

Code documentation

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The present document provides technical documentation for the Python-based routines to compute the organization metrics L_{org} and I_{org} freely available on github at https://github.com/giobiagioli/organization_indices and used in Biagioli and Tompkins (2023).

Given the distribution of a set of objects over a subregion of the plane, the routines compute all-pair separation distances and estimate the corresponding nearest-neighbor cumulative distribution function (NNCDF) and Besag's L -function. The latter is a measure of the mean number of neighbors of any element of the pattern for a range of neighborhood sizes, rescaled by the spatial density of events (Ripley, 1976; Besag, 1977; Ripley, 1979, 1981). The routines also compute the NNCDF and L -function that would be expected theoretically if the *same* number of objects were *randomly* distributed over the same region. The comparison of the theoretical and estimated functions allows to compute indices that assess the level of organization of convective cloud scenes, respectively $I_{\text{org}}/RI_{\text{org}}$ (Tompkins and Semie, 2017; Seifert and Heus, 2013; Biagioli and Tompkins, 2023) and $L_{\text{org}}/dL_{\text{org}}$ (Biagioli and Tompkins, 2023).

The present routines are designed to operate under a broad range of configurations, both in terms of domain geometry and details of the calculation algorithms. They therefore include a set of flags that specify under which conditions the organization degree of a scene is assessed (square/rectangular domains with open/periodic boundaries, use of four-connectivity clustering algorithms, Poisson/binomial models as reference for spatial randomness).

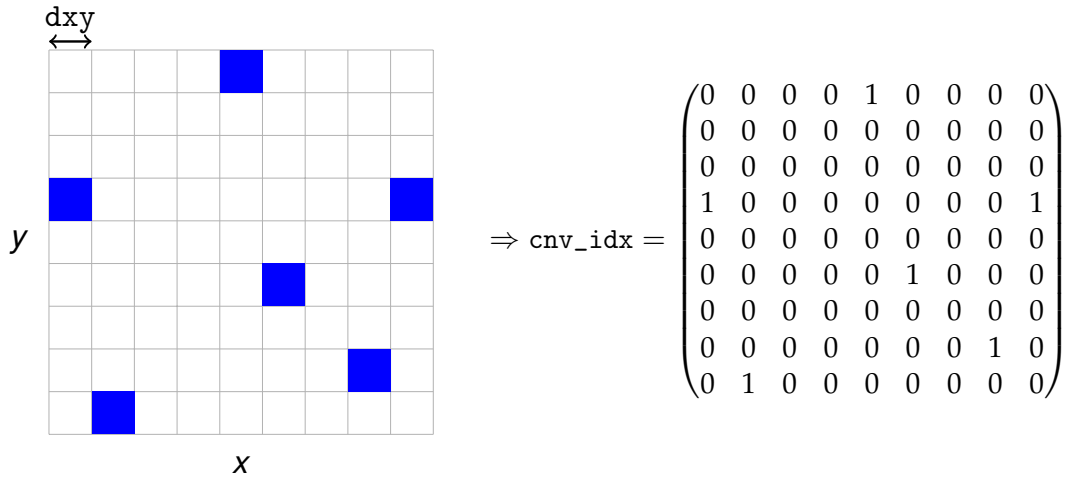
1 Input parameters

The basic syntax of the main routine `_compute_organization_indices` is

```
_compute_organization_indices(dxy, cnv_idx, rmax, bins,
    periodic_BCs, periodic_zonal, clustering_algo,
    binomial_continuous, binomial_discrete, edge_mode)
```

where `dxy`, `cnv_idx`, `rmax`, `bins` are parameters describing the specifics of the gridded dataset and the details of the binning procedure to compute the pair separation distances, and in particular:

- **dx** is the horizontal grid spacing, assumed to be the same in both directions. The user can provide it in units of length (e.g., if the resolution of the dataset is $\Delta x = 2$ km, then dx should be set to 2000) or in grid units (dx = 1). Please note that, in this latter case, the cloud-to-cloud distances as measured by the routine are in grid units as well.
- **cnv_idx** is the 2D binary field of convective (cnv_idx = 1) and non-convective (cnv_idx = 0) grid cells, obtained by applying some specific thresholding approach to relevant variables or resulting from synthetic generation of scenes. The Python object class must be `numpy.ndarray`. In the example below, the convective grid boxes are represented as blue squares and the corresponding binary matrix cnv_idx is shown.



- **rmax** is the maximum search radius (r_{\max} in the manuscript) or box size (ℓ_{\max}) for evaluation of neighbor counts. Typical values for practical cases are provided in section 2. Units must be consistent with those specified in dx.
- **bins** is the array (Python object type `numpy.ndarray`) specifying the rightmost edge of distance/box size bands in which to evaluate the event counts. Typically, bin edges range from 0 to rmax with spacing given by a multiple (or a submultiple) of the grid resolution:

```
bins = np.arange(0, rmax+bin_w*dx, bin_w*dx)
```

where `bin_w` is the parameter that sets the bin width. The user can alternatively use the command `np.linspace`.

For instance, assume the discrete organization index dL_{org} is to be computed for a dataset with spatial resolution $\text{dx} = 1$ km over a doubly-periodic square observation window of size 500 km. The user might want to investigate the spatial organization trends of a cloud pattern over the whole domain, in which case $\text{rmax} = 500\text{e3}$ (units of length). A reasonable choice for the width of box size bins would be $\text{bin_w} \cdot \text{dx} = 2$ km (i.e., $\text{bin_w} = 2$). For any point of the pattern, the

number of neighboring events are counted which are less (or equal) than 2 km, 4 km, 6 km, ..., 500 km from the base point.

The parameters `periodic_BCs` and `periodic_zonal` are boolean and relate to details of the domain geometry. In particular,

- `periodic_BCs` is to specify whether the domain is periodic in both x and y directions, in which case `periodic_BCs = True`.
- `periodic_zonal` is to specify whether the domain is periodic in the zonal direction and open in the meridional direction (`periodic_zonal = True`).

The flags `periodic_BCs` and `periodic_zonal` cannot be simultaneously True, in which case the code returns an error. If the user wants to prescribe open boundary conditions, then both `periodic_BCs` and `periodic_zonal` are to be set False.

For open gridded domains, the parameter `edge_mode` accounts for edge effect correction strategies in the computation of the estimated L -functions. Possible choices are:

- `edge_mode = 'none'`, in which case no correction strategies are employed.
- `edge_mode = 'besag'`, in which case the area-based correction method for edge effects (Besag, 1977) is adopted (cf. Fig. 3c in Biagioli and Tompkins (2023)).

Note that, in case of open domains, the routines are not designed to operate if continuous reference models for randomness (either Poisson or binomial) are assumed, namely, if one wants to consider the case of open boundaries, the flag `binomial_discrete` must be set True (see below). The user can anyway refer to the python class `RipleysKEstimator` in the package `astropy`, that contains built-in functions available for edge correction methods in case of random Poisson processes (see <https://docs.astropy.org/en/stable/stats/ripley.html>).

The parameters `clustering_algo`, `binomial_continuous`, `binomial_discrete` are boolean and are related to the specifics of the cluster identification algorithm and the choice of the theoretical model for random distributions of points (Poisson vs binomial). In detail,

- `clustering_algo` specifies if a four-connectivity clustering algorithm is to be used for identifying adjacent convective cells as a single cluster (`clustering_algo = True`). If False, contiguous convective objects are treated as separate entities.
- `binomial_continuous` uses the continuous binomial model as a standard of randomness if True. Note that `binomial_continuous = True` only works if doubly-periodic boundaries are prescribed. The analytical formulas for the reference random distributions in this case are provided in Biagioli and Tompkins (2023), equations (18) for square domains and (22) for non-square domains.
- `binomial_discrete` uses the discrete binomial model (cf. Fig. 3d in Biagioli and Tompkins, 2023) as a reference for randomness, if True.

Note that the two options `binomial_continuous` and `binomial_discrete` cannot be set simultaneously `True`, with the code returning an error message in this case. If the user wants to consider the Poisson model eqn. (10) for complete spatial randomness, the flags `binomial_continuous` and `binomial_discrete` have to be set `False`.

2 Main example cases

In this section, we first show a possible way of generating random synthetic cloud field scenes for square and rectangular domains and then, assuming both periodic and open boundaries and Poisson and binomial reference random models, we show which input parameters are needed to compute the indices $L_{\text{org}}/dL_{\text{org}}$. The user can create or analyze clustered, regular or composite scenes through input of a proper `cnv_idx` argument.

In order to create a random cloud field scene *without replacement* (i.e., no two objects can occupy the same location), the simple sequence of Python commands reported below can be used, where `nx` and `ny` are the number of grid points in the x and y directions, respectively, and `ncnv` is the number of convective elements in the domain.

```
cnv_idx=np.zeros([ny,nx],dtype=int)
coords=np.random.choice(np.arange(0,ny*nx,1),int(ncnv),
    replace=False)
new_loc=np.unravel_index(coords,(ny,nx))
cnv_idx[new_loc]=1
```

2.1 Square domains

2.1.1 Poisson model for randomness, cyclic boundaries

In this case, the theoretical L -function for spatial randomness is given by $\tilde{L}(r) = r_{\text{max}}^{-1}r$ and the estimated one is eqn. (11) in Biagioli and Tompkins (2023), rescaled by r_{max} and with $w_i = 1$. The user can select the desired maximum search radius r_{max} . Here, and in all the examples shown in Biagioli and Tompkins (2023), r_{max} is the maximum distance between any two points in a periodic square domain of size D , i.e., $r_{\text{max}} = D/\sqrt{2}$. Consequently `bins` is set as `bins = np.arange(0,rmax+dxy,dxy)`, where `dxy` is the dataset resolution. The value of `clustering_algo` is to be specified according to the description above. The other input parameters must be `periodic_BCs = True`, `periodic_zonal = False`, `binomial_continuous = False`, `binomial_discrete = False`, `edge_mode = 'none'`. The result is similar to Fig. 4a in Biagioli and Tompkins (2023).

2.1.2 Continuous binomial model for randomness, cyclic boundaries

In this case, the theoretical and estimated L -functions are given by eqns. (18) and (11) in Biagioli and Tompkins (2023), both normalized by r_{max} and with $w_i = 1$. The maximum

search radius r_{\max} is again $r_{\max} = D/\sqrt{2}$. The other arguments must be `periodic_BCs = True, periodic_zonal = False, binomial_continuous = True, binomial_discrete = False, edge_mode = 'none'`. The result is similar to Fig. 4d in Biagioli and Tompkins (2023).

2.1.3 Discrete binomial model for randomness, cyclic boundaries

In this case, the theoretical and estimated L -functions are given by eqns. (19) and (20) in Biagioli and Tompkins (2023), both normalized by ℓ_{\max} and with $w_i = 1$. In a square periodic domain of size D , ℓ_{\max} (rmax in the routine) is set to $r_{\max} = D$, which is the maximum size of a search box centered on any given object of the pattern. In view of this, a possible choice for bins is `bins = np.arange(0, rmax+2*dxy, 2*dxy)`, where the factor of 2 is due to the fact that now boxes (and not distances) are considered (cf. Figs. 3a and 3d in Biagioli and Tompkins, 2023). The other arguments must be `periodic_BCs = True, periodic_zonal = False, binomial_continuous = False, binomial_discrete = True, edge_mode = 'none'`. The result is similar to Fig. 4g in Biagioli and Tompkins (2023).

2.1.4 Discrete binomial model for randomness, open boundaries

The case of open boundaries is typical of the analysis of observational datasets. The theoretical and estimated L -functions are again given by eqns. (19) and (20) in Biagioli and Tompkins (2023), both normalized by ℓ_{\max} . The weighting factors w_i in the estimated L -function are now imposed as specified in the text (Section 4d). In a square open domain of size D , the maximum search box size is given by $\ell_{\max} = 2D$, hence $r_{\max} = 2D$, but, as usual, the user can restrict the analysis to any subregion of the observation window and select r_{\max} accordingly. The other arguments must be `periodic_BCs = False, periodic_zonal = False, binomial_continuous = False, binomial_discrete = True, edge_mode = 'besag'`. With `edge_mode = 'none'`, the correcting factors $w_i = 1$ and the estimated L -function will exhibit a regularity bias due to event undercounting in the long range.

2.1.5 Discrete binomial model for randomness, zonally cyclic boundaries

In this case, the estimated L -function is given by eqn. (20) in Biagioli and Tompkins (2023), with weights w_i obeying the area-based local correction strategy described in the manuscript. The theoretical L -function is instead

$$L(\ell_n) = \begin{cases} \sqrt{\frac{N_c - 1}{N_c}} (n^2 - 1) \Delta x^2 \sim n \Delta x \equiv \ell_n & \text{for } 0 \leq \ell_n \leq D, \\ \sqrt{\frac{N_c - 1}{N_c}} ((n^2 - 1) \Delta x^2 - n \Delta x (n \Delta x - D)) \sim \sqrt{\ell_n D} & \text{for } \ell_n > D, \end{cases} \quad (1)$$

where D is the size of the square periodic domain and the approximation holds for reasonable sample sizes. Both the theoretical and estimated functions are to be rescaled by

ℓ_{\max} (rmax in the routine), which, when considering the spatial tendencies of a pattern over the entire domain, must be $r_{\max} = 2D$. The other arguments are `periodic_BCs = False`, `periodic_zonal = True`, `binomial_continuous = False`, `binomial_discrete = True`, `edge_mode = 'besag'` (recall that the domain is open in the meridional direction).

2.2 Non-square domains

Given a rectangular region of width D_x and height D_y , in the sequel without loss of generality we will assume $D_x > D_y$, but the routines are designed to operate also in the other case.

2.2.1 Poisson model for randomness, cyclic boundaries

The theoretical and estimated L -functions are eqns. (10) and (11) in Biagioli and Tompkins (2023), both normalized by r_{\max} and with $w_i = 1$. r_{\max} is the maximum distance between any two points in a cyclic domain of width D_x and height D_y i.e., $r_{\max} = \sqrt{D_x^2 + D_y^2}/2$. The other input parameters are as in section 2.1.1.

2.2.2 Continuous binomial model for randomness, cyclic boundaries

In this case, the theoretical and estimated L -functions are given by eqns. (22) and (11) in the manuscript, normalized by r_{\max} and with $w_i = 1$. r_{\max} is again set to $r_{\max} = \sqrt{D_x^2 + D_y^2}/2$. The other arguments are as in section 2.1.2.

2.2.3 Discrete binomial model for randomness, cyclic boundaries

In this case, the theoretical and estimated L -functions are given by eqns. (23) and (20) in Biagioli and Tompkins (2023), normalized by ℓ_{\max} and with $w_i = 1$. In a rectangular periodic domain of width D_x and height D_y , the maximum size of a box for neighbor search that is centered on any given object of the pattern is $\ell_{\max} = \max\{D_x, D_y\}$ and rmax is chosen accordingly in the specification of the input parameters. The other arguments, including bins, are as in section 2.1.3.

2.2.4 Discrete binomial model for randomness, open boundaries

The theoretical and estimated L -functions are provided by eqns. (19) and (20) in Biagioli and Tompkins (2023), both normalized by ℓ_{\max} . The weighting factors w_i in the estimated L -function are computed according to the area-based edge-correction strategy (cf. Section 4d in the manuscript) if `edge_mode = 'besag'`. In a rectangular open domain of width D_x and height D_y , the maximum search box size is $\ell_{\max} = 2 \max\{D_x, D_y\}$, and rmax in the routine can be selected as such. However, the user can restrict the analysis to any subregion of the observation window and choose the desired rmax. The other arguments must be as in section 2.1.4.

2.2.5 Discrete binomial model for randomness, zonally cyclic boundaries

The estimated L -function is given by eqn. (20) in Biagioli and Tompkins (2023) rescaled by ℓ_{\max} and with weighting factors w_i calculated according to the area-based correction technique for edge effects. Regarding the computation of the theoretical L -function, distinct cases have to be considered:

- if $D_x \geq 2D_y$, the maximum search box size is $r_{\max} = D_x$ and, since the search boxes are entirely contained within the domain, no corrections for multiple counting due to the periodic continuation of the domain in the zonal direction have to be imposed. The theoretical L -function is therefore provided by eqn. (19) in the manuscript, rescaled by ℓ_{\max} (r_{\max} in the code).
- if $D_x < 2D_y$ (or if $D_y > D_x$), the maximum search box size is $r_{\max} = 2D_y$ and the search boxes can exceed the domain in the zonal direction, therefore a correction must be included to avoid multiple counting. Using simple arguments from the stochastic geometry, analogous to those used in Biagioli and Tompkins (2023), we have

$$L(\ell_n) = \begin{cases} \sqrt{\frac{N_c - 1}{N_c}} (n^2 - 1) \Delta x^2 \sim n \Delta x = \ell_n & \text{for } 0 \leq \ell_n \leq D_x, \\ \sqrt{\frac{N_c - 1}{N_c}} (n \Delta x D_x - \Delta x^2) \sim \sqrt{\ell_n D_x} & \text{for } \ell_n > D_x. \end{cases} \quad (2)$$

In any case, the other input parameters must be as in section 2.1.5.

3 Output

The main routine `_compute_organization_indices` returns the values of the indices I_{org} , RI_{org} and $L_{\text{org}}/dL_{\text{org}}$ and also the theoretical and estimated NNPDFs and L -functions to allow visual inspection of their profiles. In detail, the output consists of:

- `I_org` and `RI_org`, values of the I_{org} and RI_{org} indices as defined in Biagioli and Tompkins (2023), eqns. (2) and (3).
- `L_org`, value of the index L_{org} or dL_{org} depending on whether a continuous (Poisson/binomial) or discrete (binomial) model is assumed as a standard of randomness.
- `NNCDF_theor` and `NNCDF_obs`, theoretical and estimated nearest-neighbor cumulative distribution functions. The theoretical one is the Weibull cumulative distribution function eqn. (1) in Biagioli and Tompkins (2023), the observed one is derived from the distribution of the objects in the given scene.
- `Besag_theor` and `Besag_obs`, theoretical and estimated Besag's L -functions, computed as discussed in section 2 for the different cases.

References

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