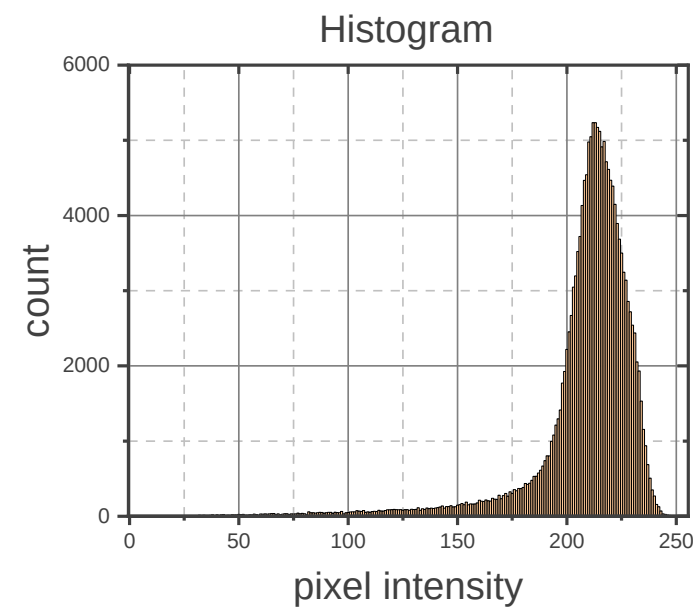
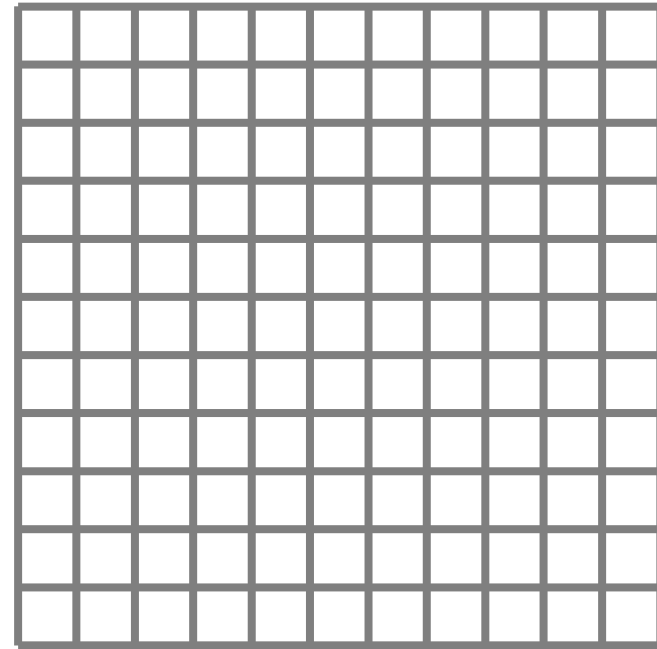


Course

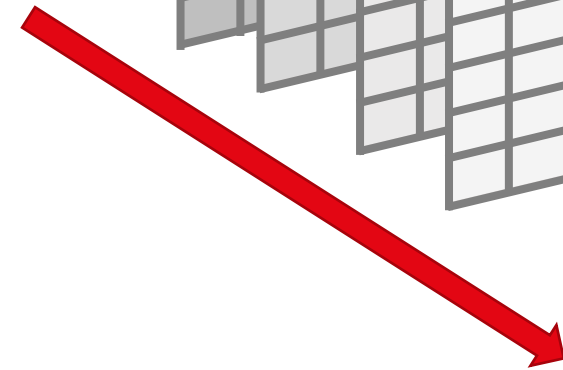
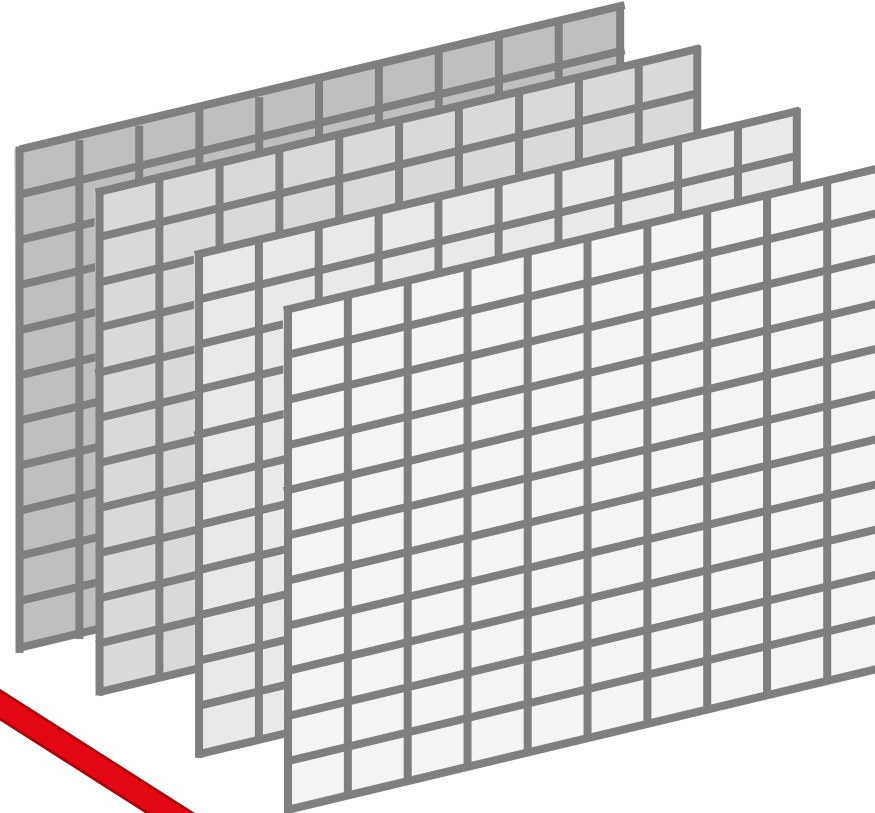
Image Statistics

Image Statistics

2D

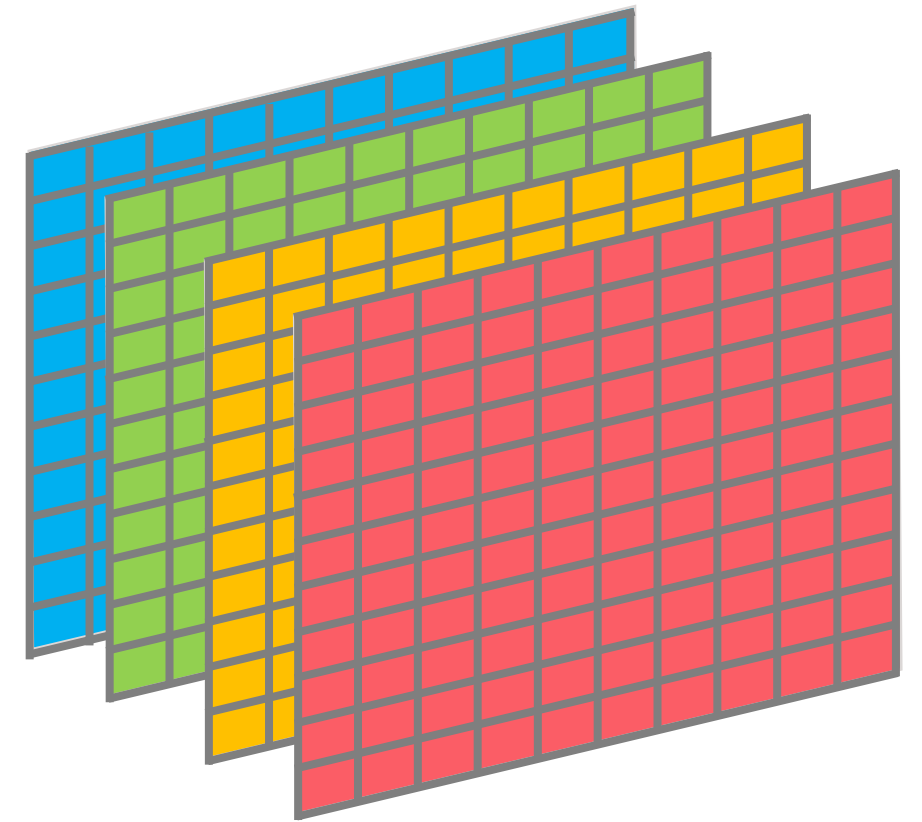


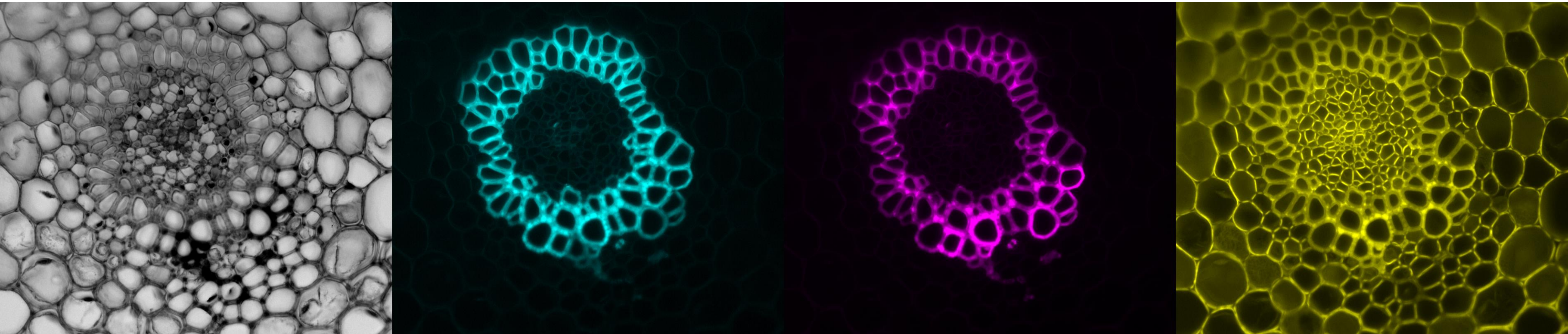
3D, 2D Time



Projection
mean, min, max,...

Multi channel





Transmission

DAPI

excitation: BP 387/11
emission: BP 447/60

Green

excitation: BP 466/40
emission: BP 525/50

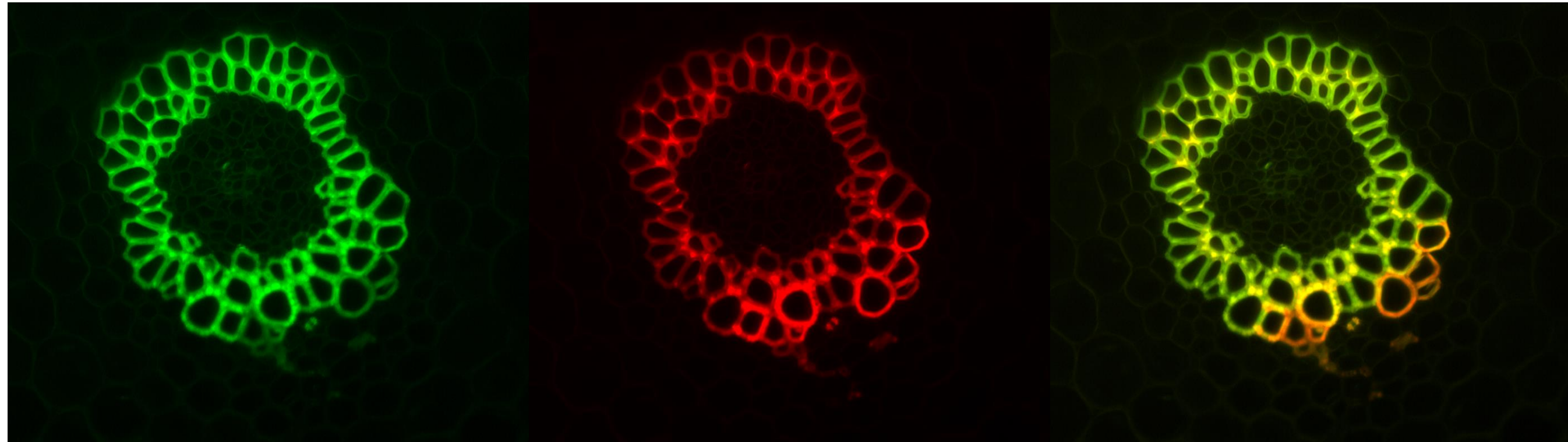
Red

excitation: BP 562/40
emission: BP 624/40

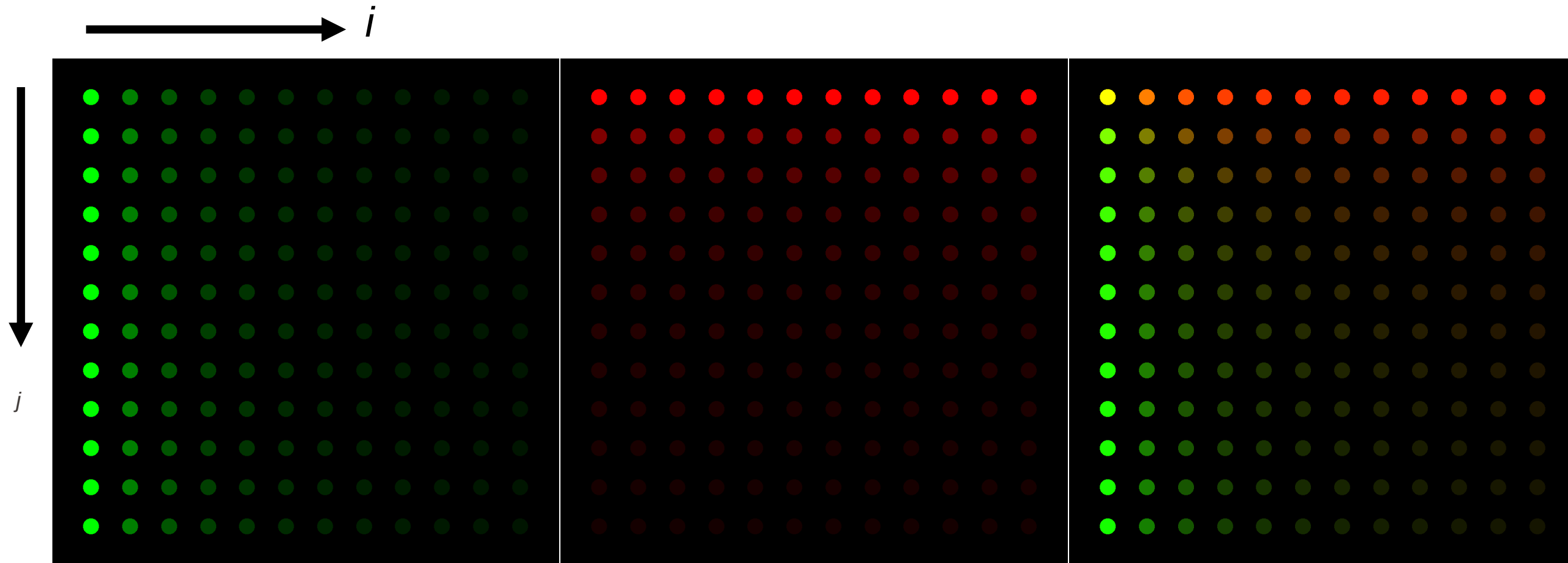
Overlap detection and quantification

Overlap detection

Color merge

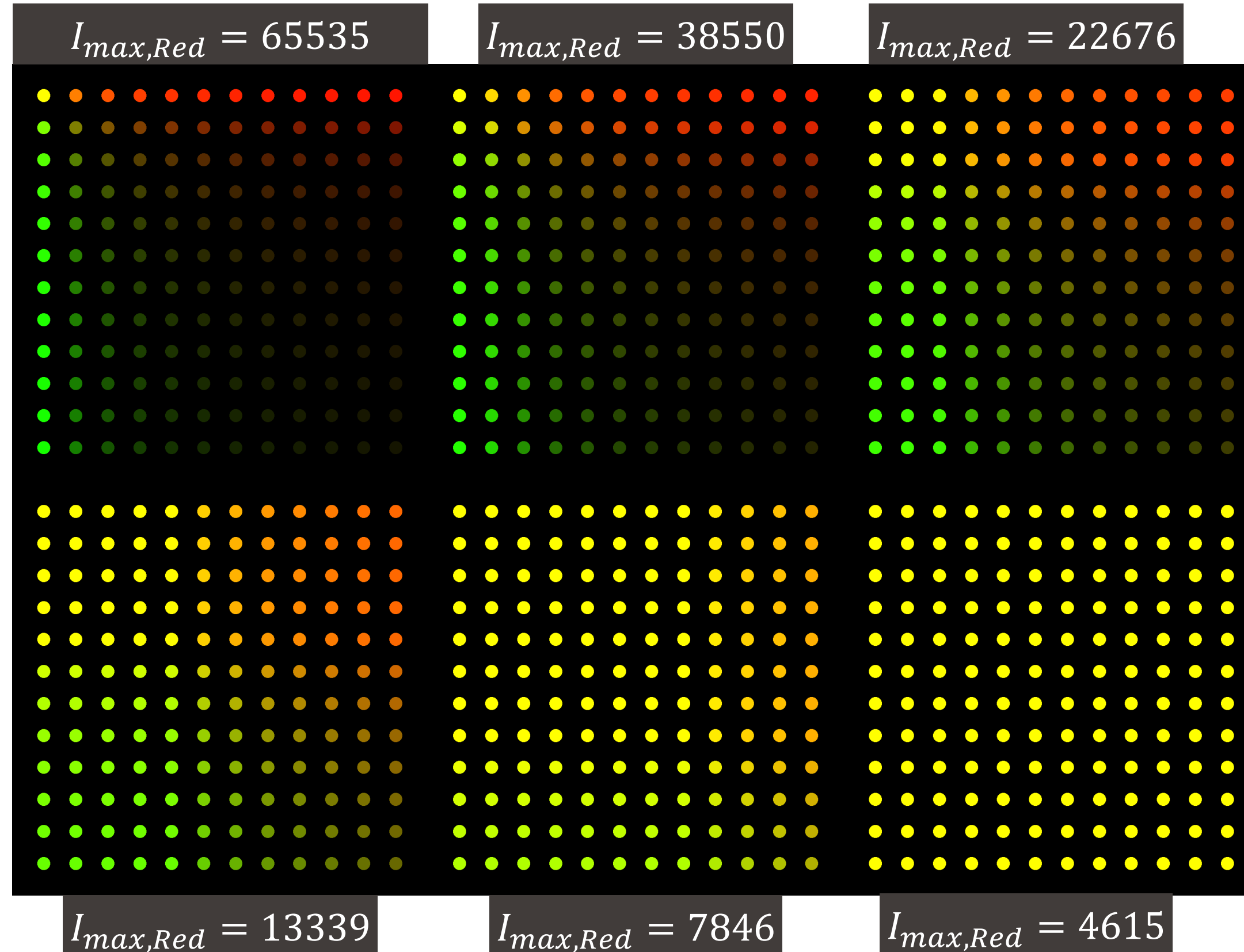


Colour is a false friend

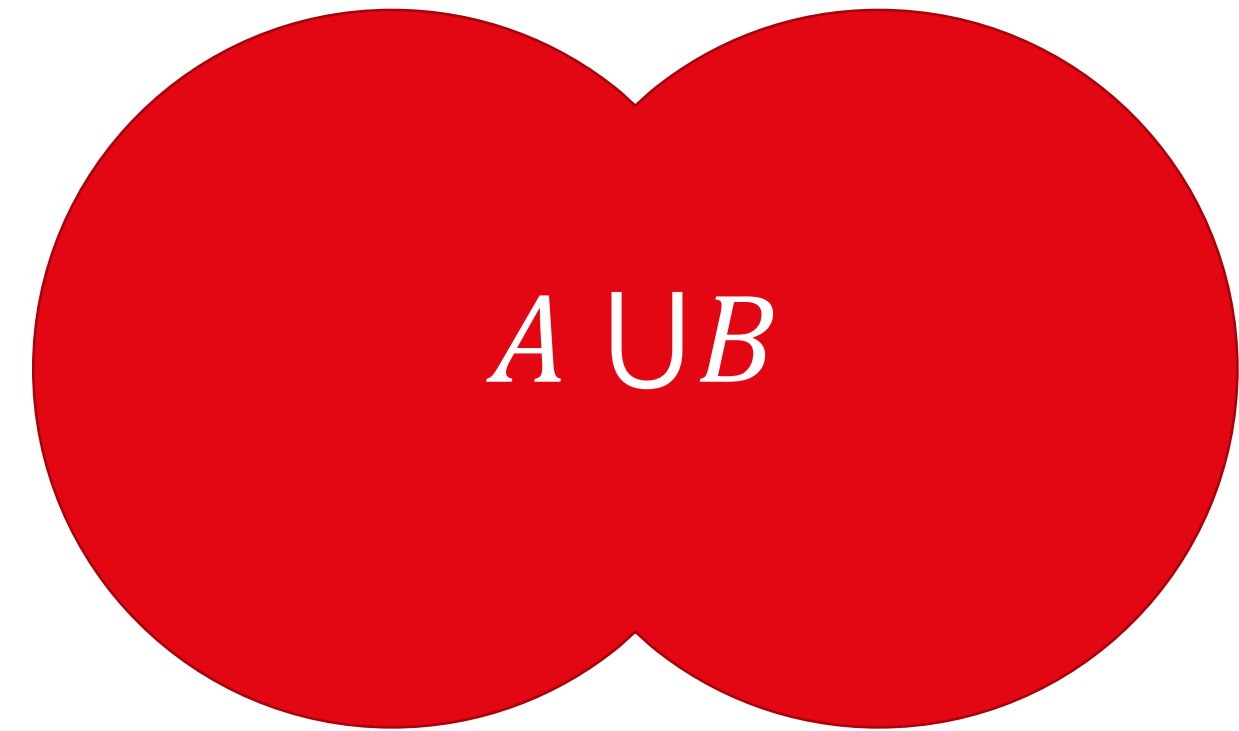
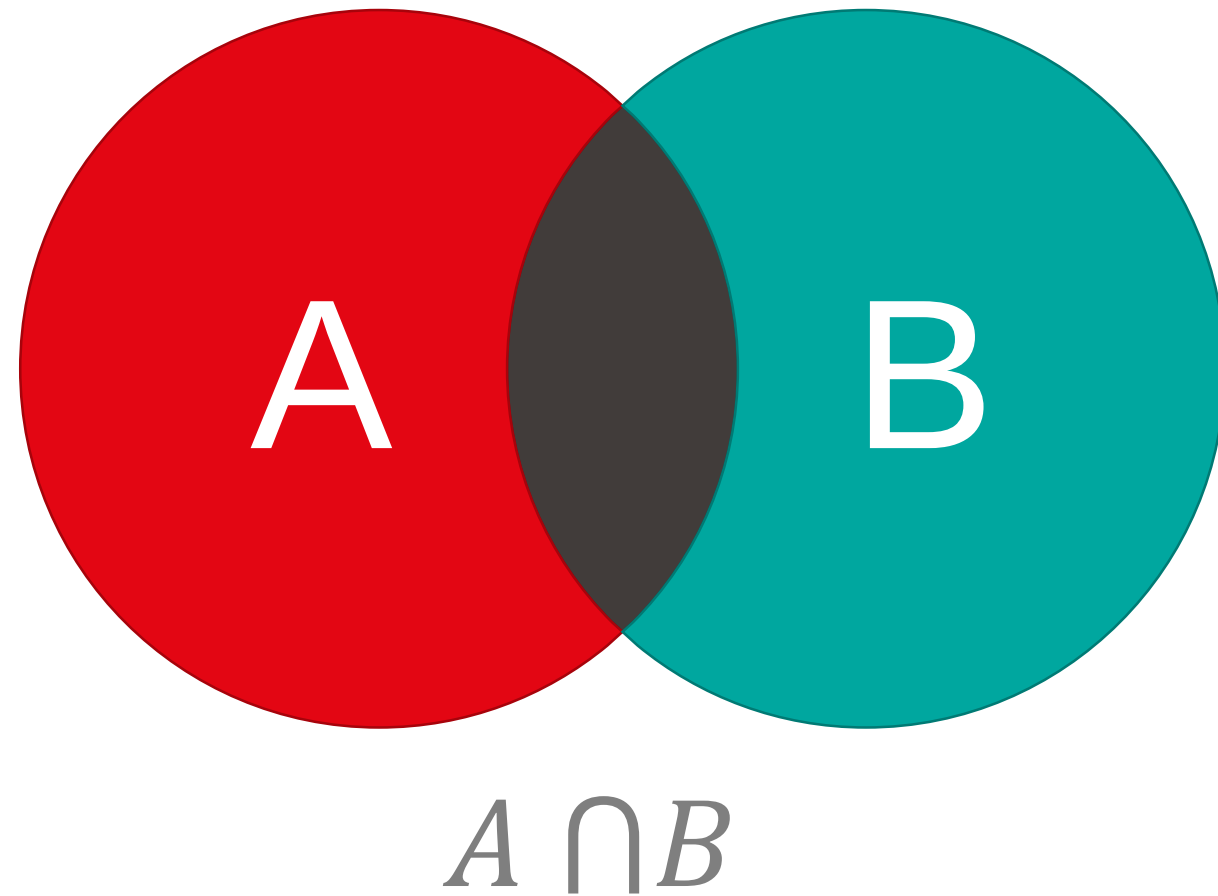


$$I_{green}(i) = \frac{1}{i} * (2^{16} - 1) \quad I_{red}(j) = \frac{1}{j} * (2^{16} - 1)$$

Colour is a false friend



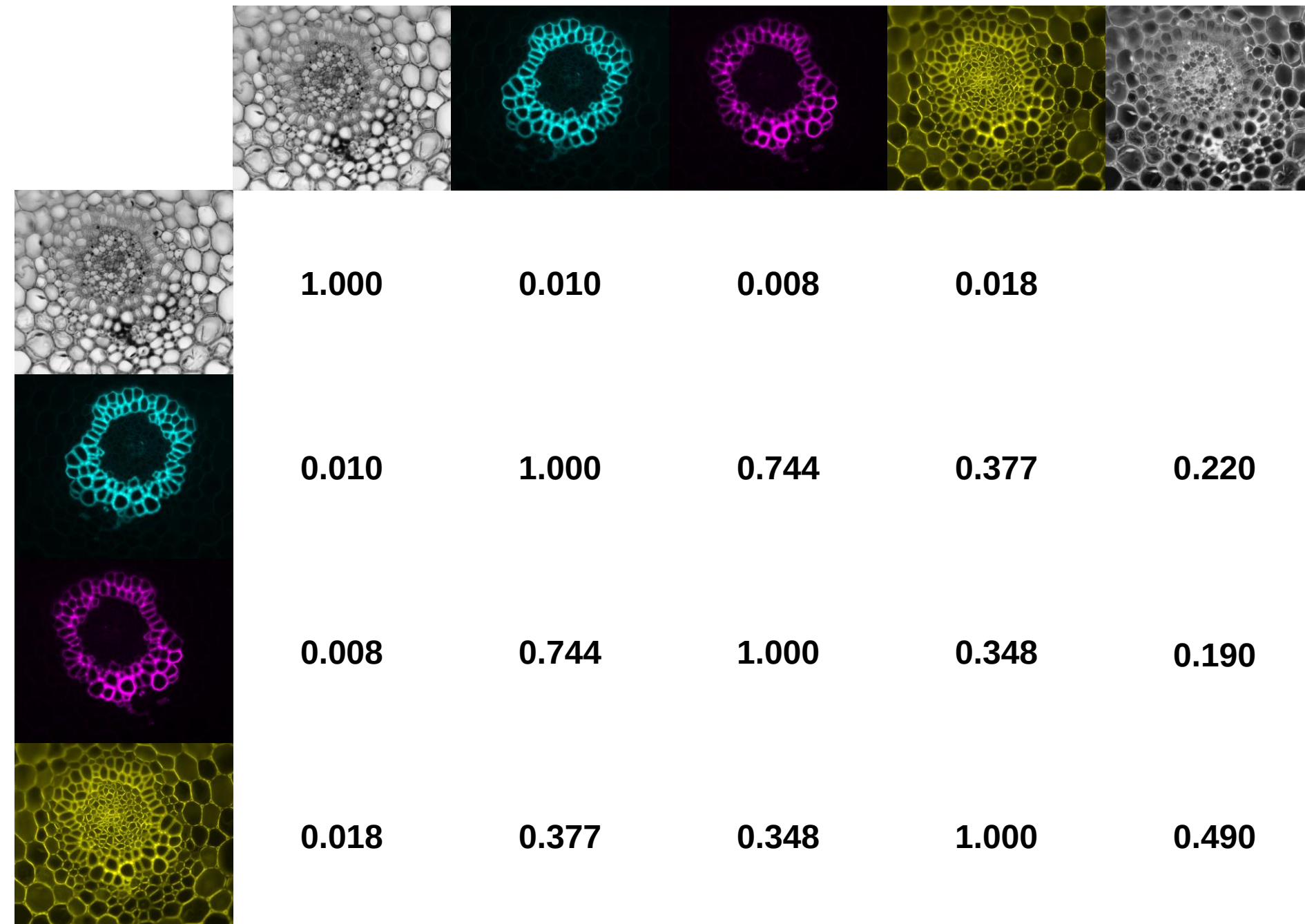
Overlap metrics



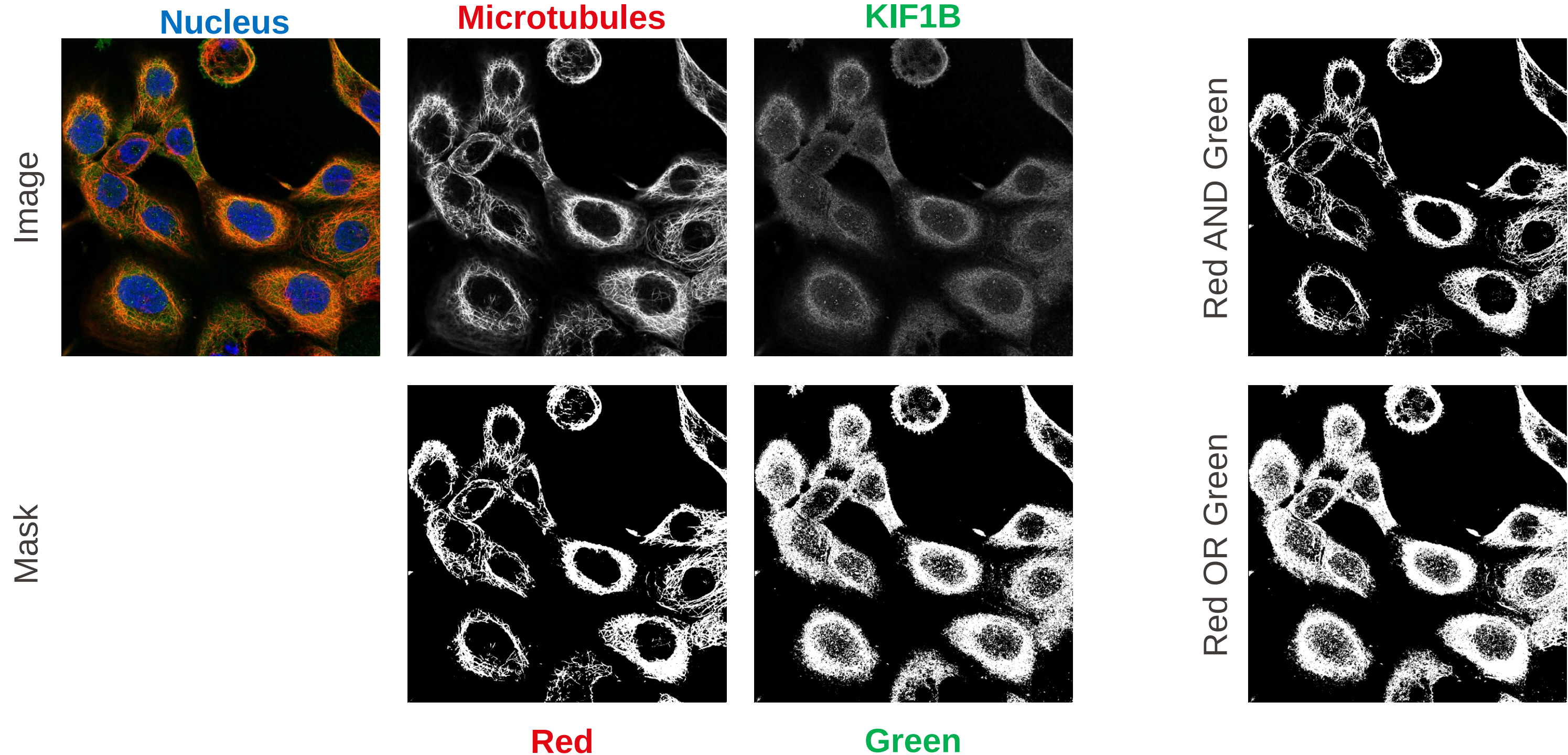
$$J(A, B) = \frac{|A \cap B|}{|A \cup B|}$$

Intersection over Union (IoU)

Overlap (IoU)

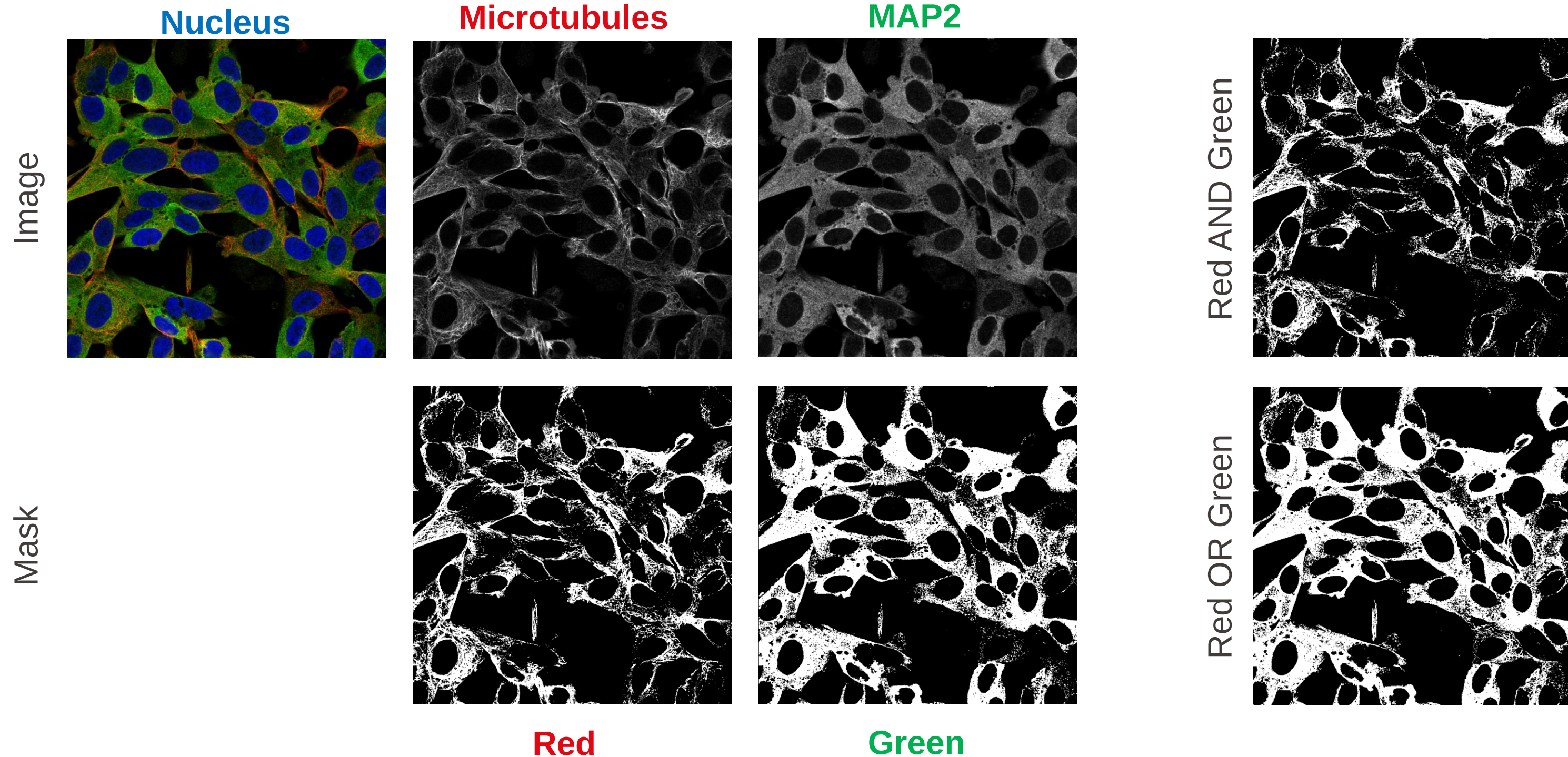


Overlap in life sciences



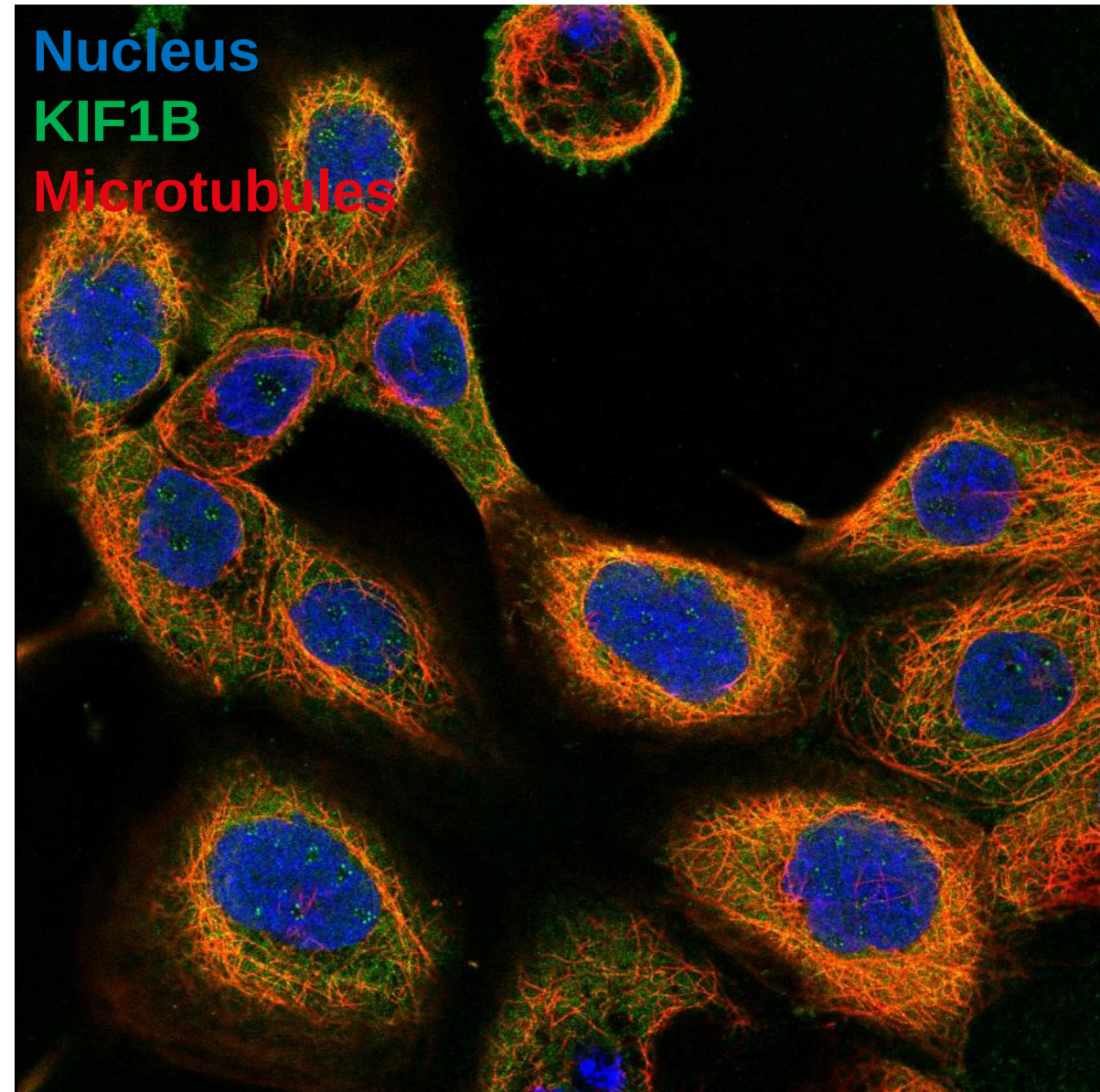
<https://www.proteinatlas.org/ENSG00000054523-KIF1B/cell#img>

Overlap in life sciences

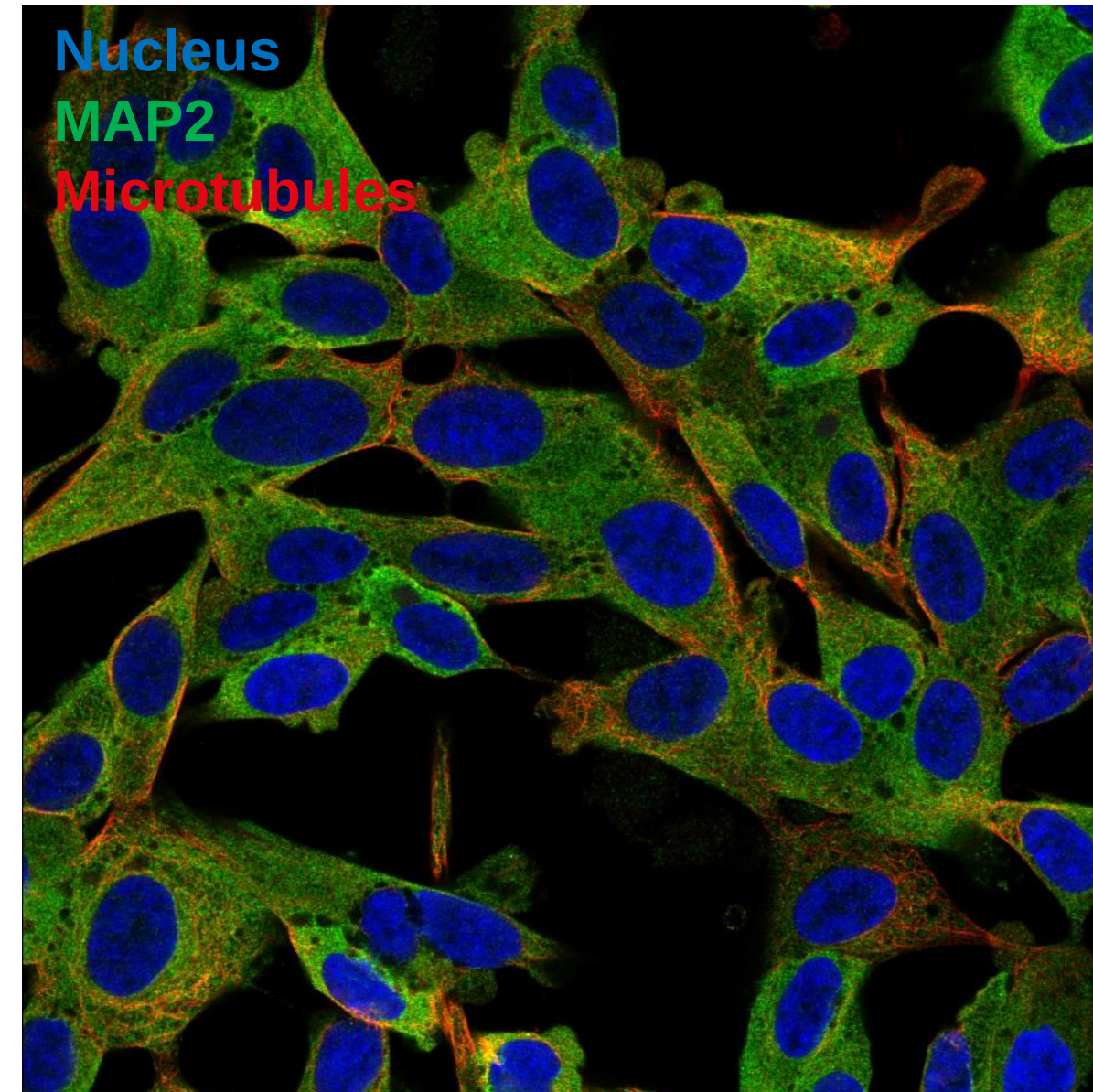


<https://www.proteinatlas.org/ENSG00000078018-MAP2/cell#img>

Overlap in life sciences

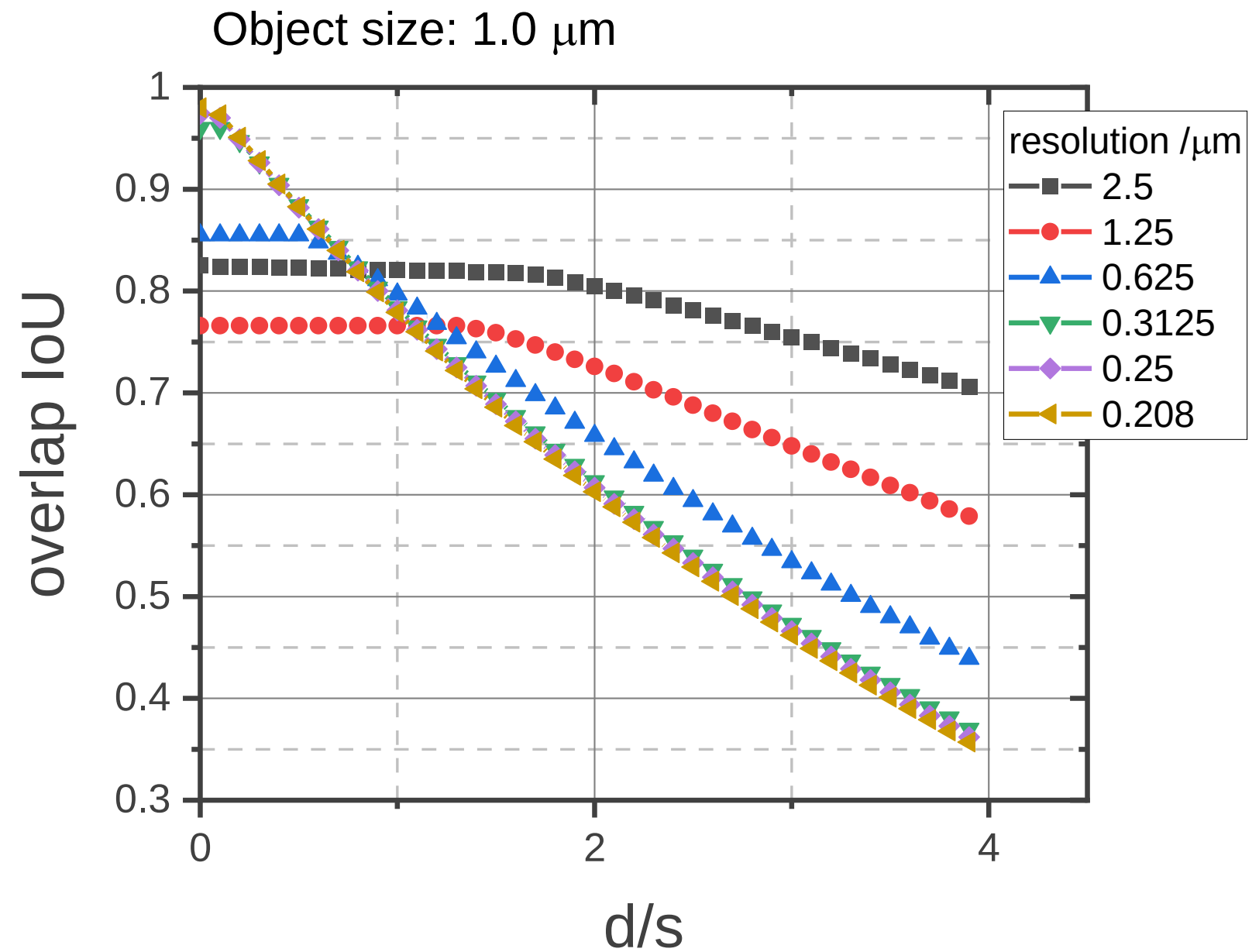
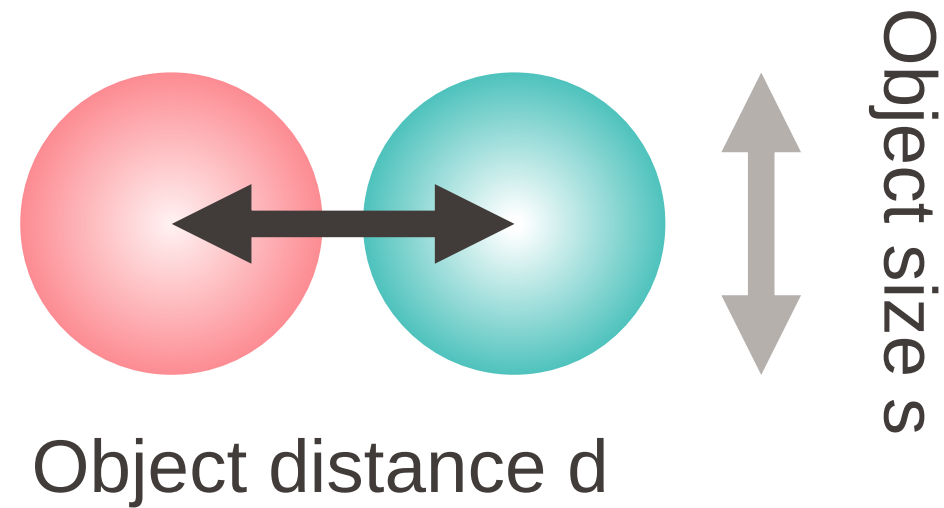


$$J(A,B)=0.498$$

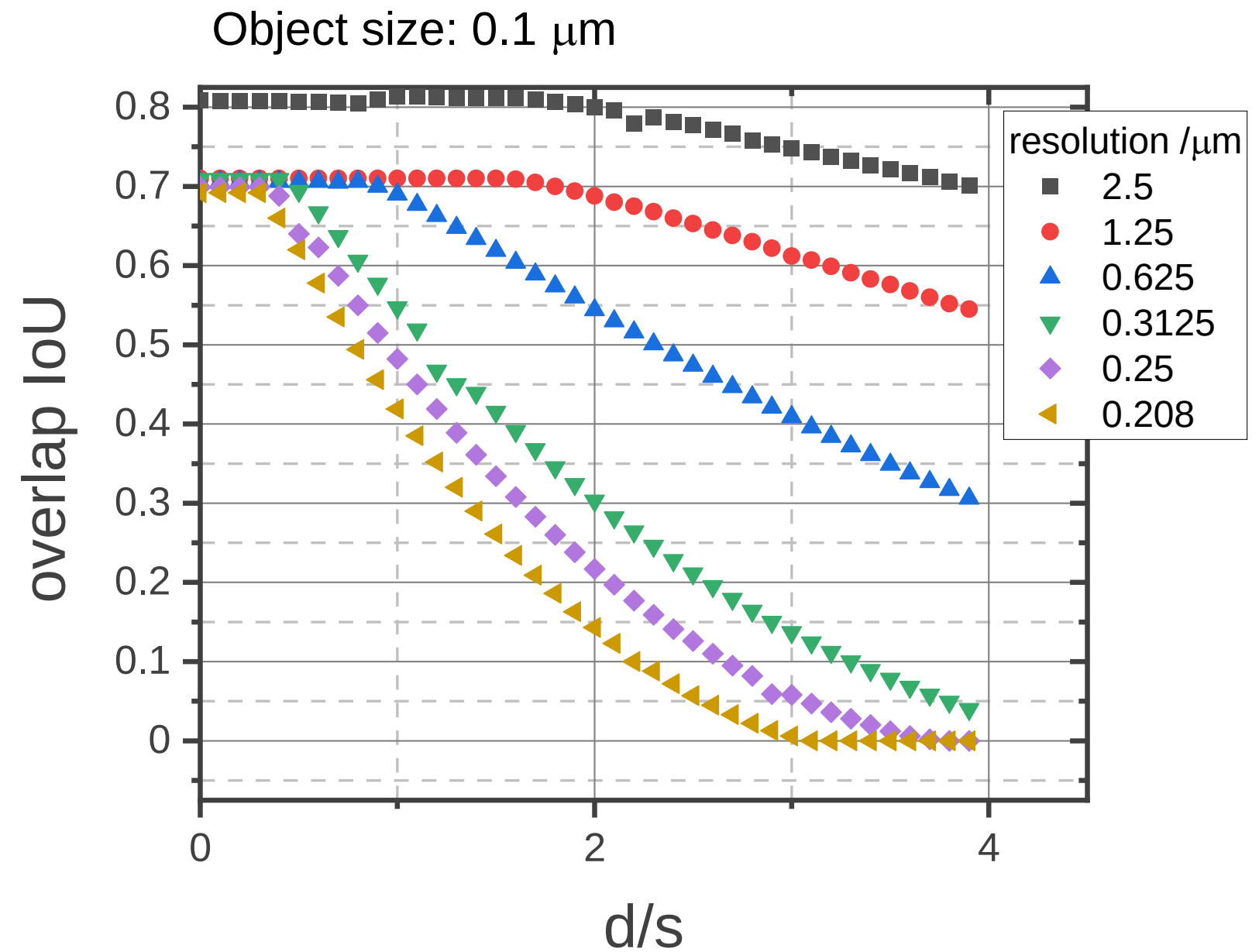
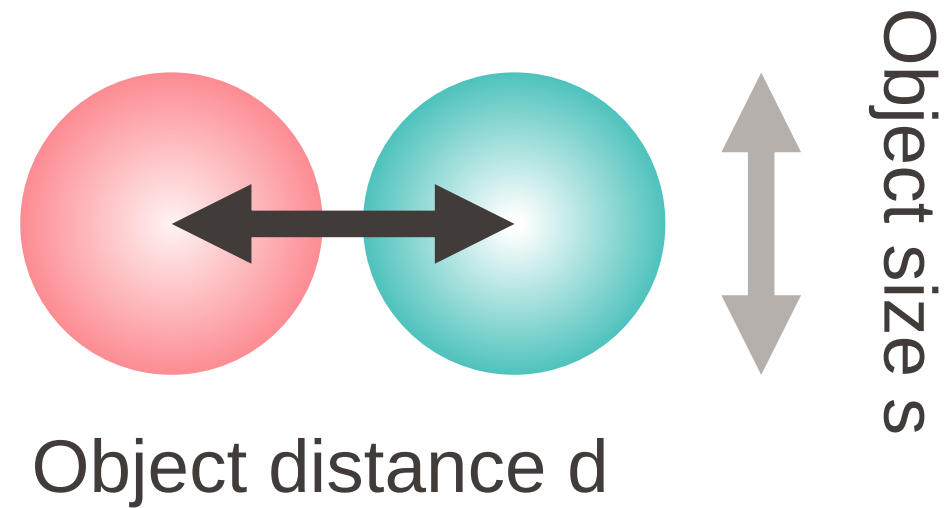


$$J(A,B)=0.437$$

Overlap Size dependence

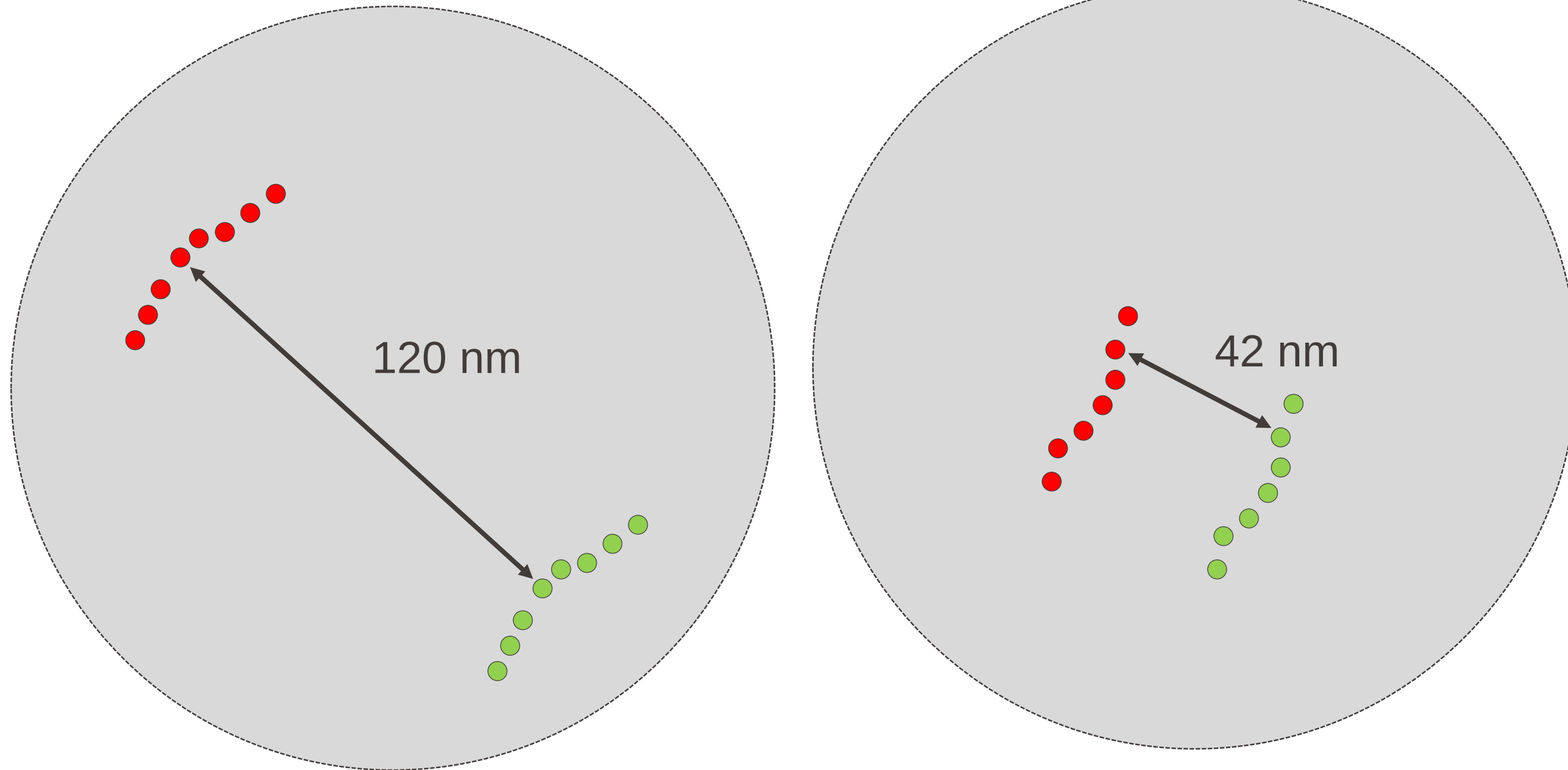


Overlap Size dependence



Low pass filtering

Diffraction limited spot (200 nm)



● ● = fluorescent proteins (to scale)

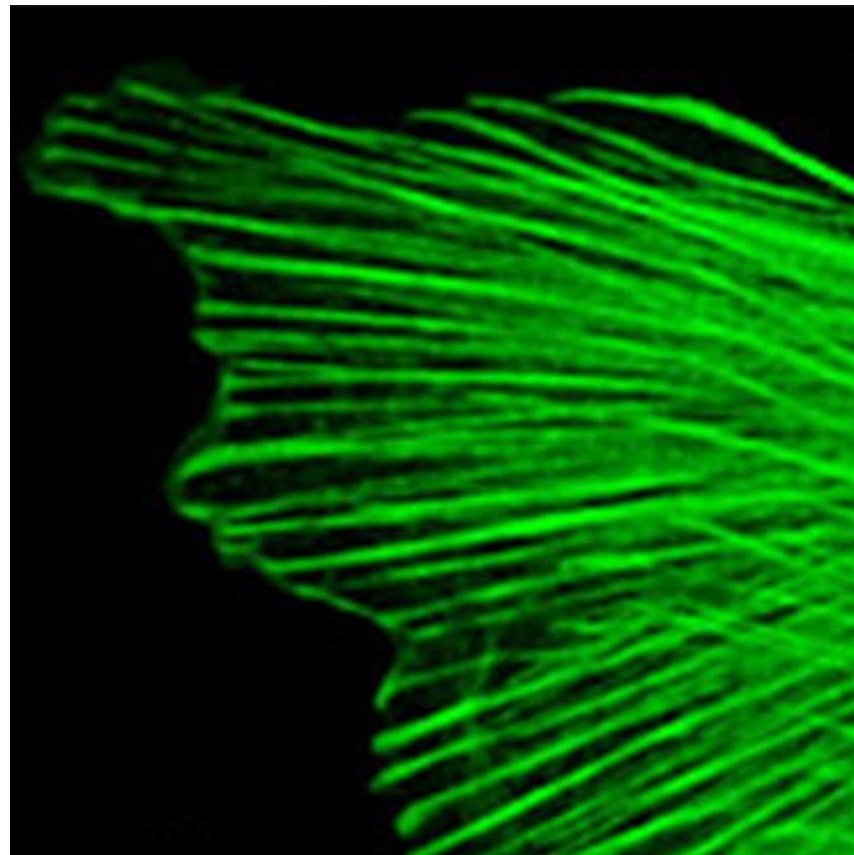
Overlap Summary

- Overlap (IoU)
quantitative metric
- Requires setting
thresholds for the
individual channels
- Size and resolution
dependence

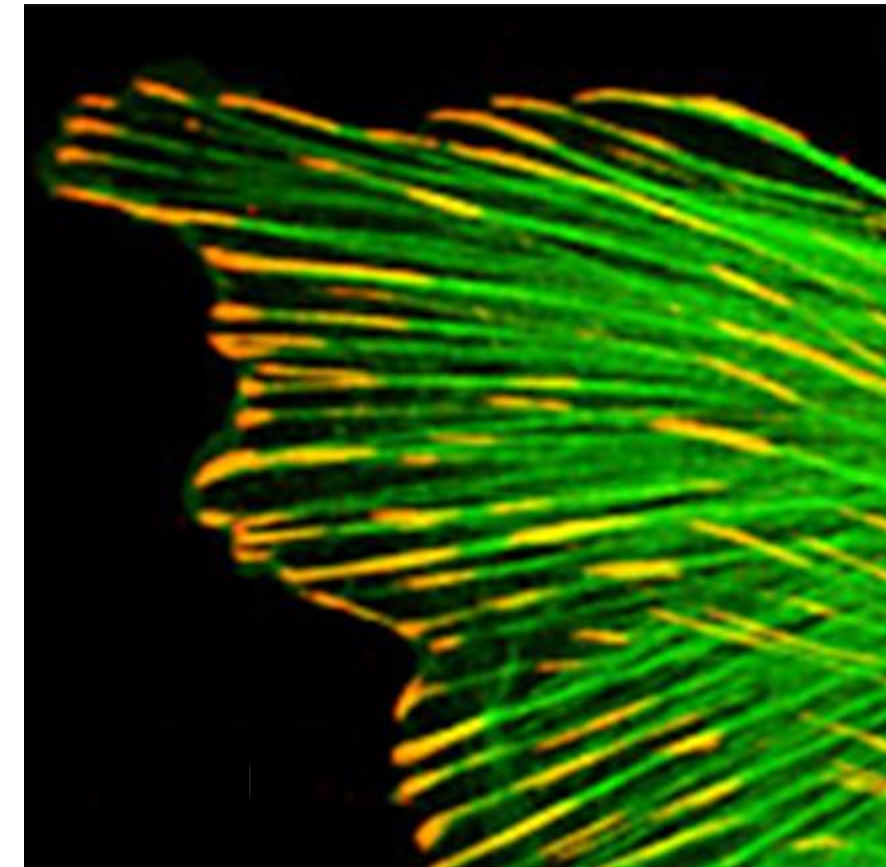
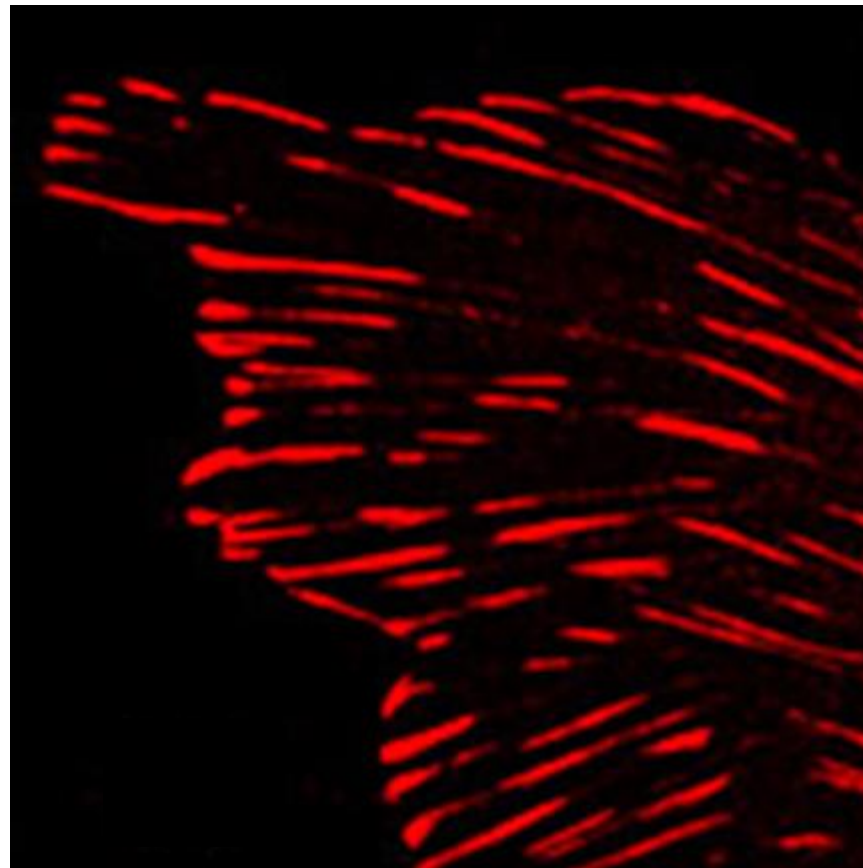
Subcellular Localization & Colocalization

Subcellular Colocalization

Vinculin
Alexa568



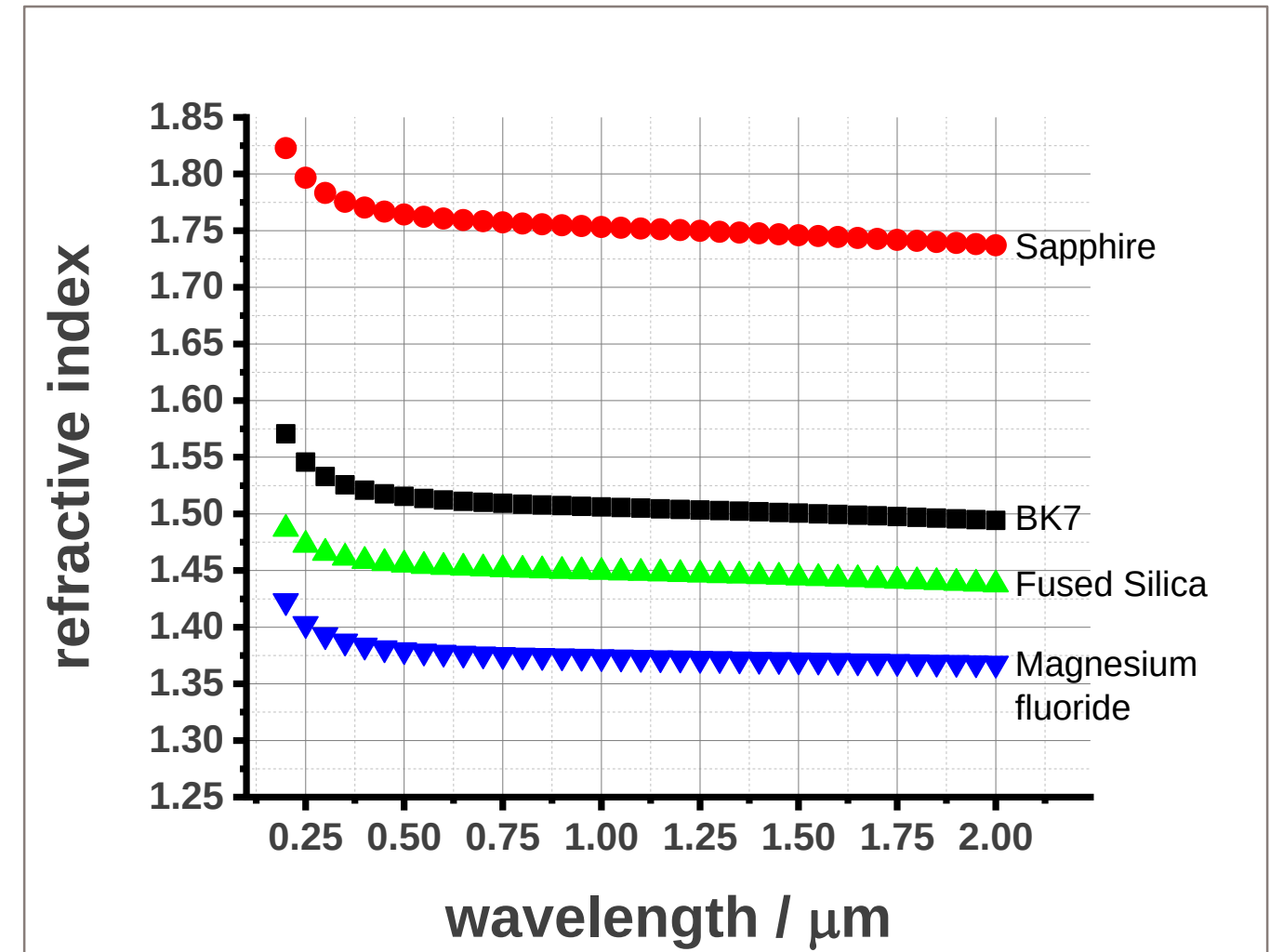
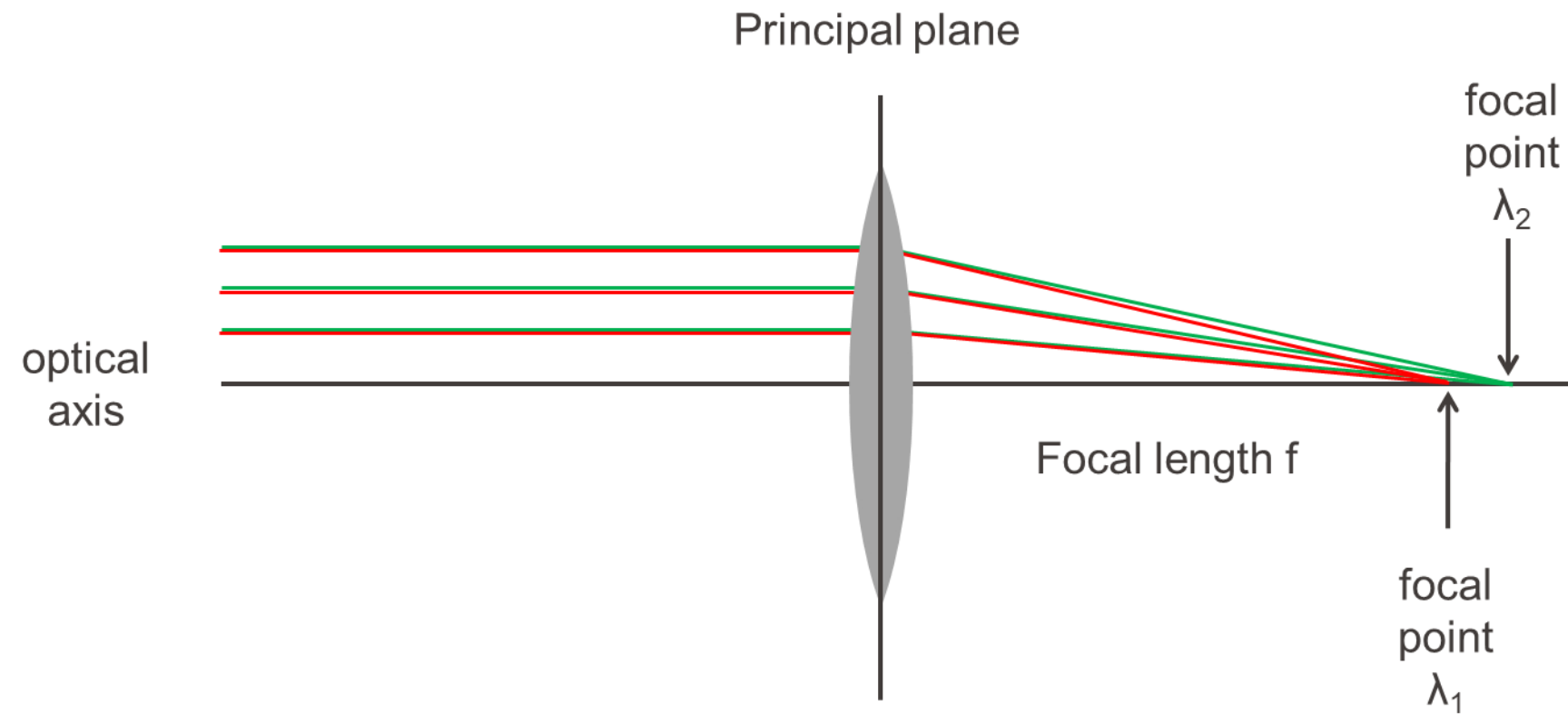
Actin
Alexa488



<http://www.olympusconfocal.com/applications/colocalization.html>

Colocalization: The presence of two or more fluorophores on the same physical structure (in a cell).

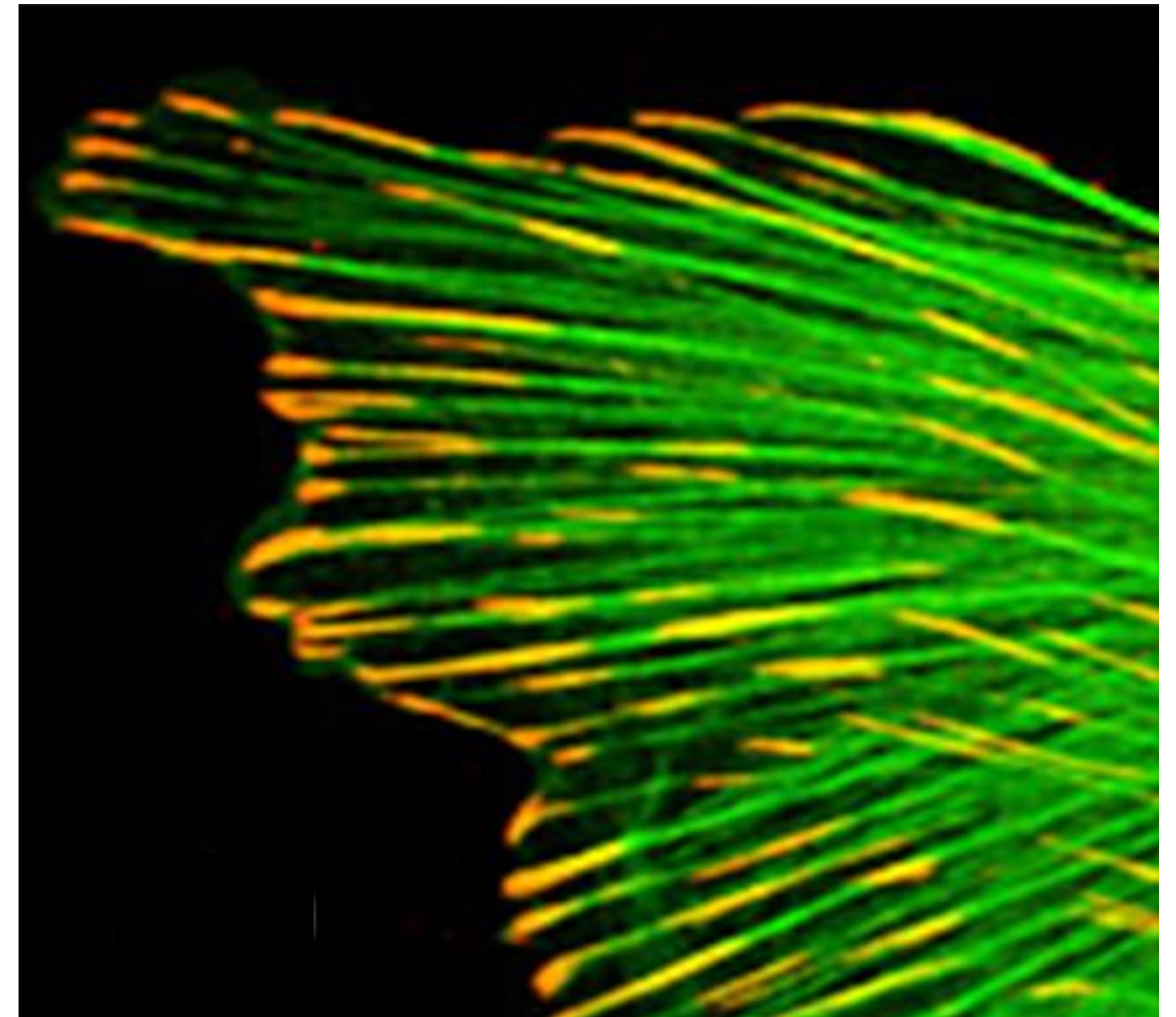
Chromatic aberrations



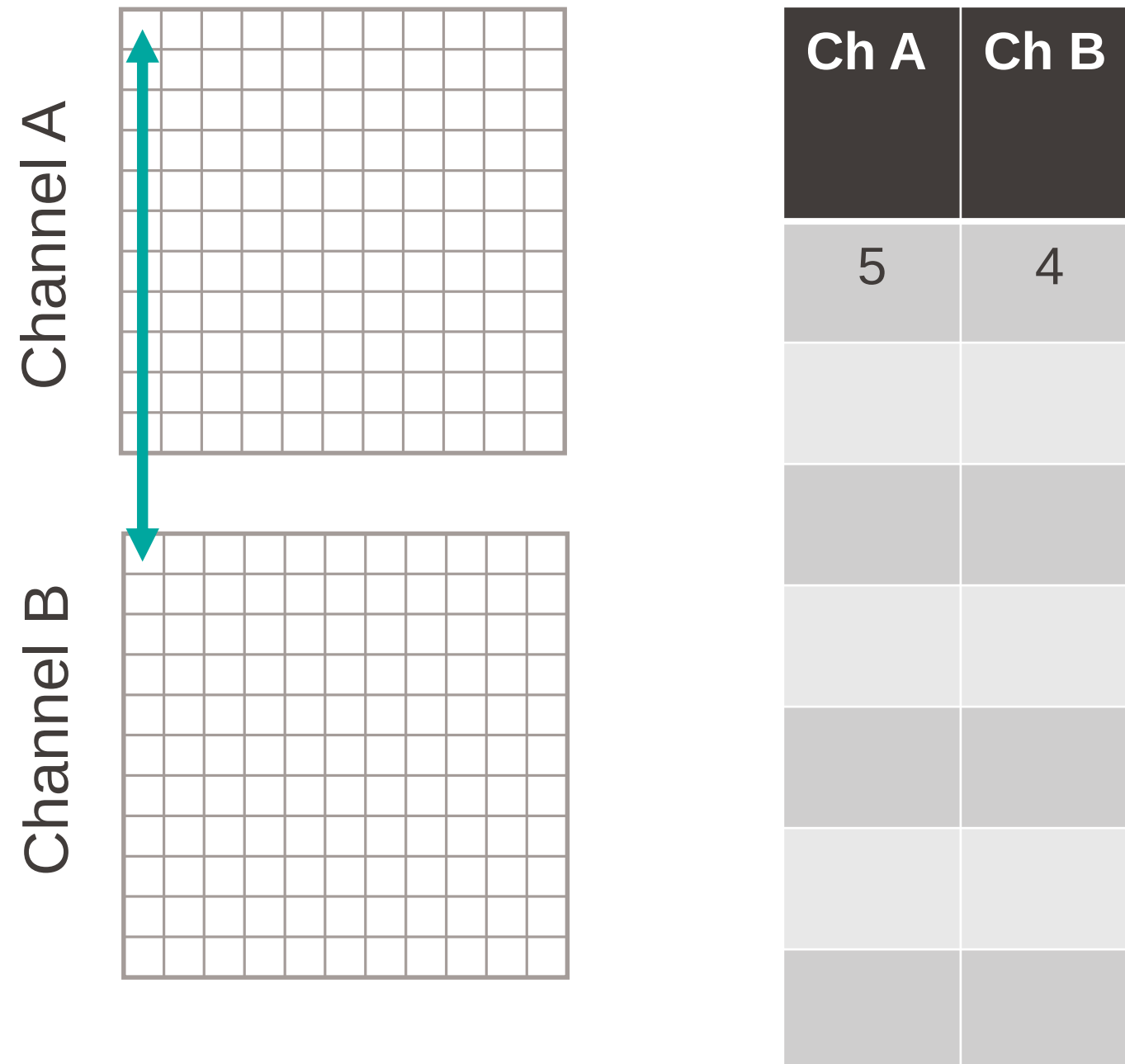
Colocalisation

Experimental requirements

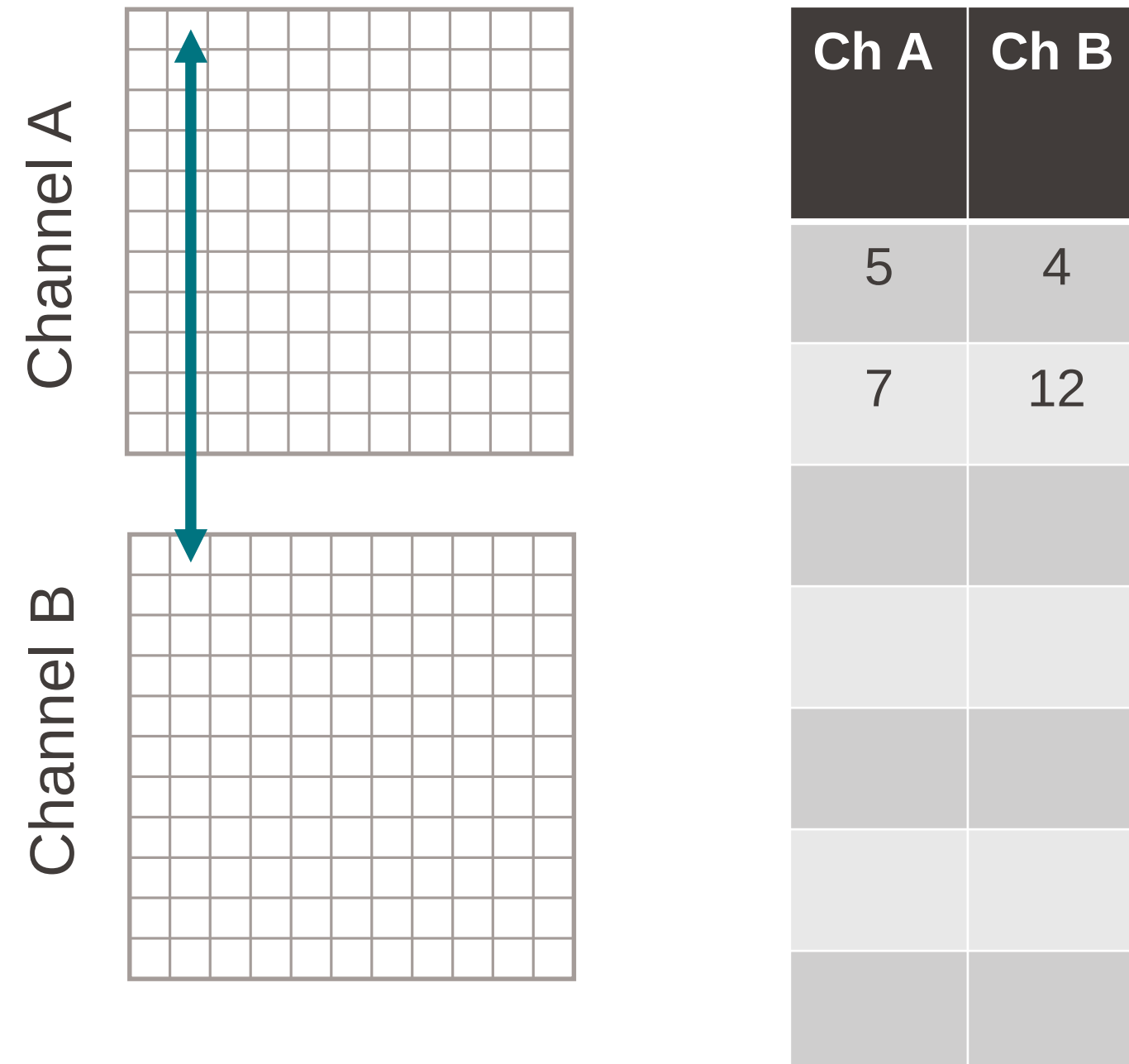
- Resolution (NA of the objective lens) needs to match the structure of interest
- Avoid chromatic aberrations
 - Correct for remaining chromatic aberrations if necessary
- Avoid channel cross-talk
 - Correct for cross-talk if necessary
- Reduce the out-of-focus background
- Use preferably high SNR images



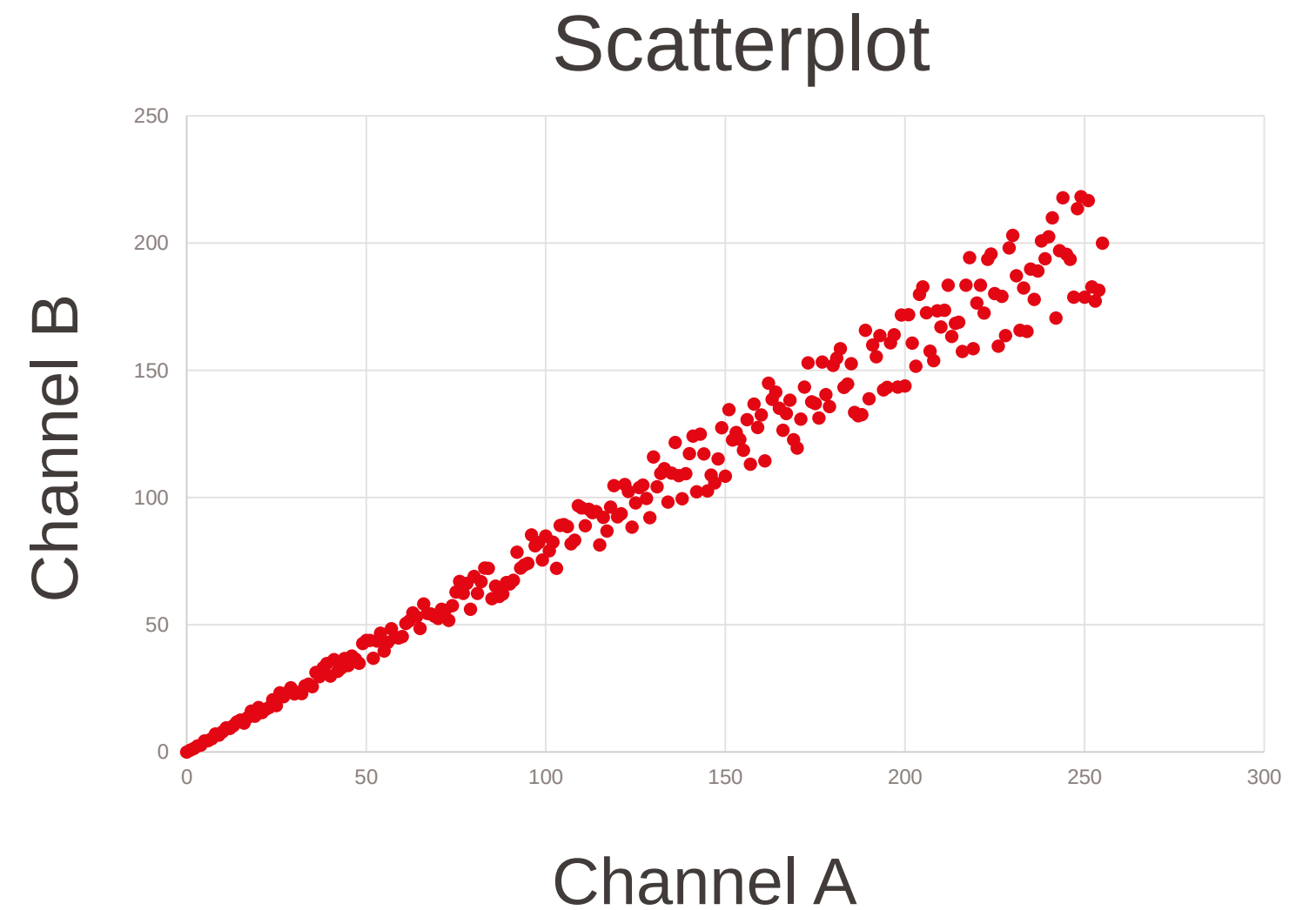
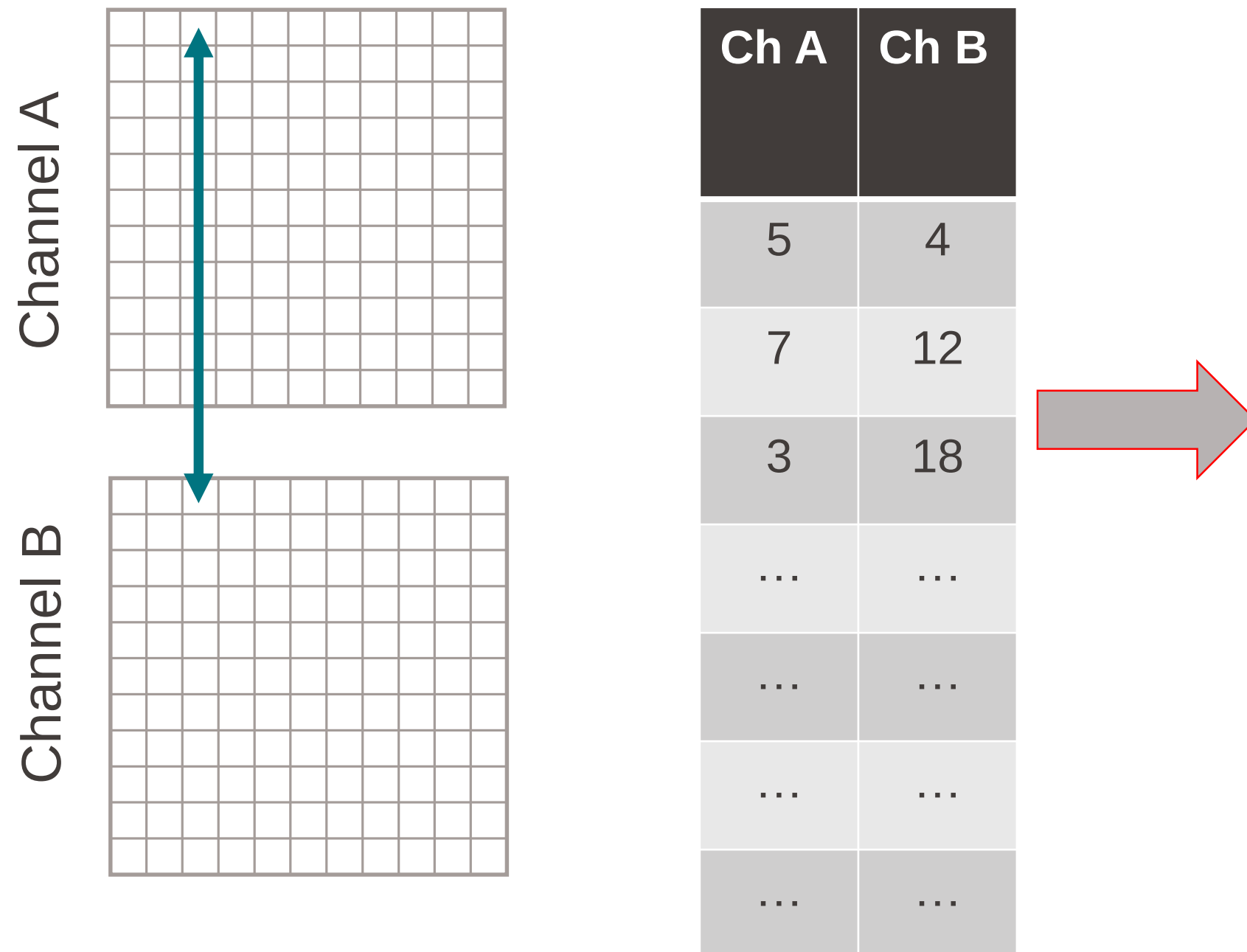
Channel Statistics



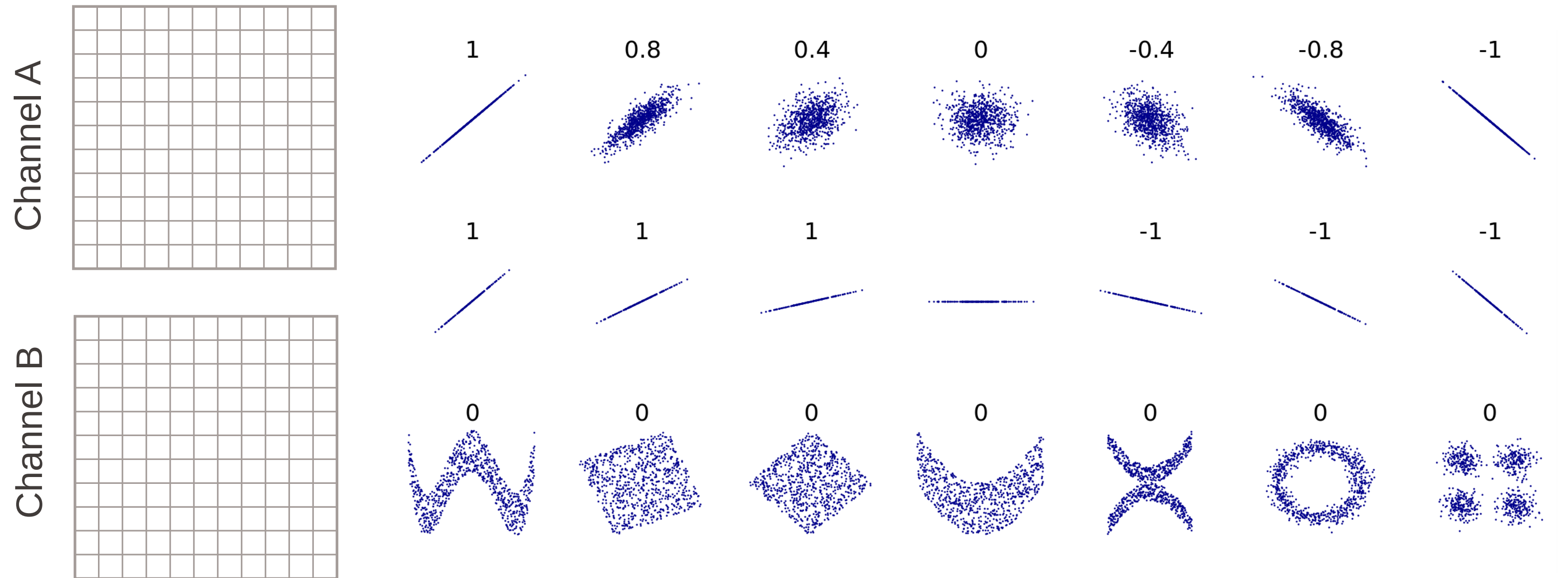
Channel Statistics



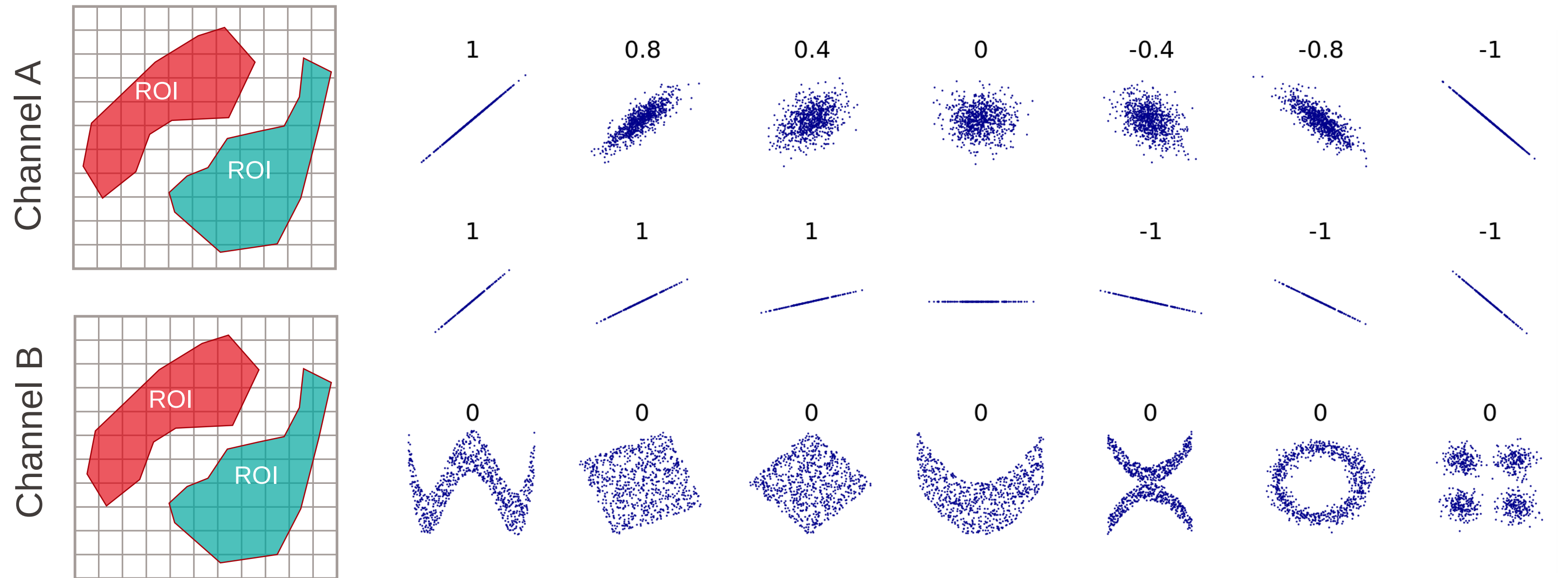
Channel Statistics



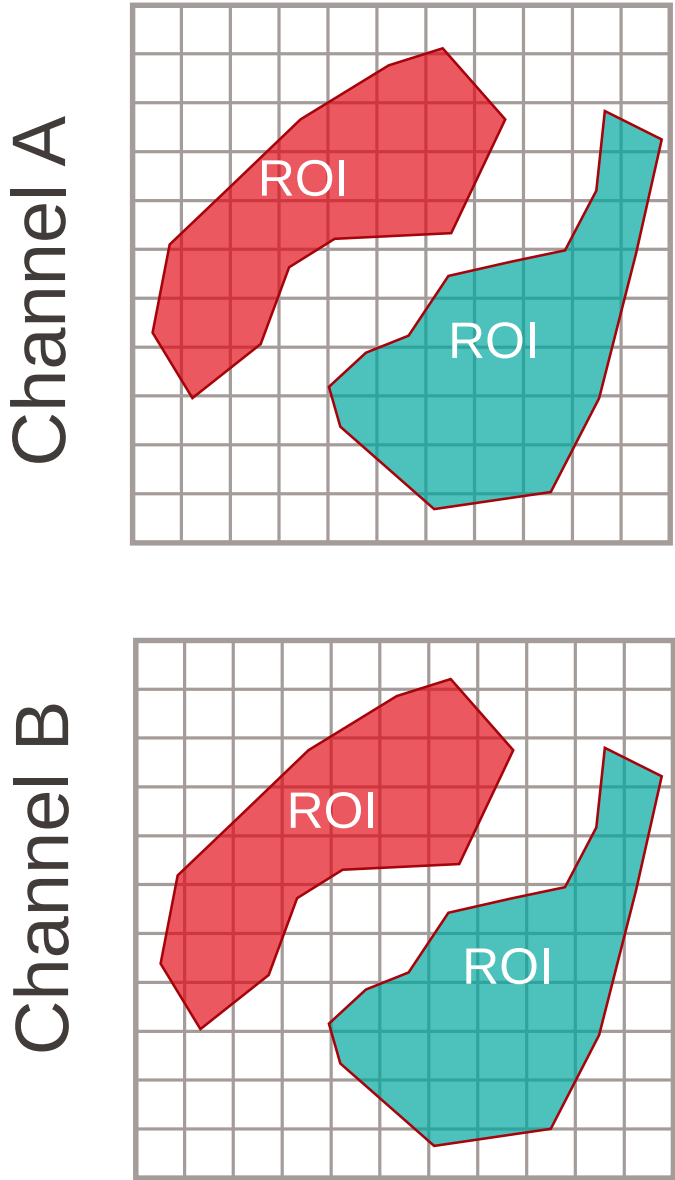
Pearson Correlation Coefficient



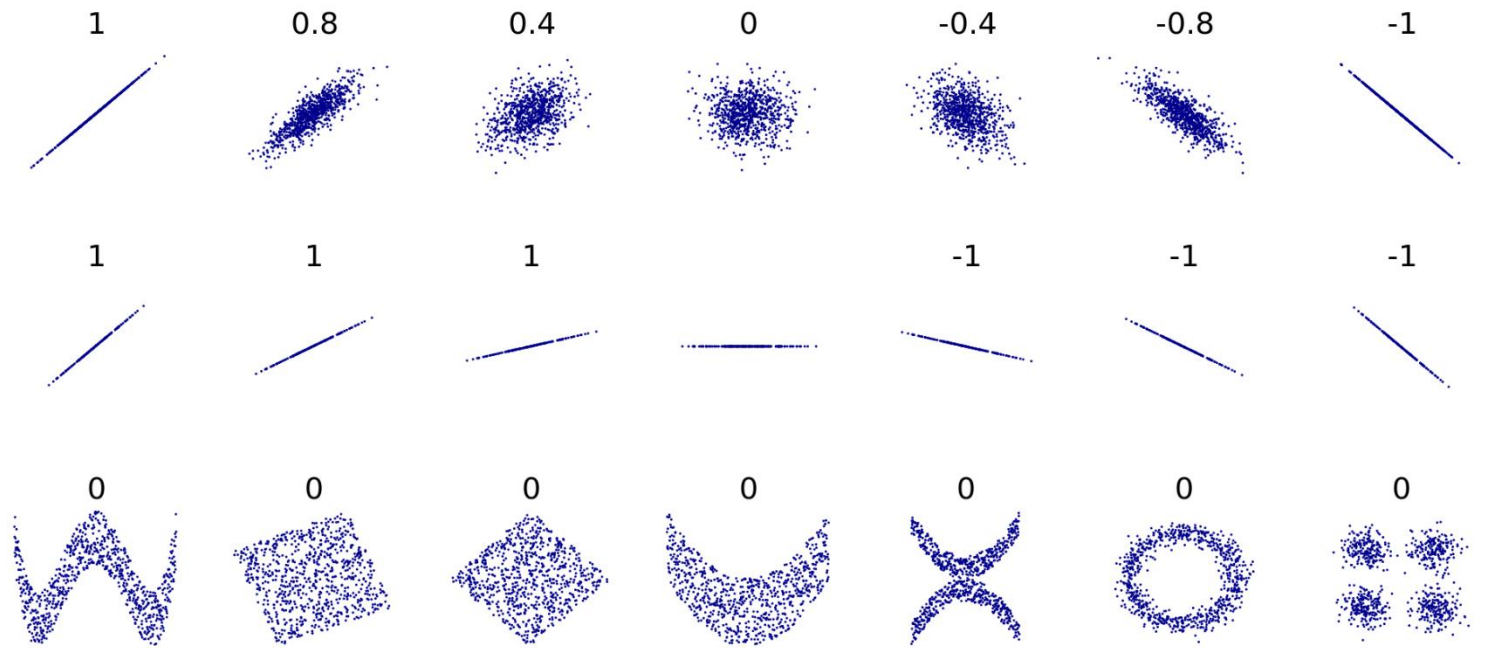
Pearson Correlation Coefficient



Pearson Correlation Coefficient



ROI	PCC
1	0.8
2	0.5
3	0.7

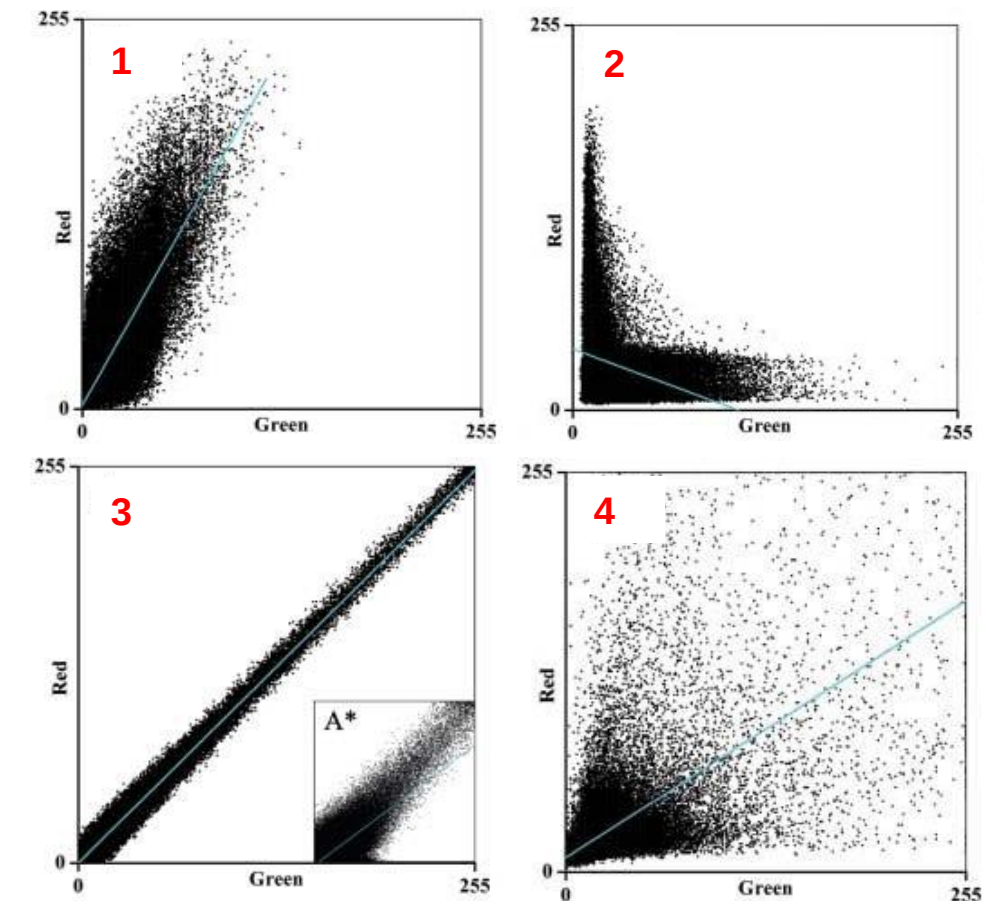
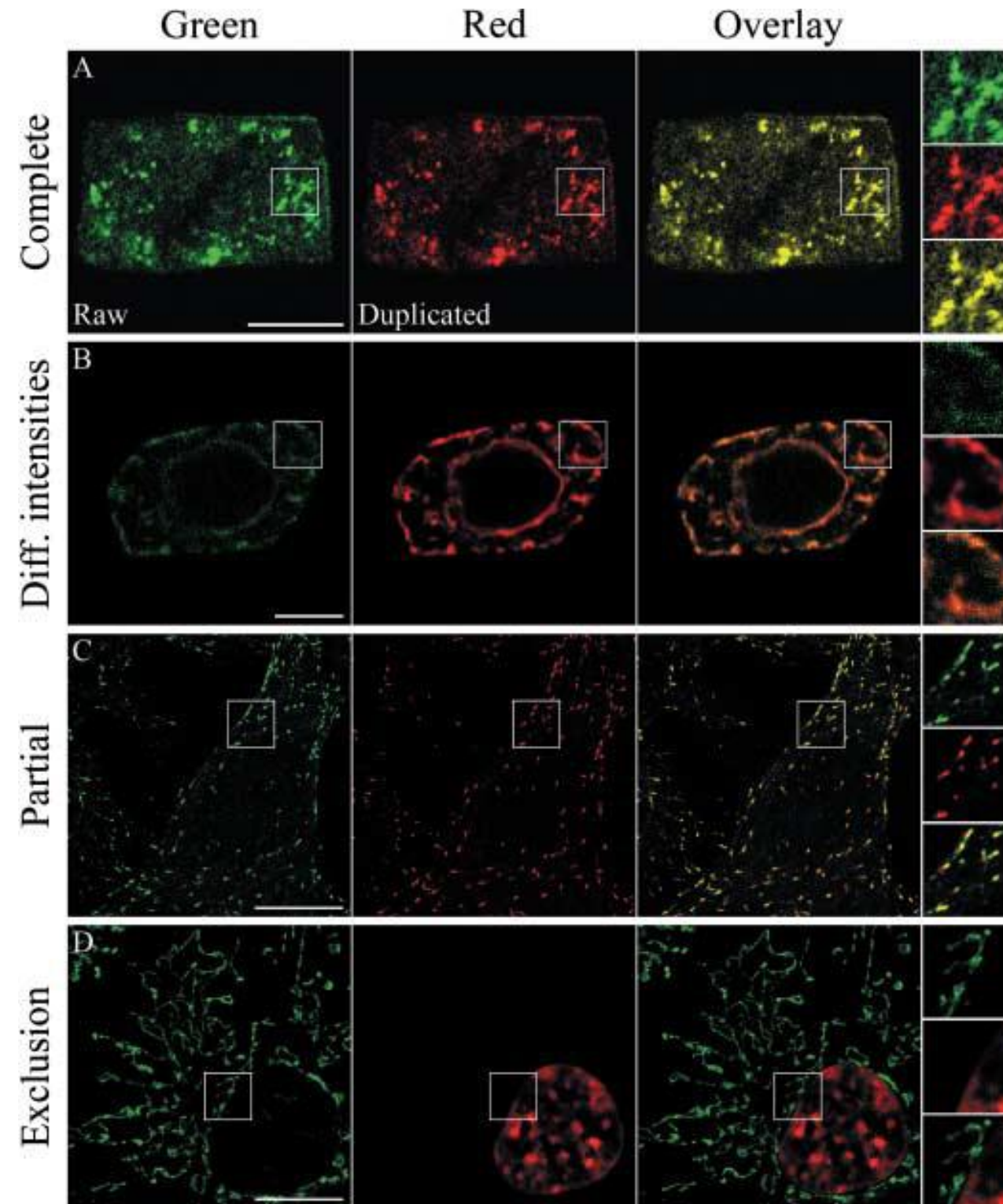


Colocalization Indicators

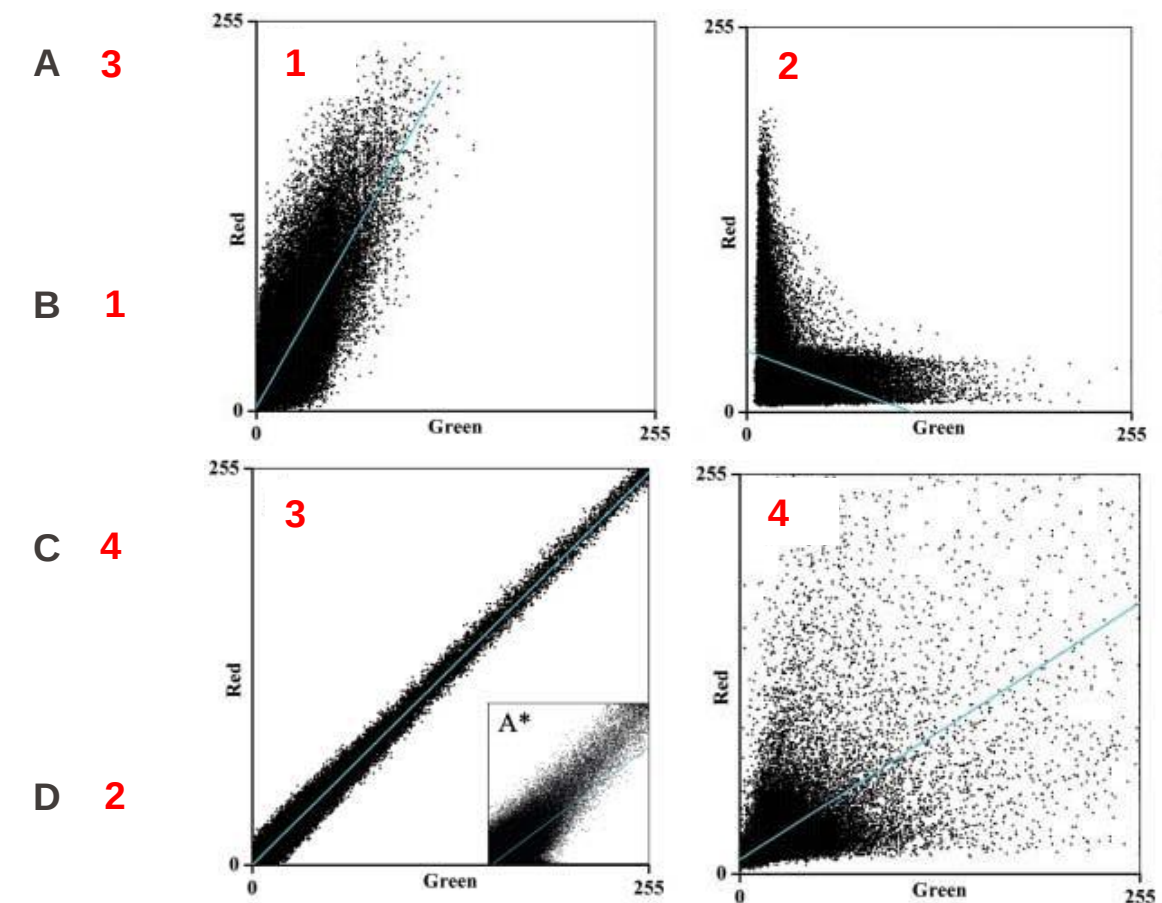
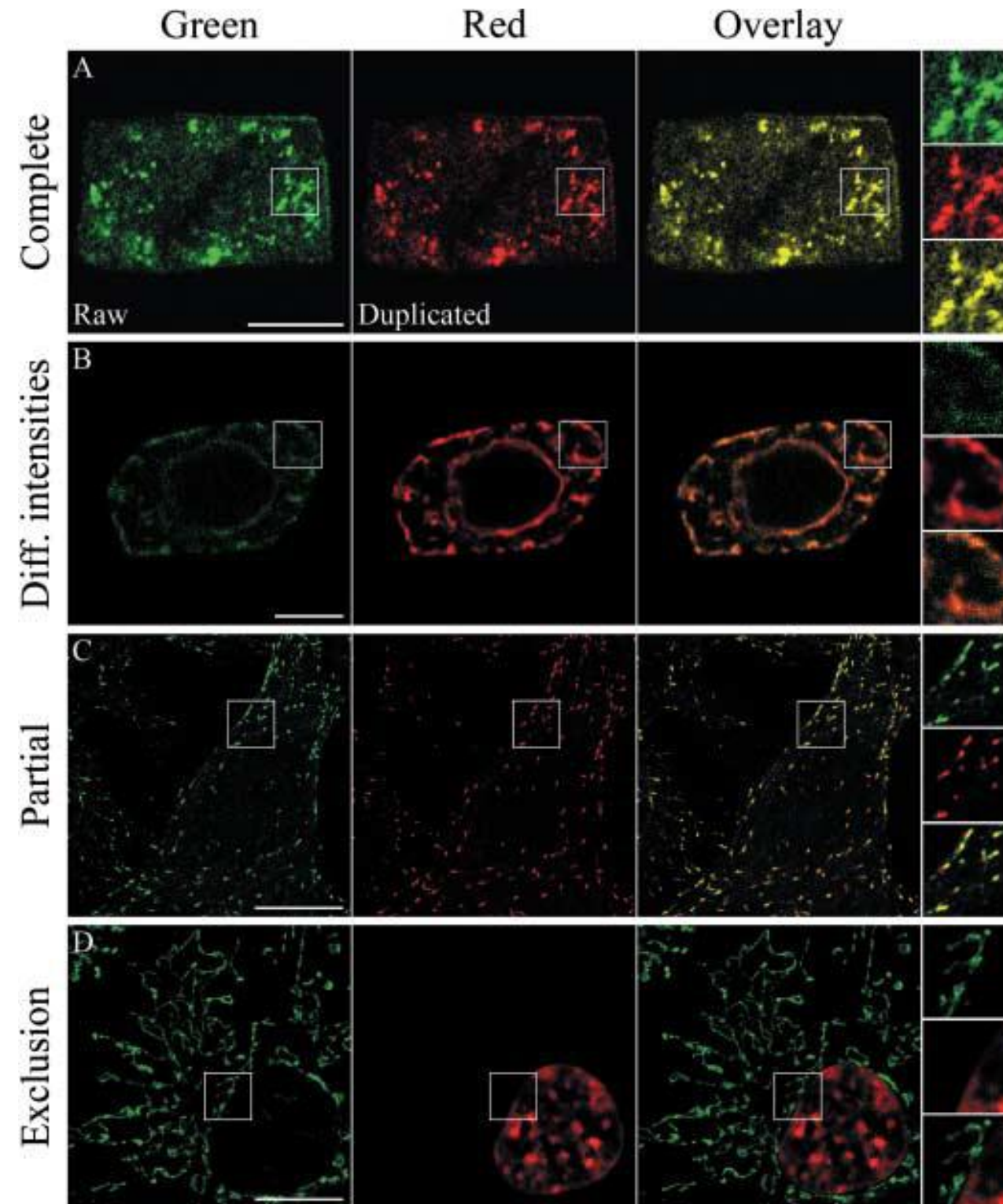
Pearson coefficient

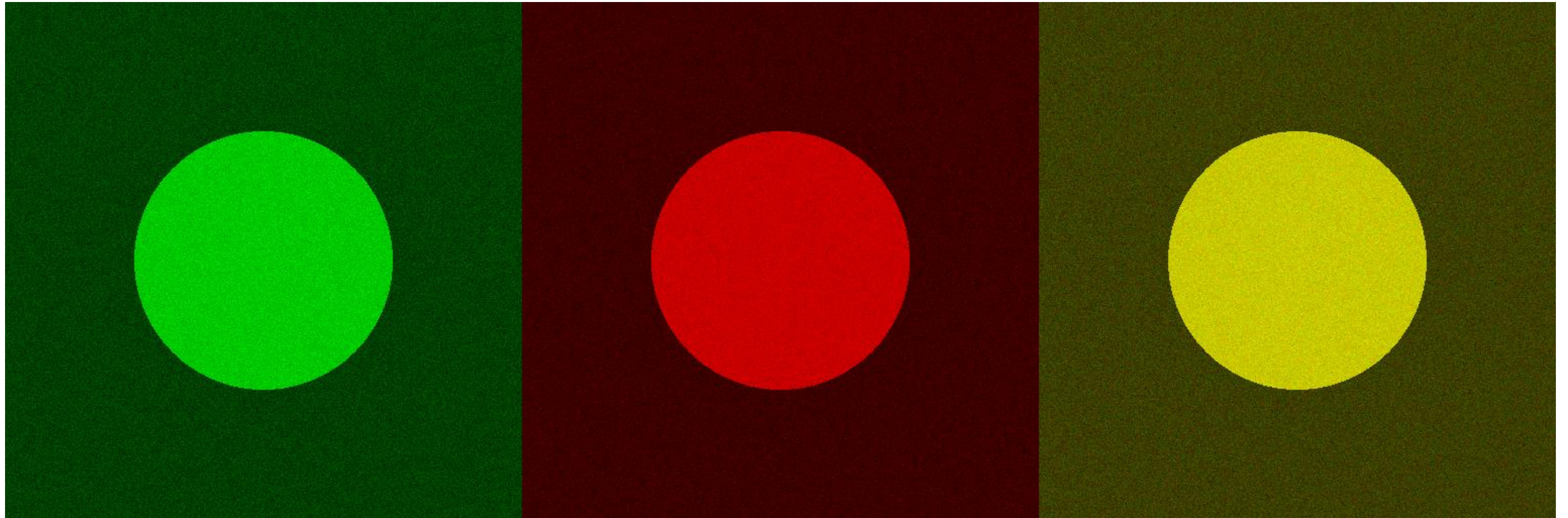
$$PCC = \frac{\sum (R_i - \bar{R}) \times (G_i - \bar{G})}{\sqrt{\sum (R_i - \bar{R})^2 \times \sum (G_i - \bar{G})^2}}$$

Which scatterplot belongs to which image sequence?



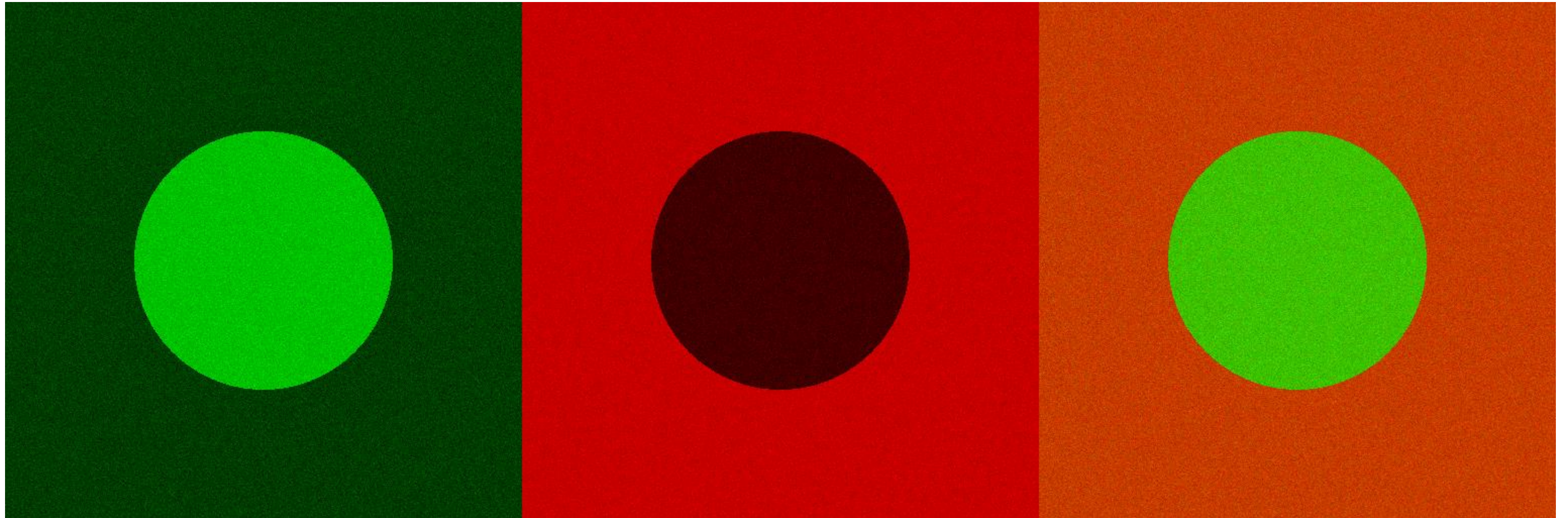
Which scatterplot belongs to which image sequence?



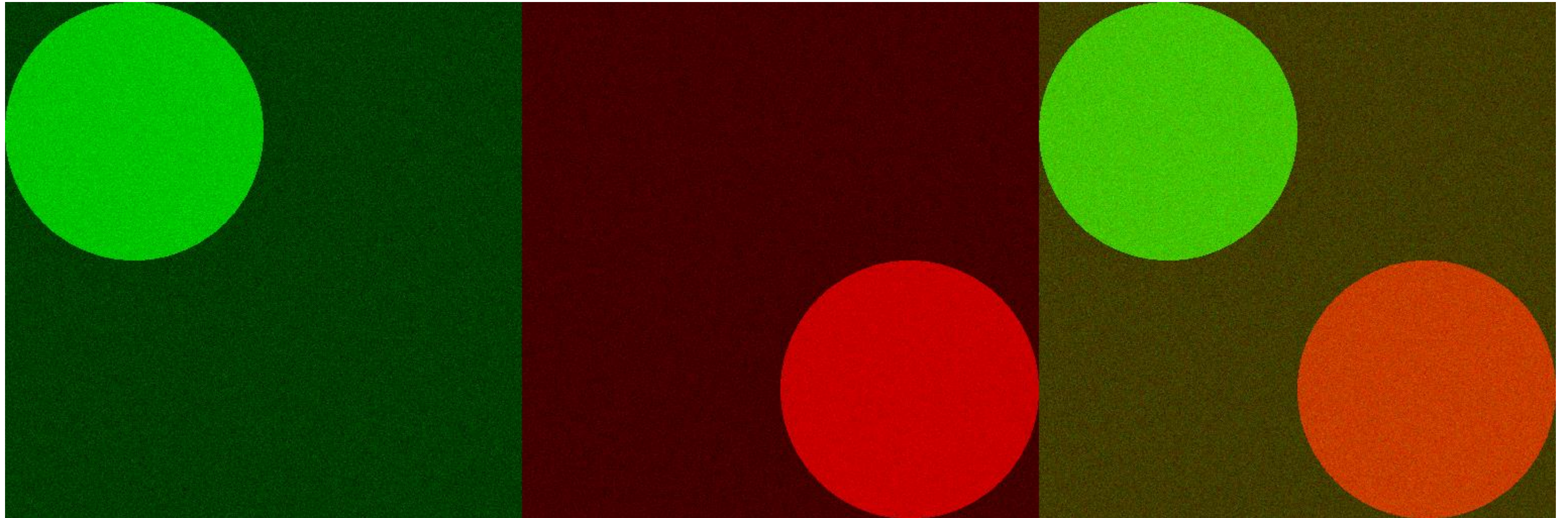


Pearson's Correlation Coefficient: 0.94

AntiCorrelation

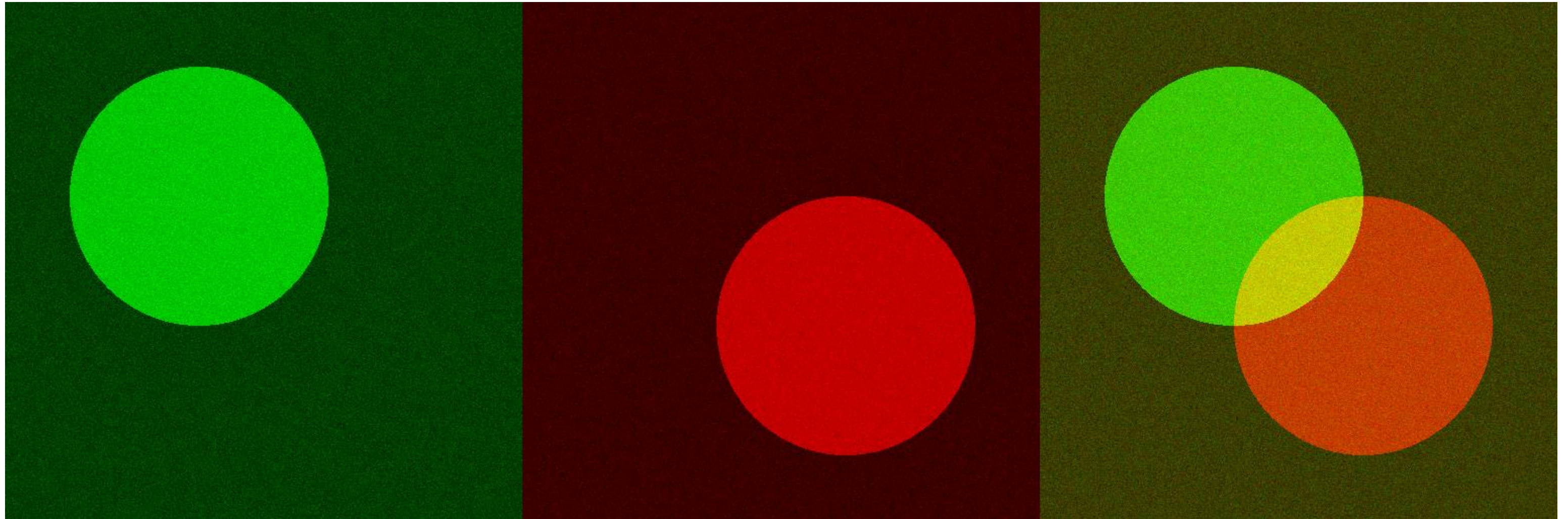


Pearson's Correlation Coefficient: -0.94



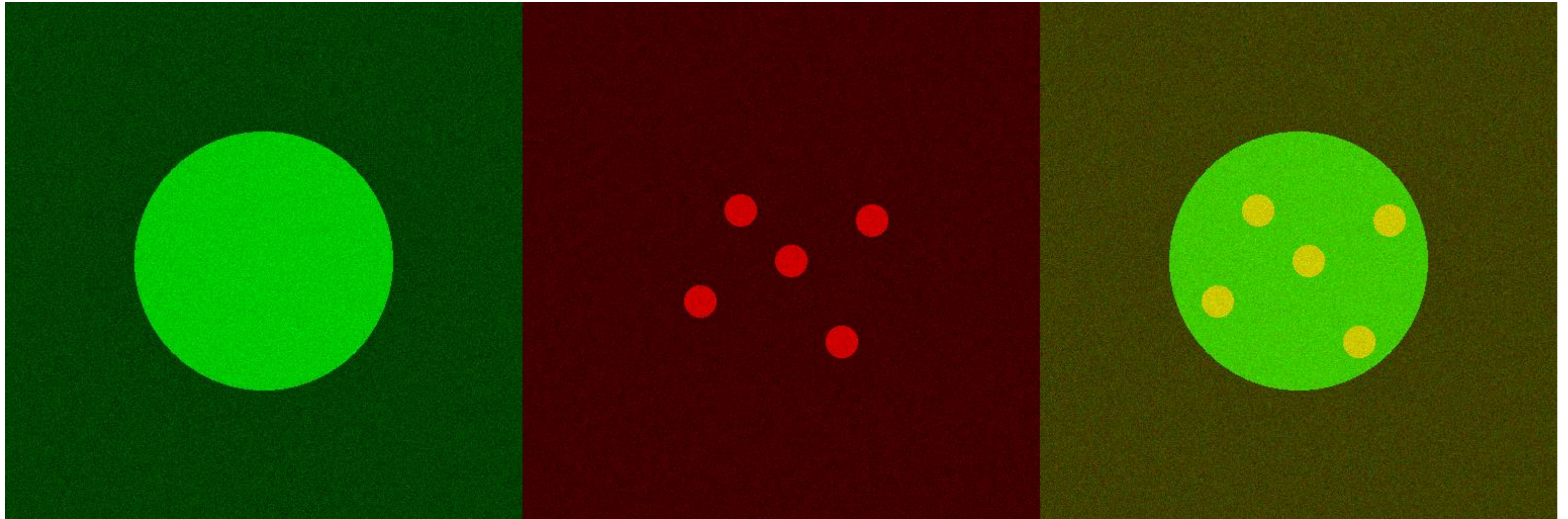
Pearson's Correlation Coefficient: -0.29

Partial Overlap



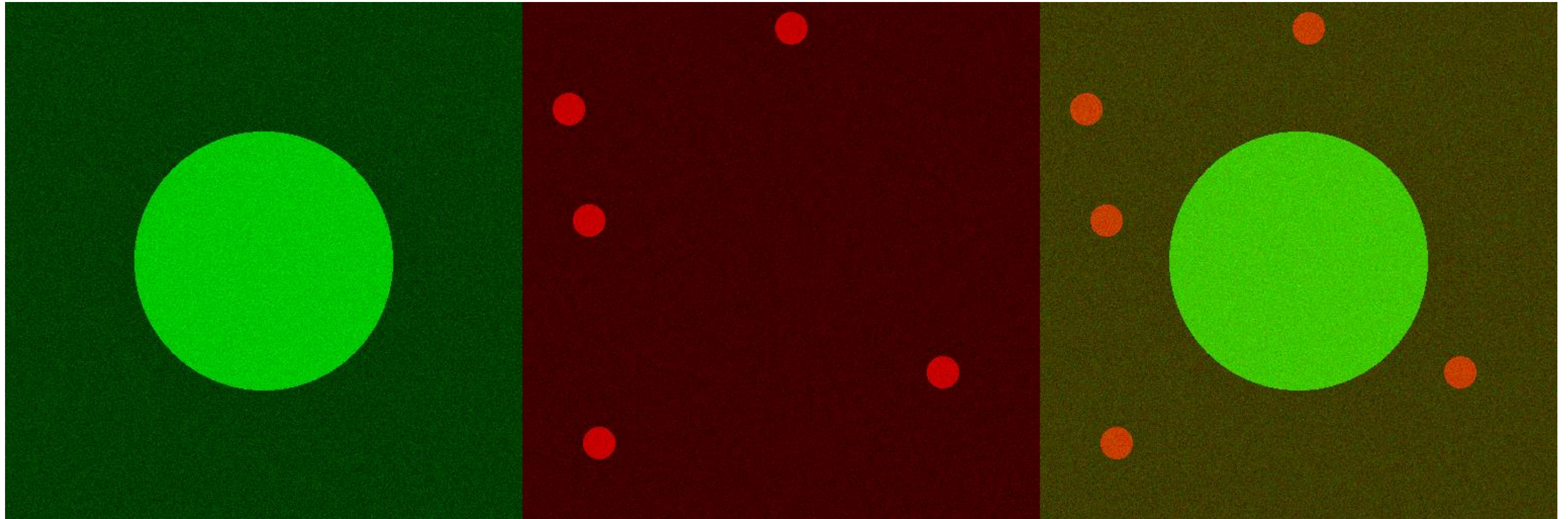
Pearson's Correlation Coefficient: -0.016

Big/Small Colocalization



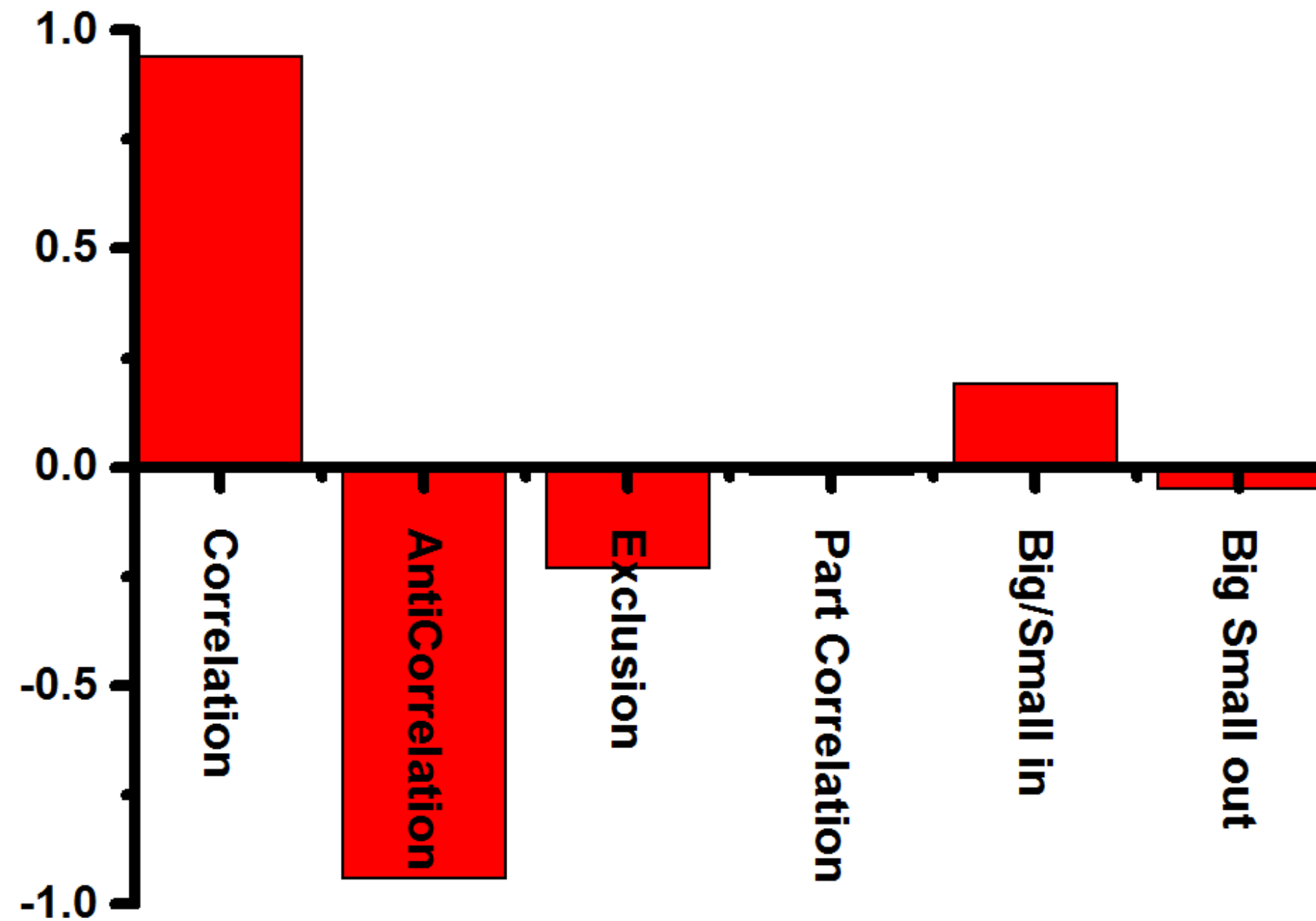
Pearson's Correlation Coefficient: 0.19

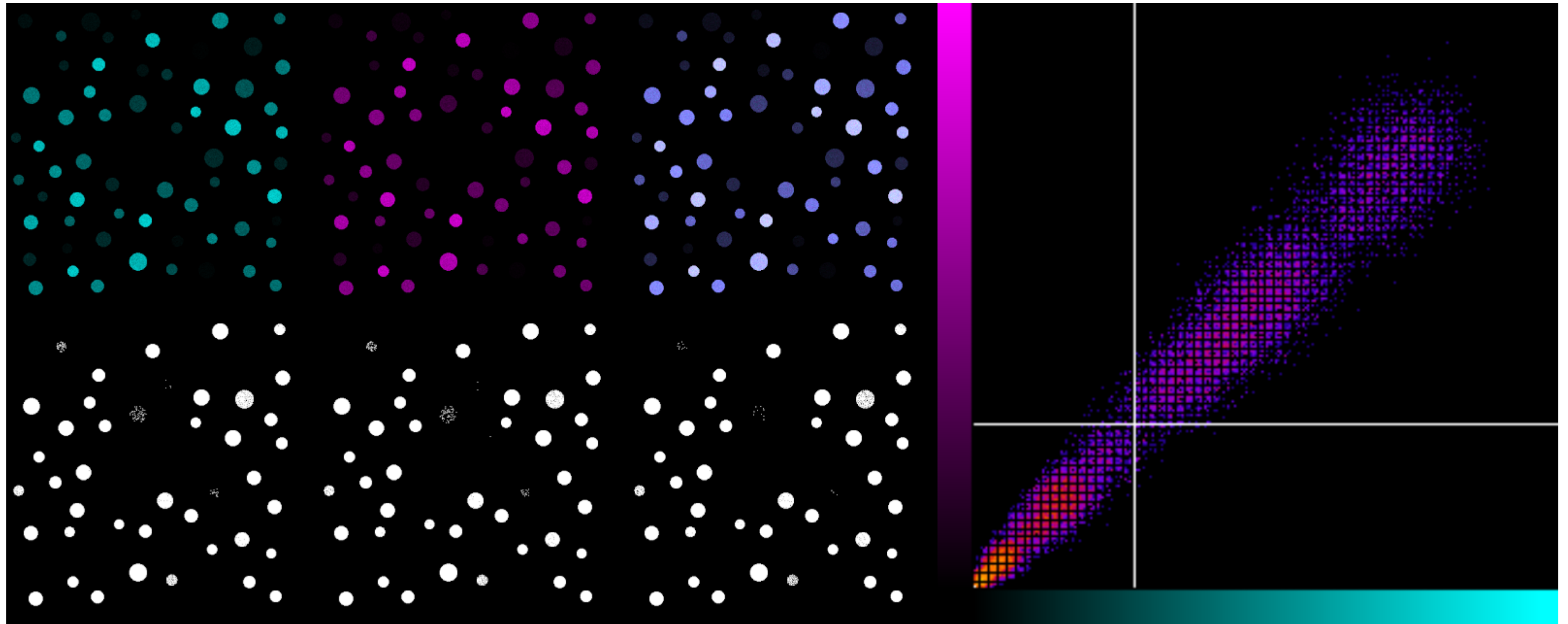
Big/Small Exclusion

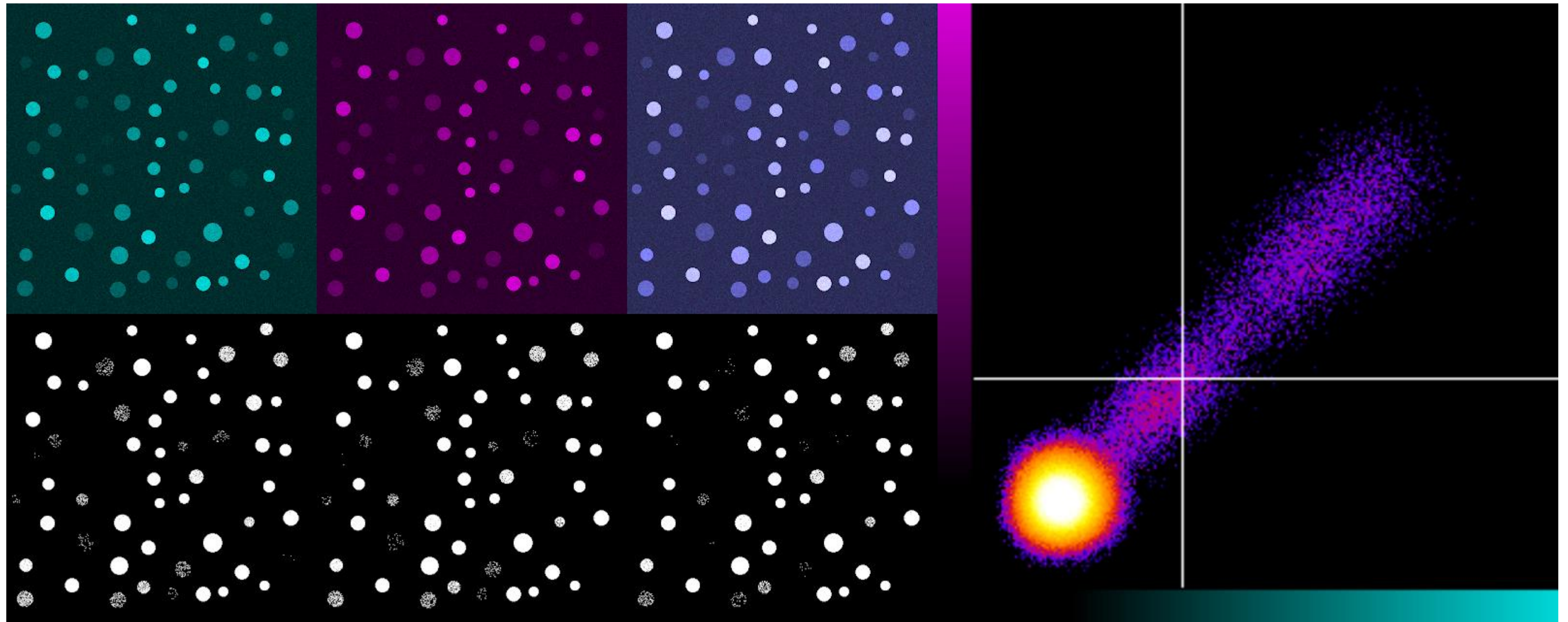


Pearson's Correlation Coefficient: -0.047

Pearson coefficient

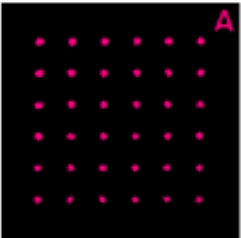
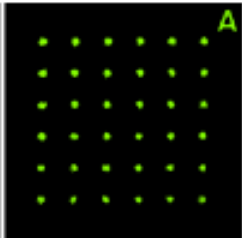
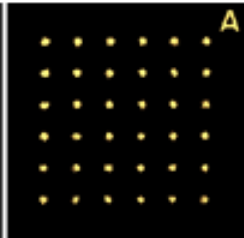
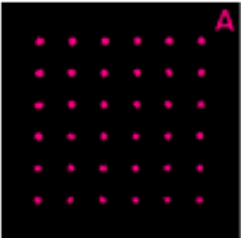
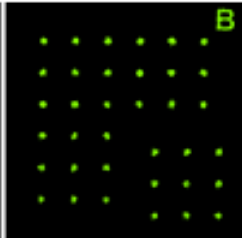
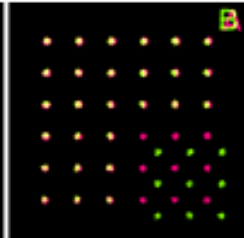
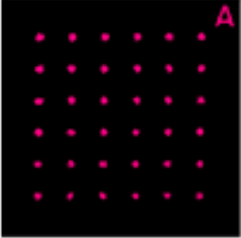
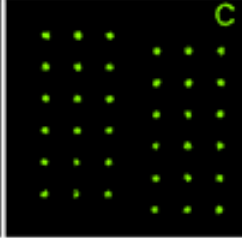
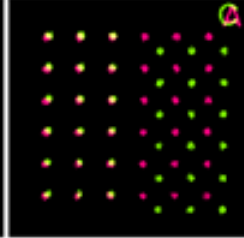
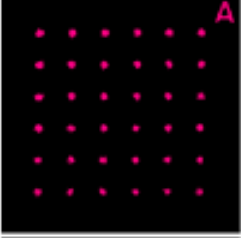
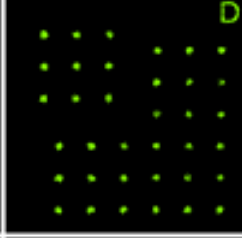
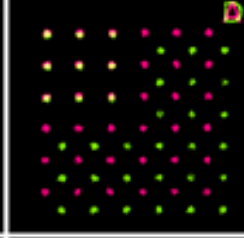
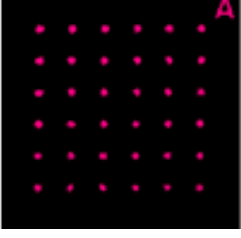
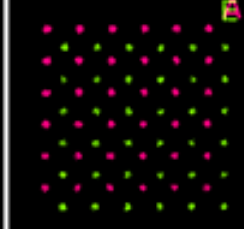




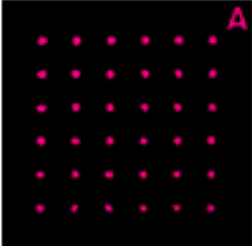
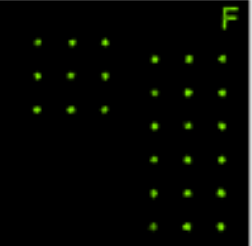
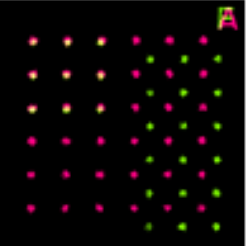
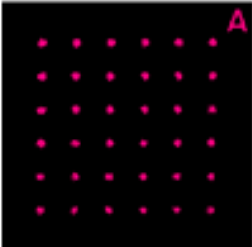
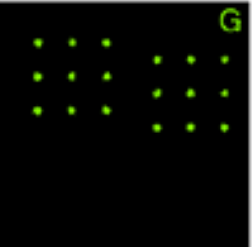
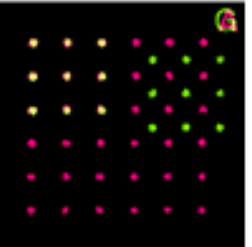
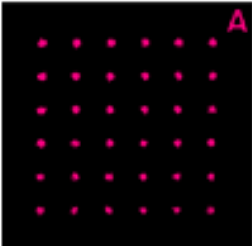
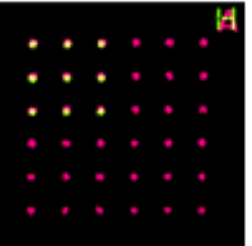
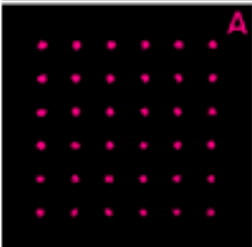
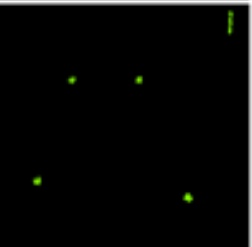
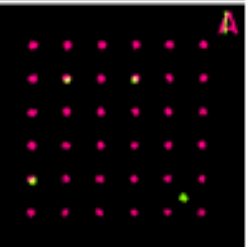


EPFL

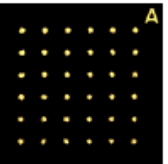
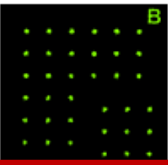
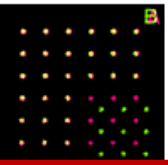
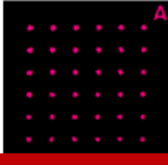
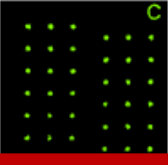
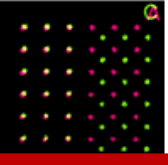
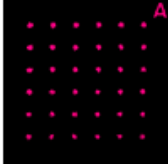
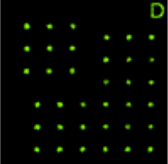
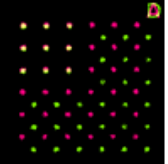
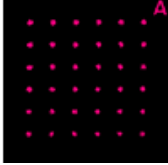
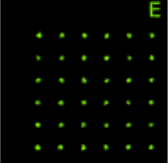
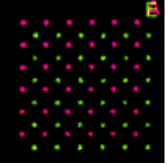
PCC limitation

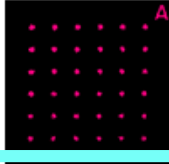
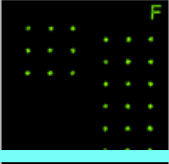
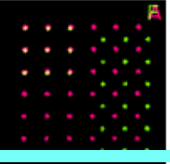
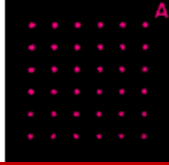
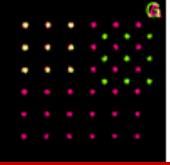
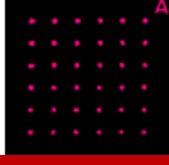
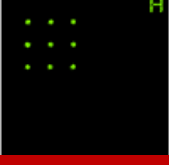
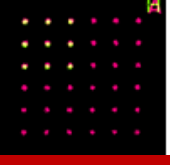
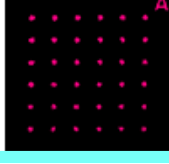
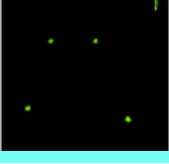
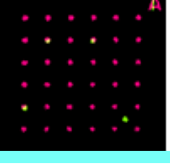
Case	Ch. 1	Ch.2	Merge	Spots Number			Pearson
				Ch. 1	Ch. 2	Co-loc.	
AA				36	36	36	1
AB				36	36	27	0.72
AC				36	36	18	0.44
AD				36	36	9	0.16
AE				36	36	0	-0.12

EPFL PCC limitation

Case	Ch. 1	Ch.2	Merge	Spots Number			Pearson
				Ch. 1	Ch. 2	Co-loc.	
AF				36	27	9	0.22
AG				36	18	9	0.30
AH				36	9	9	0.48
AI				36	4	3	0.23

PCC limitation

Case	Ch. 1	Ch. 2	Merge	Spots Number			Pearson
				Ch. 1	Ch. 2	Co-loc.	
AA				36	36	36	1
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Case	Ch. 1	Ch. 2	Merge	Spots Number			Pearson
				Ch. 1	Ch. 2	Co-loc.	
AF				36	27	9	0.22
AG				36	18	9	0.30
AH				36	9	9	0.48
AI				36	4	3	0.23

Colocalization Indicators

$$PCC = \frac{\sum (R_i - \bar{R}) \times (G_i - \bar{G})}{\sqrt{\sum (R_i - \bar{R})^2 \times \sum (G_i - \bar{G})^2}}$$

$$overlap = \frac{\sum R_i \times G_i}{\sqrt{\sum R_i^2 \times \sum G_i^2}} \quad k_1 = \frac{\sum R_i \times G_i}{\sqrt{\sum R_i^2}}; k_2 = \frac{\sum R_i \times G_i}{\sqrt{\sum G_i^2}}$$

R_i, G_i : intensity of the pixel i in image R and G respectively

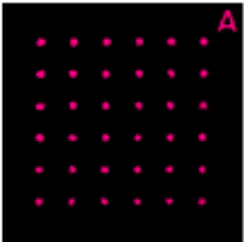
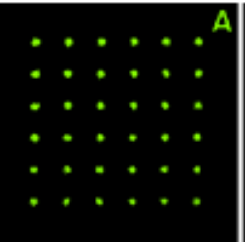
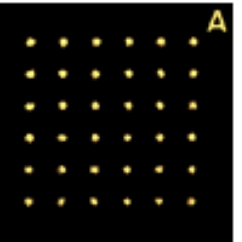
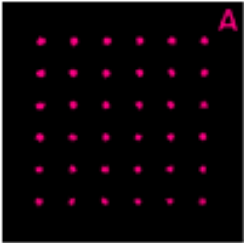
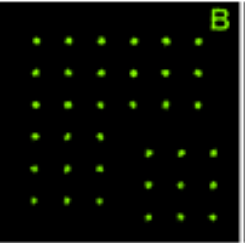
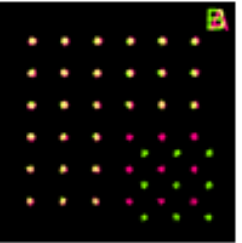
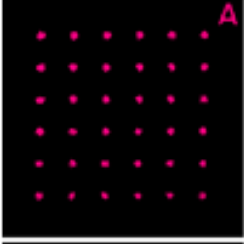
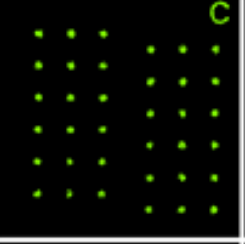
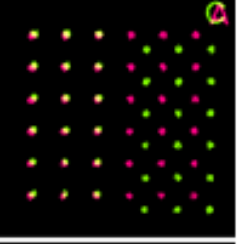
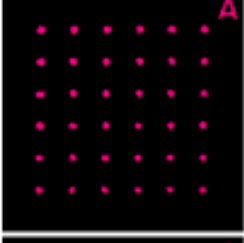
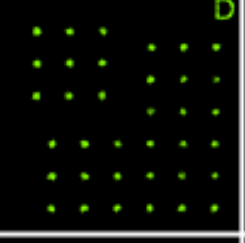
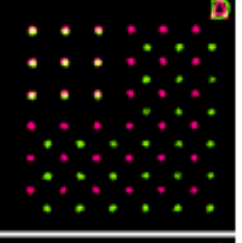
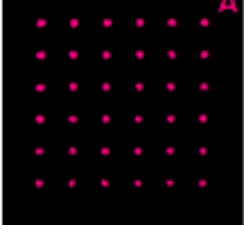
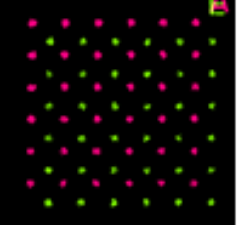
$\bar{R}, \bar{G}$: average intensity of the image R and G respectively

$$M_1 = \frac{\sum R_{i,coloc}}{\sum R_i}; M_2 = \frac{\sum G_{i,coloc}}{\sum G_i}$$

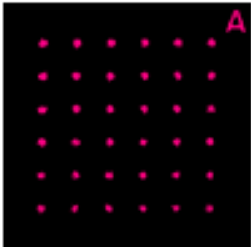
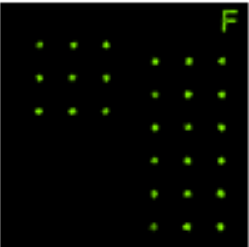
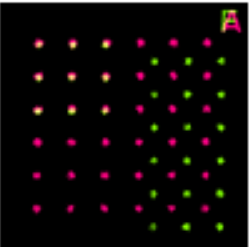
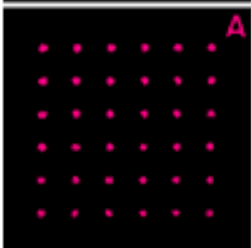
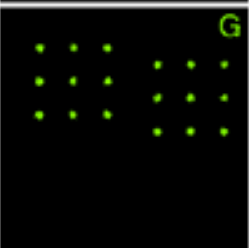
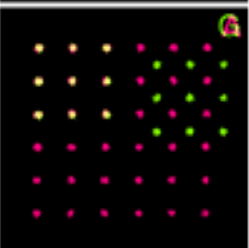
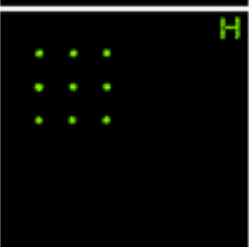
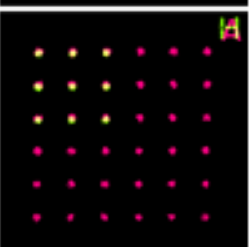
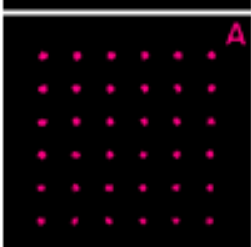
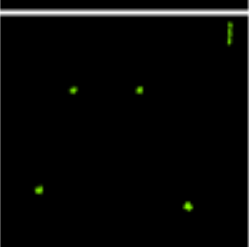
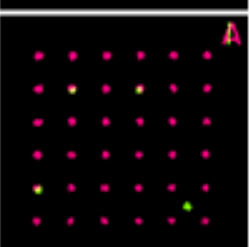
$$R_{i,coloc} = R_i \quad \text{if } G_i > 0$$

$$G_{i,coloc} = G_i \quad \text{if } R_i > 0$$

Mander's Coefficient

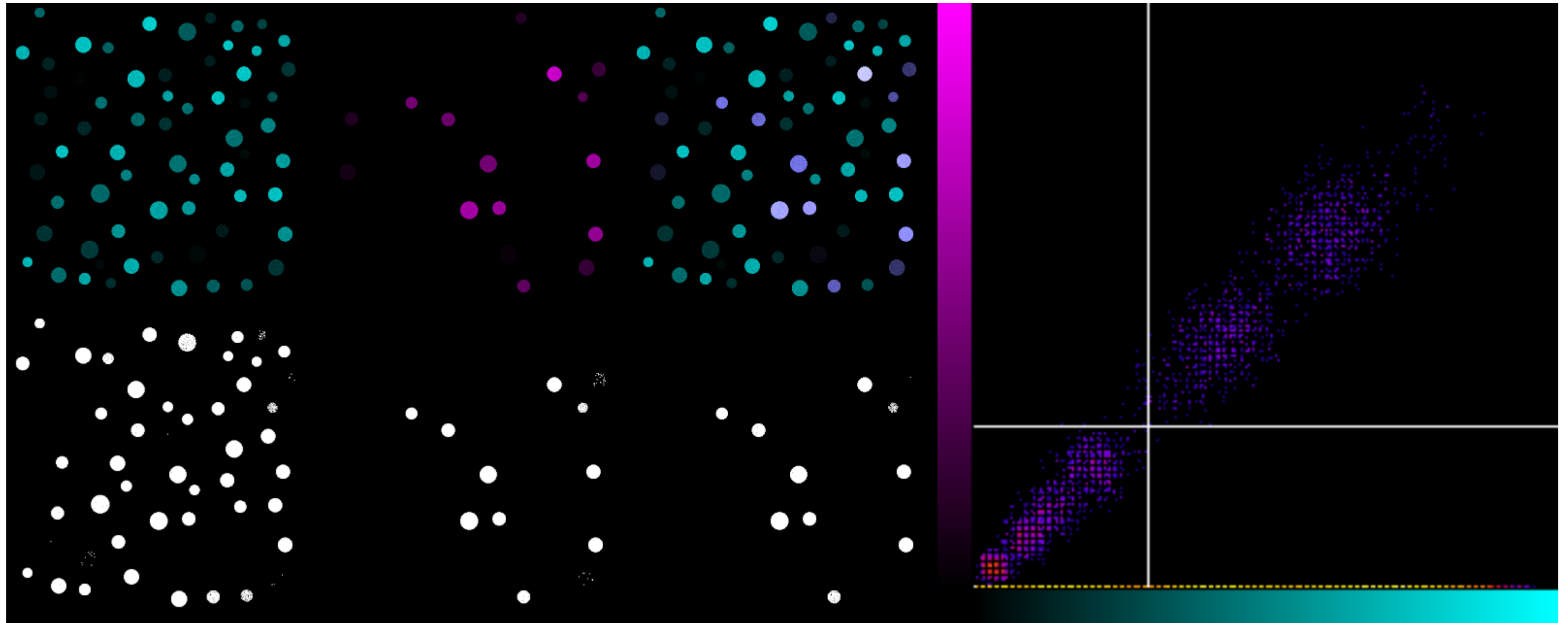
Case	Ch. 1	Ch.2	Merge	Spots Number			Pearson	Mander	
				Ch. 1	Ch. 2	Co-loc.		Ch. 1 = M1	Ch. 2 = M2
AA				36	36	36	1	1	1
AB				36	36	27	0.72	0.75	0.75
AC				36	36	18	0.44	0.5	0.5
AD				36	36	9	0.16	0.25	0.25
AE				36	36	0	-0.12	0	0

Mander's Coefficient

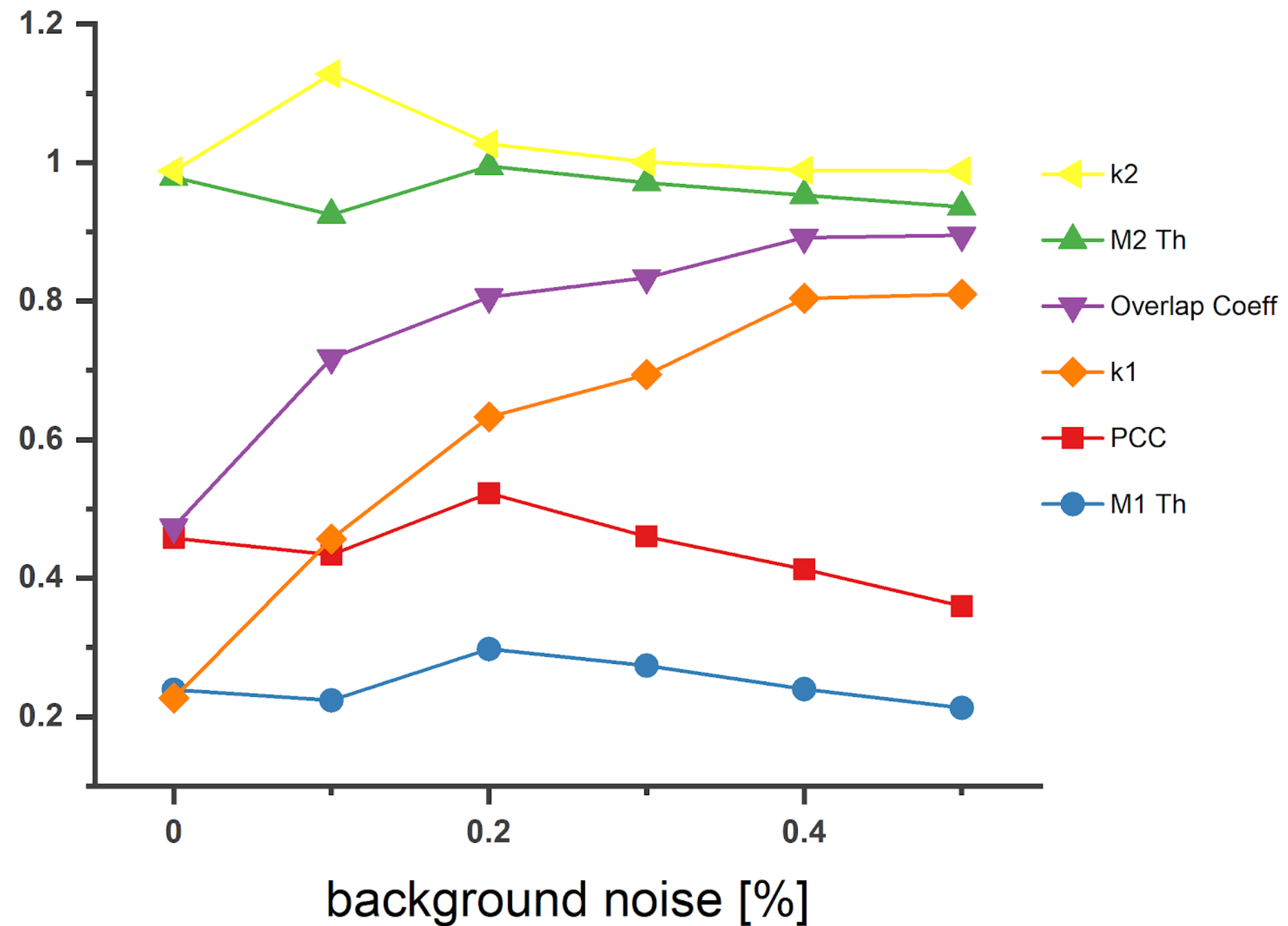
Case	Ch. 1	Ch.2	Merge	Spots Number			Pearson	Mander	
				Ch. 1	Ch. 2	Co-loc.		Ch. 1 = M1	Ch. 2 = M2
AF				36	27	9	0.22	0.25	0.33
AG				36	18	9	0.30	0.25	0.50
AH				36	9	9	0.48	0.25	1.0
AI				36	4	3	0.23	0.08	0.75

Colocalization and noise

PCC: 0.49 M_1 :0.239 M_2 :0.979



Colocalization and noise

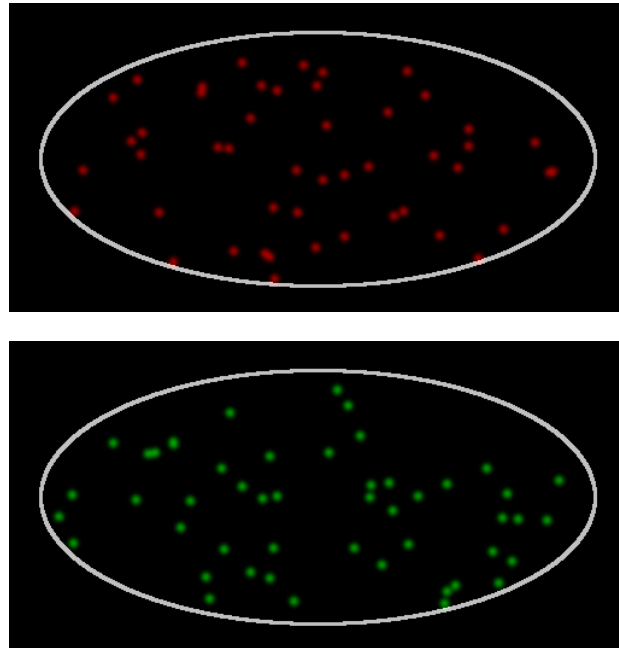


Pixel based Colocalization

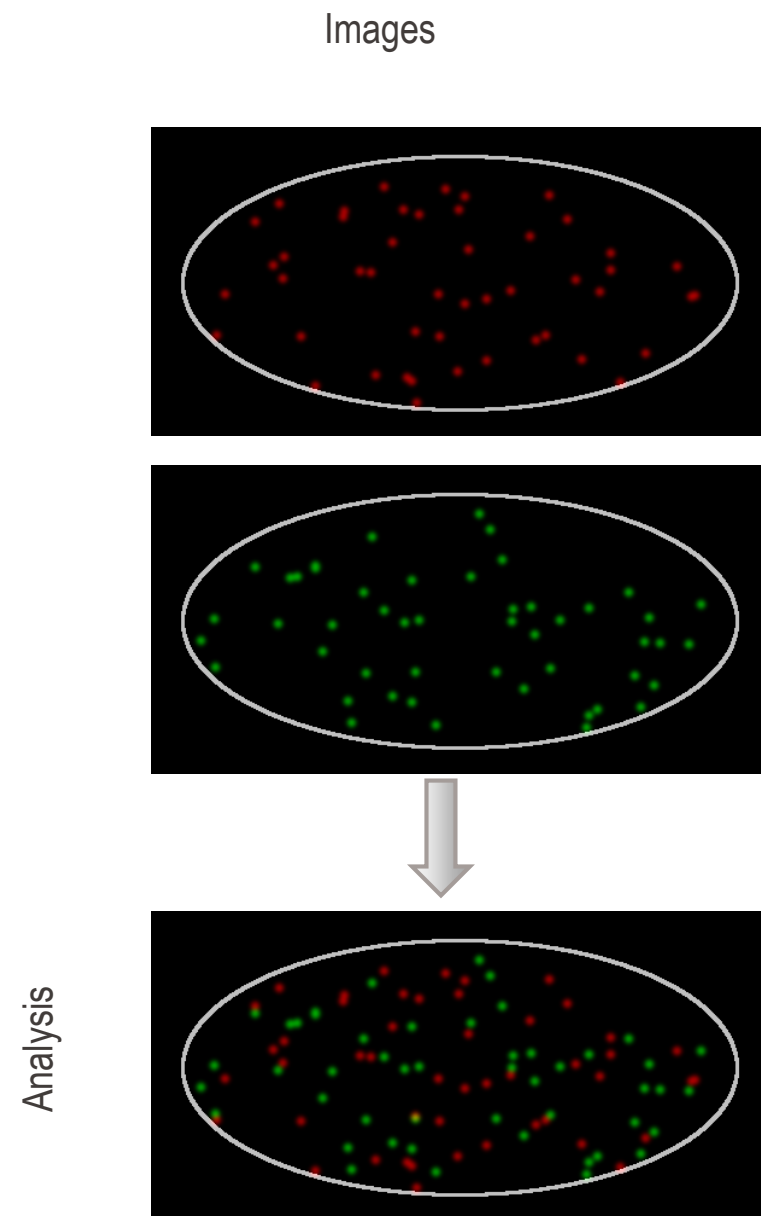
- Two different molecules can never be at the same physical spot at the same time.
- Colocalization seen in images is coming from the low-pass filtering of the image formation in light microscopy.
- Colocalization is an artificial phenomenon and therefore always relative (e.g. it scales with noise, resolution, object size).
- The coefficient used needs to fit your experiment.
- Coefficients are well suited for cell-based analyses.

Colocalisation Workflow

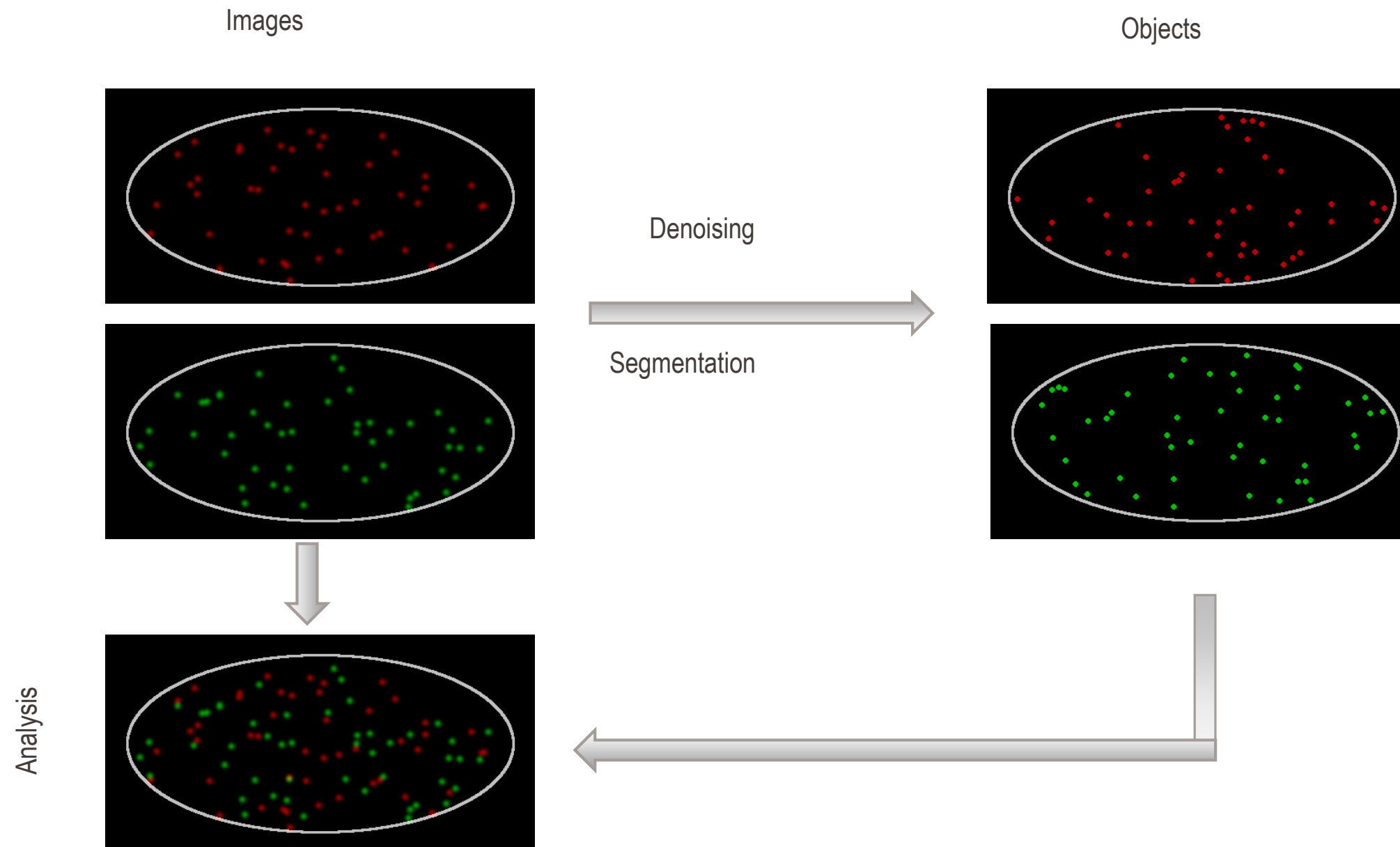
Images



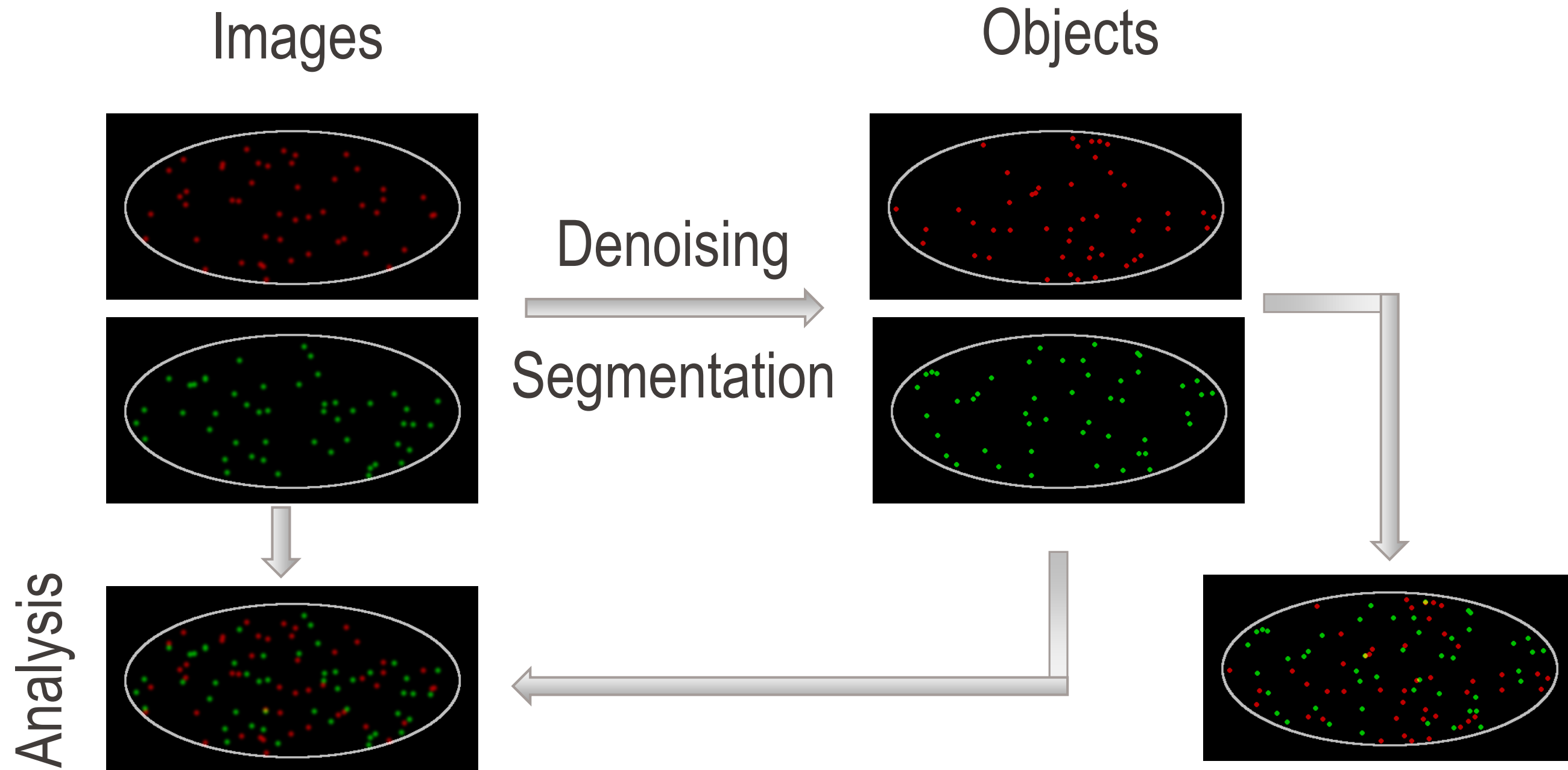
Colocalisation Workflow



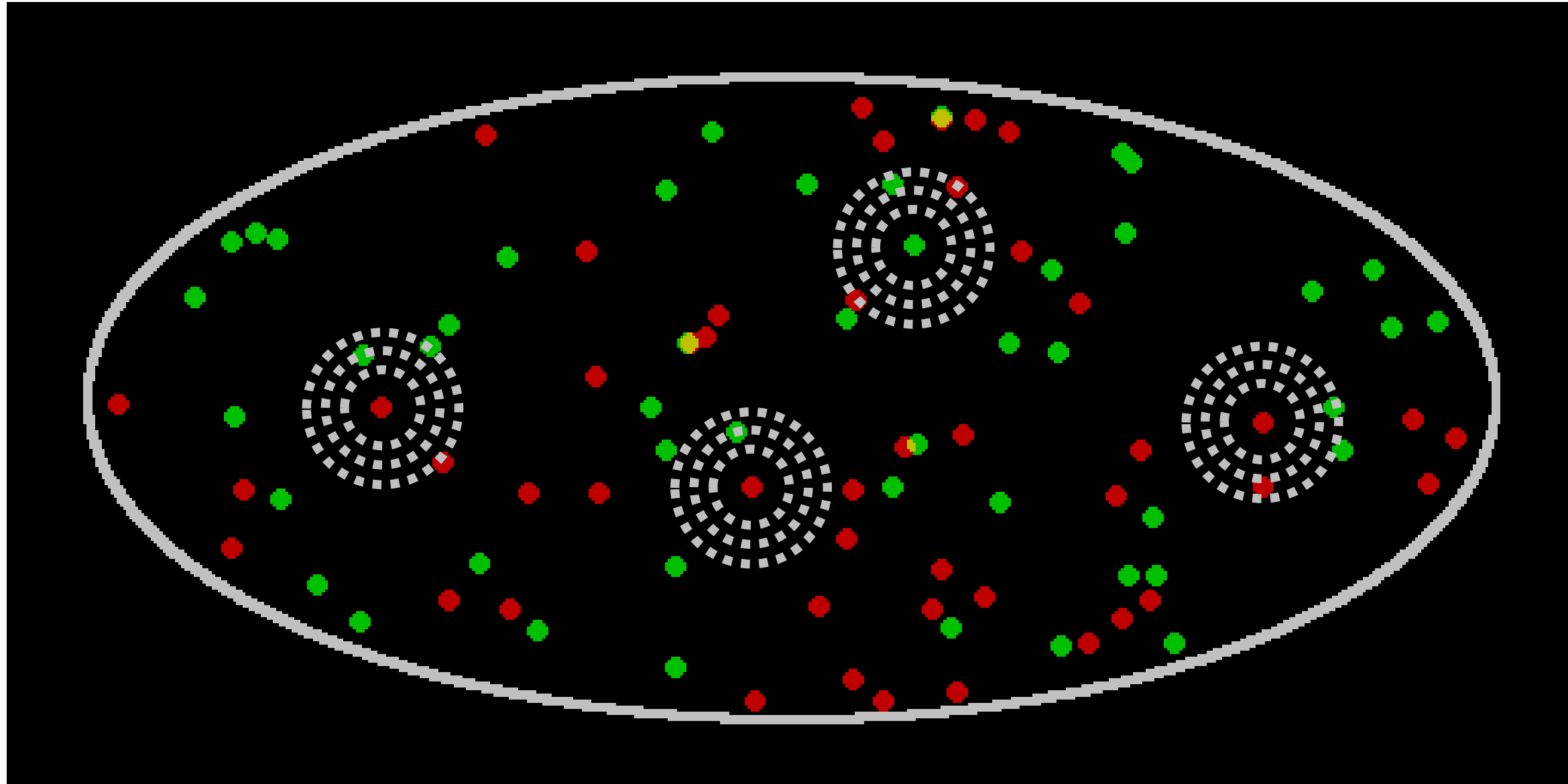
Colocalisation Workflow



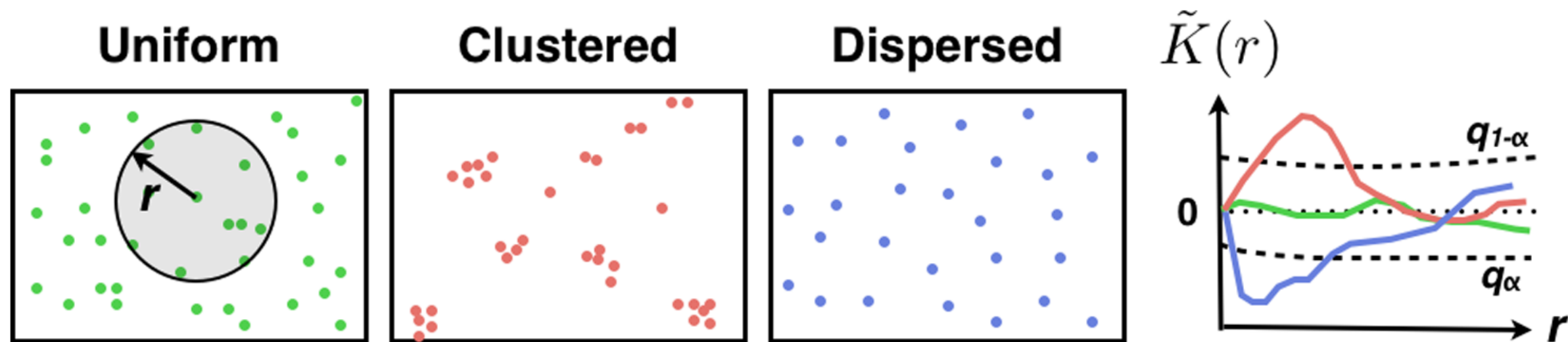
Object Based Colocalisation



Neighbourhood analysis



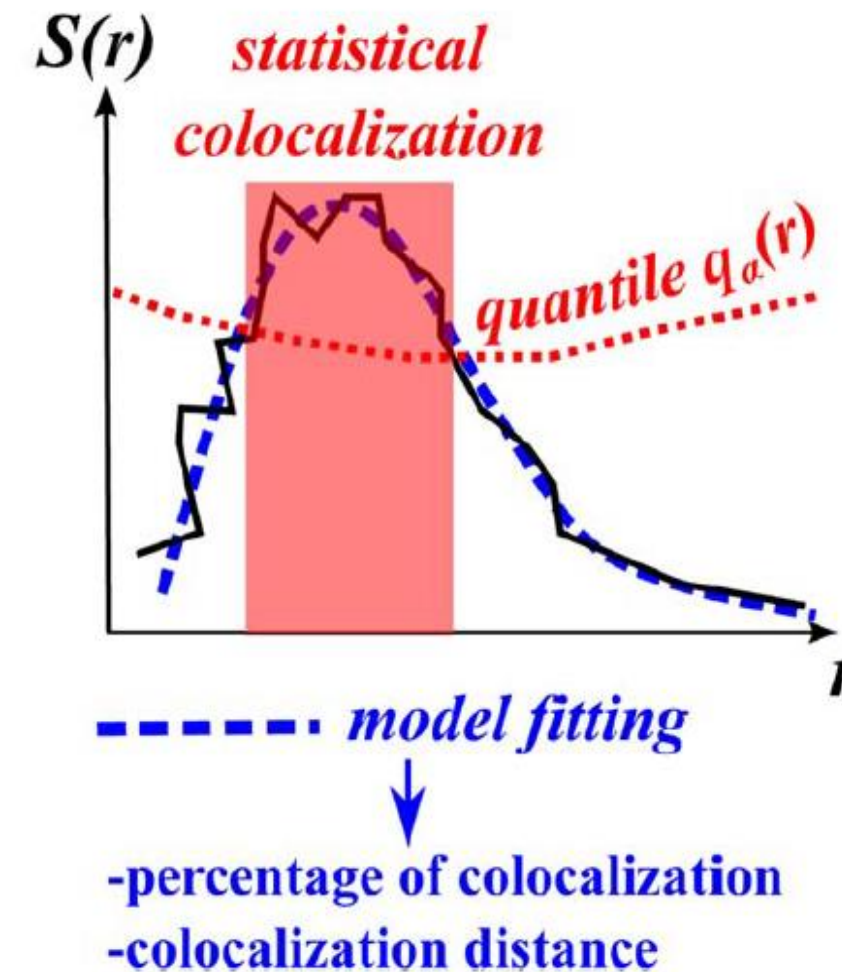
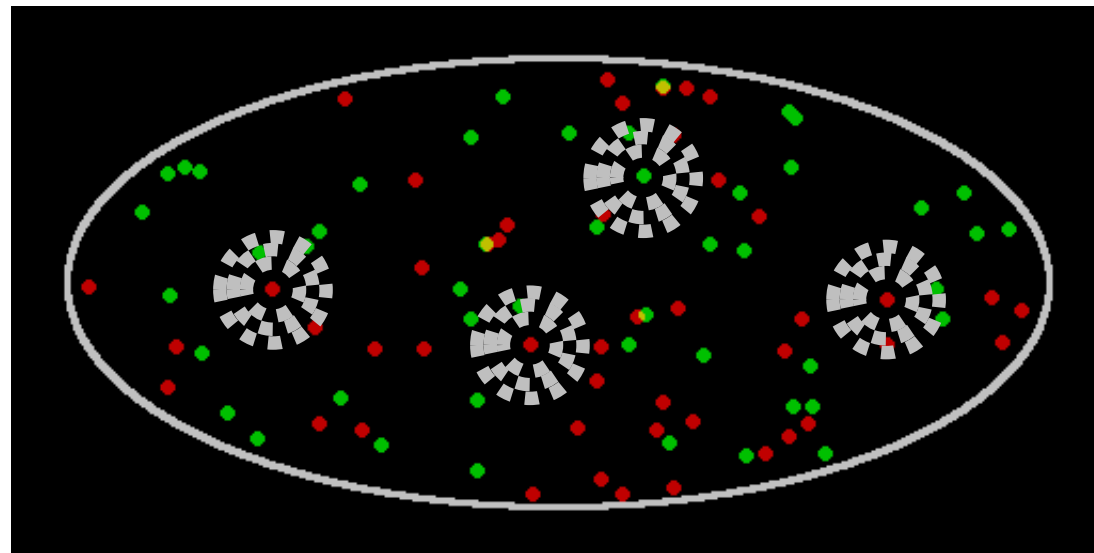
Ripley's K-function



Lagache T, Lang G, Sauvonnet N, Olivo-Marin JC.
PLoS One. 2013 Dec 4;8(12):e80914. doi: 10.1371/journal.pone.0080914. eCollection 2013.

Neighbourhood analysis

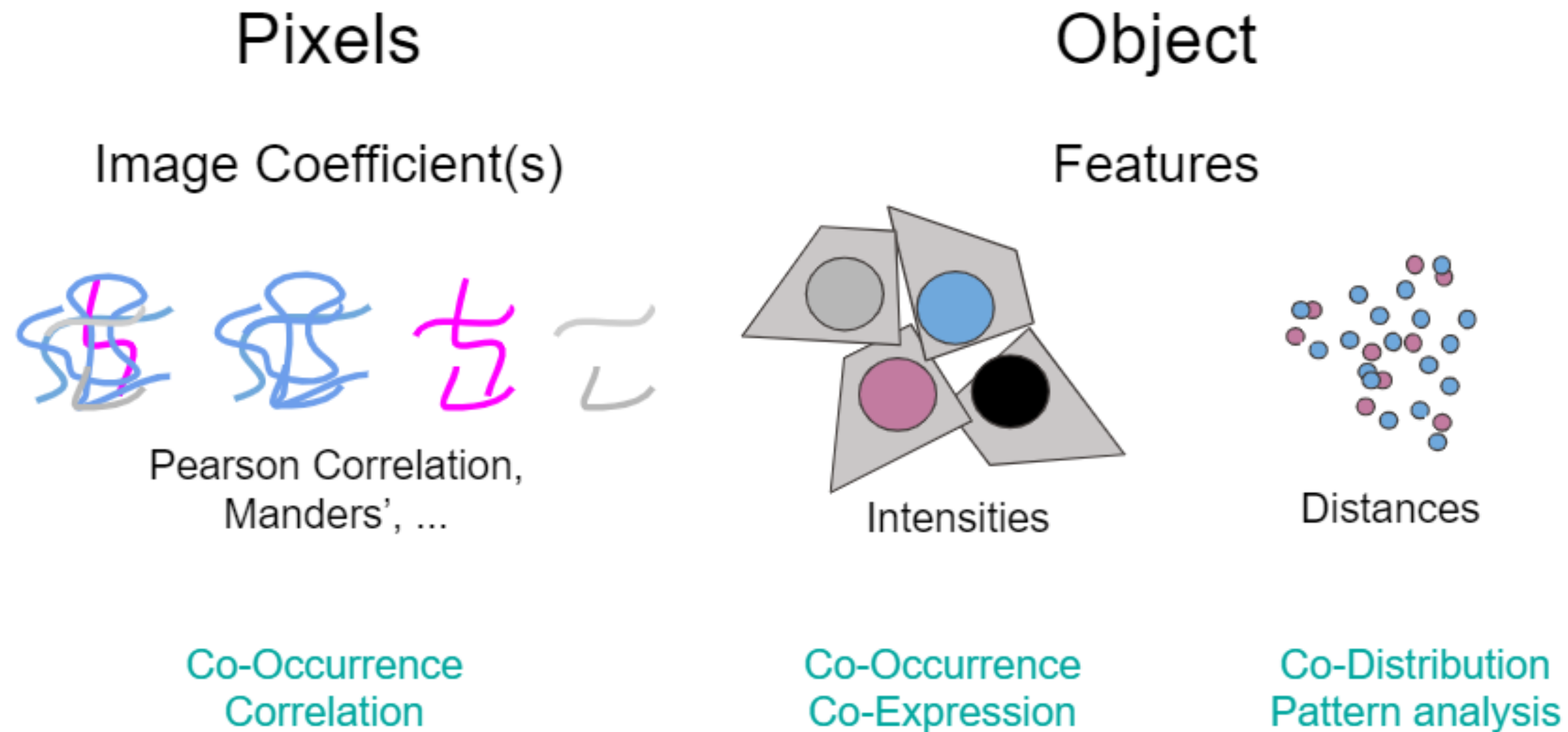
- Nearest Neighbour
- Ripley's K-function



Lagache T, Sauvonnnet N, Danglot L, Olivo-Marin JC.
Cytometry A. 2015 Jun;87(6):568-79. doi: 10.1002/cyto.a.22629.

Lagache T, Grassart A, Dallongeville S, Faklaris O, Sauvonnnet N, Dufour A, Danglot L, Olivo-Marin JC.
Nat Commun. 2018 Feb 15;9(1):698. doi: 10.1038/s41467-018-03053-x.

Co-Localization Summary



Colocalization Summary

