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Analysing spatial and temporal variability in diversity

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Diversity in diversity...

HOW STANDARDS PROLIFERATE:
(SEE: A/C CHARGERS, CHARACTER ENCODINGS, INSTANT MESSAGING, ETC.)





- Partitioning Diversity
 - Alpha, Beta and Gamma diversity
- Barro Colorado Island
 - Spatial Partitioning
 - Temporal Partitioning



Different kinds of diversity

- Diversity, even biodiversity, can mean lots of different things to different people
- Individual diversity measures can reflect many different aspects of this
- As a result measures can often disagree with one another about what is diverse
- This is often correct, but we need to understand what each measure is telling us and which measure (if any) answers the question we are interested in



Partitioning Diversity

- What do we mean when we talk about partitioning diversity?



Partitioning Diversity

- Imagine we have divided up our ecosystem into N sub-communities
- We want to know:
 - How many distinct parts are there?
 - Which are the interesting ones?
 - Which should we conserve?



Partitioning Diversity

- How do we divide up the “total” diversity into the diversity “within” the sub-communities, and the diversity “between” the sub-communities?
 - This is usually described as *partitioning* the *gamma* (global) diversity into *alpha* (average) and *beta* (between) components



Partitioning Diversity

\bar{p}_i	$\left(\sum_{i=1}^S p_i^q\right)^{1/(q-1)}$	$\sqrt[q-1]{\sum_{i=1}^S p_i p_i^{q-1}}$	mean proportional species abundance; the weighted generalised mean with exponent $q-1$ of the p_i values
${}^q\lambda_\gamma$	$\sum_{i=1}^S p_i^q$	\bar{p}_i^{q-1}	basic sum of order q in relation to the γ -classification: the sum of p_i^q values over all species i , or mean proportional species abundance raised to the power $q-1$
qD	$1 / \bar{p}_i$	${}^q\lambda_\gamma^{1/(1-q)}$	true diversity of order q : the inverse of mean proportional abundances of the types of interest, or the numbers equivalent of ${}^q\lambda_\gamma$
H'	$\log({}^1D)$	$\log(1 / \bar{p}_i)$	Shannon entropy: the logarithm of true diversity of order 1, or the logarithm of the inverse of the geometric mean of the proportional abundances of the types of interest
γ'			the raw value of a species diversity index as calculated using the entire dataset, e.g. ${}^q\lambda_\gamma$ or H'
α'			the alpha component obtained when partitioning γ'
β'			the beta component obtained when partitioning γ'
${}^qD_\gamma$	$\gamma = 1 / \bar{p}_i$	${}^q\lambda_\gamma^{1/(1-q)}$	true gamma diversity: total effective number of species in the dataset (measurement unit: effective species or sp_E)
$\bar{p}_{(ij)}$	$\sqrt[q-1]{\sum_{i=1}^S p_{ij} p_{ij}^{q-1}}$		mean proportional species abundance within sampling unit j ; the weighted generalised mean with exponent $q-1$ of the p_{ij} values corresponding to sampling unit j
${}^qD_{\gamma j}$	$1 / \bar{p}_{(ij)}$	$\gamma_j = {}^q\lambda_{\gamma j}^{1/(1-q)}$	gamma diversity (= effective number of species) within sampling unit j (measurement unit: sp_E)
$\bar{p}_{(ij)all}$	$\sqrt[q-1]{\sum_{j=1}^N \sum_{i=1}^S p_{ij} p_{ij}^{q-1}}$		mean proportional species abundance in the dataset; the weighted generalised mean with exponent $q-1$ of all p_{ij} values
α_t	${}^q\bar{D}_{\gamma j} = \bar{\gamma}_j$	$1 / \bar{p}_{(ij)all}$	mean species diversity within sampling units: weighted generalised mean with exponent $1-q$ of the ${}^qD_{\gamma j}$ values with w_j used as weights (measurement unit: sp_E)
α_d	${}^qD_\alpha$	α_t / CU	true alpha diversity: effective number of species per virtual sampling unit of mean species diversity, or per compositional unit (measurement unit: sp_E / CU)
α_R	${}^qD_{\gamma\omega} / \omega$	${}^qD_{\gamma\omega} / {}^qD_\omega$	effective number of species abundance values per effective sampling unit (measurement unit: $sp_E SU_E / SU_E$)
β_{Md}	${}^qD_\beta = {}^qD_\gamma / {}^qD_\alpha$	γ / α_d	true beta diversity: number of compositional units in the dataset (measurement unit: CU)
β_{Mt}	${}^qD_{\gamma/\gamma_j} = {}^qD_\gamma / {}^q\bar{D}_{\gamma j}$	γ / α_t	regional-to-local diversity ratio (measurement unit: sp_E / sp_E)
β_R	${}^qD_\gamma / {}^qD_\omega / {}^qD_{\gamma\omega}$	γ / α_R	two-way diversity ratio (measurement unit: $sp_E SU_E / sp_E SU_E$)
β_{At}	${}^qD_\gamma - {}^q\bar{D}_{\gamma j}$	$\gamma - \alpha_t$	regional diversity excess; absolute effective species turnover (measurement unit: sp_E)
β_{Mt-1}	$\gamma / \alpha_t - 1$	$(\gamma - \alpha_t) / \alpha_t$	Whittaker's species turnover: effective species turnover expressed in multiples of the species diversity in a single compositional unit (measurement unit: sp_E / sp_E)
β_{Pt}	$1 - \alpha_t / \gamma$	$(\gamma - \alpha_t) / \gamma$	proportional species turnover: effective species turnover expressed as a proportion of total species diversity (measurement unit: sp_E / sp_E)
H'_β	$H'_\gamma - H'_\alpha$	$\log({}^1\beta_{Md}) = \log(\gamma) - \log(\alpha_d)$	beta Shannon entropy (measurement unit: depends on the base of the logarithm)
$\bar{H}'_{\gamma-\gamma_j}$	$H'_\gamma - \bar{H}'_{\gamma_j}$	$\log({}^1\beta_{Mt}) = \log(\gamma) - \log(\alpha_t)$	regional Shannon entropy excess (measurement unit: depends on the base of the logarithm)
${}^2\bar{\lambda}_{\gamma-\gamma_j}$	${}^2\bar{\lambda}_{\gamma_j} - {}^2\lambda_\gamma$	$(\gamma - \alpha_t) / \gamma \alpha_t$	regional variance excess (measurement unit: sp_E / sp_E^2)



Partitioning Diversity

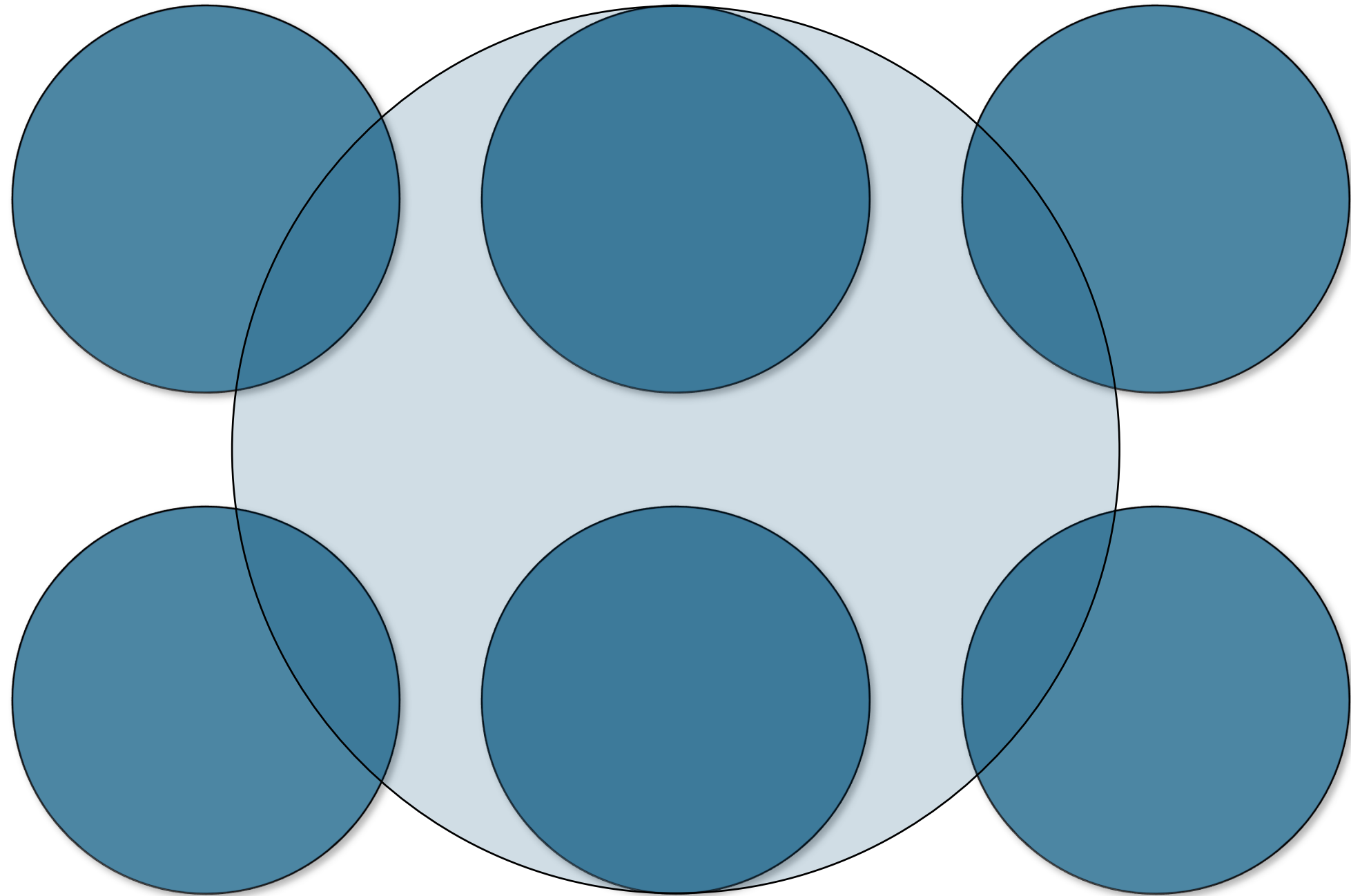
- What do we mean when we talk about beta diversity?
 - Very broadly there are two definitions...



- What do we mean when we talk about beta diversity?
 1. **Effective number of unique communities**



ALPHA DIVERSITY: 100

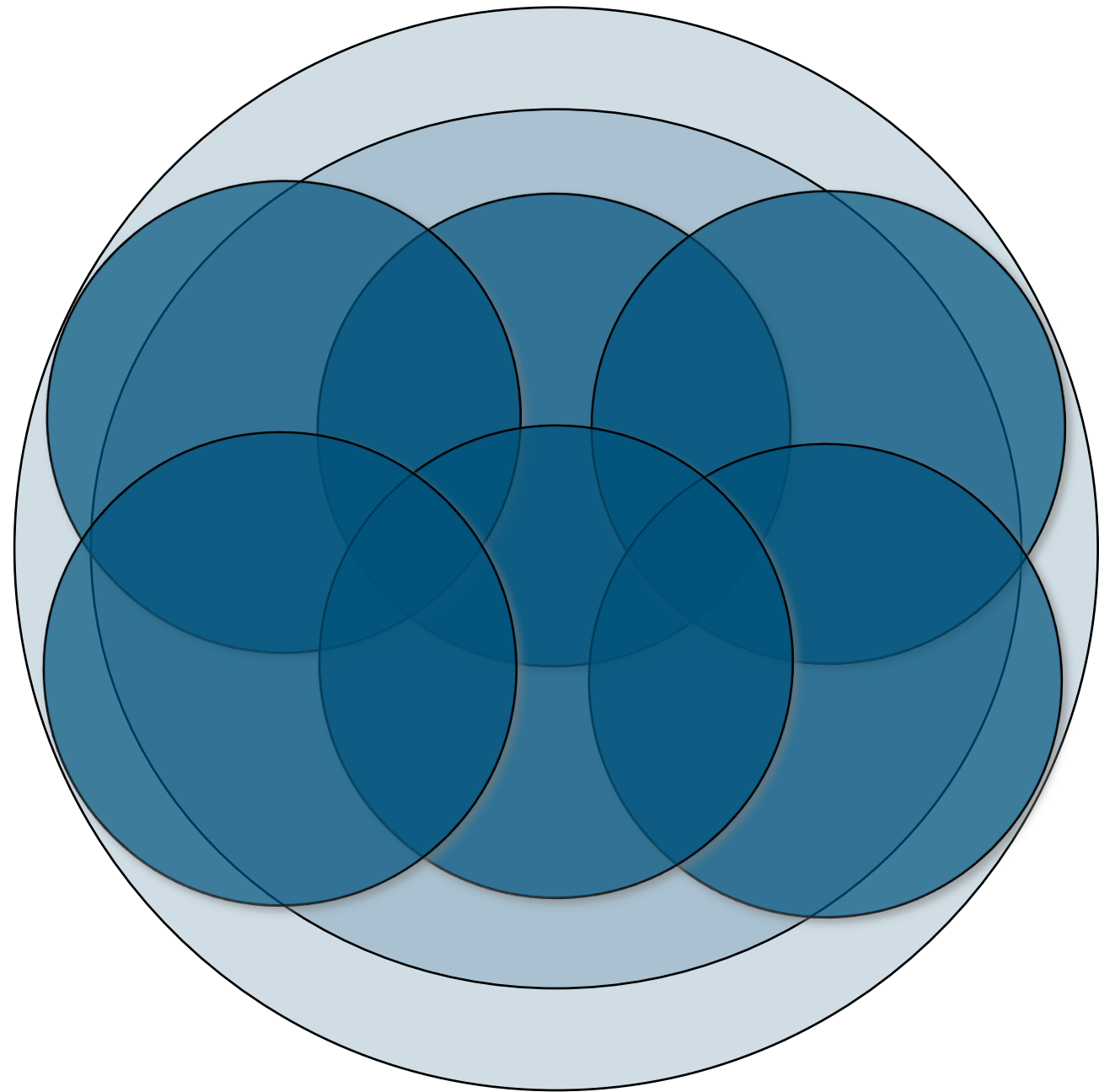


GAMMA DIVERSITY: 600



ALPHA DIVERSITY: 100

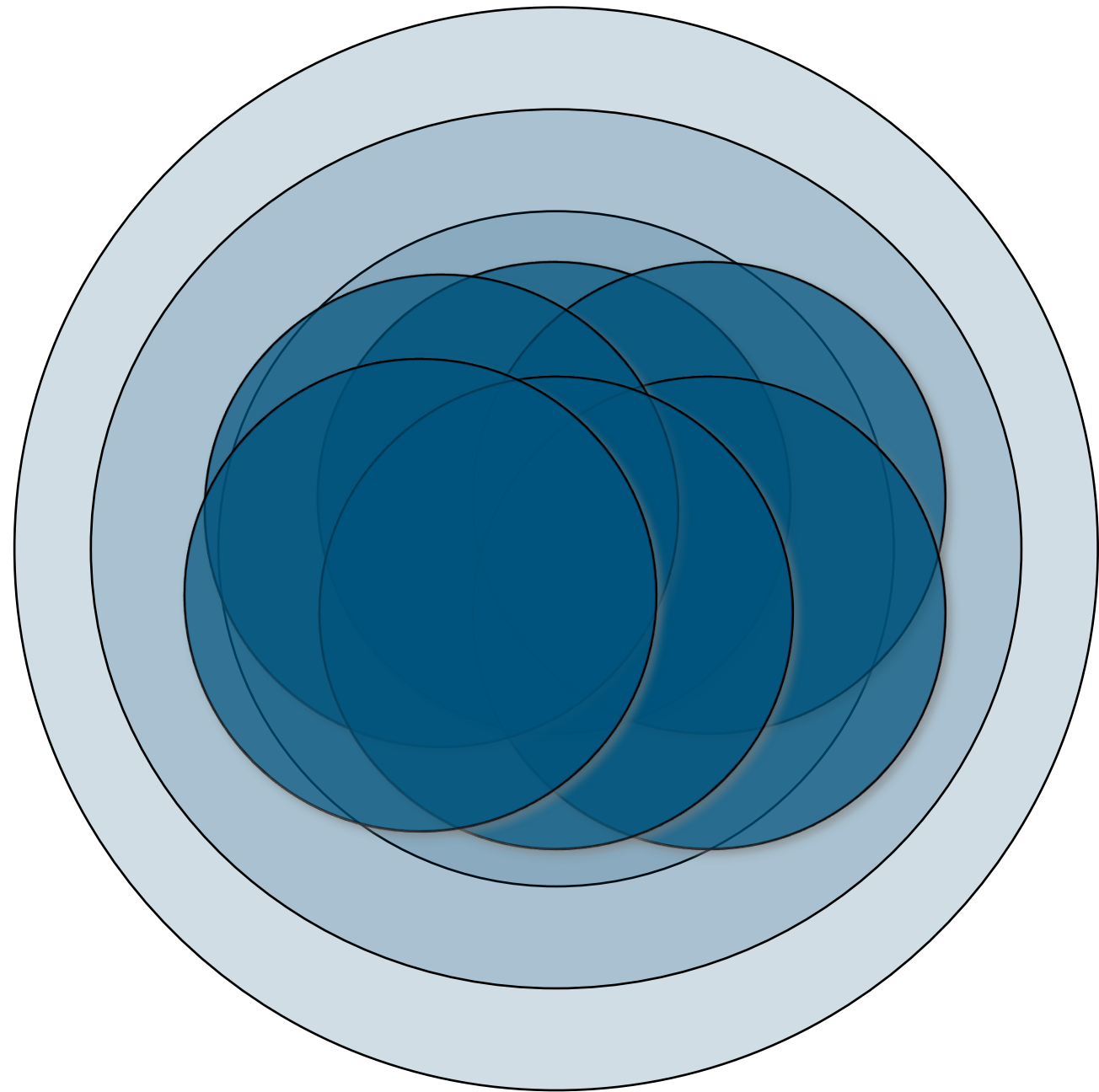
GAMMA DIVERSITY: 400





ALPHA DIVERSITY: 100

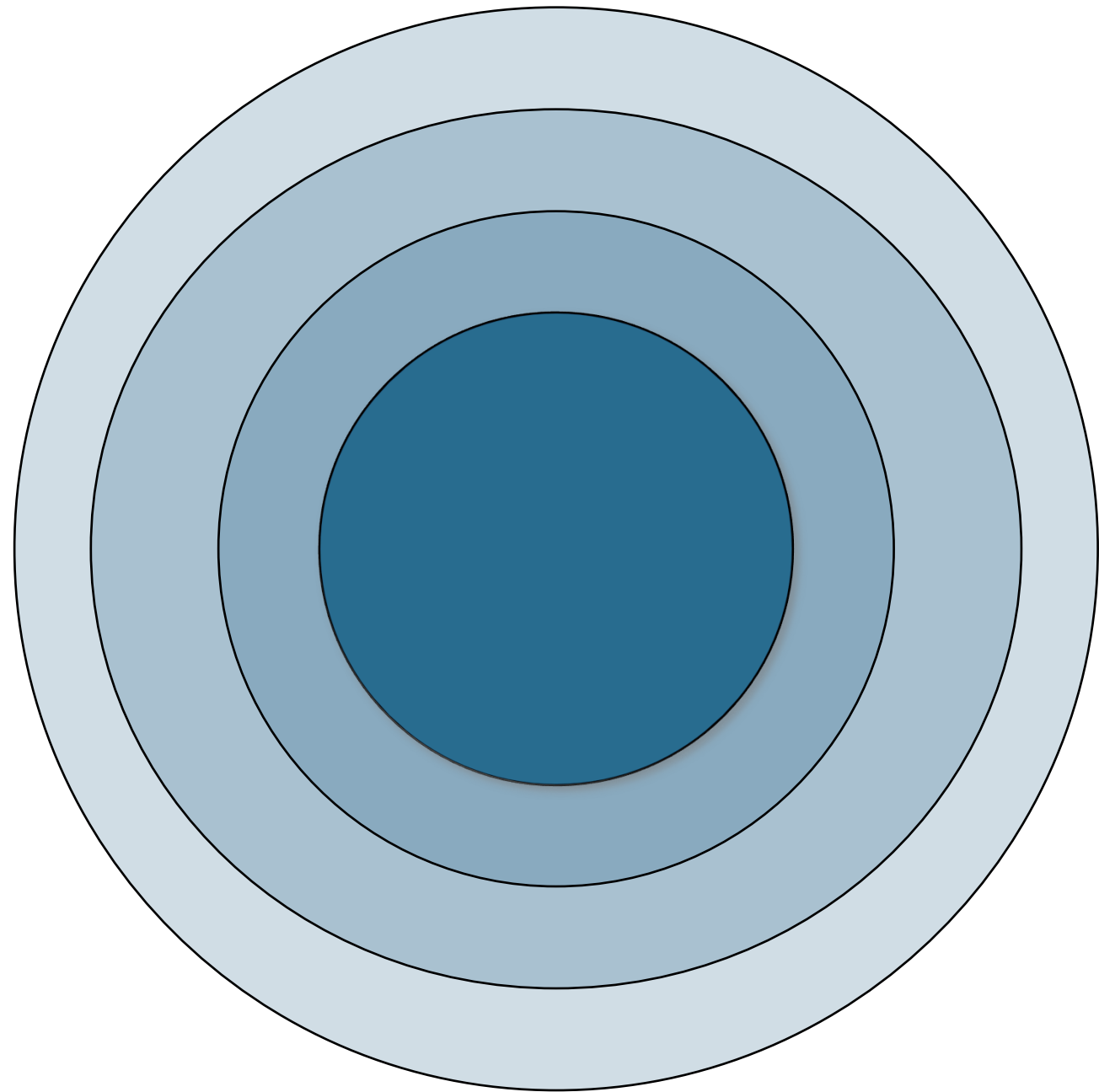
GAMMA DIVERSITY: 200





ALPHA DIVERSITY: 100

GAMMA DIVERSITY: 100



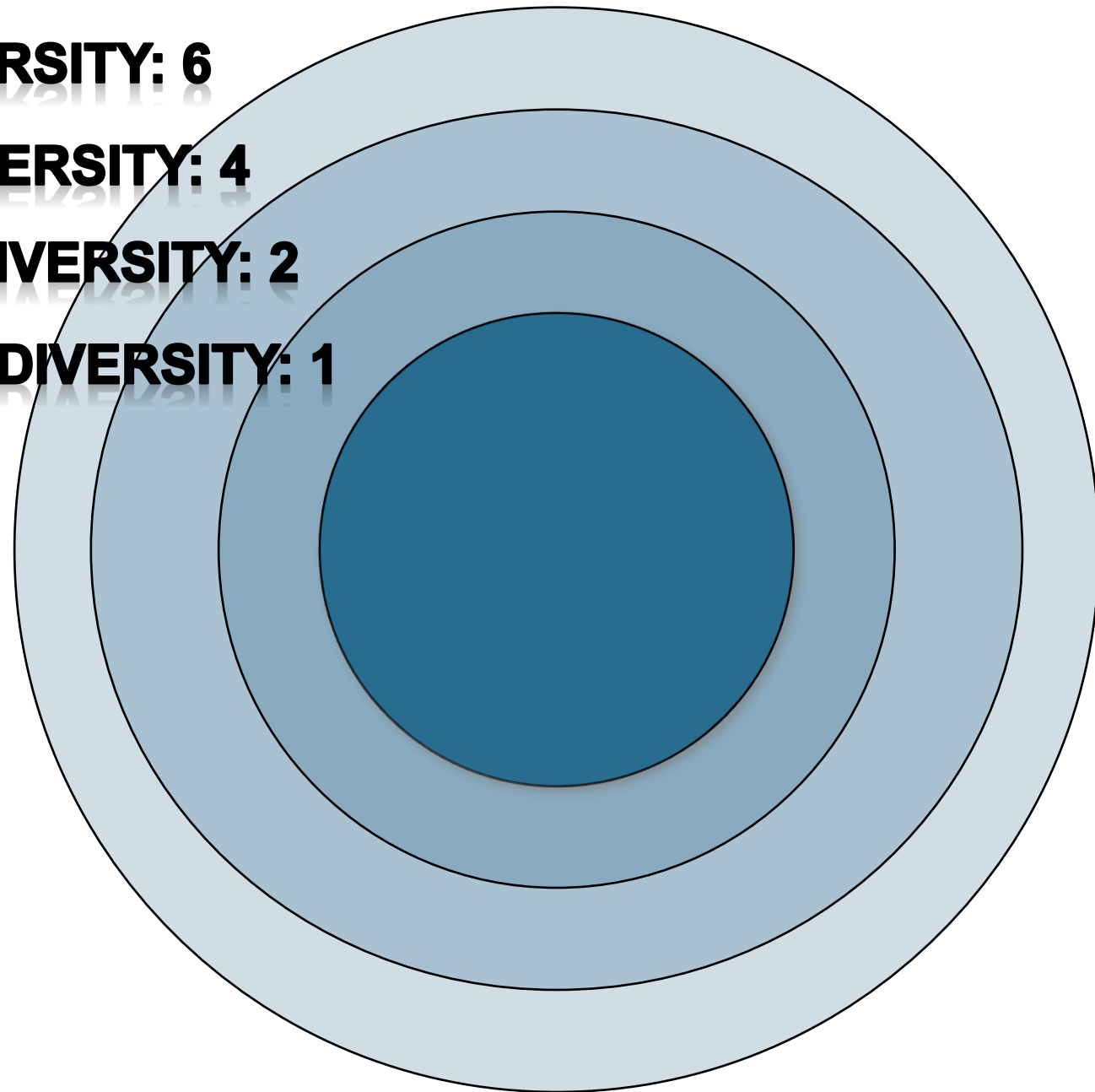


BETA DIVERSITY: 6

BETA DIVERSITY: 4

BETA DIVERSITY: 2

BETA DIVERSITY: 1





Partitioning Diversity

- What do we mean when we talk about beta diversity?
 1. Effective number of unique communities
 2. **Turnover between communities**



Partitioning Diversity



TURNOVER: 100%



Partitioning Diversity

TURNOVER: 80%



Partitioning Diversity

TURNOVER: 60%



Partitioning Diversity

TURNOVER: 40%



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Partitioning Diversity

TURNOVER: 20%



Partitioning Diversity

- What do we mean when we talk about beta diversity?
 1. Effective number of unique communities
 2. Turnover between communities
- These turn out to be the same thing...

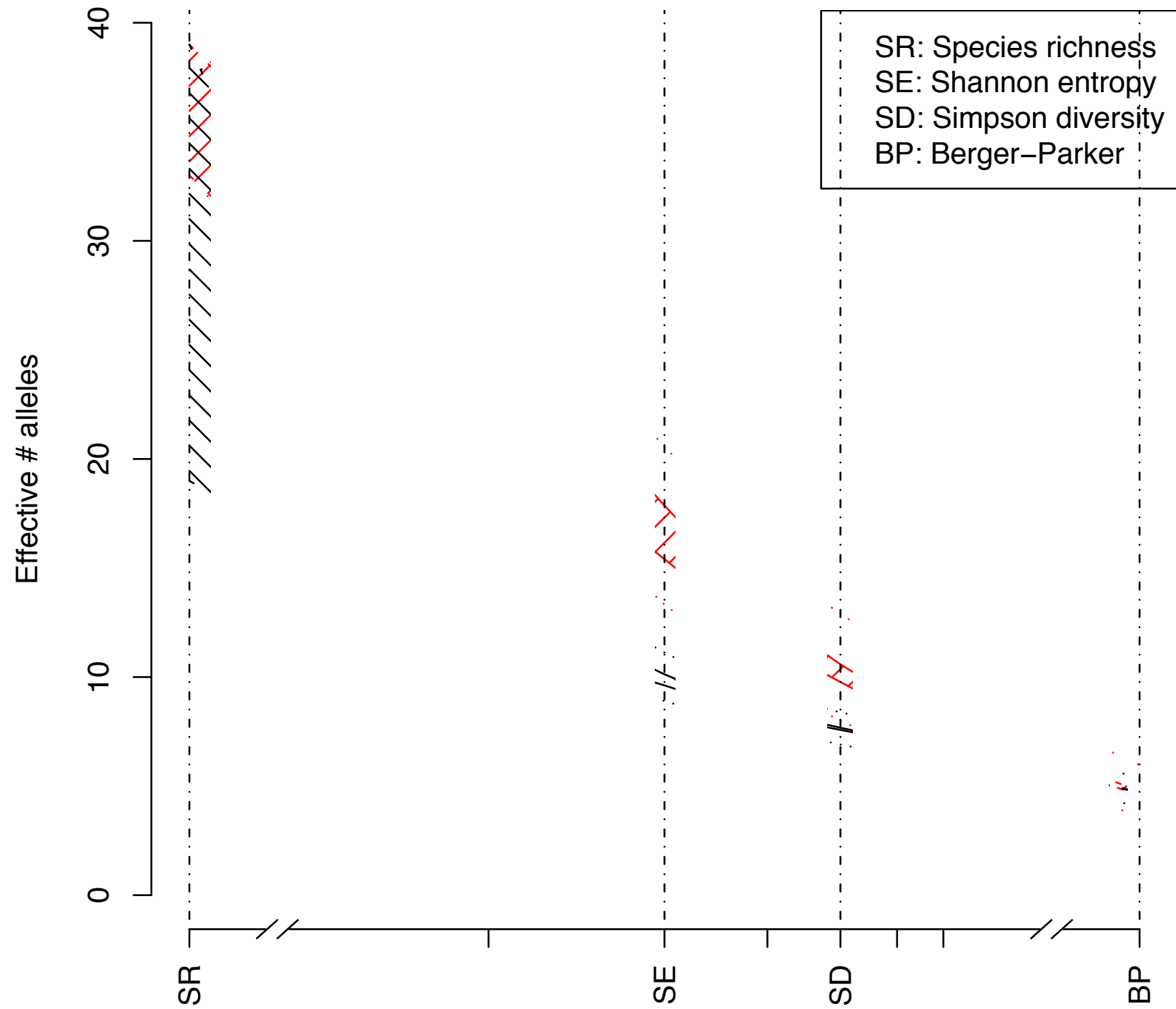


The effective number of species:

- the number of species when all species are present at equal frequency
- the more uneven the distribution, the fewer species are “effectively” present

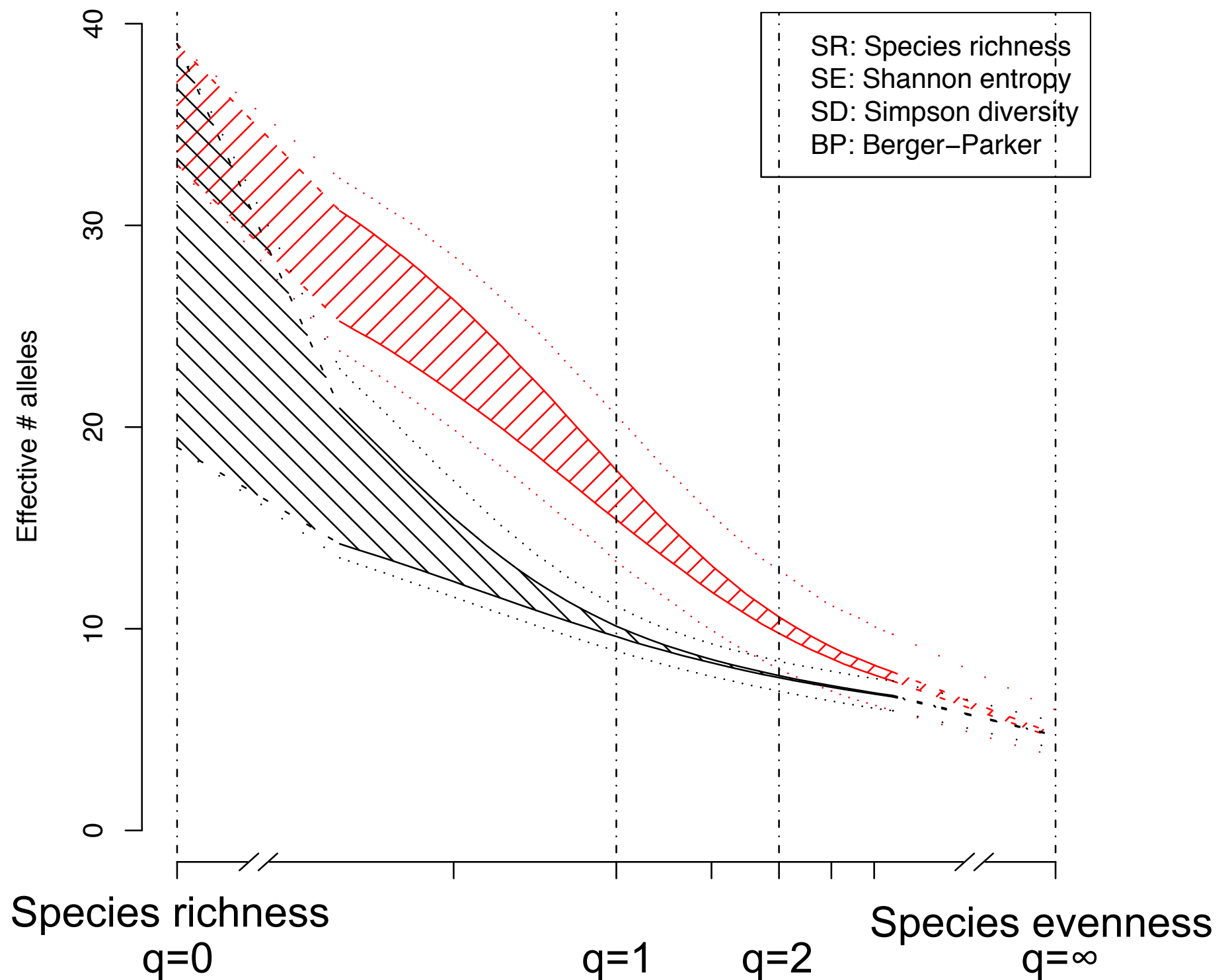


Diversity Profiles





Diversity Profiles





The effective number of species:

- the number of species when all species are present at equal frequency
- the more uneven the distribution, the fewer species are “effectively” present
- the more similar species are, the less “effectively present” each species is



Diversity Profiles with Similarity

Name	Formula
Power mean of order r of \mathbf{x} weighted by \mathbf{w}	$M_r(\mathbf{w}, \mathbf{x}) = \begin{cases} \left[\sum_{i=1}^S w_i \times x_i^r \right]^{1/r} & r \neq 0 \\ \prod_{i=1}^S x_i^{w_i} & r = 0 \end{cases}$
Hill numbers	${}^q D(\mathbf{p}) = M_{q-1}(\mathbf{p}, \mathbf{p})^{-1}$
Similarity-sensitive diversity	${}^q D^Z(\mathbf{p}) = M_{q-1}(\mathbf{p}, \mathbf{Zp})^{-1}$ $[{}^q D(\mathbf{p}) = {}^q D^I(\mathbf{p})]$



ALPHA DIVERSITY

BETA DIVERSITY

GAMMA DIVERSITY

- What's different?
 - We think in terms of individual sub-communities as much as the whole ecosystem
- Coming from that perspective allows us to answer specific questions:
 - Which sub-communities are the most diverse?
 - Which sub-communities are the most distinct?
 - Which sub-communities contribute the most to overall diversity?



Diversity measure	Sub-community diversity (d)	Ecosystem diversity (D)
Alpha diversity	${}^q\alpha_j^Z(\mathbf{P}_j) = M_{q-1}(\hat{\mathbf{P}}_j, \mathbf{Z}\mathbf{P}_j)^{-1}$ ${}^q\hat{\alpha}_j^Z(\mathbf{P}_j) = {}^qD^Z(\hat{\mathbf{P}}_j)^* = w_j \times {}^q\alpha_j^Z(\mathbf{P}_j)$	${}^qD^Z(\mathbf{P}) = M_{1-q}(\mathbf{w}, {}^q\alpha^Z(\mathbf{P}_j) _{j \in \{1 \dots N\}})$ ${}^q\hat{D}^Z(\mathbf{P}) = M_{1-q}(\mathbf{w}, {}^q\hat{\alpha}^Z(\mathbf{P}_j) _{j \in \{1 \dots N\}})$
Beta diversity	${}^q\beta_j^Z(\mathbf{P}_j, \mathbf{p}) = M_{q-1}\left(\hat{\mathbf{P}}_j, \frac{\mathbf{Z}\mathbf{P}_j}{\mathbf{Z}\mathbf{p}}\right)$ ${}^q\hat{\beta}_j^Z(\mathbf{P}_j, \mathbf{p}) = M_{q-1}\left(\hat{\mathbf{P}}_j, \frac{\mathbf{Z}\hat{\mathbf{P}}_j}{\mathbf{Z}\mathbf{p}}\right) = \frac{{}^q\beta_j^Z(\mathbf{P}_j, \mathbf{p})}{w_j}$	${}^q_{\beta}D^Z(\mathbf{P}) = M_{1-q}(\mathbf{w}, {}^q\beta^Z(\mathbf{P}_j, \mathbf{p}) _{j \in \{1 \dots N\}})$ ${}^q_{\beta}\hat{D}^Z(\mathbf{P}) = M_{1-q}(\mathbf{w}, {}^q\hat{\beta}^Z(\mathbf{P}_j, \mathbf{p}) _{j \in \{1 \dots N\}})^{\dagger}$
Gamma diversity	${}^q\gamma_j^Z(\mathbf{P}_j, \mathbf{p}) = M_{q-1}(\hat{\mathbf{P}}_j, \mathbf{Z}\mathbf{p})^{-1} [= {}^q\hat{\gamma}_j^Z(\mathbf{P}_j, \mathbf{p})]$	${}^q_{\gamma}D^Z(\mathbf{P}) = M_{1-q}(\mathbf{w}, {}^q\gamma^Z(\mathbf{P}_j, \mathbf{p}) _{j \in \{1 \dots N\}})^{\dagger}$ $[{}^q_{\gamma}\hat{D}^Z(\mathbf{P}) = {}^q_{\gamma}D^Z(\mathbf{P}) = {}^qD^Z(\mathbf{p})^*]$



Sub-community diversity

${}^q \alpha_i^Z$ Patch diversity

${}^q \beta_i^Z$ Distinctiveness from others

${}^q \gamma_i^Z$ Contribution to total diversity



Real World Example

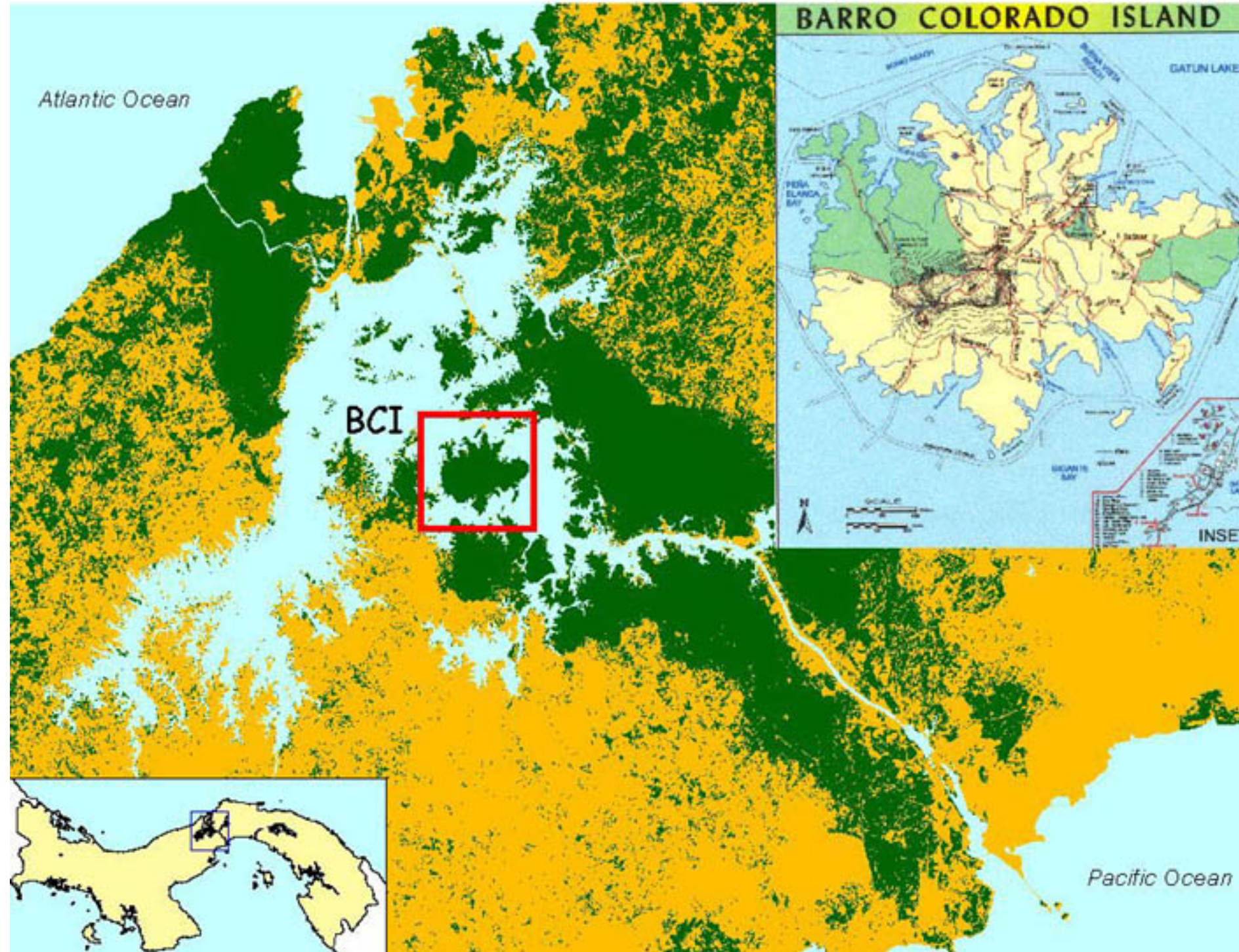


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Barro Colorado Island



Barro Colorado Island





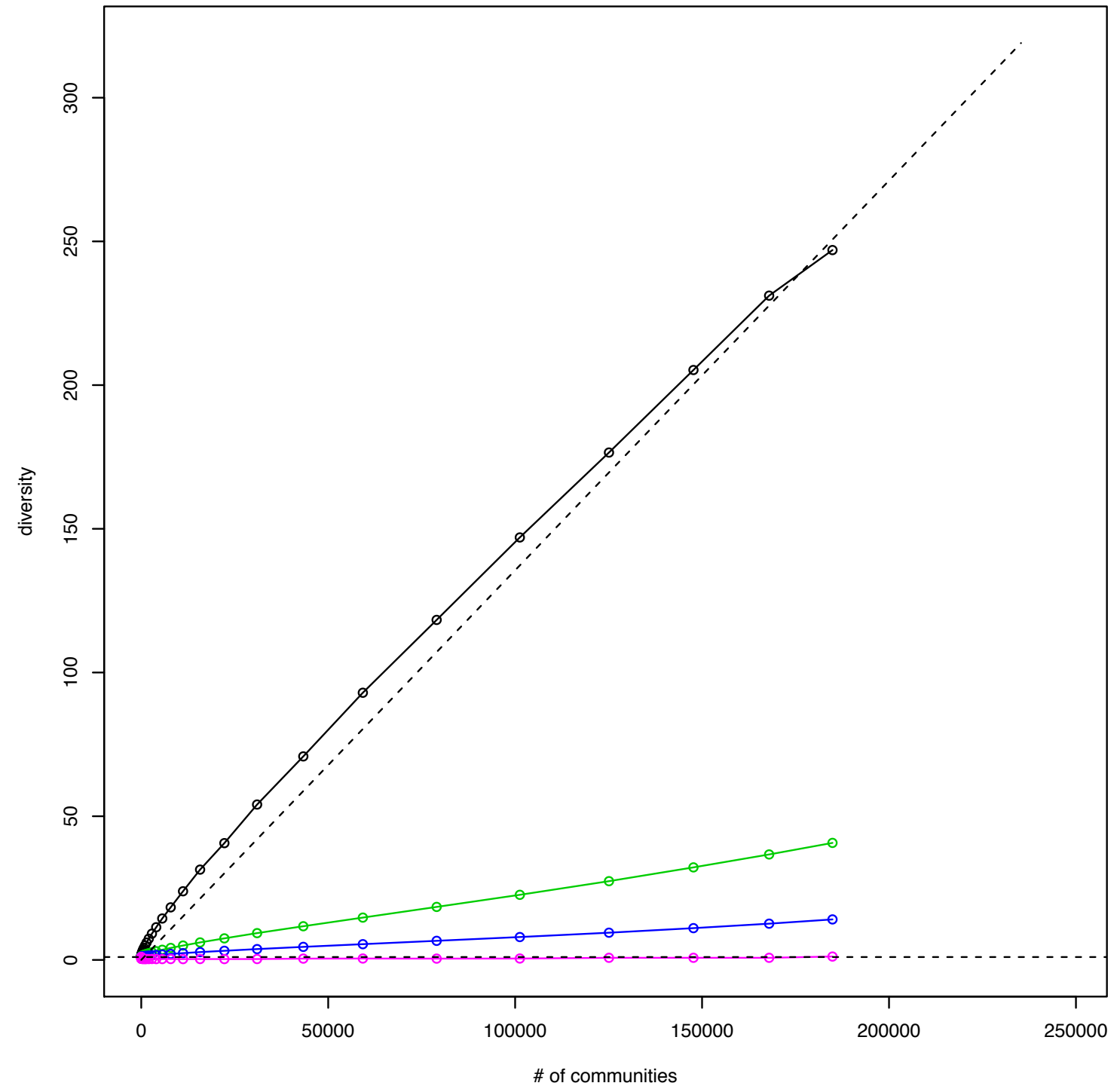
Barro Colorado Island

- 50 ha plot in Panama
 - Every individual tree and sapling recorded
 - Species, Size, Location
 - 1981-2, 1985, 1990, 1995, 2000, 2005
 - Work done by Center for Tropical Forest Science and many others, funded by multiple sources



What can we tell from total beta diversity?

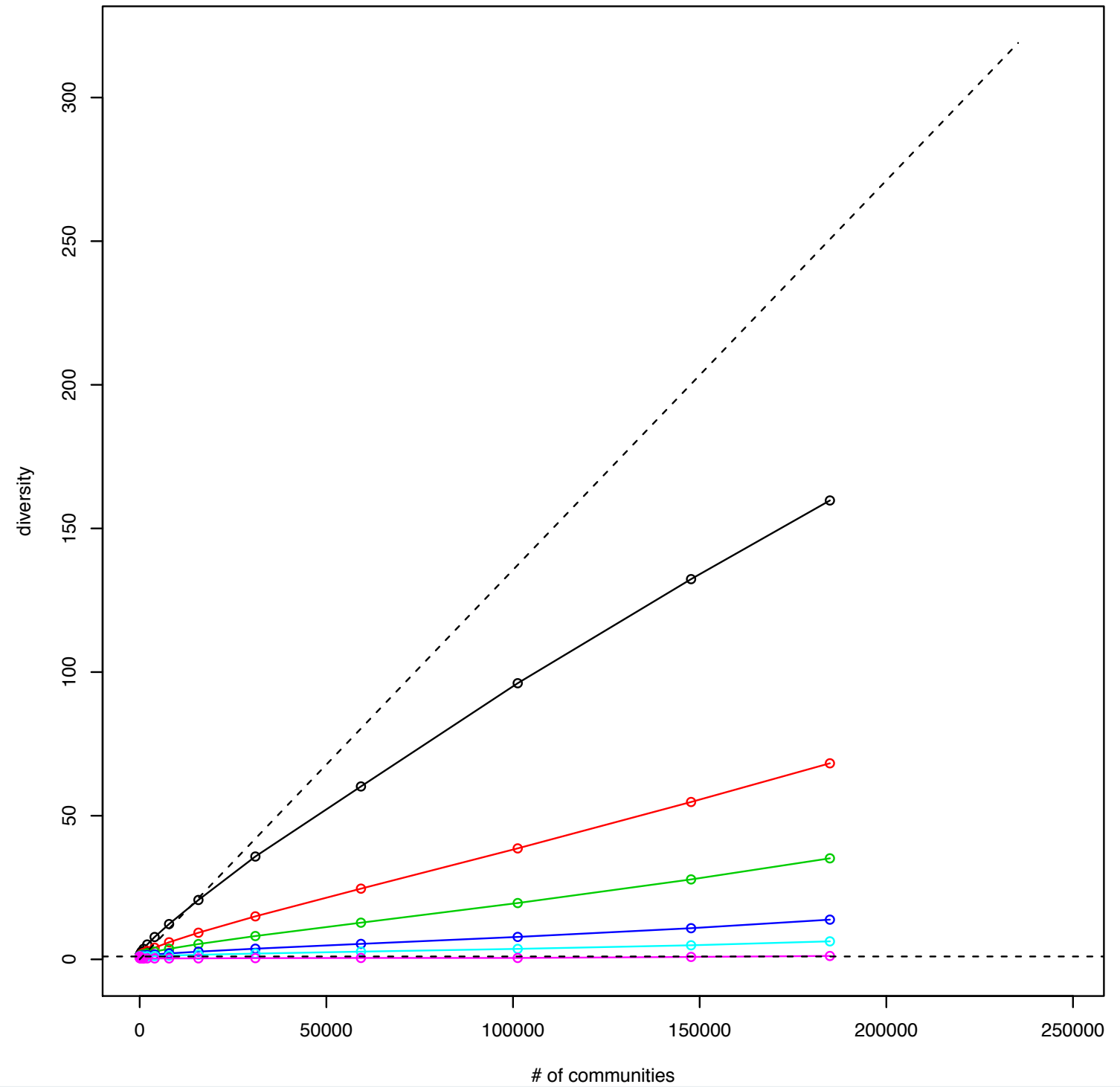
Beta diversity vs communities





What can we tell from total beta diversity?

Beta diversity vs communities





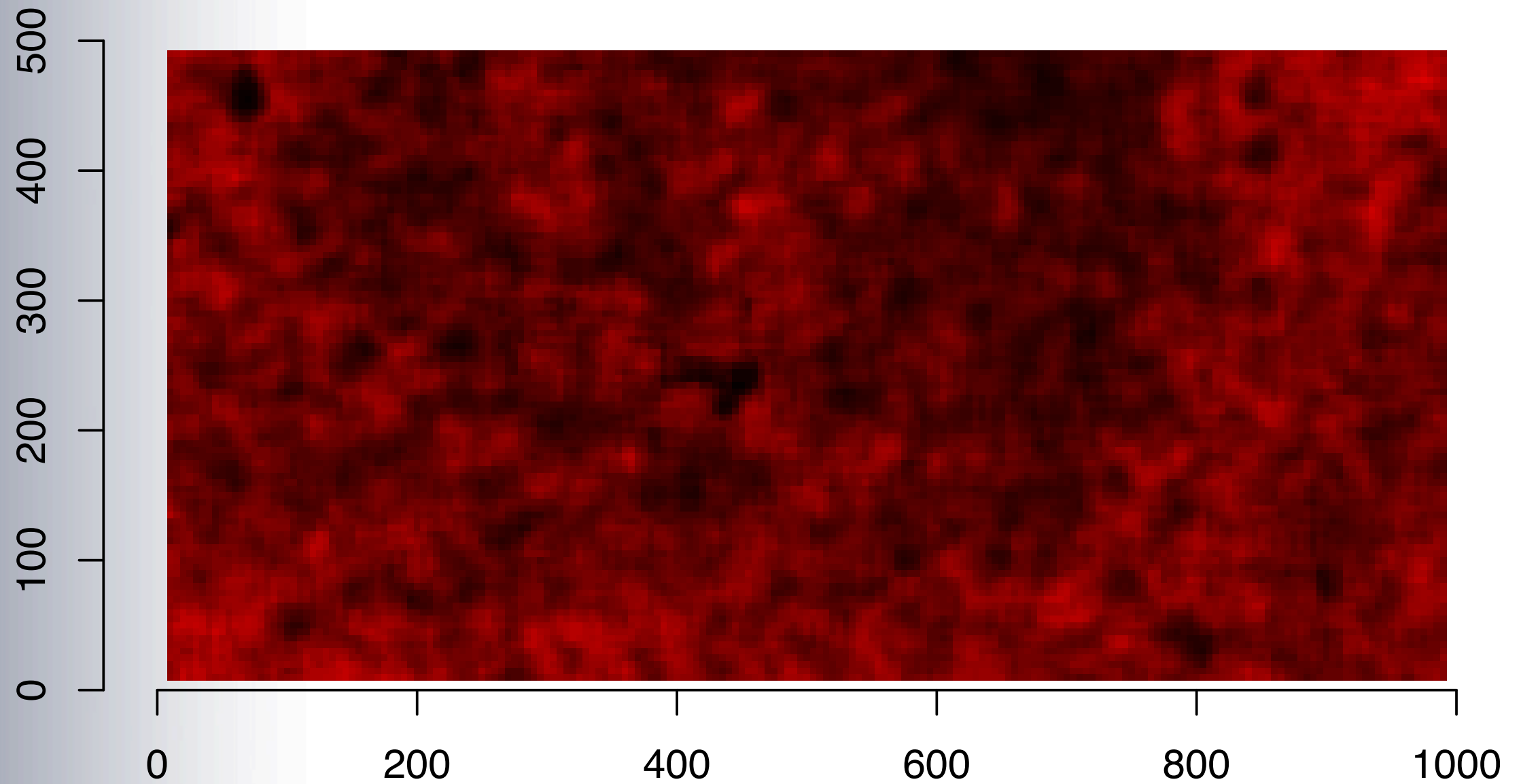
What can we tell from total beta diversity?

- Not much!
 - and total alpha and gamma diversity are similar...



How about individual alpha diversity?

Alpha diversity



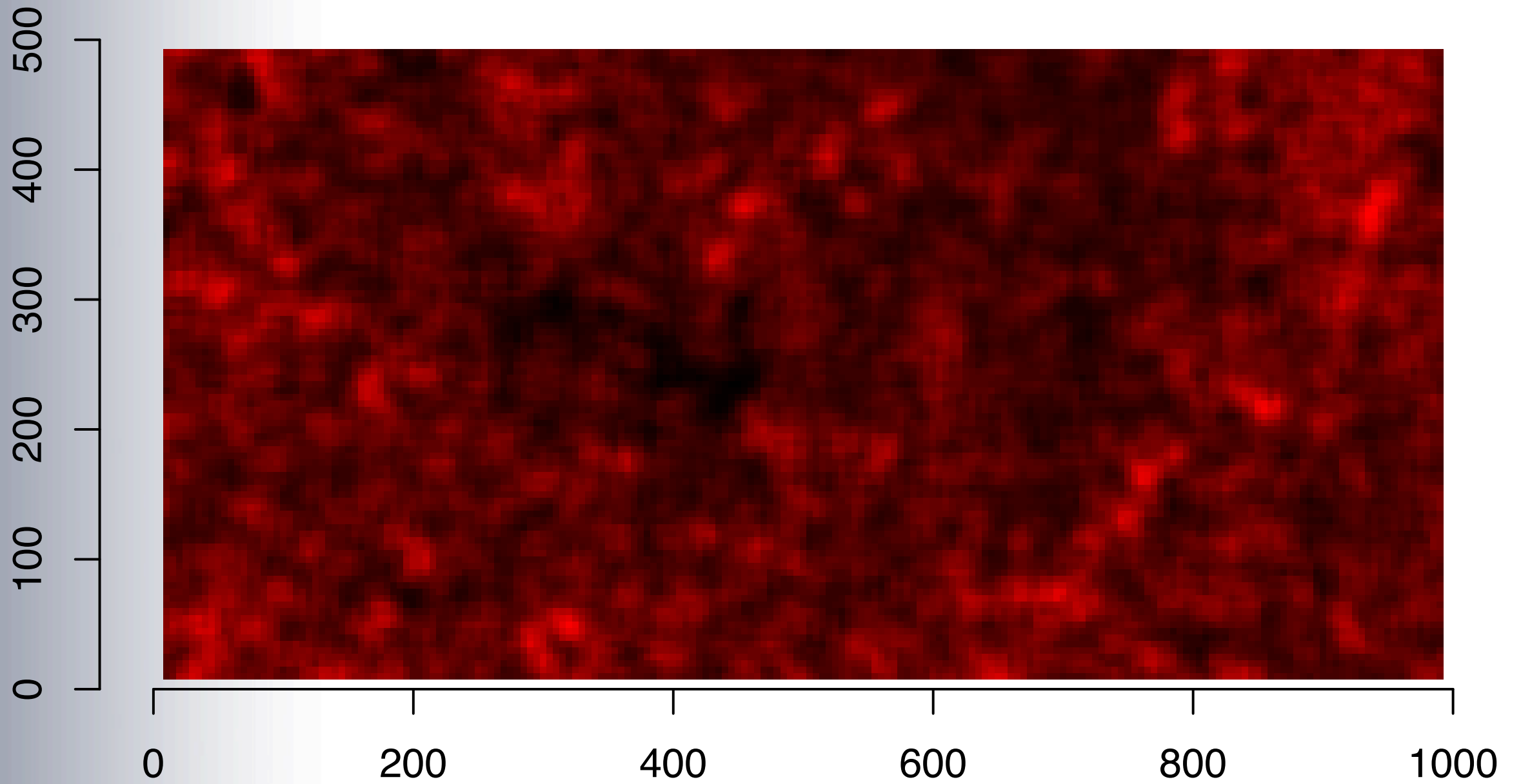


What can we assess?

- ${}^q\alpha_i^Z$ Value in terms of patch diversity
- ${}^q\beta_i^Z$ Value in terms of distinctiveness
- ${}^q\gamma_i^Z$ Value in term of contribution to total diversity



Value in terms of patch diversity



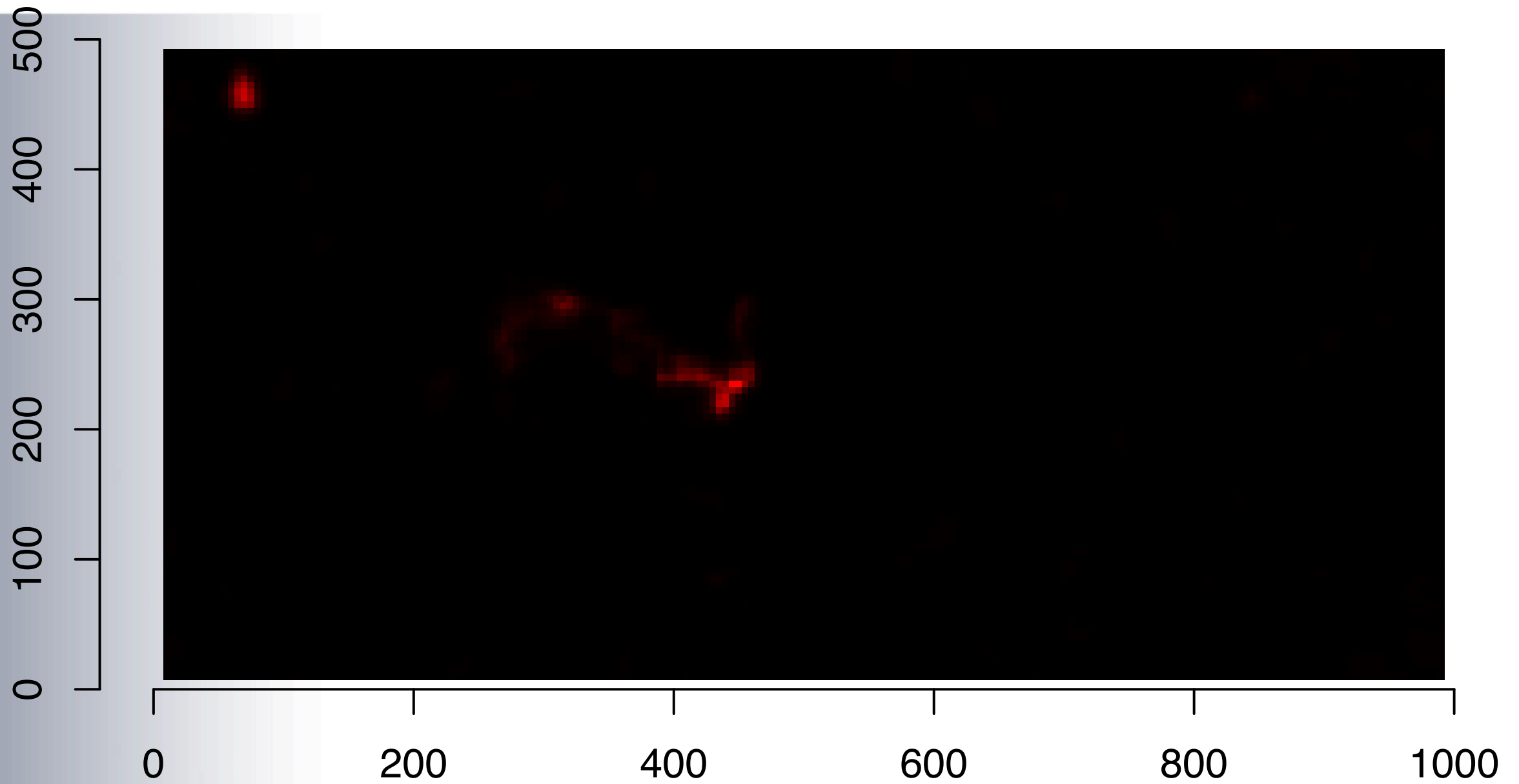


Value in terms of patch diversity

- We are beginning to see patterns here...
 - Picking out area with a lot of trees of high diversity
 - But not necessarily distinct from other areas, so selecting these, we may repeat ourselves with some species and miss others



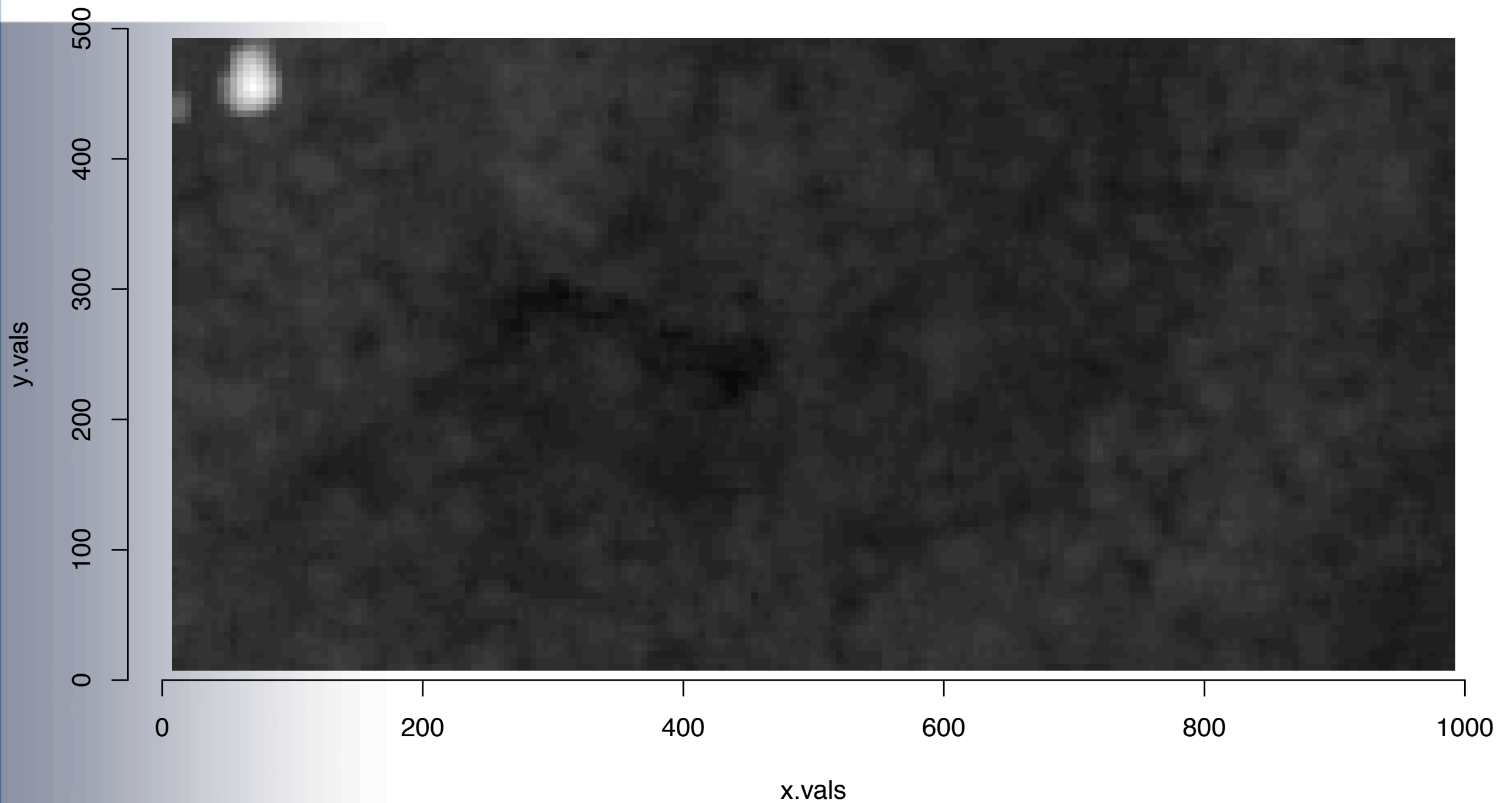
Beta diversity





What are these areas?

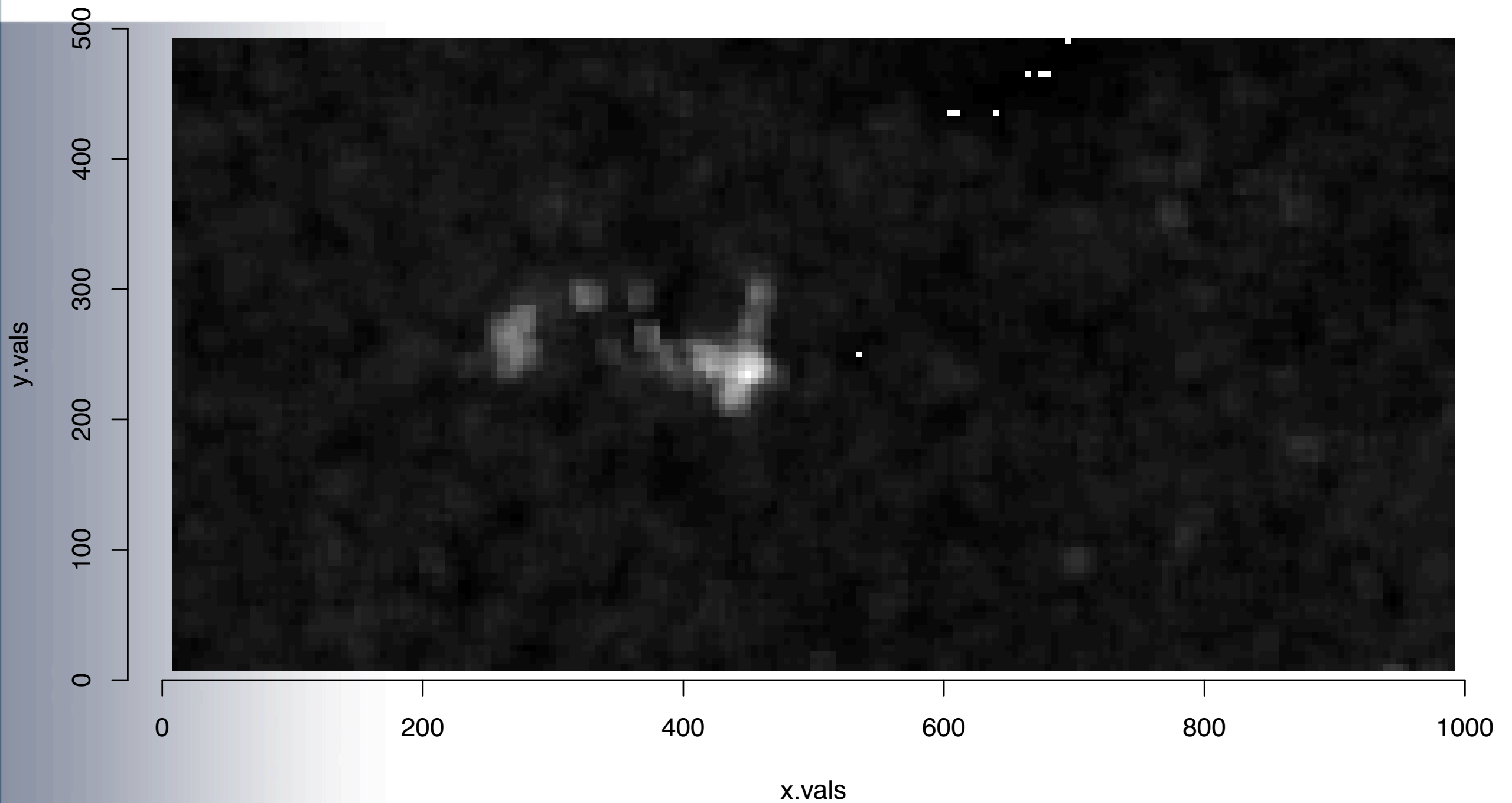
Similarity





What are these areas?

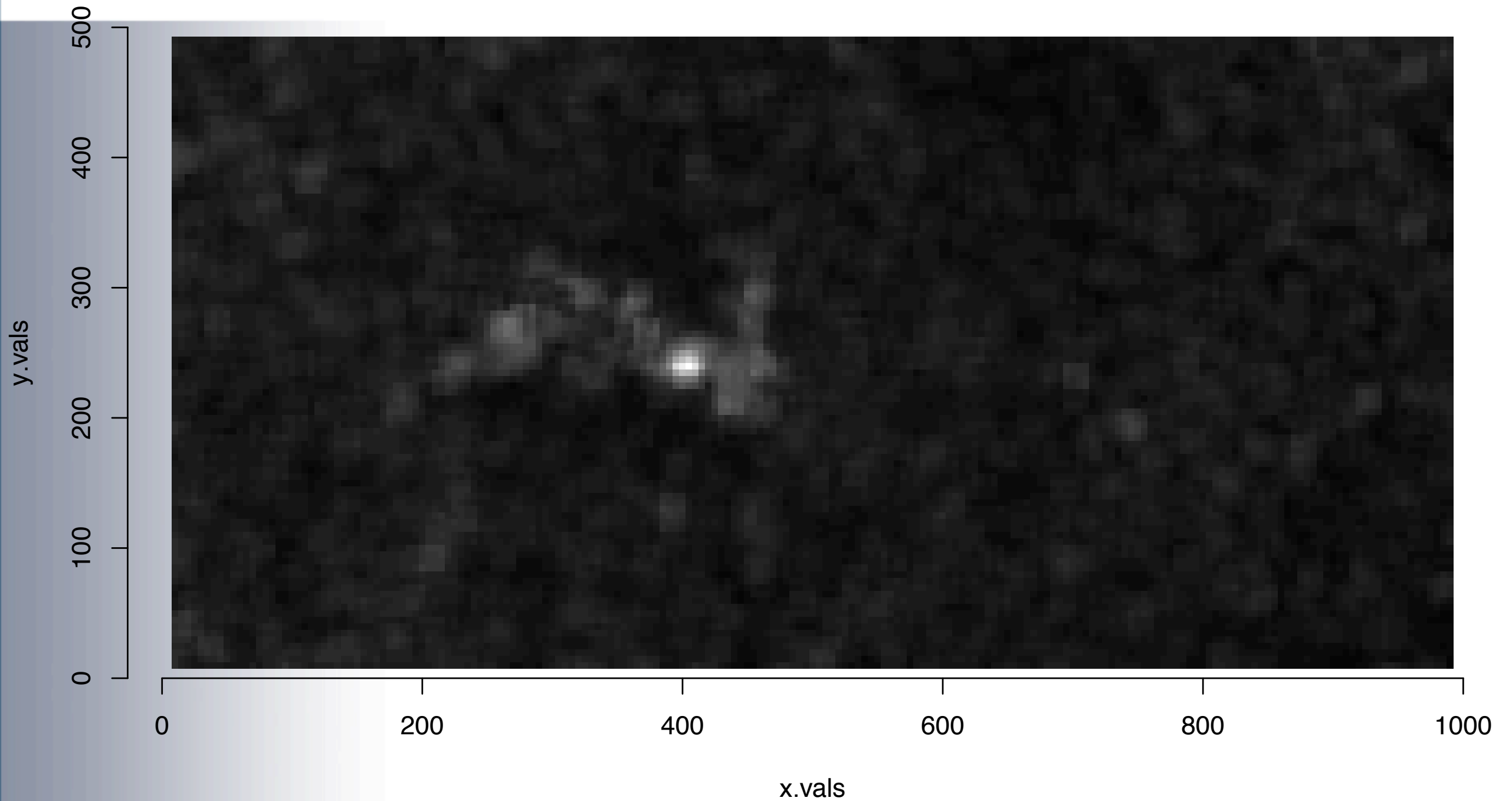
Similarity





What are these areas?

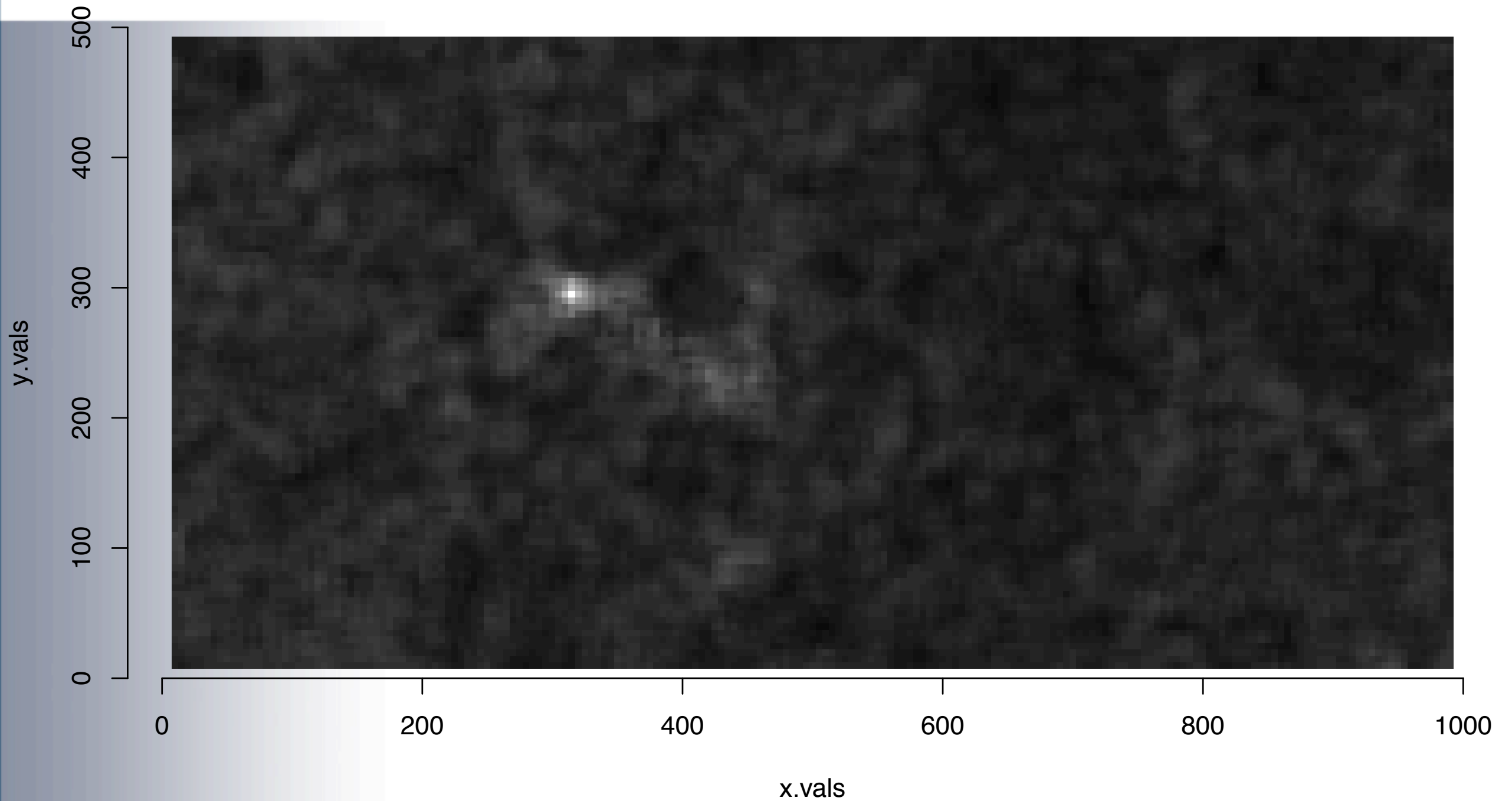
Similarity





What are these areas?

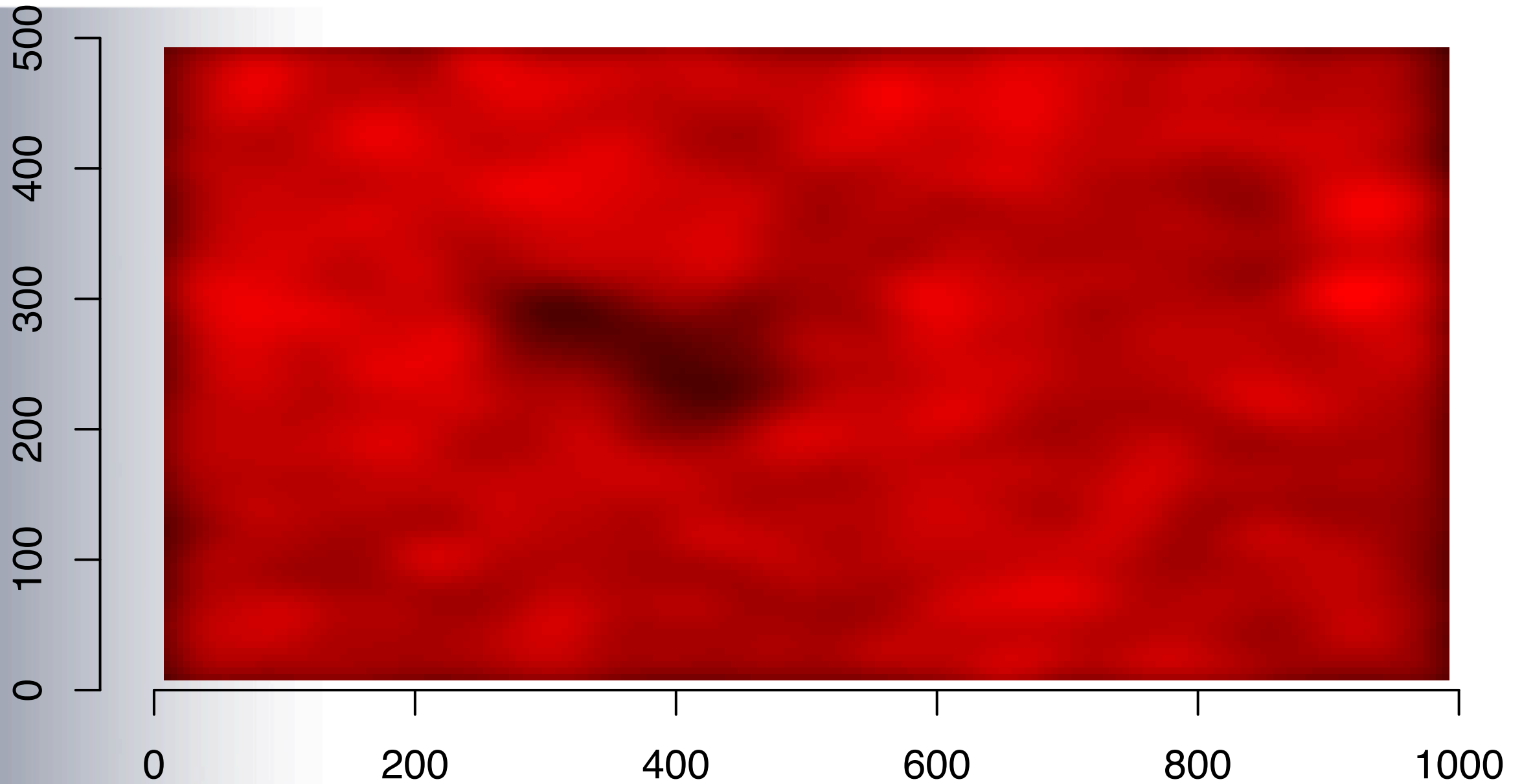
Similarity





What does Beta Diversity mean though?

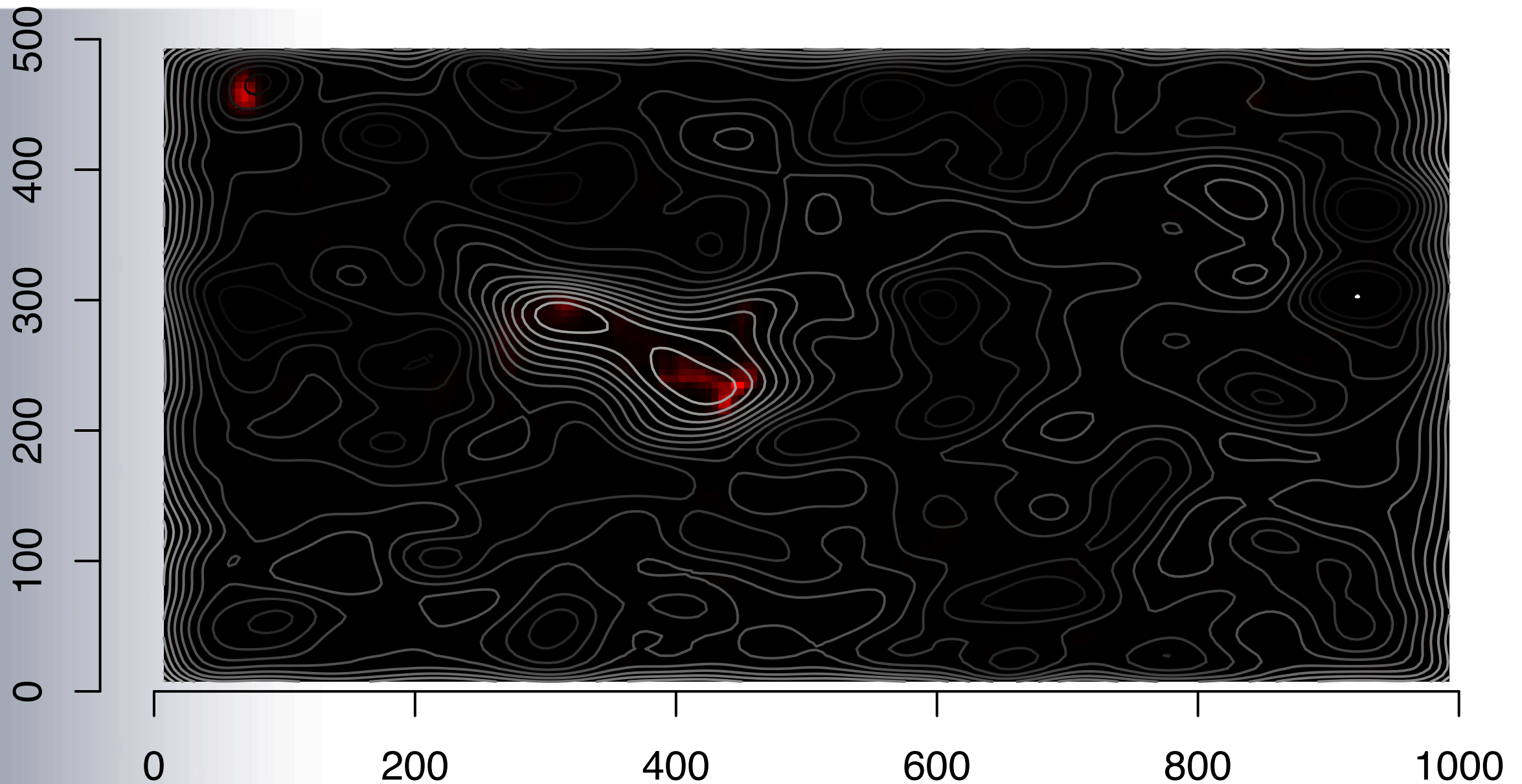
Tree density





What does Beta Diversity mean though?

Beta diversity





What does Beta Diversity mean though?

- Areas of high “distinctiveness”, but no regard for diversity or number of trees
- It identifies two areas, but one of them is largely devoid of trees...

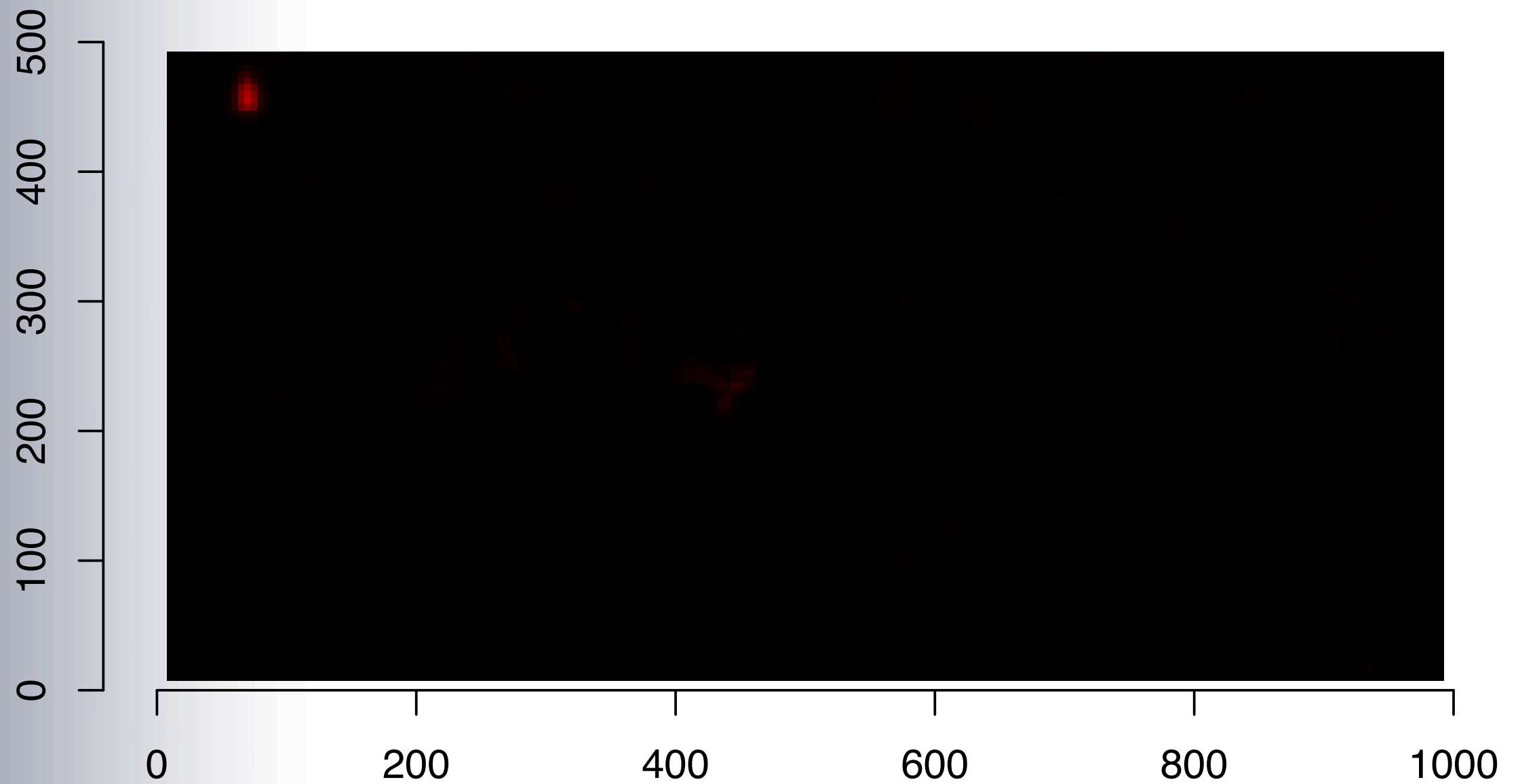


What can we assess?

- ${}^q \alpha_i^Z$ Value in terms of patch diversity
- ${}^q \beta_i^Z$ Value in terms of distinctiveness
- ${}^q \gamma_i^Z$ Value in term of contribution to total diversity



Value in terms of distinctiveness





Value in terms of distinctiveness

- It identifies areas that contribute a lot to total beta diversity
- Specifically finds one area, the area with high individual beta diversity but which also has trees in it!



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But is this what we want?



What can we assess?

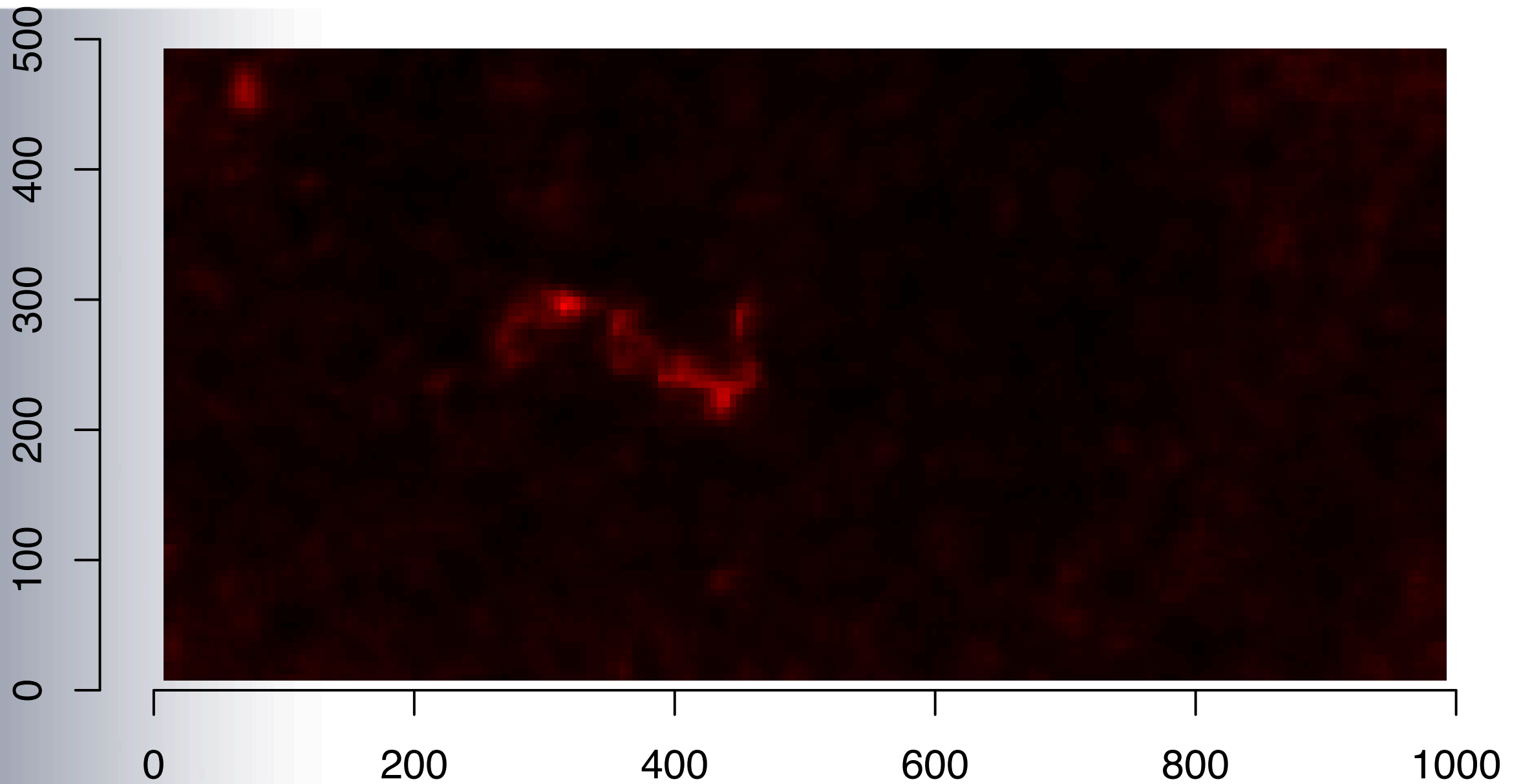
${}^q \alpha_i^Z$ Value in terms of patch diversity

${}^q \beta_i^Z$ Value in terms of distinctiveness

${}^q \gamma_i^Z$ Value in term of contribution to total diversity

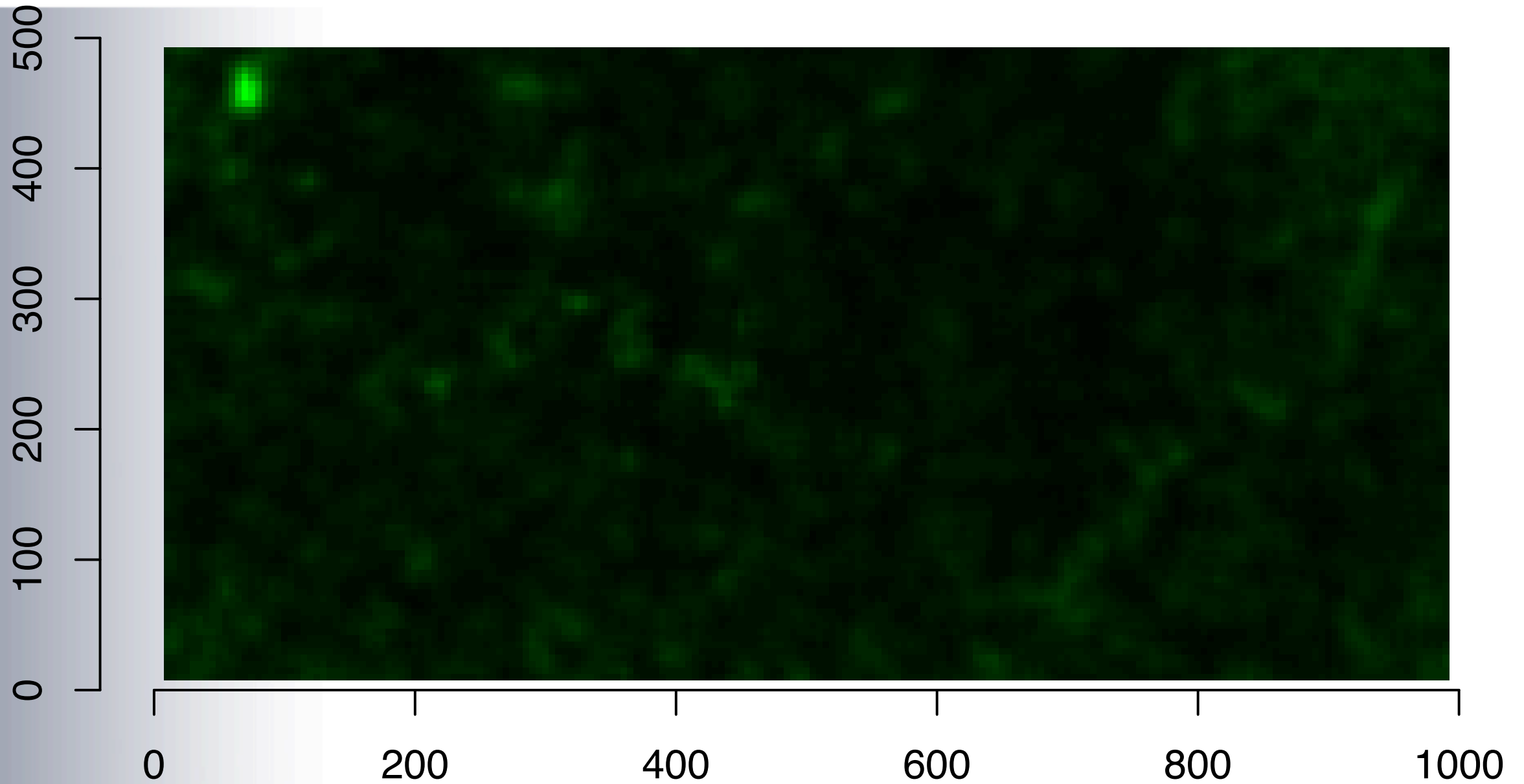


Gamma diversity





Value in terms of contribution to total diversity





Value in terms of contribution to total diversity

- It identifies areas that contribute a lot to the total gamma diversity of the whole plot
- Which areas this encompasses varies depending on your meaning of diversity!



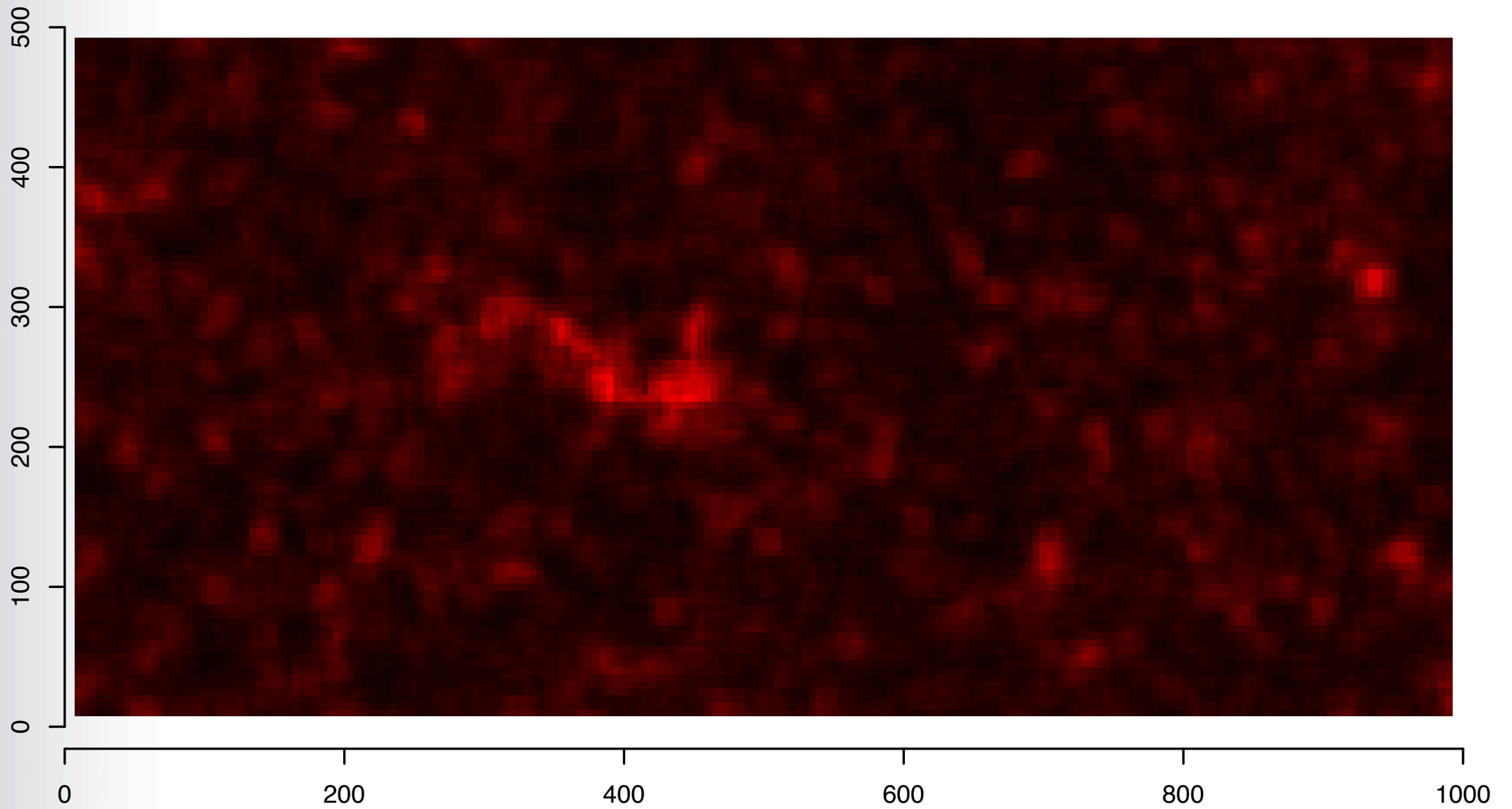
Temporal Beta Diversity

- Comparing a community at one timepoint to the “ecosystem” of that community measured at all timepoints



Temporal Beta Diversity

True beta, grid 20



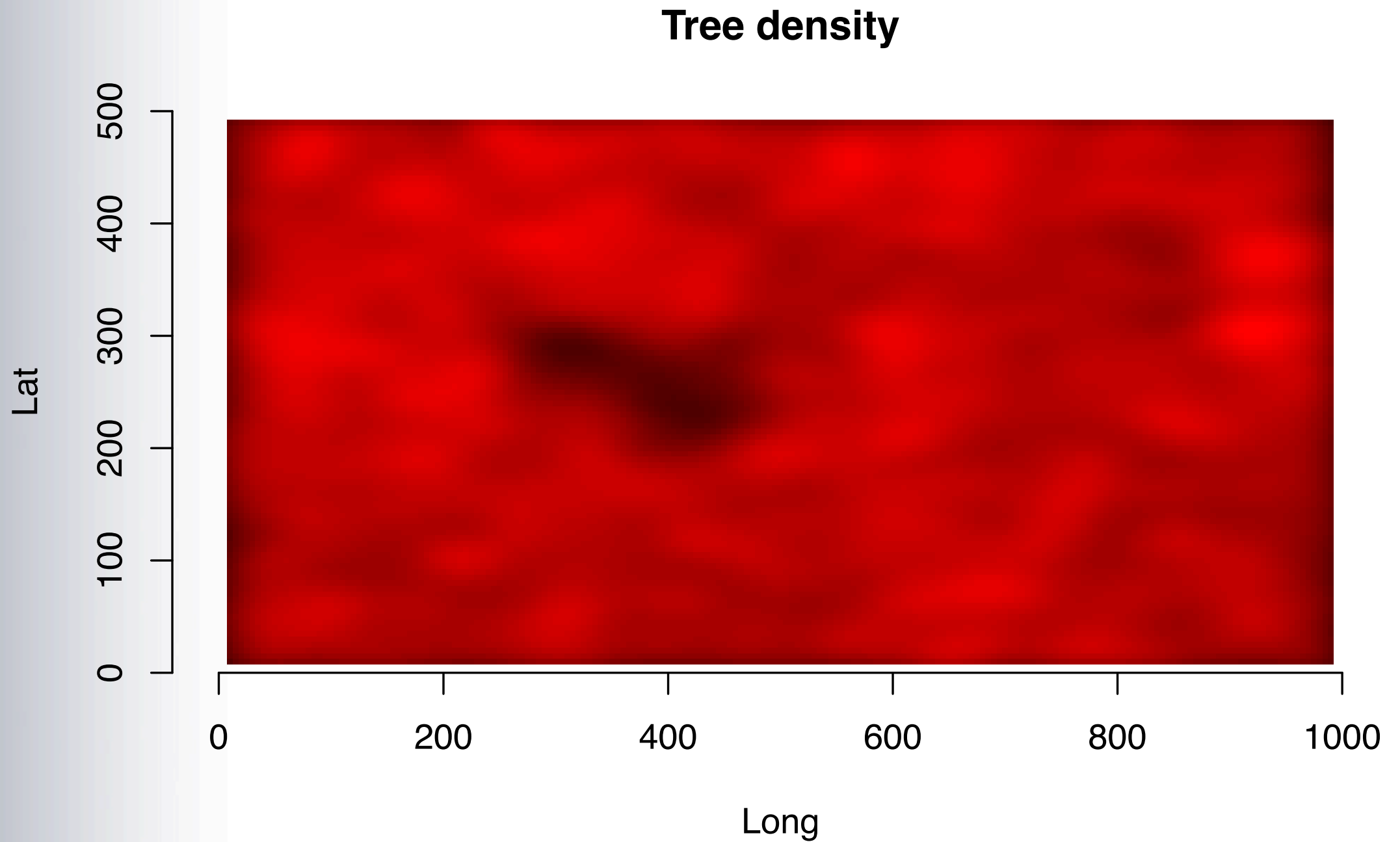


Temporal beta diversity

- One area has particularly high temporal beta diversity
 - and we've seen it before...

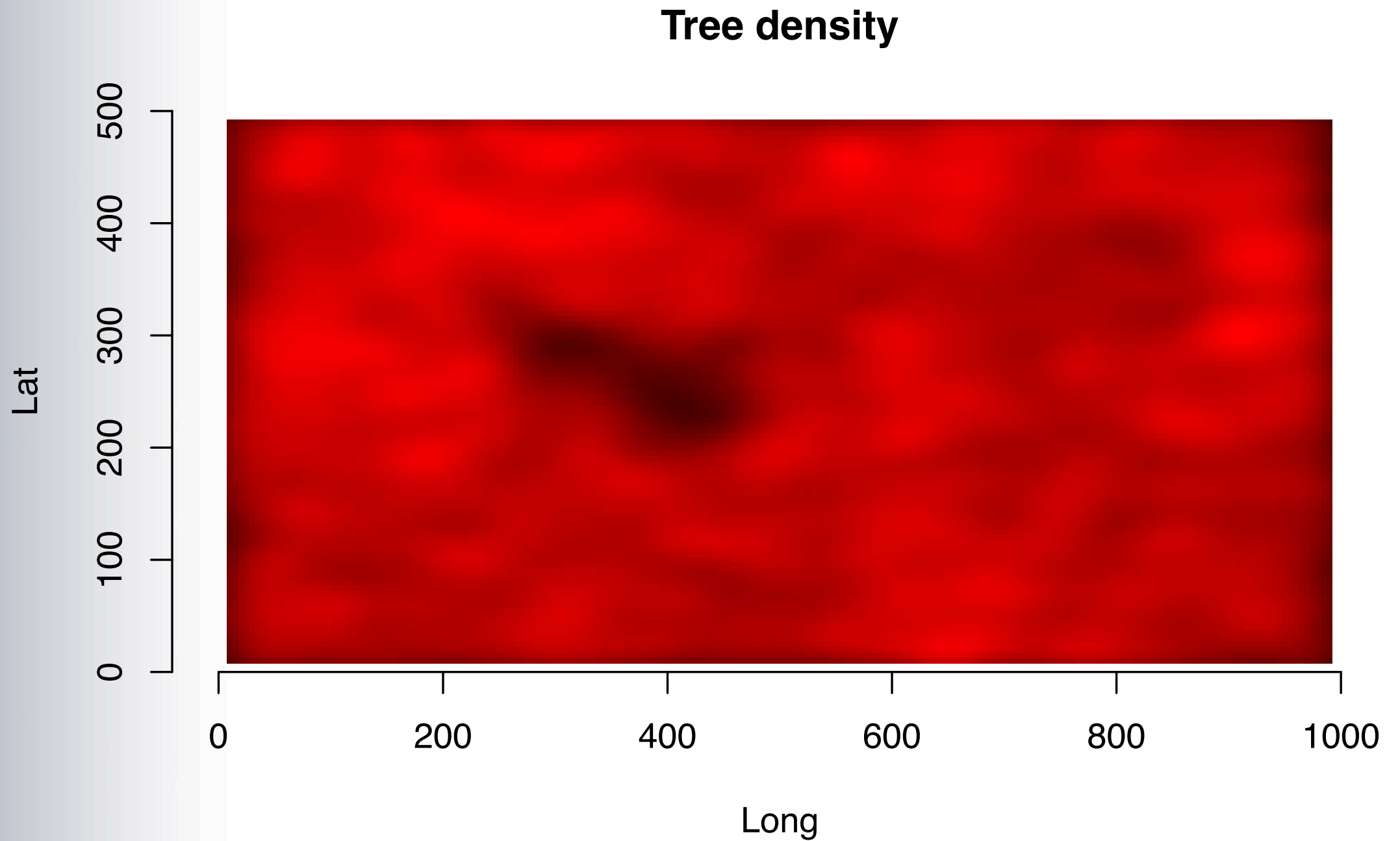


What's happening in this patch – 1981



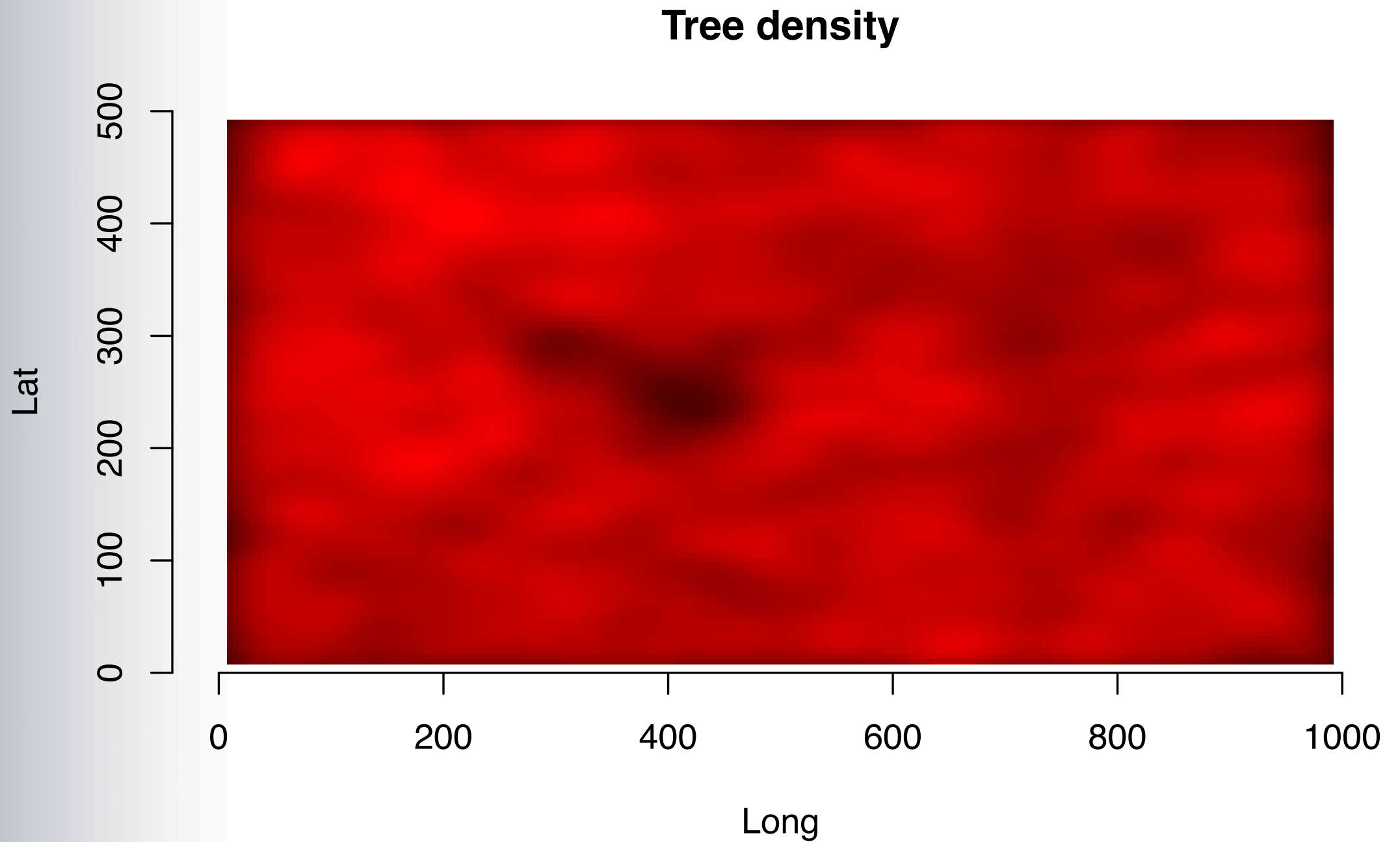


What's happening in this patch – 1985



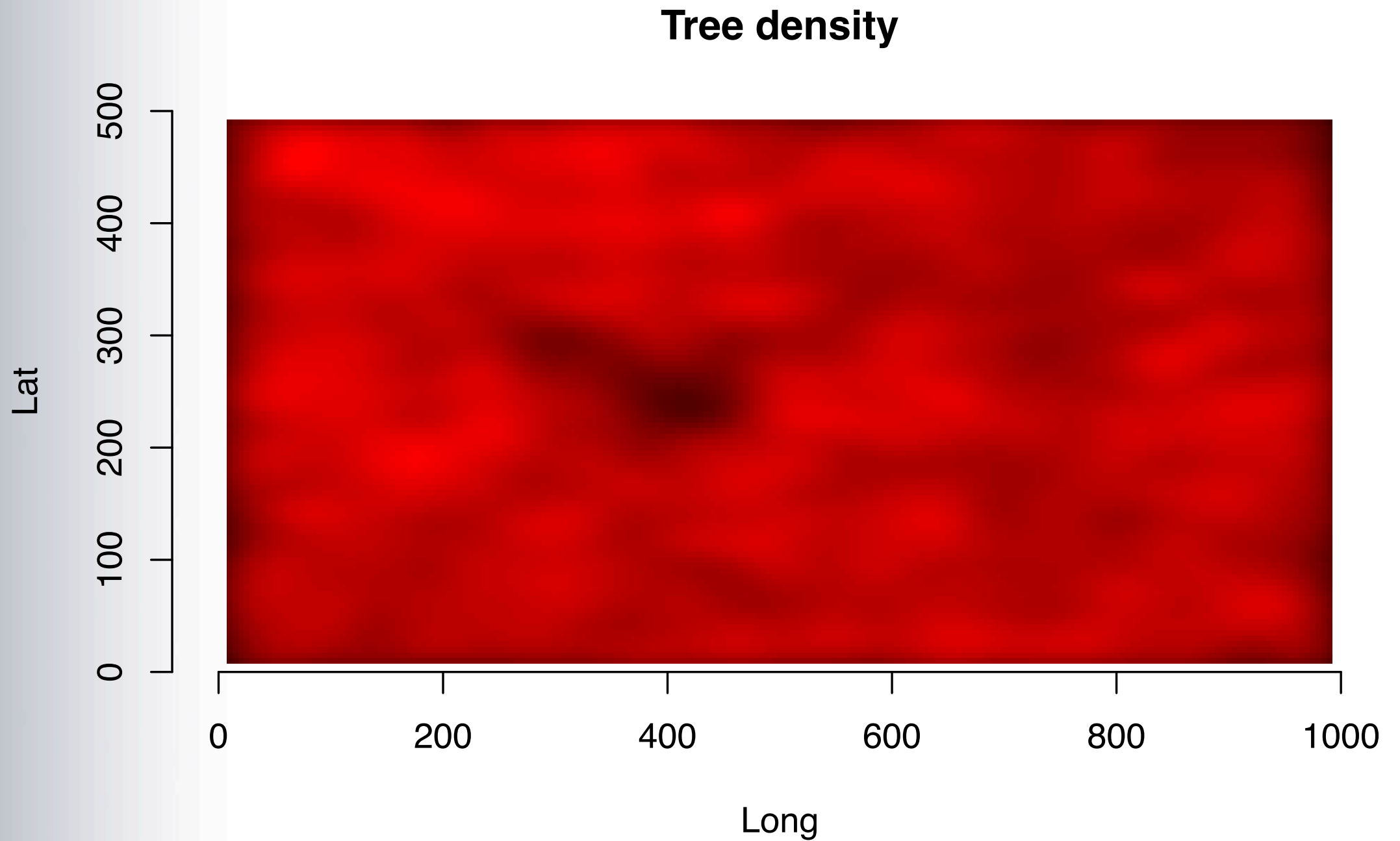


What's happening in this patch – 1990



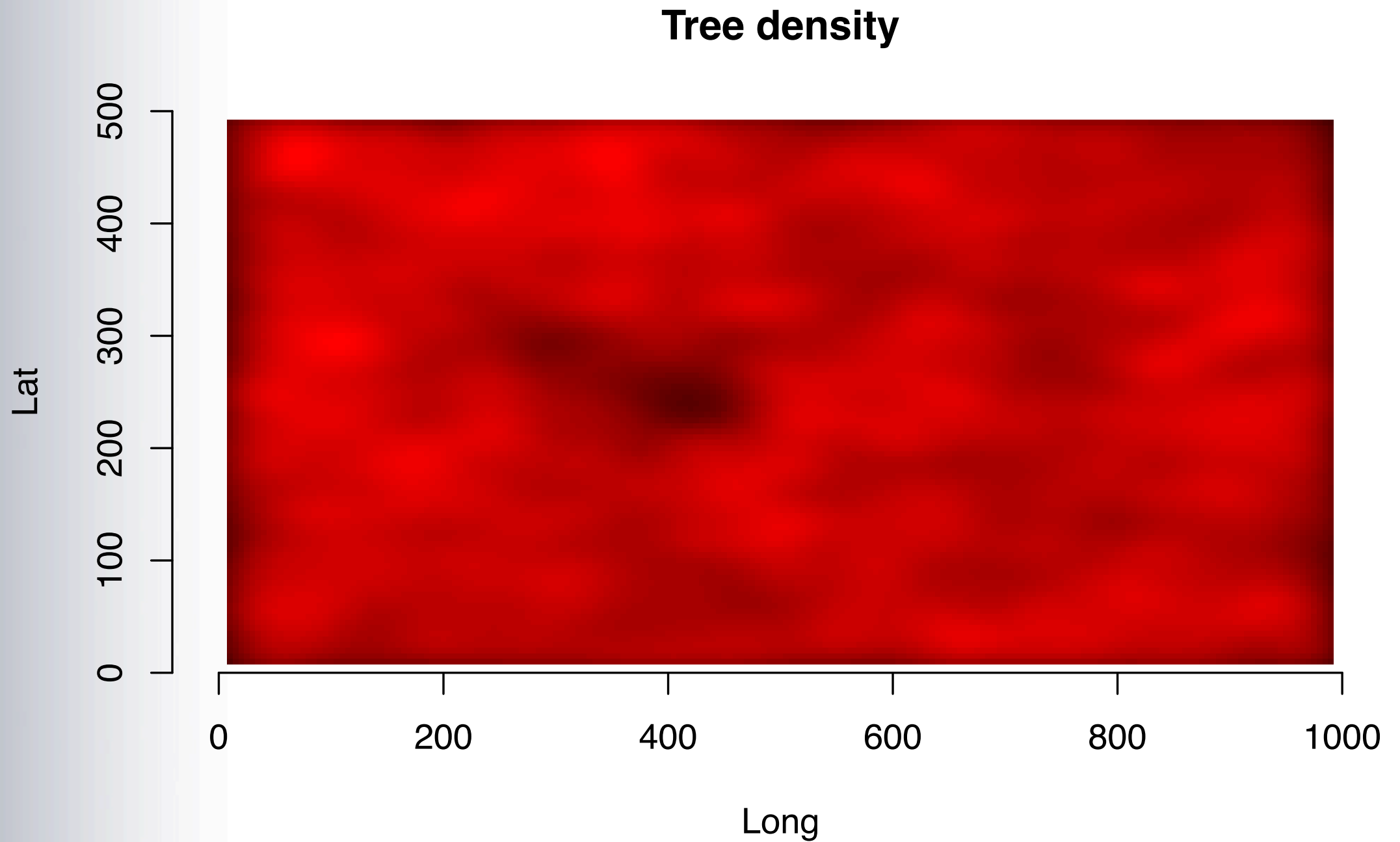


What's happening in this patch – 1995



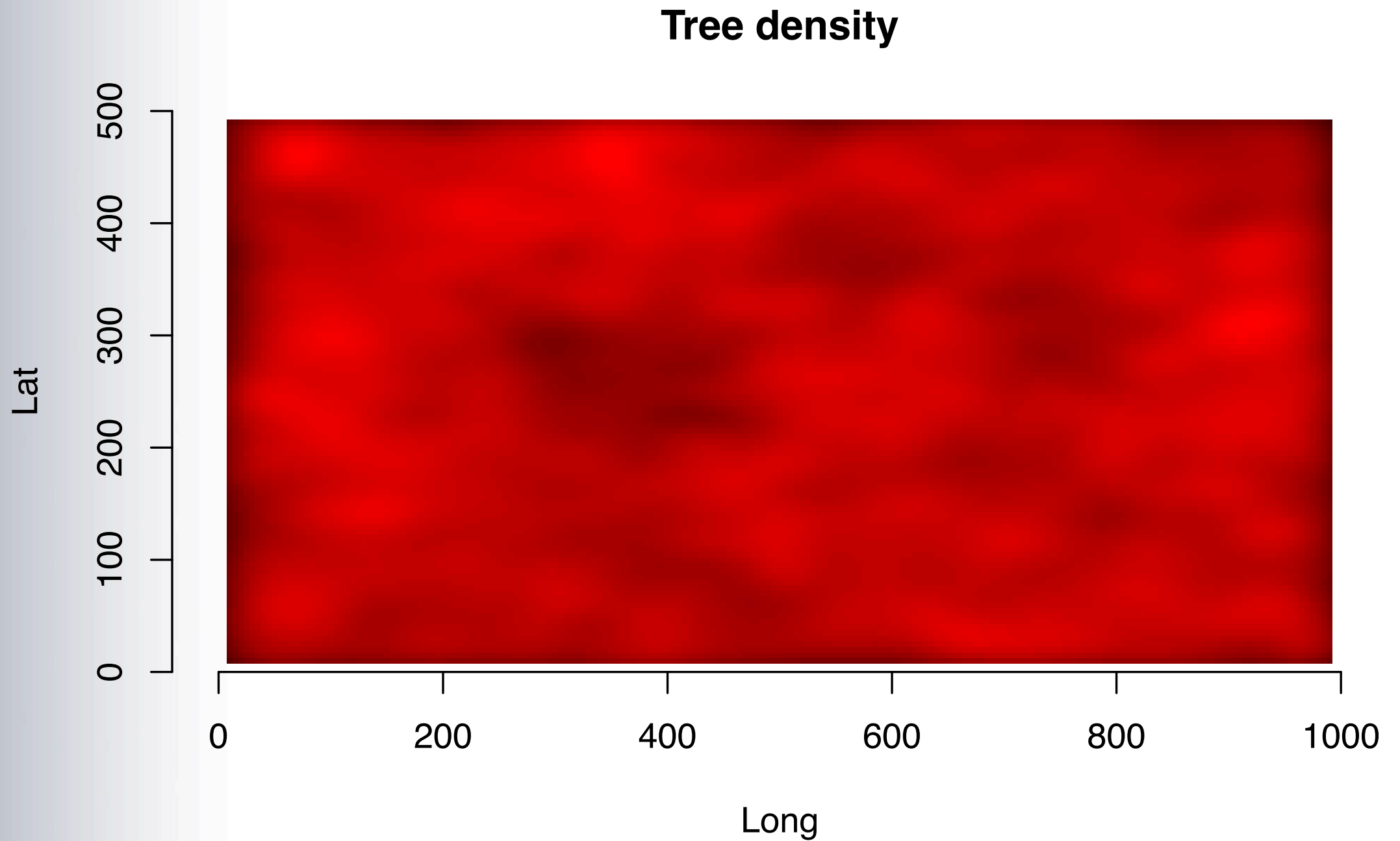


What's happening in this patch – 2000





What's happening in this patch – 2005





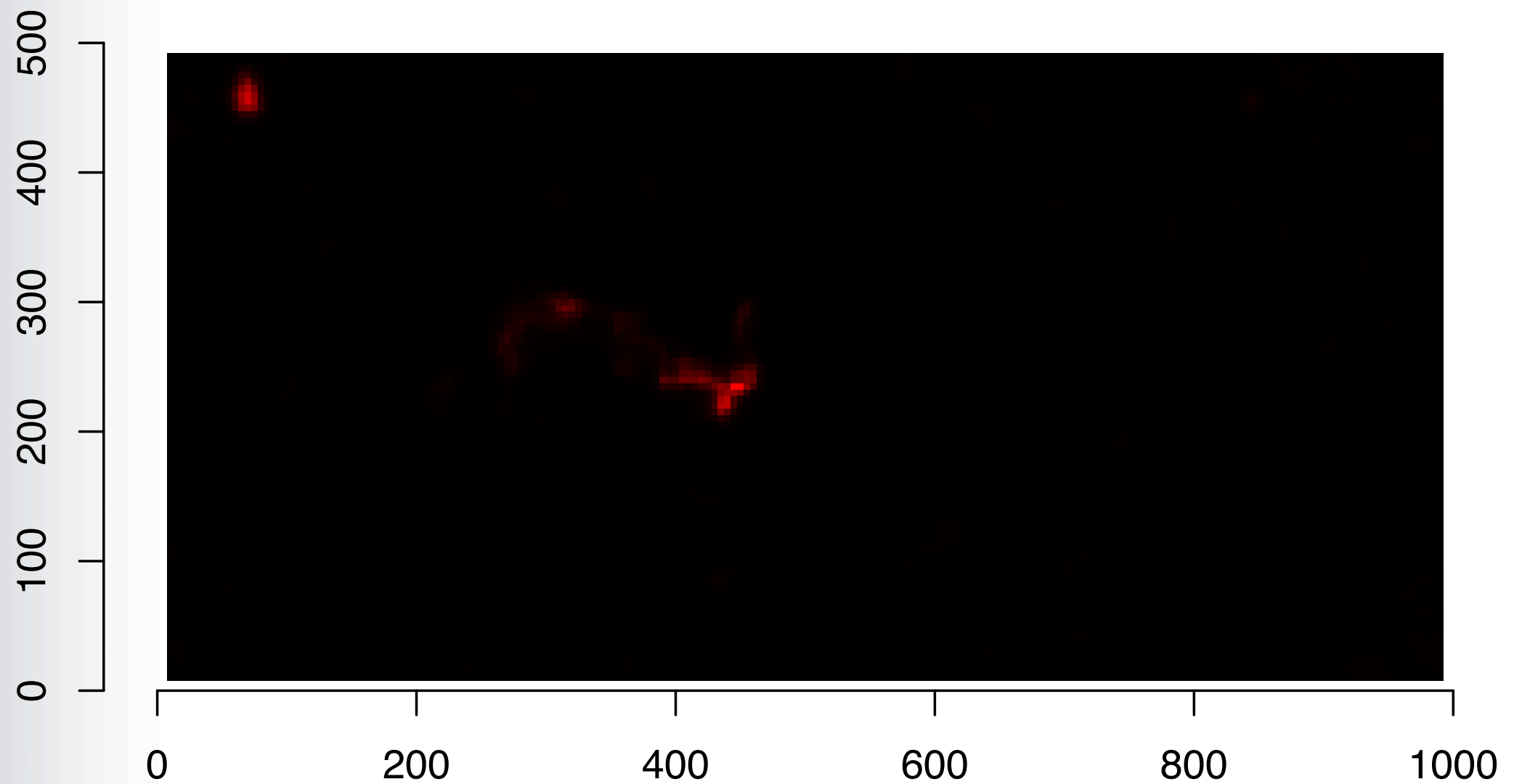
What's happening in this patch?

- The trees are filling in...



Revisiting Beta Diversity – 1981

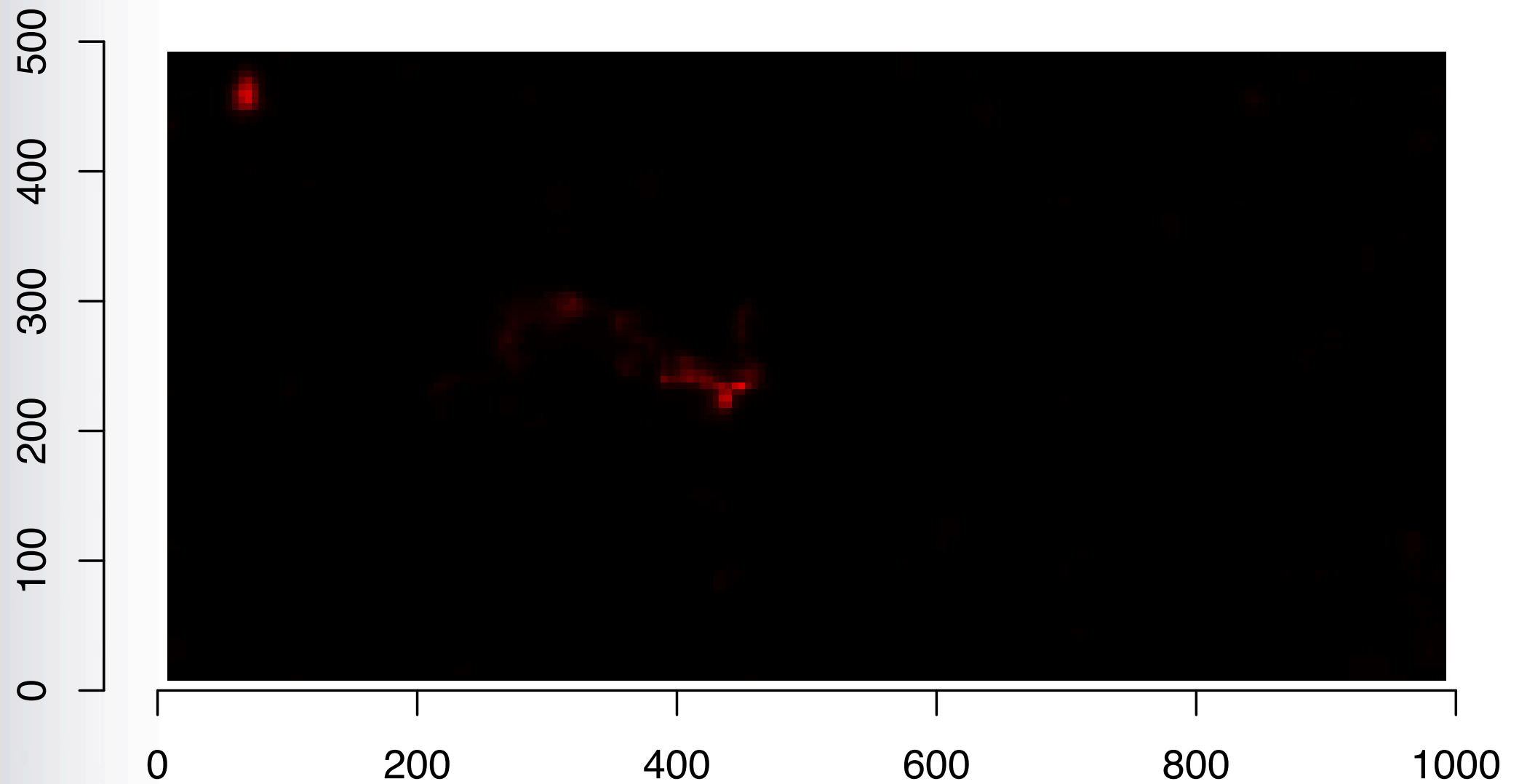
Beta diversity





Revisiting Beta Diversity – 1985

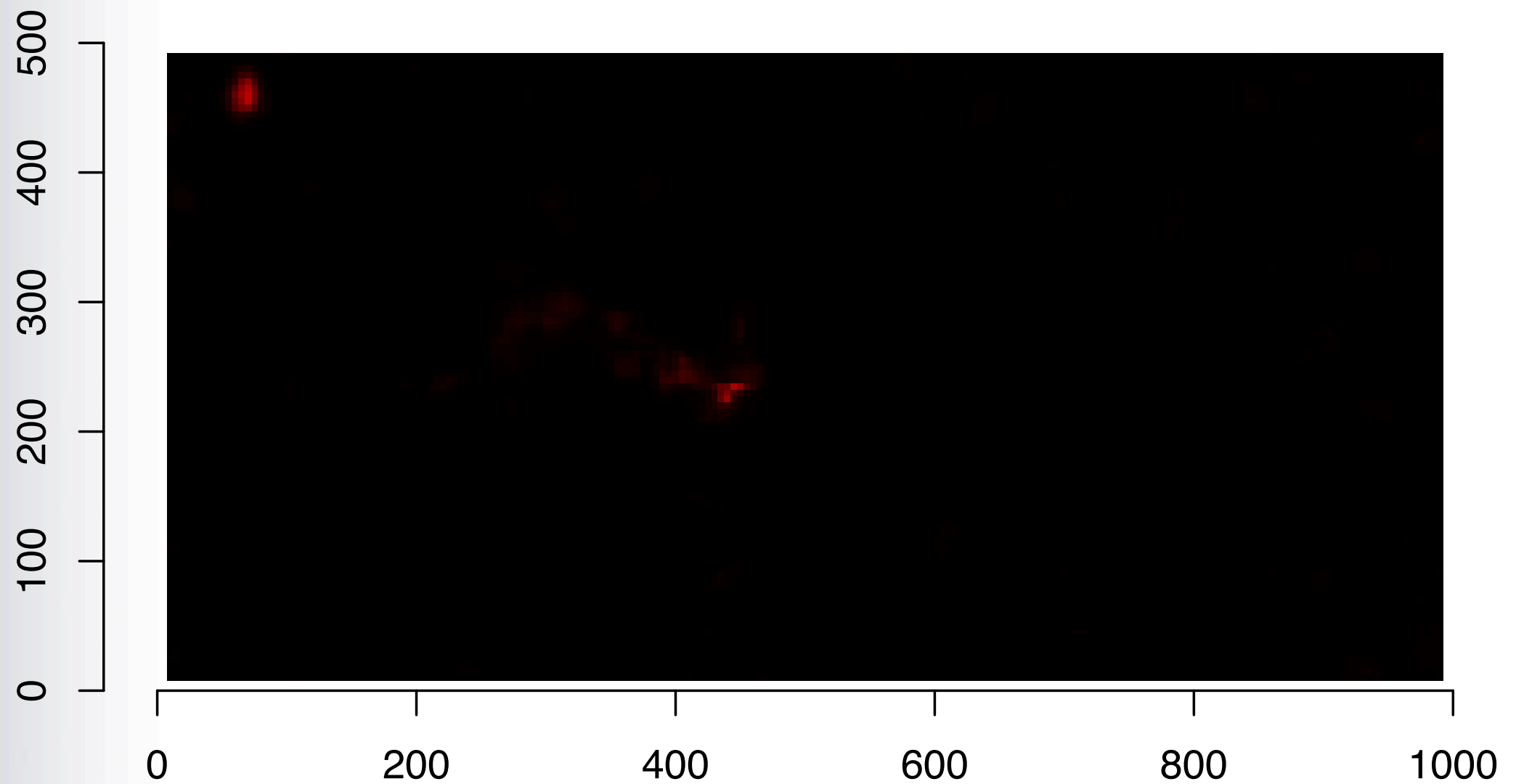
Beta diversity





Revisiting Beta Diversity – 1990

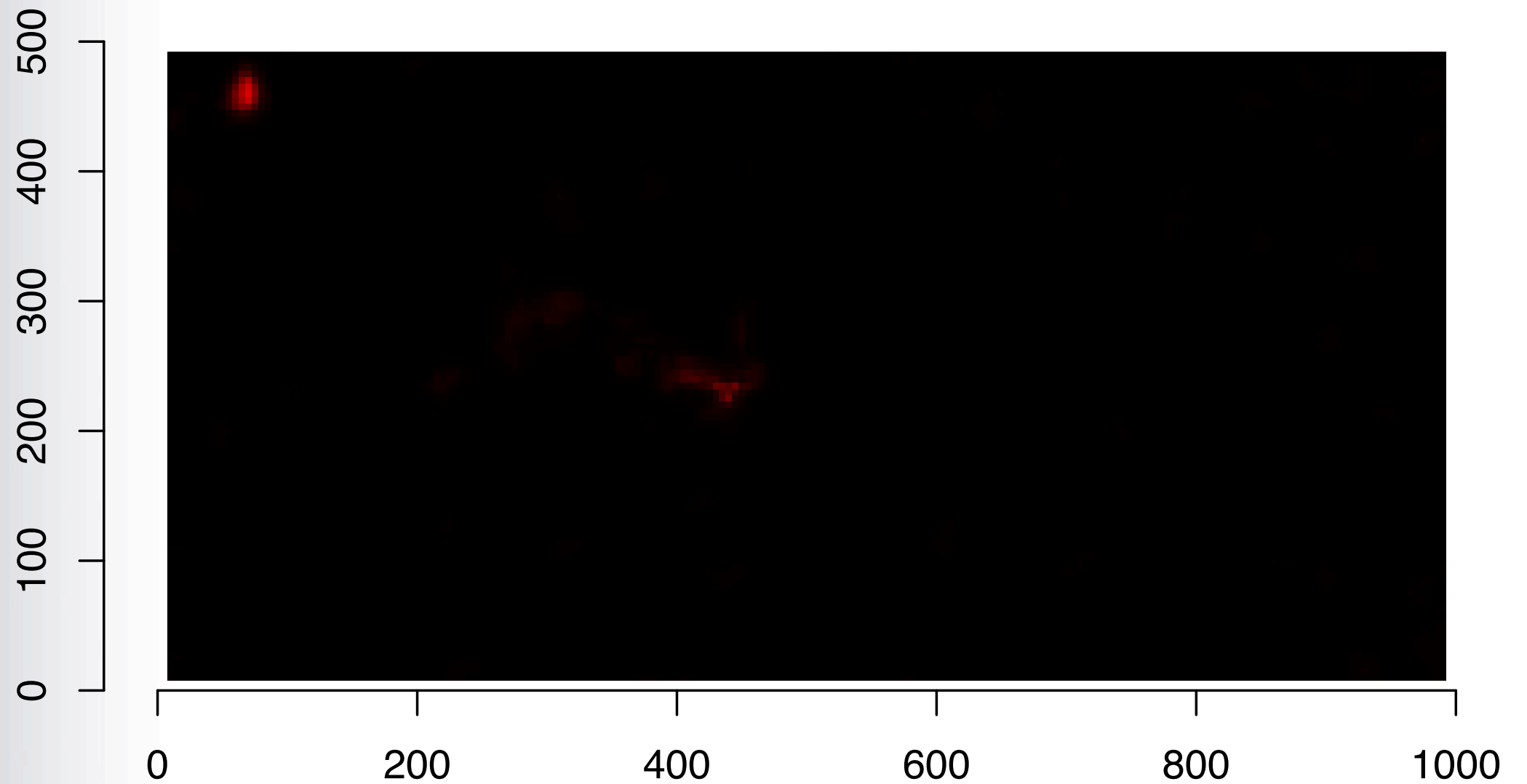
Beta diversity





Revisiting Beta Diversity – 1995

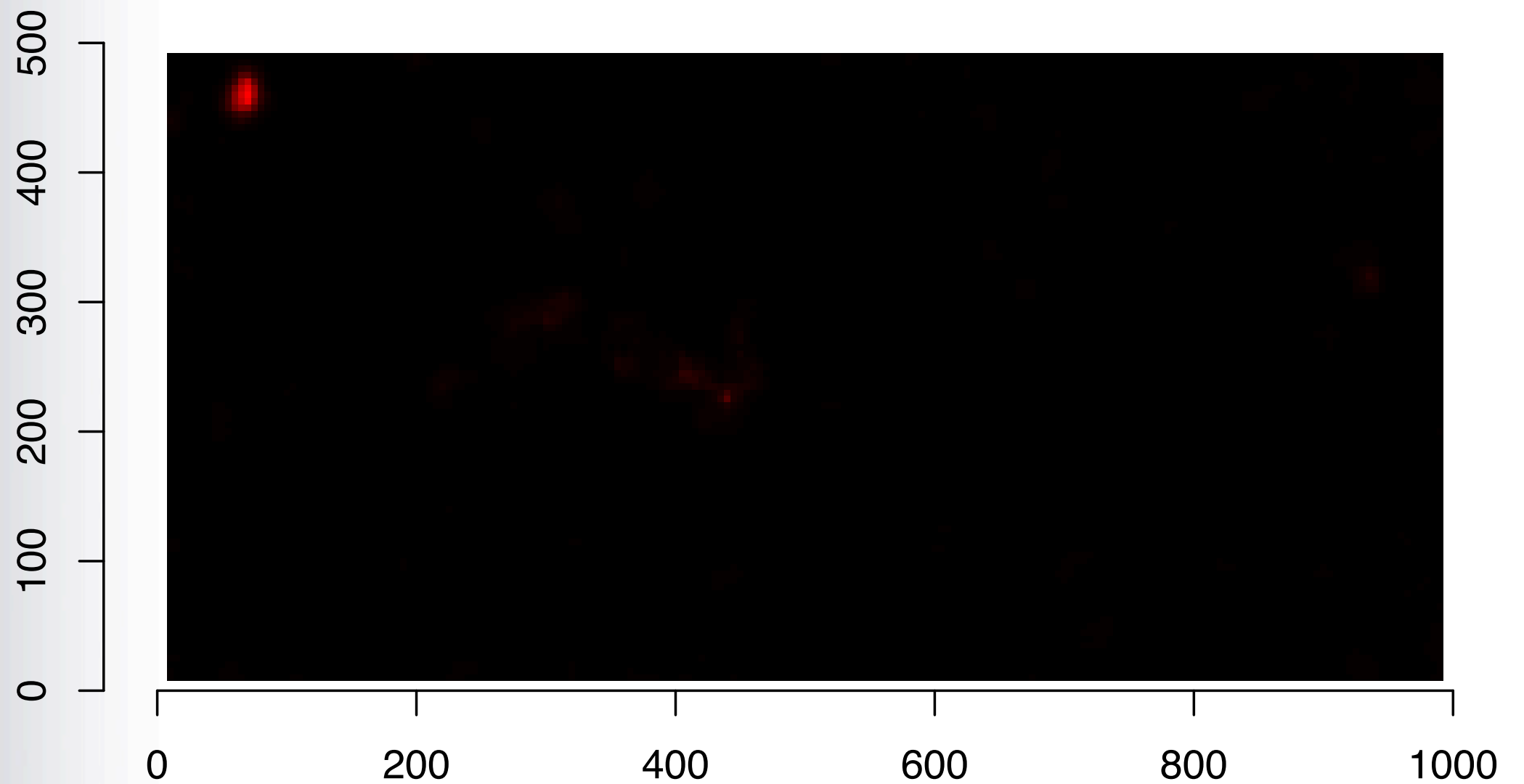
Beta diversity





Revisiting Beta Diversity – 2000

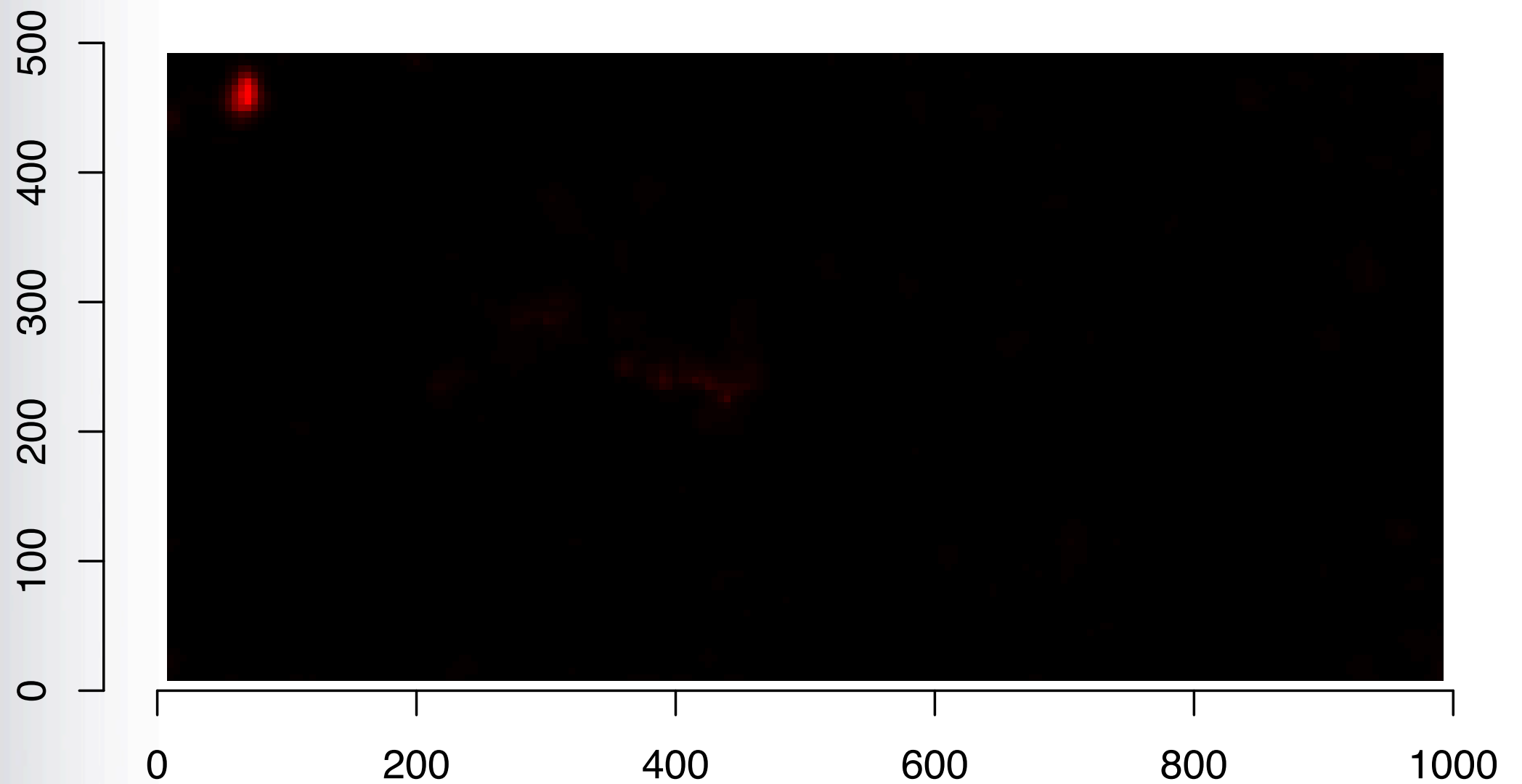
Beta diversity





Revisiting Beta Diversity – 2005

Beta diversity





Temporal beta diversity

- Correctly identifies an interesting area where there were few, but distinct, trees in the 1980s, but now the area has filled in with lots of more normal trees...



Conclusions

- We have a generalisation of existing measures of beta diversity that incorporates similarity
- We have a new concept of individual alpha, beta and gamma diversity, which tell us
 - which specific communities are most distinct
 - which contribute most to overall diversity
 - which are changing most over time
- We can use these to identify areas of interest and well as measuring overall diversity, turnover and beta diversity...



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