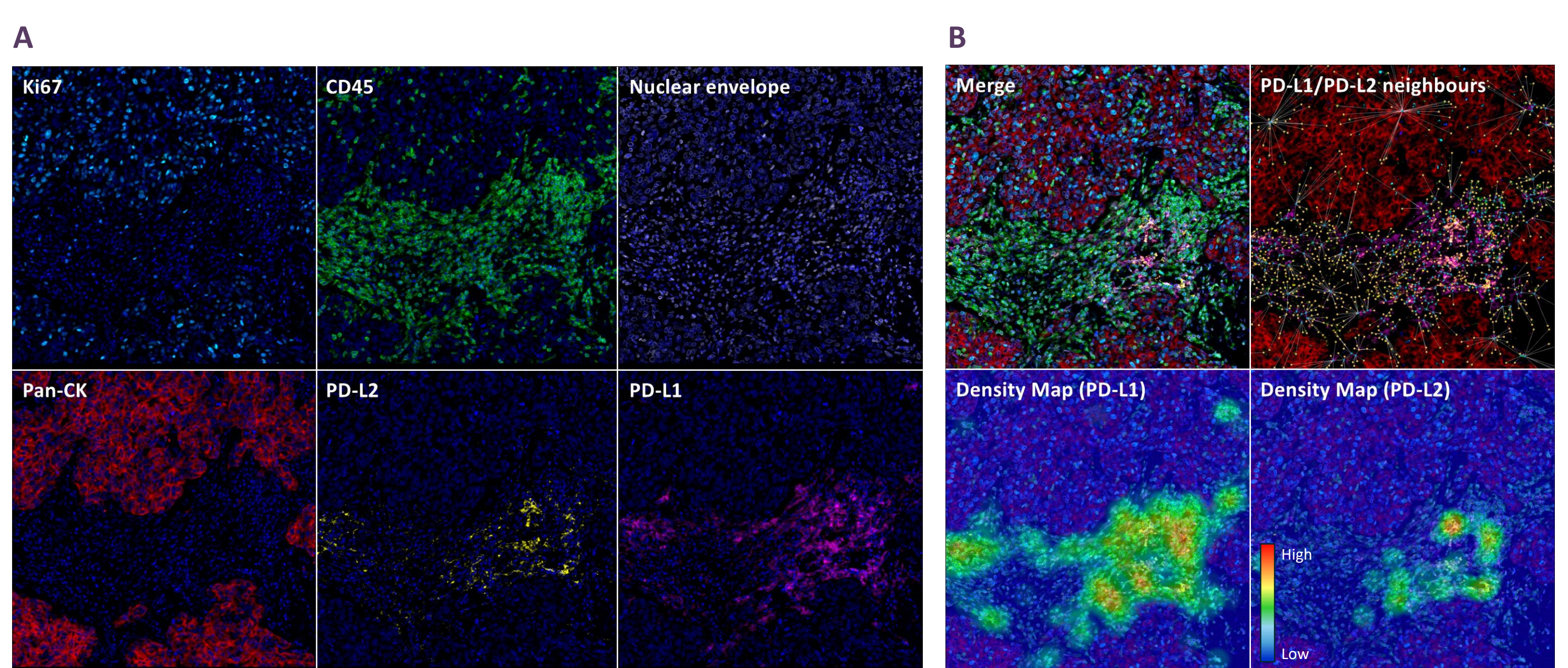
### Antibody staining concordance by *isPLA*

*In situ* PLA technology was used to build additional confidence in PD-L2 protein expression pattern and antibody specificity (see <https://navinci.se/technology/>)



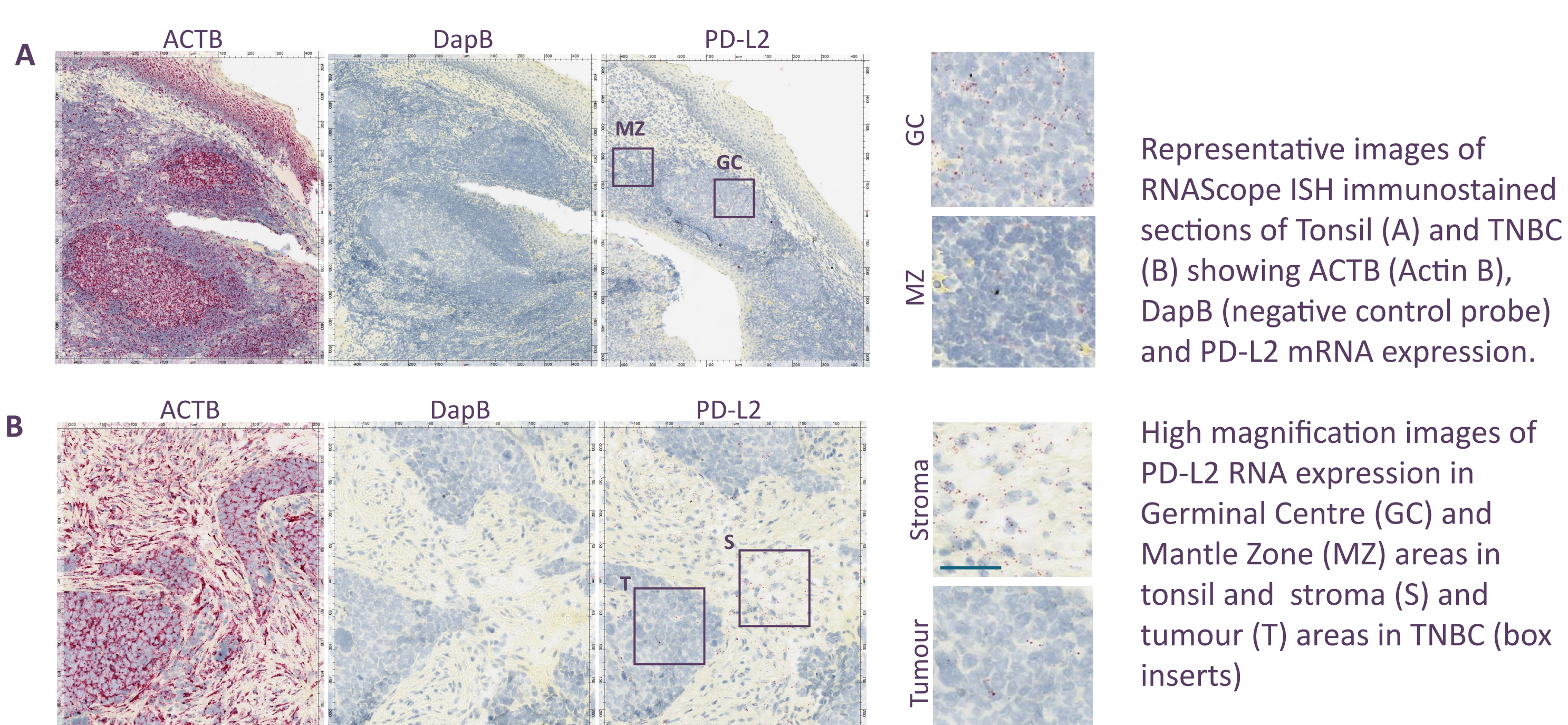
### mIF panel development and spatial image analysis

Iterative optimisation of antibody position, antigen retrieval conditions, and Opal fluorophore assignments resulted in clean multiplex signal without bleed-through (Opal 480, 520, 570, 620, 690, 780). Fluorescent channels display distinct immune (CD45), epithelial (Pan-CK), proliferative (Ki67), nuclear envelope staining and immune-checkpoint (PD-L1/PD-L2) compartments (Panel A). Panel B shows the spatial distribution, cell density and nearest neighbour analysis for PD-L1 and PD-L2 positive cells in TNBC.



### RNA-protein concordance

RNAscope *in situ* hybridisation reveals stromal-predominant PD-L2 RNA expression in TNBC, mirroring PD-L2 IHC patterns with the CST antibody and validating antibody specificity.



### Methods

All immunostaining was performed on a Leica Bond RX Autostainer. For each technique, the optimal assay conditions were determined in-house.

- Two commercial anti-PD-L2 antibodies (CST and Merck)
- Singleplex IHC (DAB endpoint) immunostaining using Leica Bond Polymer Refine kit
- isPLA* (DAB endpoint) using the NaveniBright BOND RX HRP kit
- RNAscope *in situ* hybridisation and Leica BOND Red detection kit
- Selection of optimal anti-PD-L2 antibody for development of multiplex immunofluorescence panel using Akoya classic reagents, including MOTIF opal fluorophores and Akoya HRP polymer
- Spatial image analysis using HALO® Image Analysis software. HighPlex FL algorithm and rainbow forest tissue classifier were used for cell segmentation, marker detection and tissue classification

## Conclusion

- ✓ The development approach successfully identified the **optimal PD-L2 antibody** for Triple-Negative Breast Cancer samples
- ✓ Multi-modal testing confirmed **high specificity, low background, and true biological signal**
- ✓ Robust multiplex immunofluorescence development workflow and control generated a final **validated, reproducible and fit-for purpose** mIF panel for spatial analyses of TNBC
- ✓ HALO phenotype mapping provides **high-resolution spatial insight**, supporting clinical translation