

9th SSIAB Workshop, Avignon - May 9-11, 2012 Testing of mark independence for marked point patterns

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Outline

The talk is based on the paper

P. Grabarnik, M. Myllymäki and D. Stoyan (2011). Correct testing of mark independence for marked point patterns. *Ecological Modelling* 222, 3888–3894.

and discusses

- the conventional envelope test
- the refined envelope test
- the deviation test

through two marked point pattern data examples.



Tharandter Wald: These data observed in a 56 m \times 38 m rectangle come from a Norway spruce forest in Saxony (Germany).



Frost shake of oaks: 392 oak trees observed in a 100 m \times 100 m square at Allogny in France (Courtesy to Goreaud & Pelissier, 2003).



Circles are proportional to the diameters of trees at breast height (=marks).

White circle = 1, a sound oak; Black circle = 2, an oak suffering from frost shake



Our question here:

are the marks independently assigned for the points in an originally non-marked point pattern?

"Random labeling" hypothesis



How is the hypothesis typically tested?

Monte Carlo significance tests (Besag and Diggle, 1977)

- makes s = 99 simulations under the null hypothesis (How?)
- chooses a summary function F(r) and calculated its estimate F(r) for data and each simulated marked point pattern
- Then either 1) calculates the minimum and maximum for each r in [r_{min}, r_{max}]

$$F_{up}(r) = \max_{i=2,\dots,s+1} \hat{F}_i(r),$$

$$F_{low}(r) = \min_{i=2,\dots,s+1} \hat{F}_i(r).$$

and compared the data function to the envelopes, or,

2) summarizes the information contained in the functional summary statistic F(r) into a scalar test statistic

Consider first 1)!





The summary function?

Here the summary functions

$$L_{mm}(r) = \sqrt{rac{K_{mm}(r)}{\pi}}$$
 (Tharandter Wald data)

and

$$L_{12}(r) = \sqrt{\frac{K_{12}(r)}{\pi}}$$

(Frost shake of oaks data)

are used, which both are generalizations of Ripley's *K*-function to marked or bivariate point patterns.



Envelopes for the Tharandter Wald data

s = 99



Conclusions?



Problem of the envelope test

The spatial correlations are inspected for a range of distances simultaneously.

- ► Ripley (1977)
 - introduced envelope tests
 - mentioned that the frequence of committing the type I error in the envelope test may be higher than for a single distance test
- Diggle (1979, 2003)
 - proposed the deviation test
- Loosmore and Ford (2006)
 - adopted the deviation test
 - demonstrated the multiple testing problem of envelope test by estimating the type I error probability by simulation for the complete spatial randomness hypothesis based on the nearest neighbour distance distribution function
 - rejected the envelope test



Envelopes for the Tharandter Wald data

s= 99, type I error approximation pprox 0.48



Conclusions?



Type I error approximation?



In the case of minimum and maximum envelopes, the type I error is approximated by t/s where

- t is the number of those simulations that take part in forming the envelopes
- s is the total number of simulations



Towards the refined envelope test

A natural way to make the envelope method valid, i.e. to obtain a reasonable type I error, is to increase the number of simulations from which the envelopes are calculated.



Envelopes for the Tharandter Wald data

s = 1999, type I error approximation ≈ 0.04



Conclusions?



The refined envelope test

The refined testing procedure = the envelope test, where

- the type I error probability is evaluated and taken into account in making conclusions
- if the choice of the number of simulations s leads to an unacceptably large type I error, s can be increased so that the type I error comes close to a desired value

The refined envelope test is then a rigorous statistical tool.



Deviation test

A deviation test

• summarizes information on F(r) into a *single* number

$$u_i = \max_{r_{\min} \leq r \leq r_{\max}} |\hat{F}_i(r) - F_{H_0}(r)|,$$

$$u_i = \int_{r_{\min}}^{r_{\max}} (\hat{F}_i(r) - F_{H_0}(r))^2 dr,$$

- is based on the rank of the data statistic
- provides the exact type I error probability, i.e. the null hypothesis is declared false, when it is true, precisely with the prescribed probability (Barnard, 1963; Besag &Diggle, 1977)



The data example 1

s = 1999, type I error approximation \approx 0.04



Max-deviation: $\hat{p} = 0.31$; Int-deviation: $\hat{p} = 0.20$. Conclusions?



Frost shake of oaks: 392 oak trees observed in a 100 m \times 100 m square at Allogny in France (Courtesy to Goreaud & Pelissier, 2003).



White circle = 1, a sound oak; Black circle = 2, an oak suffering from frost shake



Earlier studies:

Goreaud & Pelissier (2003) and Illian et al. (2008):

- ▶ used the *L*₁₂-function and the envelope test
- ► G & P: 0.5%-lower and -upper envelopes based on s = 10000 simulations
- Illian et al.: minimum and maximum envelopes from s = 99 simulations
- came to the conclusion to reject the random labeling hypothesis



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Type I error approximation: 1) 0.21 2) 0.41



s= 999, type I error approximation pprox 0.04



Conclusions?



s= 999, type I error approximation pprox 0.04



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Discussion

The deviation test

- + do not need so many simulations
- + p-values can be easily estimated
- + different forms
- says only little about the reason of rejection
- says nothing on the scales at which there is behavior of *F*(*r*) leading to rejection
- ► performance depends on the behavior of the variance of *F*(*r*) over the range of chosen distances (→ more sophisticated edge correction methods, Ho & Chiu, 2006)



Discussion

The refined envelope test

- + help to detect reasons why the data contradict the null hypothesis (important when ecologists seek for alternative hypothesis!)
- ► + also raw estimators can be used (as long as the same estimator is used for $F_1(r)$ and $F_i(r)$, i = 2, ..., s + 1
- needs many simulations
- -(?) no p-values

We recommend to couple formal testing with diagnostic tools using non-cumulative functions.



References

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Thank you!

