Pseudo-code for simulating the Snapshot model

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Reminder: The method here implemented is the one described in general terms in Carrizo Vergara et al. (2024, Section 6.1). The underlying trajectory model is supposed to depend upon a parameter θ_X . The R function present in the repository is a vectorized implementation of this code, which can only be applied for square snapshot regions.

Algorithm 1 Algorithm to simulate the Snapshot model

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Input: N, \theta_X, p, the times of snapshot takes t_1, ..., t_n, and for every k = 1, ..., n, the disjoint regions A_{k,1}, ..., A_{k,m_k} over which the snapshots are taken at time t_k.

1: for j = 1, ..., N do

2: Simulate the individual trajectory (x_j(t_1), ..., x_j(t_n)) \sim \mu_{X(t_1), ..., X(t_n)|\theta_X}.

3: end for
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- 4: **Define** $Q = (\vec{Q}_k)_{k \in \{1,...,n\}}$ a list of n null vectors, each vector $\vec{Q}_k = (Q_{k,1},...,Q_{k,m_k})$ being of dimension m_k (Q can be a null matrix if $m_1 = ... = m_k$).
- 5: **for** k = 1, ..., n **do**6: **for** $l = 1, ..., m_k$ **do**7: $Q_{k,l} \leftarrow \sum_{j=1}^{N} \mathbb{1}_{A_{k,l}}(x_j(t_k))$ 8: $Q_{k,l} \leftarrow Binomial(Q_{k,l}, p)$
- 9: end for
- 10: **end for**
- 11: return Q

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References

Carrizo Vergara, R., Kéry, M., & Hefley, T. (2024). Movement-based models for abundance data. *arXiv* preprint arXiv:2407.13384.