

Optimizing Imaging Flow Cytometry with BD CellView™ Technology for Higher-Parameter Data Generation

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Abstract

Background: Imaging flow cytometry merges quantitative flow cytometry with high-resolution single-cell imaging. BD FACSDiscover™ S8 and A8 systems, featuring BD CellView™ Image Technology, enable real-time imaging that expands data dimensionality beyond fluorescence. BD CellView™ captures spatial and morphological features, supporting subcellular classification and higher-parameter analysis. However, standardized optimization guidelines remain limited.

Objective: To optimize fluorochrome selection and imaging parameters using BD CellView™ technology, demonstrated through a 12-color spectral panel with 2-color imaging for enhanced T-cell analysis.

Methods: PBMCs were stained with 55 CD4-conjugated fluorochromes and acquired on a BD FACSDiscover™ S8 using default imaging gains to rank brightness and spillover across three channels. A 12-color panel targeting surface and intracellular markers (NucSpot 488, T-bet-RB705, CD3-BUV805, CD4-APC-H7, CD8-BUV496, CCR7-BUV395, CD45RA-BV421, CD27-R718, CD28-BUV615, PD-1-RY586, TIM-3-BV480, LAG-3-AF647) was used to stain resting and stimulated PBMCs (CD3/CD28 beads, 48h). Cells were fixed, permeabilized, and resuspended in wash buffer prior to acquisition. FMO controls validated imaging detection of NucSpot and T-bet, and confirmed “Correlation” as a parameter for T-bet trafficking.

Results: Stimulation increased PD-1 and T-bet co-expression in CD8+ T-cell subsets: Naïve (5.86%→88.1%), CM (23.8%→93.6%), EM (42.8%→86.9%), EMRA (36.1%→87.5%). Imaging revealed T-bet localization patterns—cytoplasmic, nuclear, or both—with nuclear T-bet predominating post-stimulation (73.7%). Imaging-enhanced analysis identified 48 functional T-cell subsets, tripling the resolution of traditional analysis without imaging using the same panel.

Conclusion: This study provides a practical framework for optimizing BD CellView™ imaging and fluorochrome assignment. Integrating spectral and image-derived parameters enables deeper functional classification and higher-parameter data generation, advancing cytometric analysis.

Methods

PBMCs were isolated from whole blood obtained from healthy volunteer donors using Ficoll™ gradient centrifugation.

Isolated PBMCs were cultured with or without Dynabeads Human T-cell Activator CD3/CD28 (Thermo Fisher Scientific) for 48 hours. CD4 stained samples were stained fresh.

Freshly isolated PBMCs were surface stained with single CD4-conjugated antibodies (Table 1). Resting and activated cells were surface stained with a cocktail of antibodies (Table 2).

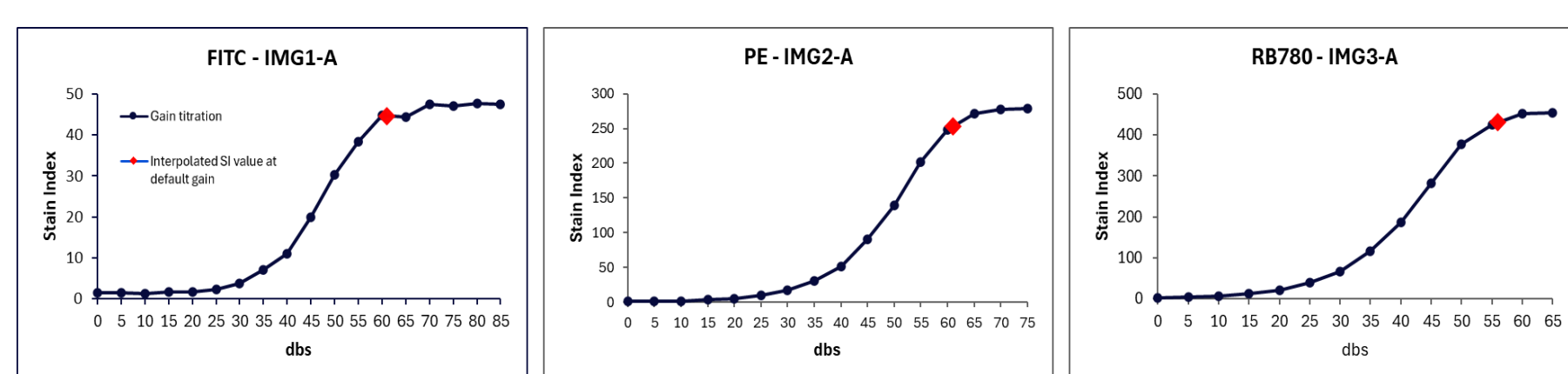
Upon surface staining, resting and activated cells were fixed and permeabilized with BD Pharmingen™ Transcription Factor Buffer Set. CD4-stained cells were acquired right after surface staining (no fixation/permeabilization).

Cells were then intracellularly stained with BD Horizon™ RB705 Mouse Anti-T-bet antibody.

Cells were resuspended in wash buffer containing NucSpot 488 (resting and activated cells) or wash buffer only (CD4-stained cells) and stored on ice, protected from light, until acquisition on the BD FACSDiscover™ S8 Cell Sorter

Data were analyzed using FlowJo_v10.10.0

Gain Titration and Interpolation of Default Gains



A gain titration of the three imaging channels was performed using freshly isolated PBMCs single stained with different CD4-conjugates (FITC for IMG1, PE for IMG2 and RB780 for IMG3). Samples with corresponding CD4-conjugates were acquired at intervals of 5db until reaching detector saturation. Stain index at each gain was calculated and gain titration curve generated for each imaging channel. Corresponding stain indices at default gains were calculated by interpolation (red dot).

MFI Ranking and Spillover Assessment on Imaging Channels



Table 1. CD4 – conjugated antibodies. Freshly isolated PBMCs were single stained with CD4-conjugated antibodies (indicated in table 2). Samples were acquired on the FACSDiscover™ S8 set up at its default optimal gains. Histogram overlays show Mean Fluorescence Intensity (MFI) of each fluorochrome in each imaging channel. Fluorochromes were ranked from highest to lowest MFI. Fluorophores marked with asterisk (*) are early candidates from the BD new dye technology team.

2) Design and Optimization of a 12-color spectral + 2-color Imaging Flow Cytometry Panel

Spectral + Imaging Panel Design

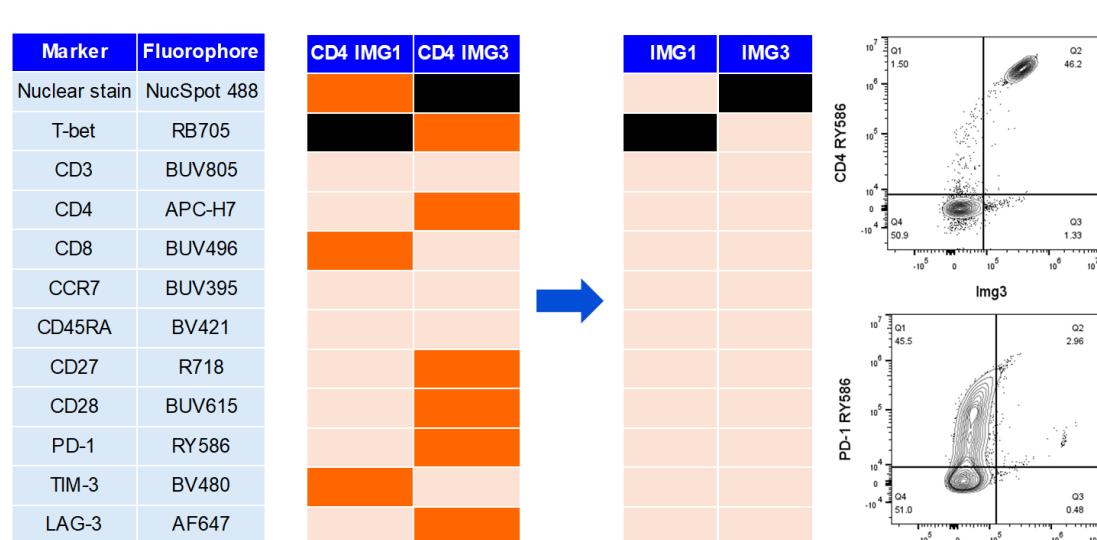
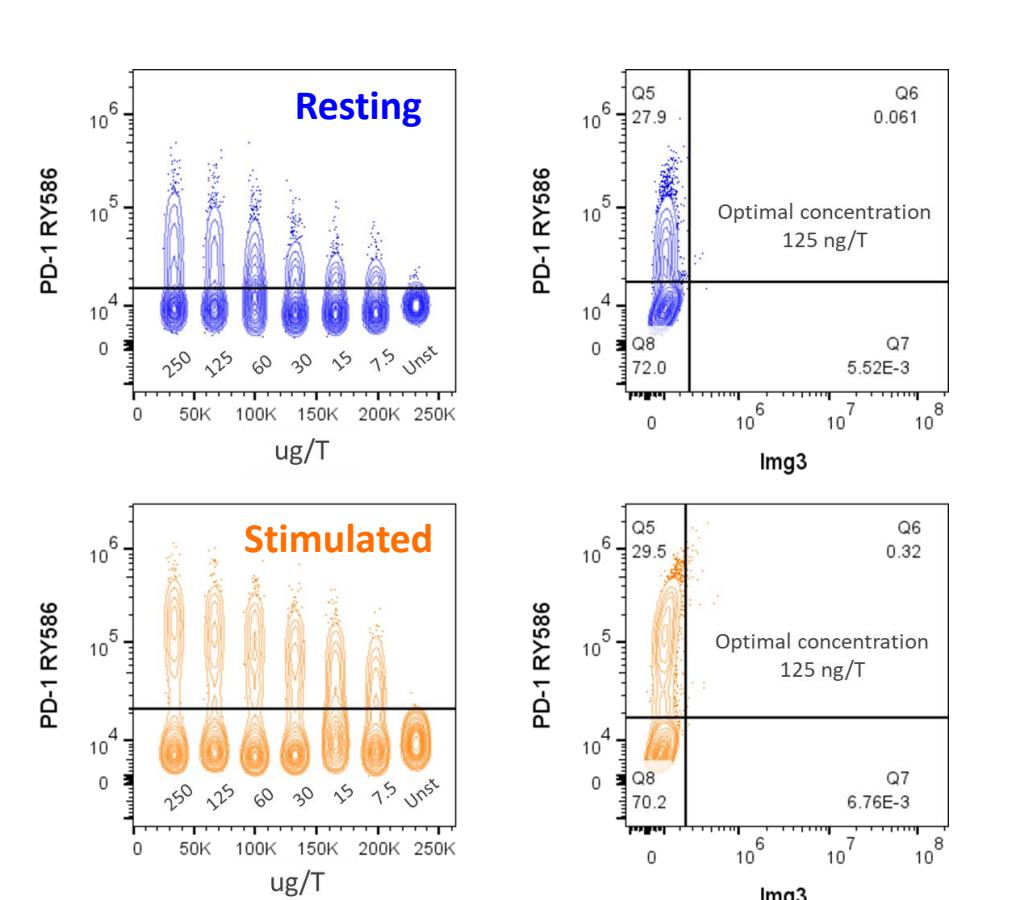


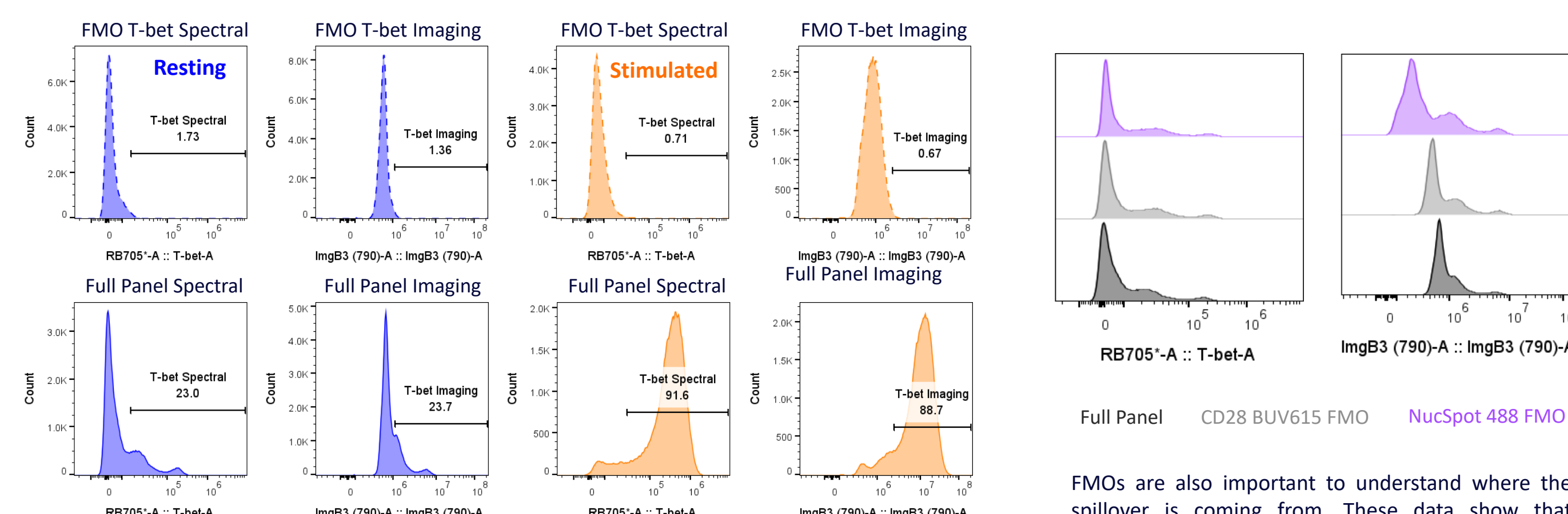
Table 2. 12-color fluorescent + 2-color imaging panel to localize T-bet in activated lymphocytes. Panel was designed by selecting fluorochromes with no or mild spillover into the imaging channels based on CD4 stain (Results 1D). Fluorochromes with mild spillover into imaging channels were assigned to lower density antigens. Table on the right represents imaging spillover expectation after antigen matching (expression level vs CD4) and reagents titration.

Reagent Titration and Spillover



Reagents titration is critical for determining optimal concentration for best resolution and to minimize spillover. It is important to assess spillover in all the experimental conditions since marker expression may change.

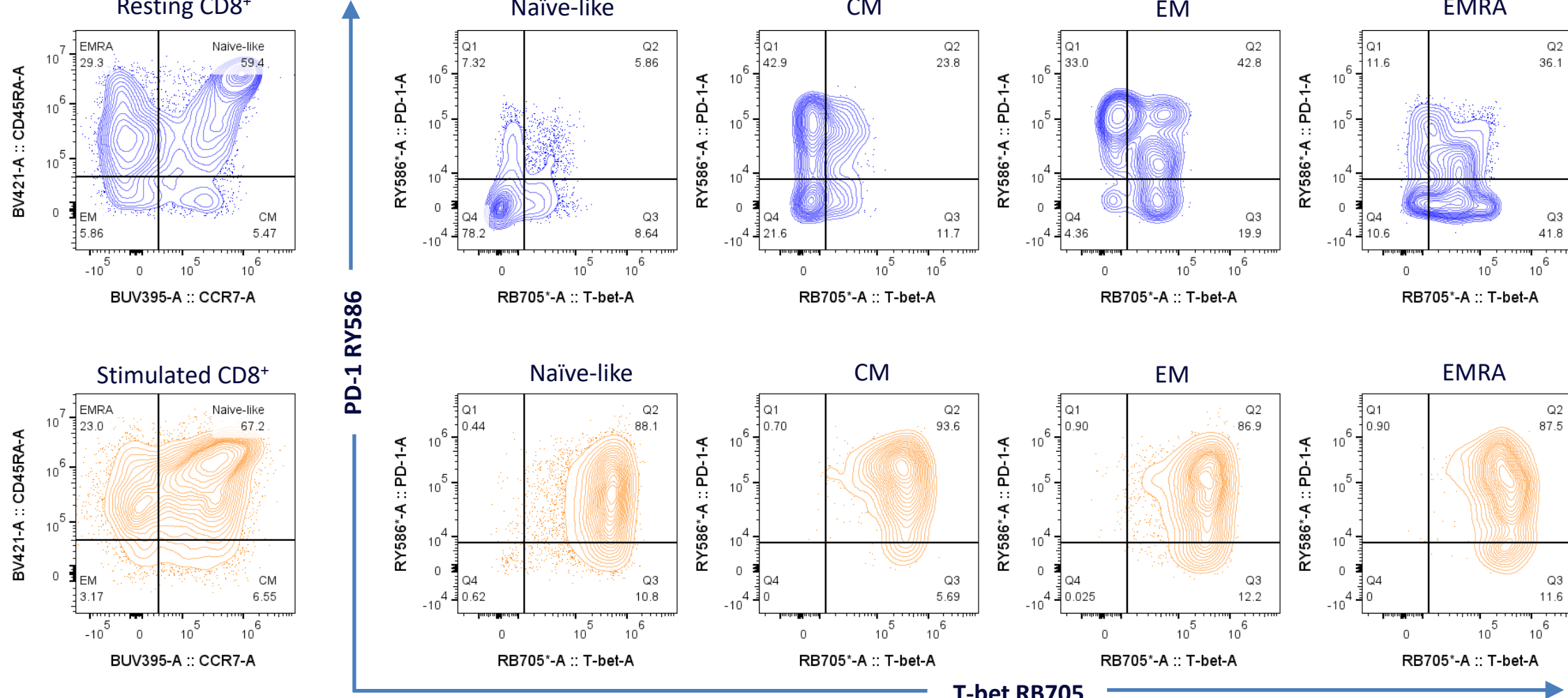
Evaluating Biological Impact Due to Spillover



FMOs are critical to assess spillover impact into imaging channels. FMOs are used to set gates for T-bet, both spectral and imaging detectors. Spectral data are unmixed and represent the biological standard. We can see spillover in IMG3 as showed by the high negative population. However, based on FMO gates, we can still resolve a dim and bright population. This data confirm we were able to minimize spillover and maintain resolution.

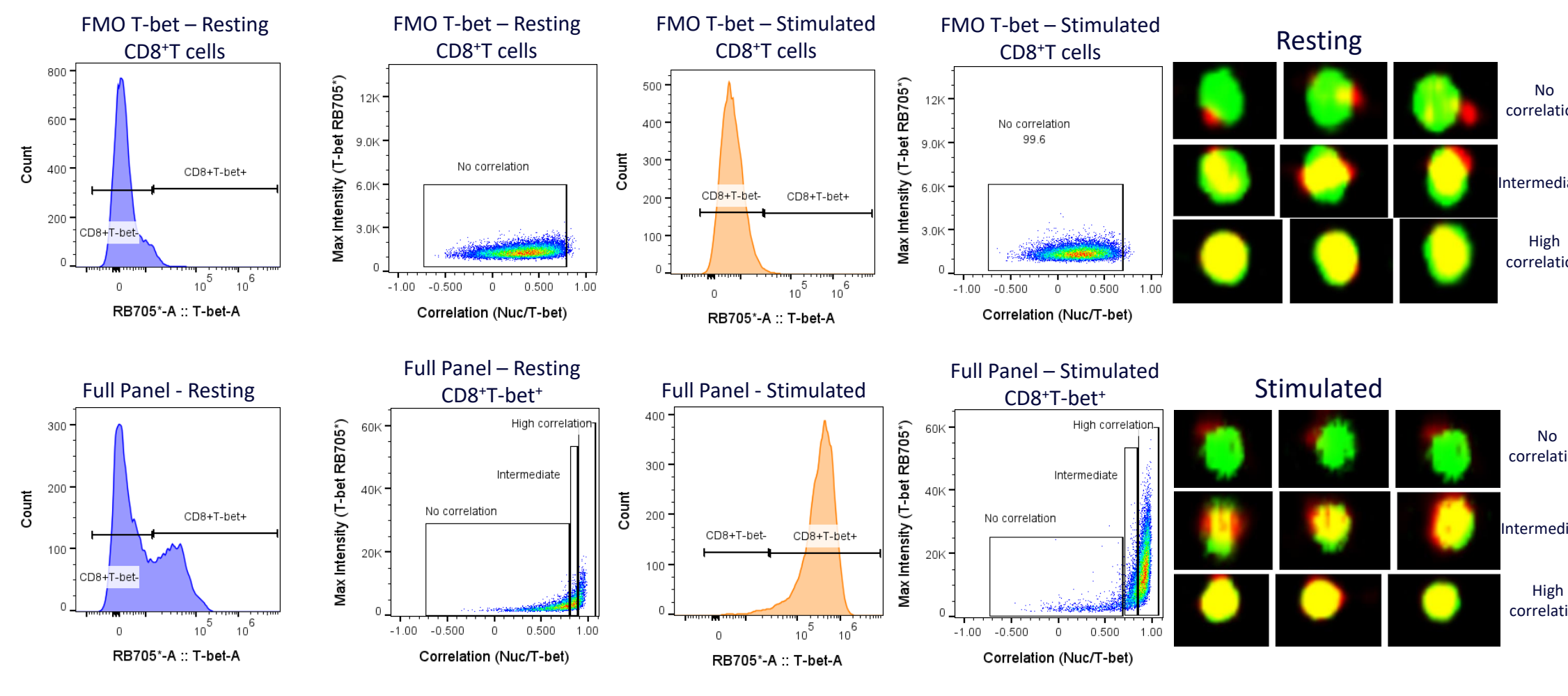
3) Integrating Spatial Localization and Expression Levels of T-bet with Flow Cytometry

T-bet and PD-1 Expression in CD8+ T cells



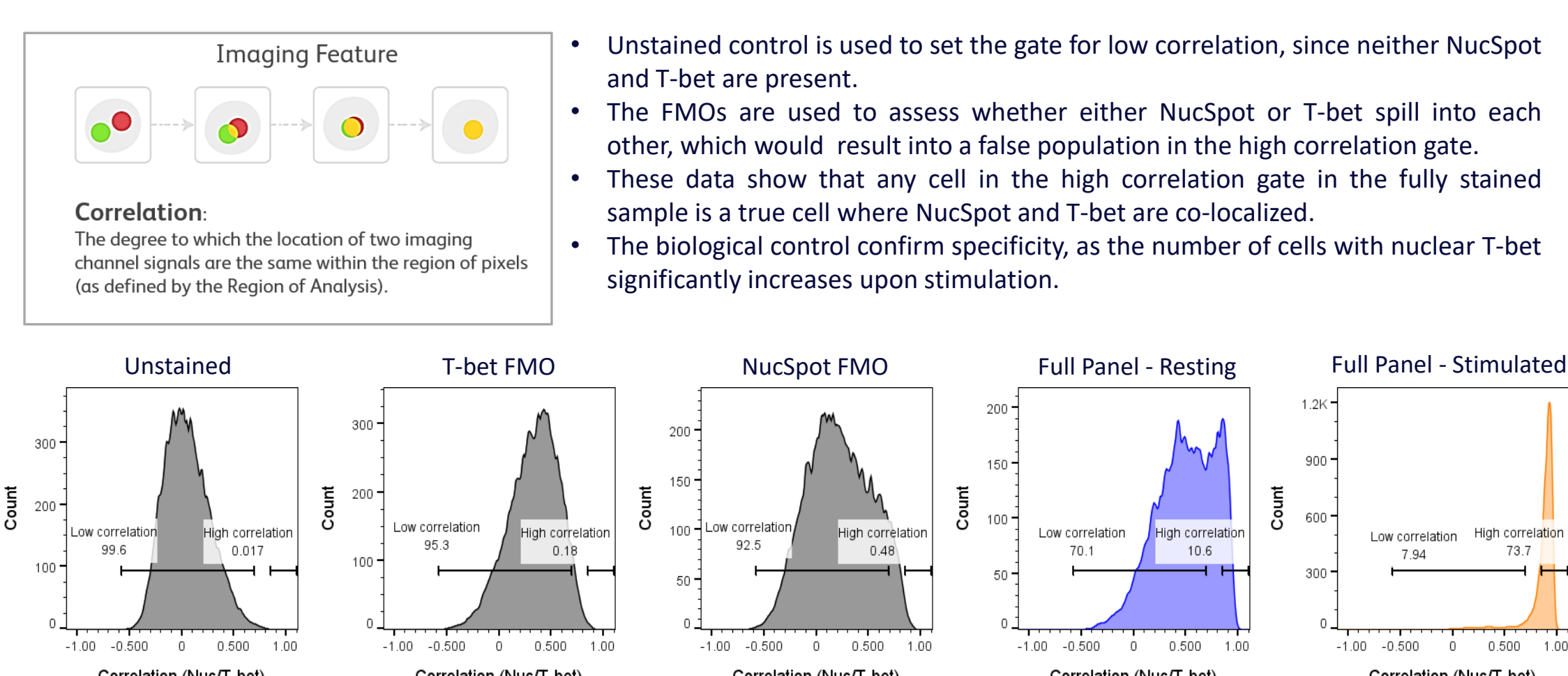
The panel successfully indicates the differential expression levels of PD-1 and T-bet in all evaluated CD8+ T-cell subsets. Upon activation, CD8+ T cells present upregulation of both PD-1 and T-bet increasing the % of double-positive subsets.

T-bet Spatial localization within CD8+ T cells

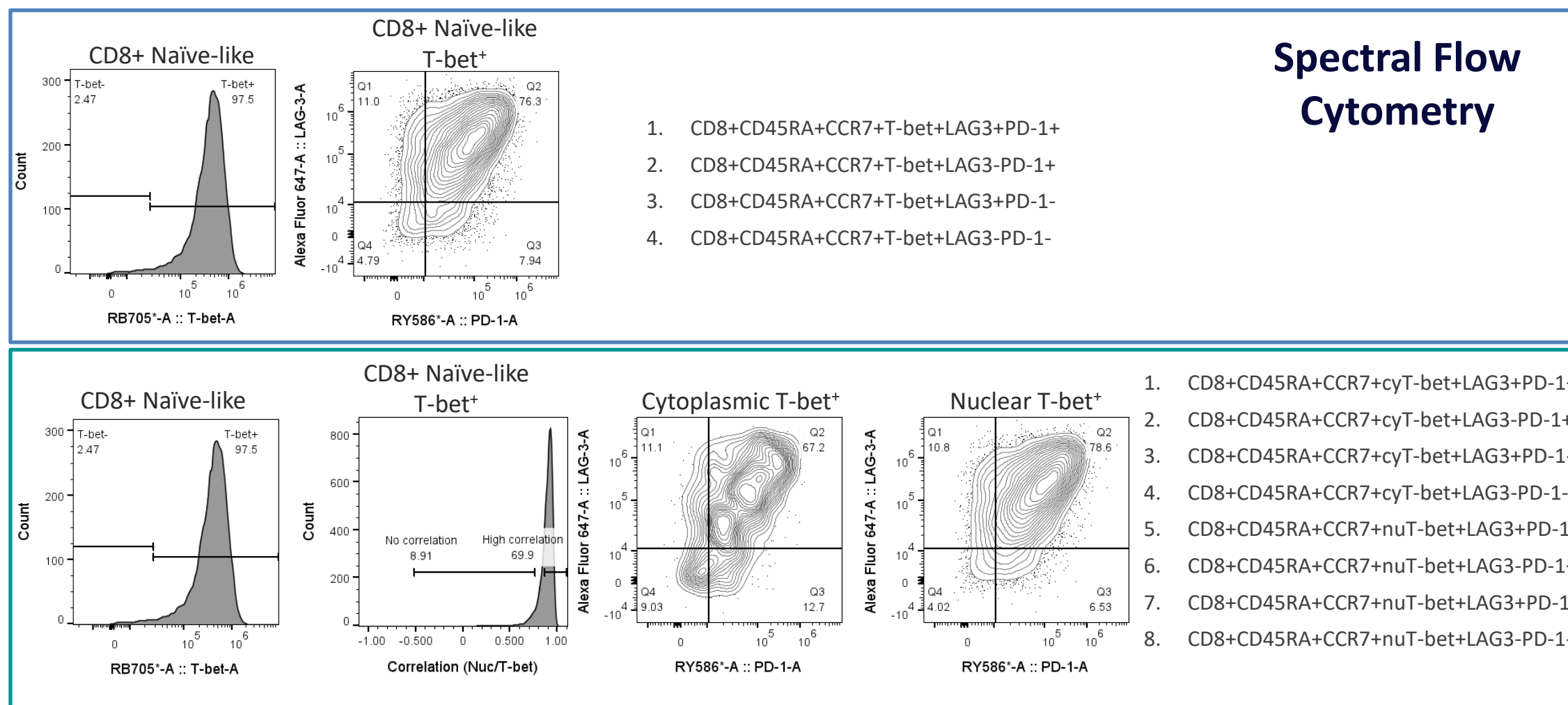


Cell Wall images confirm T-bet localization based on Correlation (Nuc/T-bet). No correlation gate shows T-bet localization within the cytoplasm; Intermediate correlation gate shows T-bet localization both in the cytoplasm and in the nucleus; High correlation gate shows T-bet localization within the nucleus. **Red** T-bet in the cytoplasm **Yellow** T-bet in the nucleus **Green** Nucleus (NucSpot)

Measurement of the “Correlation” Imaging Feature



More subsets can be detected when adding spatial biology



Spectral Flow Cytometry

1. CD8+CD45RA+CCR7+cyT-bet+LAG3+PD-1+
2. CD8+CD45RA+CCR7+cyT-bet+LAG3+PD-1+
3. CD8+CD45RA+CCR7+cyT-bet+LAG3+PD-1+
4. CD8+CD45RA+CCR7+cyT-bet+LAG3+PD-1+
5. CD8+CD45RA+CCR7+cyT-bet+LAG3+PD-1+
6. CD8+CD45RA+CCR7+cyT-bet+LAG3+PD-1+
7. CD8+CD45RA+CCR7+cyT-bet+LAG3+PD-1+
8. CD8+CD45RA+CCR7+cyT-bet+LAG3+PD-1+

Real-Time Imaging + Spectral Flow Cytometry

Conclusions

- Gain titration of imaging channels demonstrate that default gains are the optimal gains for best Stain Index.
- Performance (Stain Index) of a given fluorochrome is likely to be different in the spectral and in the imaging detectors.
- It is crucial to titrate reagents to ensure best resolution and to evaluate spillover.
- FMOs are critical to assess spillover impact into imaging channels.
- Spillover should be assessed using data plots, since signal visualization on the wall depends on post-acquisition imaging adjustments.
- We were able to visualize T-bet translocation upon PBMCs stimulation by applying best practices in panel design and lessons learnt in Imaging Cytometry.
- T-bet translocation was successfully visualized both on the Cell Wall and using imaging parameters (Correlation).
- When adding imaging to spectral cytometry more subsets can be visualized.

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