

Spatial statistics: Object-based colocalization





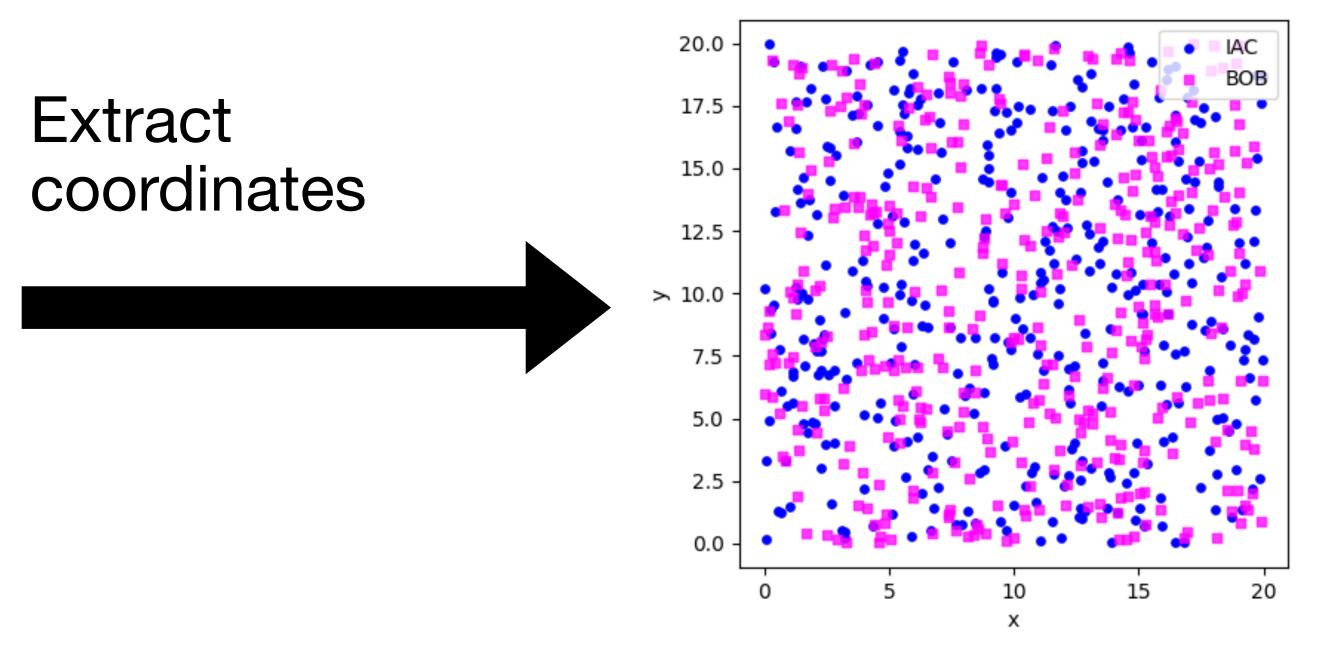


The data



IAC and BOB are proteins in the eastern spruce budworm (*Choristoneura fumiferana*) epidermis

You hypothesize that the spatial interaction between IAC and BOB changes with temperature and season.

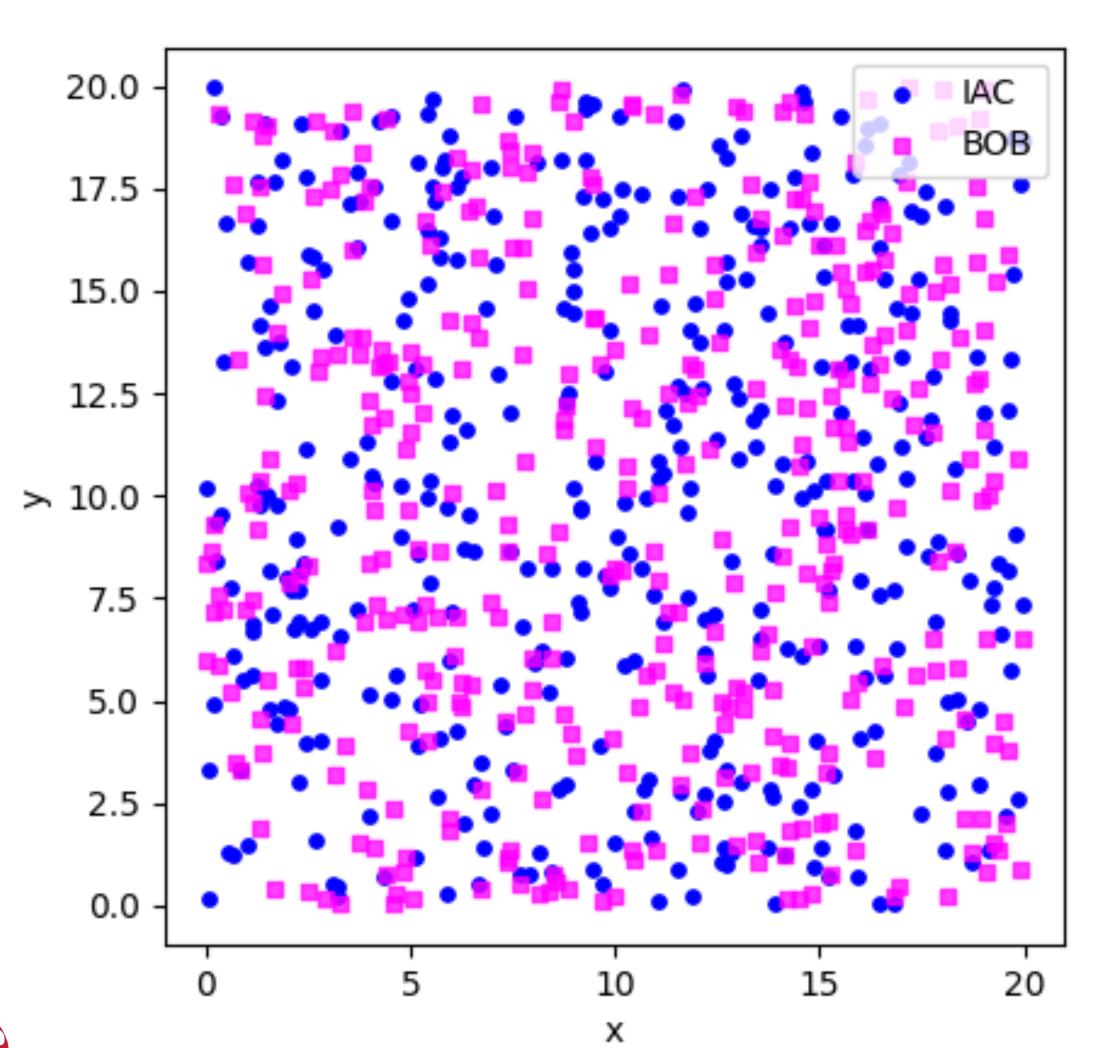




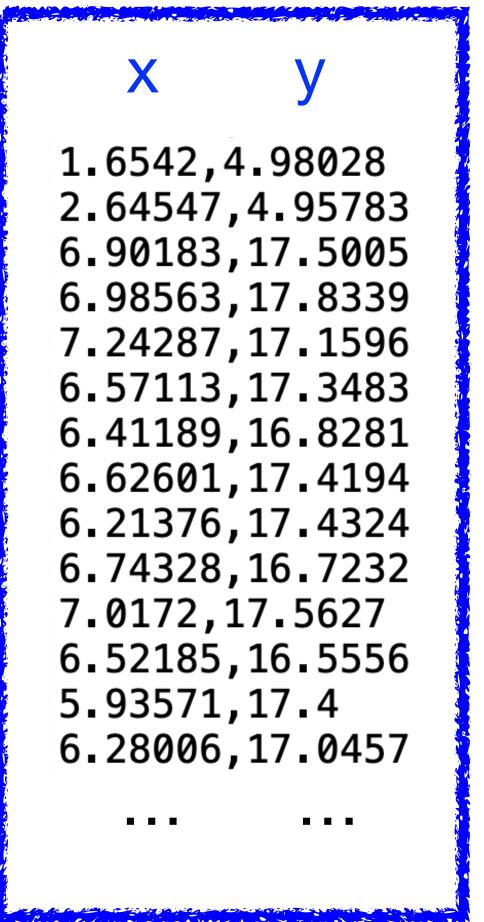




The data



IAC



BOB

X	y
2.59176, 2.35522, 3.60981, 2.91734, 2.46703, 2.60448, 1.24649, 3.67557, 2.63406, 2.77047, 2.90153, 4.02456,	12.7033 12.9357 12.0081 12.7667 11.849 12.6463 11.4218 4.29607 4.7991 4.19997 4.83014 4.98462







The data — Fall epidermis samples

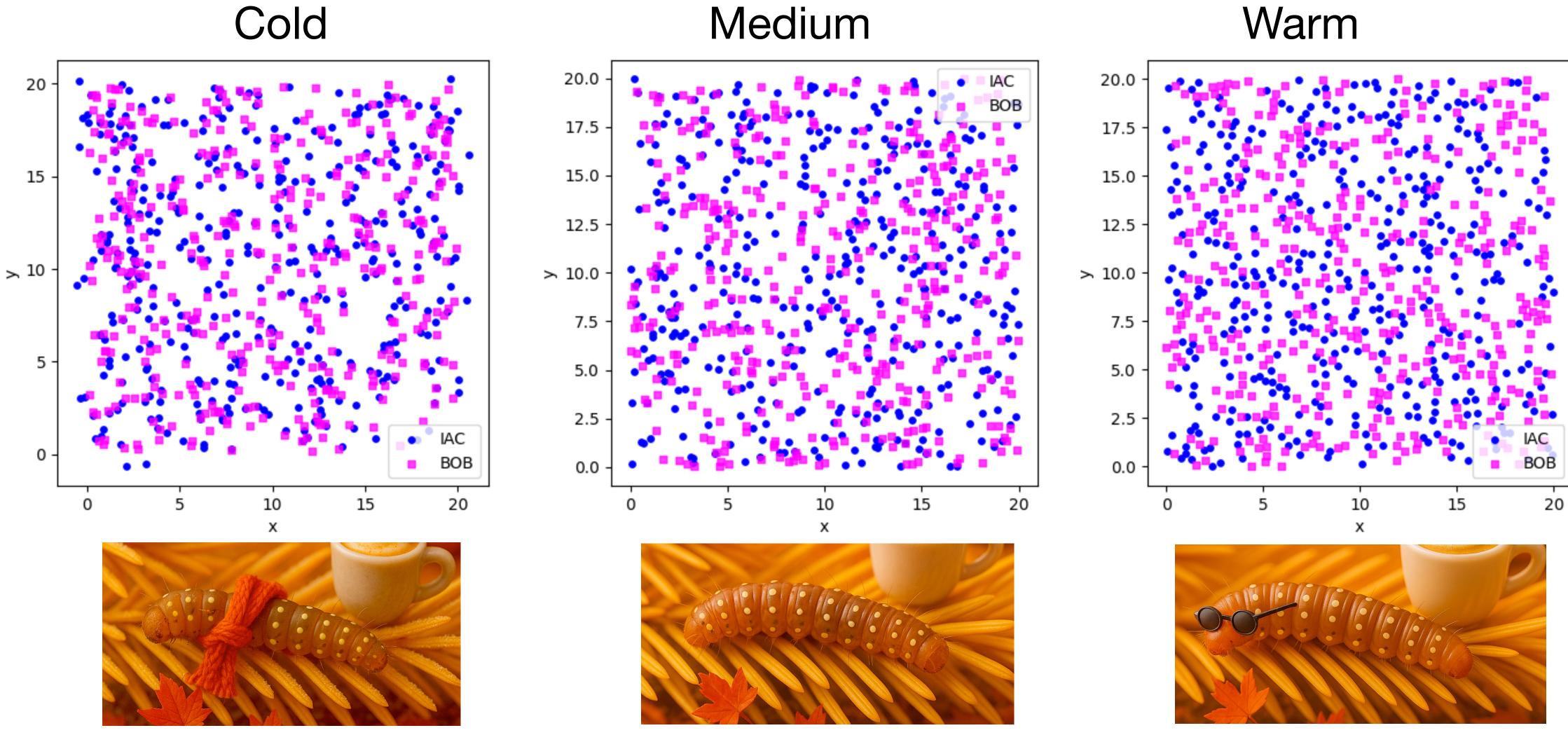








The data — Fall epidermis samples

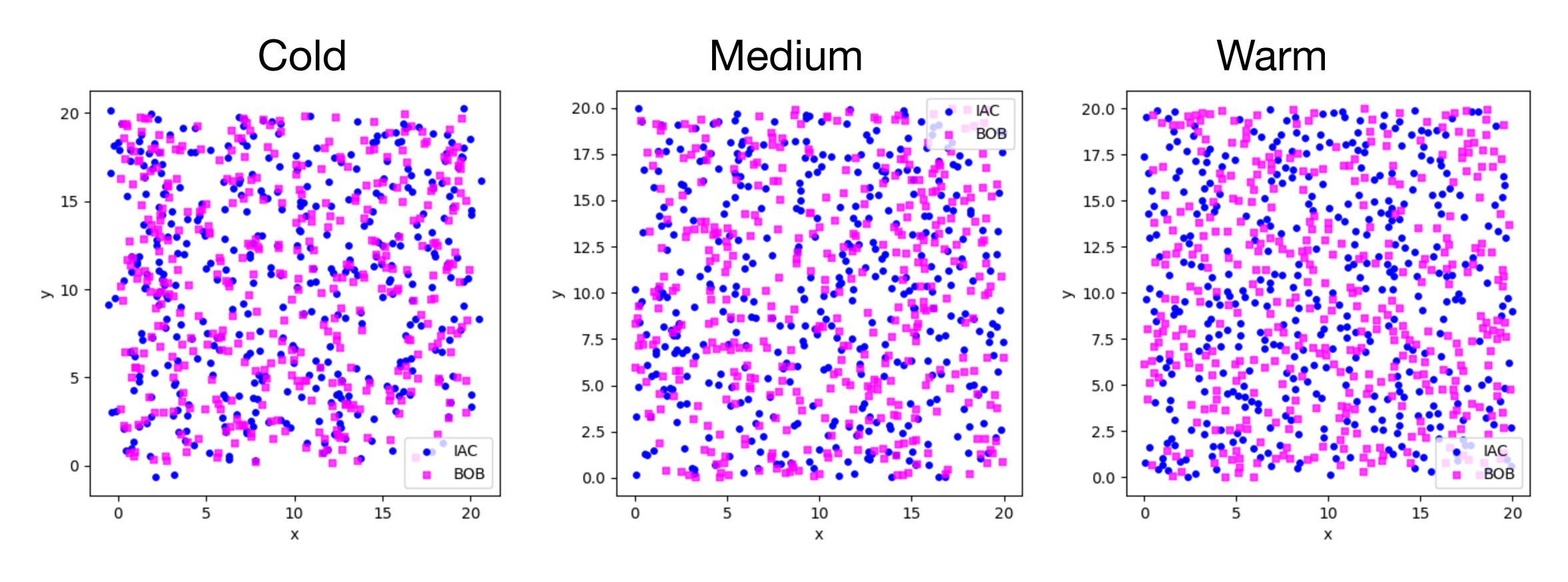








The data — Fall epidermis samples



Do IAC and BOB attract or repulse each other depending on temperature? Is there an association between attraction and repulsion and temperature?







The data — Winter epidermis samples

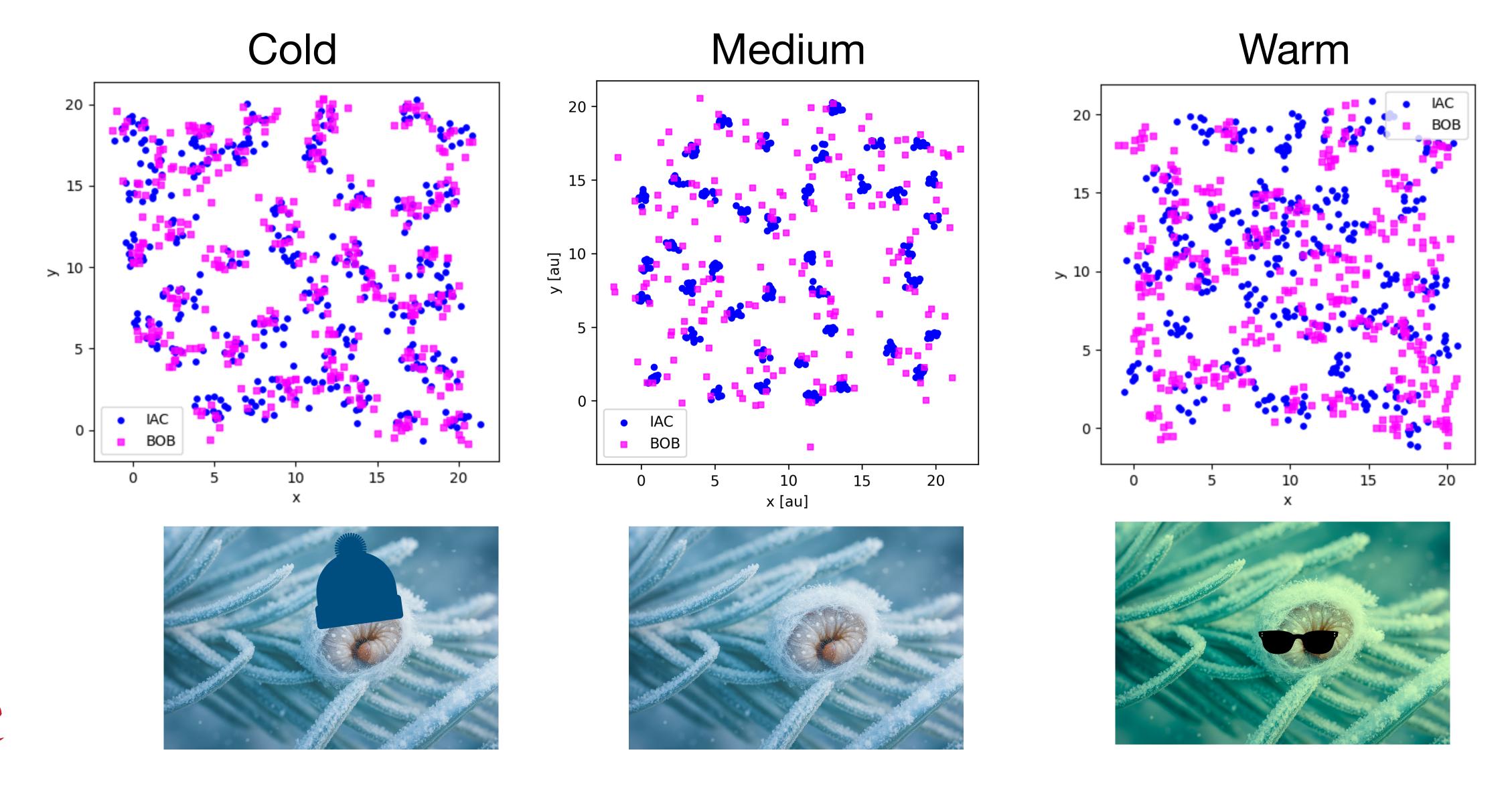








The data — Winter epidermis samples

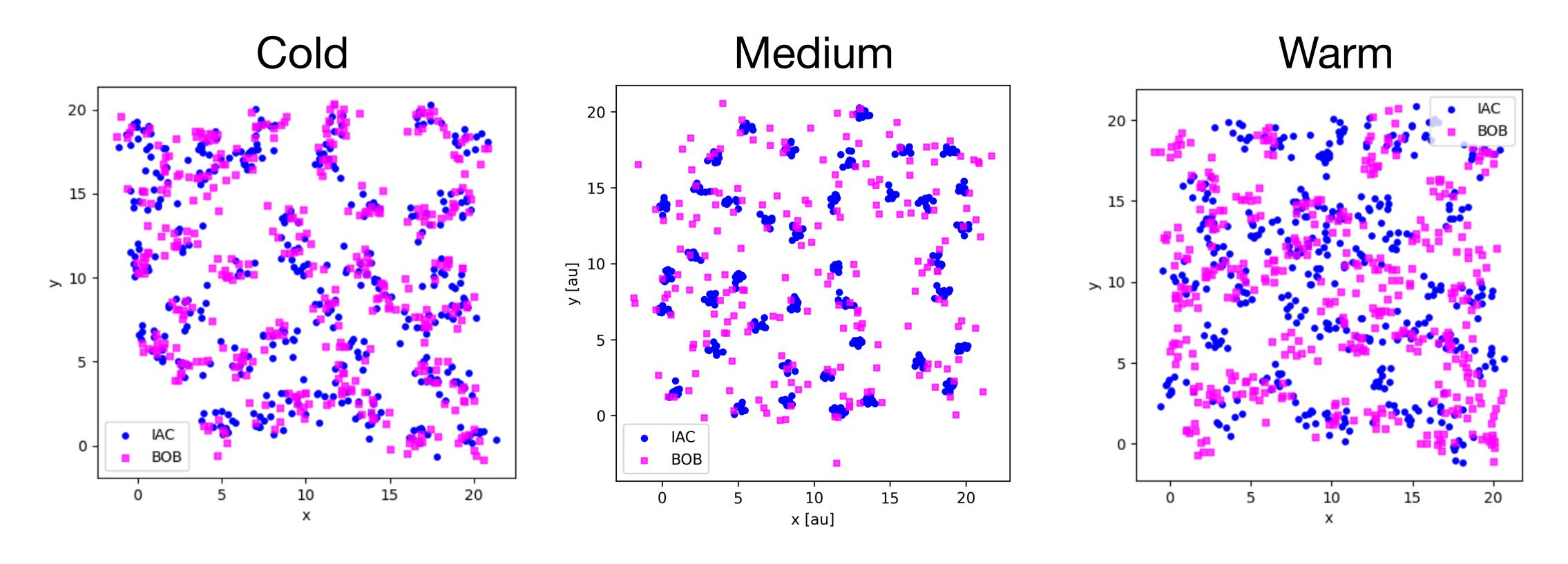








The data — Winter epidermis samples



Do IAC and BOB attract or repulse each other depending on temperature? Is there an association between attraction and repulsion and temperature?













Q: Do you see patterns?

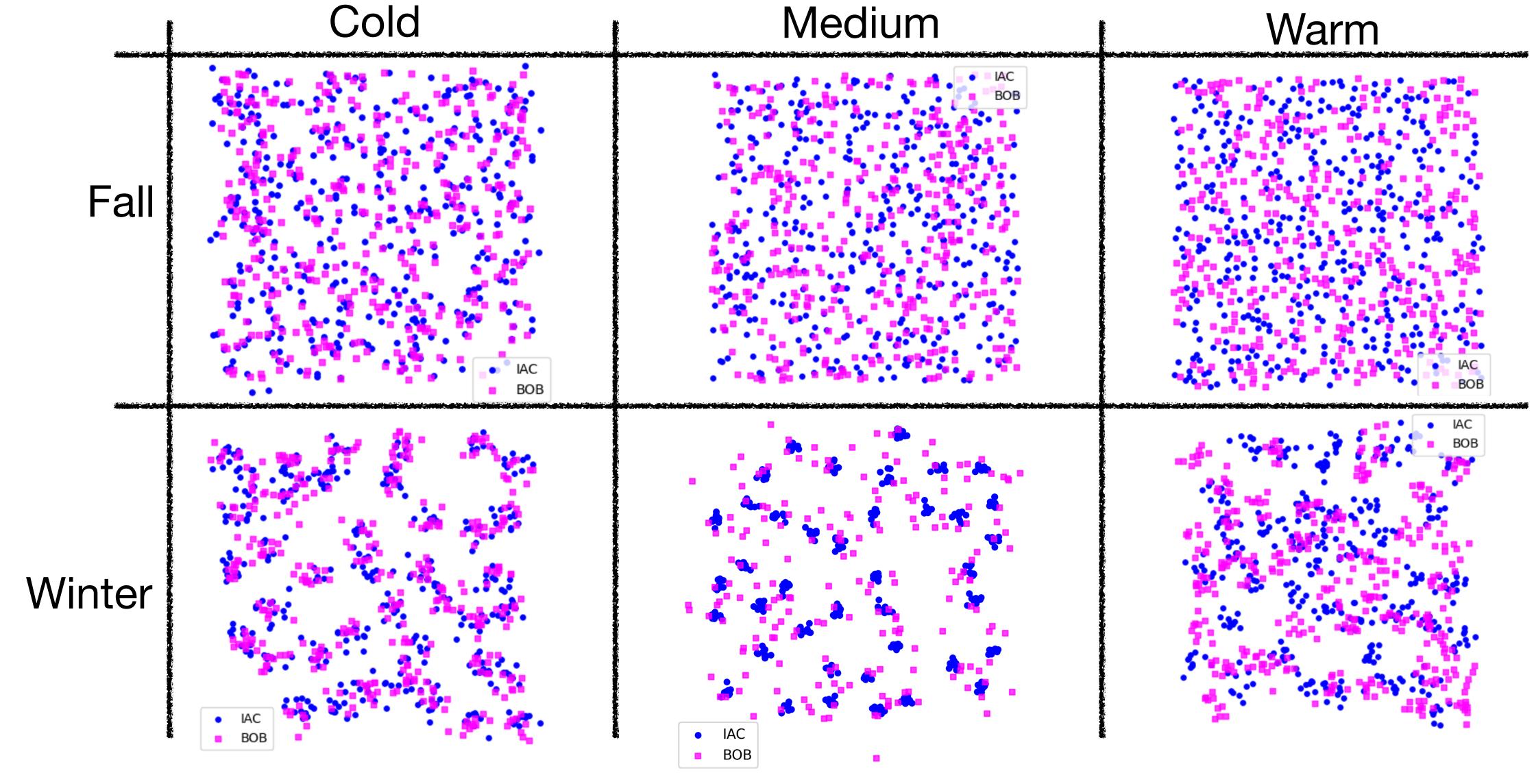








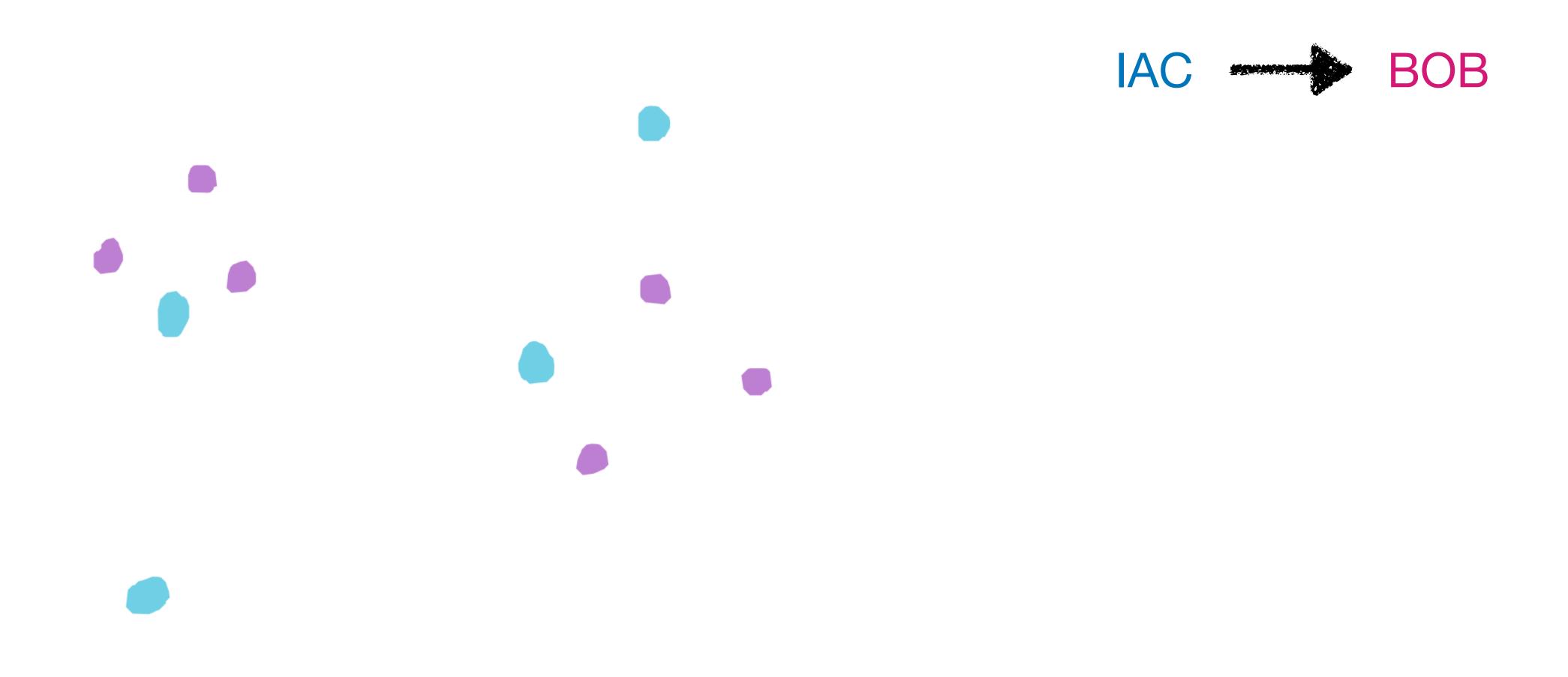
How would you analyze the data?







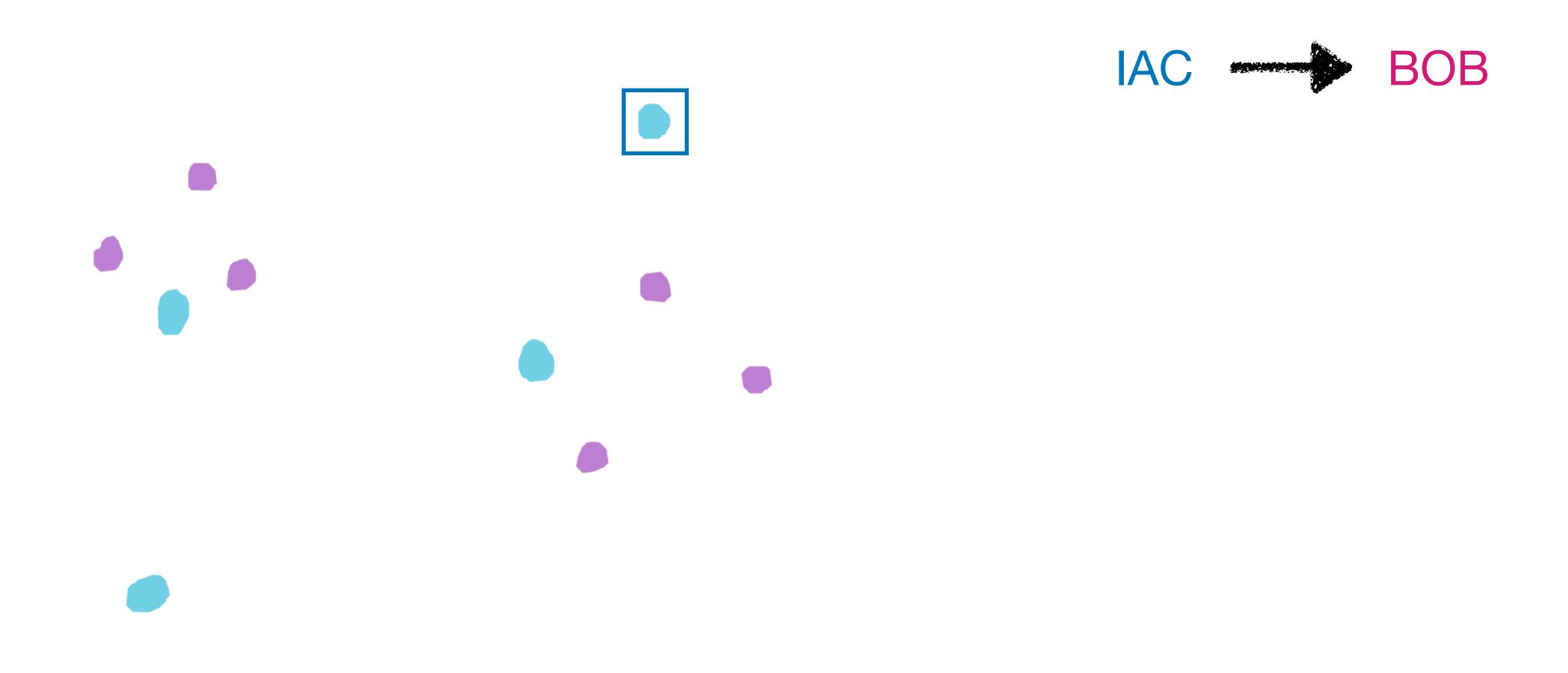








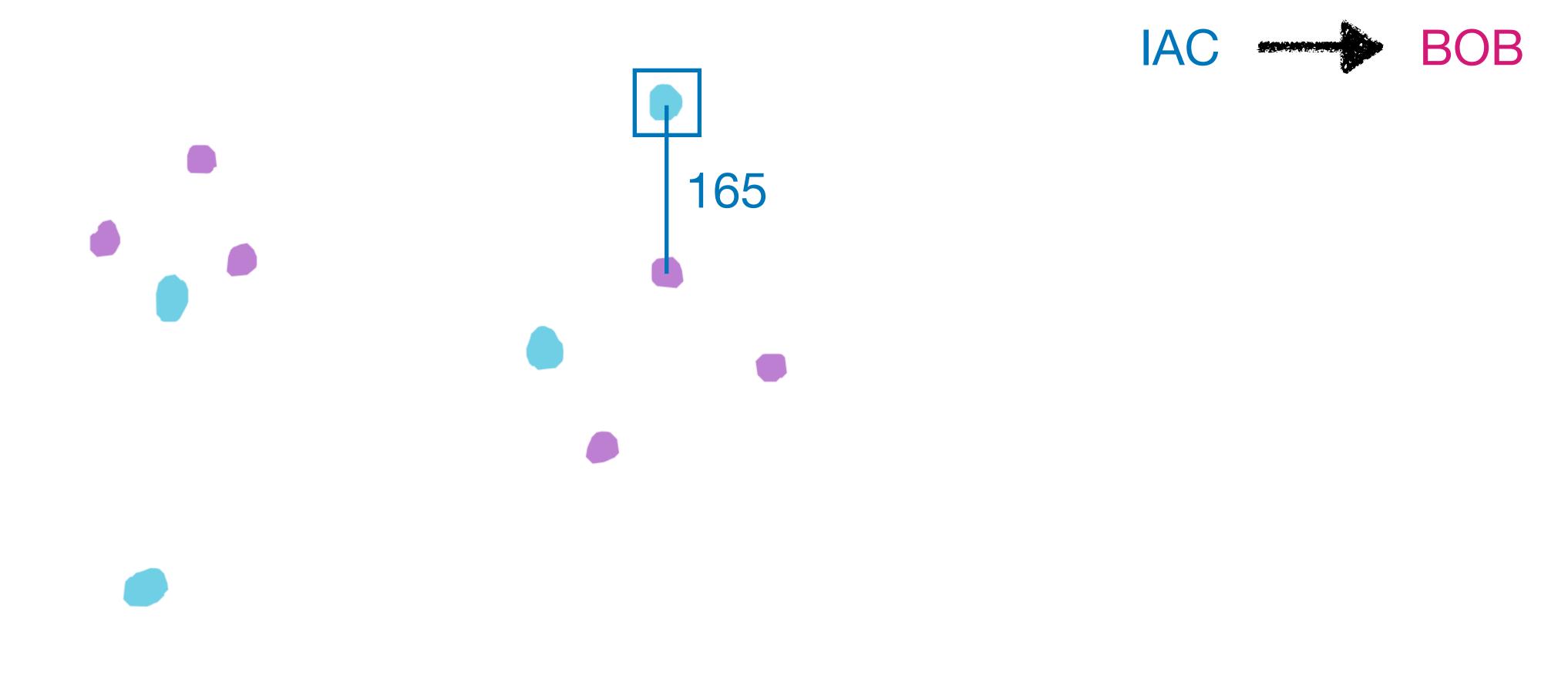








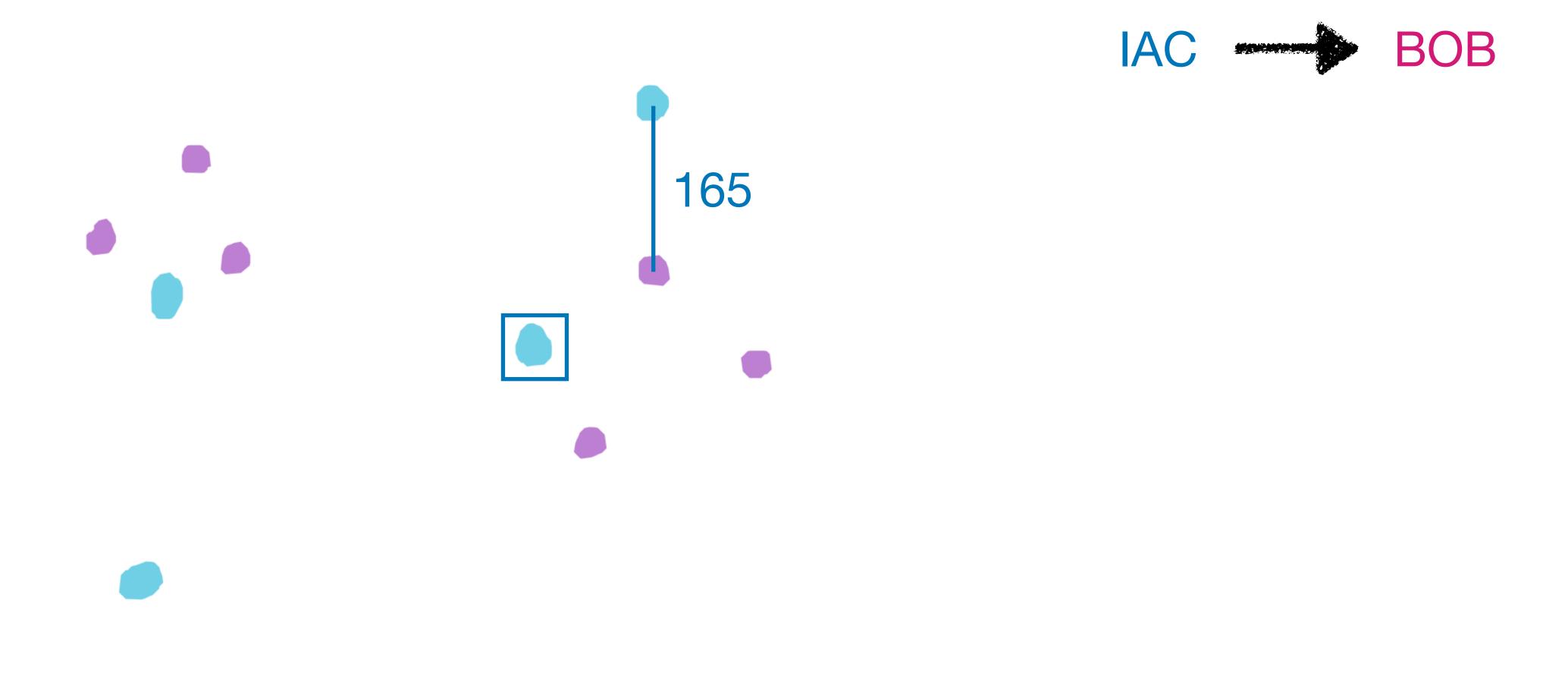








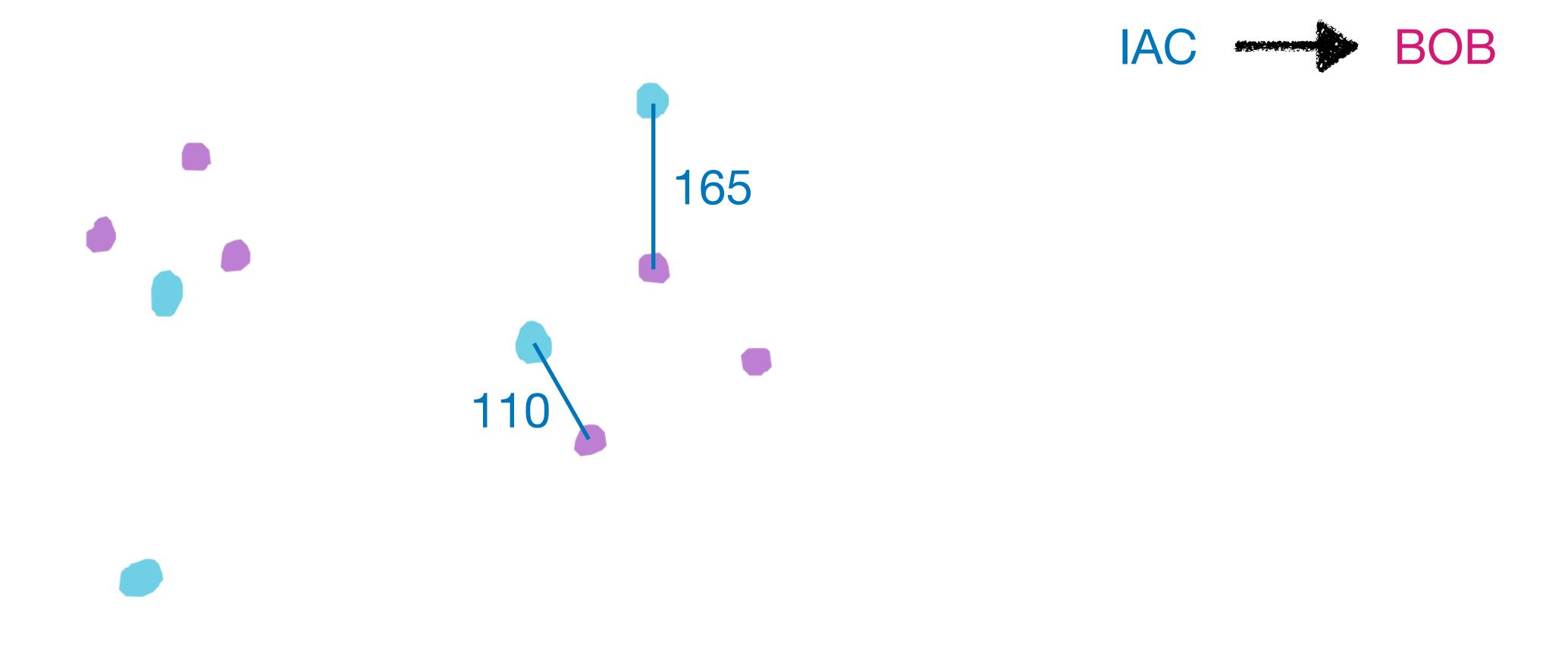








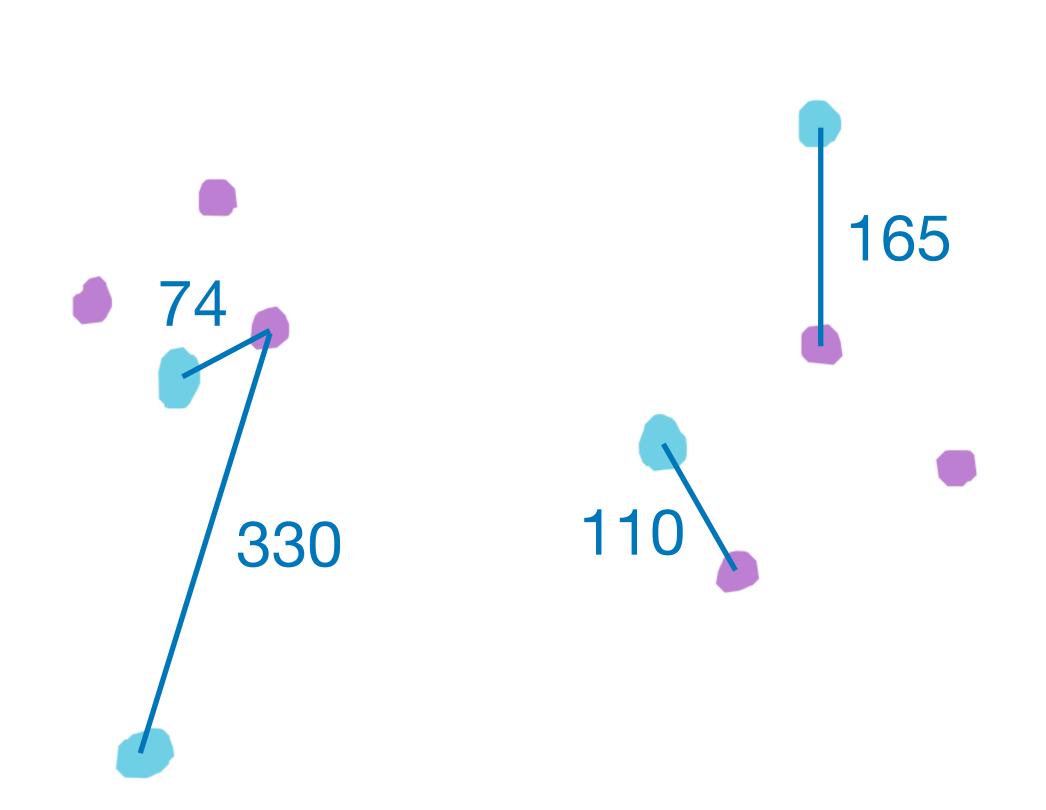


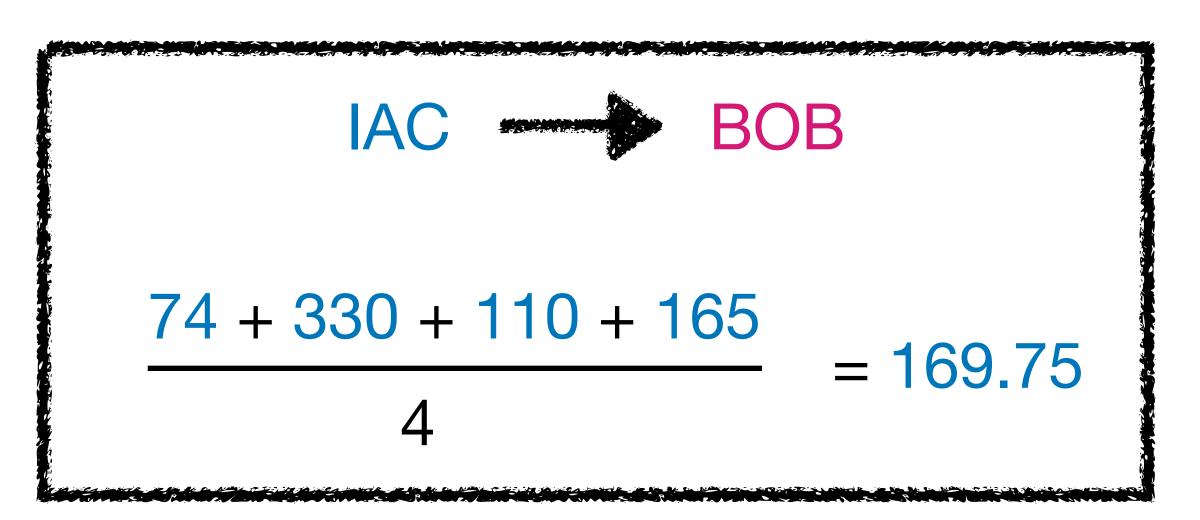








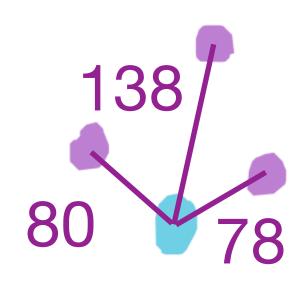


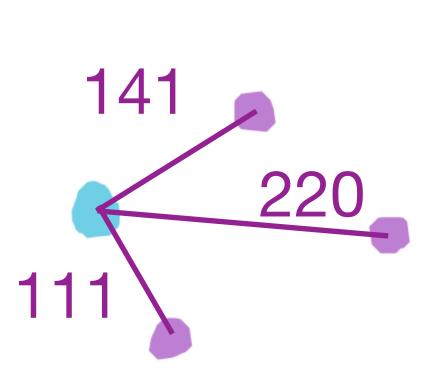














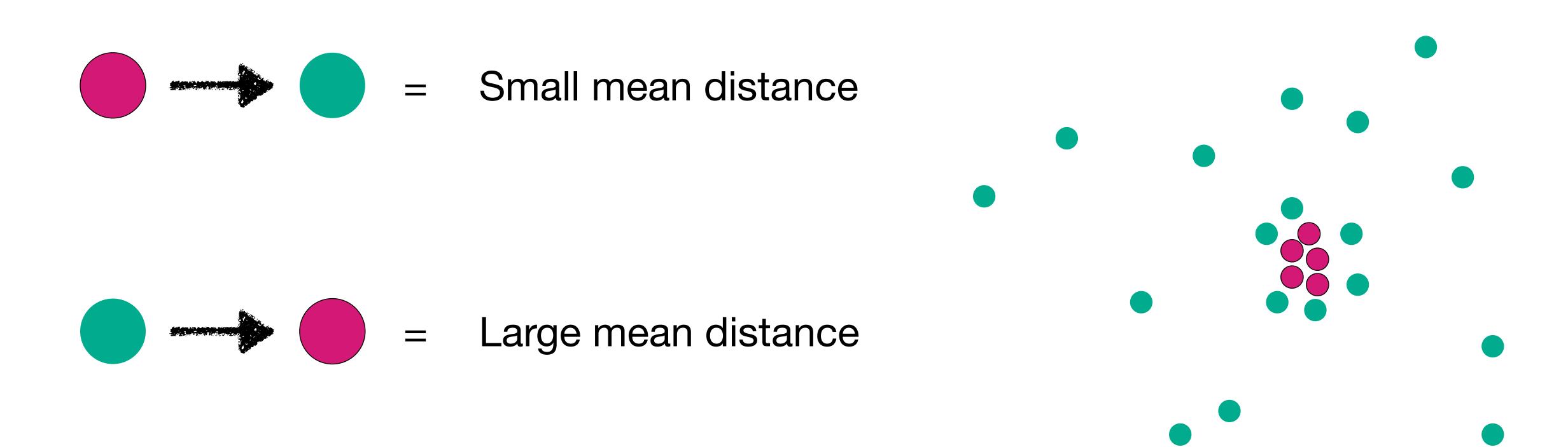
$$\frac{74 + 330 + 110 + 165}{4} = 169.75$$

BOB IAC
$$80+138+78+111+141+220 = 121$$













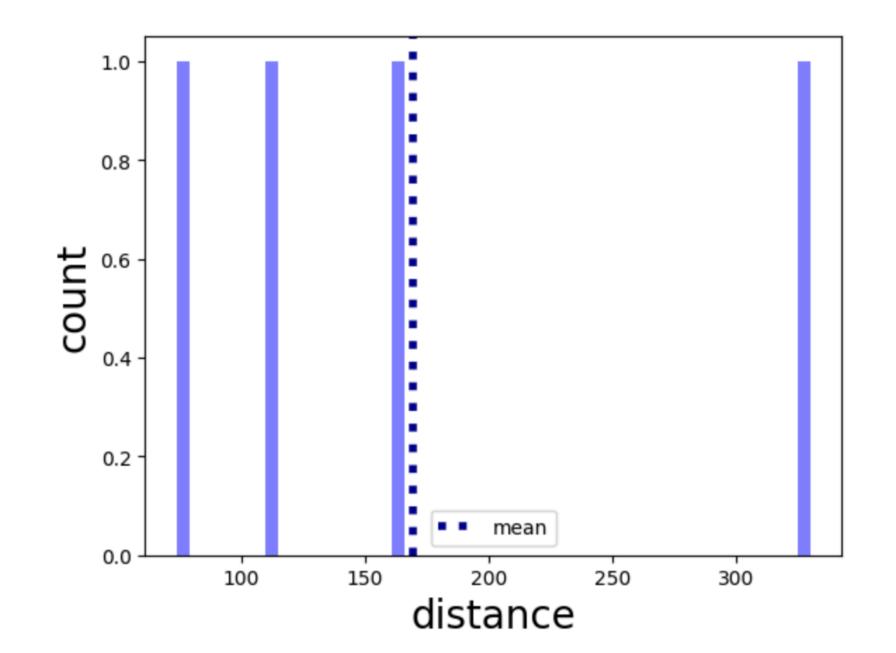


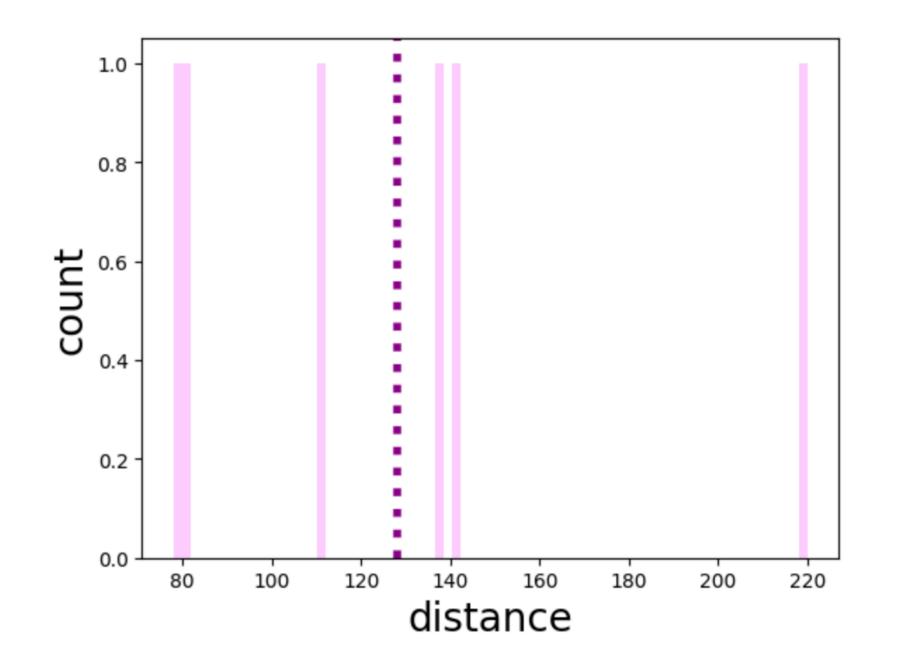




Distances	Mean	
74, 330, 110, 165	169.75	

Distances					Mean	
80,	138,	78,	111,	141,	220	121



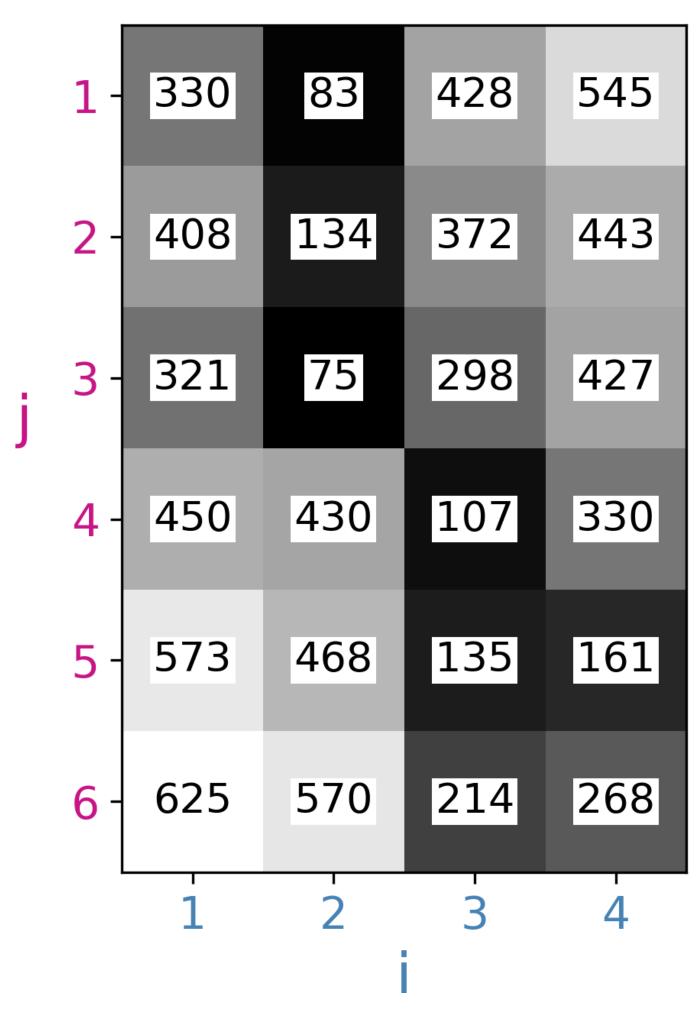


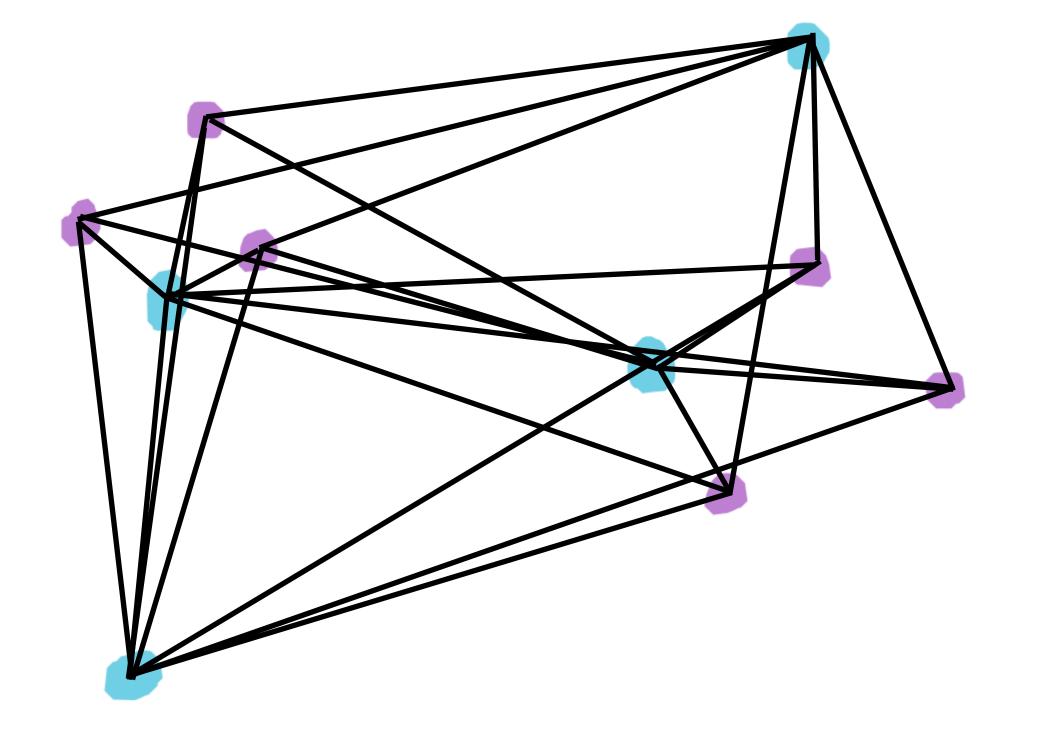






dist_matrix



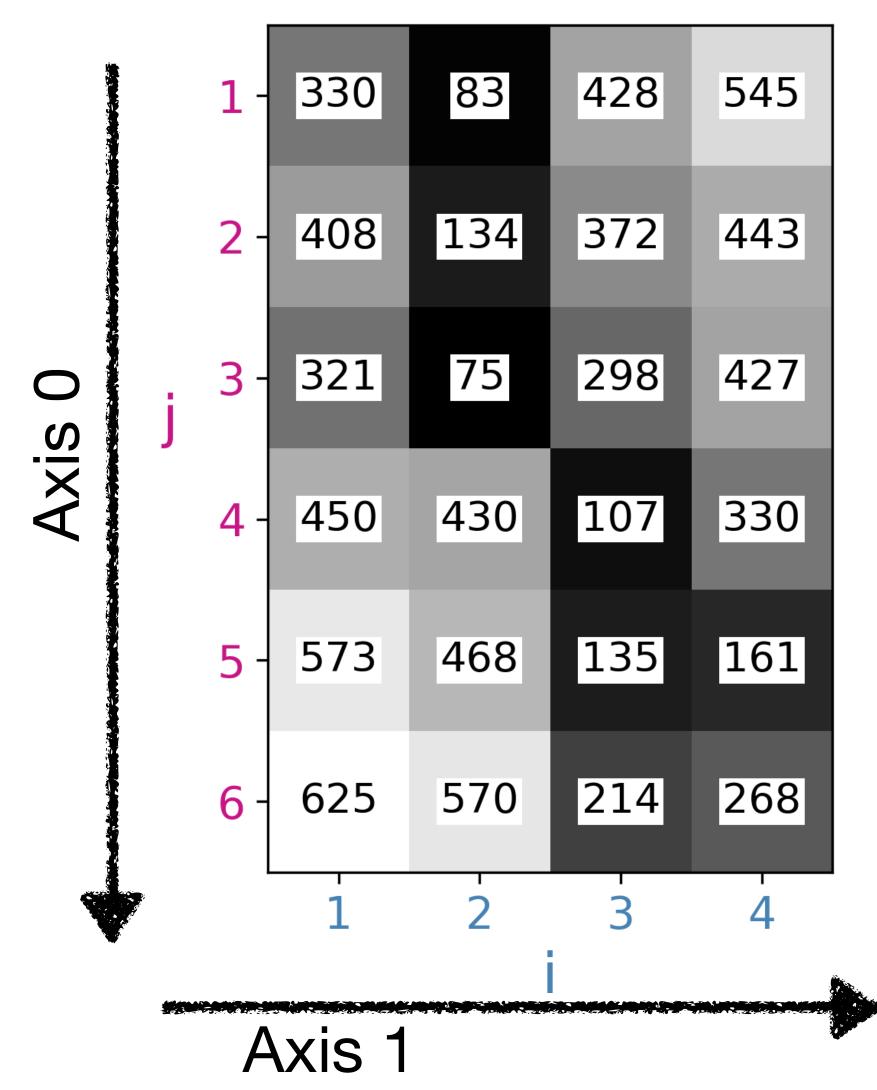




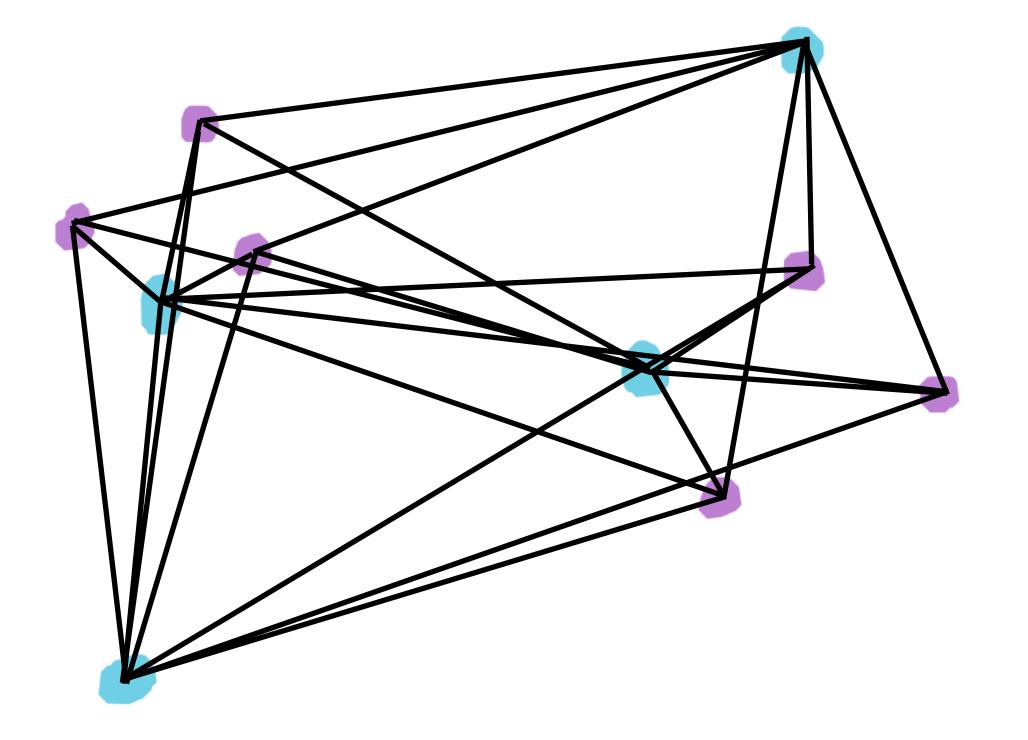




dist_matrix



np.min(dist_matrix, axis = 1)

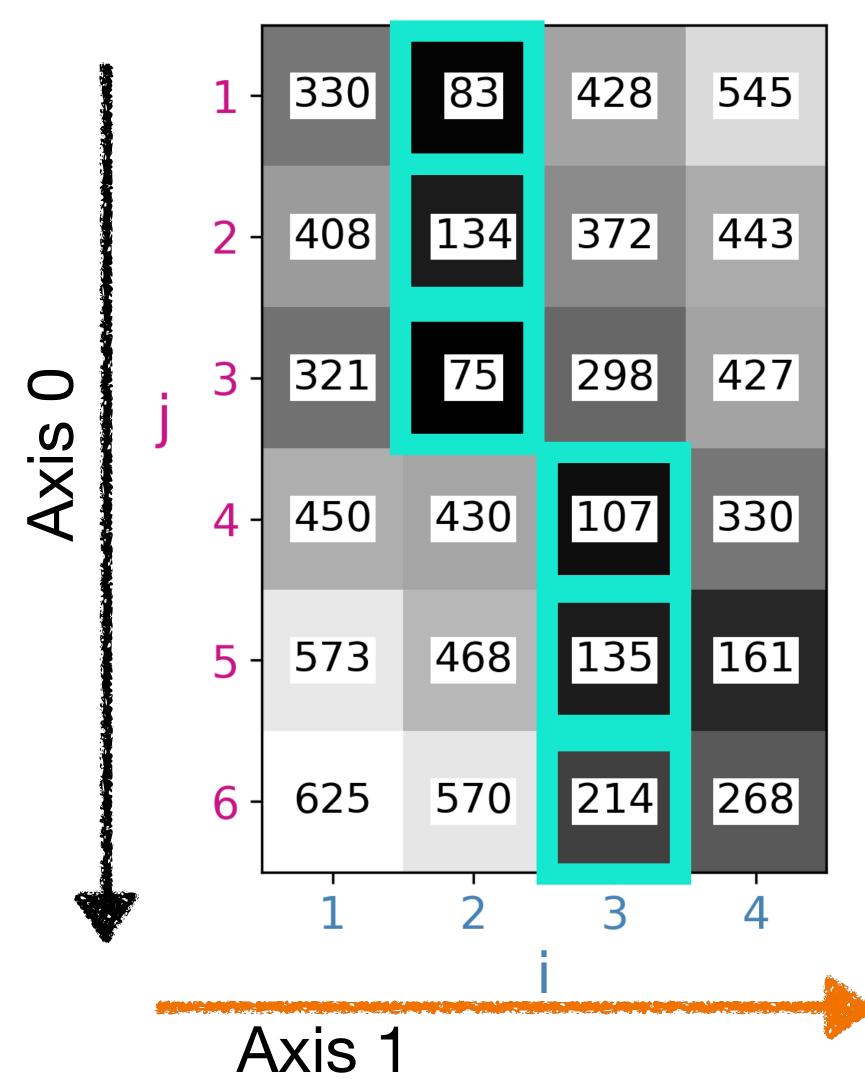




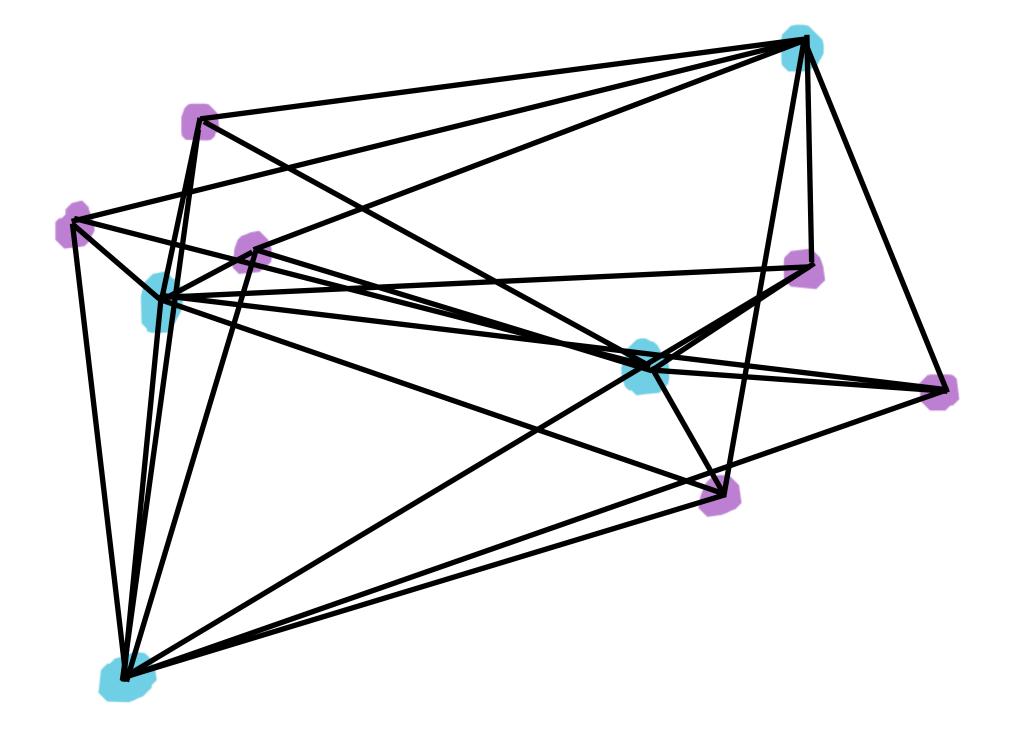




dist_matrix



np.min(dist_matrix, axis = 1)

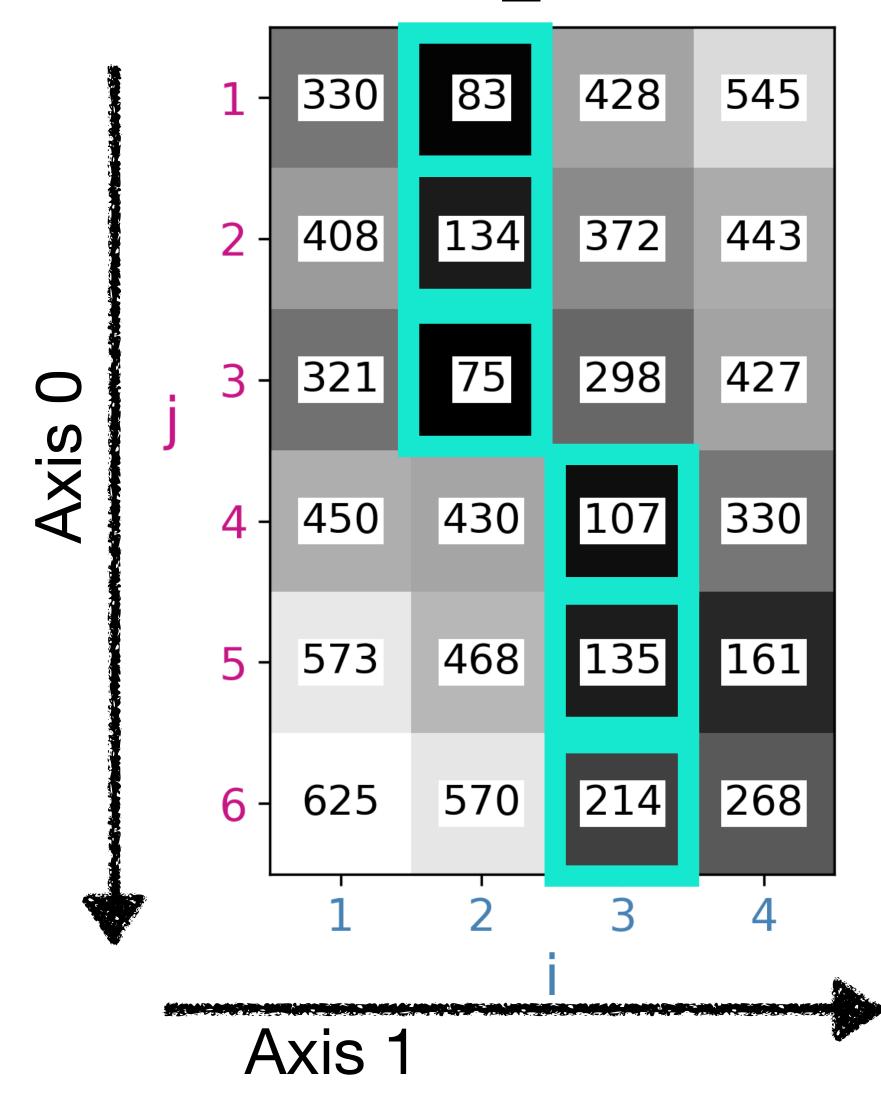




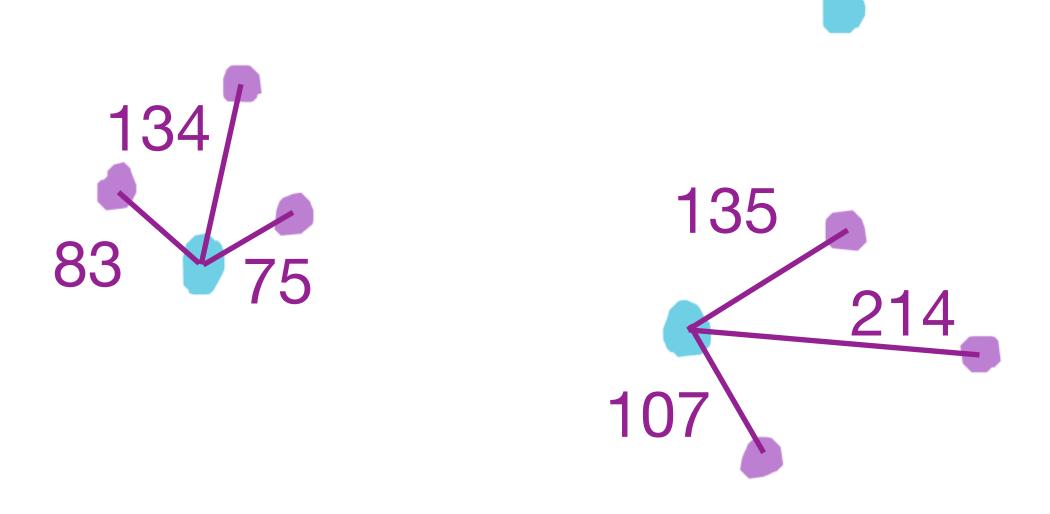




dist_matrix



np.min(dist_matrix, axis = 1)

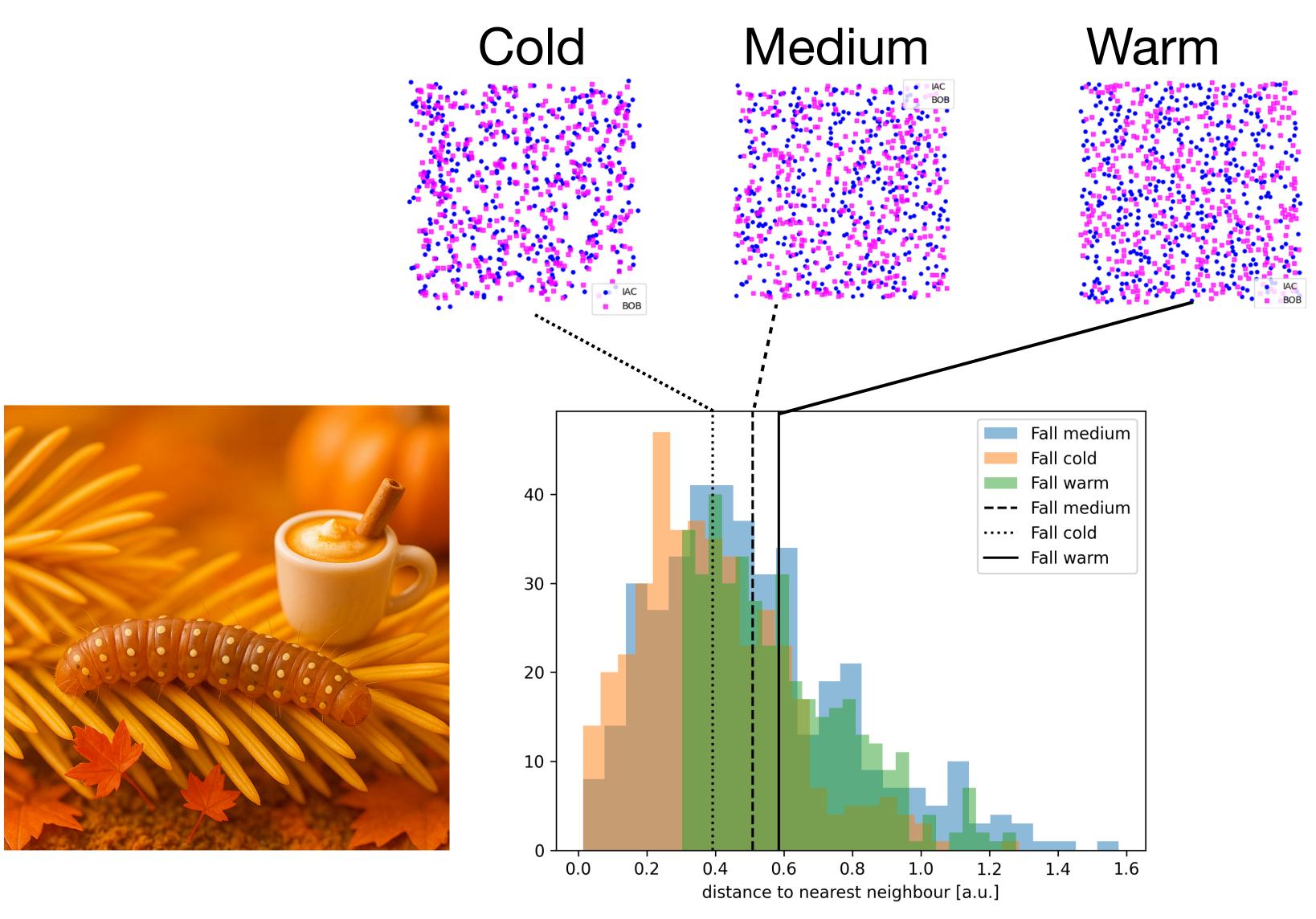








Results: Mean distance | AC -> BOB

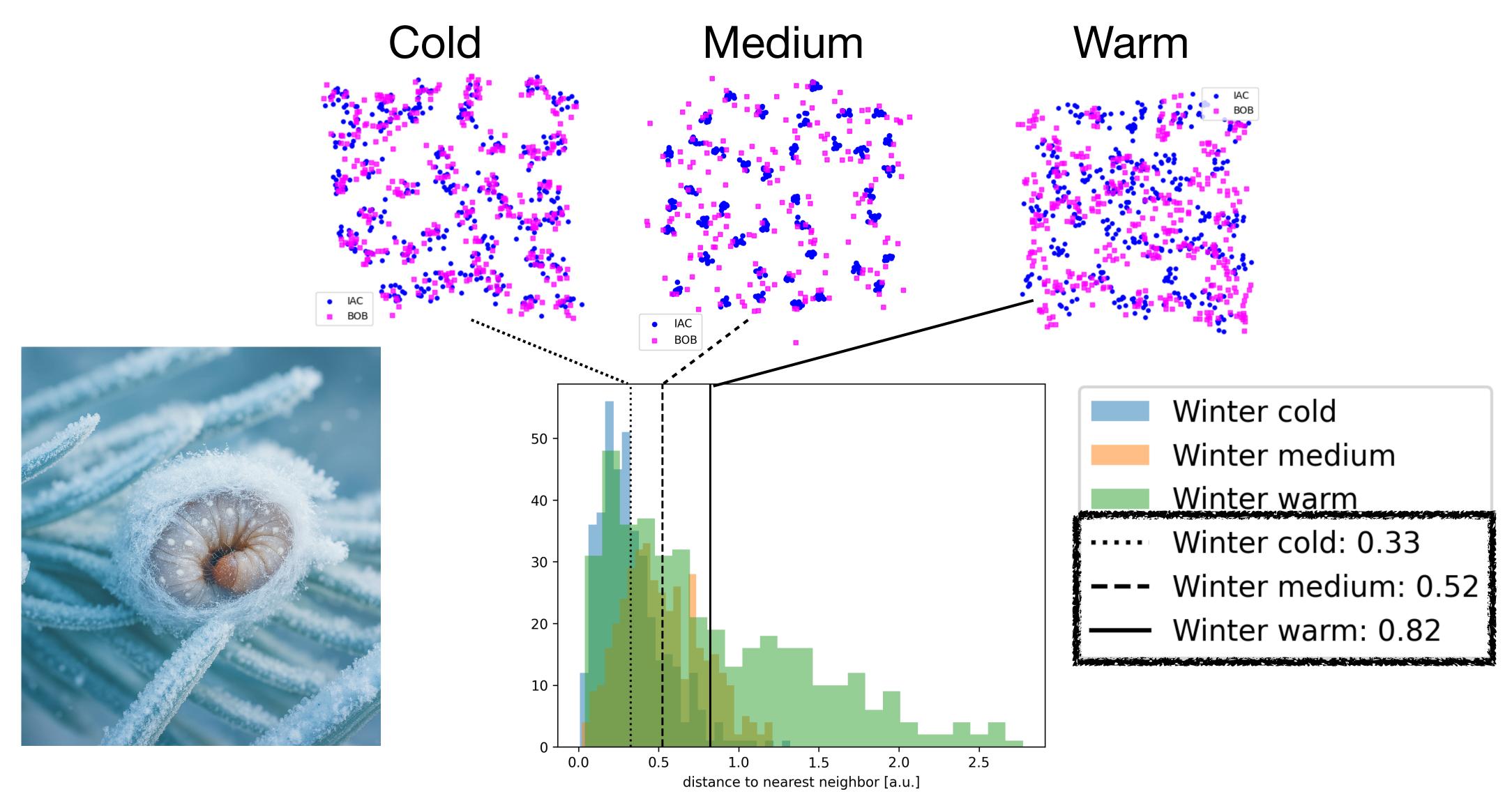








Results: Mean distance | AC -> BOB

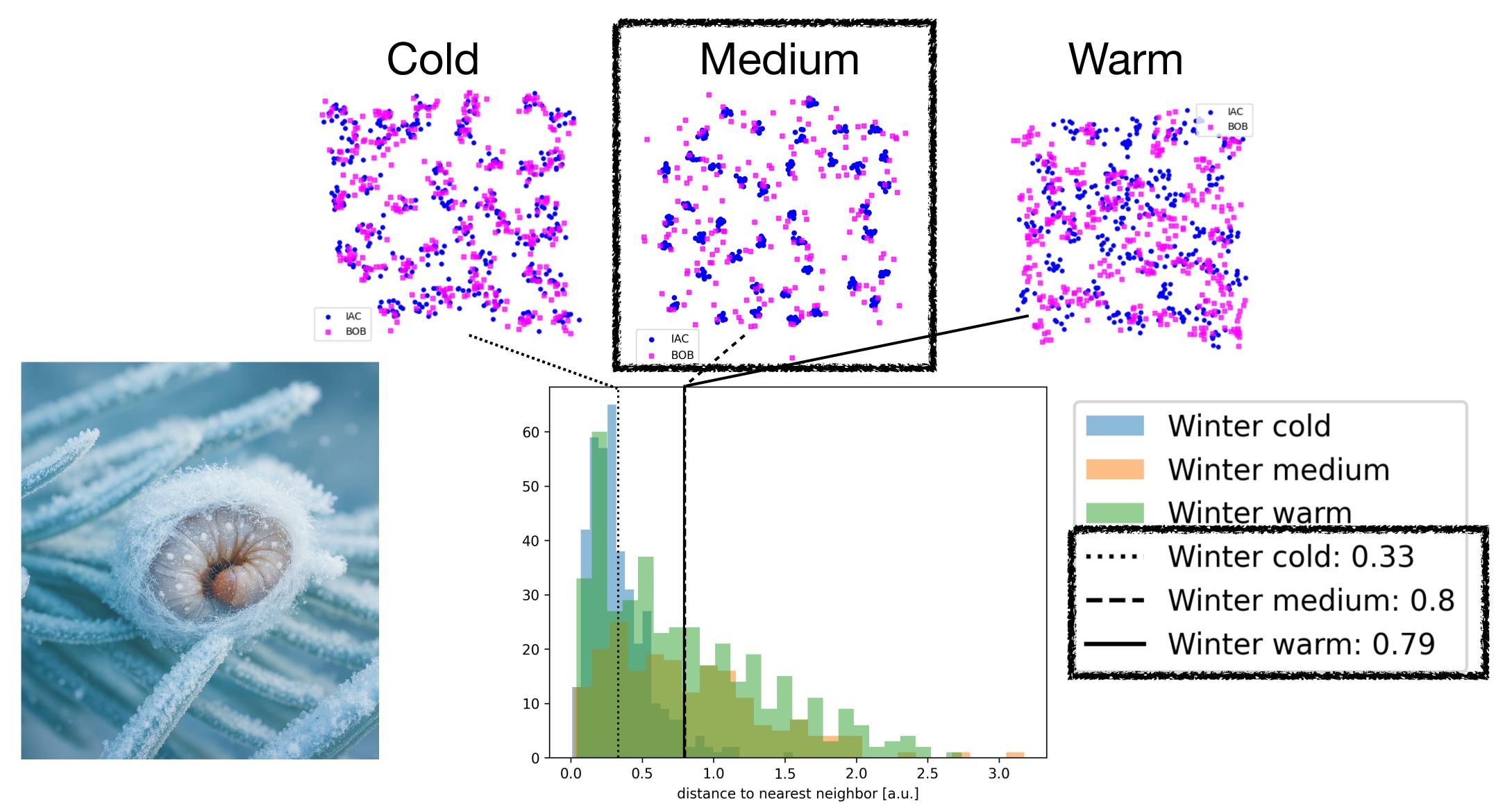








Results: Mean distance BOB -> IAC















- Asymmetric: BOB → IAC ≠ IAC → BOB
- Returns: One number
- Range: Short

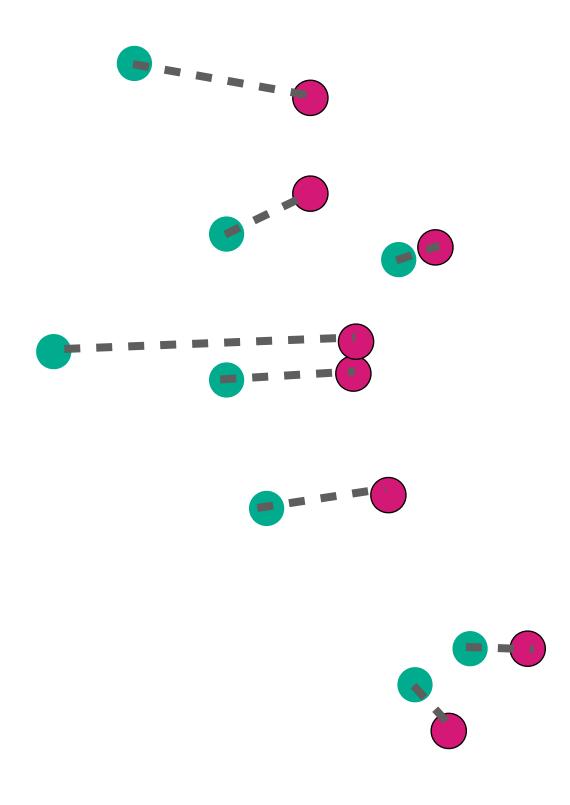






Beyond the mean distance to nearest neighbor

Similar concepts hold true beyond the realm of just points



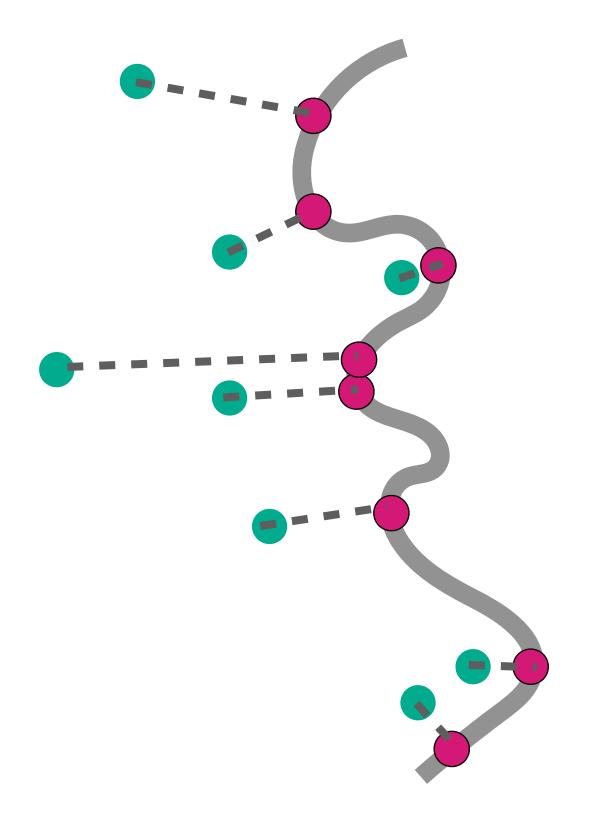






Beyond the mean distance to nearest neighbor

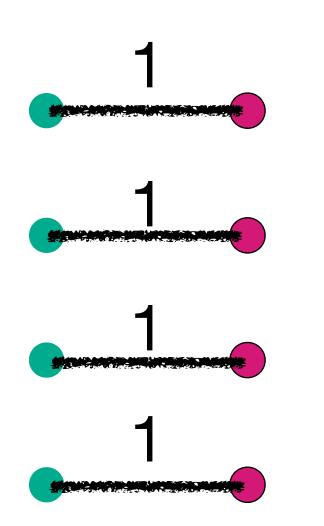
Similar concepts hold true beyond the realm of just points

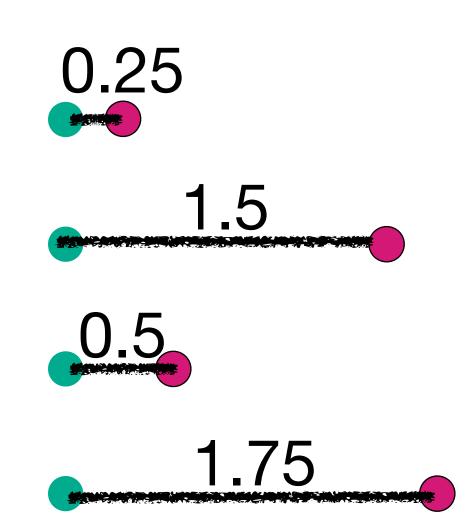












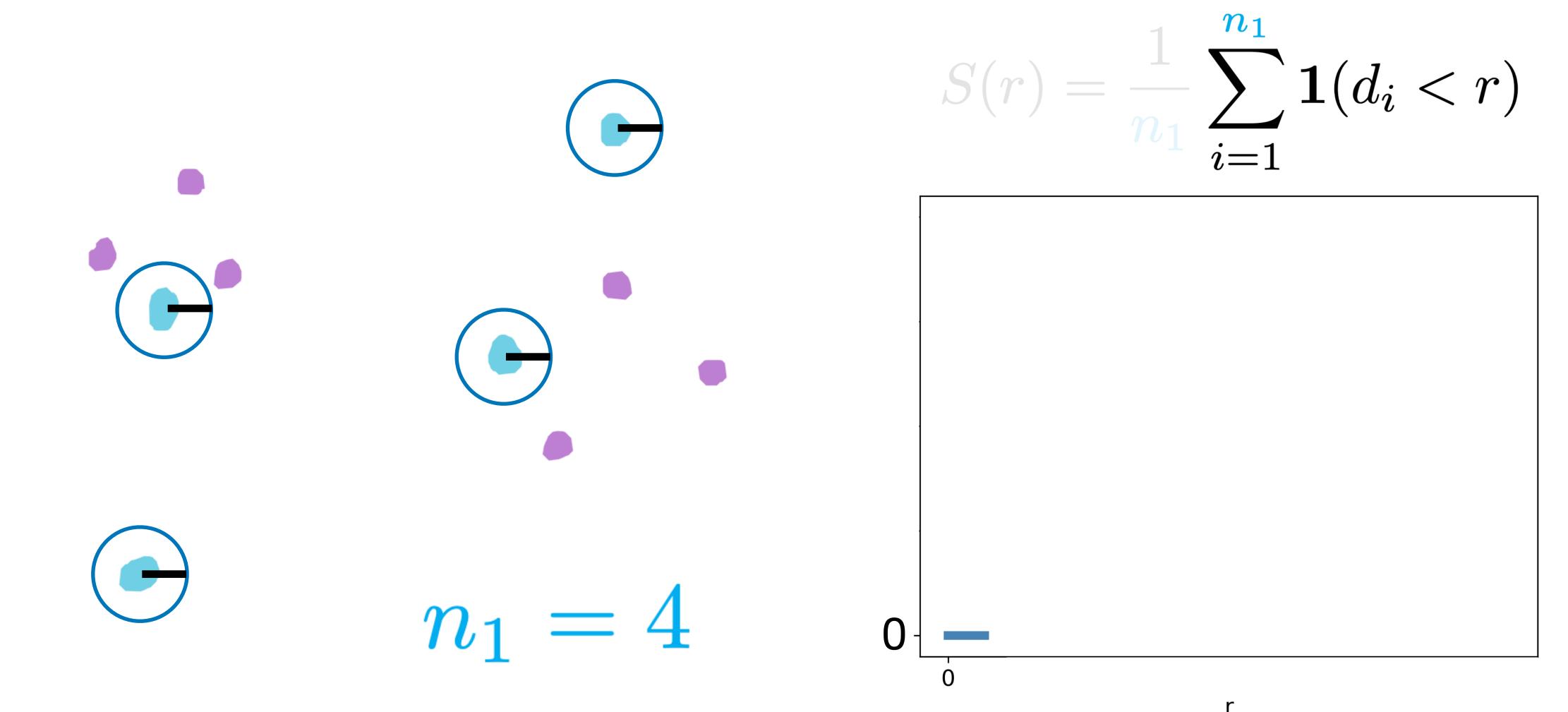
Mean dist: 1

Mean dist: 1





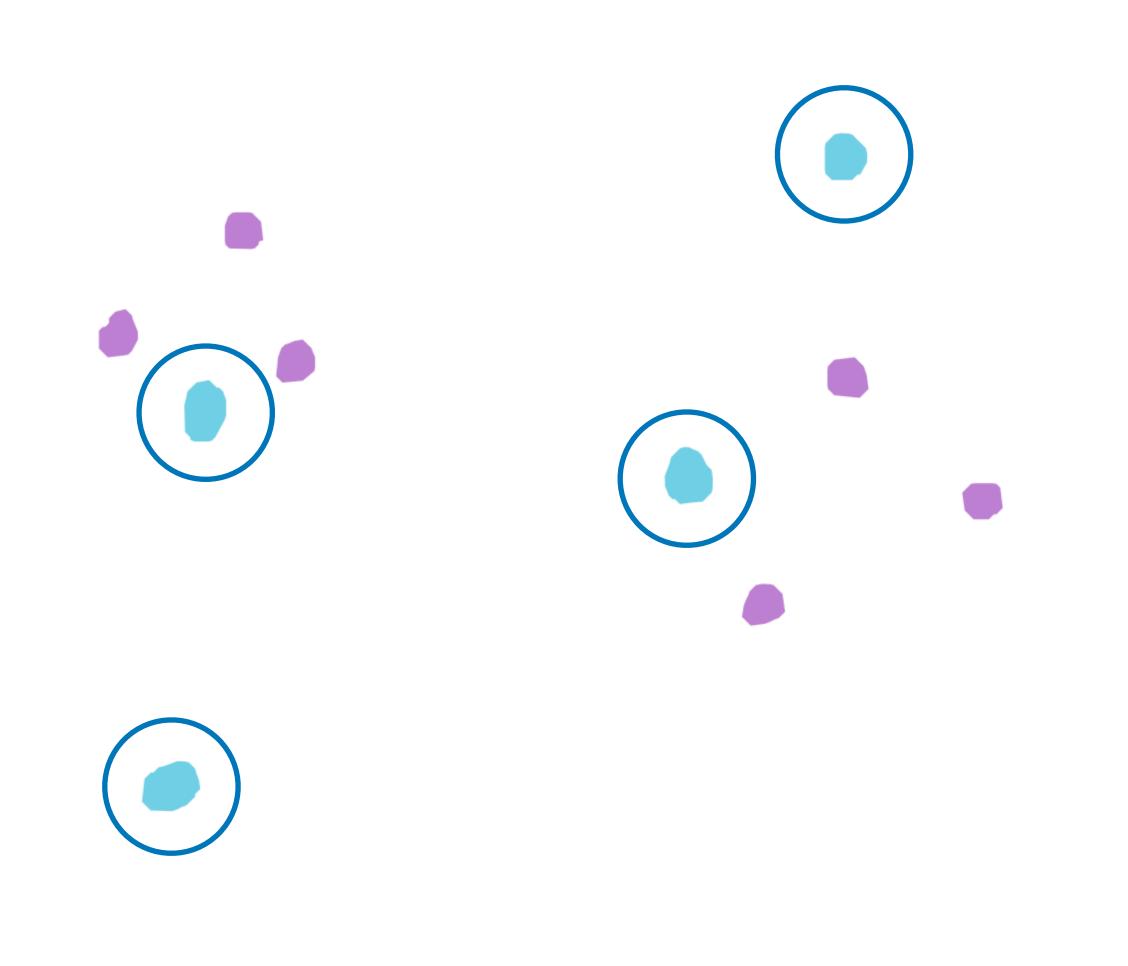




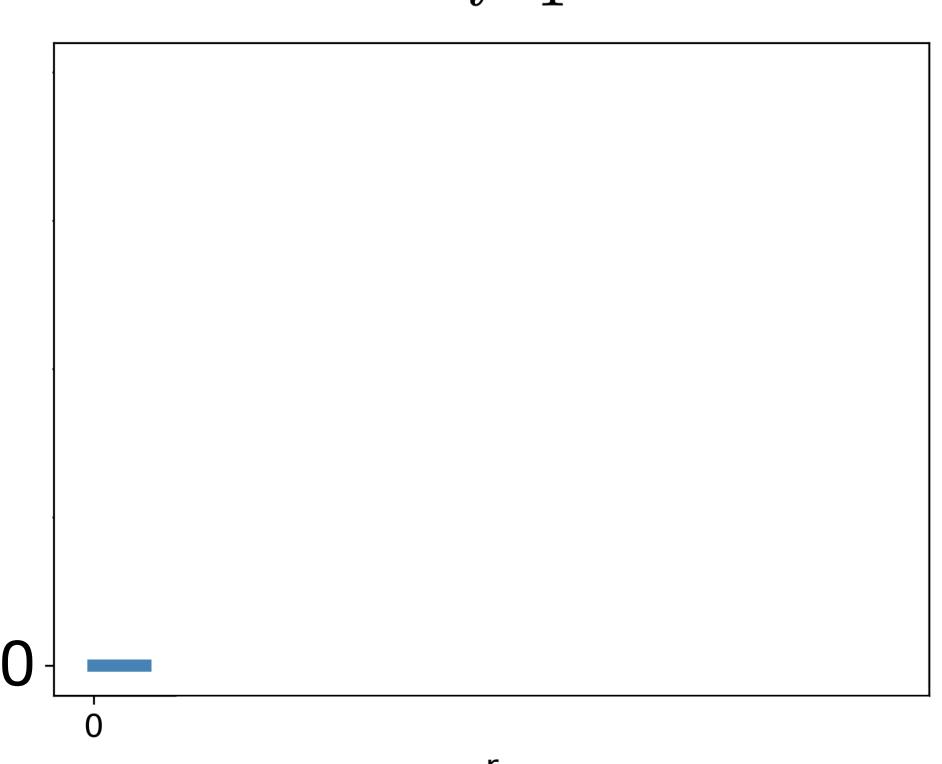








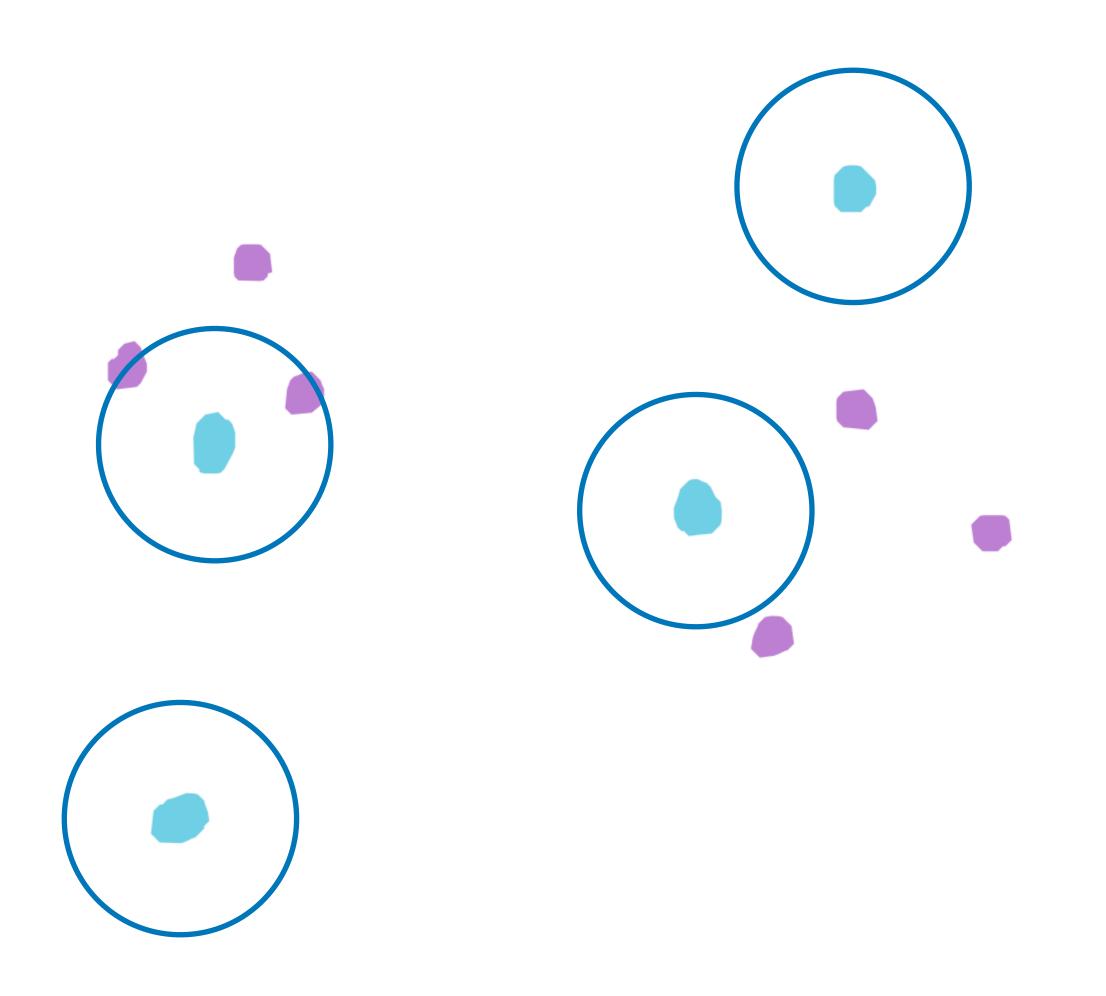
$$S(r) = \frac{1}{n_1} \sum_{i=1}^{n_1} \mathbf{1}(d_i < r)$$



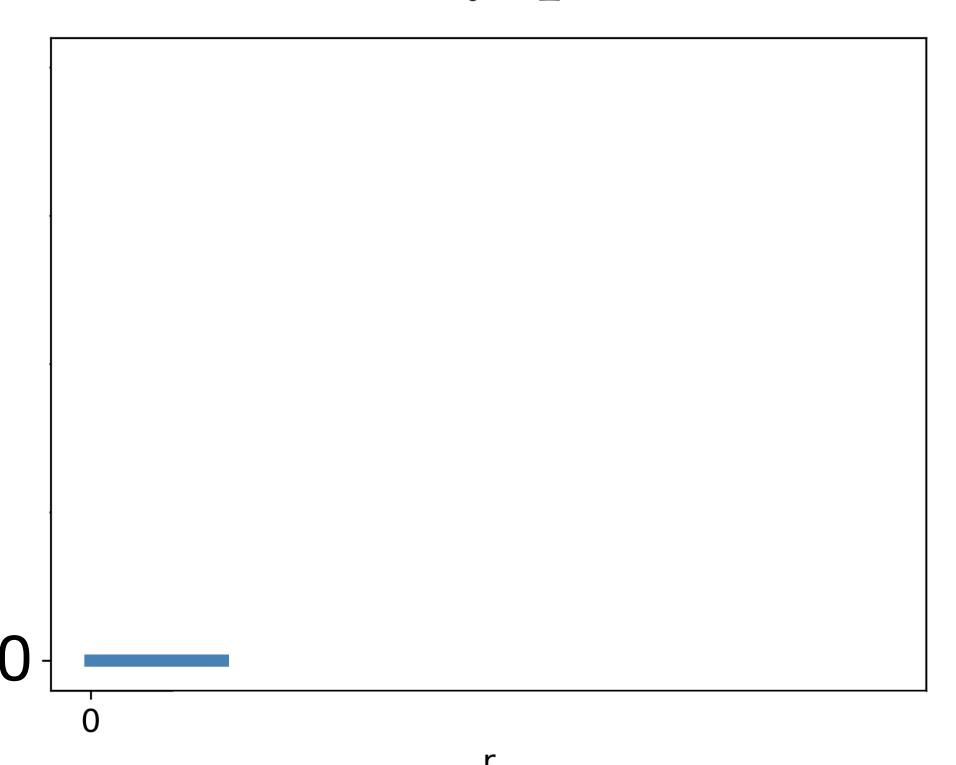








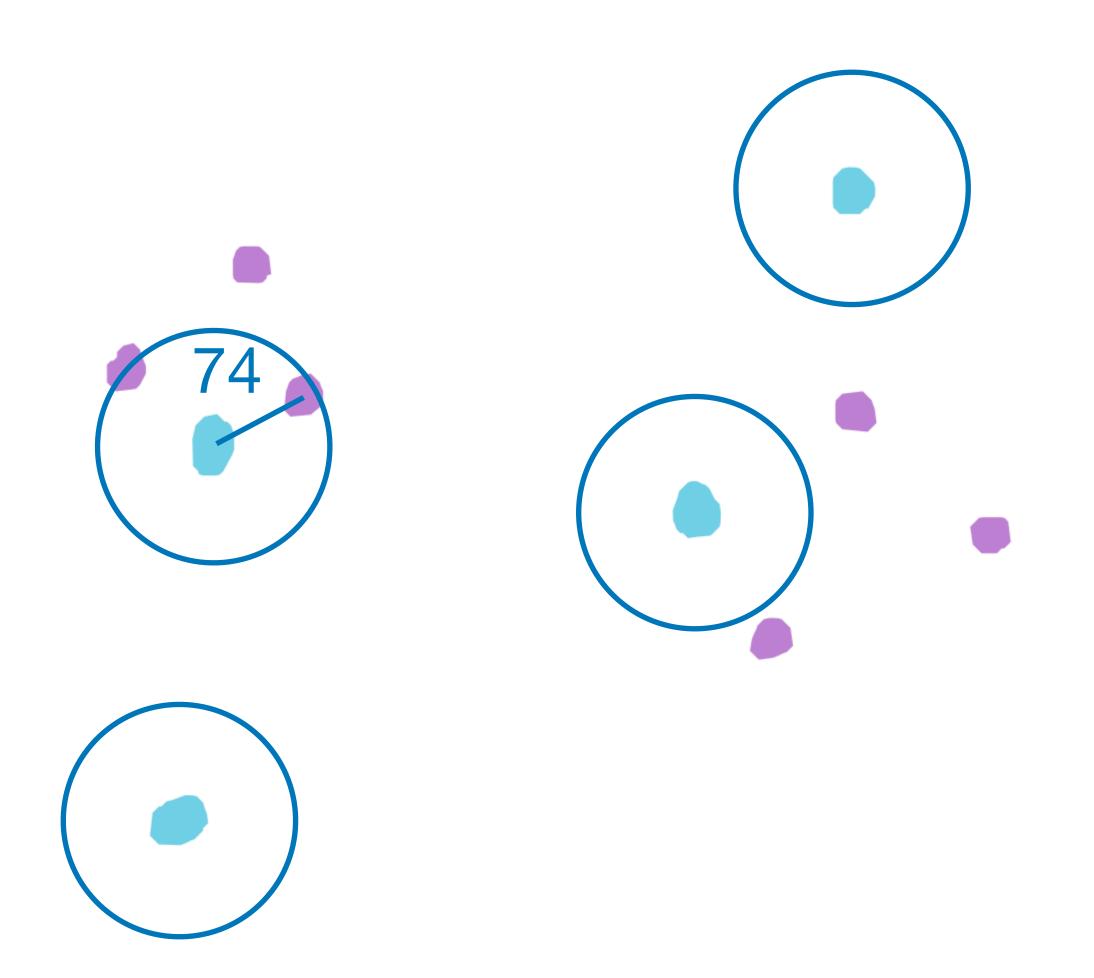
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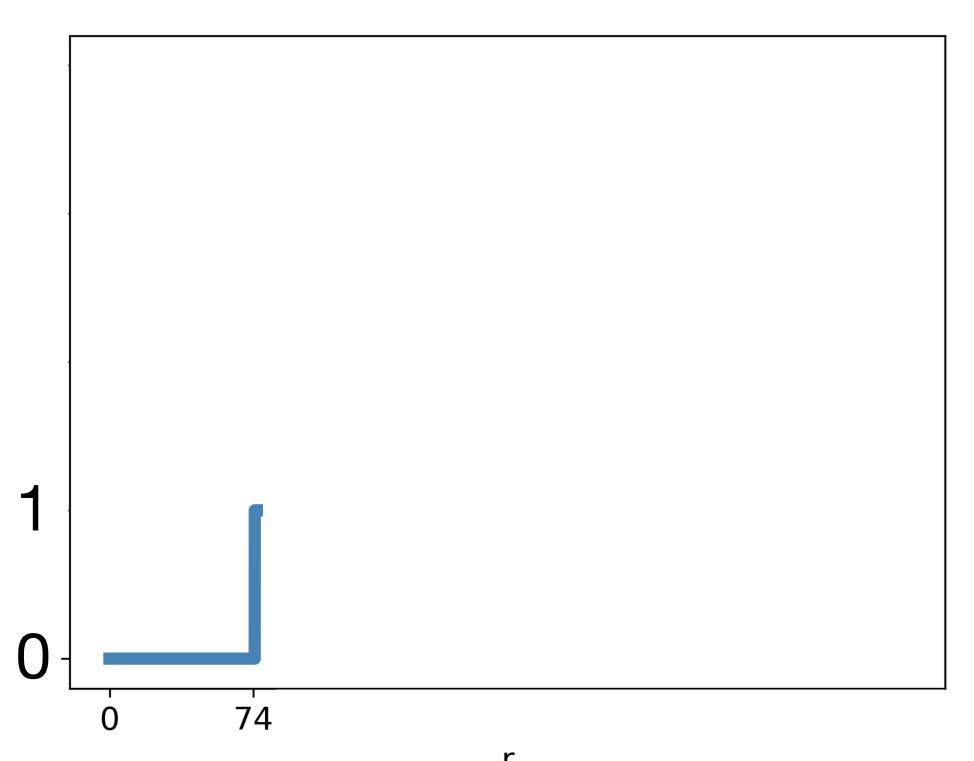








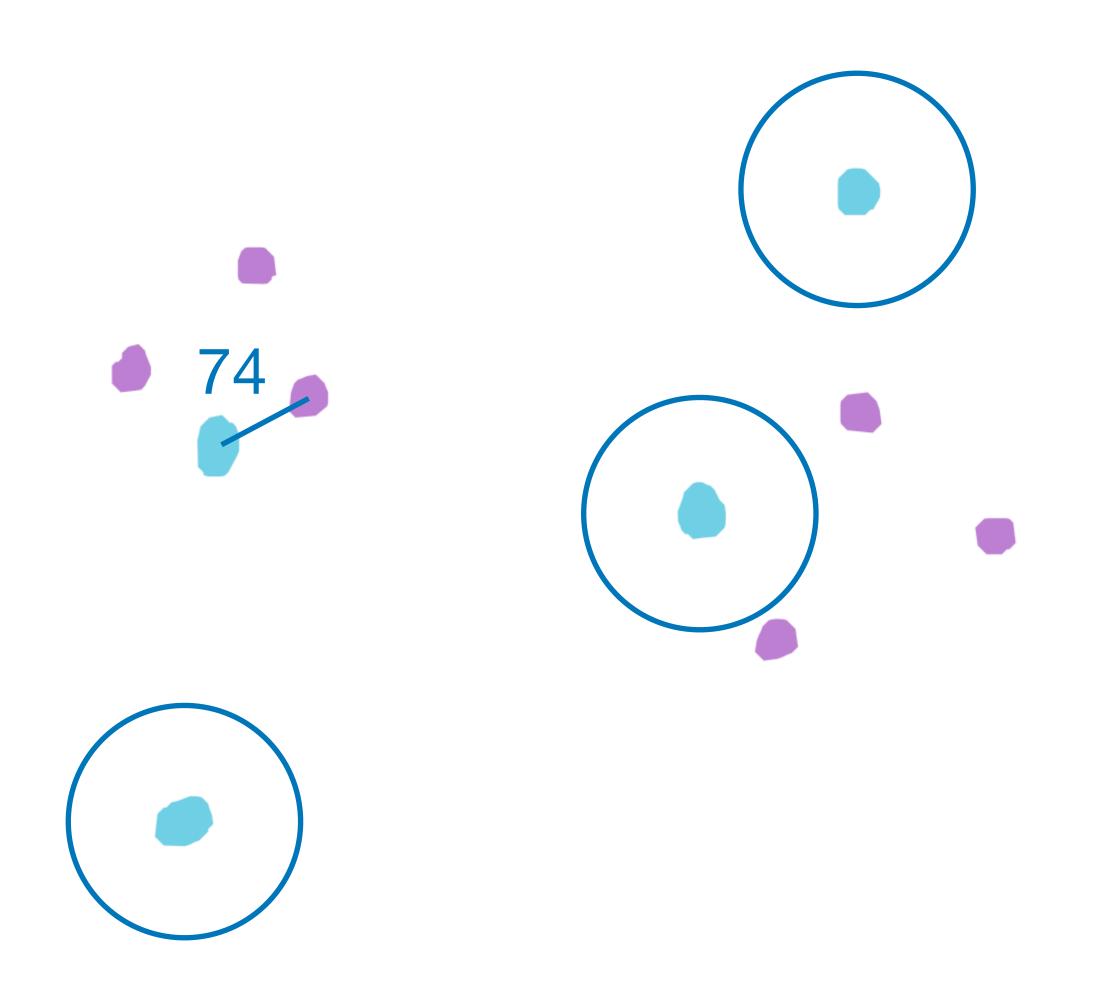
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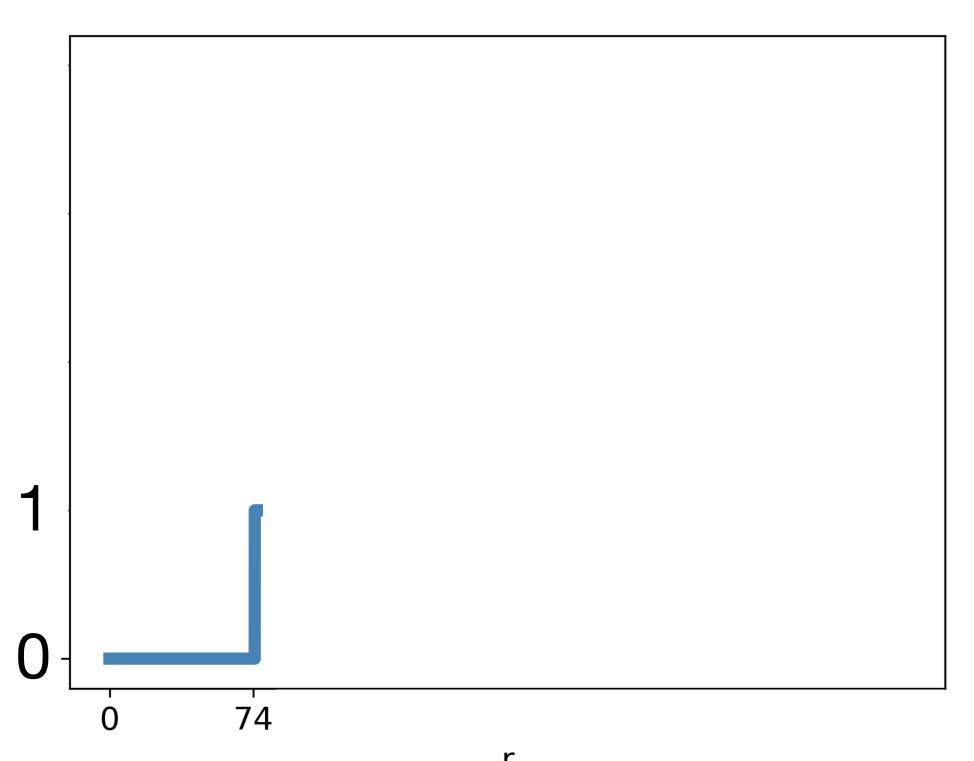








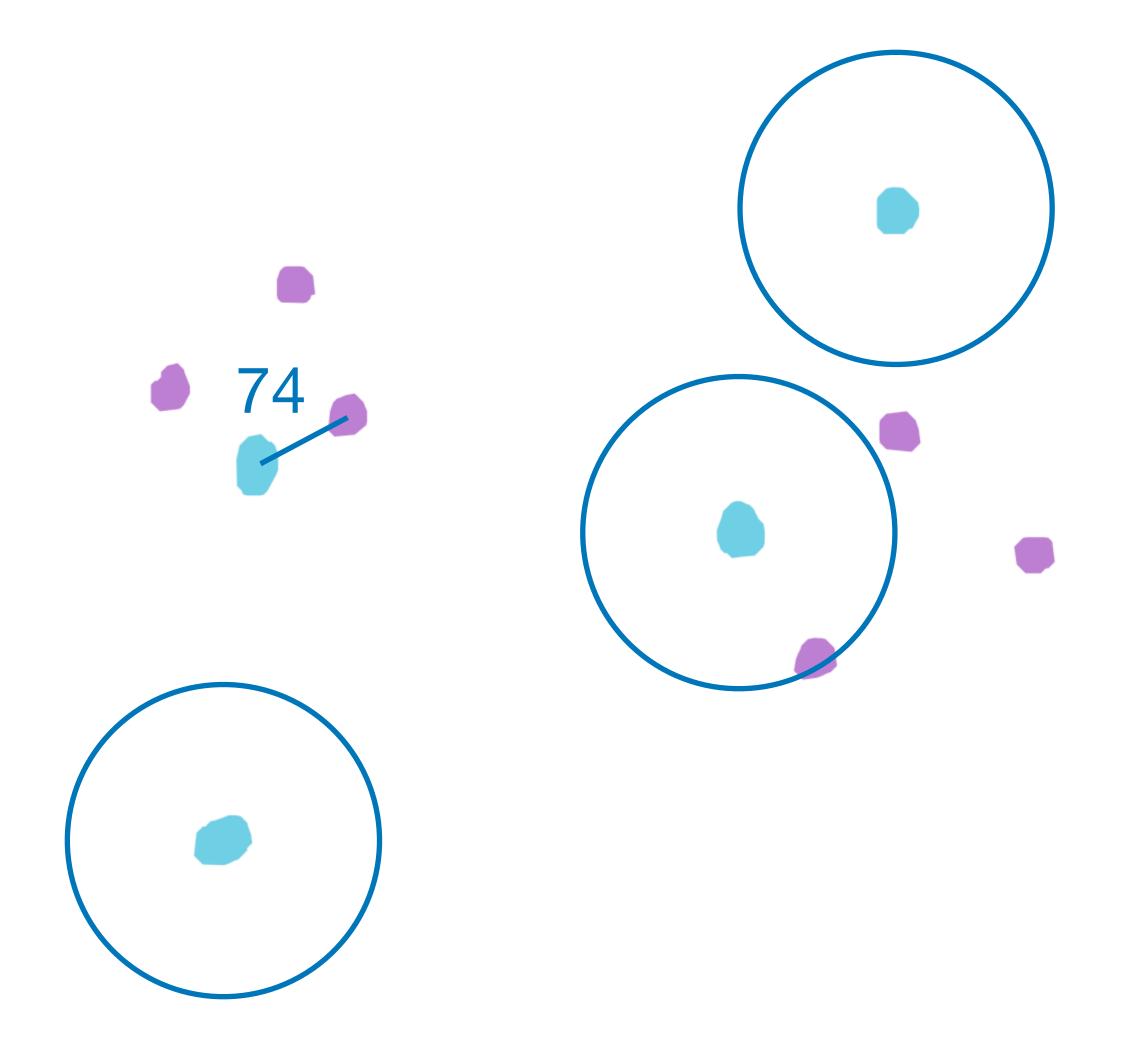
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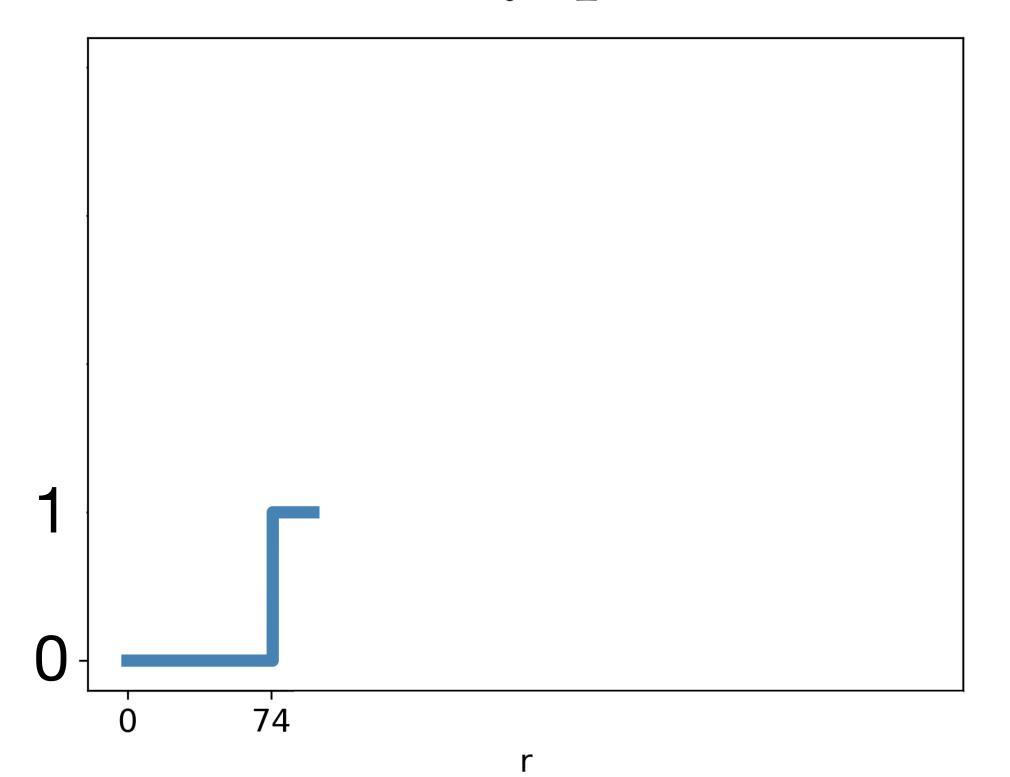








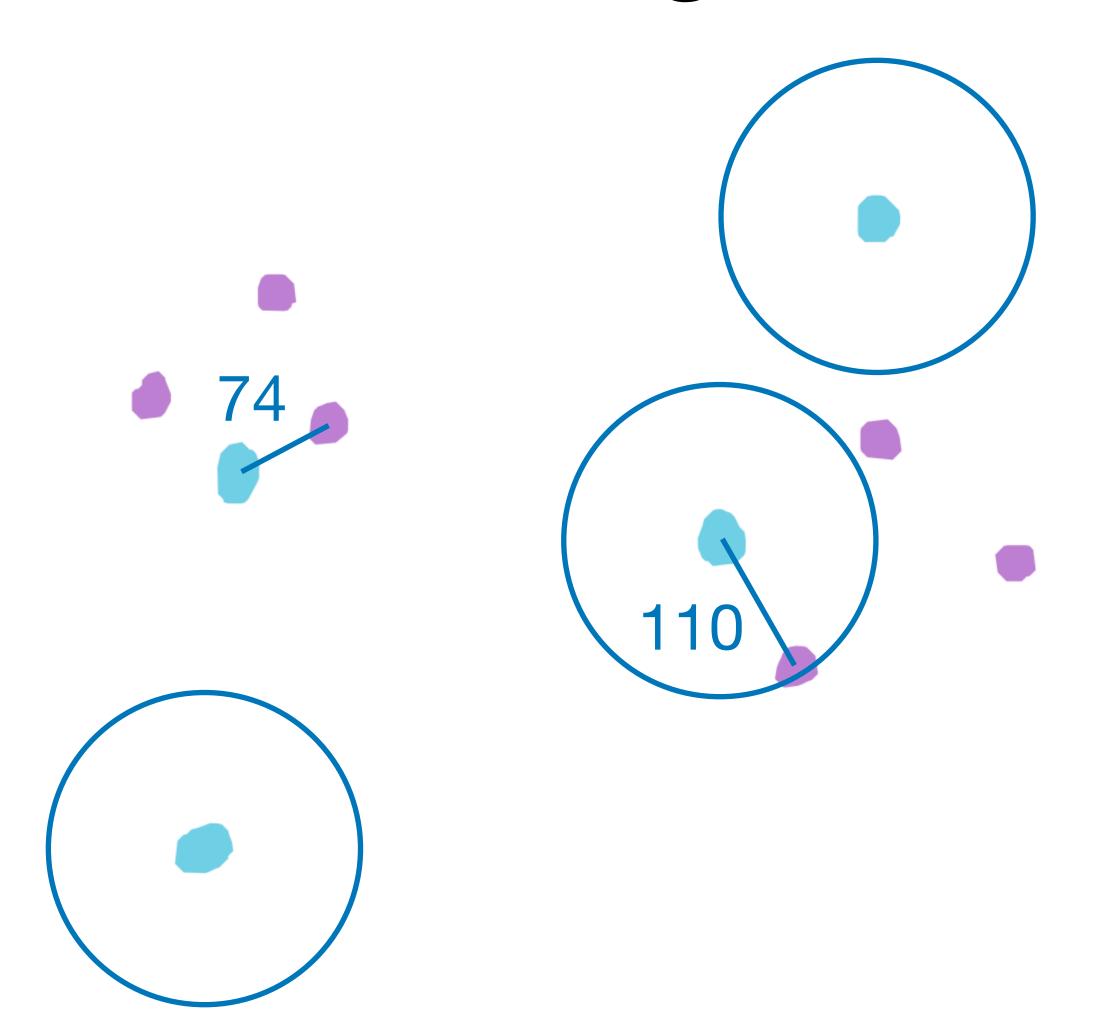
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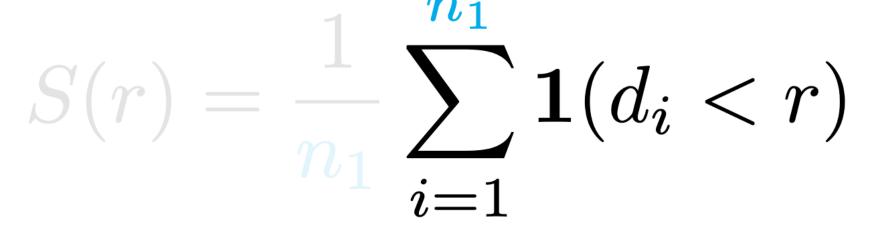


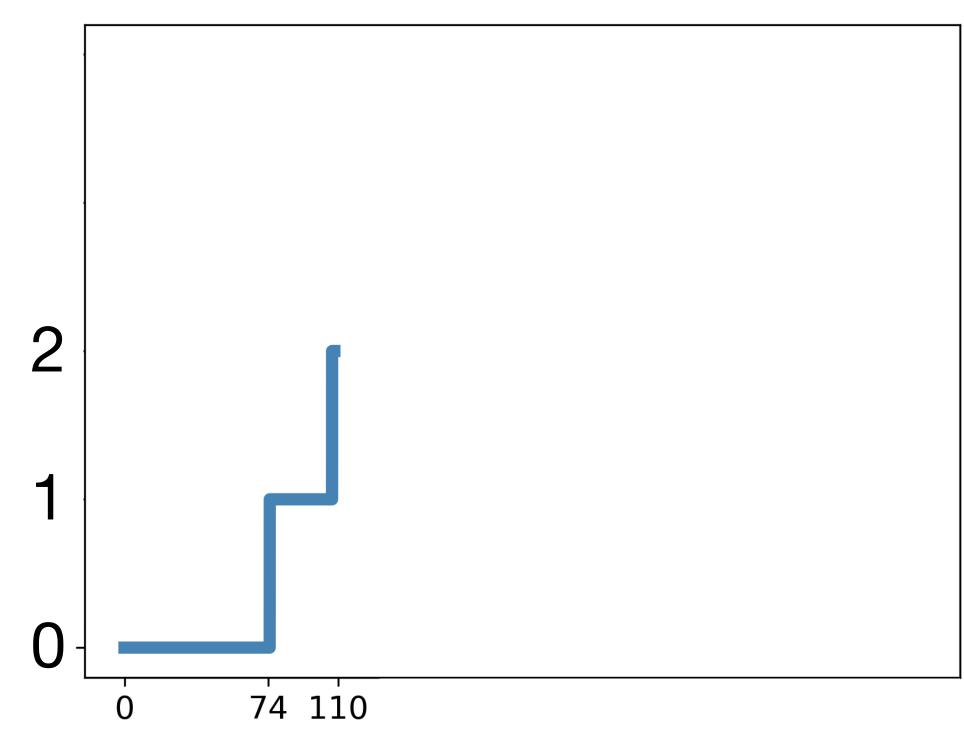








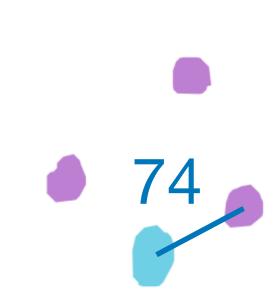


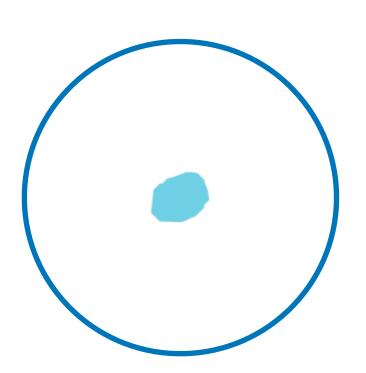


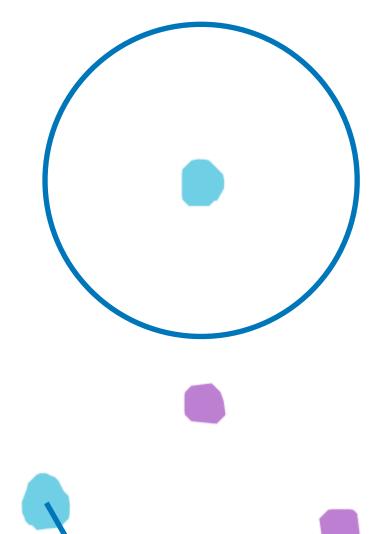




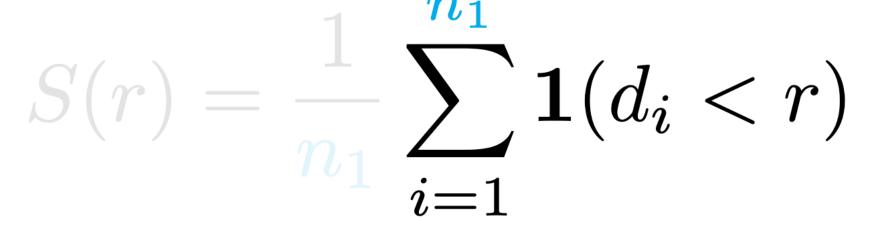


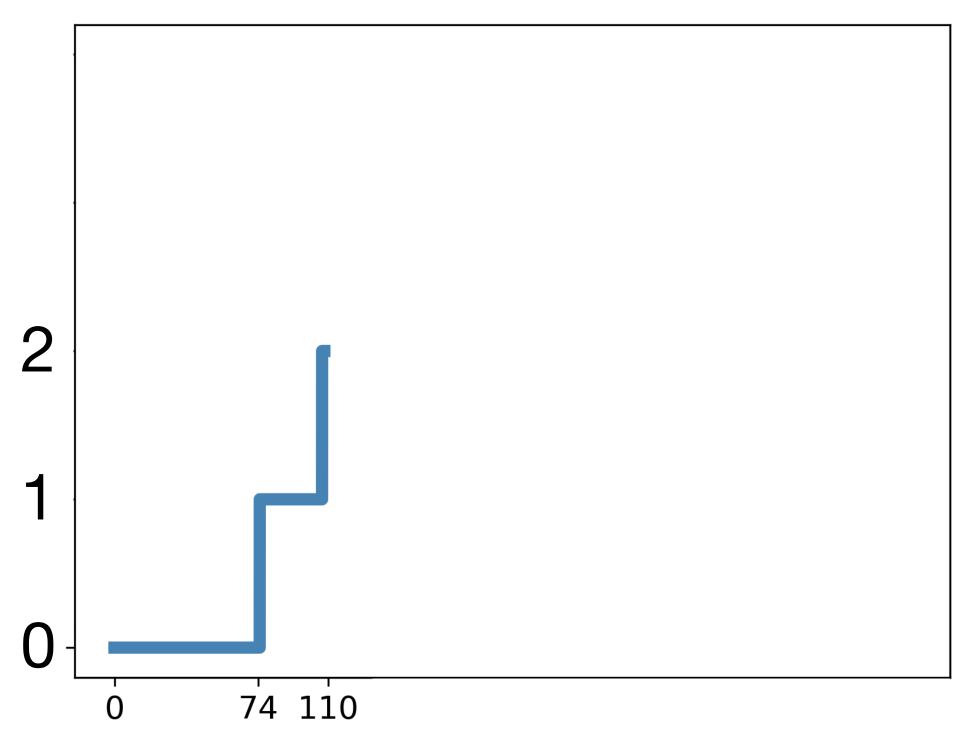








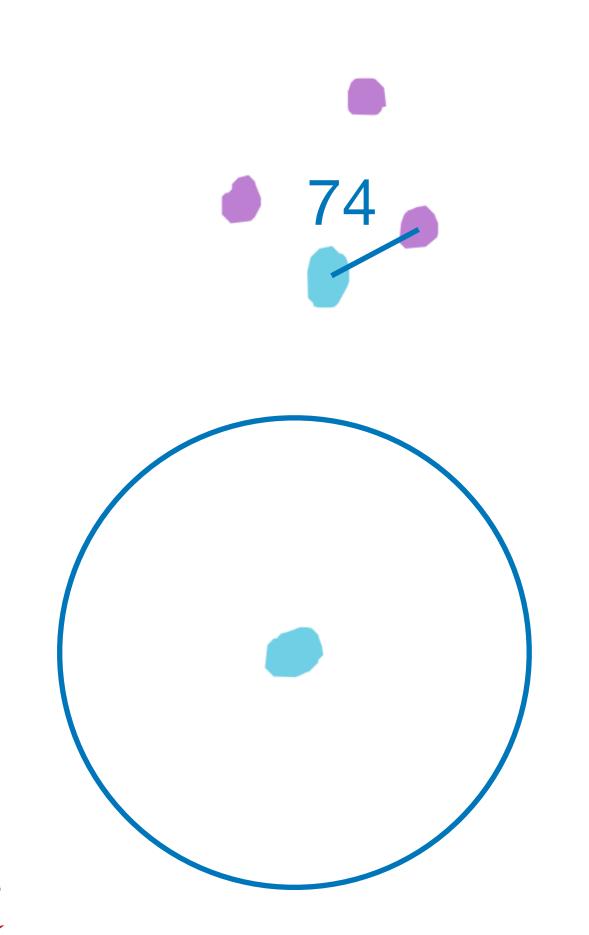


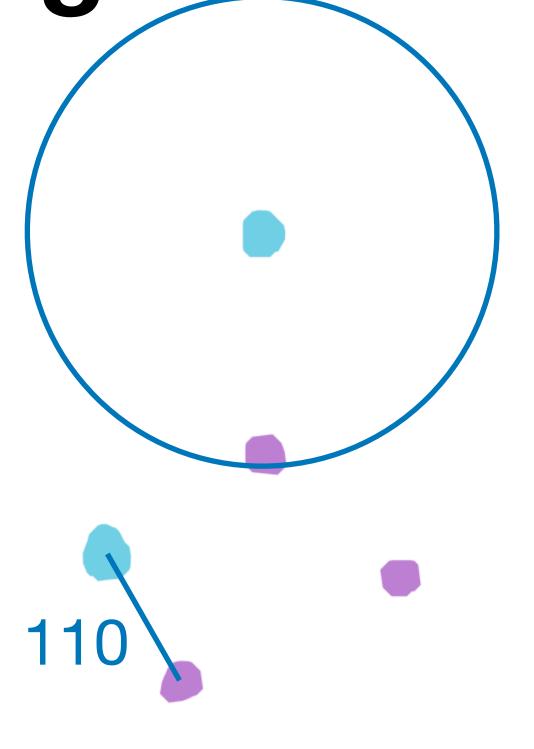


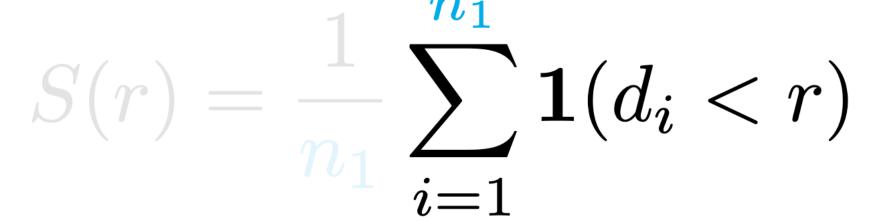


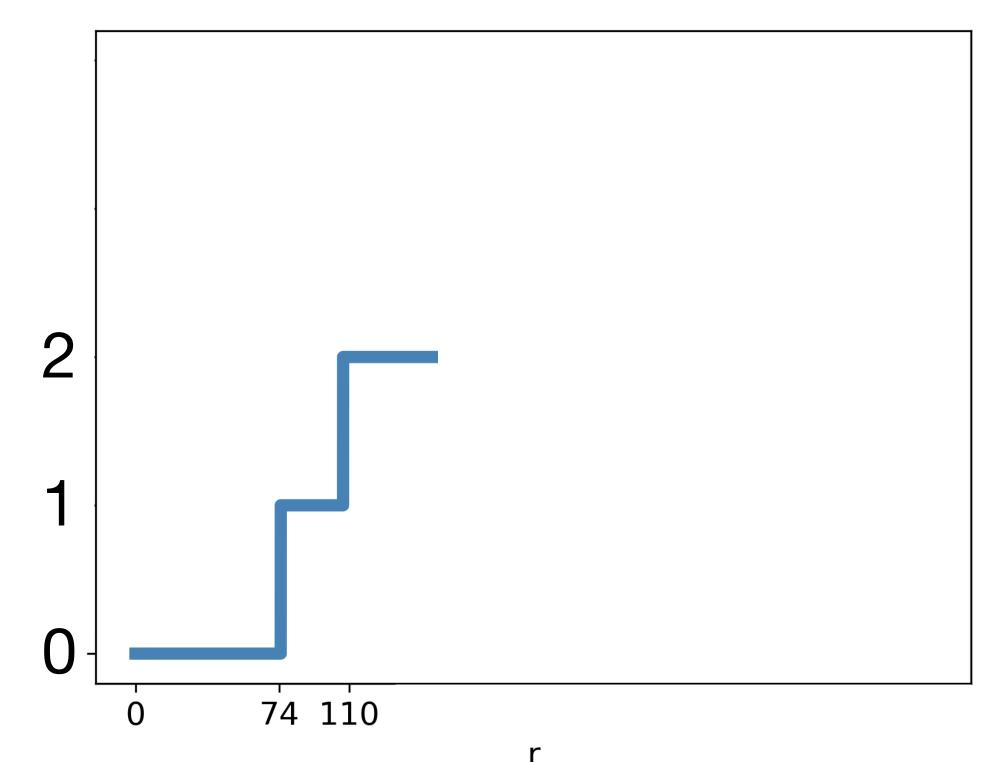








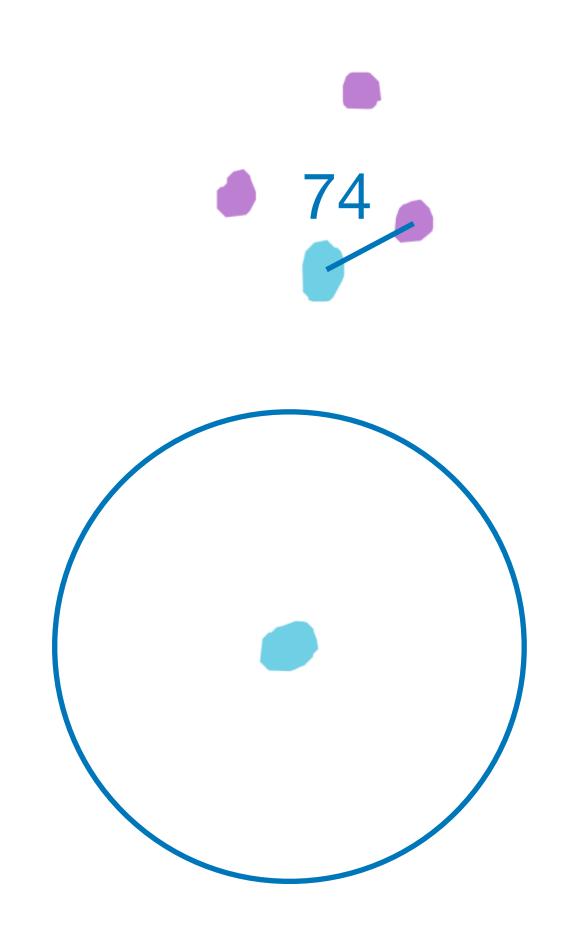


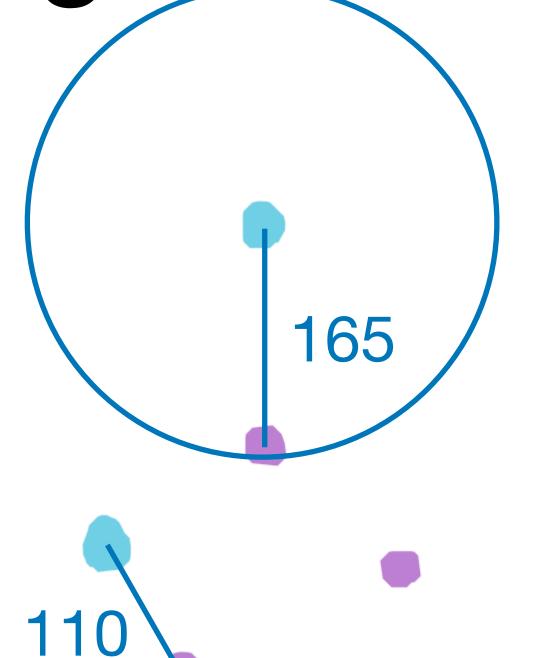


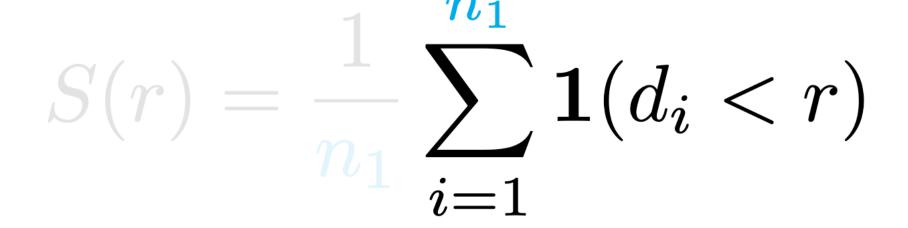


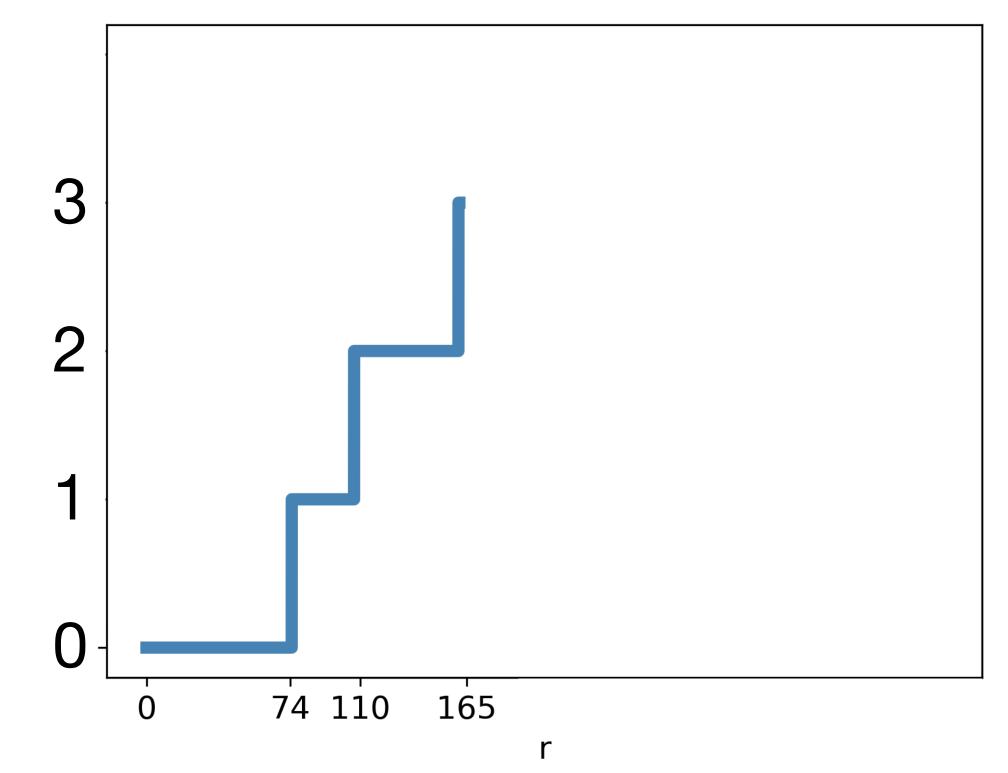








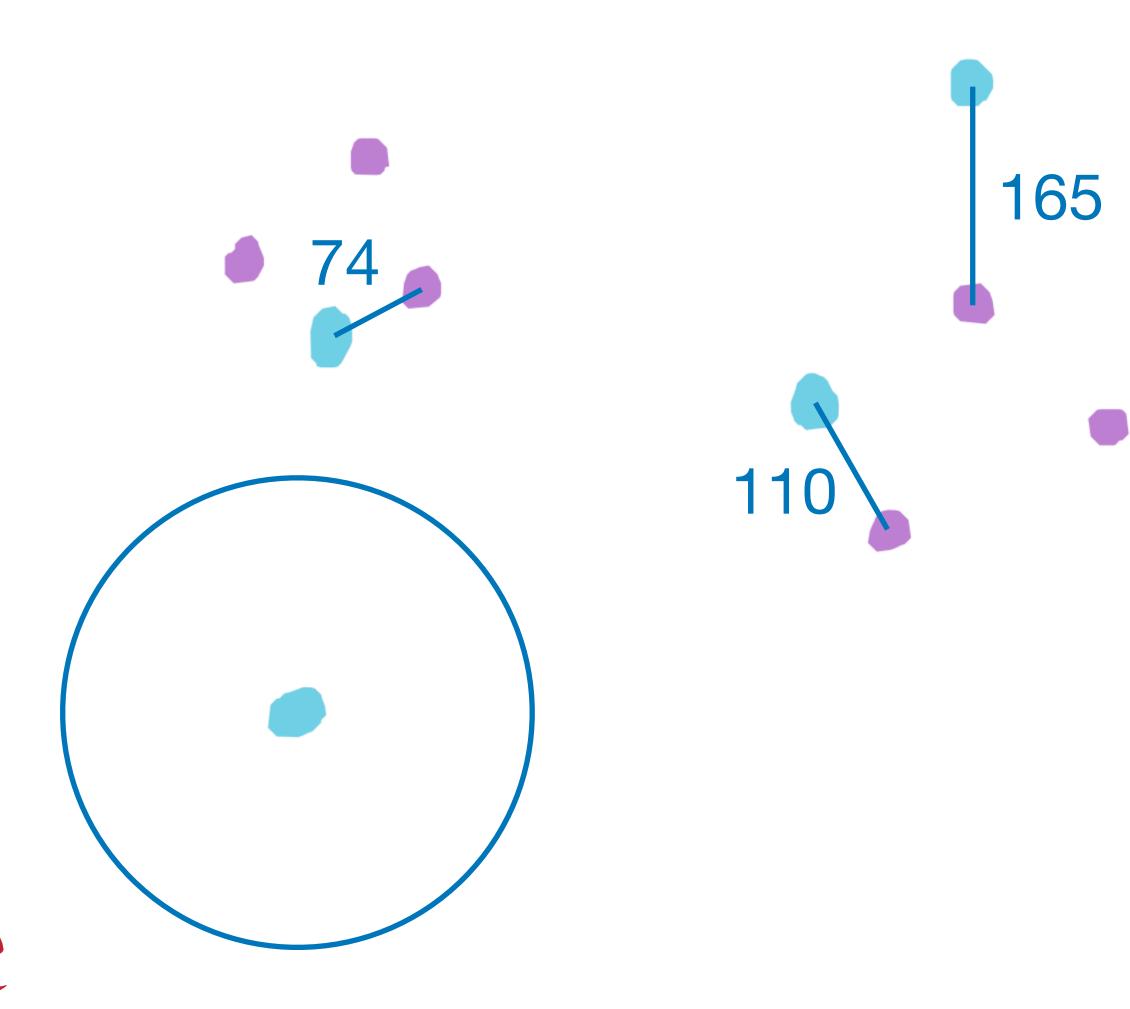




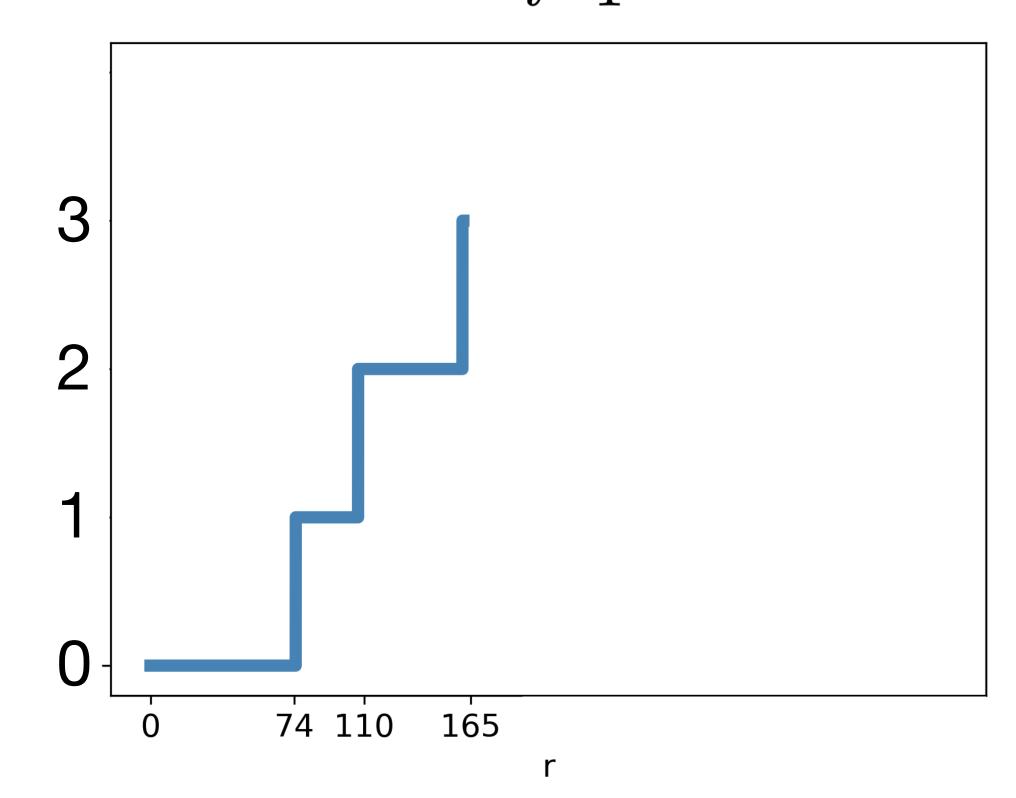








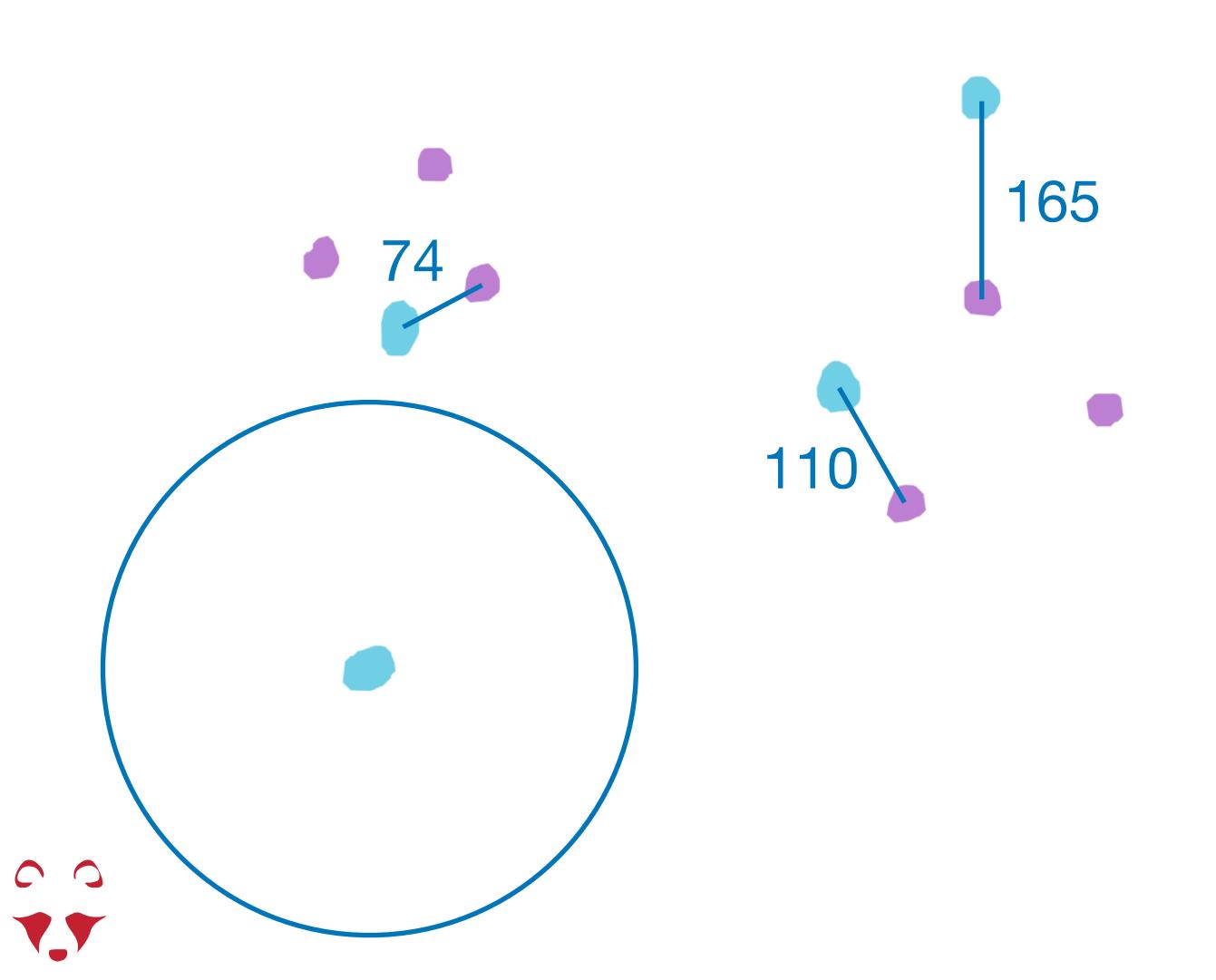
$$S(r) = \frac{1}{n_1} \sum_{i=1}^{n_1} \mathbf{1}(d_i < r)$$



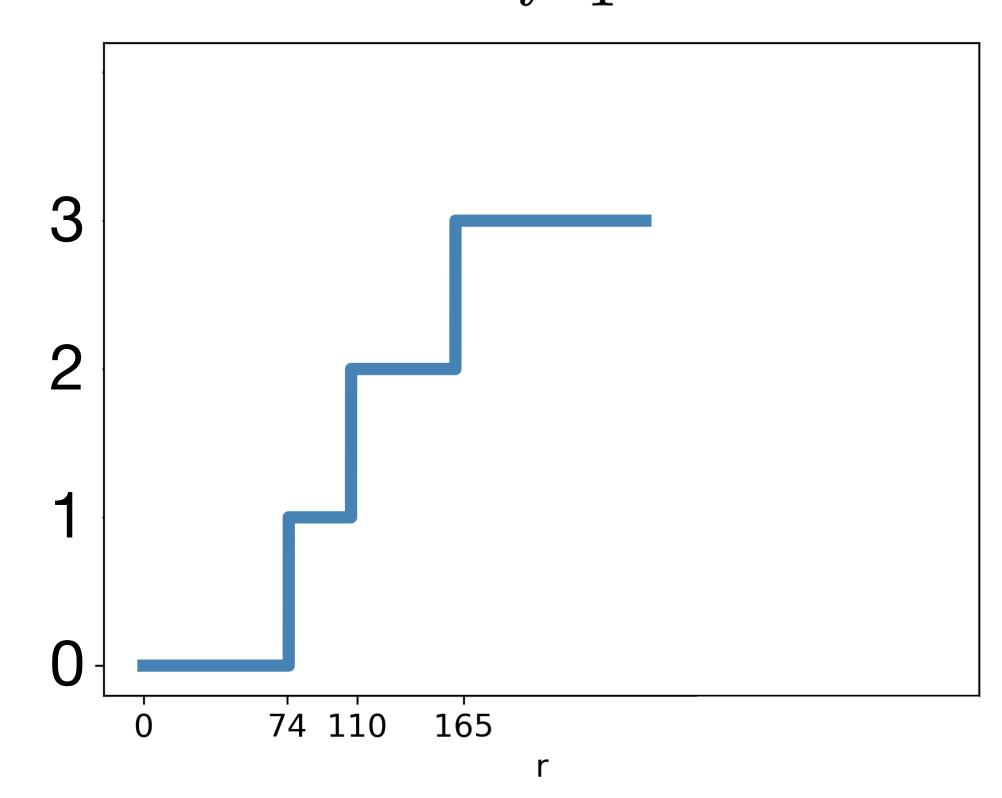






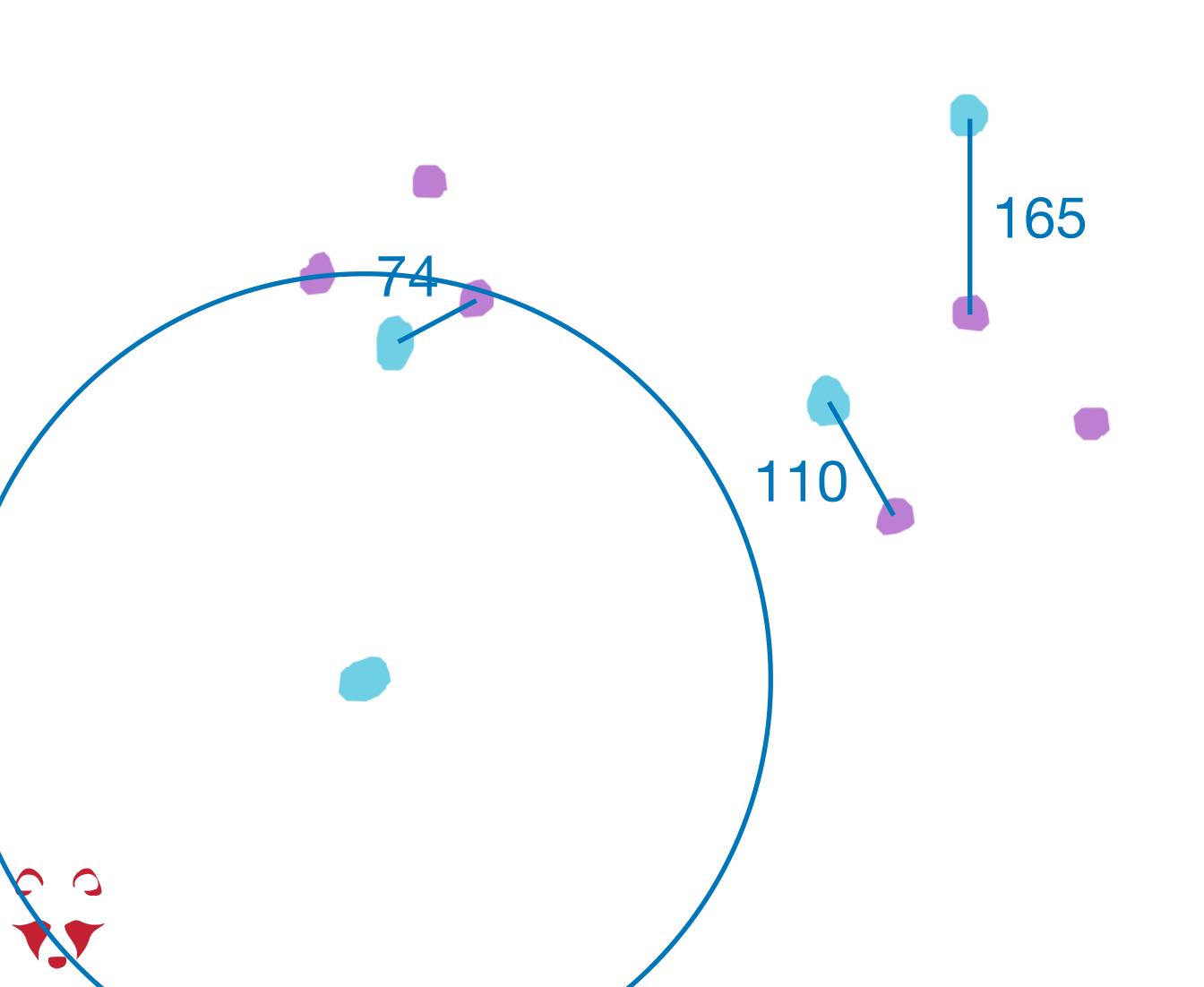


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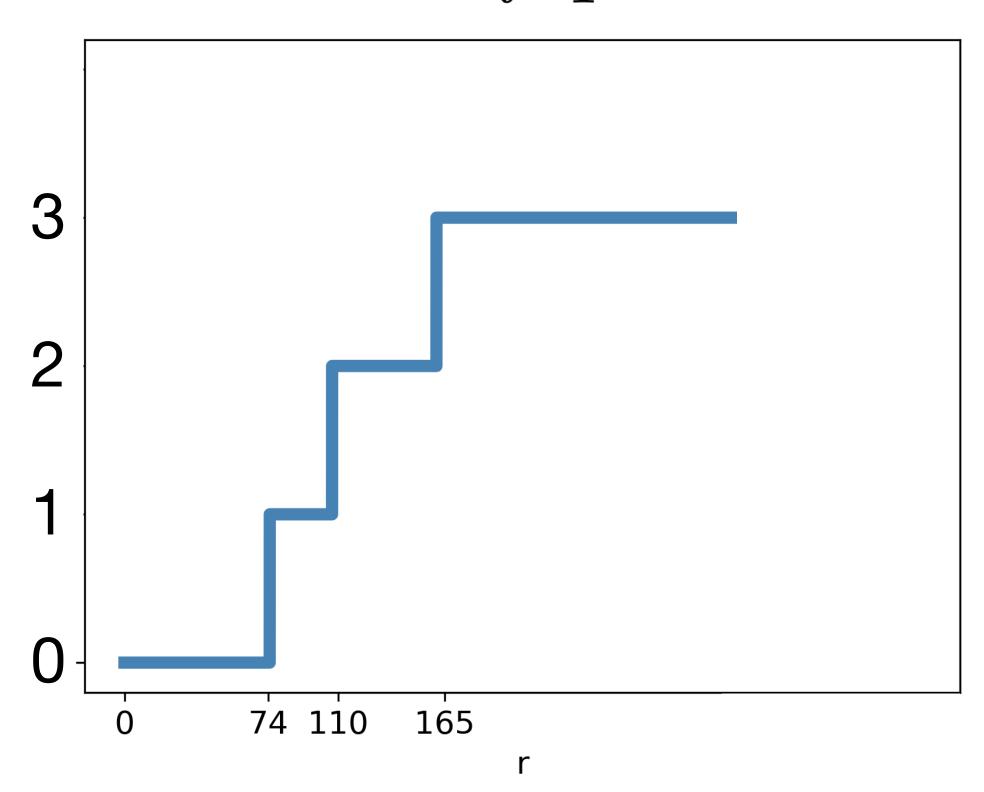






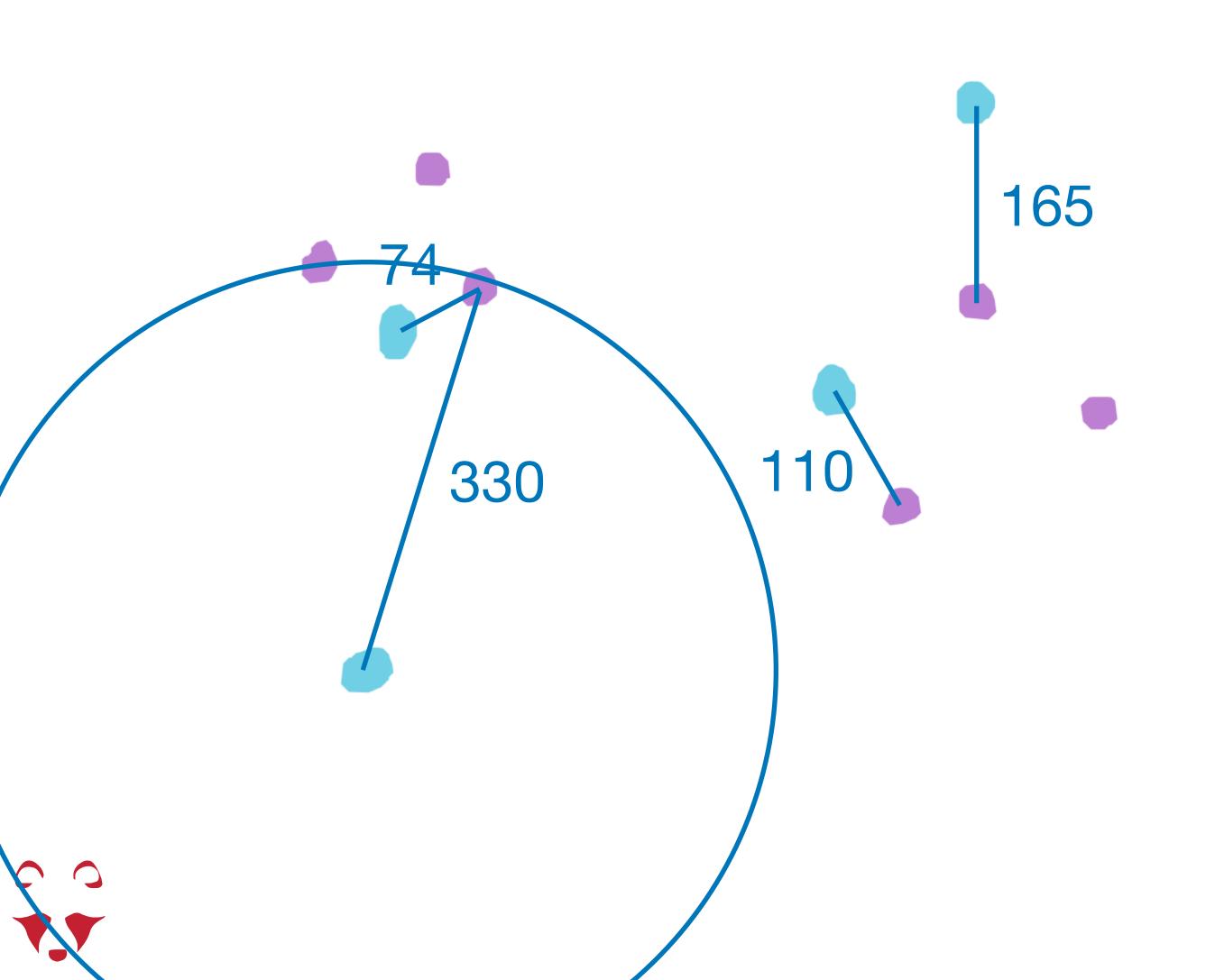


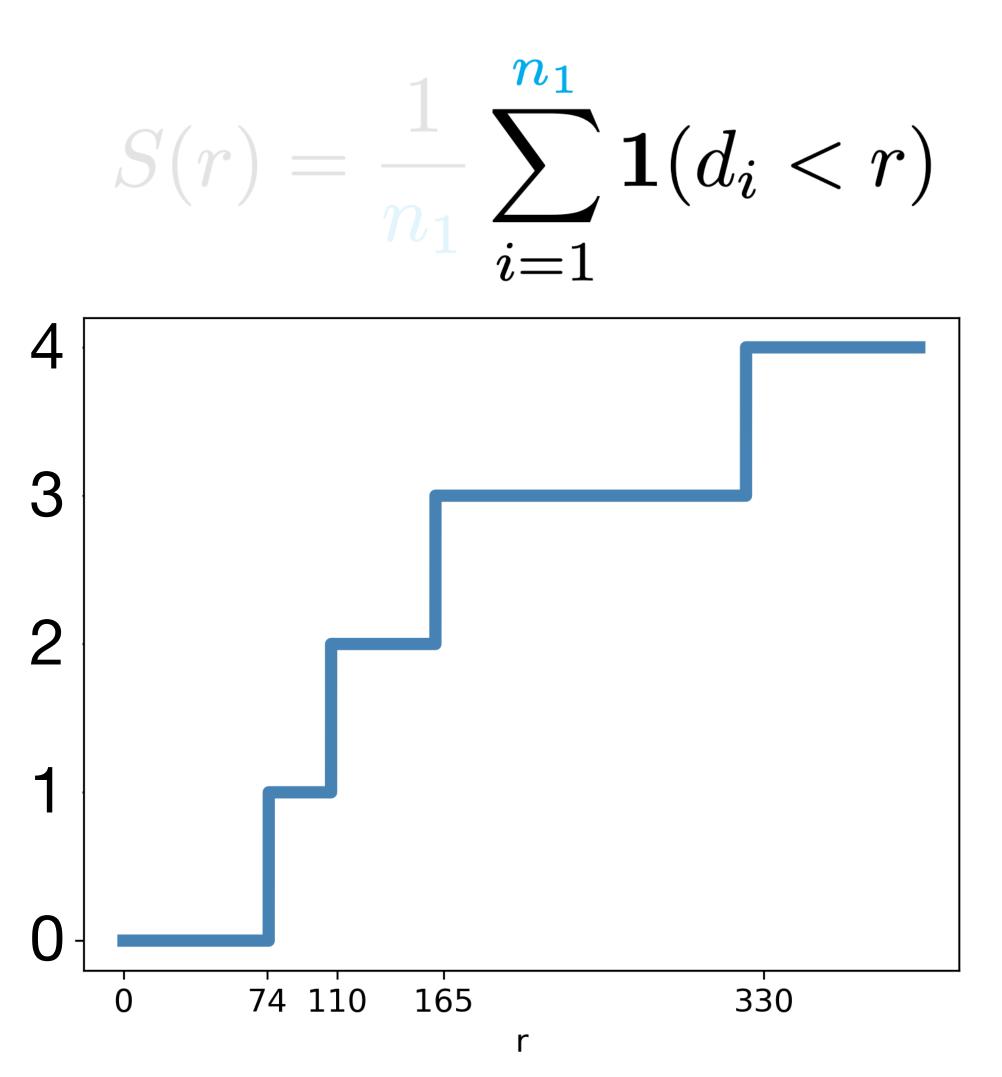
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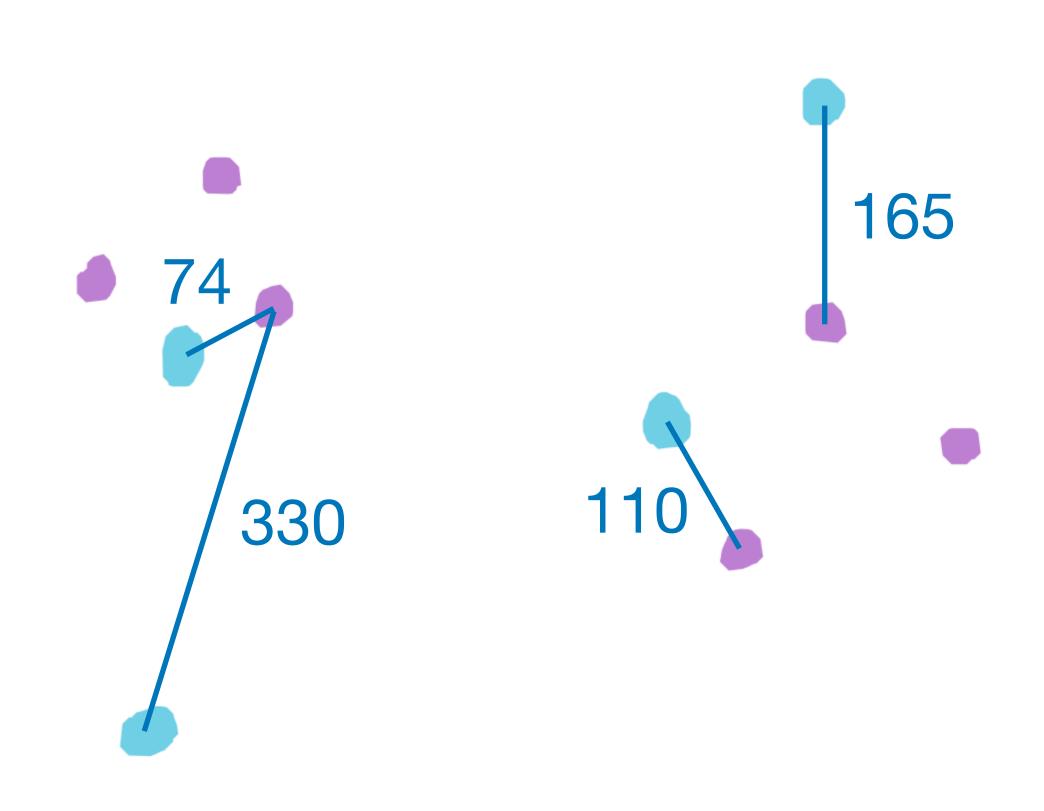


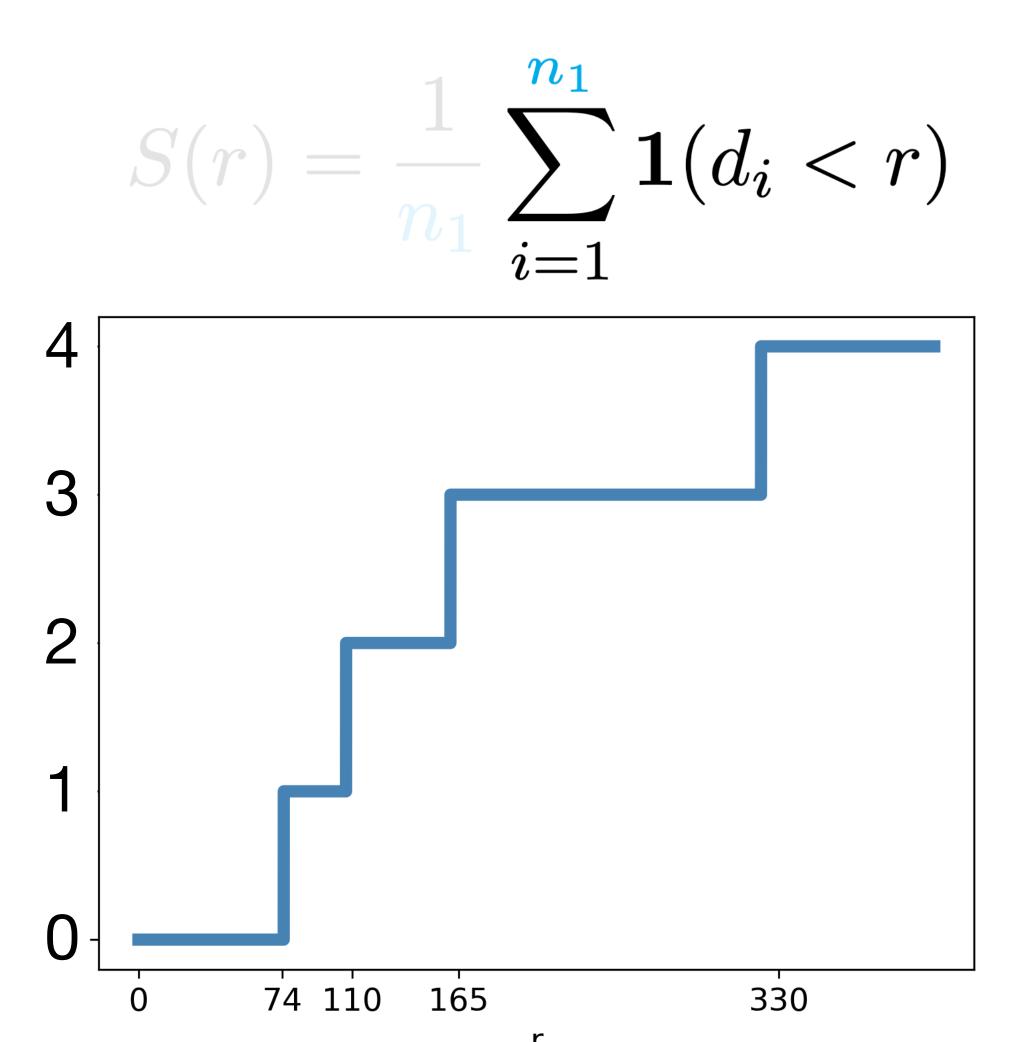








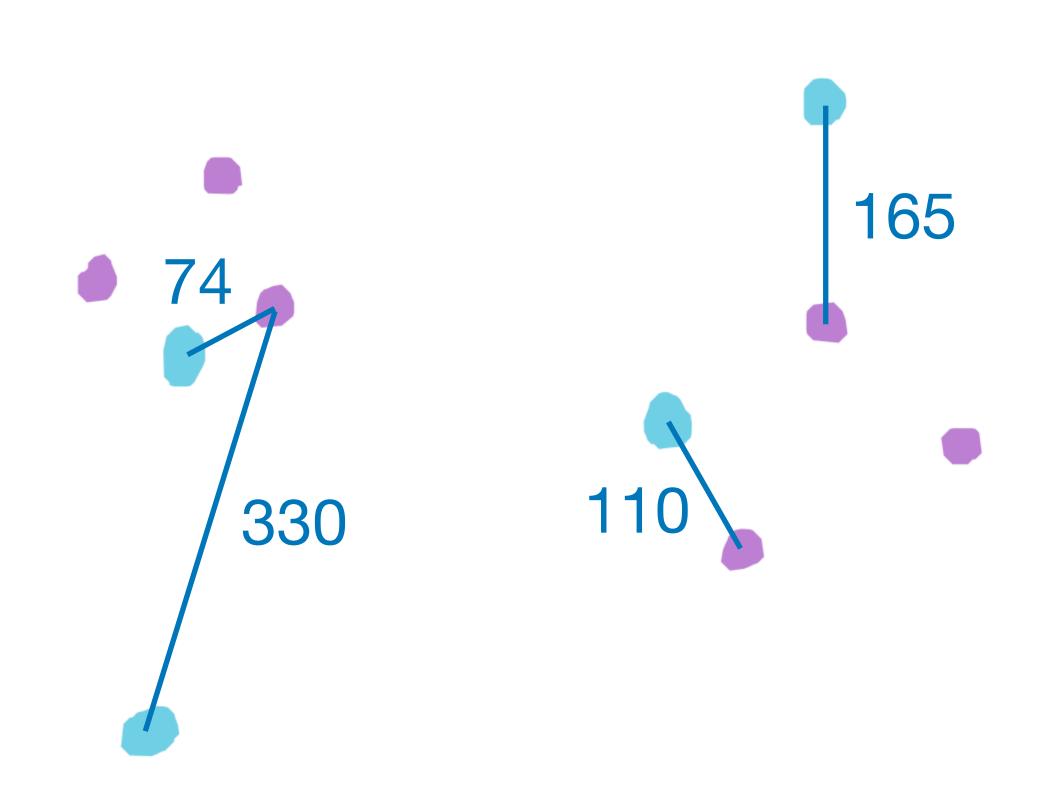


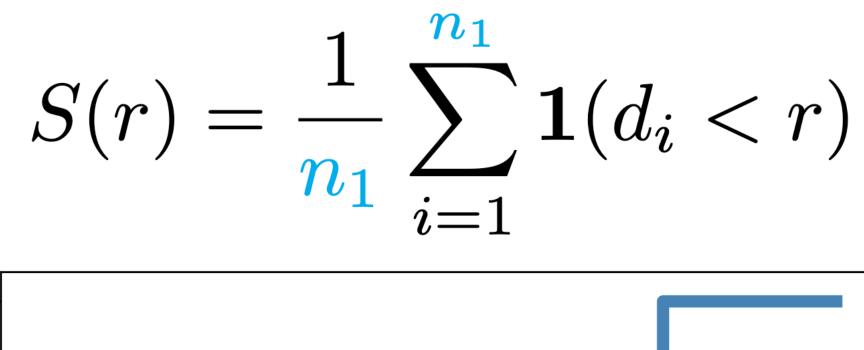


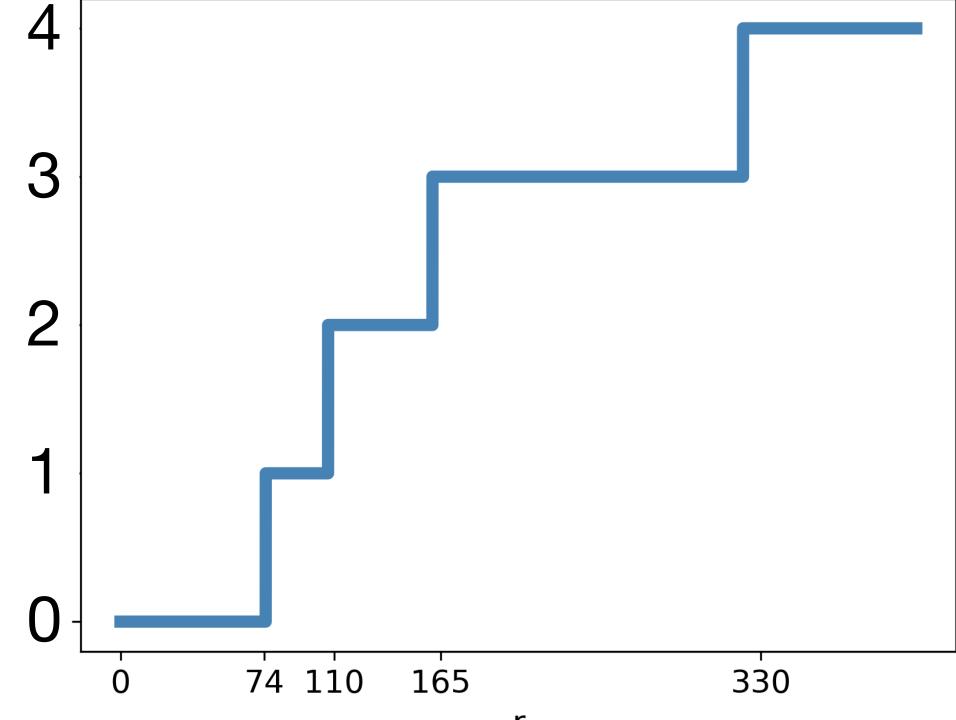








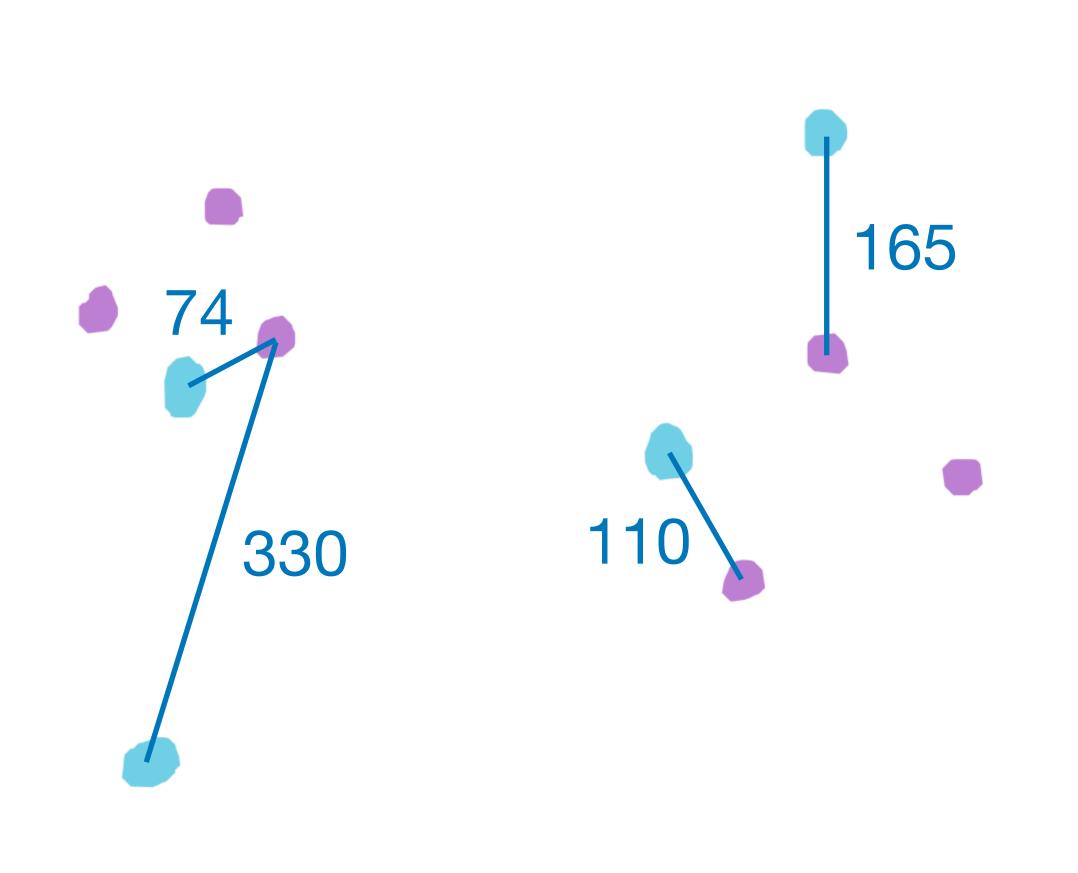


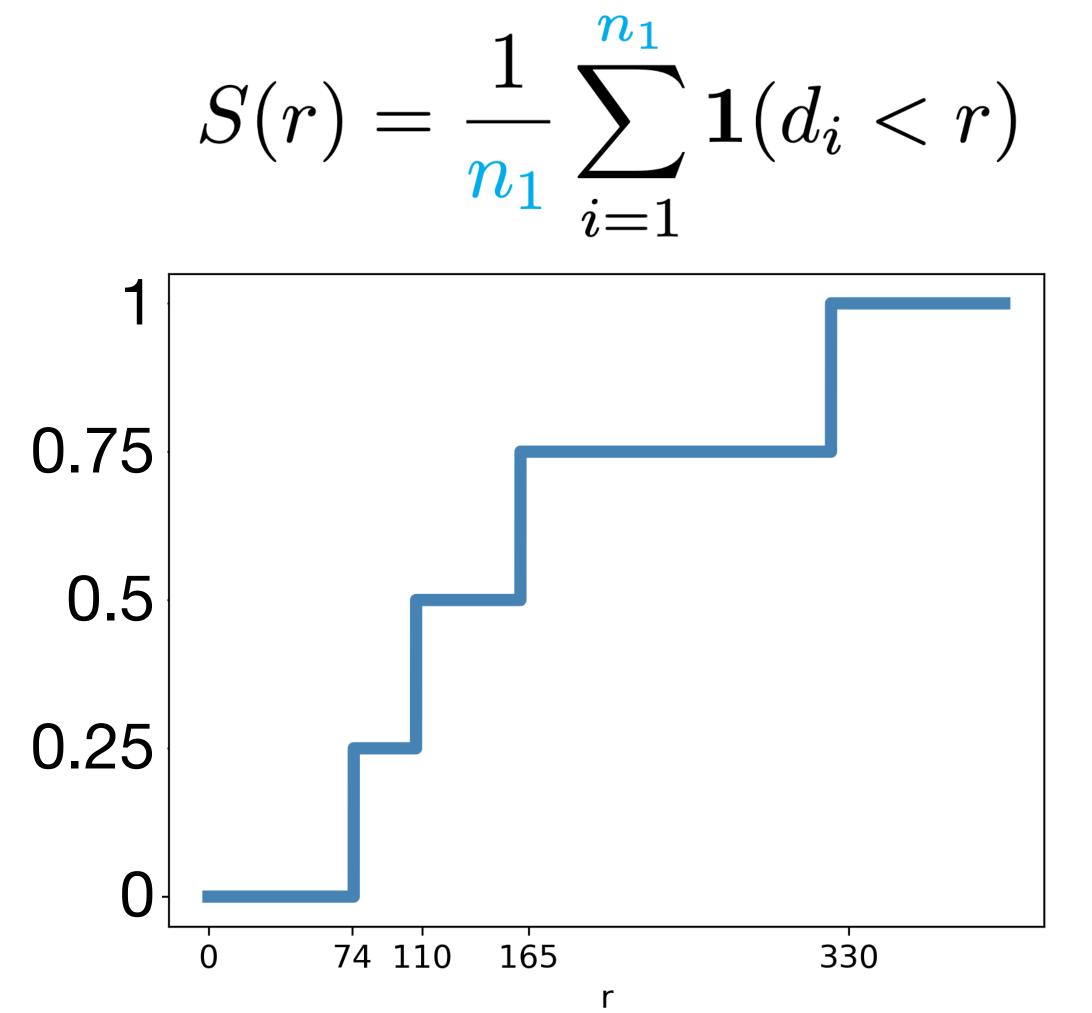








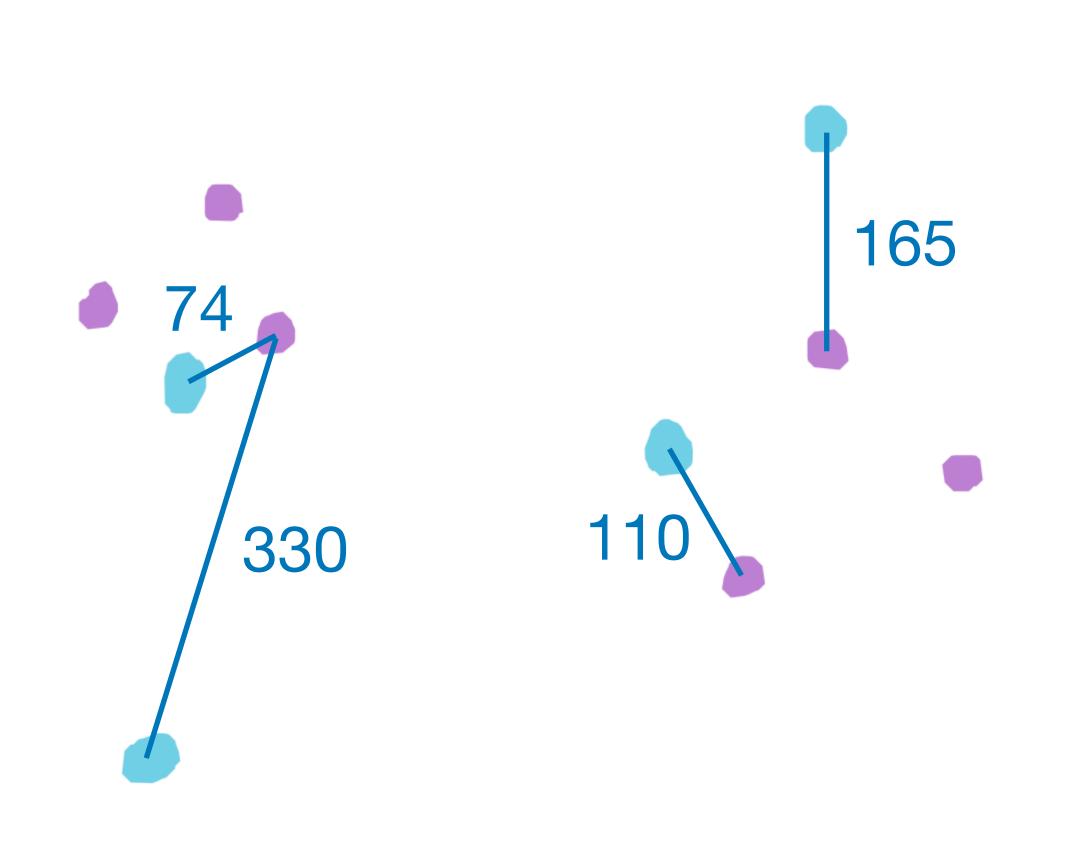


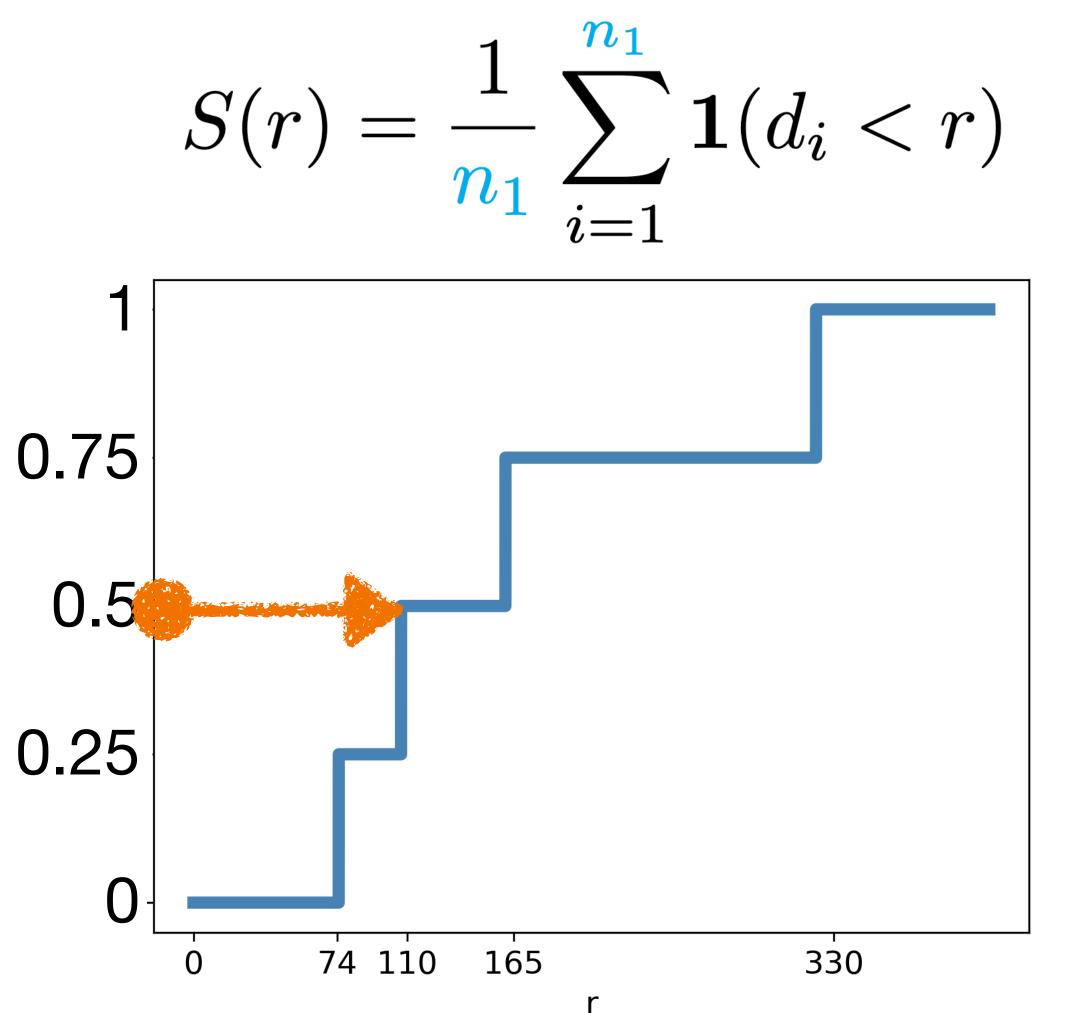




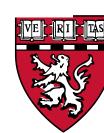




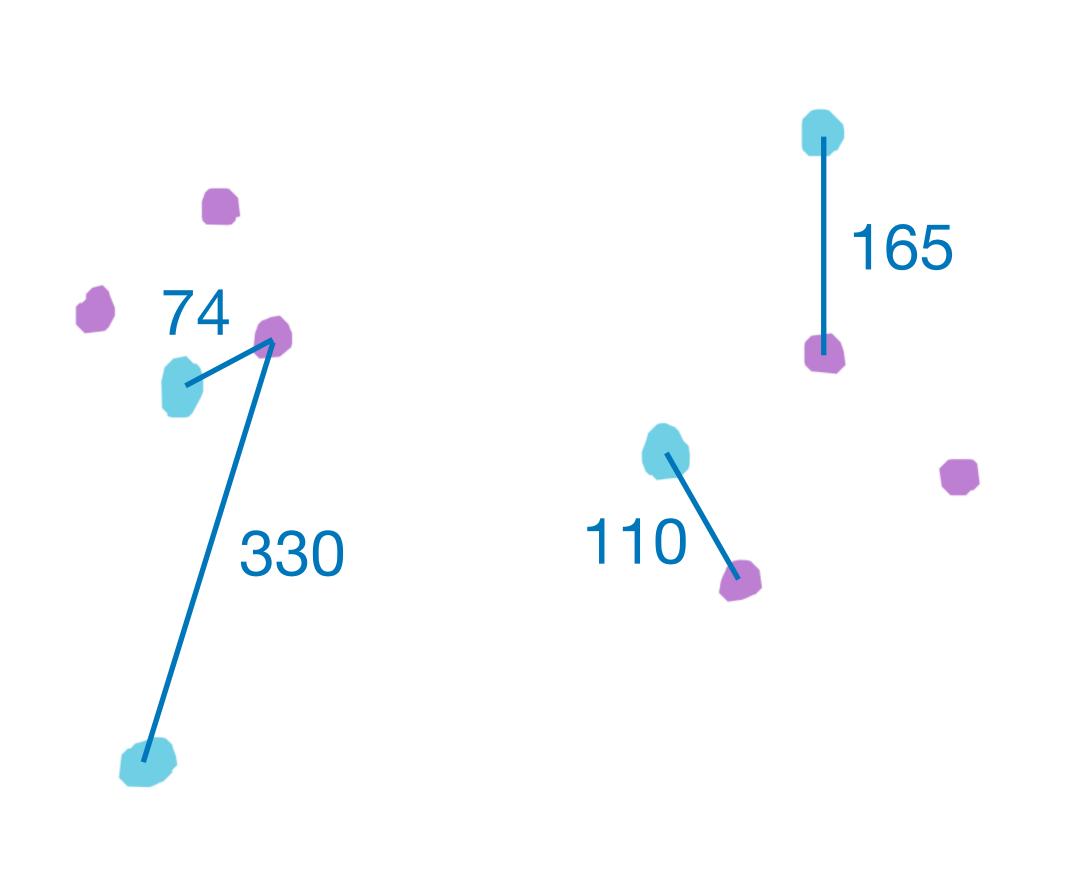


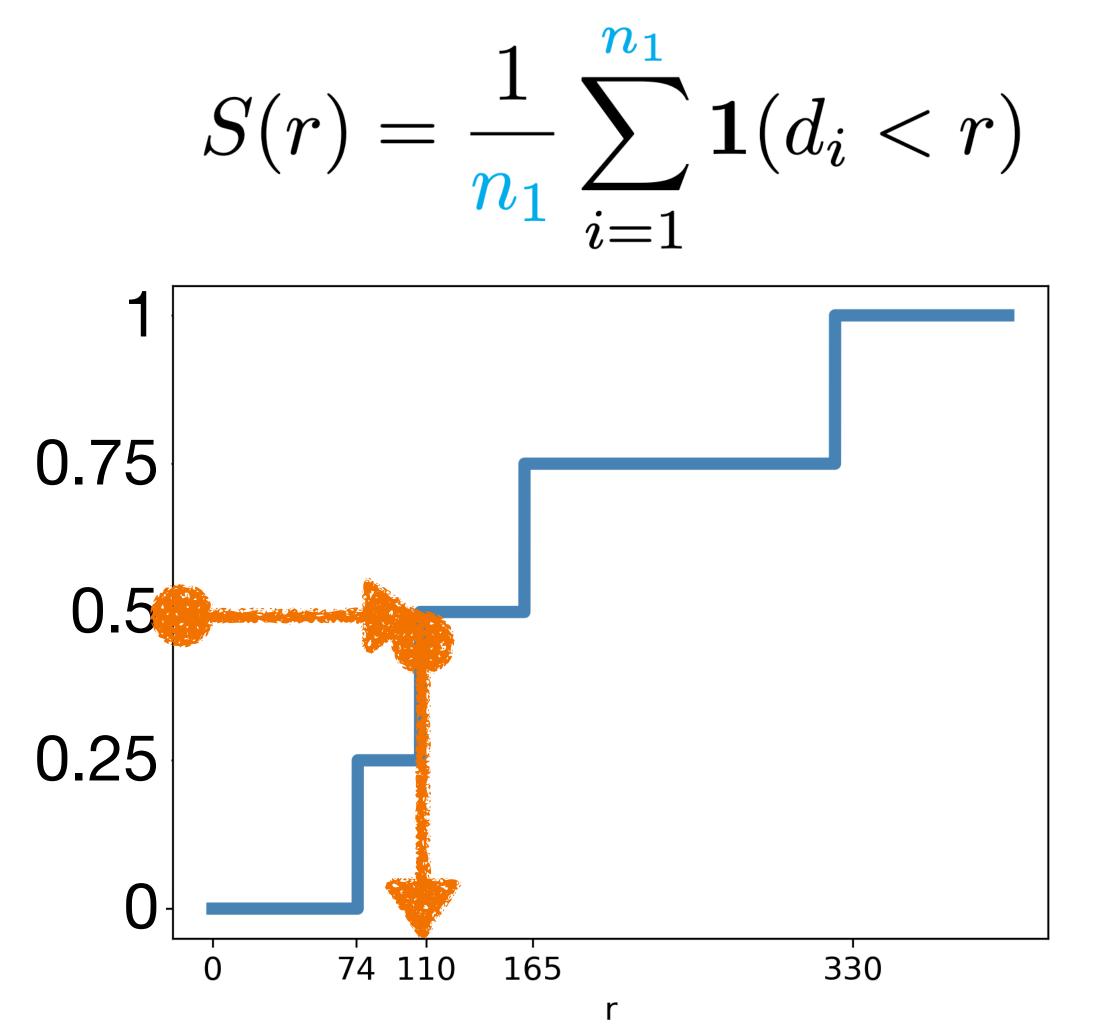








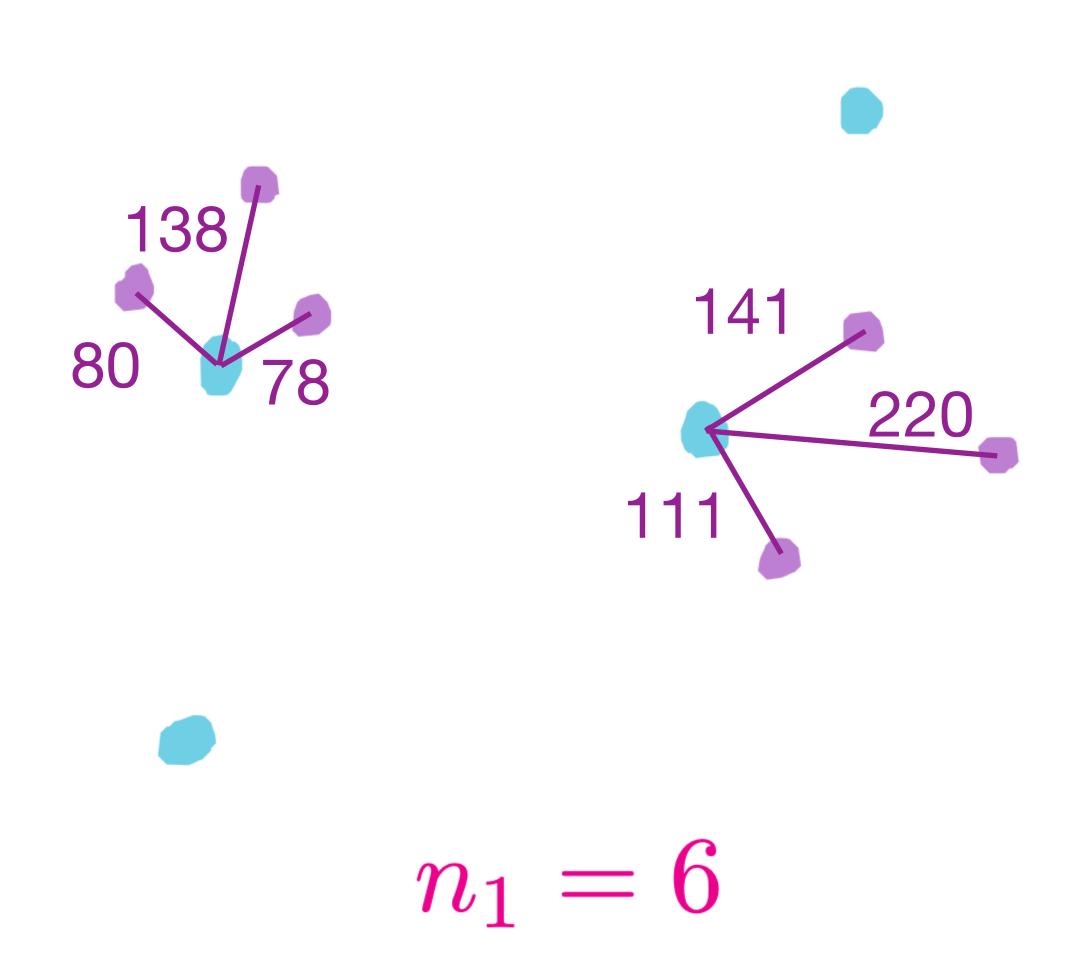




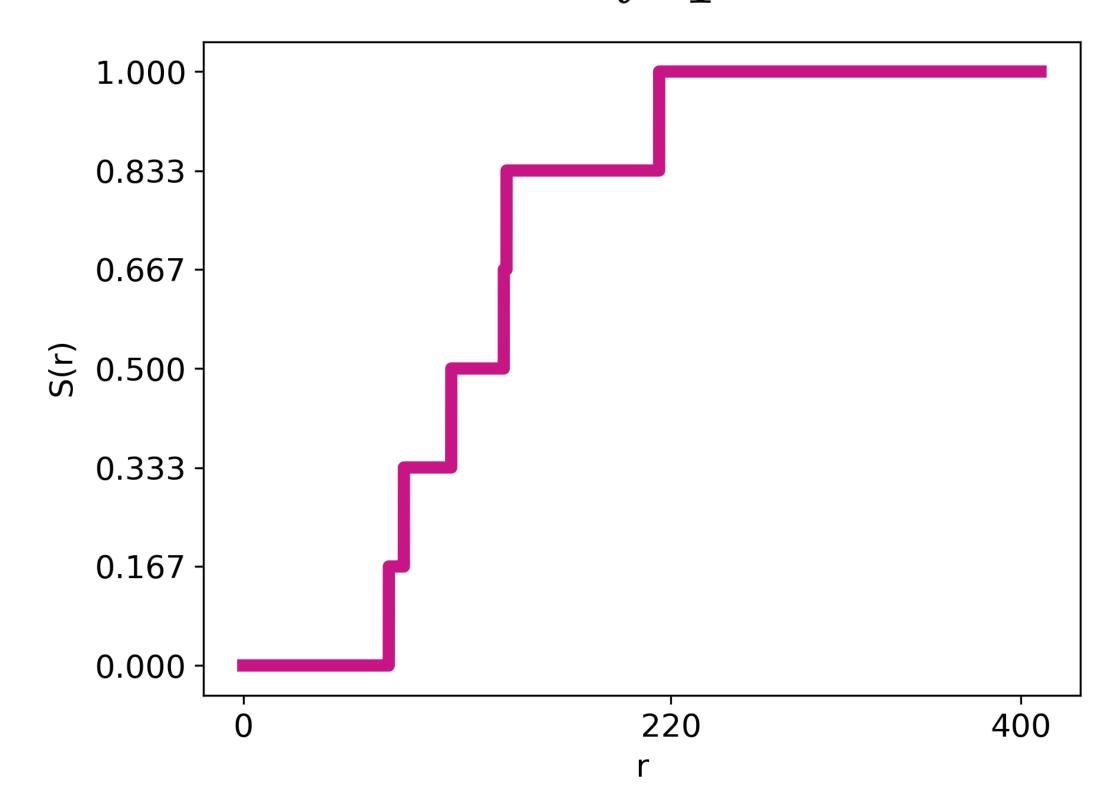








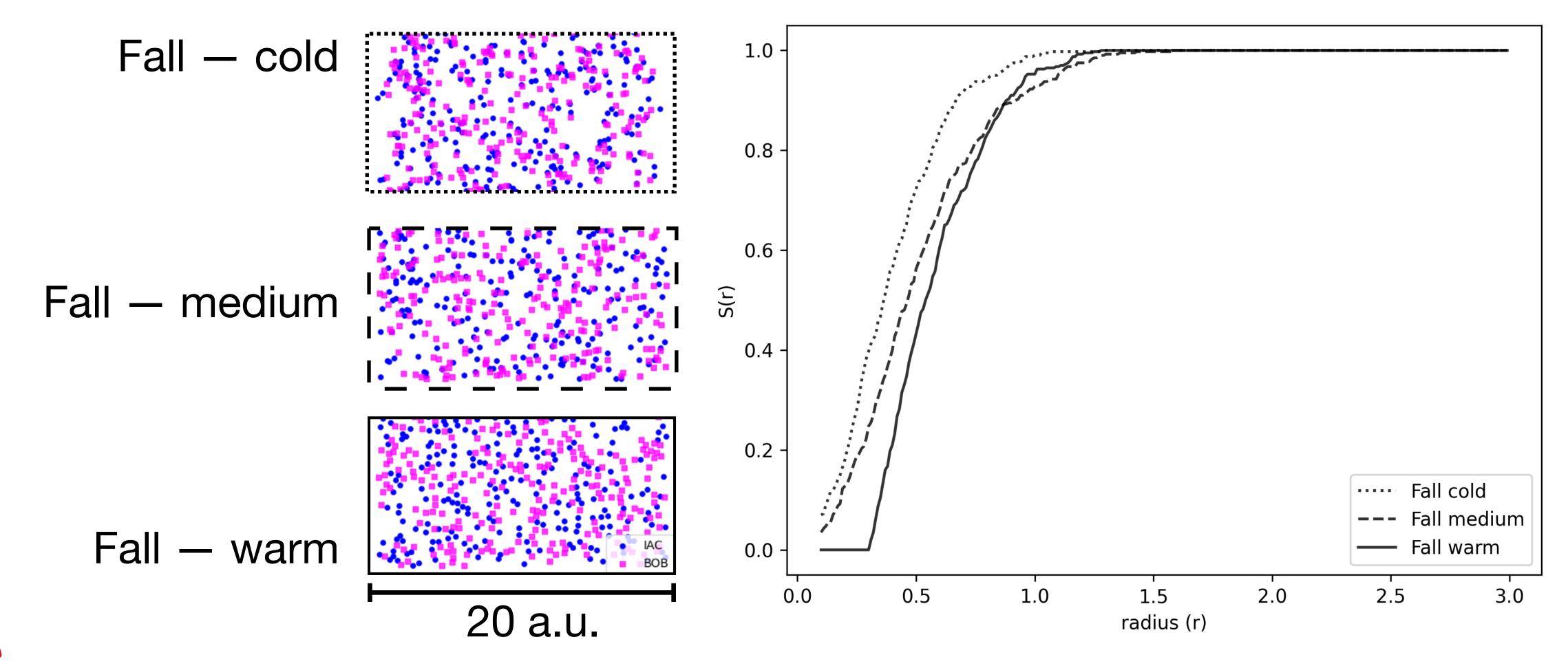
$$S(r) = \frac{1}{n_1} \sum_{i=1}^{n_1} \mathbf{1}(d_i < r)$$







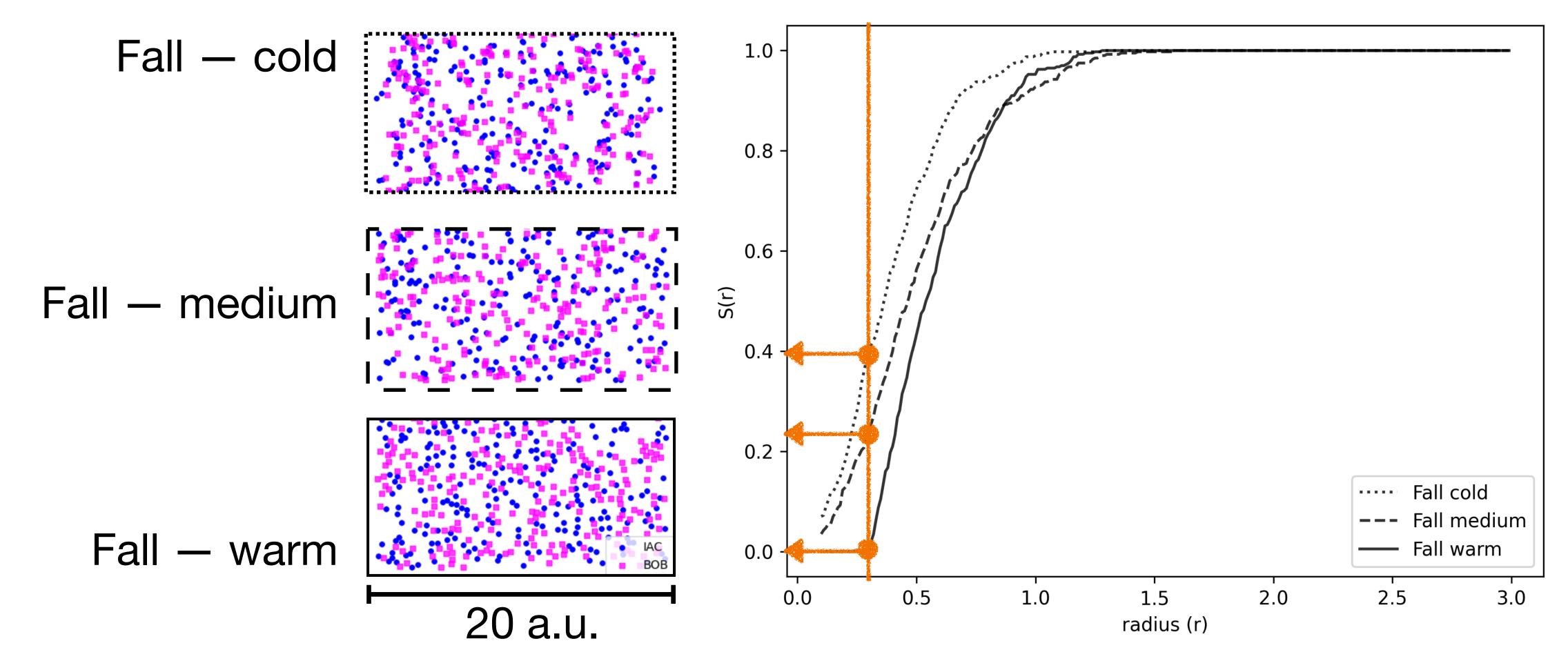








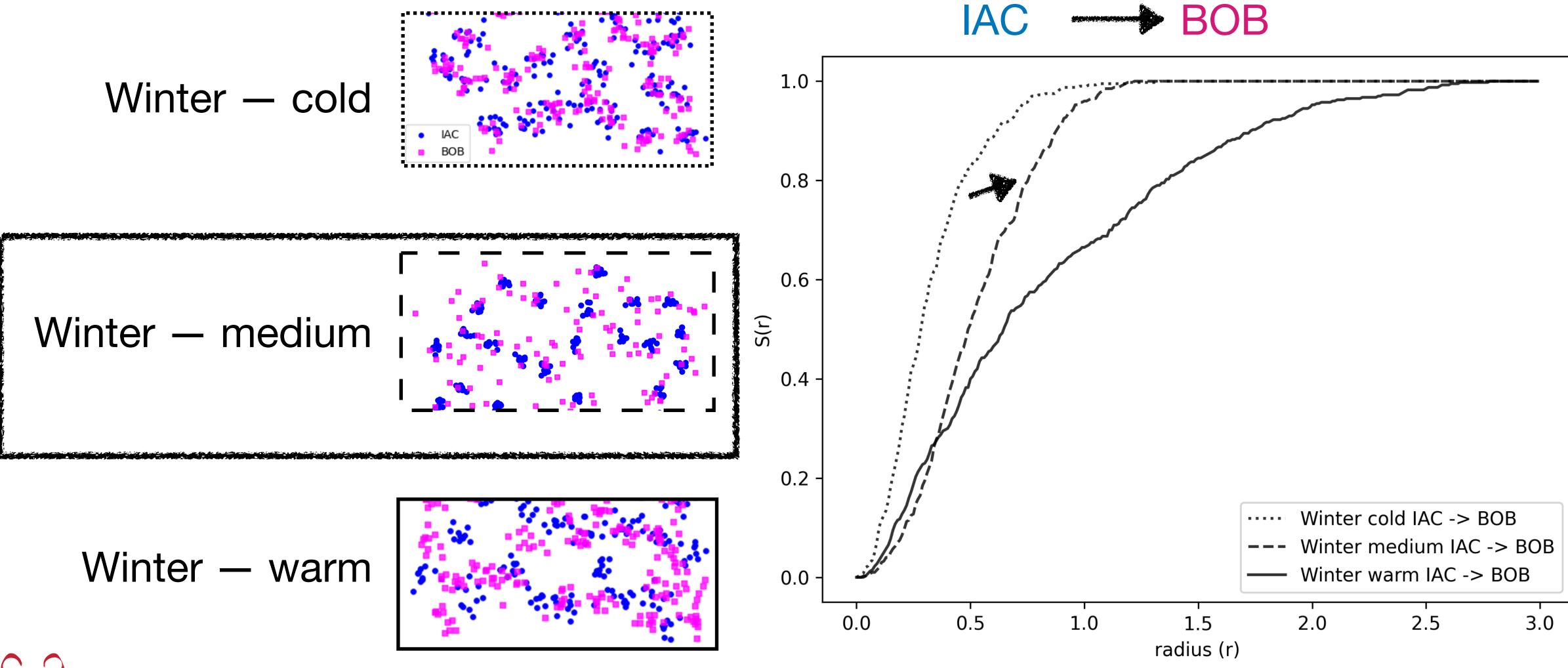








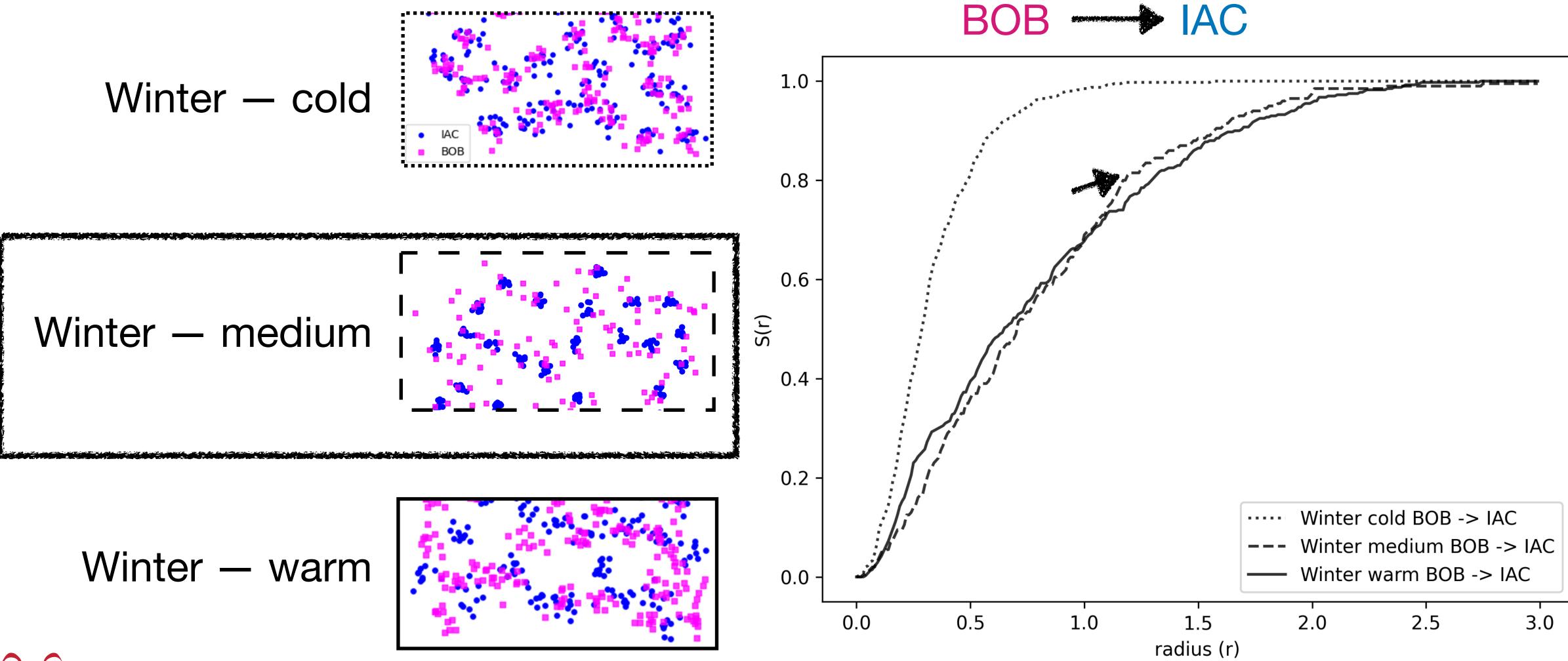






















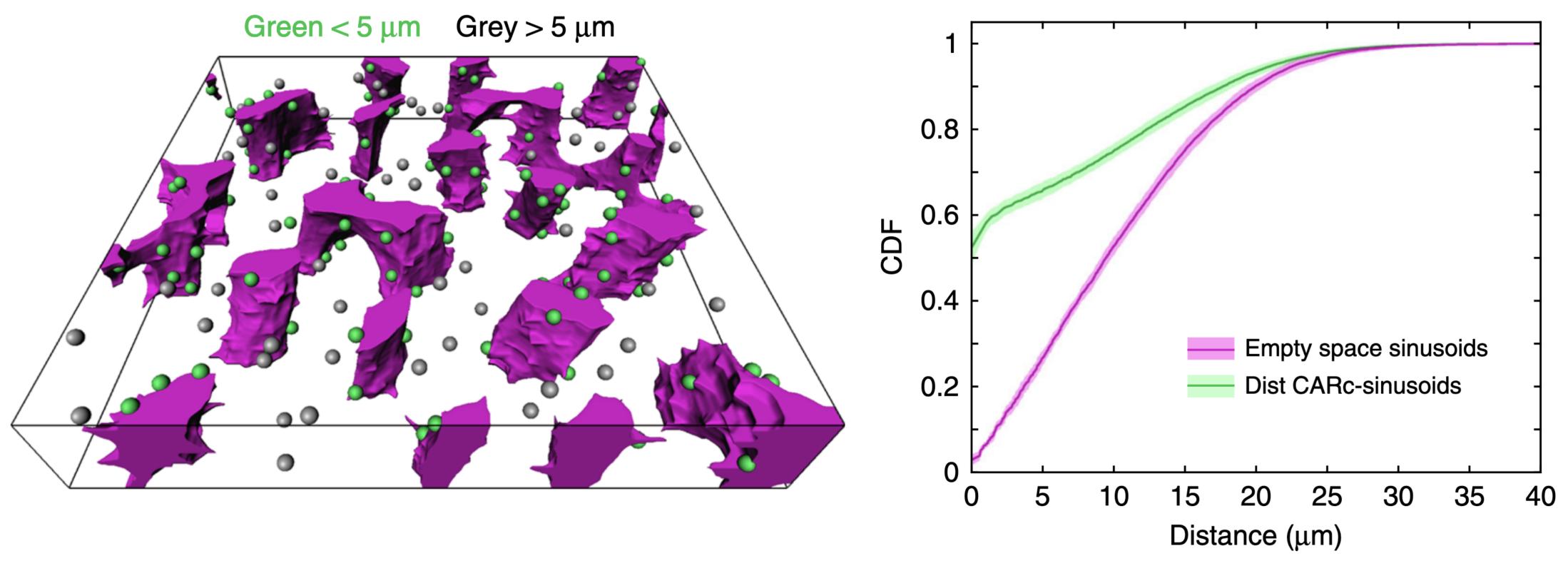
- Asymmetric: BOB → IAC ≠ IAC → BOB
- Returns: A number for each radius
- Range: Short







Beyond the nearest neighbor function

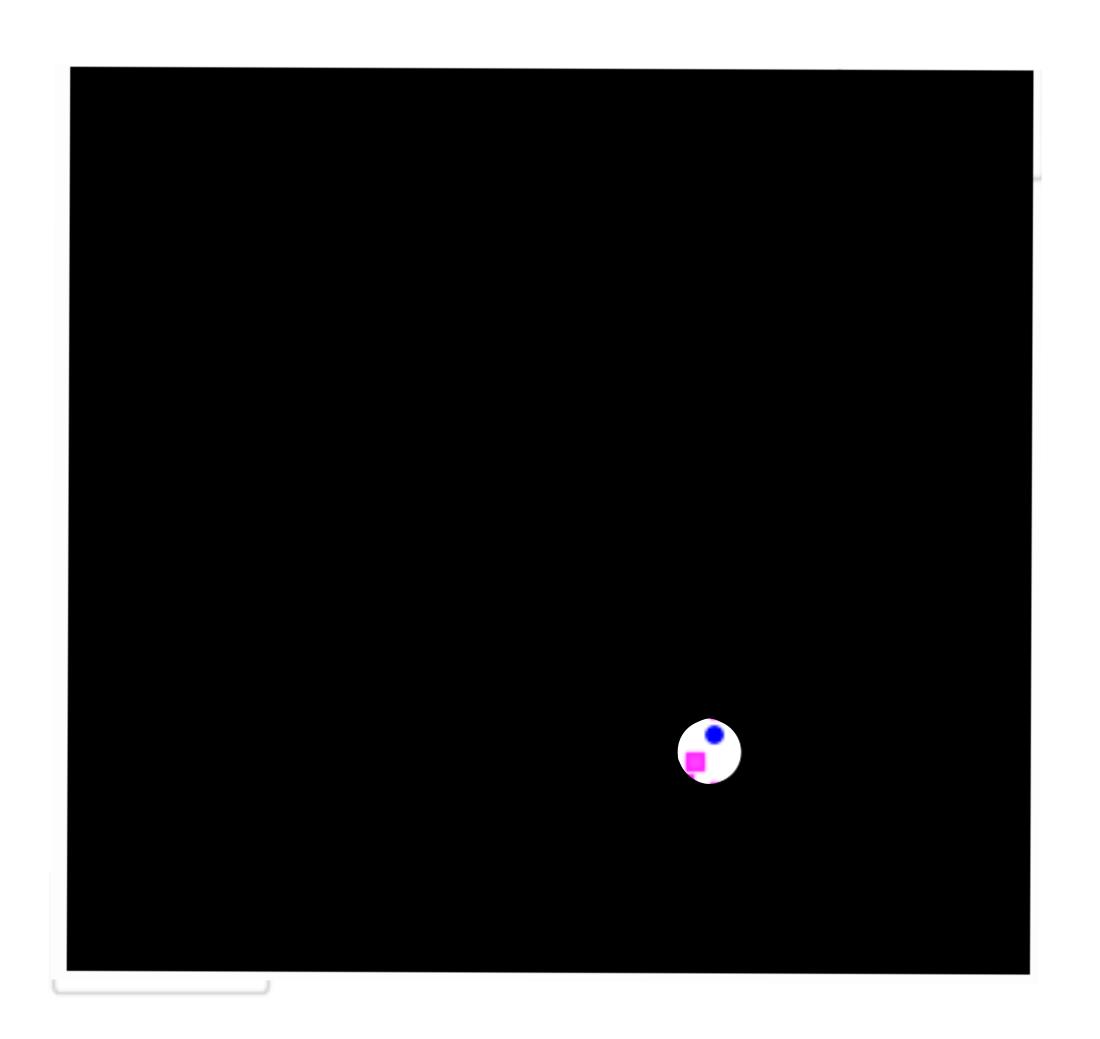










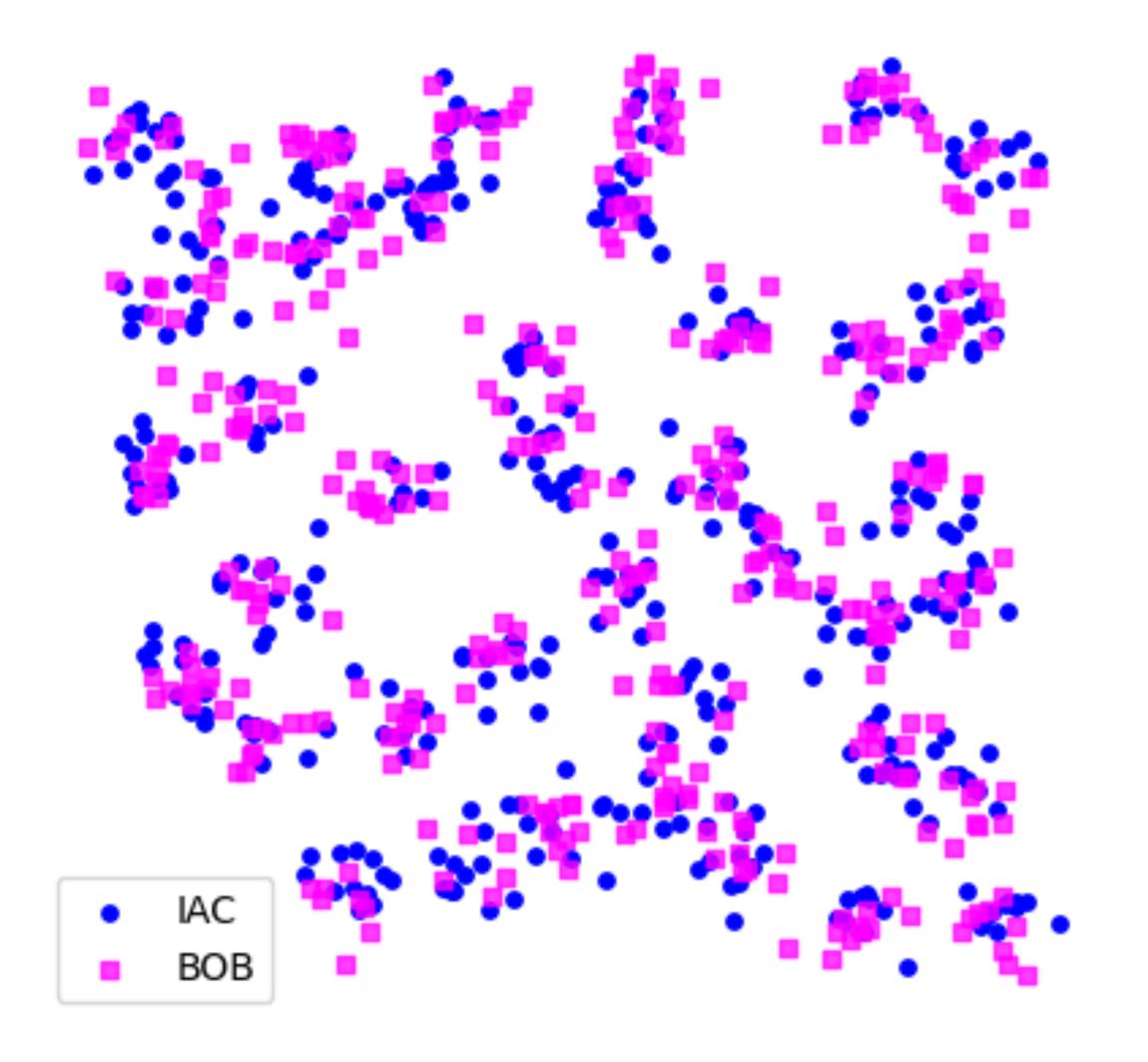










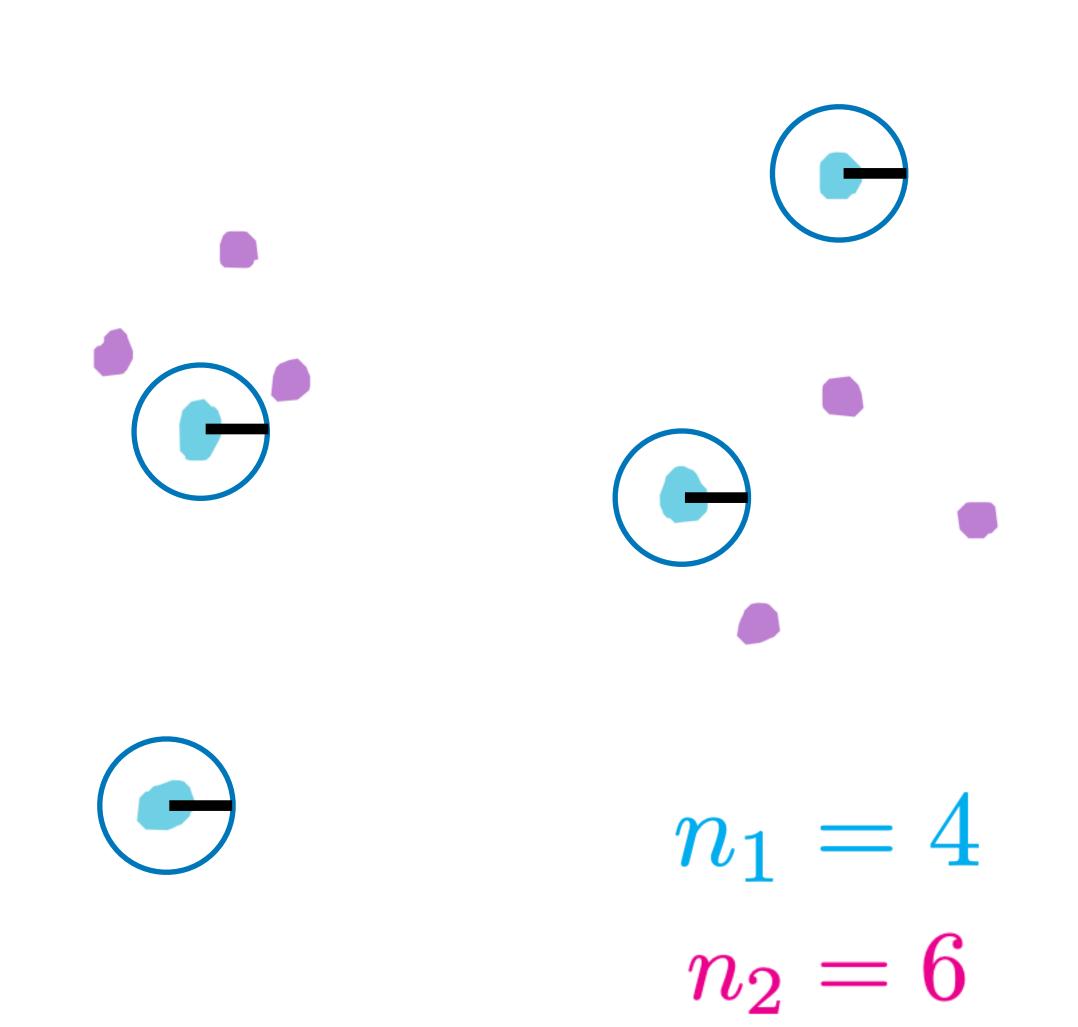




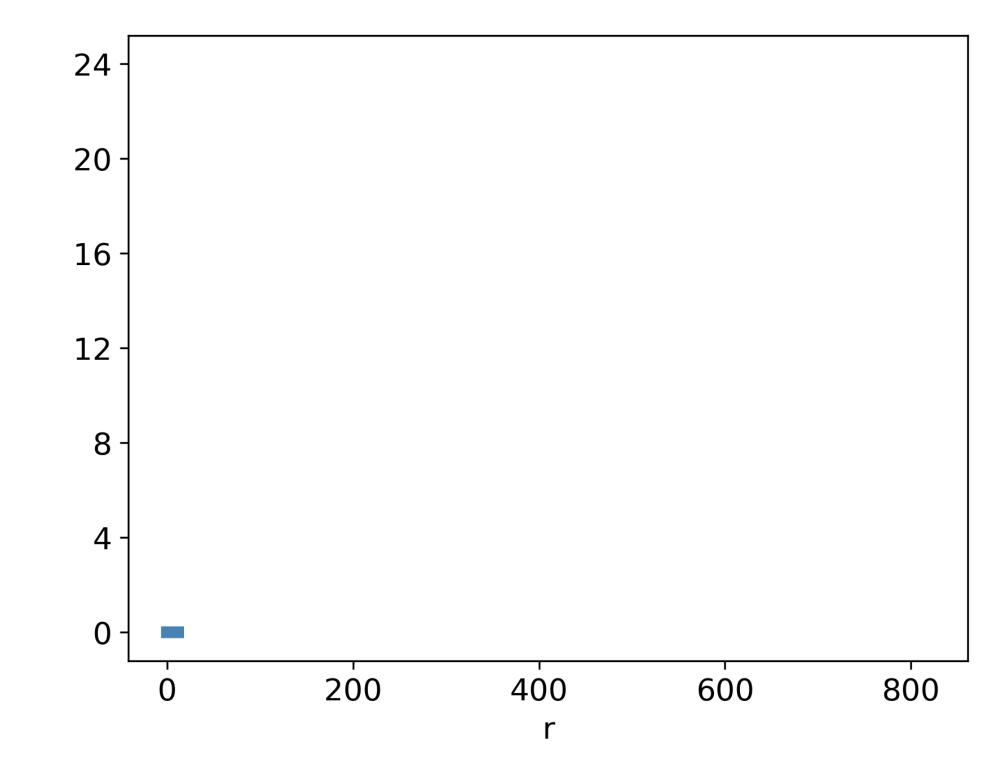








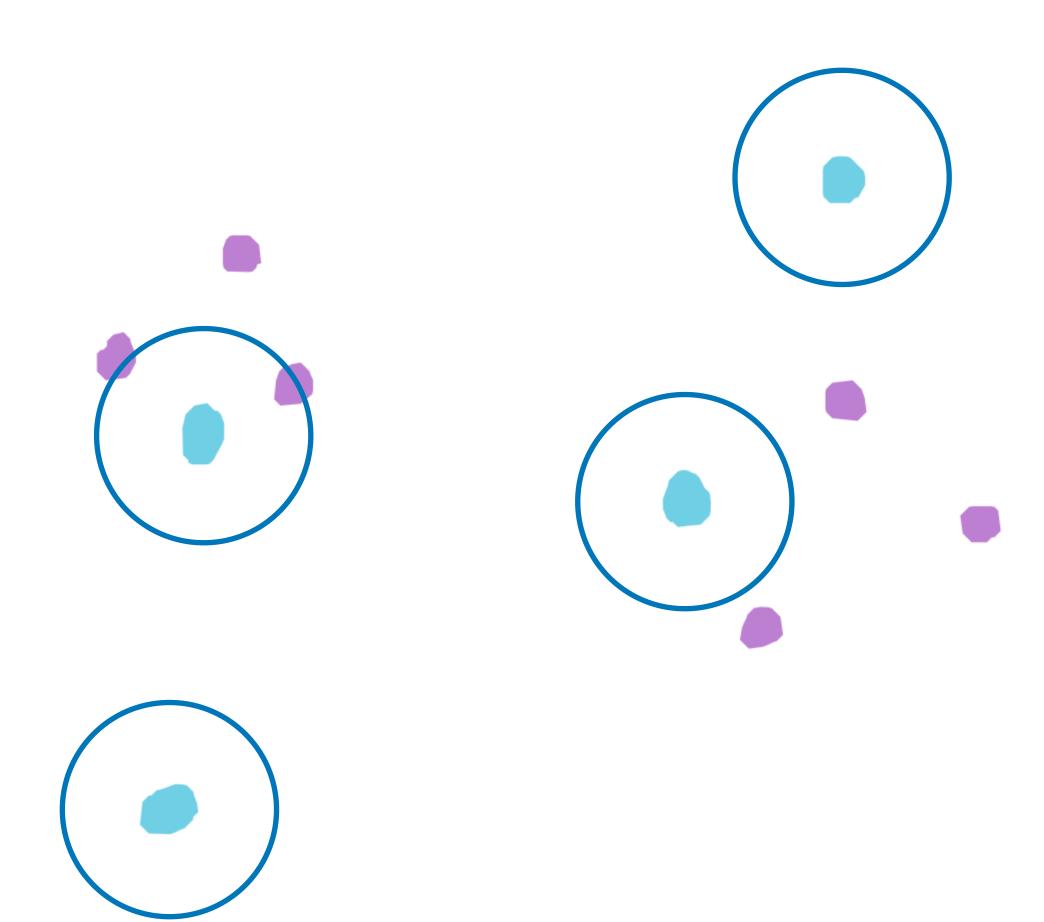
$$S(r) = \frac{|\Omega|}{n_1 n_2} \sum_{i=1}^{n_1} \sum_{j=1}^{n_2} \mathbf{1}(d_{ij} < r)b(i, j, r)$$



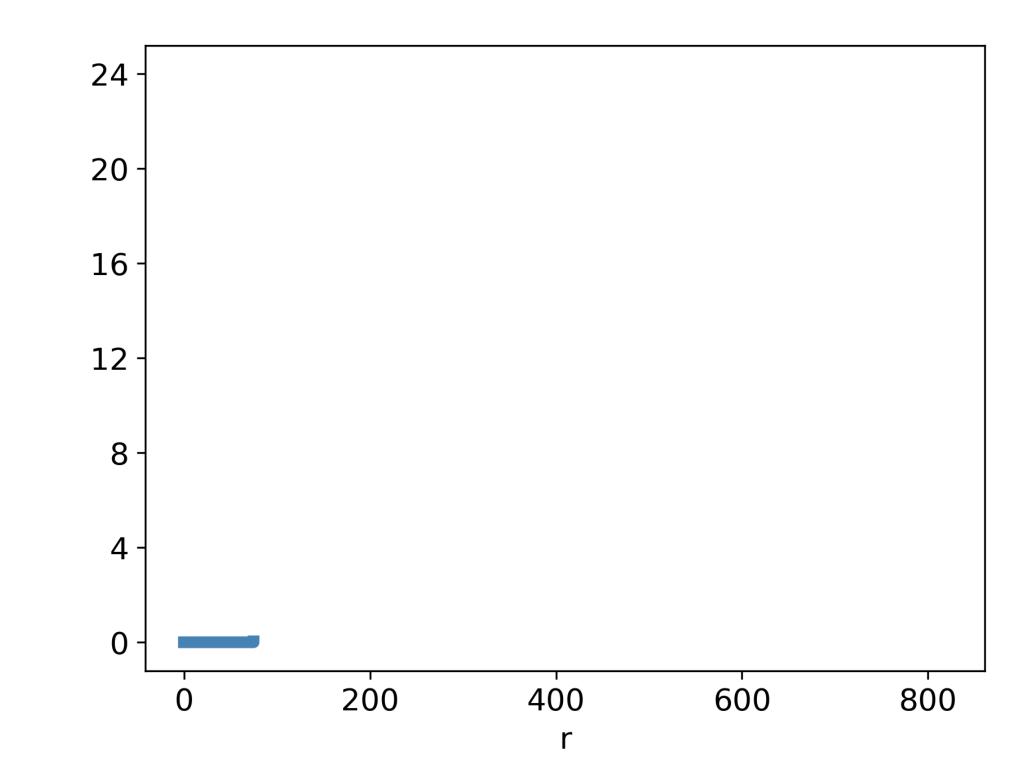








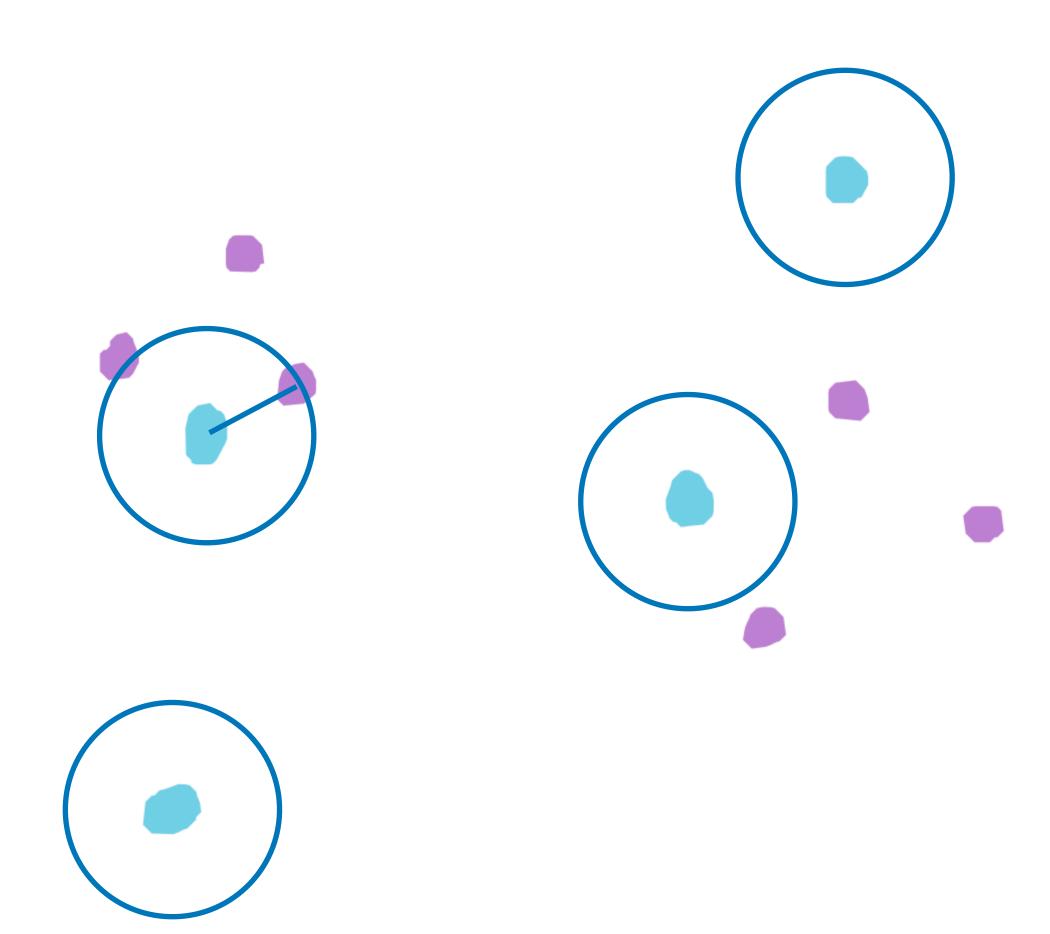
$$S(r) = \frac{|\Omega|}{n_1 n_2} \sum_{i=1}^{n_1} \sum_{j=1}^{n_2} \mathbf{1}(d_{ij} < r)b(i, j, r)$$



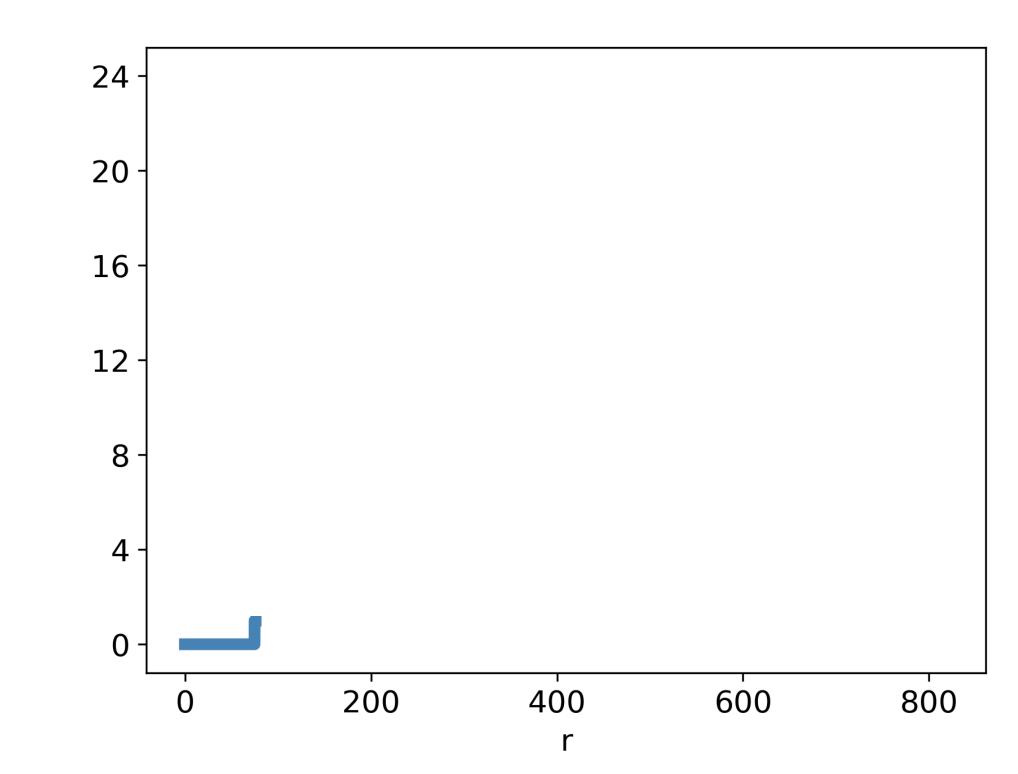








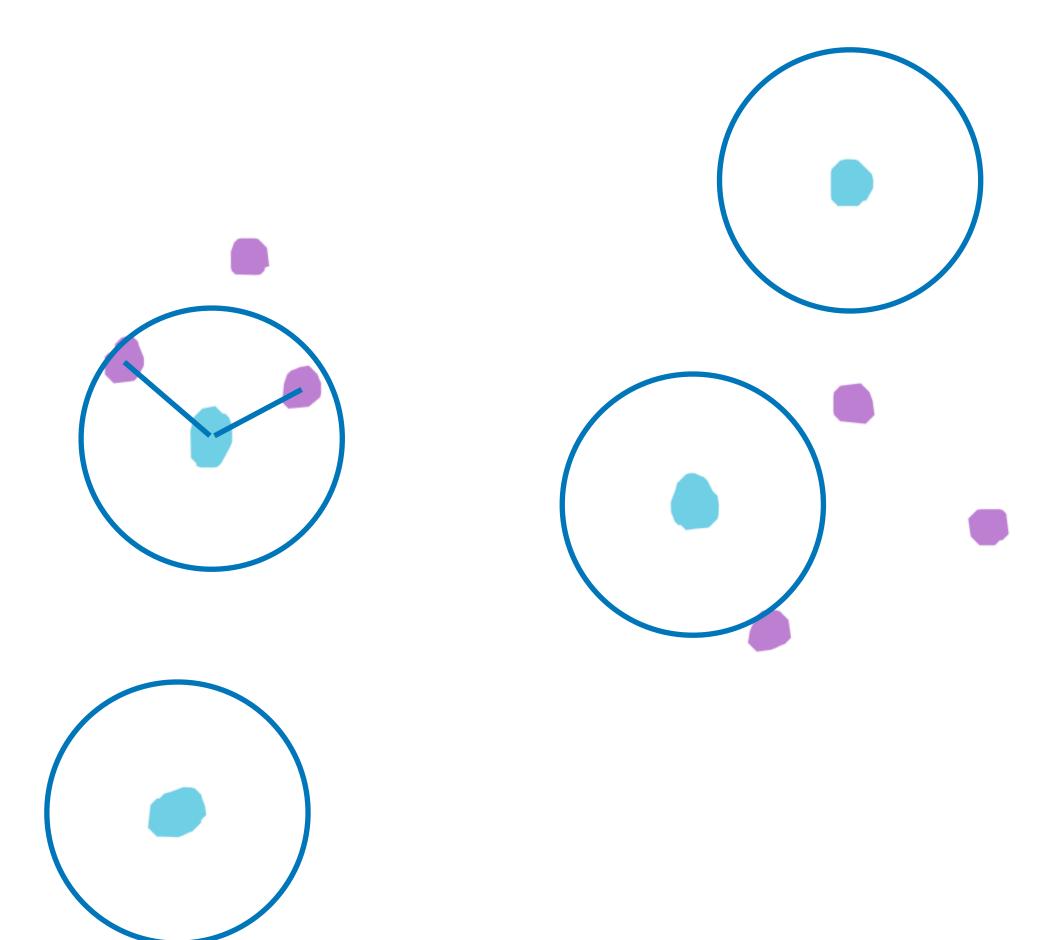
$$S(r) = \frac{|\Omega|}{n_1 n_2} \sum_{i=1}^{n_1} \sum_{j=1}^{n_2} \mathbf{1}(d_{ij} < r)b(i, j, r)$$



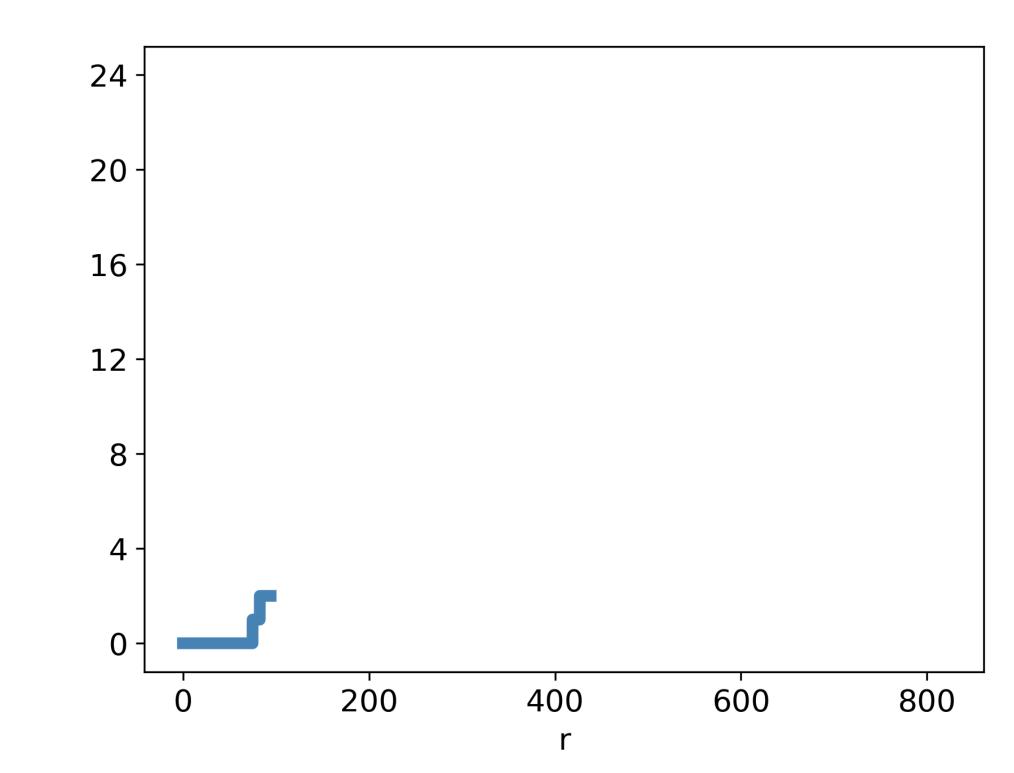








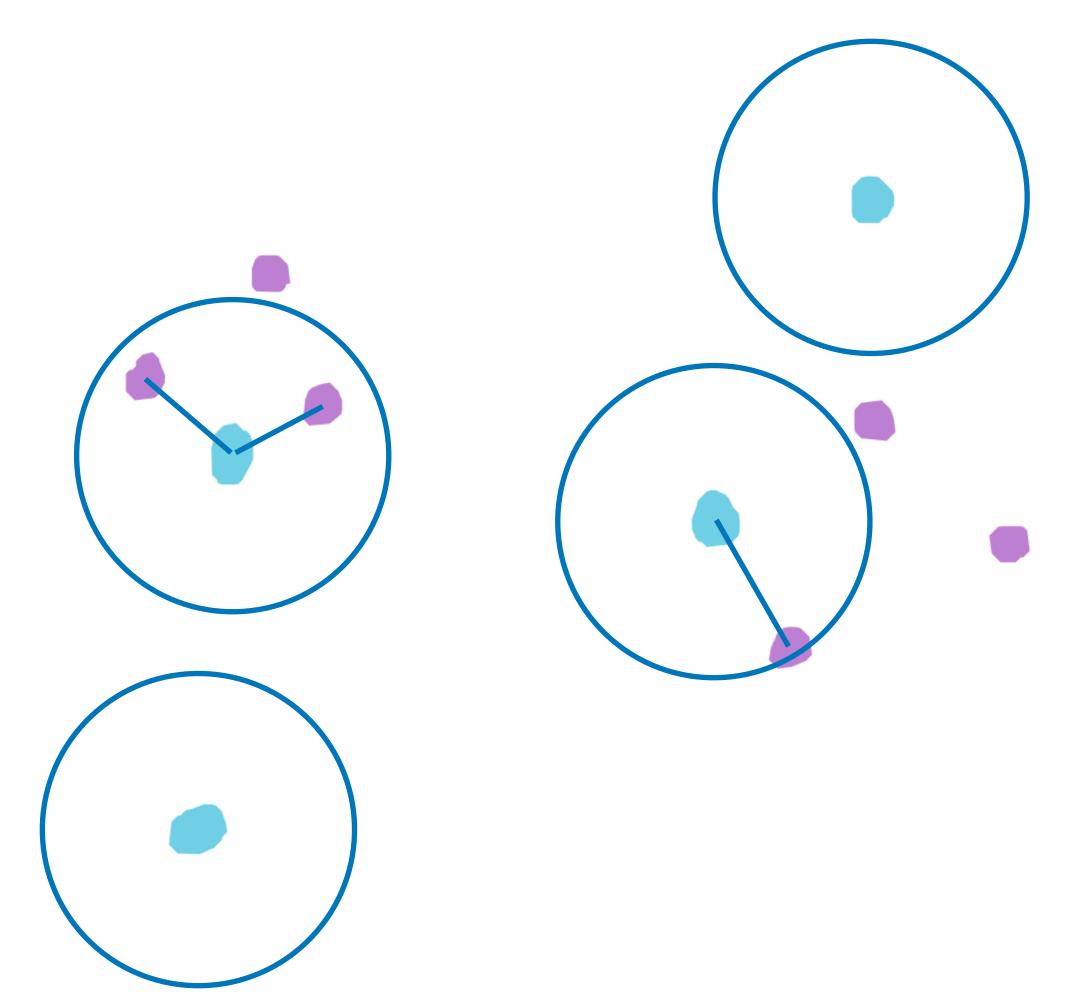
$$S(r) = \frac{|\Omega|}{n_1 n_2} \sum_{i=1}^{n_1} \sum_{j=1}^{n_2} \mathbf{1}(d_{ij} < r)b(i, j, r)$$



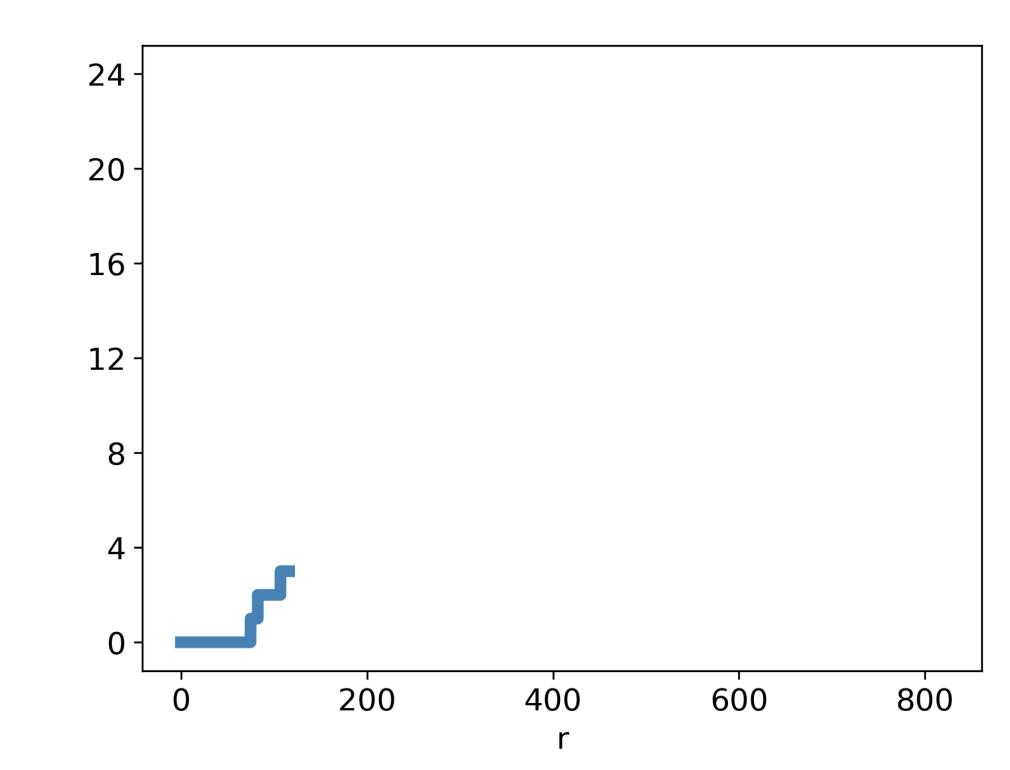








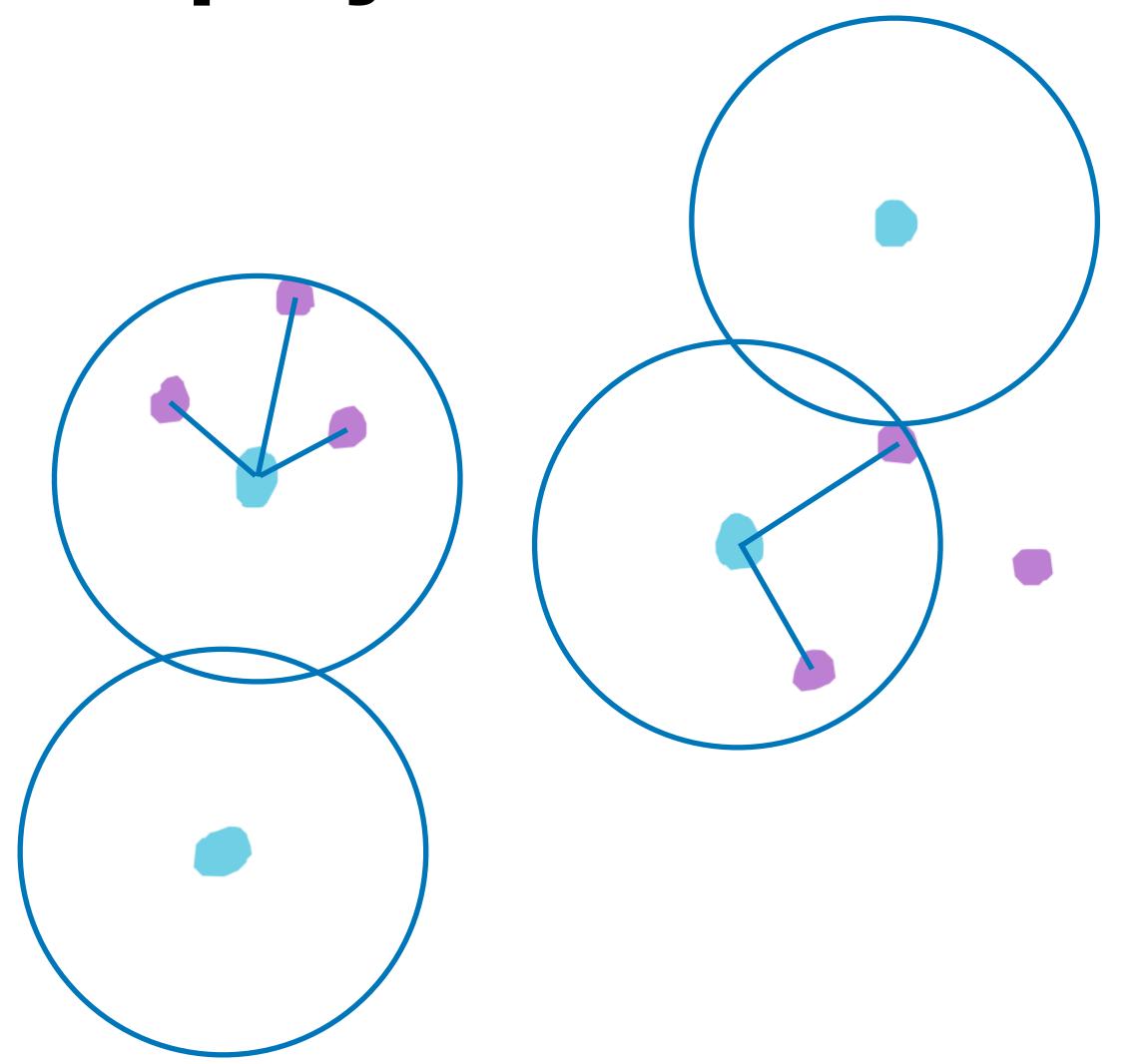
$$S(r) = \frac{|\Omega|}{n_1 n_2} \sum_{i=1}^{n_1} \sum_{j=1}^{n_2} \mathbf{1}(d_{ij} < r)b(i, j, r)$$



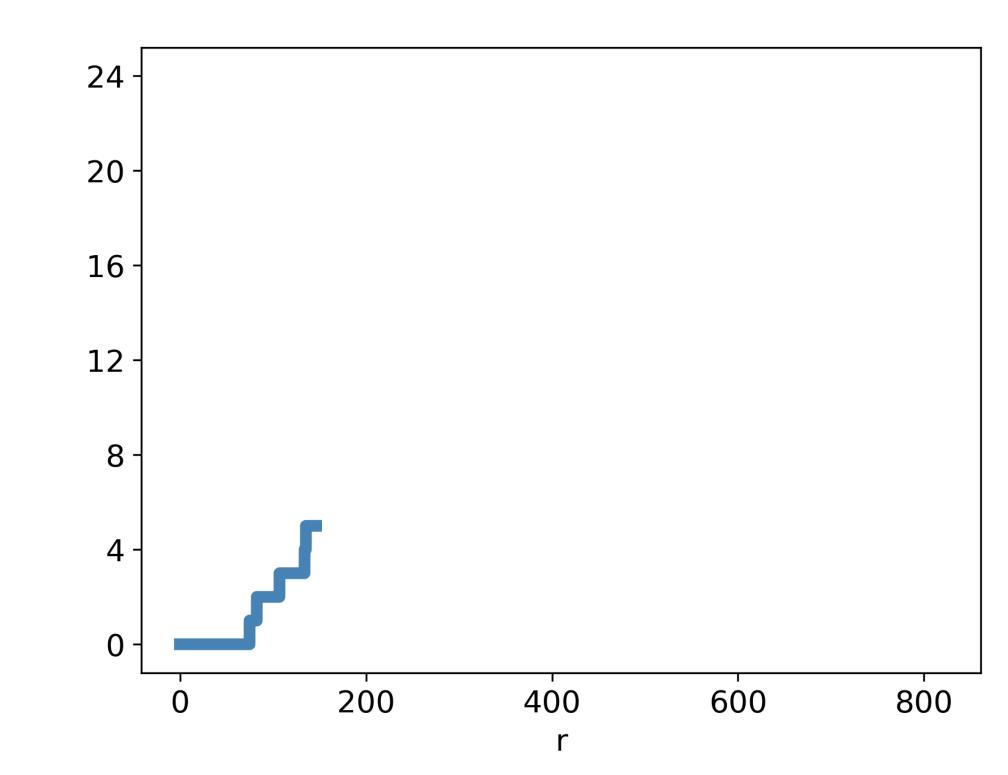








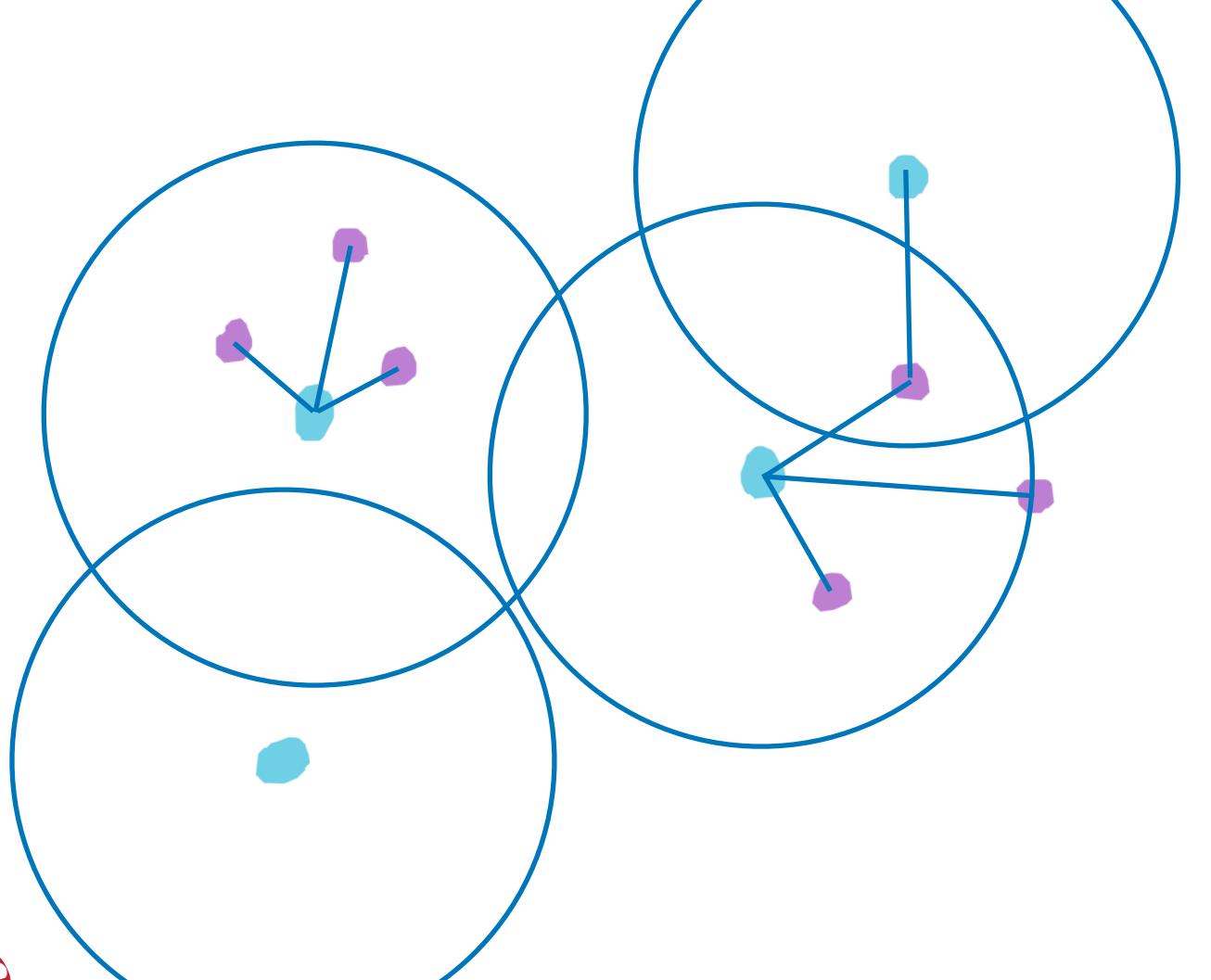
$$S(r) = \frac{|\Omega|}{n_1 n_2} \sum_{i=1}^{n_1} \sum_{j=1}^{n_2} \mathbf{1}(d_{ij} < r)b(i, j, r)$$



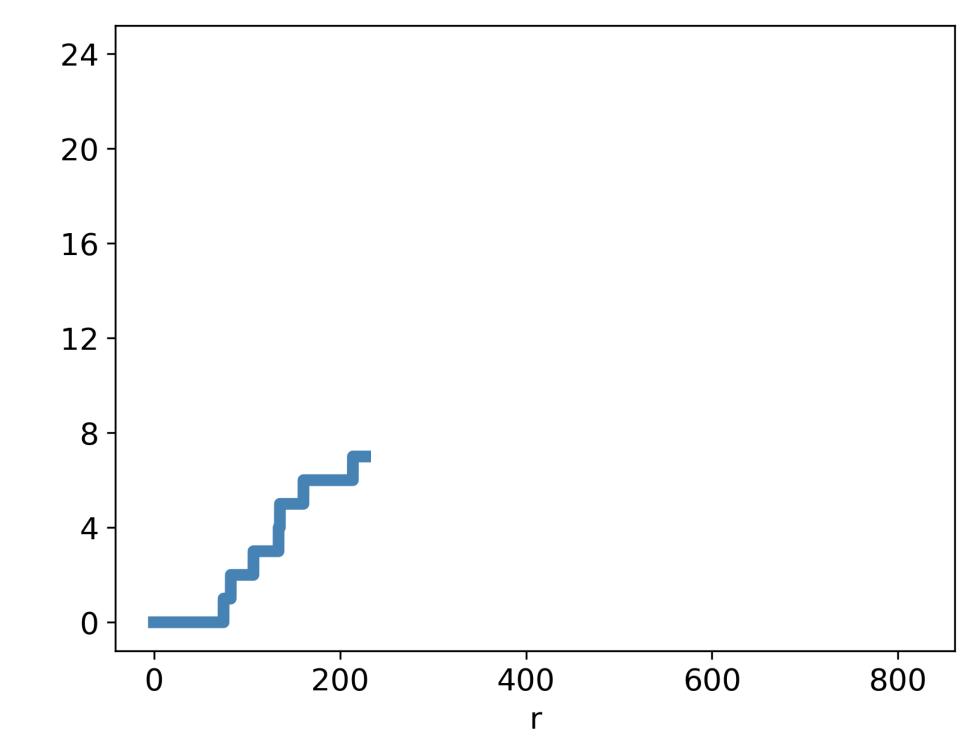






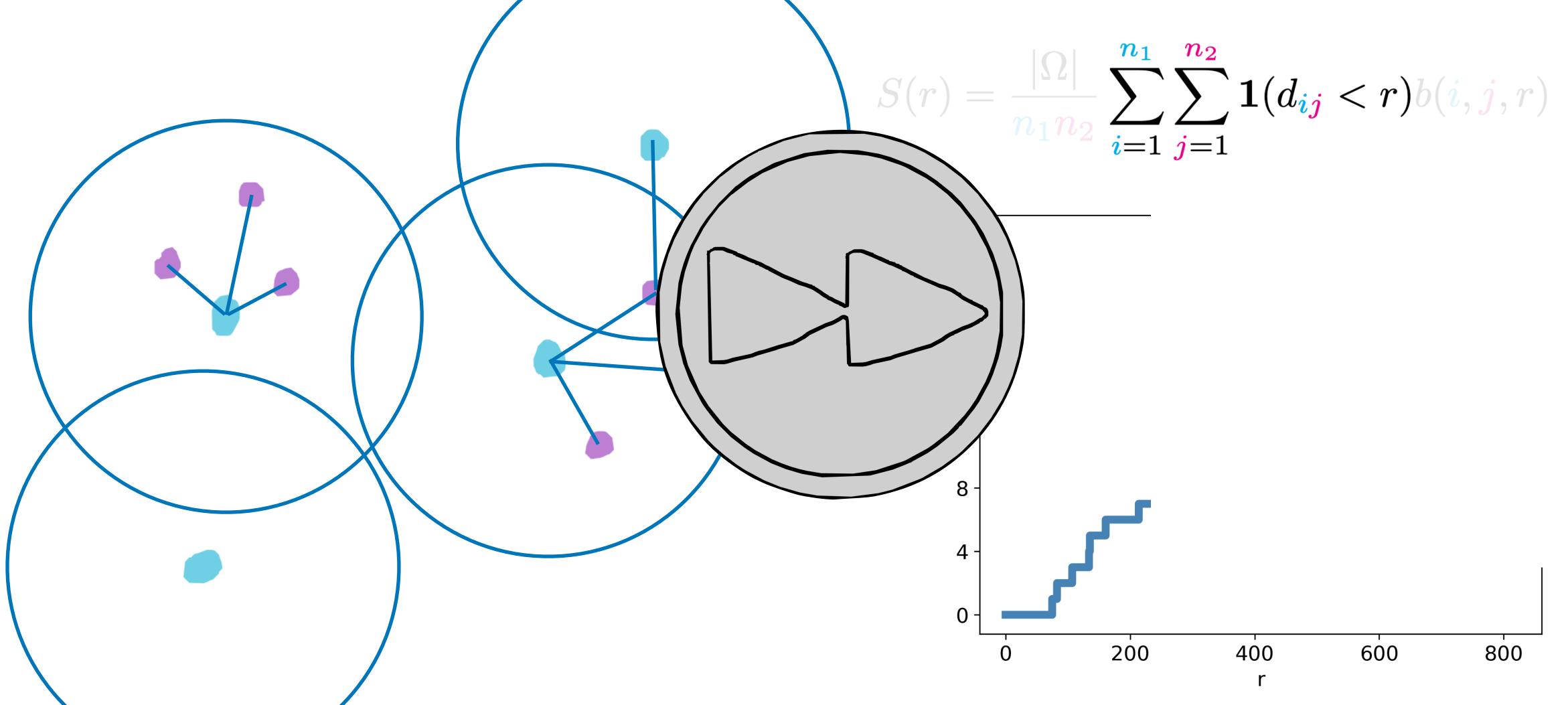


$$S(r) = \frac{|\Omega|}{n_1 n_2} \sum_{i=1}^{n_1} \sum_{j=1}^{n_2} \mathbf{1}(d_{ij} < r)b(i, j, r)$$





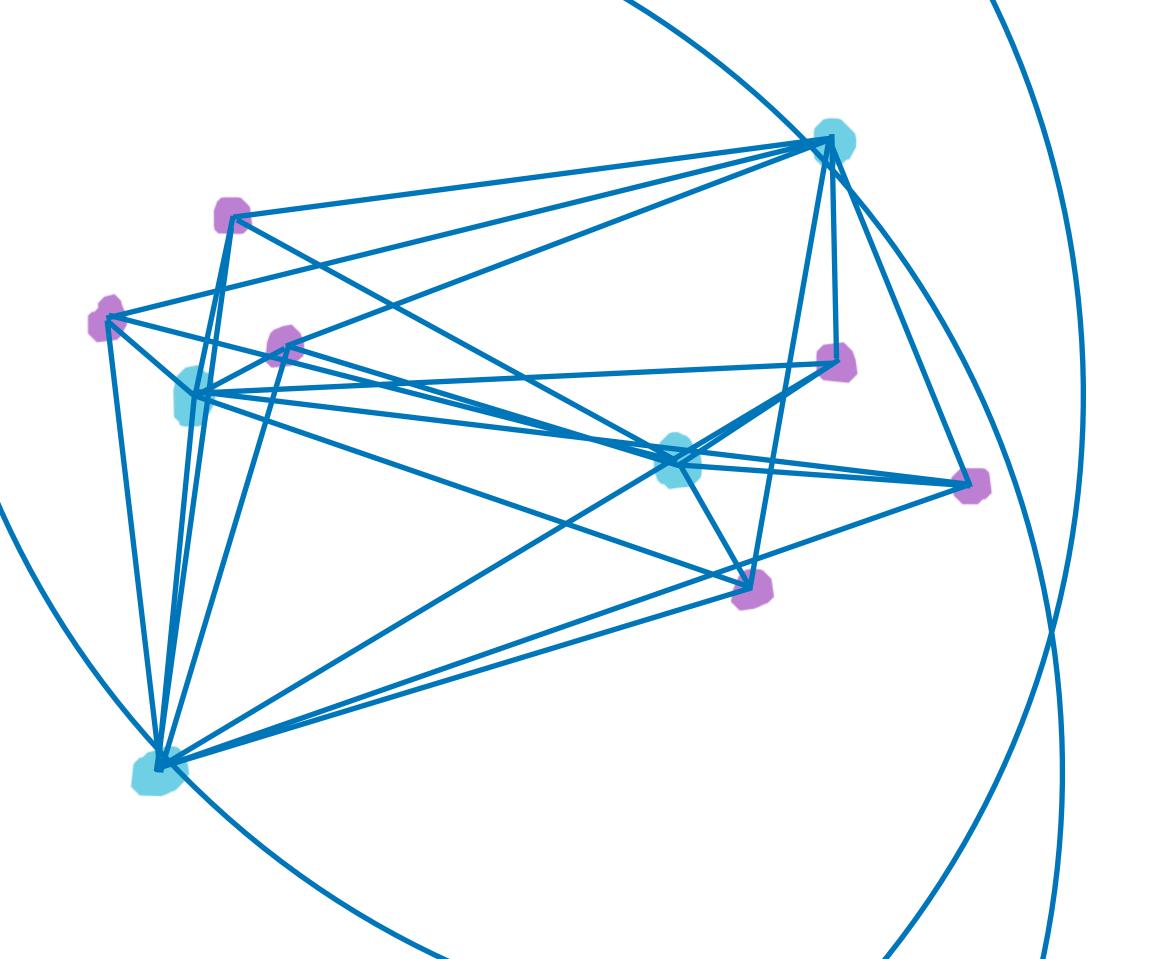


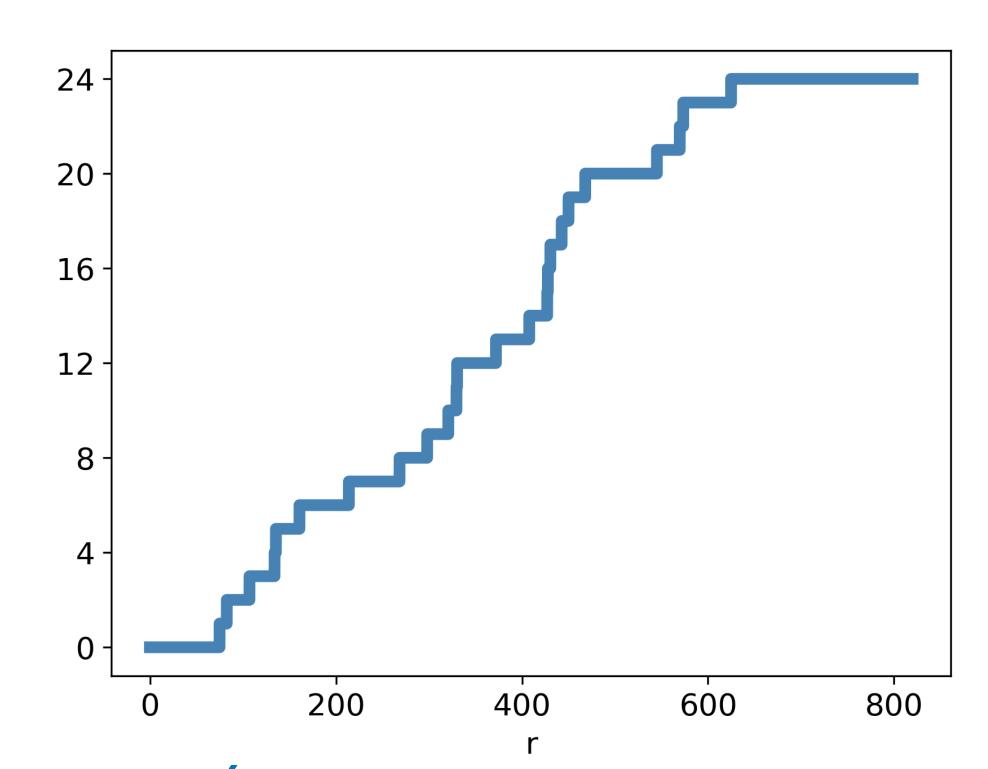








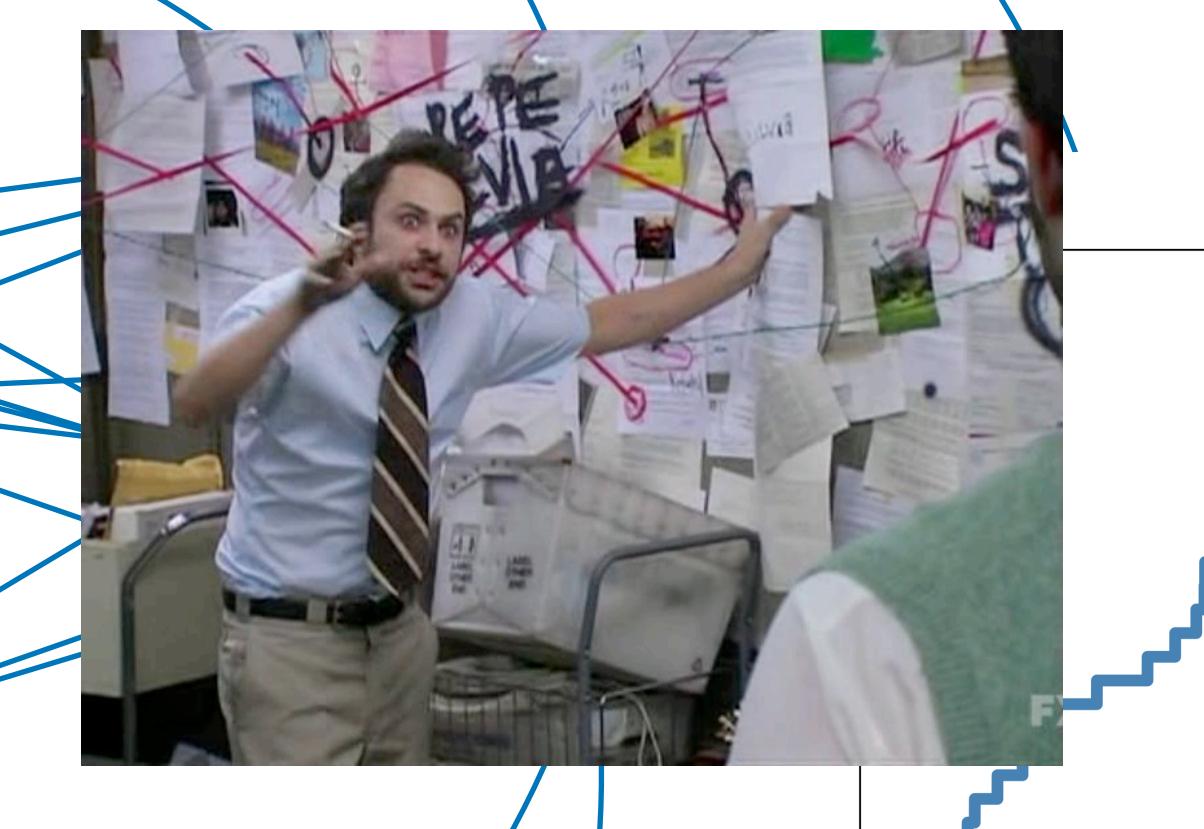








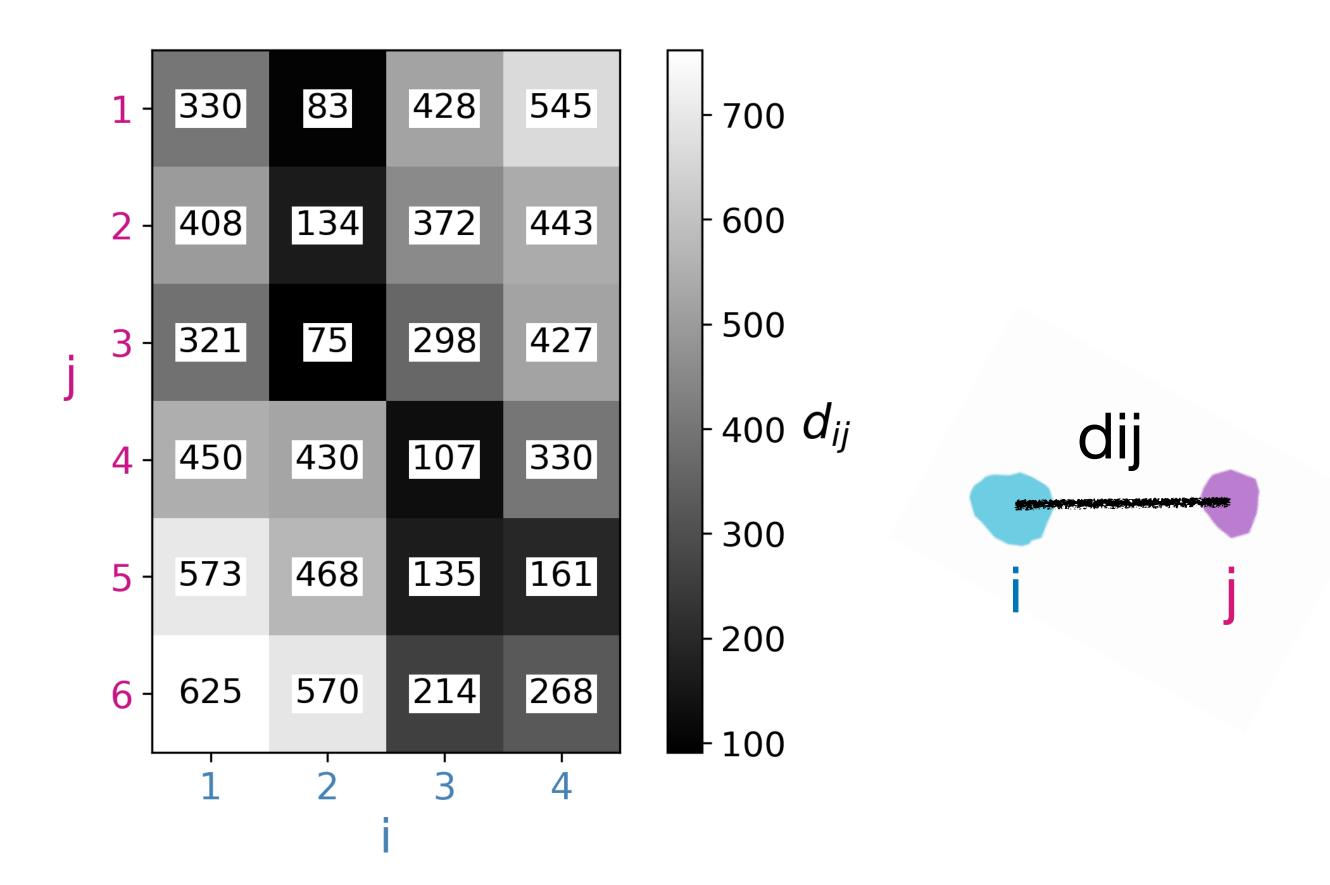




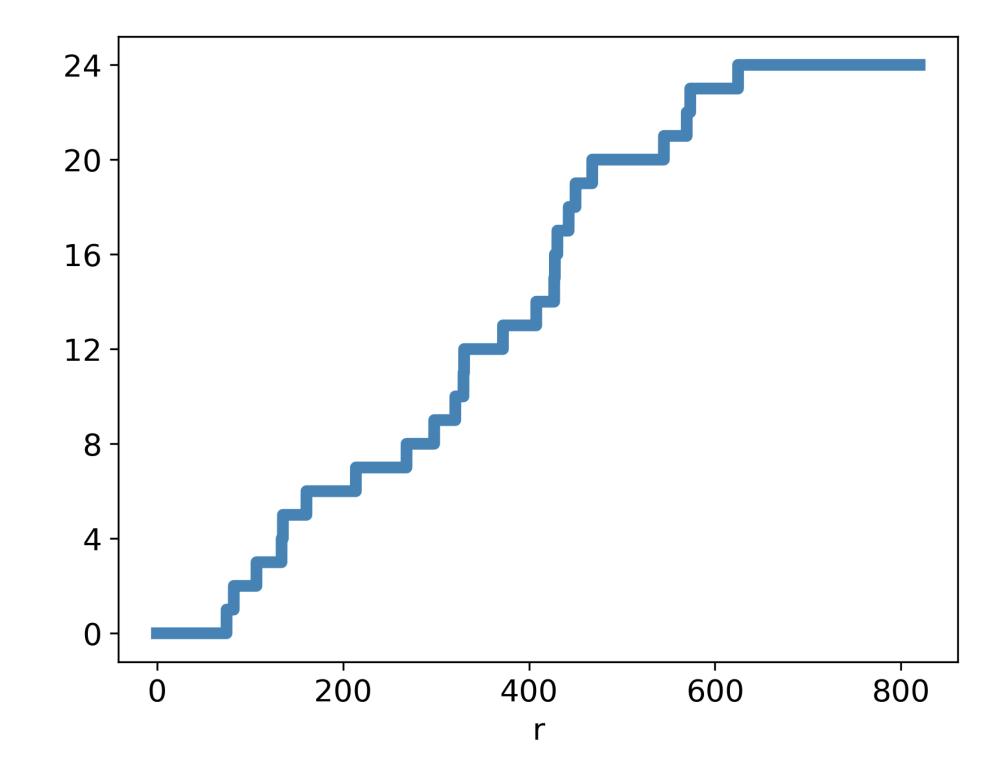








$$S(r) = \frac{|\Omega|}{n_1 n_2} \sum_{i=1}^{n_1} \sum_{j=1}^{n_2} \mathbf{1}(d_{ij} < r)b(i, j, r)$$



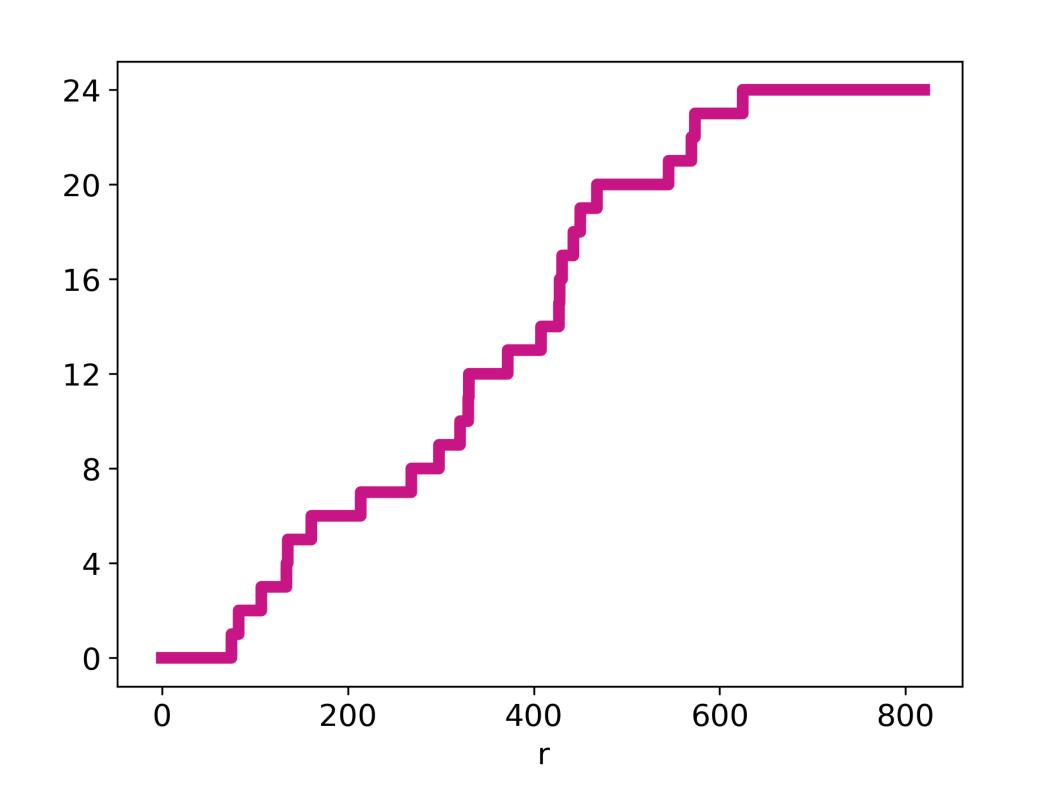


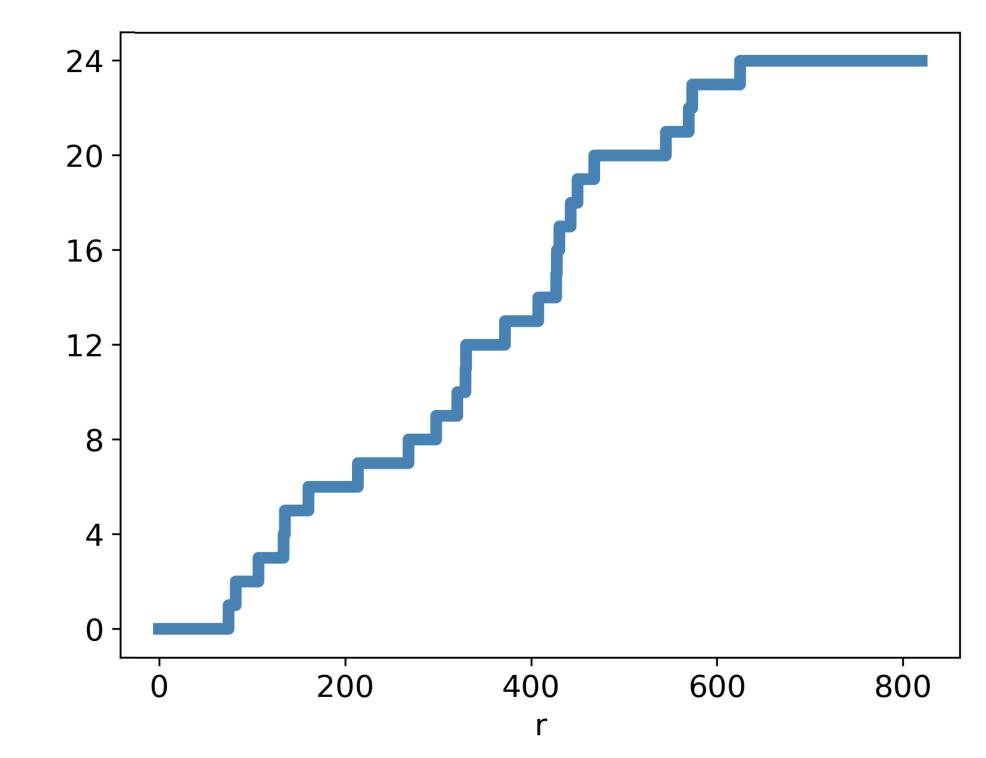




$$S(r) = \frac{|\Omega|}{n_1 n_2} \sum_{i=1}^{n_1} \sum_{j=1}^{n_2} \mathbf{1}(d_{ij} < r)b(i, j, r) \qquad S(r) = \frac{|\Omega|}{n_1 n_2} \sum_{i=1}^{n_1} \sum_{j=1}^{n_2} \mathbf{1}(d_{ij} < r)b(i, j, r)$$

$$S(r) = \frac{|\Omega|}{n_1 n_2} \sum_{i=1}^{n_1} \sum_{j=1}^{n_2} \mathbf{1}(d_{ij} < r)b(i, j, r)$$

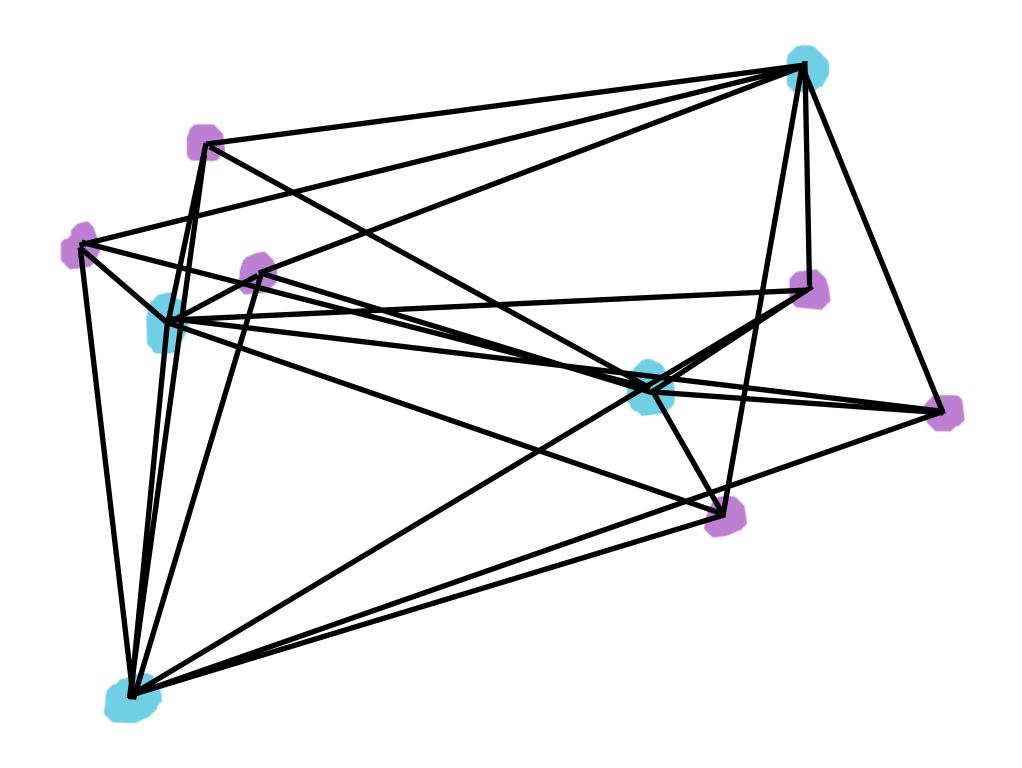




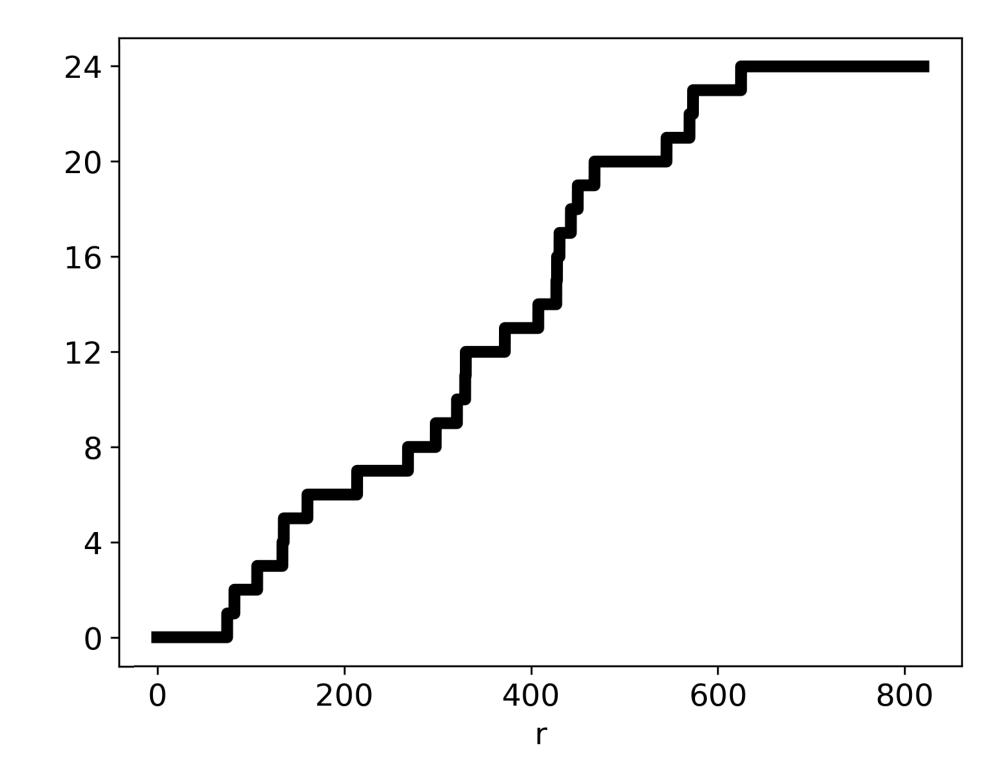








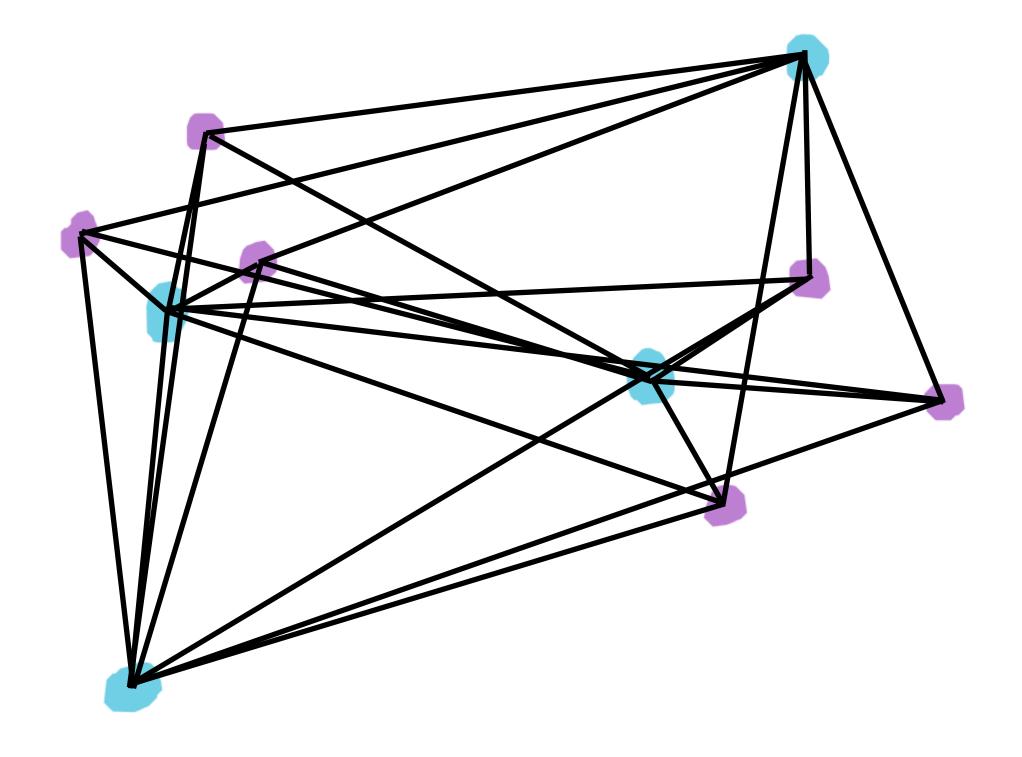
$$S(r) = rac{|\Omega|}{n_1 n_2} \sum_{i=1}^{n_1} \sum_{j=1}^{n_2} \mathbf{1}(d_{ij} < r)b(i, j, r)$$



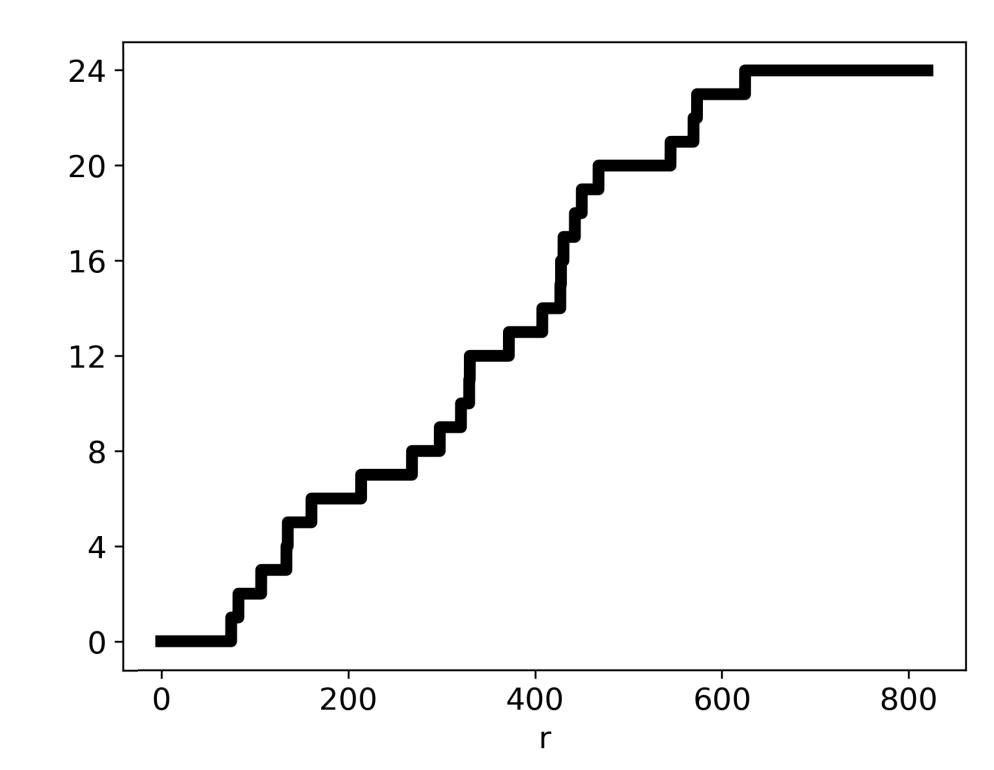








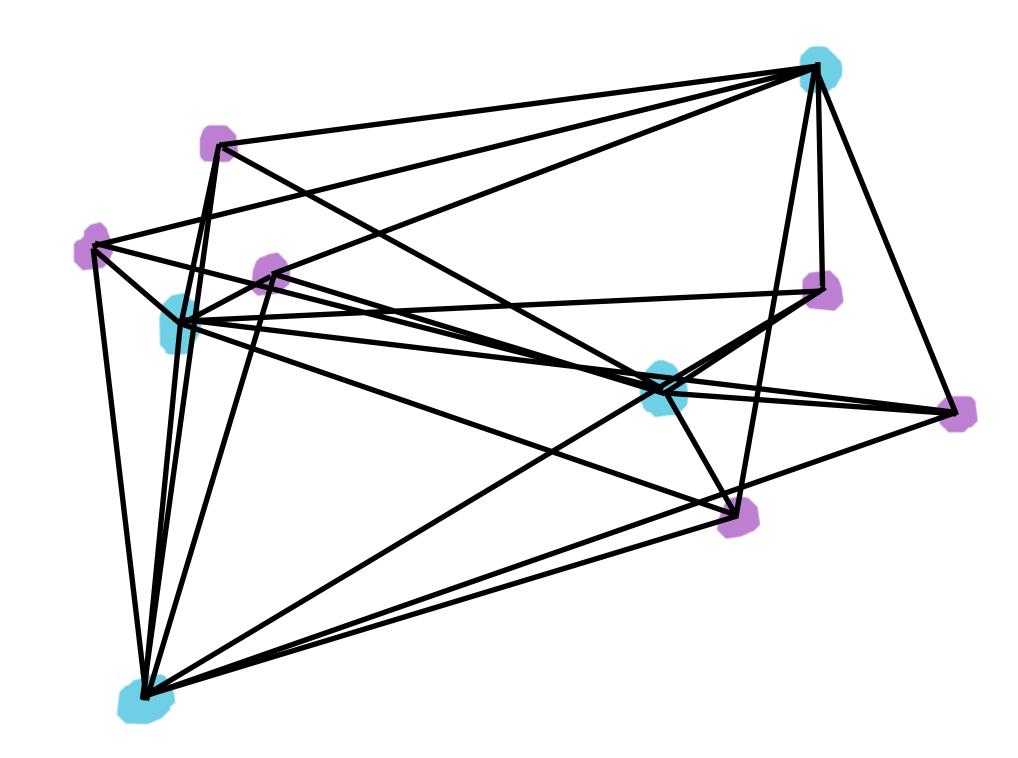
$$S(r) = rac{|\Omega|}{n_1 n_2} \sum_{i=1}^{n_1} \sum_{j=1}^{n_2} \mathbf{1}(d_{ij} < r)b(i, j, r)$$



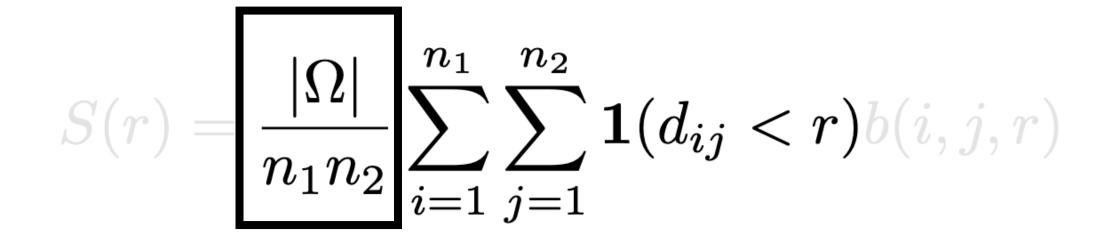


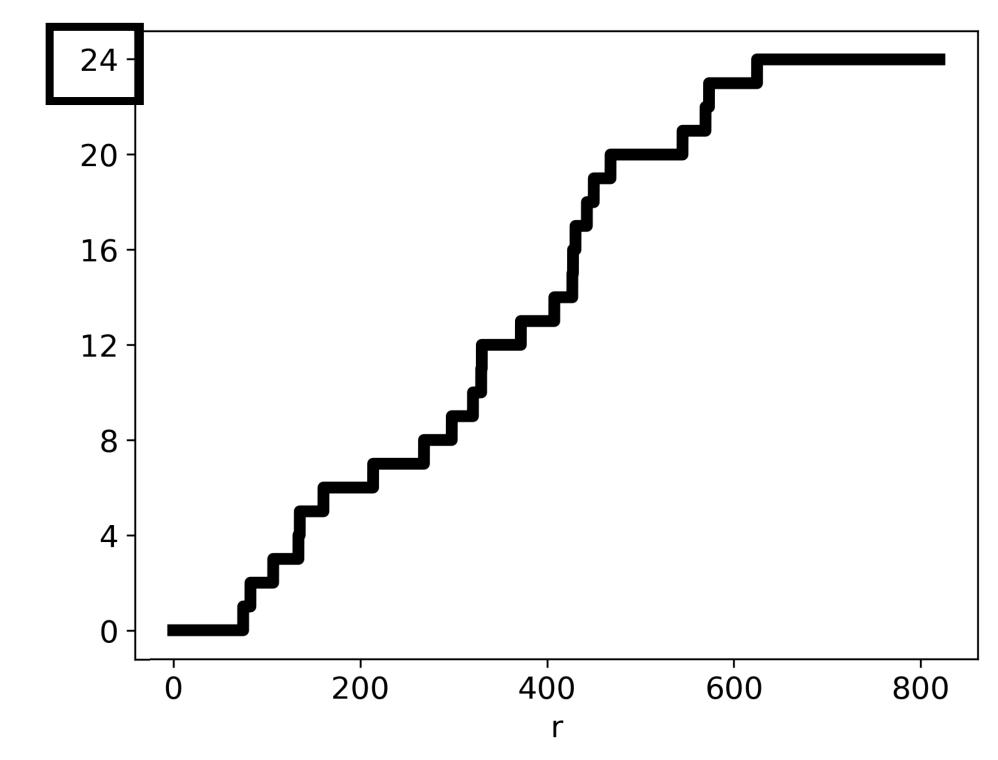






$$n_1 n_2 = n \text{ connections} = 24$$

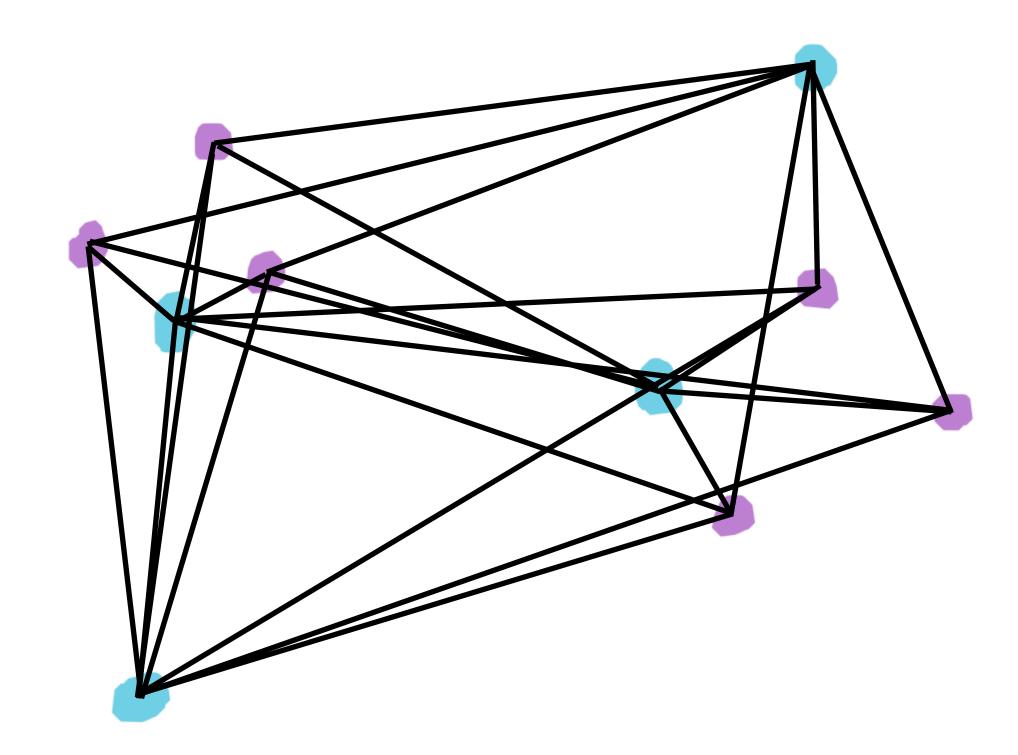




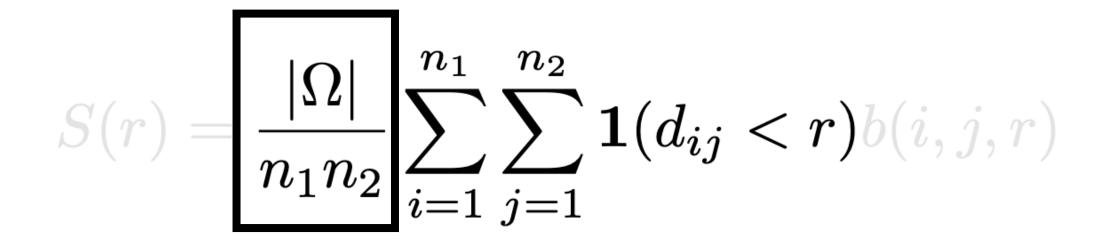


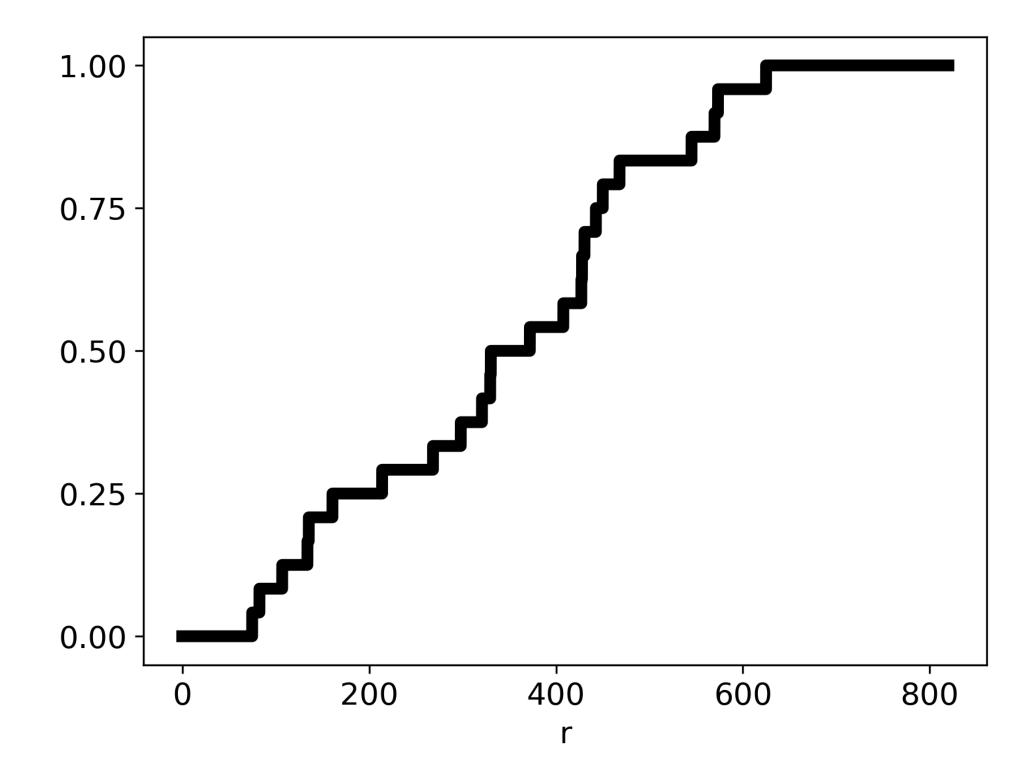




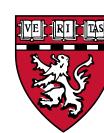


$$n_1 n_2 = n \text{ connections} = 24$$

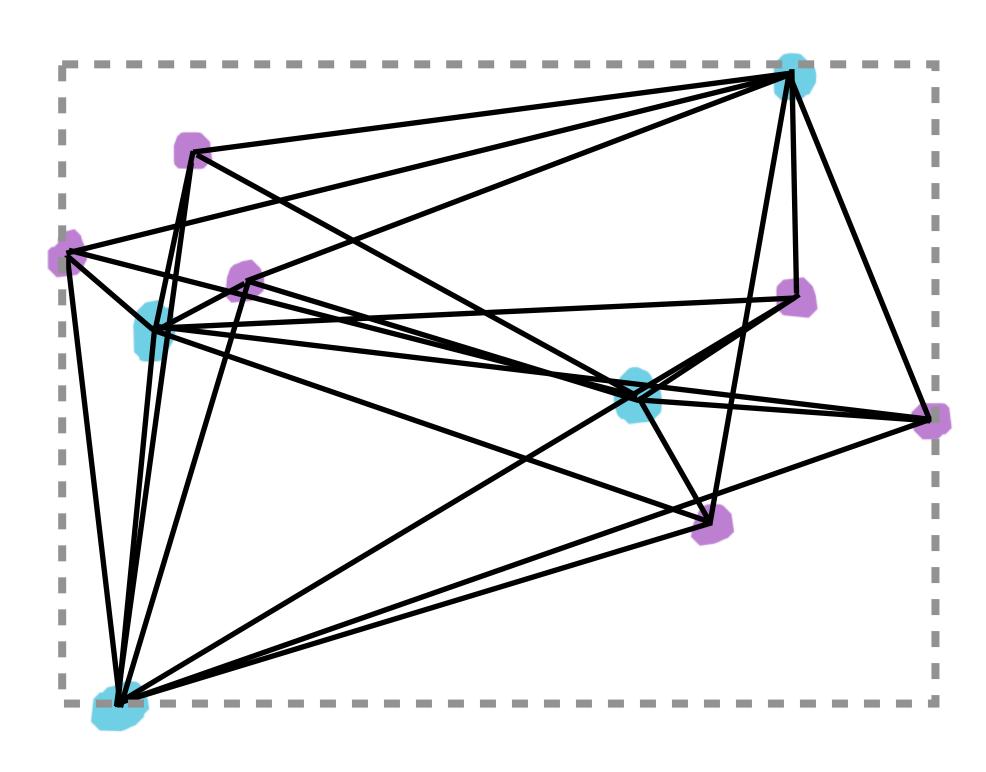




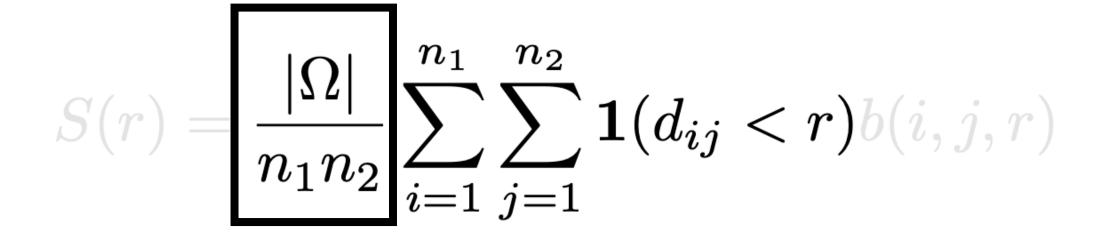


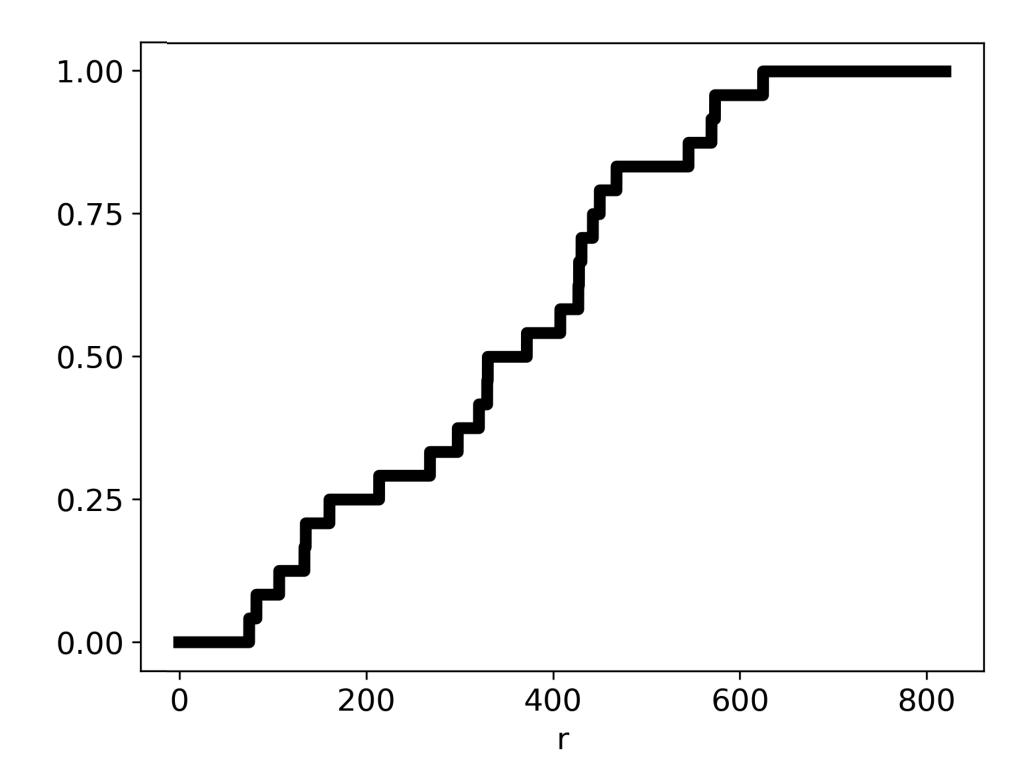






$$|\Omega|$$
 = Area of FOV

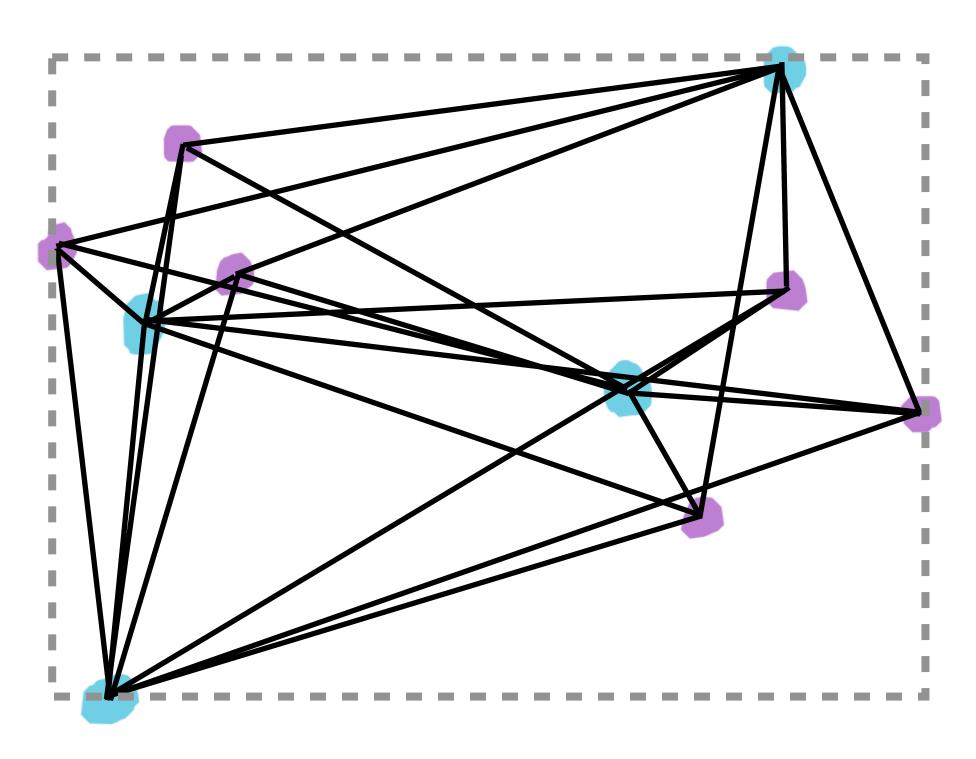






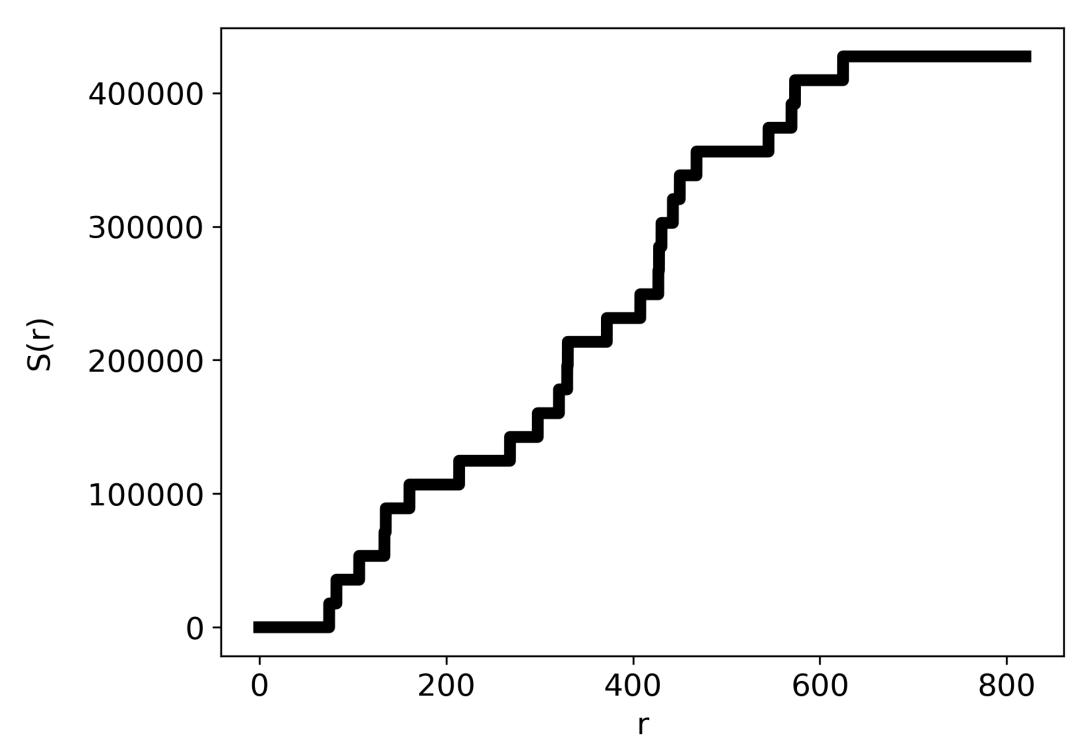






$$|\Omega|$$
 = Area of FOV

$$S(r) = \frac{|\Omega|}{n_1 n_2} \sum_{i=1}^{n_1} \sum_{j=1}^{n_2} \mathbf{1}(d_{ij} < r)b(i, j, r)$$





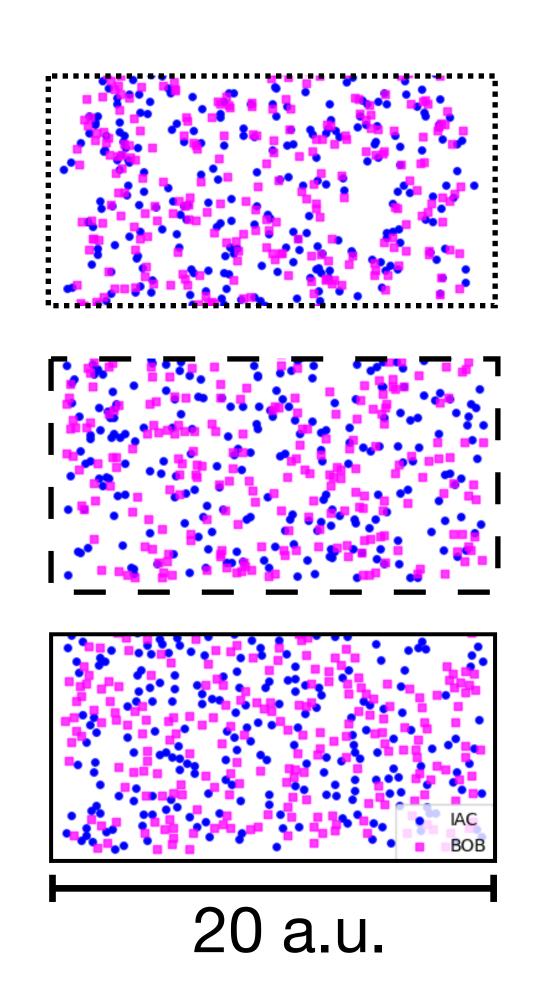


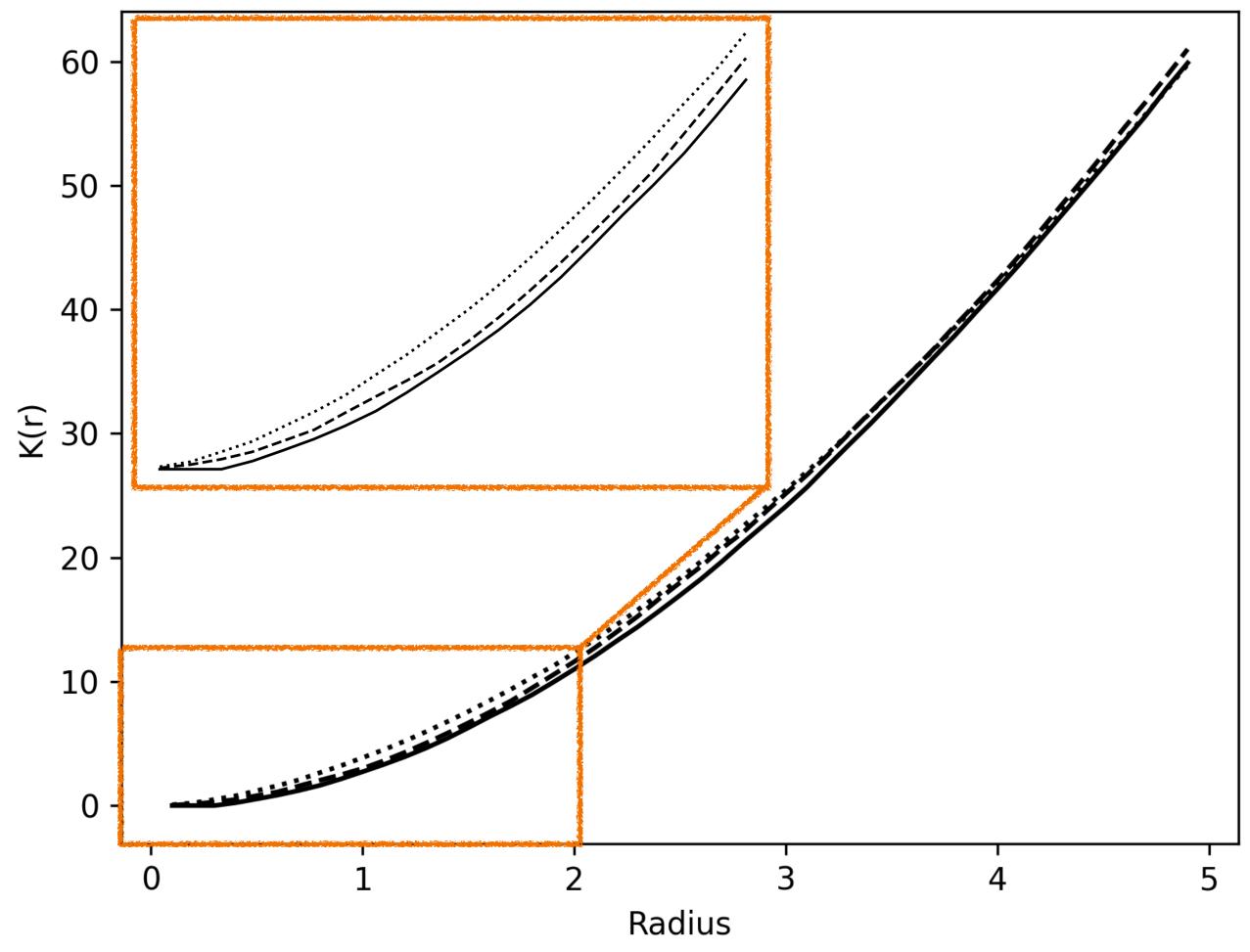


Fall — cold

Fall — medium

Fall — warm

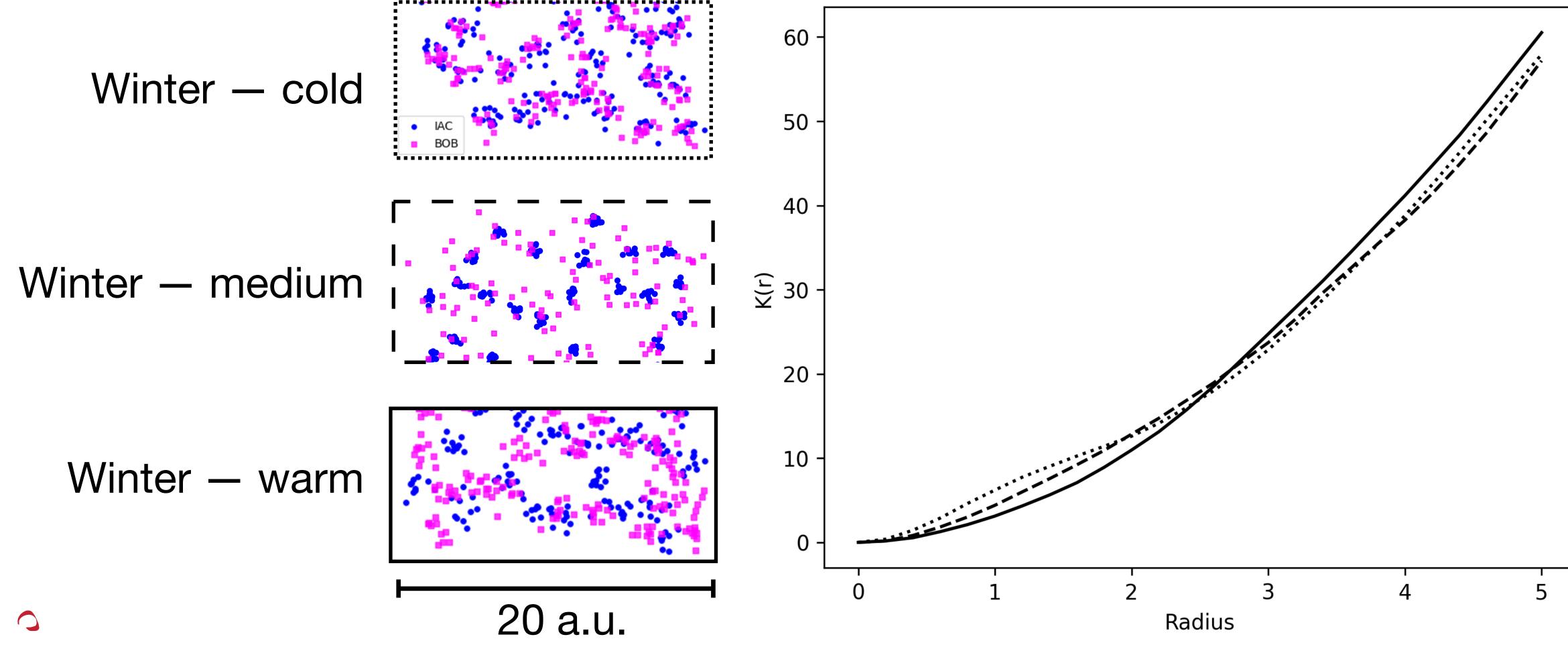








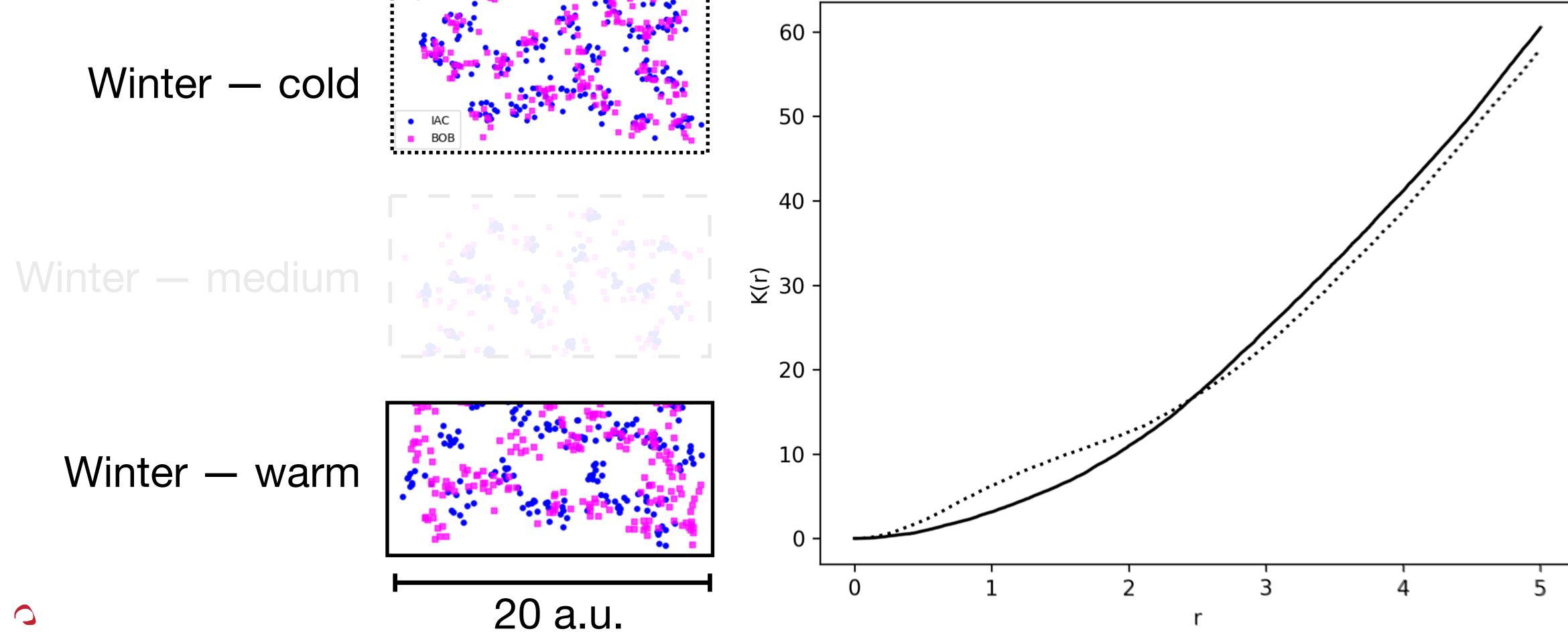








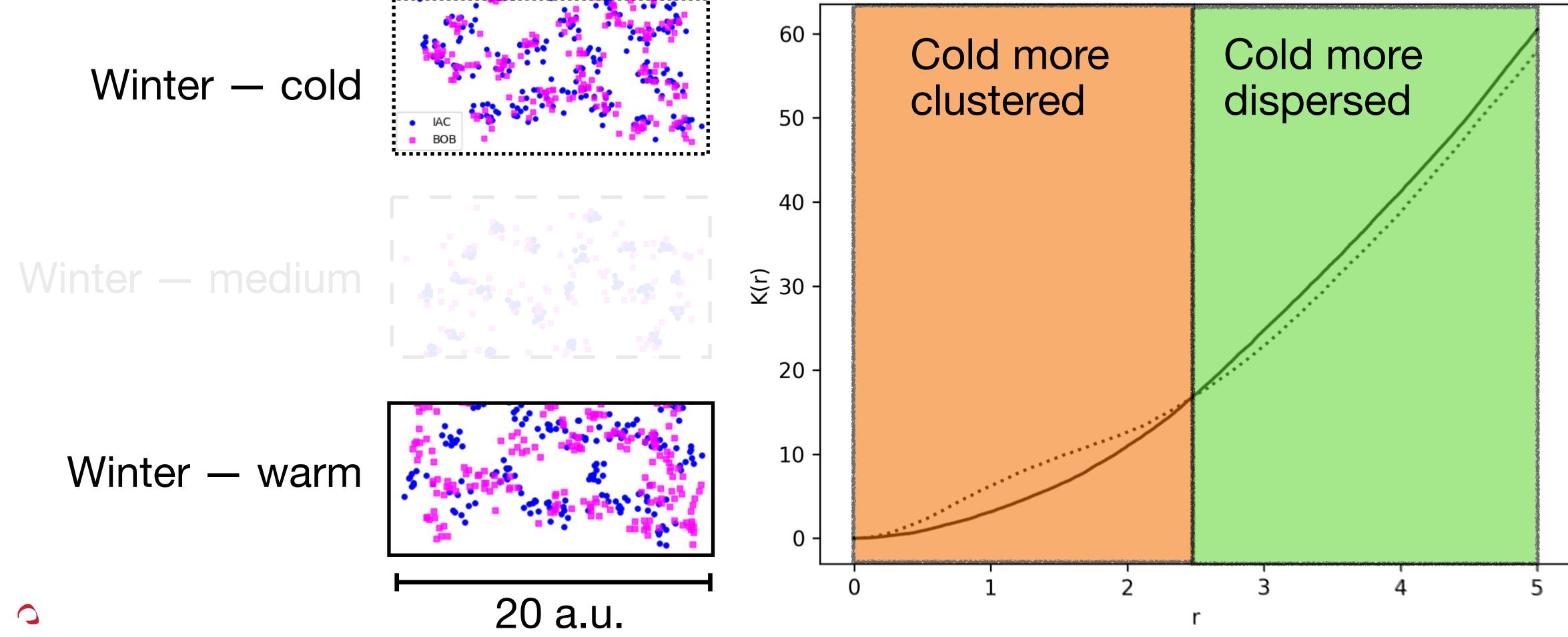














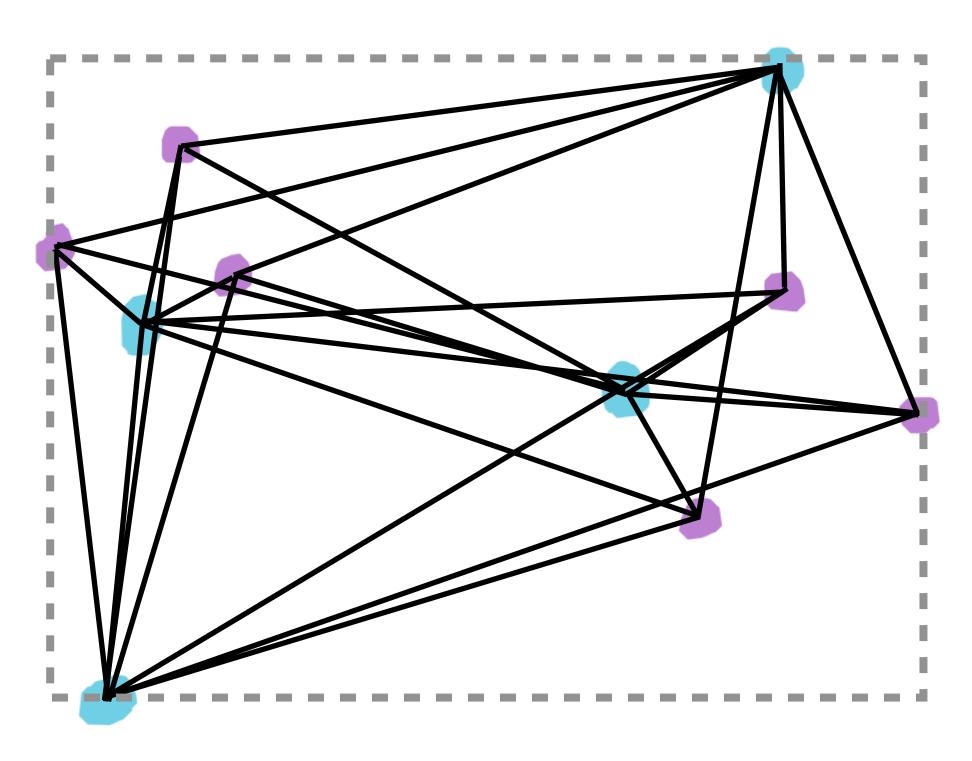






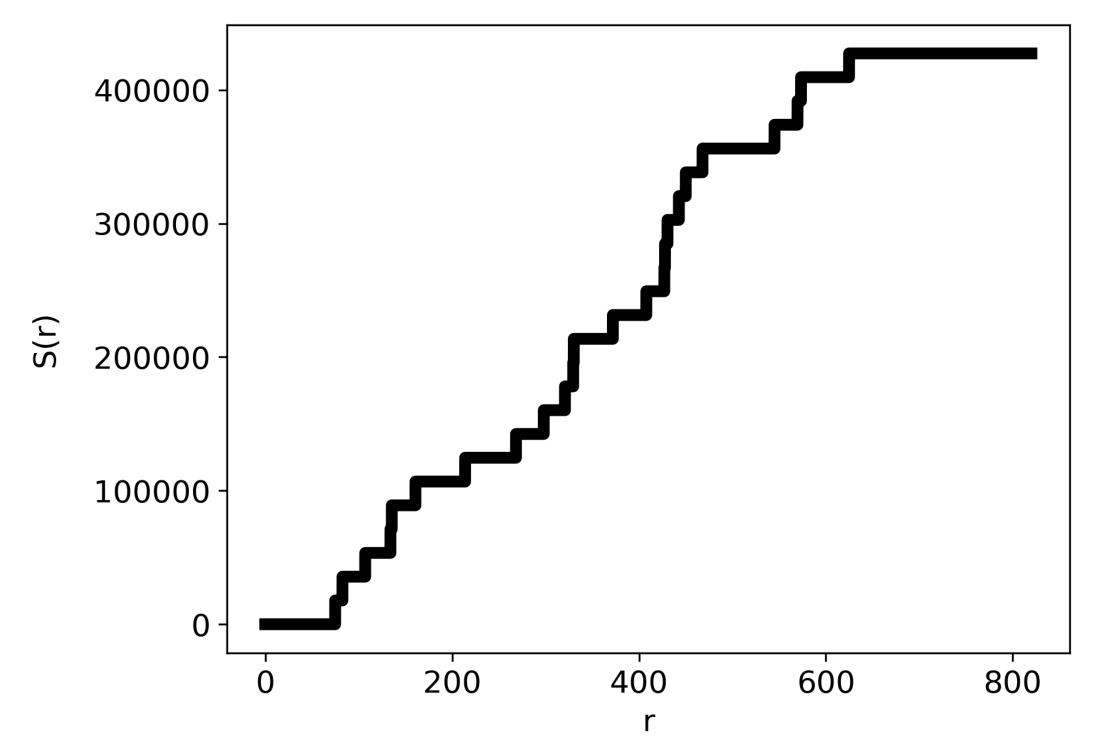






$$|\Omega|$$
 = Area of FOV

$$S(r) = \frac{|\Omega|}{n_1 n_2} \sum_{i=1}^{n_1} \sum_{j=1}^{n_2} \mathbf{1}(d_{ij} < r) b(i, j, r)$$









$$S(r) = \frac{|\Omega|}{n_1 n_2} \sum_{i=1}^{n_1} \sum_{j=1}^{n_2} \mathbf{1}(d_{ij} < r) b(i, j, r)$$

$$b(i,j,r) = \frac{|c(i,d_{ij})|}{|c(i,d_{ij}) \cap \Omega|}$$







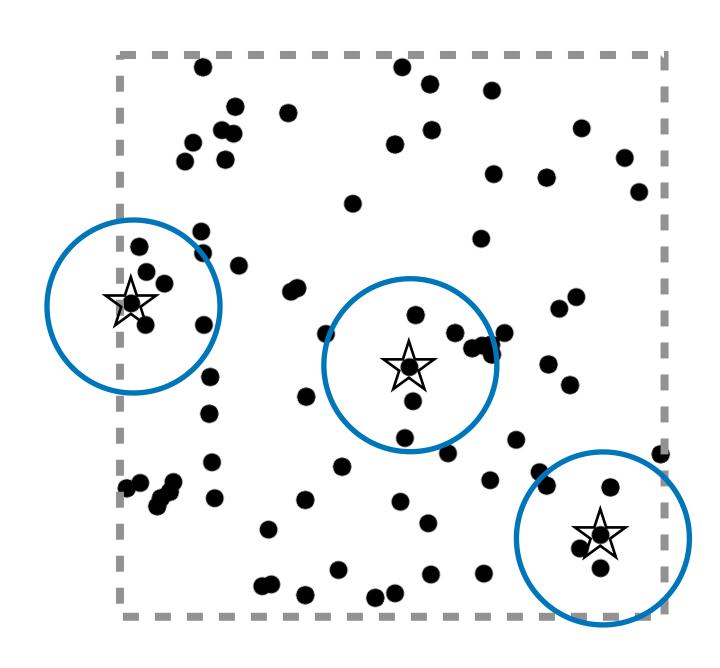
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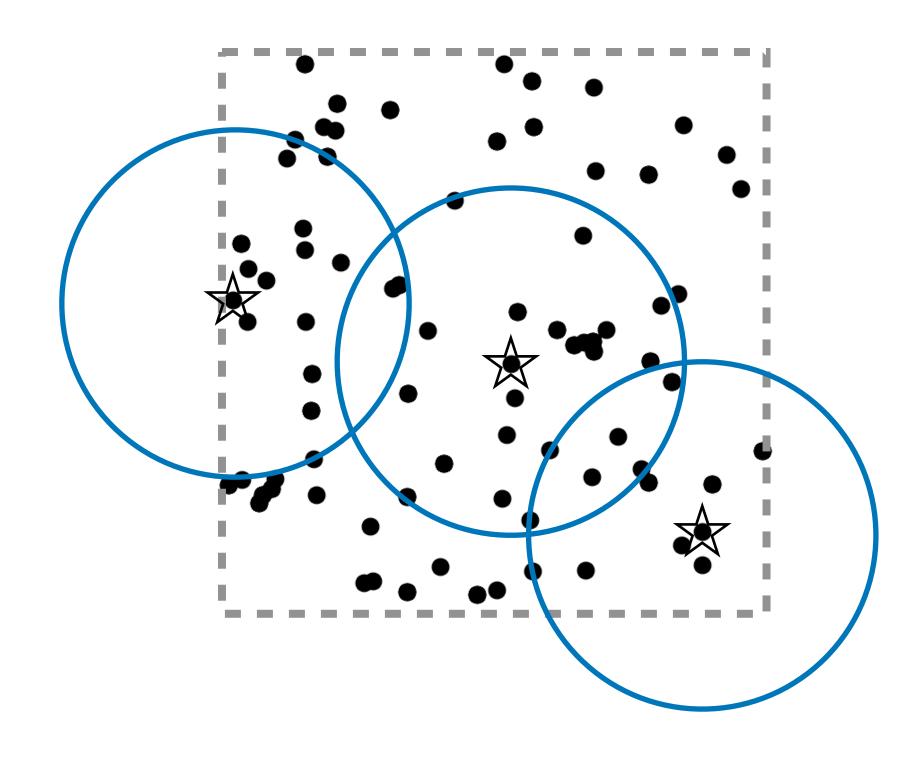
$$S(r) = \frac{|\Omega|}{n_1 n_2} \sum_{i=1}^{n_1} \sum_{j=1}^{n_2} \mathbf{1}(d_{ij} < r) b(i, j, r)$$

$$b(i,j,r) = \frac{|c(i,d_{ij})|}{|c(i,d_{ij}) \cap \Omega|}$$









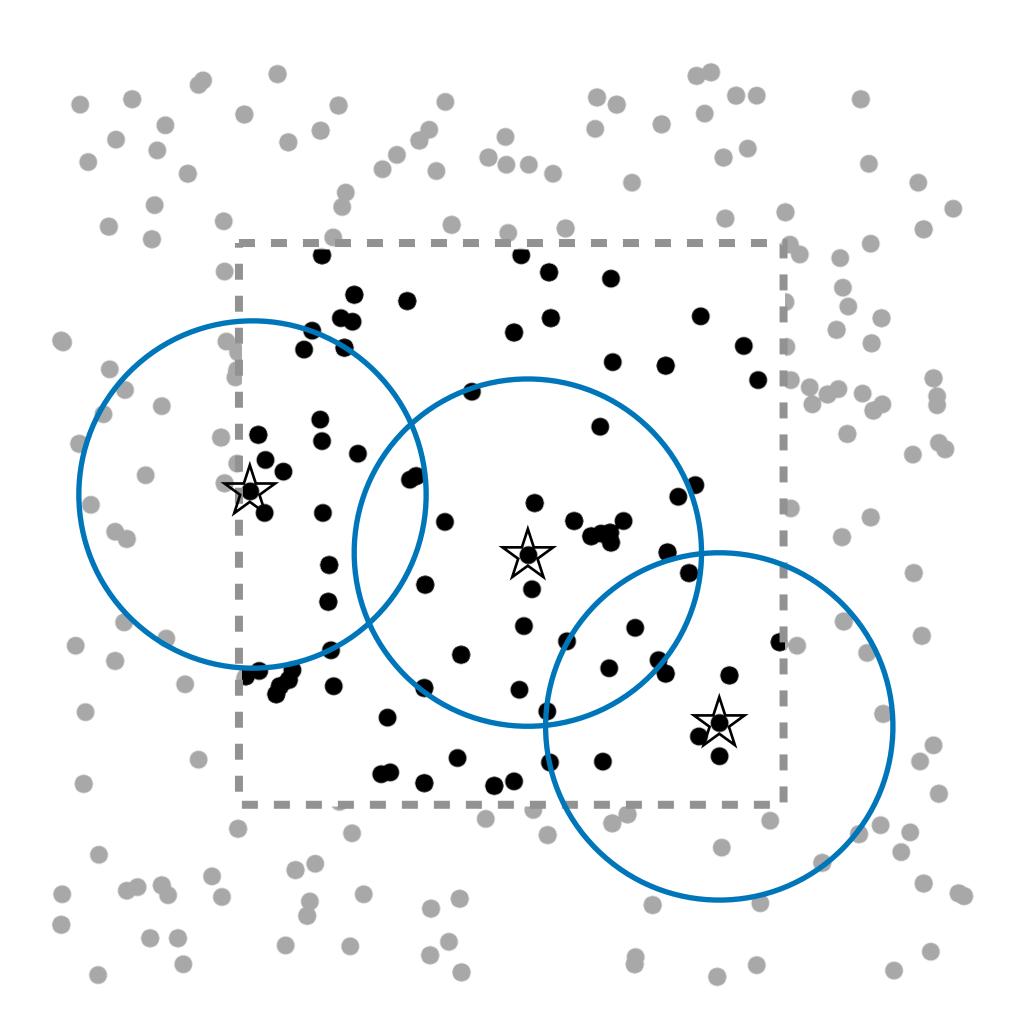
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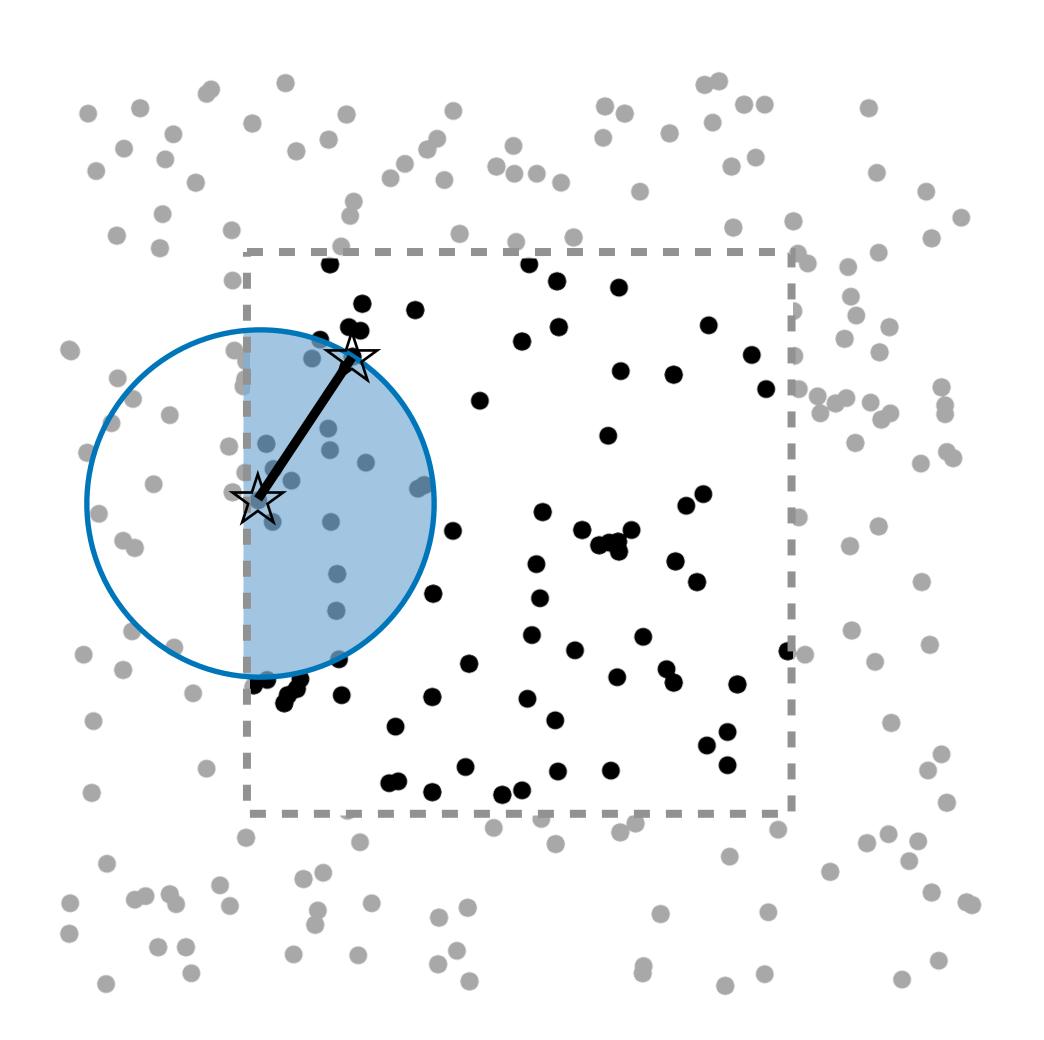
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$$b(i,j,r) = \frac{|c(i,d_{ij})|}{|c(i,d_{ij}) \cap \Omega|}$$









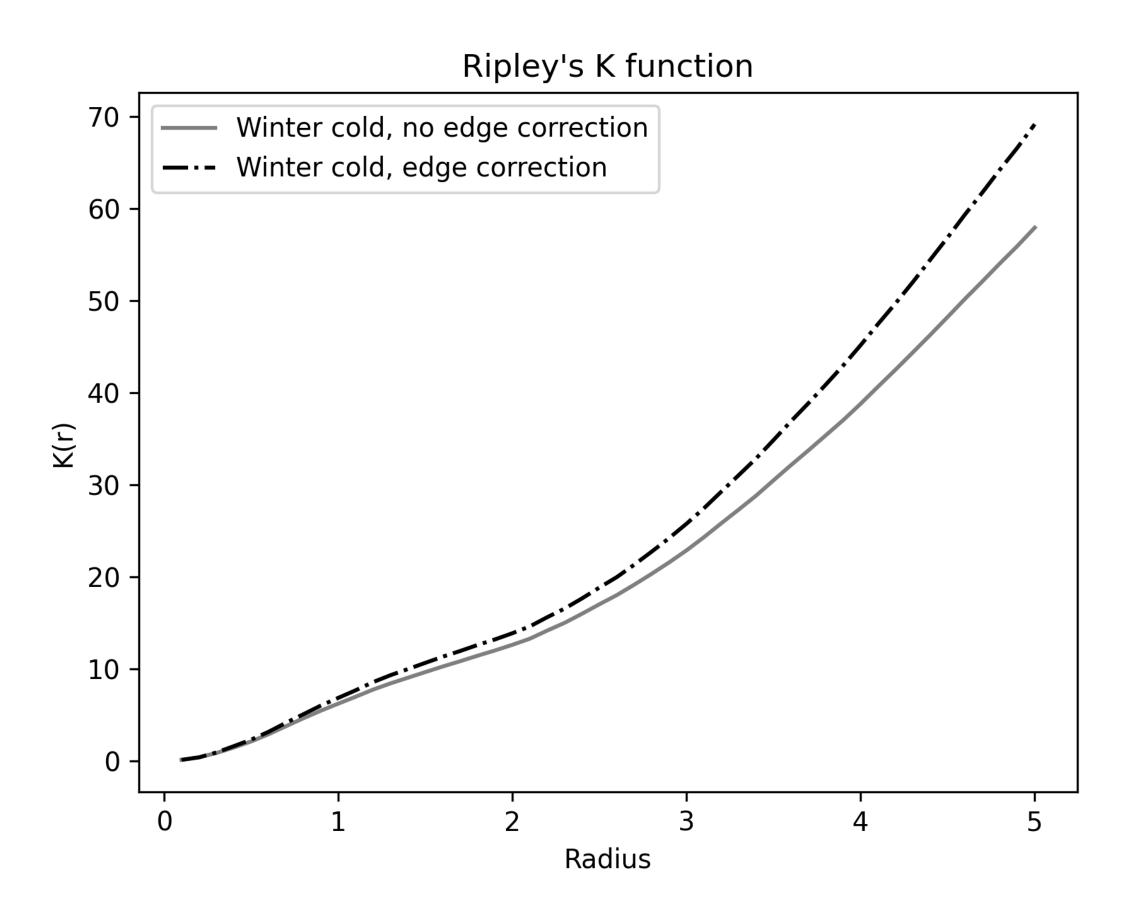
$$S(r) = \frac{|\Omega|}{n_1 n_2} \sum_{i=1}^{n_1} \sum_{j=1}^{n_2} \mathbf{1}(d_{ij} < r) b(i, j, r)$$

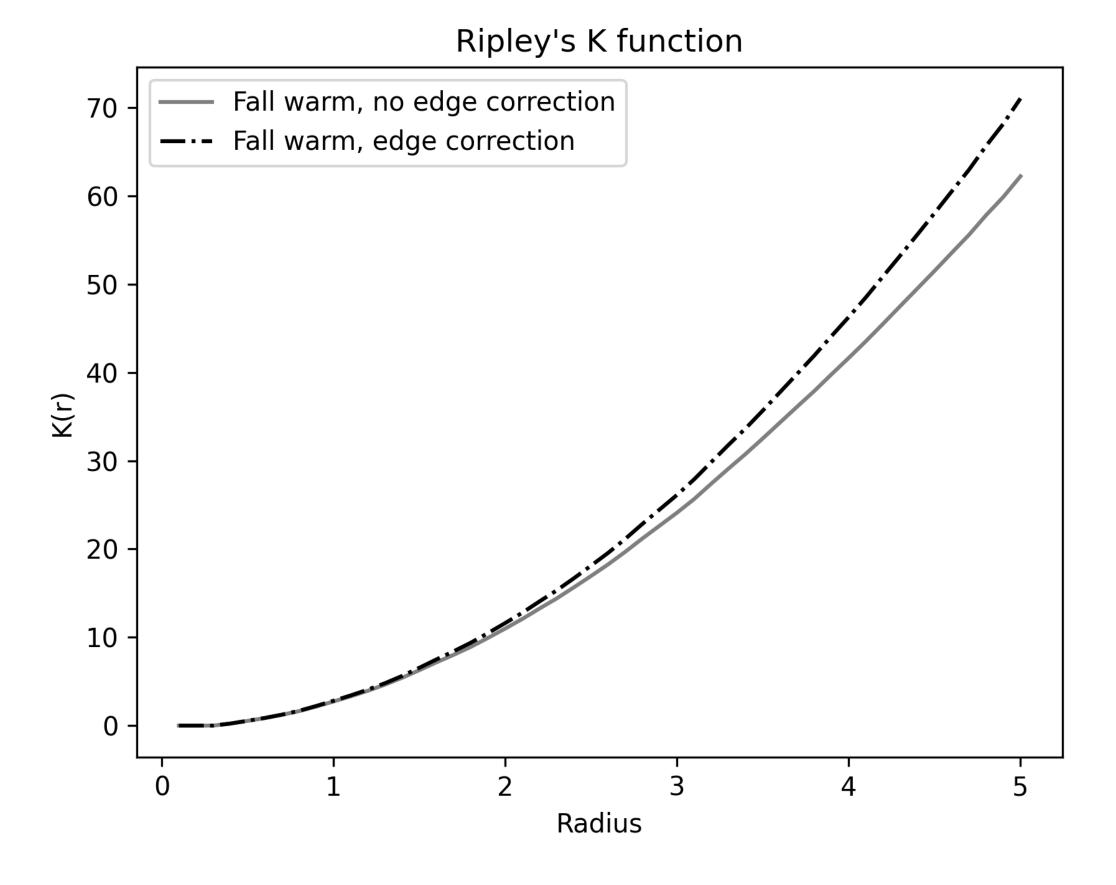
$$b(i,j,r) = \frac{|c(i,d_{ij})|}{|c(i,d_{ij}) \cap \Omega|}$$









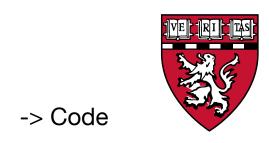














- Symmetric: BOB → IAC = IAC → BOB
- Returns: A number for each radius
- Range: Long

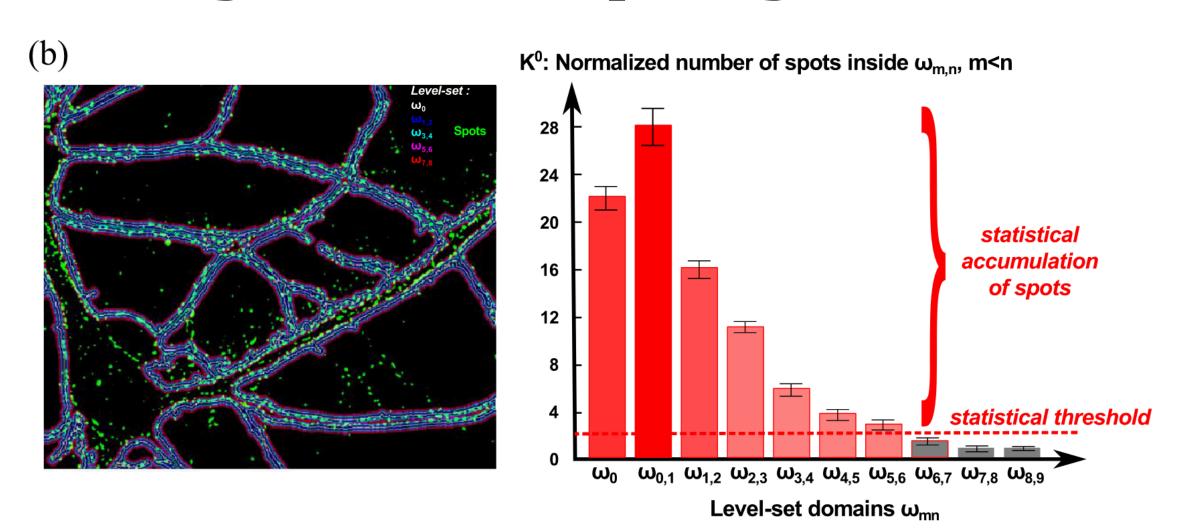




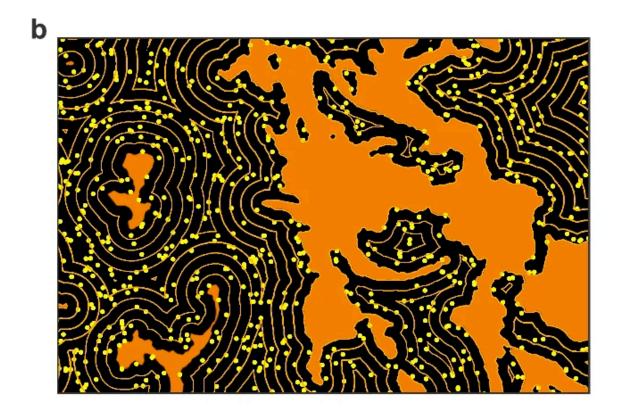


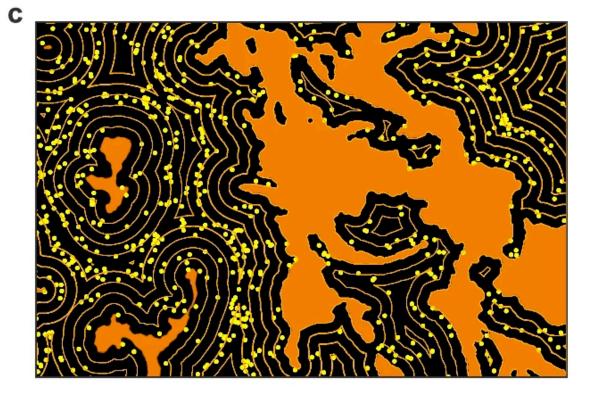


Beyond Ripley's K function



S. Mukherjee, C. Gonzalez-Gomez, L. Danglot, T. Lagache and J. -C. Olivo-Marin, "Generalizing the Statistical Analysis of Objects' Spatial Coupling in Bioimaging," in *IEEE Signal Processing Letters*, vol. 27, pp. 1085-1089, 2020, doi: 10.1109/LSP.2020.3003821.





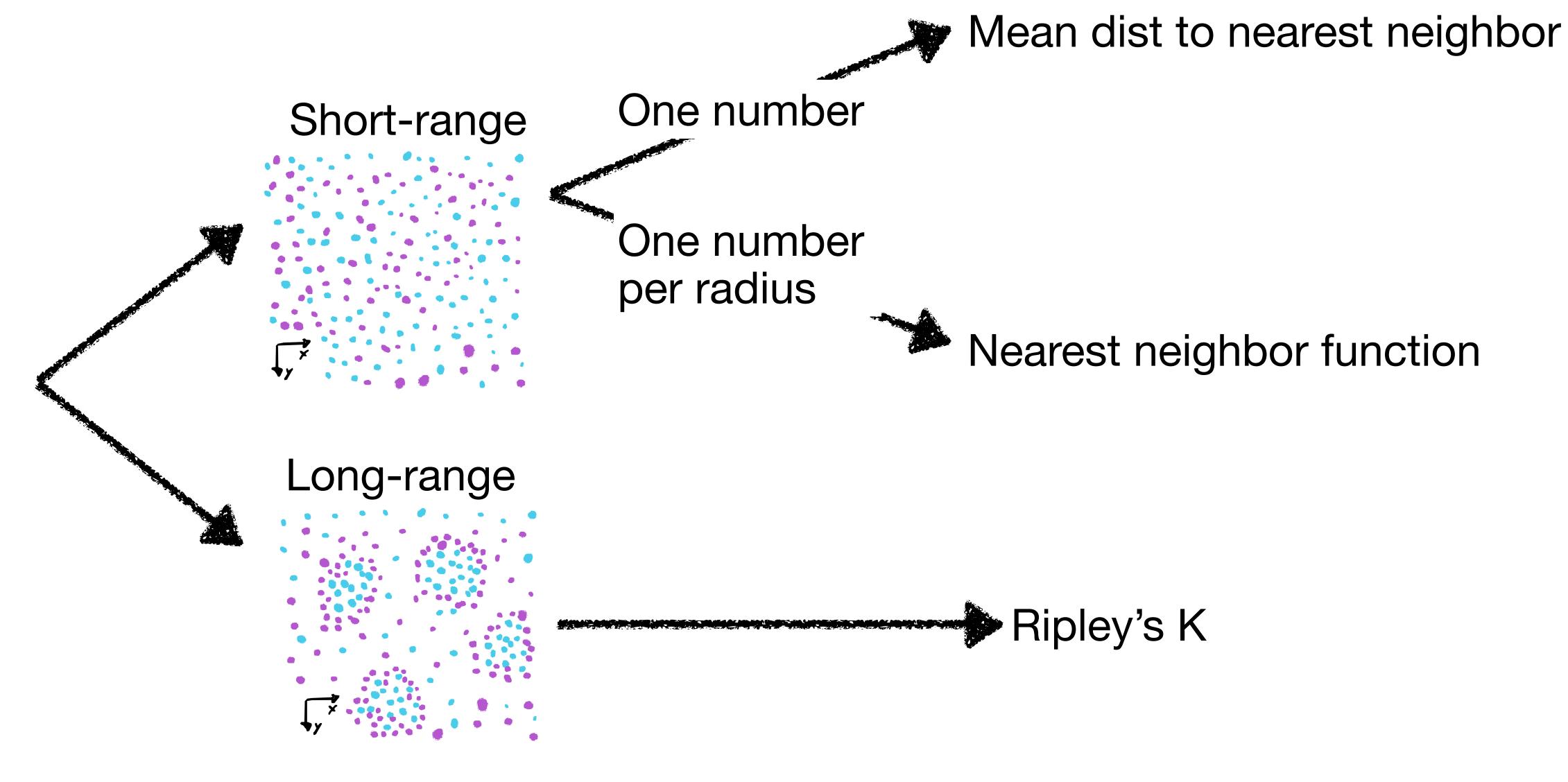
Benimam, M.M., Meas-Yedid, V., Mukherjee, S. *et al.* Statistical analysis of spatial patterns in tumor microenvironment images. *Nat Commun* **16**, 3090 (2025). https://doi.org/10.1038/s41467-025-57943-y







Summary

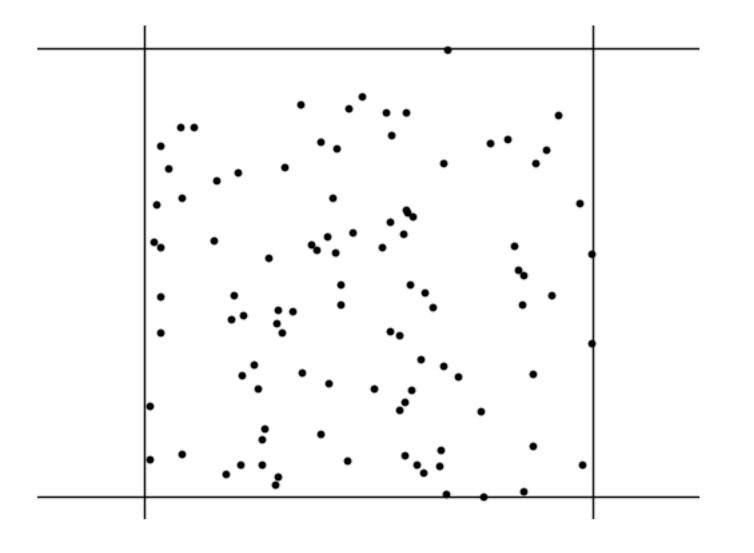








- How are proteins distributed that
 - Don't interact with each other
 - Or their surrounding



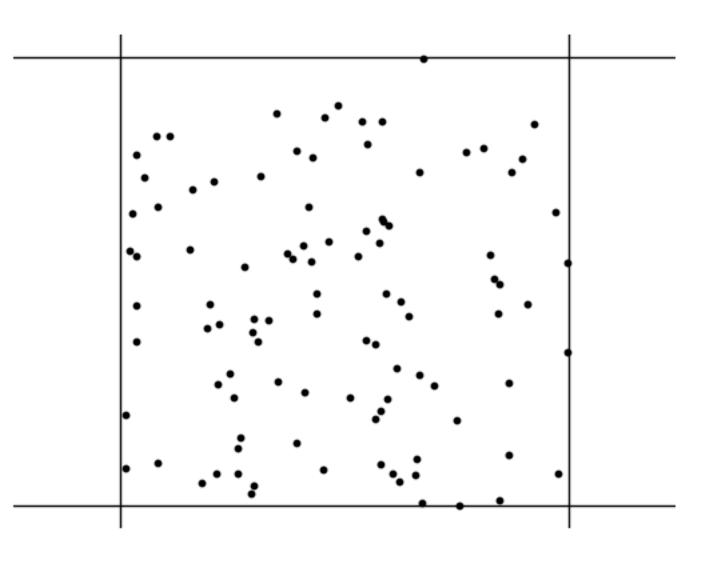
What is the evidence that my points may or may not be distributed like this?







- How are proteins distributed that
 - Don't interact with each other
 - Or their surrounding
- To find out, you blindly throw darts at a board



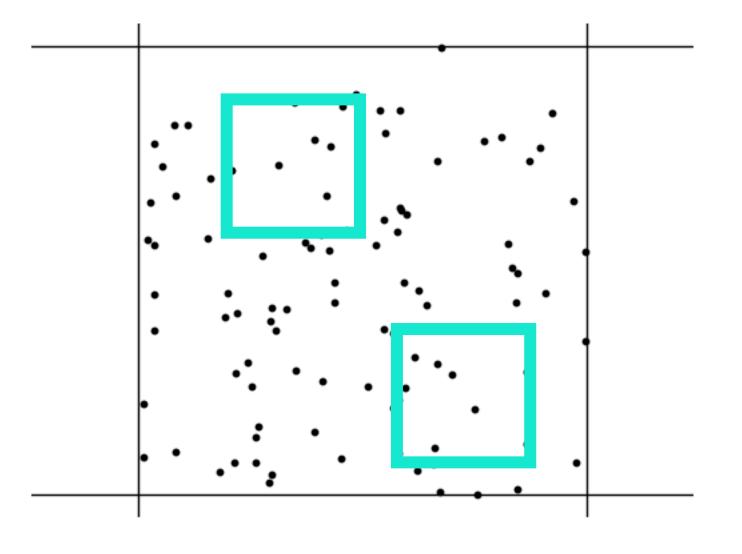
What is the evidence that my points may or may not be distributed like this?







- How are proteins distributed that
 - Don't interact with each other
 - Or their surrounding
- To find out, you blindly throw darts at a board
- The chance of a dart landing is the same, no matter where on the board



What is the evidence that my points may or may not be distributed like this?



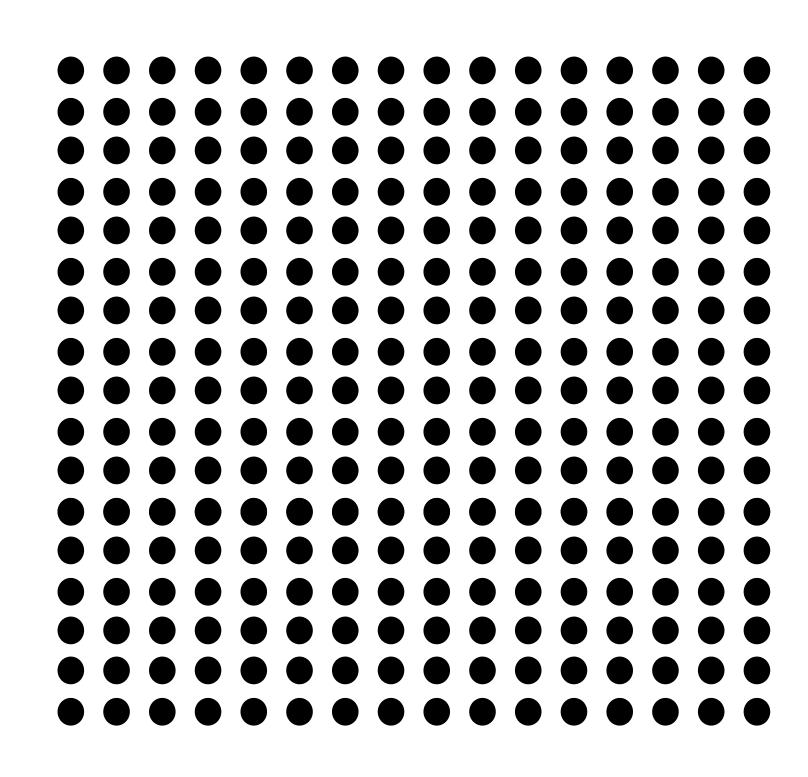




"Uniformly distributed"



"Uniformly spaced"

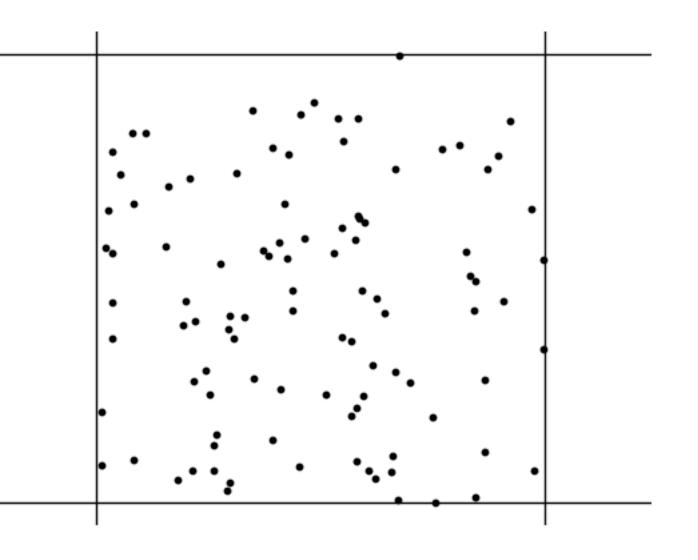








- How are proteins distributed that
 - Don't interact with each other
 - Or their surrounding
- To find out, you blindly throw darts at a board
- The chance of a dart landing is the same, no matter where on the board

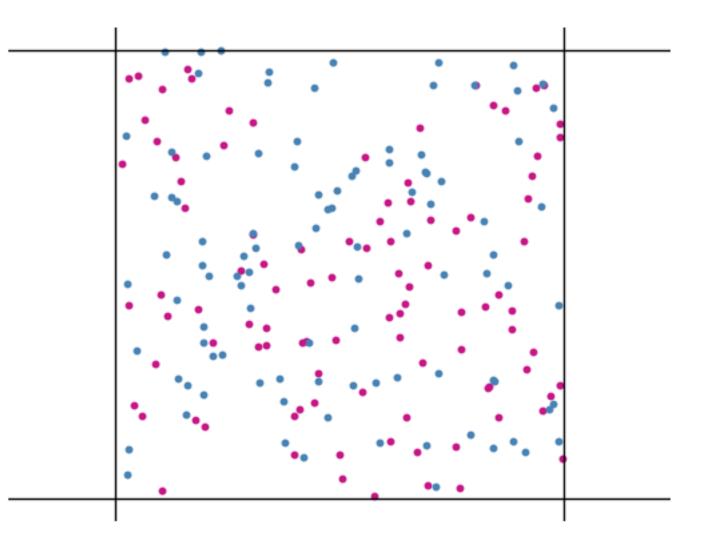








- How are proteins distributed that
 - Don't interact with each other
 - Or their surrounding
- To find out, you blindly throw darts at a board
- The chance of a dart landing is the same, no matter where on the board
- The darts can have multiple colors

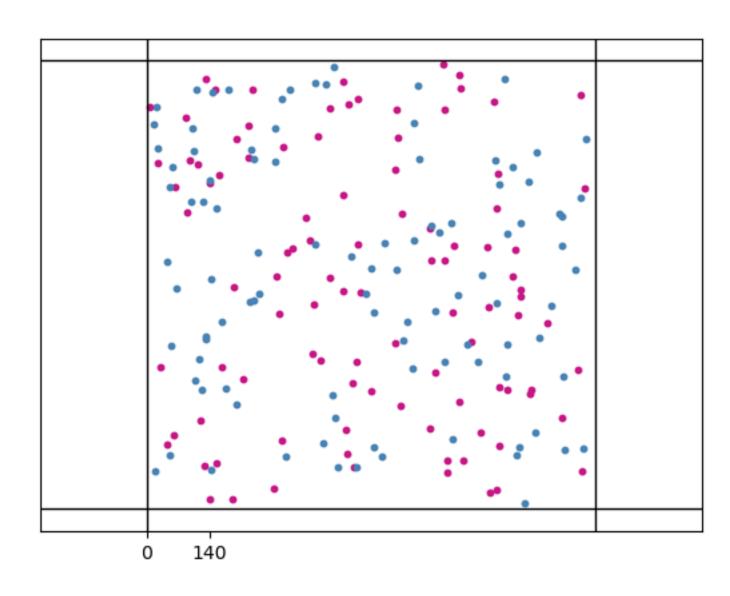


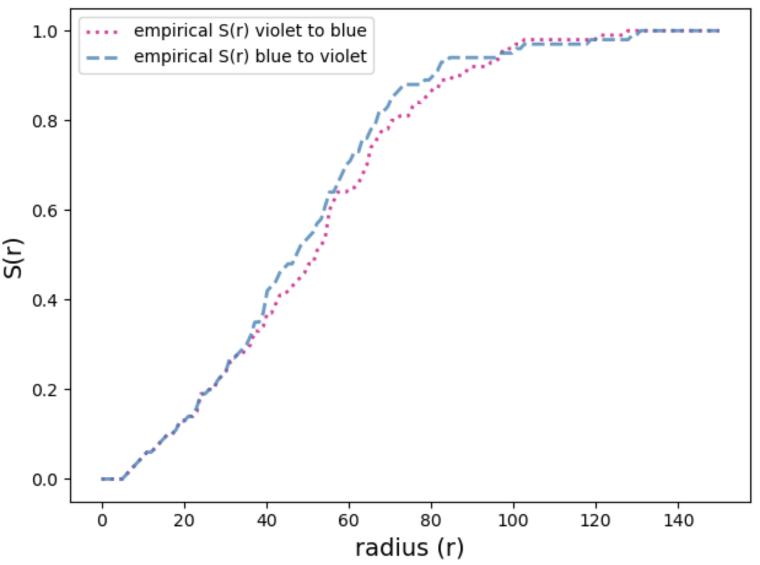






Empirical null distributions



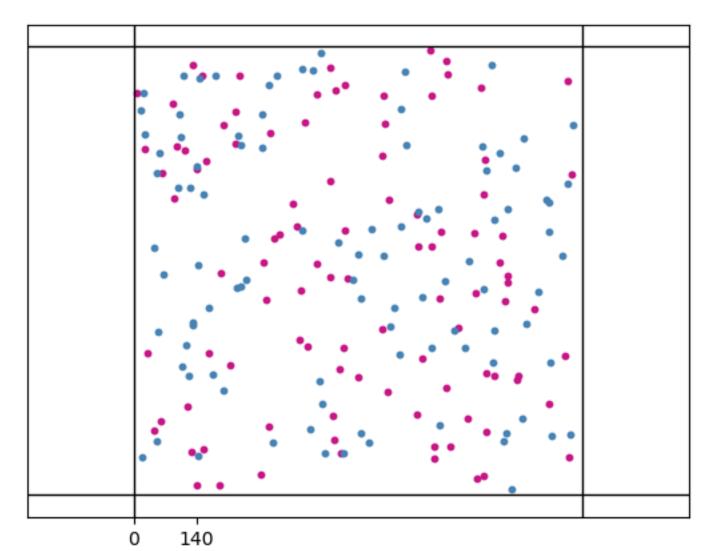


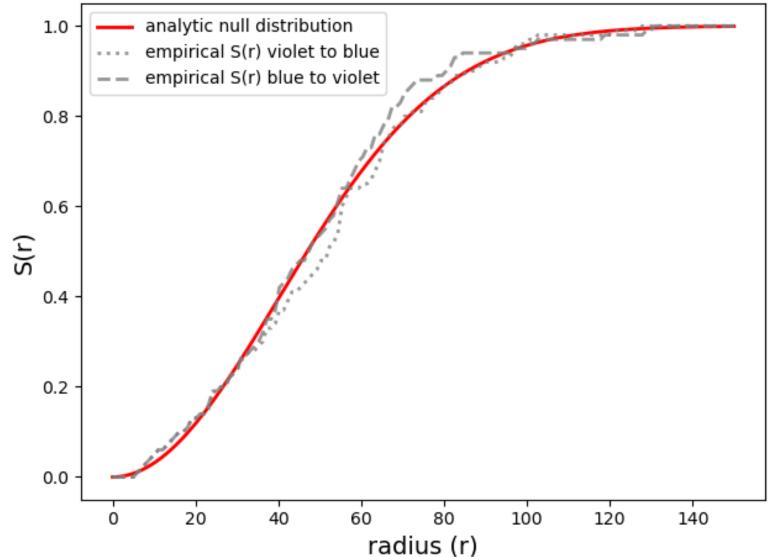


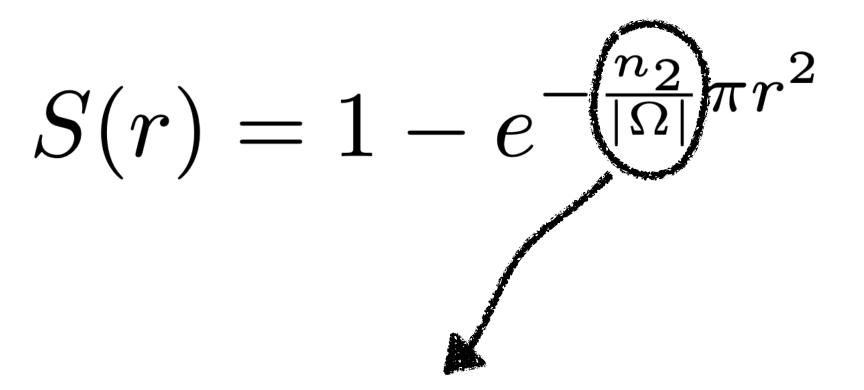




Analytic null distribution







Density of points n₂

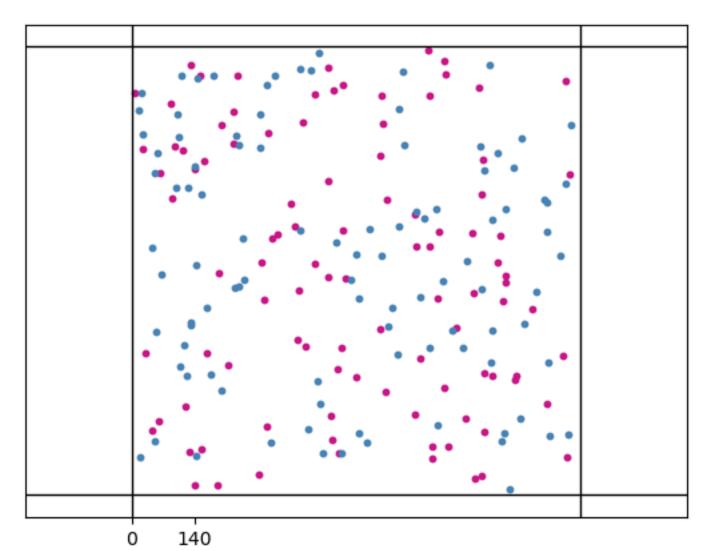
You don't need any data to compute the analytic null distribution

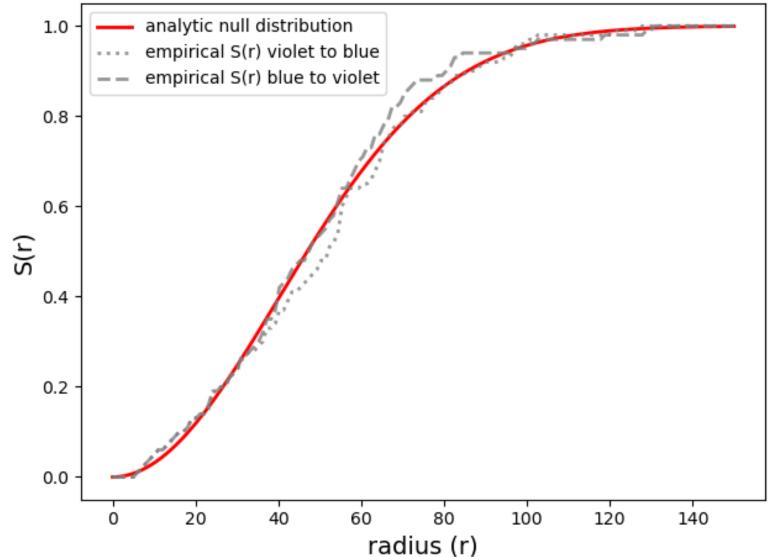






Analytic null distribution



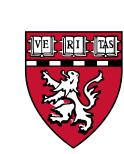


$$S(r) = 1 - e^{-\frac{n_2}{|\Omega|}\pi r^2}$$

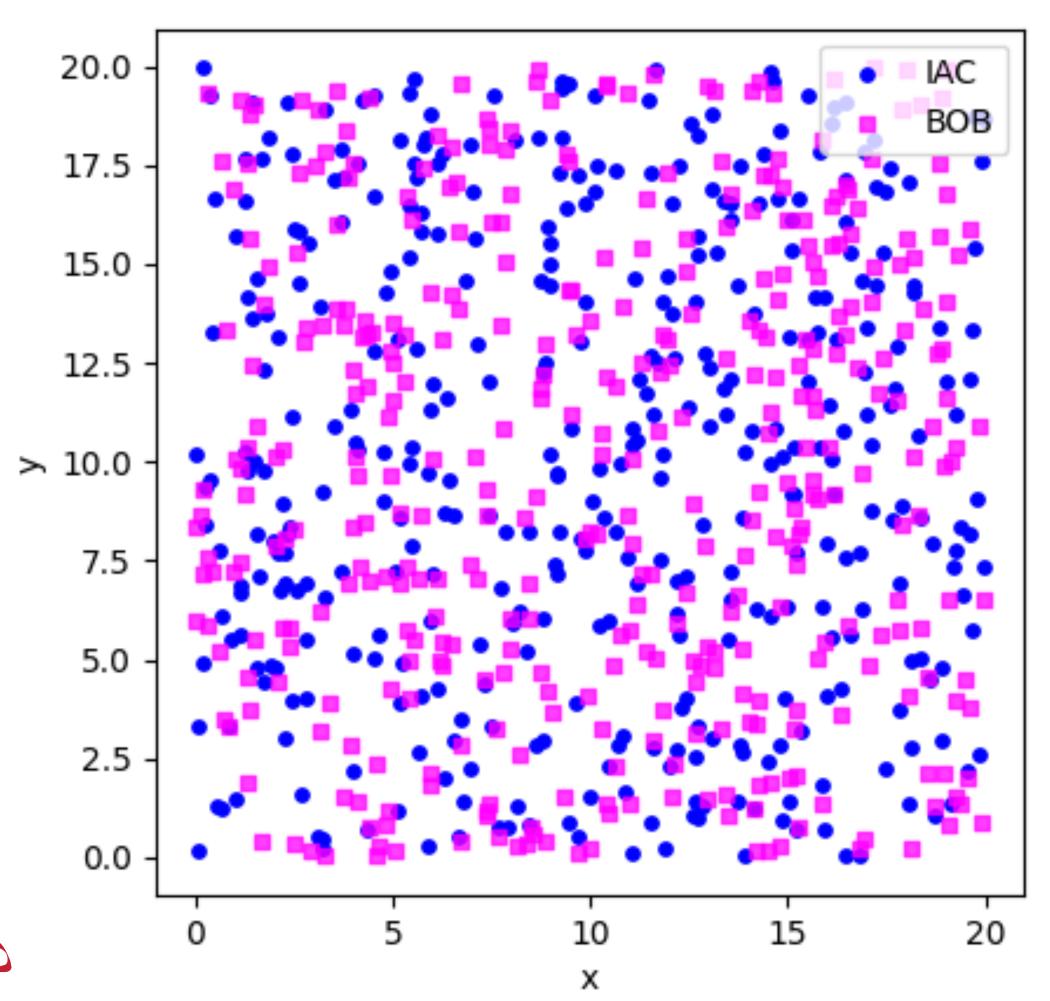
Area of a circle with radius r

You don't need any data to compute the analytic null distribution









BOB = 400

IAC = 400

$$S(r) = 1 - e^{-\frac{n_2}{|\Omega|}\pi r^2}$$

Exercise: Find good values for n2 and $|\Omega|$

```
n = 1345345 # number of points in the dataset
area = 4056780 # area of the FOV

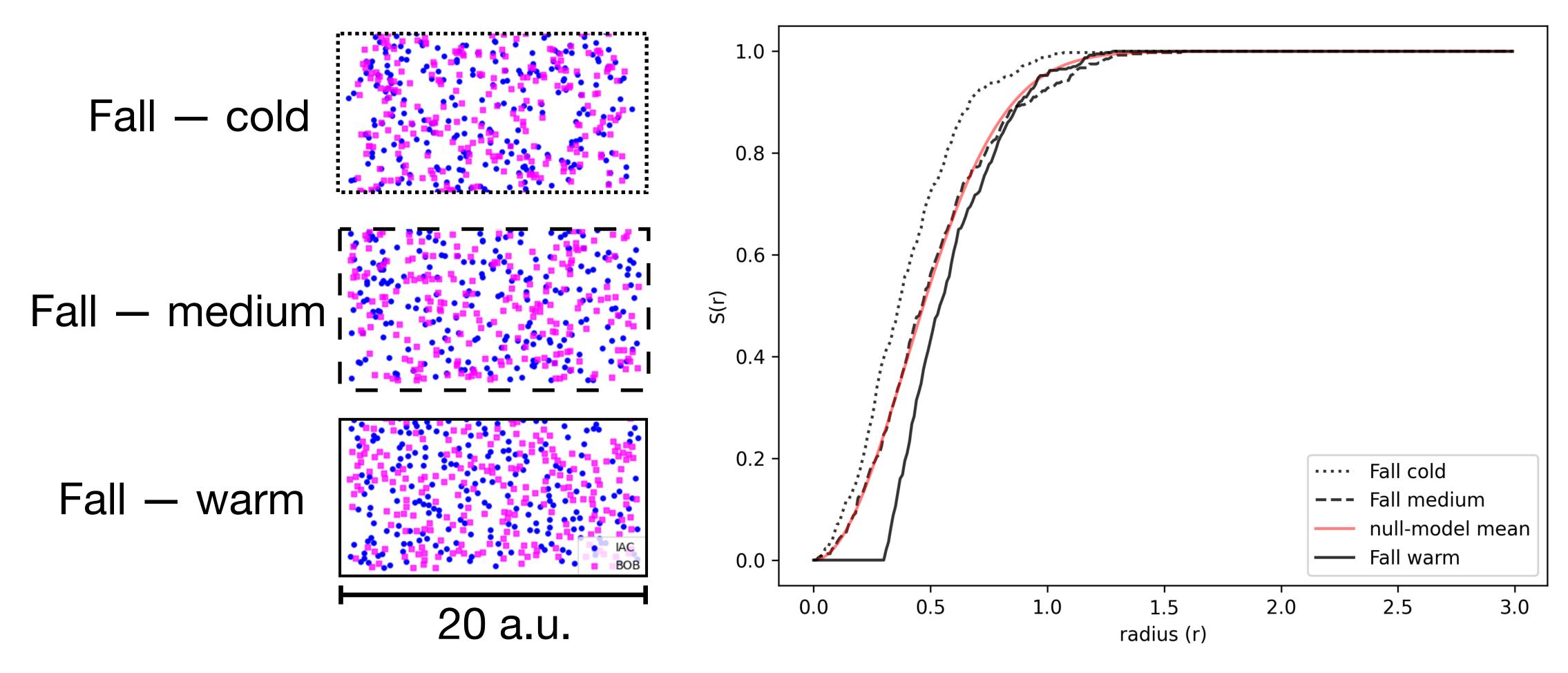
nulldist = getnulldist(
    n=n, area=area, radii=radii
) # These are not good values for n and area. Change them!
```







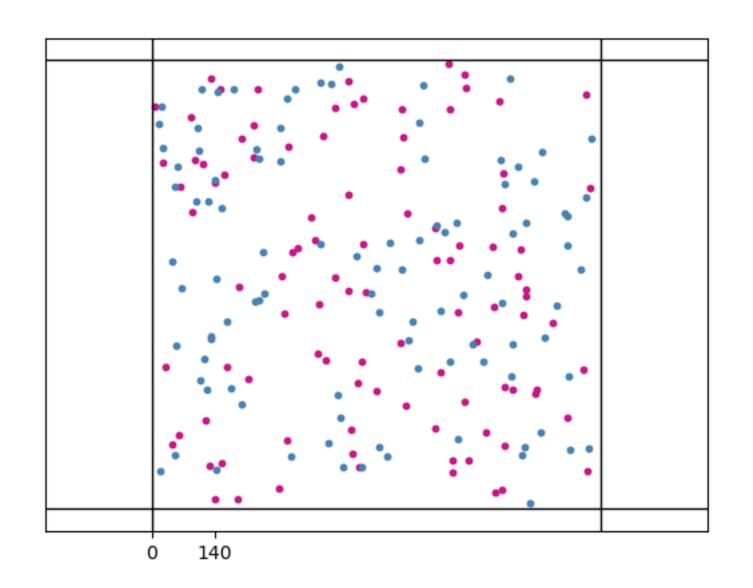
Results — the null distribution

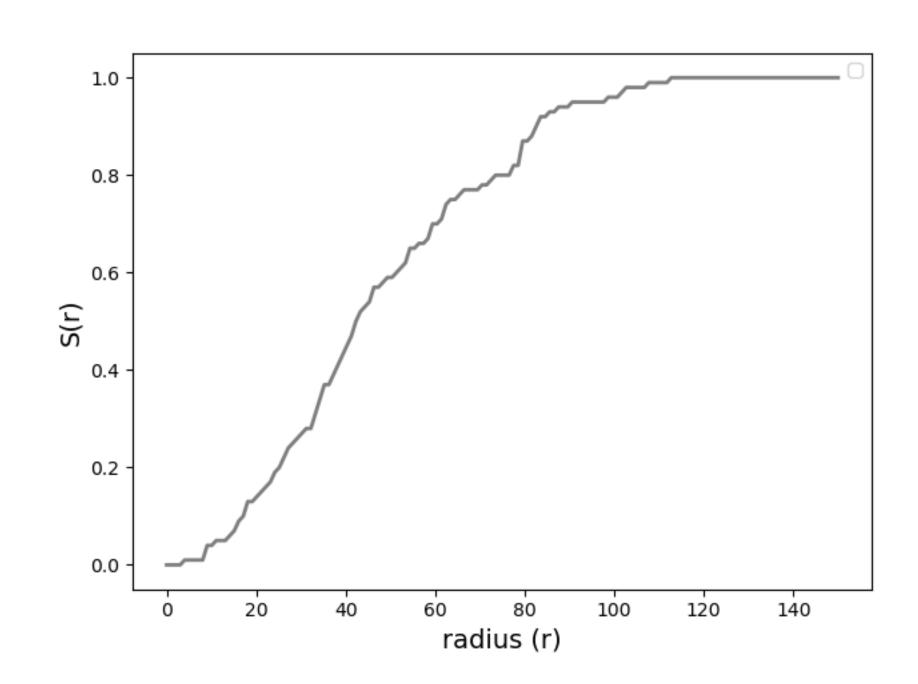










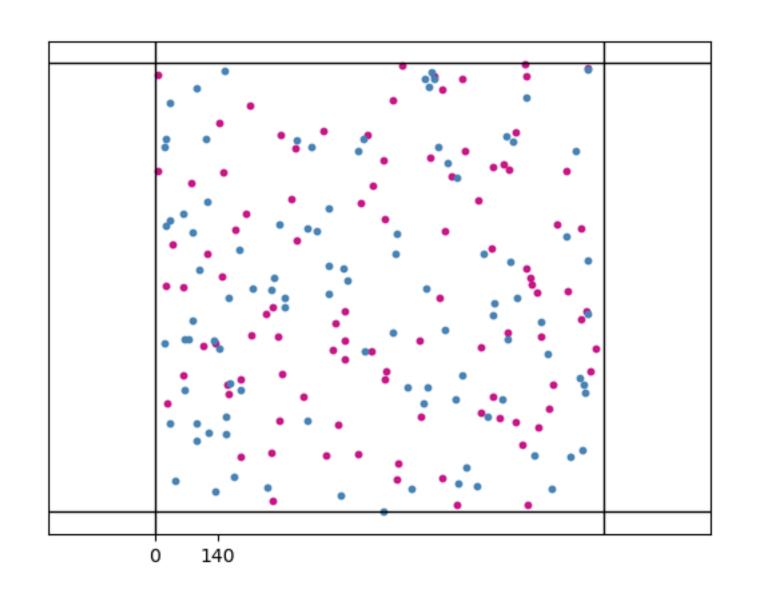


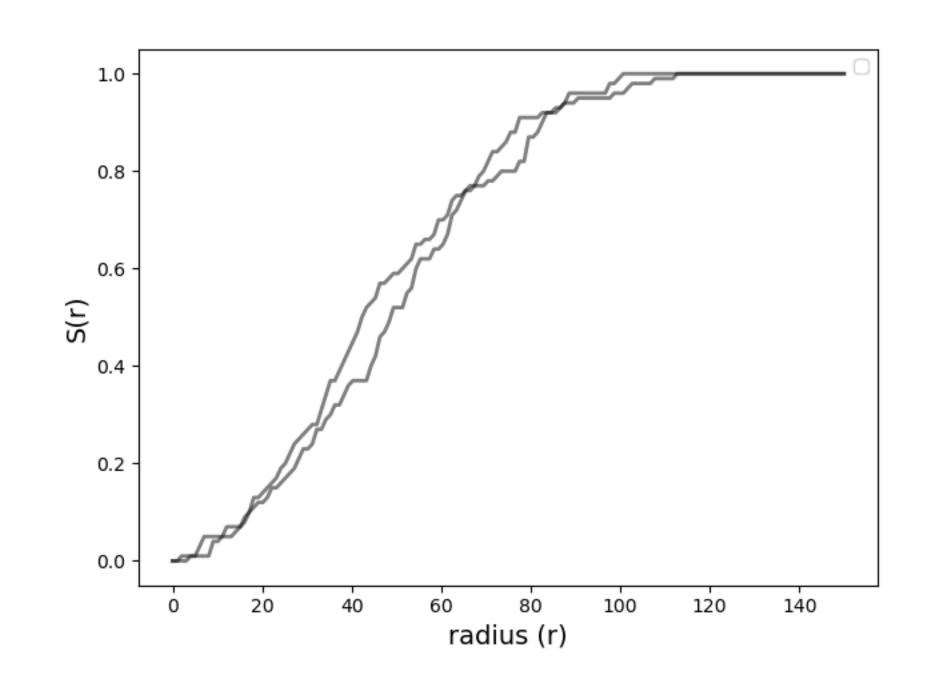








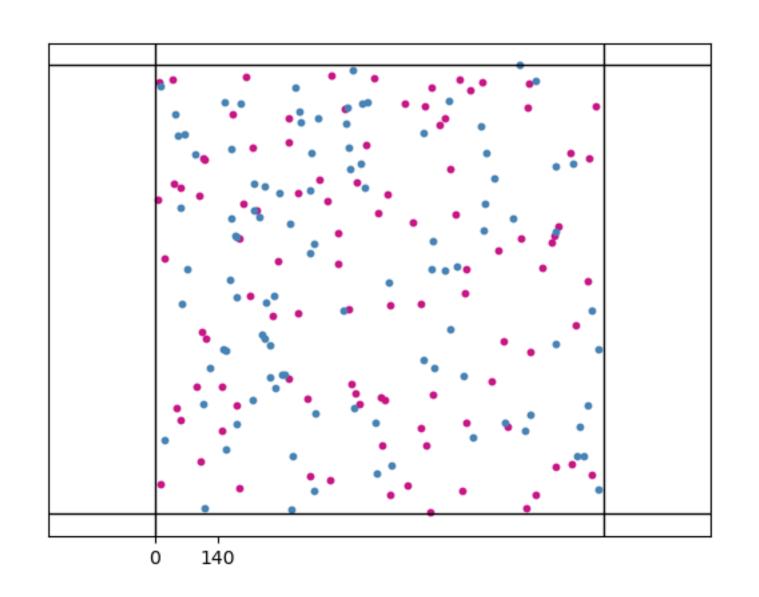


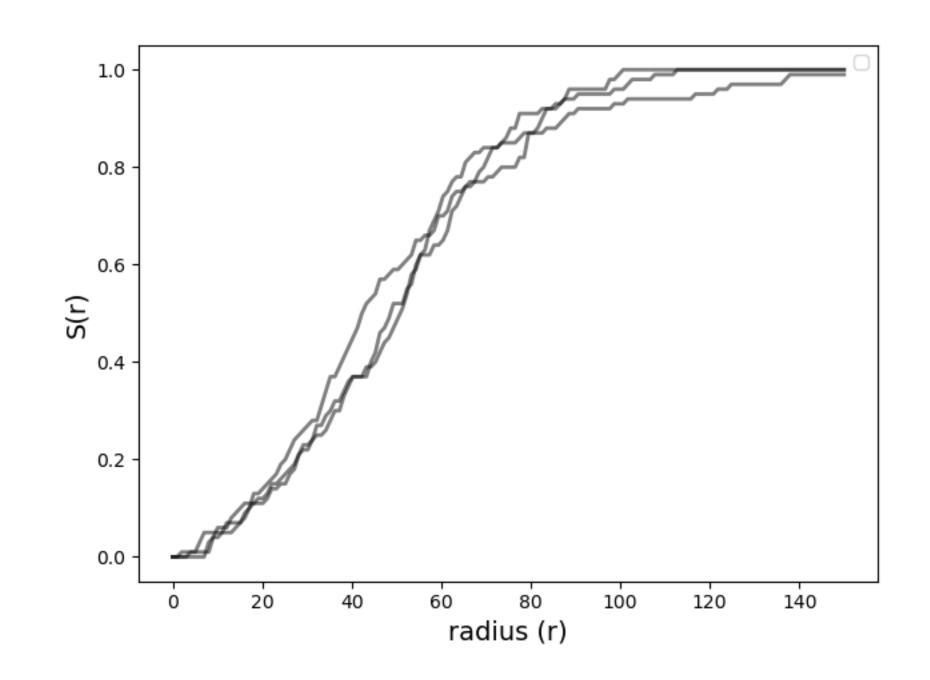










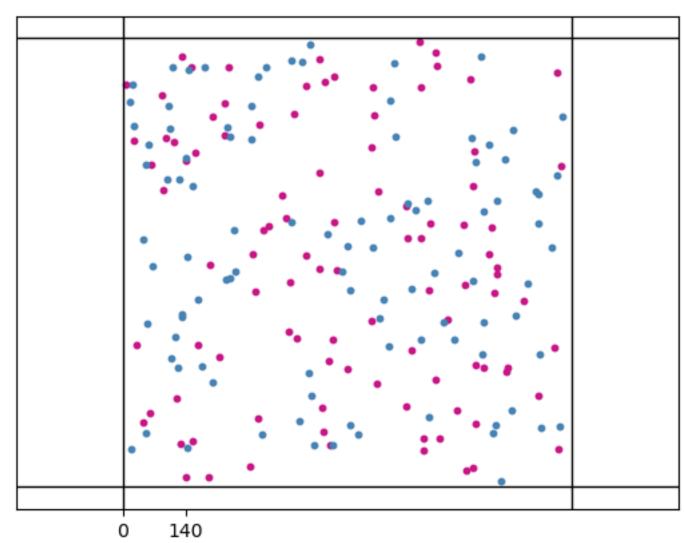


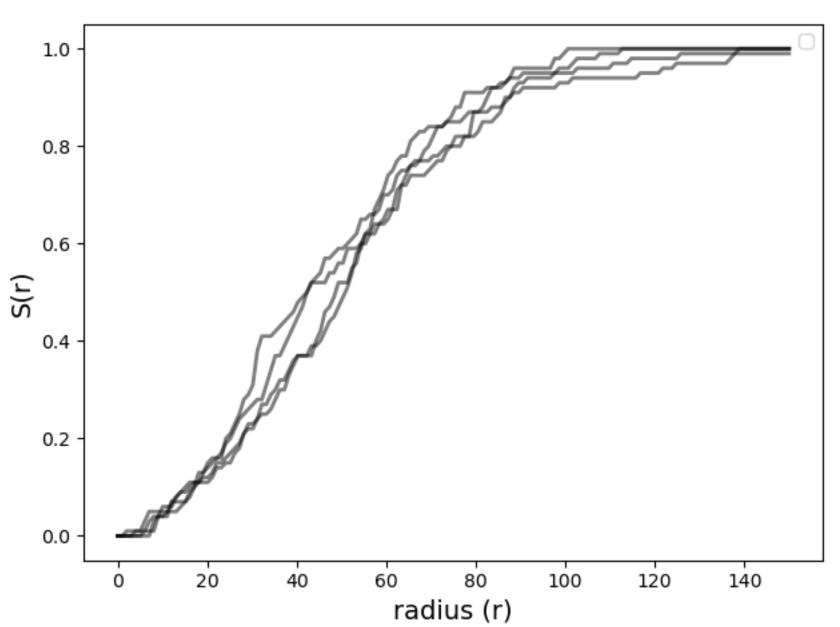










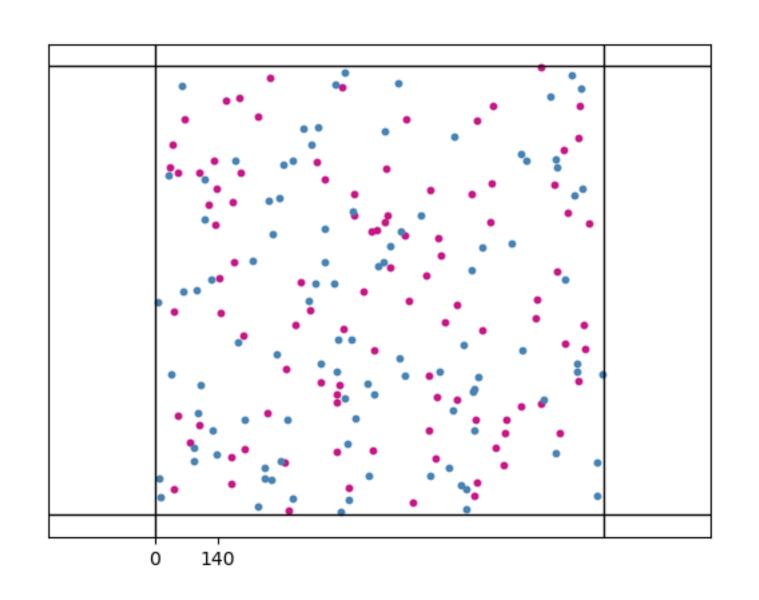


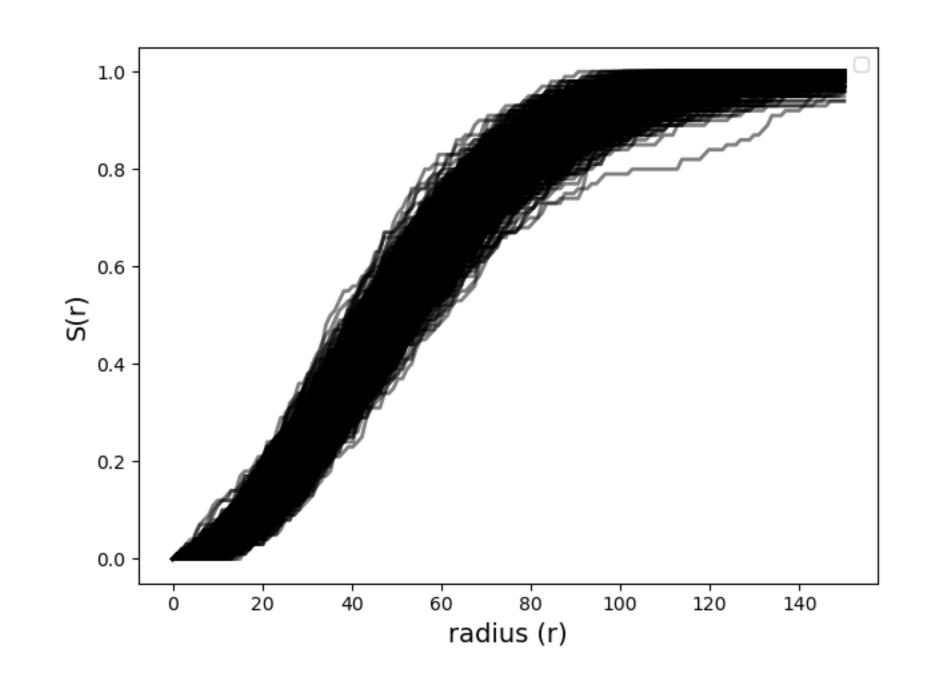










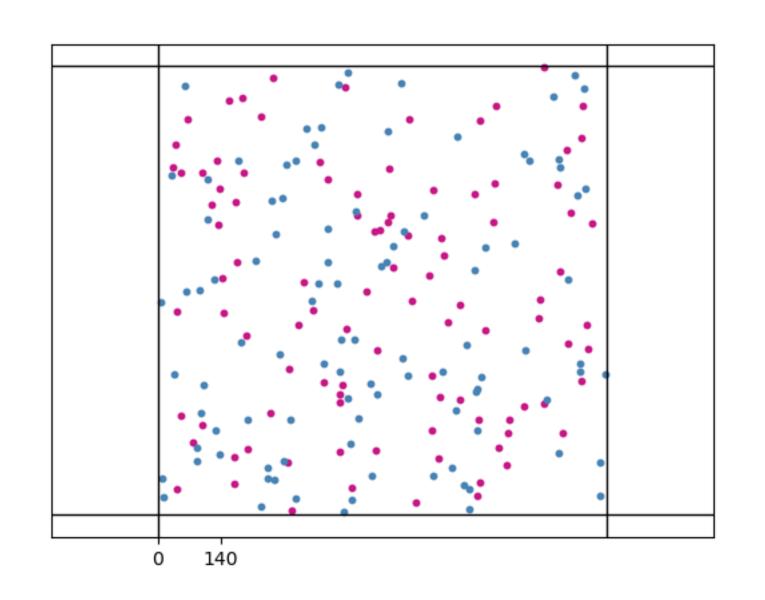


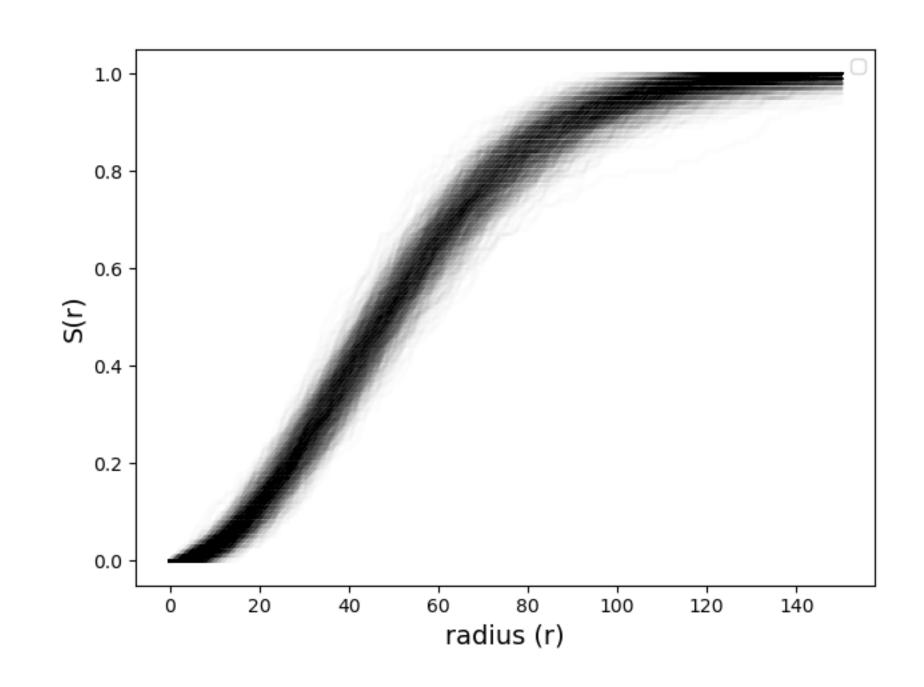
1000









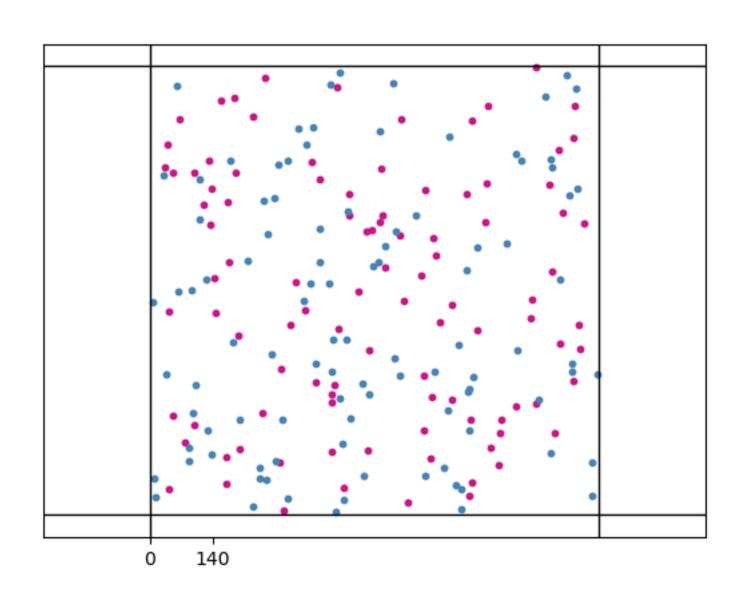


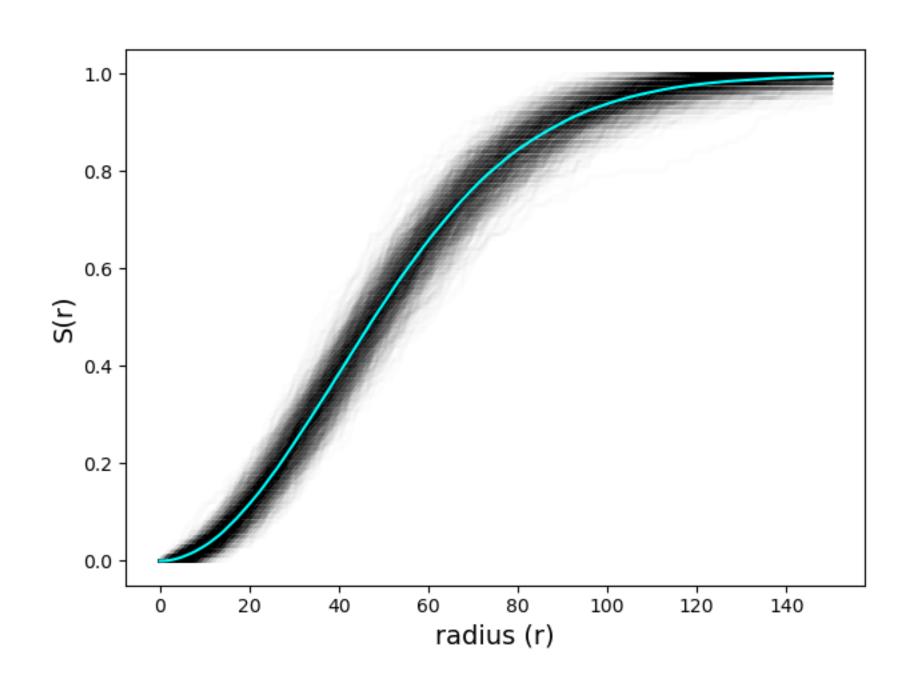
1000









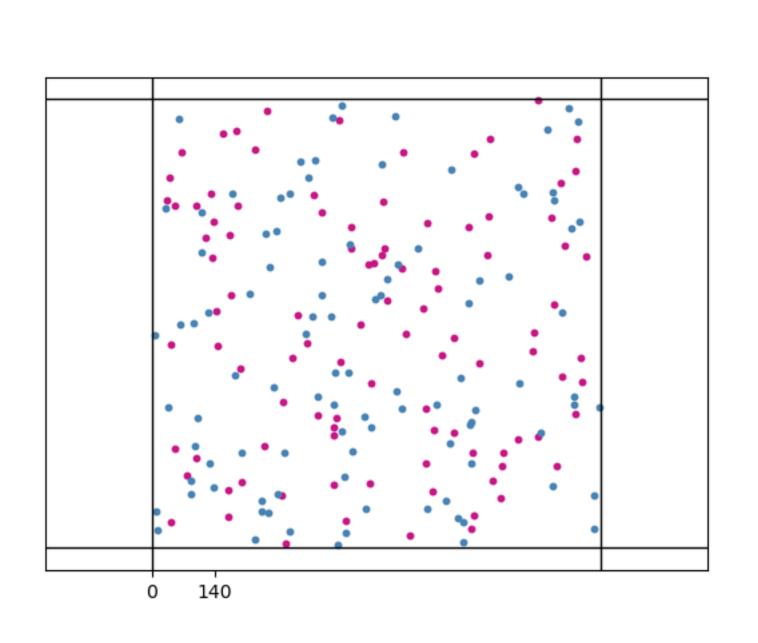


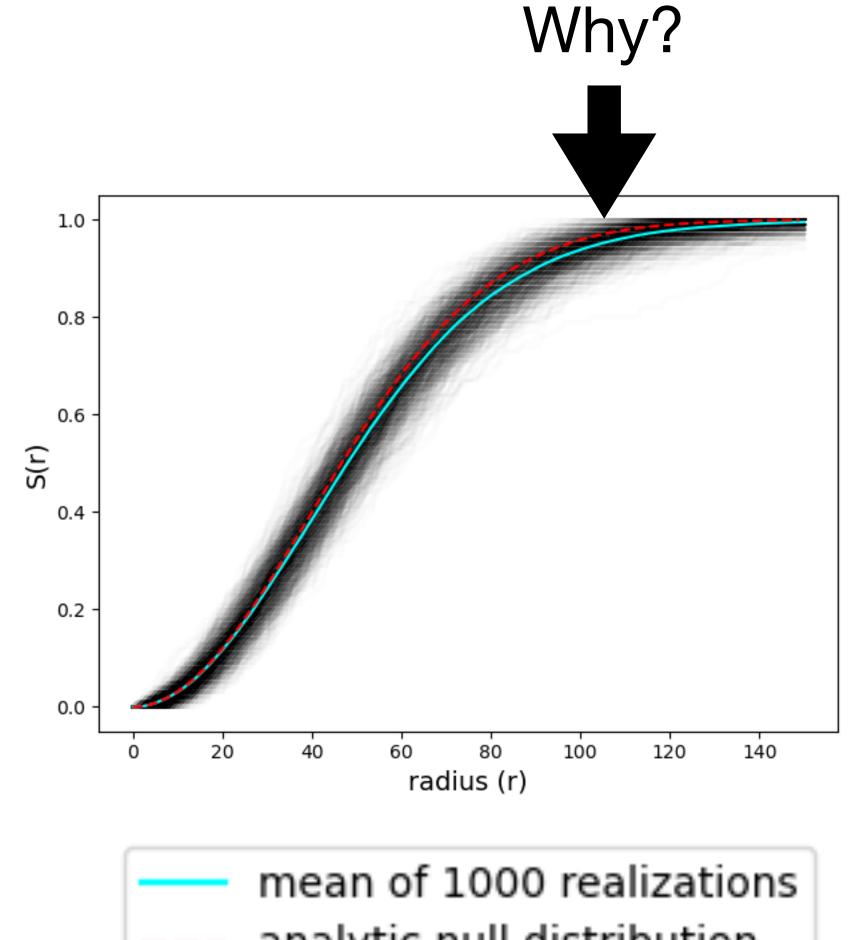
mean of 1000 realizations









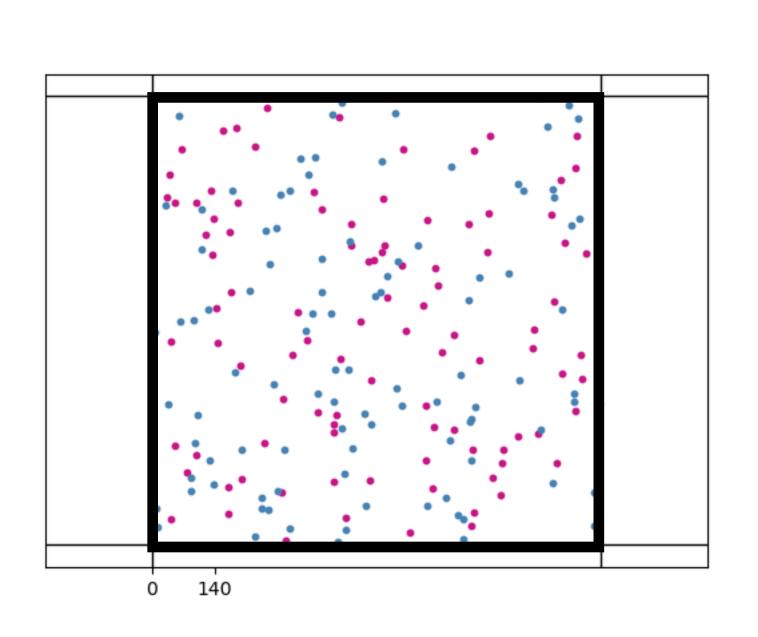


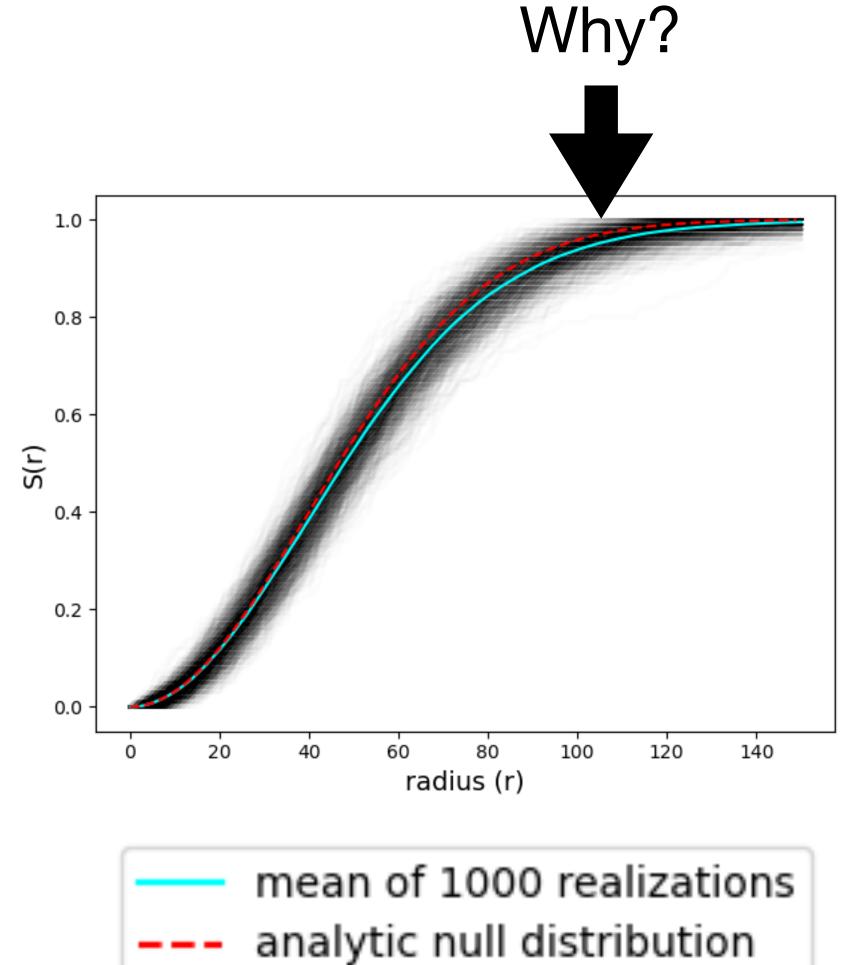
analytic null distribution







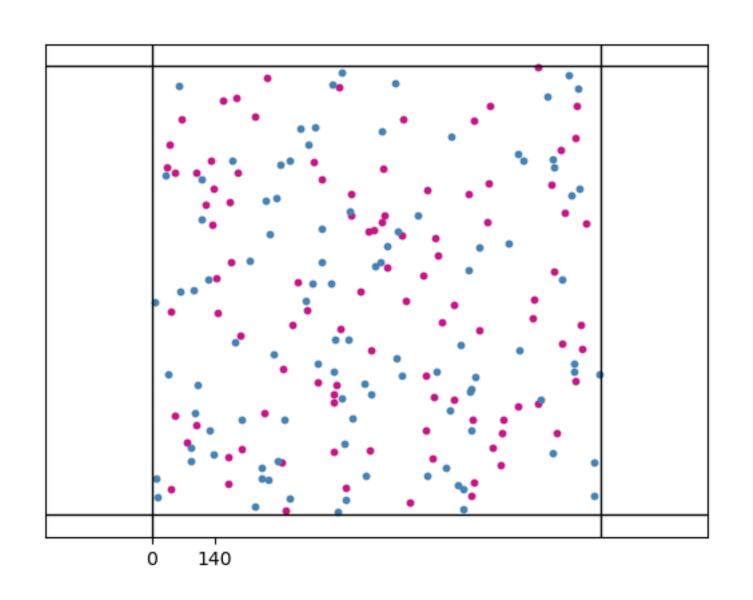


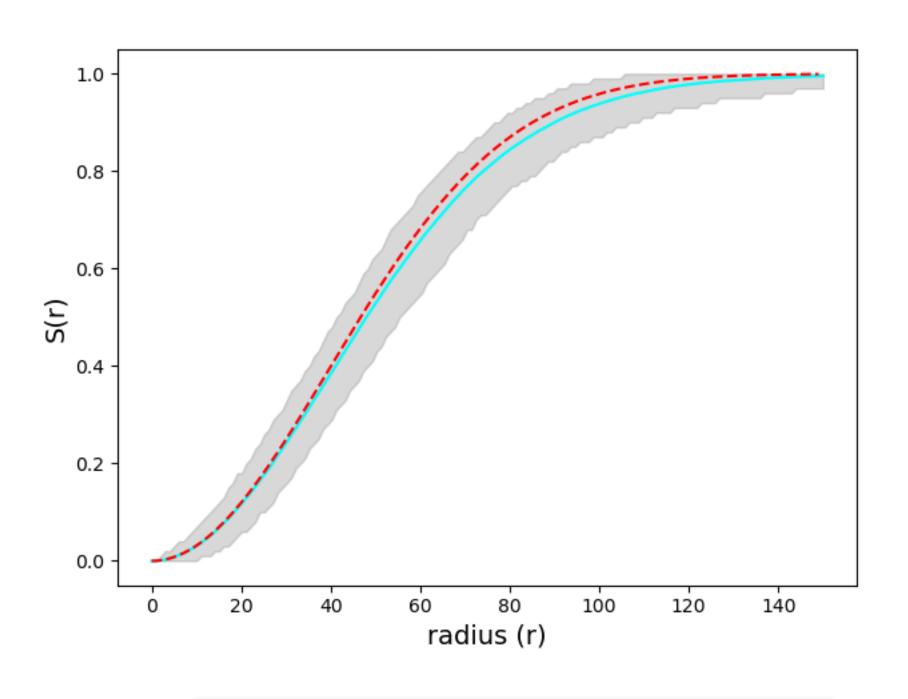












mean of 1000 realizations
2.5-97.5% quantile range
analytic null distribution







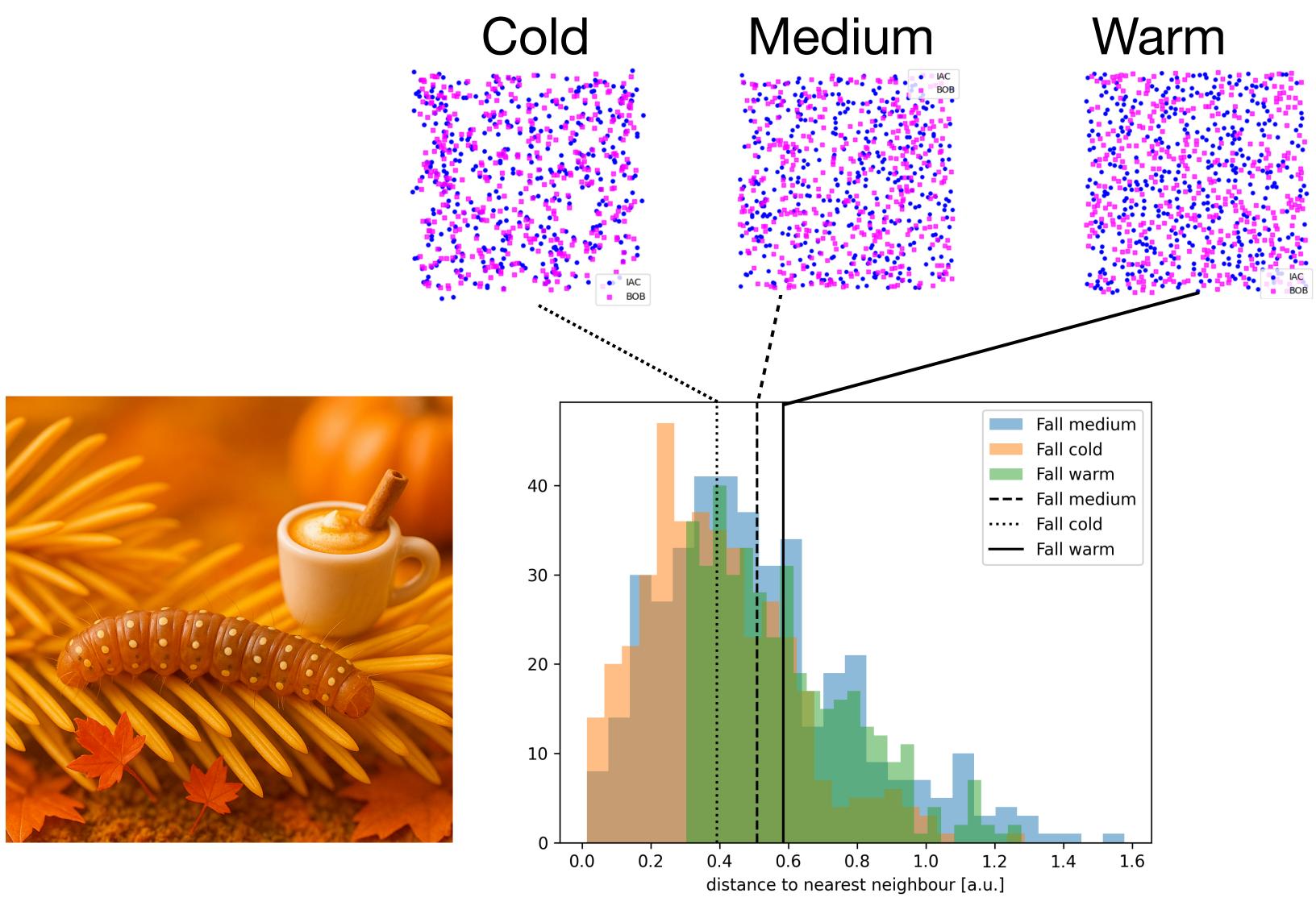
Exercise: throw darts at a board







Results: Mean distance IAC -> BOB

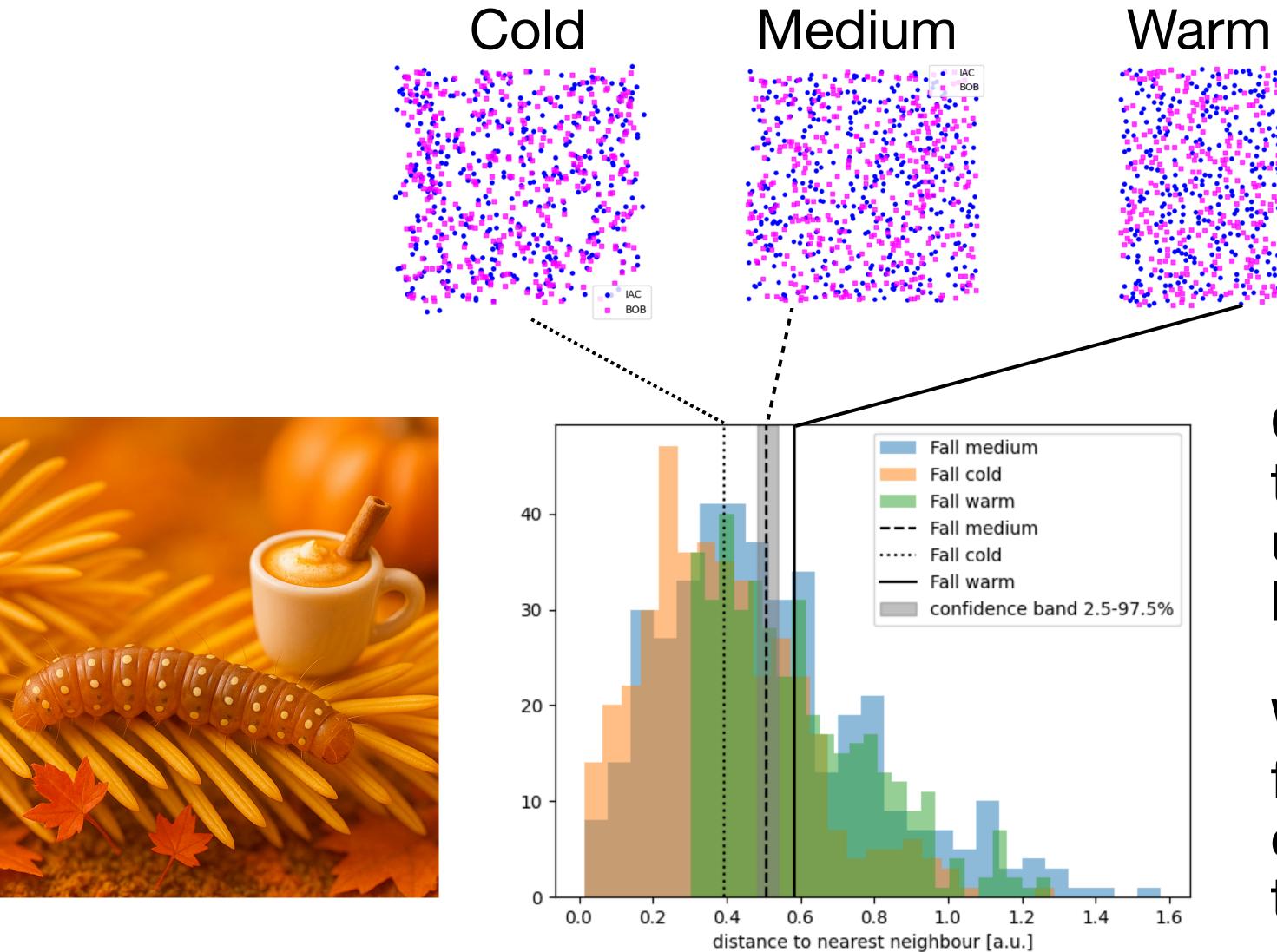








Results: Mean distance IAC -> BOB



Cold IAC->BOB are closer to each other than random under the 95% significance level

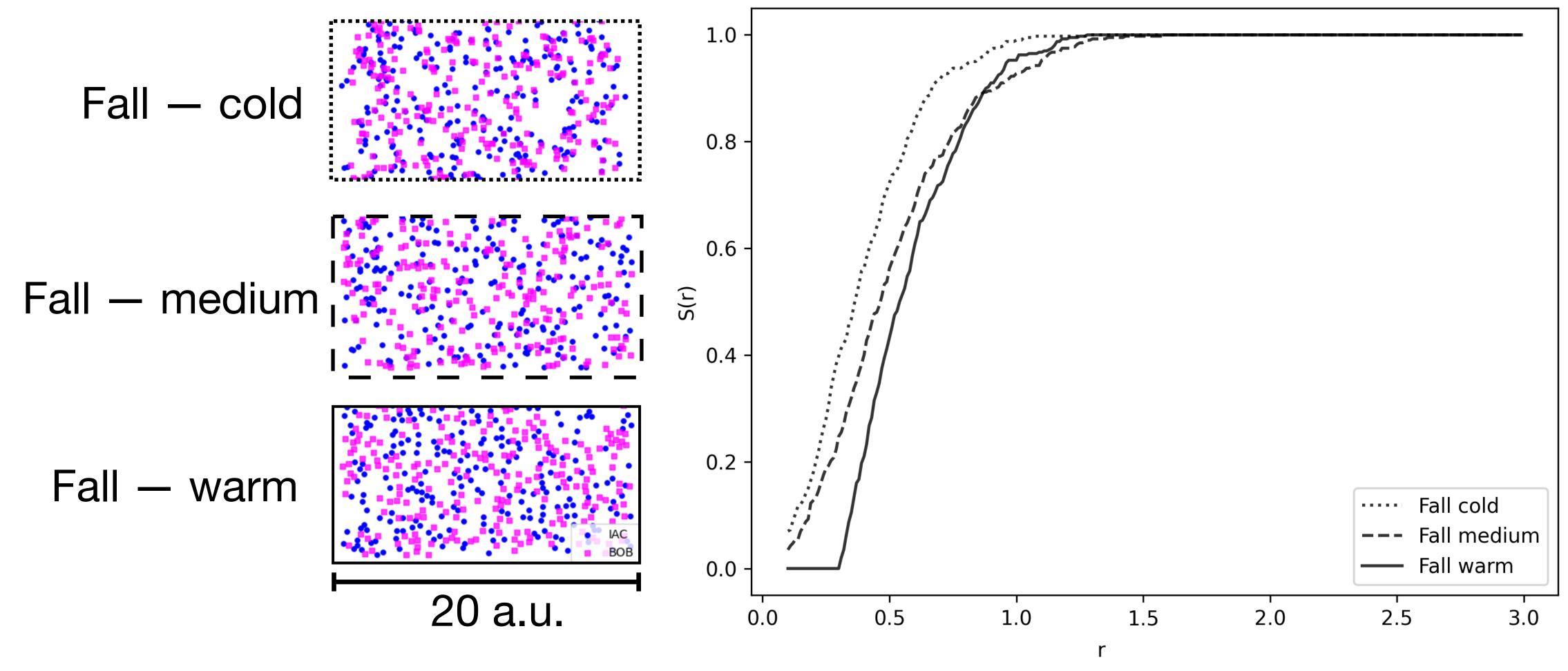
Warm IAC->BOB are further apart from each other than random under the 95% significance level







Results: Nearest neighbor function

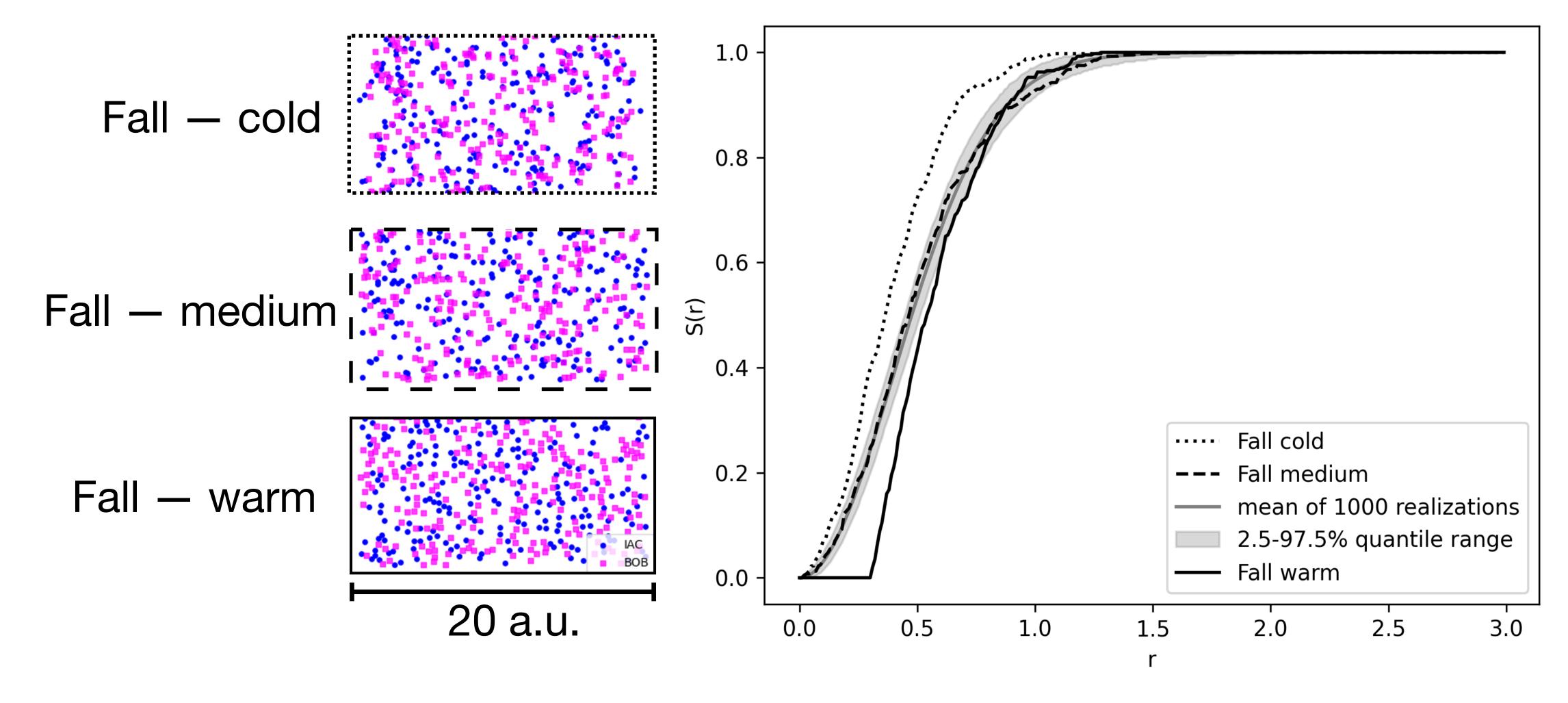








Results: Nearest neighbor function



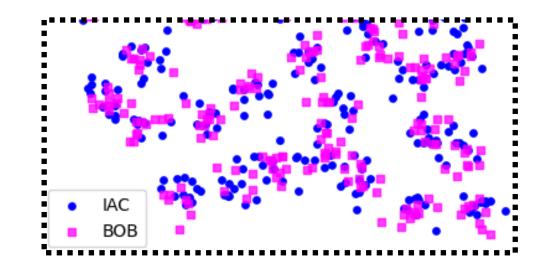






Results: Ripley's K function

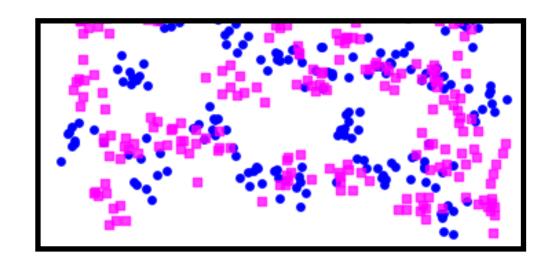
Winter — cold

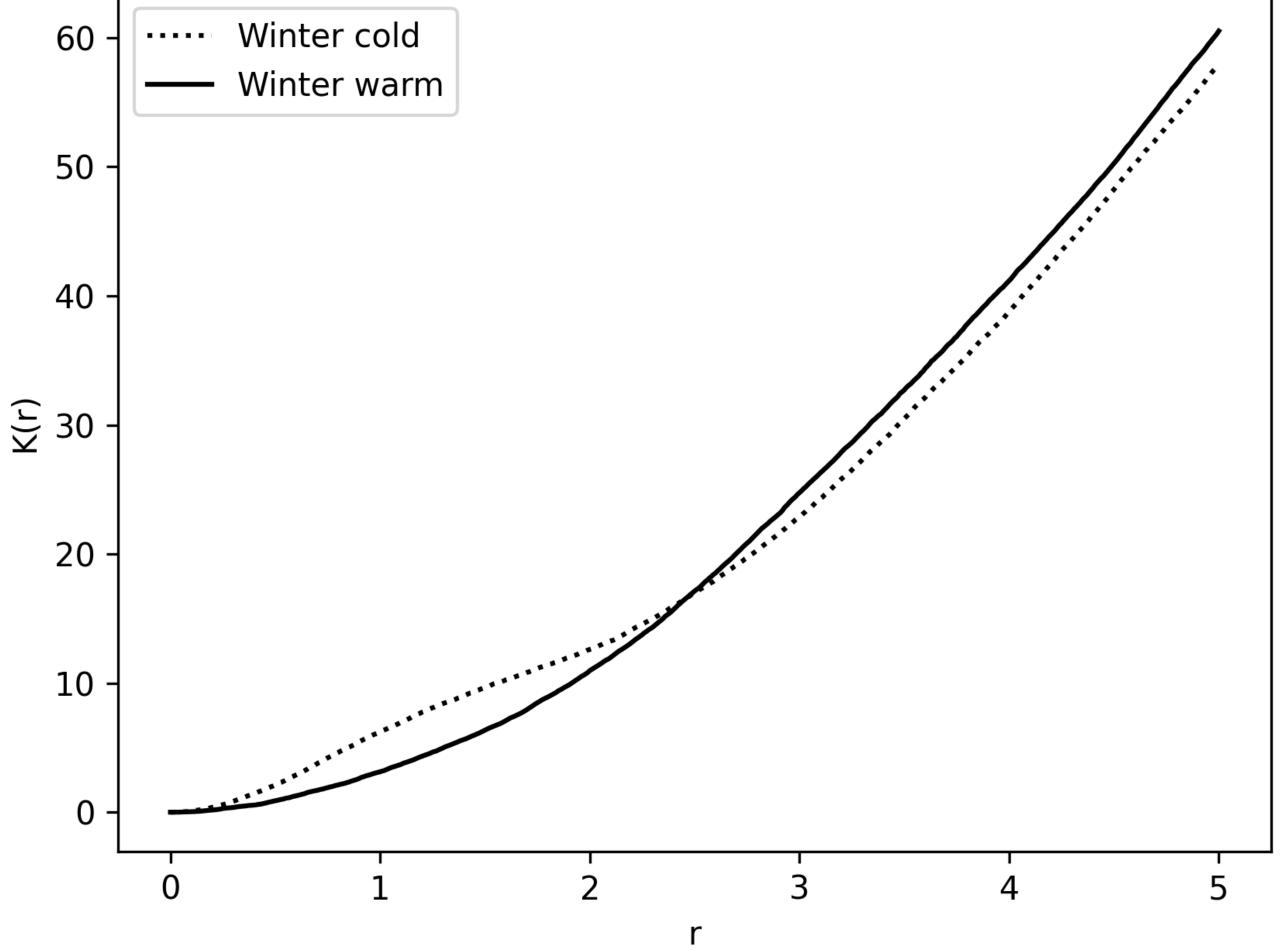


Winter — medium



Winter — warm





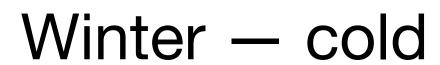


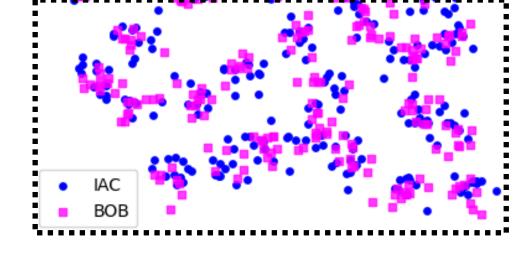




Results: Ripley's K function

Winter cold mean of 1000 realizations 2.5-97.5% quantile range Winter warm





Statistically Statistically 60 significant significant clustering dispersion 50

Winter — medium

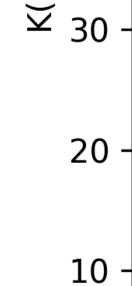


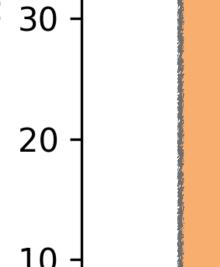


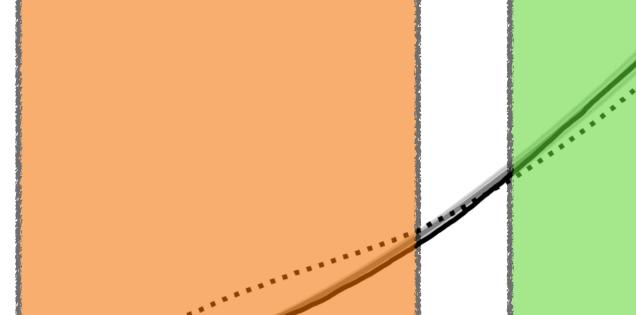


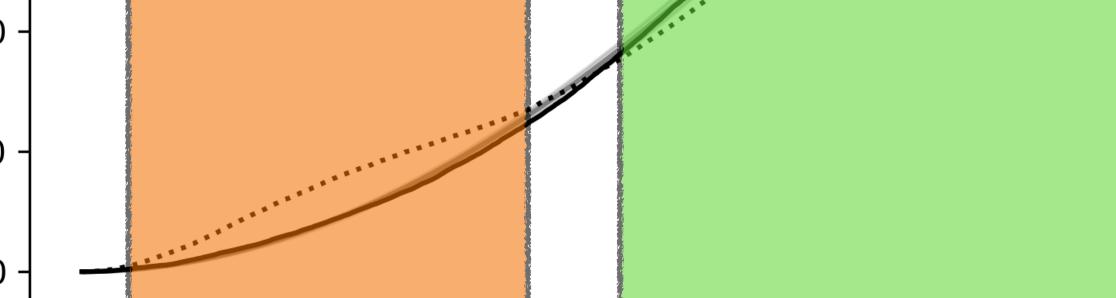


40 -

















The bigger picture: Monte-Carlo-based significance testing

- Just because your sample isn't uniformly distributed doesn't mean it is biologically meaningful!
- It is possible to simulate hypotheses beyond uniform distributions.







Python notes



Our implementation of Ripley's K was tested against the Locan library implementation:

https://locan.readthedocs.io/en/latest/tutorials/notebooks/Analysis_Ripley.html#







References

Lagache T, Sauvonnet N, Danglot L, Olivo-Marin JC. Statistical analysis of molecule colocalization in bioimaging. Cytometry A. 2015 Jun;87(6):568-79. doi: 10.1002/cyto.a.22629. Epub 2015 Jan 20. PMID: 25605428.

Ripley, B. D. "The Second-Order Analysis of Stationary Point Processes." *Journal of Applied Probability*, vol. 13, no. 2, 1976, pp. 255–66. *JSTOR*, https://doi.org/10.2307/3212829. Accessed 13 July 2025.



