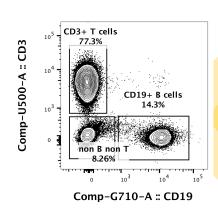
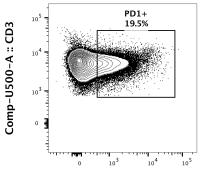
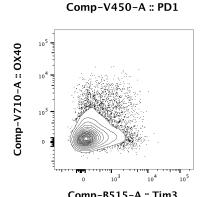
### ANTIGEN CATEGORIZATION

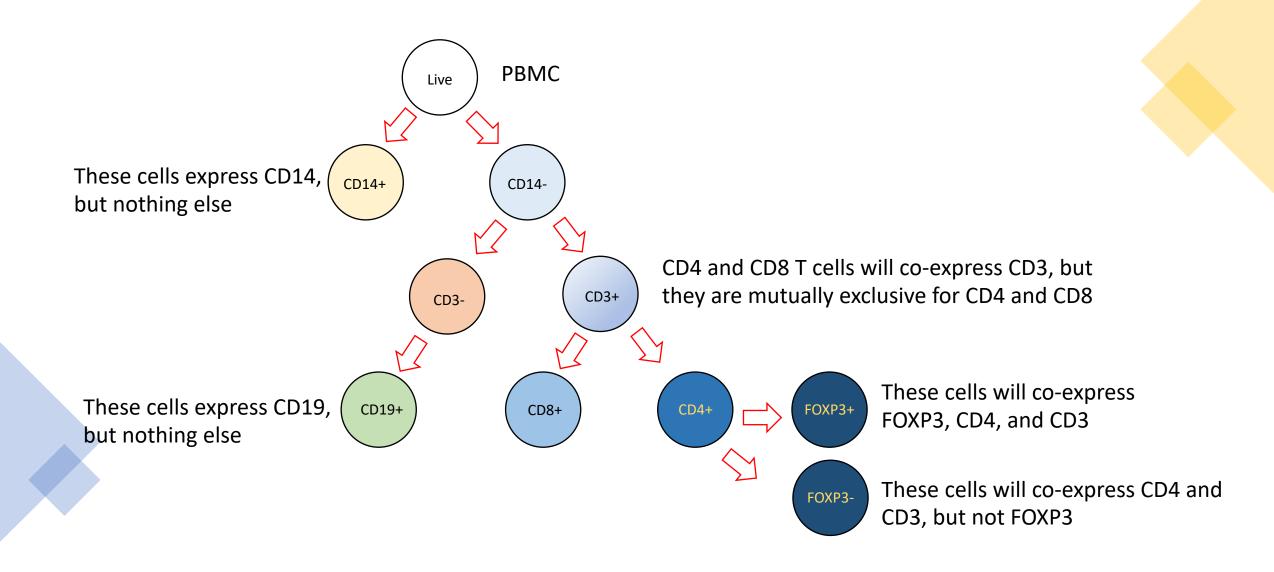
- Marker expression usually comes in three different patterns:
- On/Off expression, usually lineage markers (CD3, CD4, CD19)
- Intermediate or continuous expression patterns (CD45RA, CD38, CD57, cytokines, many more)
- **Dimly** expressed or rare markers
  - ➤ Choose detectors that receive low spillover and pair with bright fluorochromes







### Use a gating tree to assess co-expression of markers



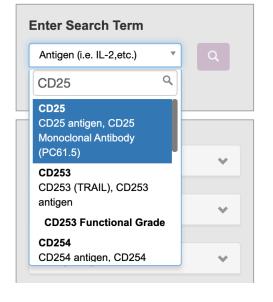
#### Assess commercial availability of conjugated fluorochromes

- Fluorofinder online resource/database that shows all commercially available antibody-fluorochrome conjugates
  - >www.fluorofinder.com
  - > Also has an online panel builder and spectra viewer
- Use filter to narrow results for
  - ➤ Target species
  - ➤ Company (or you will get false positives from Biorbyt)
  - > Fluorochrome (if you want)
  - ➤ Clone (if you want)

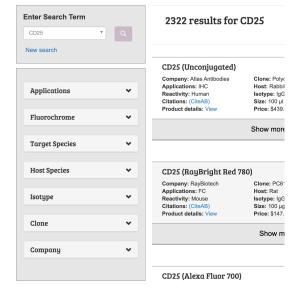
### Using Fluorfinder



Click on "Search Antibodies"

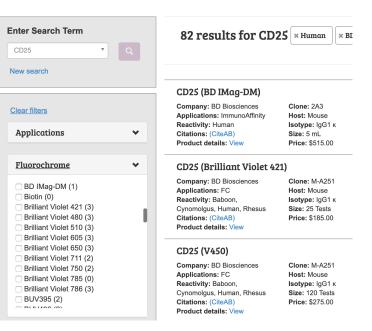


Enter antigen of interest



Narrow down results using the filters

# Scroll through fluorochromes to check availability

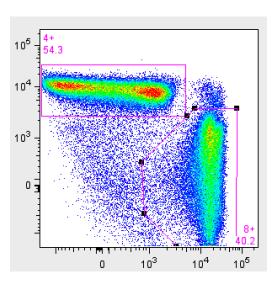


### Use the SSM and gating tree to guide antibodyfluorochrome pairings

- Assign bright markers (or highly/broadly expressed antigens) to channels that contribute little spillover
- >Assign critical or dimly expressing makers to channels that accept little spillover
- ➤ Place mutually exclusive combinations on channels with high spillover/spread values
- ➤ Use the spillover/spread matrix and gating tree to guide placement of co-expressing markers

	B515	B610	B660	B710	B780	G575	G610	G660	G710	G780
B515	0	0.668	0.638	0.763	0.319	0.237	0.236	0	0.649	0.0928
B610	0.251	0	4.66	5.11	1.41	1.35	3.71	2.08	4.76	0.659
B660	0.918	2.02	0	7.09	1.98	2.59	1.04	3.25	6.1	0.977
B710	0.848	0.677	2.73	0	4.34	2.05	0.205	1.53	10.1	3.62
B780	0.713	0.538	0.637	1.34	0	0.537	0.335	0.342	1.22	2.38
G575	0.203	4.1	2.3	2.87	0.676	0	2.1	2.01	2.95	0.55
G610	0.162	5.08	4.33	6.28	1.54	2.17	0	3.71	6.55	1.39
G660	0	0.439	4.38	6.35	2.35	1.95	0.506	0	7.69	2.06
G710	0.362	1.38	3.54	17.9	5.04	5.27	0.953	4.41	0	6.3
G780	0	0.304	0.383	0.598	6.64	0.476	0.331	0.43	0.824	0
R660	0	0.218	1.19	1.49	0.62	0.622	0.376	1.1	1.76	0.561
R710	0	0	0.465	1.74	0.844	0.511	0.14	0	2.07	0.884

Ex: CD4 and CD8 on G710 and B710 are still distinguishable as both markers are mutually exclusive

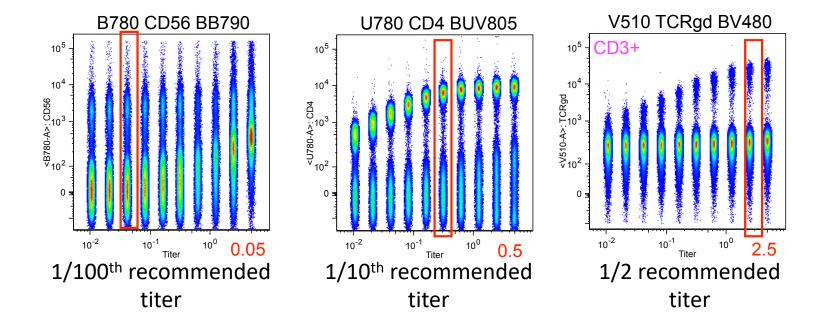


# ANTIBODY TITRATION(1)

- ALWAYS TITRATE!!!!
  - > Every clone will behave differently
  - ➤ Manufacturers vial at different concentrations
- Titrate under the conditions in which the antibody will be used in the full panel
  - ➤ i.e. surface antibodies that are part of an intracellular assay must be fixed/permed

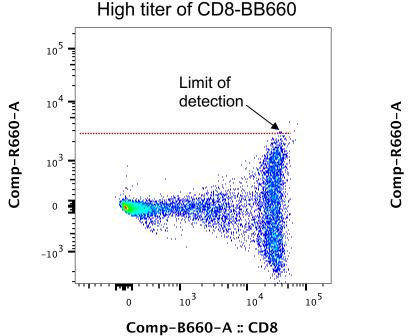
# ANTIBODY TITRATION(2)

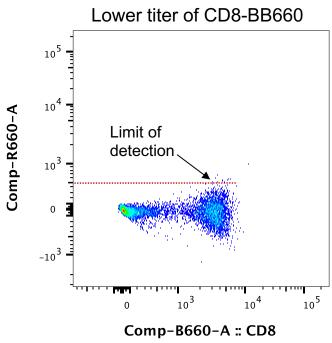
- Titrating will identify the optimal concentration at which to use the antibody
  - ➤ It will (almost always) save reagent (money)



# ANTIBODY TITRATION(3)

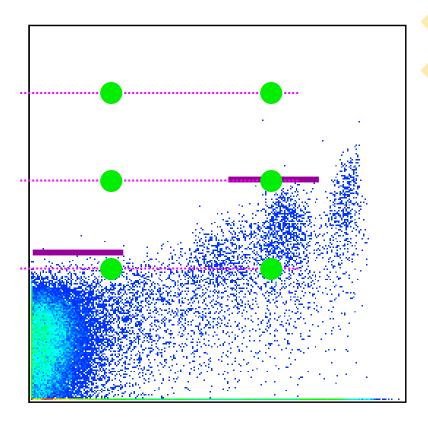
- Spreading error can be reduced if a saturating concentration is not needed (lineage markers)
  - ➤ Spreading is proportional to signal intensity





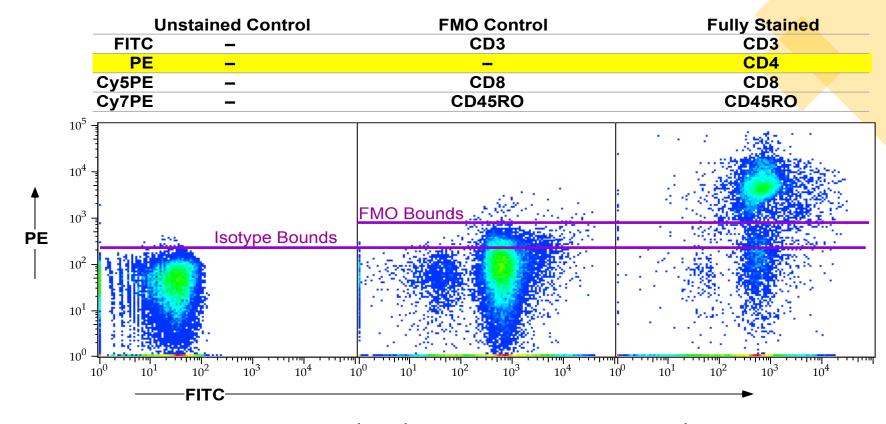
### STAINING CONTROLS

- Necessary to identify cells which do or do not express a given antigen
- Threshold for positivity may depend on the amount of fluorescence in other channels
- Unstained cells or isotype controls stains are improper controls



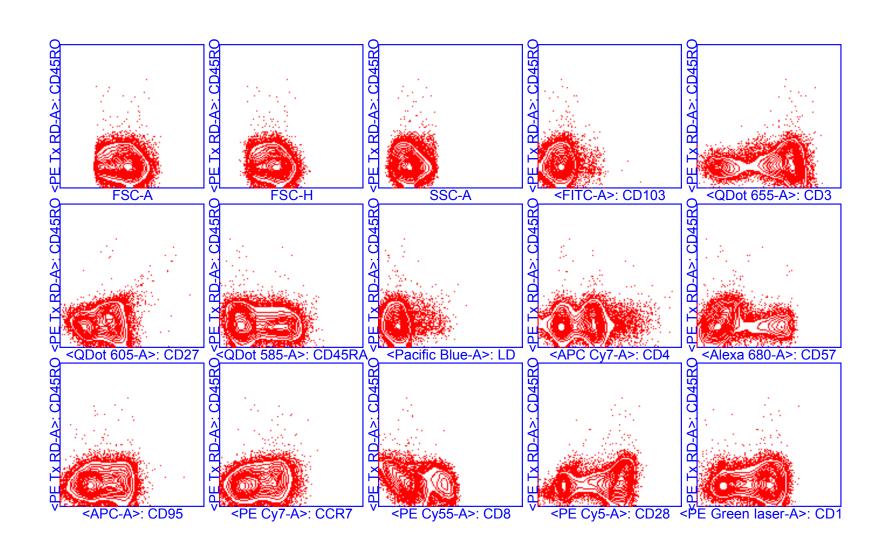
### FMO CONTROLS

- FMO = Fluorescence
  Minus One
  - Cells are stained with all reagents EXCEPT the one of interest
- Essential for complex panels
- Reveal unnoticed or unexpected issues with spreading
- Should be used for setting correct gates



PBMC stained as shown. Compensation properly set.

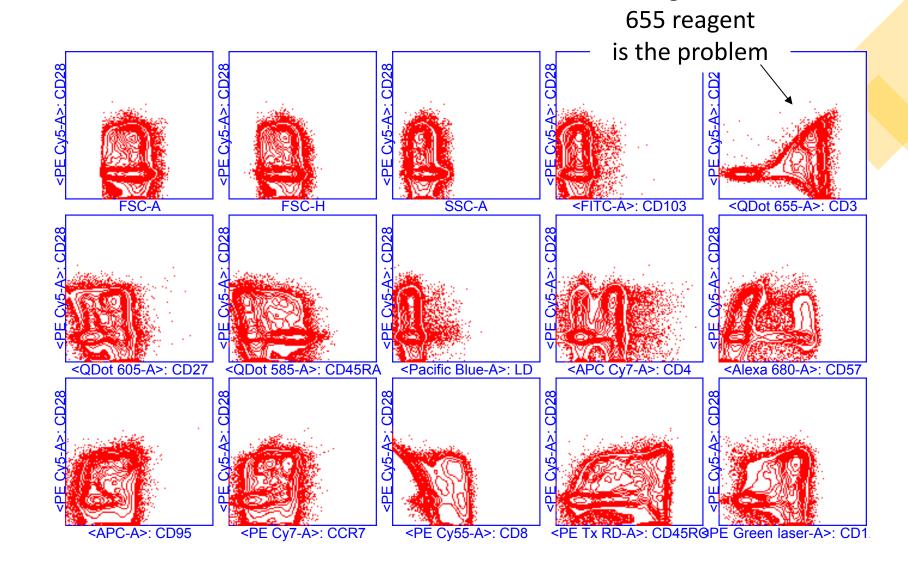
### FMO EXAMPLE - MISSING PE-TR



#### FMO Example – Missing PE-Cy5

 CD28 PE-Cy5 was not added to the staining cocktail but there appears to be a positive signal

 Staining artefact comes from large spreading of Qdot reagent



A bright Qdot

### Panel assessment using multigraph overlays

- Overlay different populations to see where subsets exist
- Identify potential spreading issues



#### **SUMMARY**

#### Compensation

- > Compensation values are arbitrary
- > All panels (but especially large) require appropriate compensation controls
- > Analyze properly transformed (and compensated) data
- > Properly compensated data reveals errors, it does not cause them

#### Spillover/Spreading

- > Spillover/spreading error is the single most important contributor to background and loss of resolution
- > Spreading error is instruments specific
- > Spreading error is proportional to signal intensity

#### Panel Development

- > A standardized, optimized instrument is key to successful panel development
- > Use the SSM matrix and antigen co-expression to guide marker placement
- > Titrate antibodies
- > Use appropriate controls and QC checks when assessing a new panel
- > The process is iterative and sometimes frustrating

### **ACKNOWLEDGMENTS**



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