

Towards machine ecoregionalization of Earth's landmass using pattern segmentation method

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Abstract

We present and evaluate a quantitative method for delineation of ecophysigraphic regions throughout the entire terrestrial landmass. The method uses the new pattern-based segmentation technique which attempts to emulate the qualitative, weight-of-evidence approach to a delineation of ecoregions in a computer code. An ecophysigraphic region is characterized by homogeneous physiography defined by the cohesiveness of patterns of four variables: land cover, soils, landforms, and climatic patterns. It is expected that such a region is likely to be characterized by a single ecosystem. In this paper, we focus on the first-order approximation of the proposed method - delineation on the basis of the patterns of the land cover alone. We justify this approximation by the existence of significant spatial associations between various physiographic variables. Resulting ecophysigraphic regionalization (ECOR) is shown to be more physiographically homogeneous than existing global ecoregionalizations (Terrestrial Ecoregions of the World (TEW) and Bailey's Ecoregions of the Continents (BEC)). The presented quantitative method has an advantage of being transparent and objective. It can be verified, easily updated, modified and customized for specific applications. Each region in ECOR contains detailed, SQL-searchable information about physiographic patterns within it. It also has a computer-generated label. To give a sense of how ECOR compares to TEW and, in the U.S., to EPA Level III ecoregions, we contrast these different delineations using two specific sites as examples. We conclude that ECOR yields regionalization somewhat similar to EPA level III ecoregions, but for the entire world, and by automatic means.

Keywords: Global ecoregions, Environmental variables, Regionalization, Segmentation, Pattern

1. Introduction

Terrestrial ecoregions (hereafter referred to as ecoregions) are the result of regionalization of land into areal units of homogeneous ecosystem which contrast from surroundings. Because the means of such regionalization are not the part of their definition, ecoregions are an umbrella term with a clear general intent, but with specifics depending on how ecosystems are described and compared (Gonzales, 1966; Jax, 2006; Haber, 2011), on the spatial scale considered, and on the approach to the regionalization procedure.

The need for ecoregions was initially driven by conservation planning (Larsen et al., 1994), but their usage has since expanded to tabulating environmental information in general. Ecoregions are mapped at different scales from global to local. At the broadest scale regionalization of ecoregions relies on climatic, geologic, and geomorphologic divisions (Bailey, 2014). With decreasing spatial scale more attention is given to landscape patterns, vegetation types and biodiversity, and, eventually, at the local scale, attention shifts to specific species of flora and fauna (see, for example, Blasi et al. (2014)).

Several different approaches have been applied to a delineation of ecoregions on the broad scale. Bailey (1989, 2014)

developed a deductive approach wherein delineation of ecoregions follows from identifying environmental variables responsible for differentiating between ecosystems and drawing boundaries where these variables change significantly. Resulting regionalization is known as Bailey's Ecoregions of the Continents (BEC). Olson et al. (2001) applied a synthetic approach wherein ecoregions are delineated based on a large body of previous biogeographical studies. Existing information was refined and synthesized using expert judgment. Resulting regionalization is referred to as Terrestrial Ecoregions of the World (TEW). The similar synthetic methodology was applied on a regional scale to develop the Digital Map of European Ecological Regions (DMEER) (Painho et al., 1996) and the Interim Biogeographic Regionalisation for Australia (IBRA) (EA, 2000). Omernik (1987) used a weight-of-evidence approach to delineate ecoregions in the conterminous U.S. In this approach maps of environmental variables are overlaid and ecoregions are delineated by expert judgment through reconciling differences between variability of individual variables. The difference between Bailey's deductive approach and the weight-of-evidence approach is that whereas in the first the reconciliation follows an a priori determined scheme while in the second it is done on the case-by-case basis.

The issue with the synthetic approach to ecoregionalization (TEW, DMEER, IBRA) lies in the lack of quantitative framework. TEW is a compilation of local regions taken from pre-

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49 existing, independently conducted studies. On one hand, this¹⁰⁶
50 may be viewed as a positive because TEW combines expert¹⁰⁷
51 knowledge of the broad community. On the other hand, there¹⁰⁸
52 are no straightforward means to inspect materials and protocols¹⁰⁹
53 that contributed to the creation of TEW. As there is no under-¹¹⁰
54 lying quantitative framework, there are no criteria to assess the¹¹¹
55 quality of TEW. Therefore, no systematic checks, modifications¹¹²
56 or objective updates to TEW are possible. Moreover, although¹¹³
57 many individual regions in TEW may be well-delineated, as a¹¹⁴
58 whole, TEW lacks overall consistency. A user has no means of¹¹⁵
59 knowing which regions are well-delineated and which are not.¹¹⁶
60 TEW legend conveys a short description of a region which usu-¹¹⁷
61 ally pertains to a combination of region's geography, climate,¹¹⁸
62 and flora. Because regions in TEW lack quantitative descrip-¹¹⁹
63 tion, the inter-regions comparison is limited to contrasting their¹²⁰
64 short descriptions in the legend. ¹²¹

65 The weight-of-evidence approach (Omernik, 1987; Omernik¹²²
66 and Griffith, 2014) also lacks quantitative framework, but, it is¹²³
67 rooted in a clear conceptual framework – “Ecoregions should¹²⁴
68 depict areas of similarity in the collective patterns of all biotic,¹²⁵
69 abiotic, terrestrial, and aquatic ecosystem components with hu-¹²⁶
70 mans being part of the biota.” (Omernik and Griffith, 2014).¹²⁷
71 Regions are delineated manually by experts on the basis of vi-¹²⁸
72 sually perceived breaks in aforementioned patterns. In this ap-¹²⁹
73 proach the resulting ecoregionalization may be consistently de-¹³⁰
74 lined (to a degree that humans perception can be consistent),¹³¹
75 but, like in the case of TEW, a user has no means of determin-¹³²
76 ing the quality of the regionalization. Omernik's legend has the¹³³
77 character similar to that in TEW, the inter-regions comparison¹³⁴
78 is limited to contrasting their descriptions in the legend. ¹³⁵

79 In BEC a delineation of regions follows the Köppen-¹³⁶
80 Trewartha climate classification modified by land cover infor-¹³⁷
81 mation (Bailey, 2014). BEC legend conveys regions' climatic¹³⁸
82 and floristic character. Because of its reliance on the climate,¹³⁹
83 BEC offers only the broadest scale regionalization.

84 An attempt to automate the ecoregionalization process using¹⁴⁰
85 a multivariate k -means clustering algorithm was made by Har-¹⁴¹
86 grove and Hoffman (2005) and followed up by Kumar et al.¹⁴²
87 (2011). In such framework vectors of environmental variables¹⁴³
88 are associated with each pixel (a tract of land corresponding to¹⁴⁴
89 the resolution of the data) and pixels agglomerated into larger¹⁴⁵
90 zones (ecoregions) on the basis of the Euclidean distance be-¹⁴⁶
91 tween these vectors. Such automated approach addresses issues¹⁴⁷
92 related to objectivity, consistency, and inter-region comparabil-¹⁴⁸
93 ity (see our discussion above), however, its ability to yield a¹⁴⁹
94 useful ecoregionalization is limited by the choice of clustering¹⁵⁰
95 as a technique enabling the automation. Clustering leads to a¹⁵¹
96 delineation of non-contiguous, highly fragmented zones, with¹⁵²
97 the fragments spread over wide areas. Clustering may be well-¹⁵³
98 suited for classification but it is ill-suited for mapping. Mapping¹⁵⁴
99 needs to be based on characteristics which are macroscopically¹⁵⁵
100 recognizable (Klijn et al., 1995), which environmental variables¹⁵⁶
101 measured on the scale of an individual pixel are not. ¹⁵⁶

102 In this paper, we propose and describe an approach to data-¹⁵⁷
103 driven machine regionalization of the entire terrestrial landmass¹⁵⁸
104 capable of producing a useful global map of ecophysiological¹⁵⁹
105 regions. We call the resultant regions “ecophysiological” be-¹⁵⁹

cause they are mapped based on physiography but aim at de-
lineating ecosystems as well. This is consistent with the notion that ecoregionalization on larger scales should be based on physiography (Klijn et al., 1995; Sayre et al., 2014). Following Omernik and Griffith (2014), our mapping is based on macroscopically recognizable *patterns* of physiographic categorical variables, but a decision on where to put boundaries between the regions is made by a segmentation algorithm instead of a committee of experts. Segmentation is a natural choice for machine delineation of regions because it is an algorithmic implementation of regionalization. Quantitative assessment of segmentation quality corresponds directly to the qualitative notion (McMahon et al., 2001; Loveland and Merchant, 2004; Omernik and Griffith, 2014) that regions should be internally as homogeneous as possible with respect to the environment, and they should stand out from adjacent regions.

Pattern-based segmentation is the enabling technology behind our proposed method but it also presents a big challenge. This recently developed technology (Jasiewicz et al., 2015, 2017) works at present only with patterns of a single variable, not with patterns of multiple variables as our proposed framework calls for. However, we find a high level of spatial association between categories of various physiographic variables, thus we can achieve a viable regionalization by segmenting the landmass on the basis of patterns of the land cover alone. The quality of such approximation is checked a posteriori.

The goals of this paper are as follows. (1) To describe how pattern-based segmentation technique can be used for automatic creation of a global map and the legend of ecophysiological regions. (2) To demonstrate that a segmentation based only on patterns of land cover yields a viable ecoregionalization. (3) To compare such ecoregionalization with TEW. (4) To provide a spatial database of delineated regions with a detailed quantitative description of patterns in each region.

2. Data and Methods

Table 1 lists four global physiographic datasets we used to calculate associations between categories of land cover, climate, topography, and soils, and to calculate homogeneity of delineated regions. Our choice of environmental variables is very similar to that made by Sayre et al. (2014) except we use newly available (Hengl et al., 2017) soil types data (reclassified to 12 orders) instead of lithology used by Sayre et al. (2014) as a proxy for soils. Note that all variables are categorical. Land cover is arguably the most ecologically important of the four variables because it was demonstrated to provide the first-order information about geographical distribution of biodiversity and ecological processes (Siriwardena et al., 2000; Maes et al., 2003; Eyre et al., 2004; Heikkinen et al., 2004; Fuller et al., 2005; Luoto et al., 2006). Details about the land cover dataset (CCI-LC) including its accuracy can be found in the Land Cover CCI Product User Guide V.2 (ESA, 2017).

2.1. Pattern-based segmentation of Earth's landmass

Segmentation was performed using the Geospatial Pattern Analysis Toolbox (GeoPAT) (Jasiewicz et al., 2015, 2017) – a

Table 1: Global environmental datasets

| Variable | Dataset | Data type | Res. | Source |
|------------|-------------------------------|-------------------------------|-------|---|
| land cover | CCI-LC 2010 | categorical grid (22 classes) | 300 m | http://maps.elie.ucl.ac.be/CCI |
| climate | bioclimatic classification | categorical grid (37 classes) | 250 m | Sayre et al. (2014) modified from Metzger et al. (2013) |
| topography | landforms classification | categorical grid (17 classes) | 250 m | Karagulle et al. (2017) |
| soil | SoilGrids soil classification | categorical grid (12 classes) | 250 m | Hengl et al. (2017) |

collection of GRASS GIS (GRASS Development Team, 2016) modules for carrying out pattern-based analysis of large categorical grids. Pattern-based segmentation differs from the standard pixel-based segmentation by agglomerating sites (tracts of land much larger than an individual pixel) on the basis of patterns of variable rather than agglomerating pixels on the basis of at-pixel values and texture of variables.

Fig. 1 illustrates the basic concept of the pattern-based segmentation algorithm. First, the landmass is tessellated into sites – square blocks (of the size $k \times k$ of CCI-LC cells) to form a new, k^2 coarser, grid of sites (Fig. 1A) Sites are tracts of land large enough to encompass patterns of physiographic variables but small enough to be building blocks of regions. Sites of size $k = 100$ (30 km) are shown in Fig. 1A. A site holds a local pattern (mosaics of pixels assigned different land cover categories); a pattern of the land cover in a selected site is shown in Fig. 1B. Those patterns are numerically described using a co-occurrence histogram (Jasiewicz et al., 2015; Niesterowicz et al., 2016). Co-occurrence histogram encapsulates composition and configuration of the pattern. A level of dissimilarity between two sites is a dissimilarity between their corresponding co-occurrence histograms and is measured by the Jensen-Shannon divergence (Lin, 1991). For more details on the concept of pattern-based segmentation see Supplement S2 as well as Niesterowicz et al. (2016) and Niesterowicz and Stepinski (2017). The number of segments and thus a character of regionalization depend on parameters of the segmentation algorithm. Here we use a default set of parameters derived in Jasiewicz et al. (2017). The size (k) of individual sites relates to the level of physiographic pattern generalization, larger values of k leads to a smaller number of segments. We segmented terrestrial landmass assuming three different site’s sizes: $k = 30$ (9 km), $k = 50$ (15 km), and $k = 100$ (30 km). The smallest chosen size is dictated by a requirement of having enough pixels in a site to form a meaningful pattern, and the largest chosen size is dictated by a desire for not having over-generalized patterns. We refer to resulting regionalizations as ecophysiographic regionalizations (ECORs).

Our pattern-based segmentation algorithm is based on the concept of seeded region growing (Fig.1C). A segment starts from a single site and grows by adding sites from its current perimeter until growth stopping criterion is met; for details see

Jasiewicz et al. (2017). The end result of the segmentation is the landmass divided into regions of cohesive land cover patterns (Fig.1D). We also expect that due to the high level of association between categories of land cover and the categories of the remaining variables (see section 3.1) these regions have cohesive patterns of the remaining variables as well. Calculating quality metrics of obtained regionalization will be able to confirm or confute this expectation.

2.2. Assessing the quality of regionalizations

We assess the quality of ECORs through statistics of regions homogeneity and isolation metrics with respect to patterns of all physiographic variables. We compare these statistics with analogous statistics for regions in BEC, and TEW. In ECORs a single region is associated with each segment. In BEC and TEW regions are individual polygons (land units) in their respective shapefiles. Note that the term “ecoregion” in BEC and TEW does not necessarily refer to a contiguous land unit, instead it refers to a class of such units. There are 96 ecoregions containing 623 land units in BEC, and there are 825 ecoregions containing 14,458 land units in TEW. A regionalization has a good quality if regions are pattern-homogeneous and different from their neighbors (isolated).

To assess homogeneity of a region with respect to a pattern of land cover, landforms, and soils we calculate an inhomogeneity metric. Region’s inhomogeneity is a mutual dissimilarity between all sites within this region. A detailed explanation of inhomogeneity metric is given in Supplement S2 or in Jasiewicz et al. (2017). Inhomogeneity of BEC regions is calculated assuming site’s size of $k = 100$ because of their large sizes, and inhomogeneity of TEW regions is calculated assuming site’s size of $k = 30$ because of their smaller sizes. Inhomogeneity metric has a range 0 to 1, smaller values are better (they indicate larger homogeneity).

Climate changes on large spatial scales, thus climate categories do not form patterns over extents of most regions. Therefore, to assess homogeneity of a region with respect to climate we calculate its Shannon’s entropy, $H = -\sum_{i=1}^m p(i) \log_2 p(i)$, where $p(i)$ is a fraction of region’s area occupied by the category i of the climate variable. The summation is over all $m = 37$ categories of bioclimate (see SupplementS3). Minimum possible value of H is zero and it occurs when a segment is com-

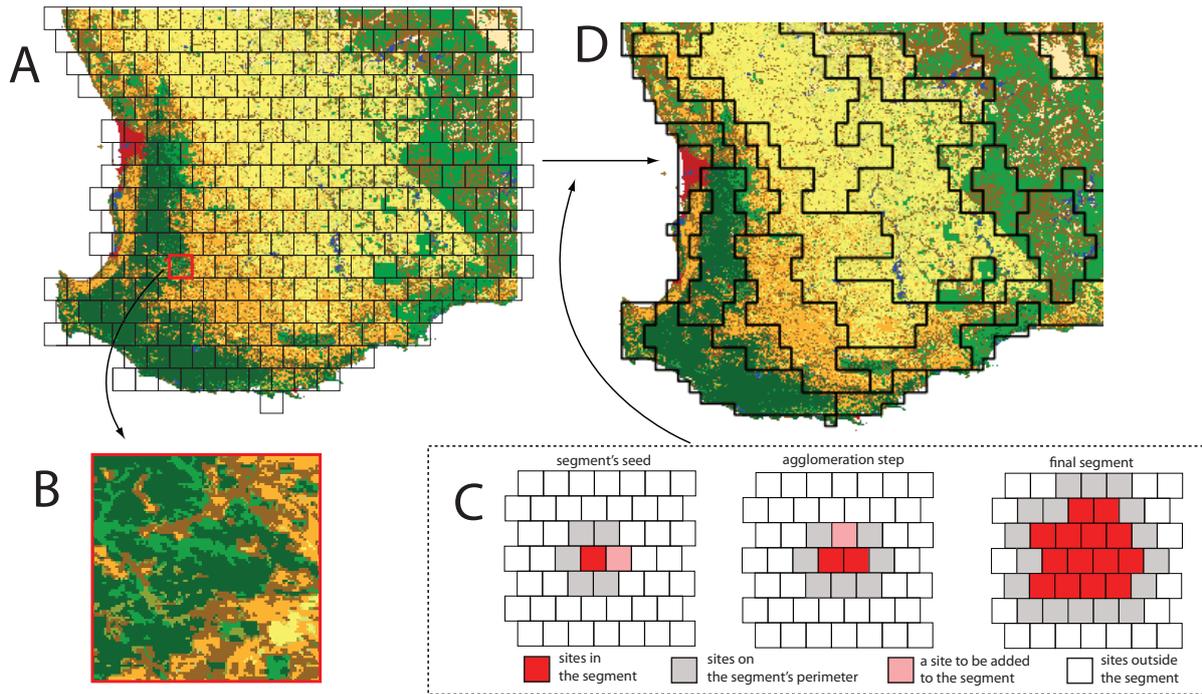


Figure 1: Basic concept of pattern-based segmentation using a fragment of landmass located in the southwestern Australia around the city of Perth. (A) A grid of sites. (B) A zoom-in onto a single $30\text{km} \times 30\text{km}$ site to show its pattern. (C) The concept of seeded region growing algorithm; see the main text for a description. (D) The result of the segmentation algorithm is the regionalization of land cover patterns. The background map is the CCI-LC, different colors indicate different categories of land cover (see Supplement S3 for the legend).

243 pletely within a single climate category (it is completely homo-268
 244 geneous). The larger the value of H the more inhomogeneous269
 245 the segment is with respect to climate. 270

246 To assess how much a pattern in a given region differs from
 247 patterns in neighboring regions we calculate an isolation metric.
 248 To obtain a value of region's isolation metric we calculated an
 249 average dissimilarity (JSD) between the focus region and all of
 250 its immediate neighbors. The average is weighted by the per-
 251 centage of region's perimeter shared with different neighbors.
 252 See Supplement S2 or Jasiewicz et al. (2017) for details. To
 253 calculate isolation with respect to climate, percentages of re-
 254 gion's area occupied by different climate types are used instead
 255 of the co-occurrence histograms in the calculation of JSD. Iso-
 256 lation metric has a range 0 to 1, larger values are better (regions
 257 are more distinct).

258 3. Results

259 3.1. Associations between physiographic variables 271

260 We first estimate a degree of association between our four273
 261 physiographic variables in order to provide a priori rationale for274
 262 using land cover patterns as the only input to the segmentation275
 263 algorithm. We want to check to what degree categories of dif-276
 264 ferent variables co-occur on the scale of our sites. To start we277
 265 regrid the four variables from their native resolutions (see278
 266 Table 1) to grids with $9\text{km} \times 9\text{km}$ and $30\text{km} \times 30\text{km}$ cells us-279
 267 ing the mode values method. Because we deal with categorical280

variables we use Cramér's V measure of association (Cramér,
 2016). Table 2 shows the values of Cramér's V for all combi-
 nations of variables.

Table 2: Degree of association between physiographic variables

| | LC | BC | LF | S | Mean | St.Dev. |
|--|------|------|------|------|------|---------|
| $9\text{km} \times 9\text{km}$ sites | | | | | | |
| LC | n/a | 0.34 | 0.20 | 0.40 | 0.32 | 0.10 |
| BC | 0.34 | n/a | 0.13 | 0.50 | 0.32 | 0.19 |
| LF | 0.20 | 0.13 | n/a | 0.09 | 0.14 | 0.05 |
| S | 0.40 | 0.50 | 0.09 | n/a | 0.33 | 0.21 |
| $30\text{km} \times 30\text{km}$ sites | | | | | | |
| LC | n/a | 0.34 | 0.19 | 0.40 | 0.31 | 0.11 |
| BC | 0.34 | n/a | 0.13 | 0.51 | 0.33 | 0.19 |
| LF | 0.19 | 0.13 | n/a | 0.1 | 0.14 | 0.05 |
| S | 0.40 | 0.51 | 0.1 | n/a | 0.34 | 0.21 |

LC-land cover, BC-bioclimate, LF-landforms, S-soils.

271 Interpretation of Cramér's V values is as follows (Corbett and
 272 LeRoy, 2003): $V < 0.2$ – a weak relation, $V = 0.2$ – 0.25
 – a moderate relationship, $V = 0.25$ – 0.30 – a moderately
 strong relationship, $V = 0.30$ – 0.35 – a strong relationship,
 $V = 0.35$ – 0.40 – a very strong relationship, and $V > 0.4$ –
 a worrisomely strong relationship (two variables measure the
 same concept). Our results in Table 2 indicate that associations
 between land cover, soils and climate are strong, very strong, or
 worrisomely strong. However, landforms are found to be less
 associated with the remaining three variables, although they are

the most associated with land cover (at the edge of the moderate level). Thus, an association analysis reveals that land cover is the best choice of the variable to be used as a sole input to the segmentation algorithm. A priori analysis suggests that obtained regions should be homogeneous with respect to land cover, soils, and climate, but maybe less homogeneous with respect to landforms.

3.2. Regionalizations

ECORs based on 30km × 30km sites, 15km × 15km sites, and 9km × 9km sites yield 9,942, 36,284, and 101,274 regions, respectively. Areas of regions vary greatly from as little as the size of a single site to as much as 1.2×10⁷ km². Those ecoregionalizations are in the form of SQL-searchable spatial databases. The list of attributes for each region includes an ID number, region’s area, the physiography (the area shares of land cover, bioclimate, landforms, and soils categories), values of inhomogeneity and isolation metrics, and the numerical code, which encapsulates a short overall description of a region. The shares of categories provide a detailed numerical description of physiography in each region. A database could be used to search for regions which are similar to each other on the basis of any combinations of categories.

The numerical code gives an information about a region’s physiography compressed to a single, 16-digit number; the list of deciphered codes form a legend to the ECOR map. To make such a compact representation possible we first analyzed statistics of regions’ categories shares (histograms of categories present in a region). It turns out that for all four variables, histograms are either predominantly monothematic or predominantly bi-thematic.

Table 3 shows data in support of this finding. The entries in the table are (percentage of all regions in a given type of histogram (monothematic or bi-thematic) / average percentage of region’s area in either a top category (for monothematic) or in top two categories (for bi-thematic). For example, the entry 14/89 means that 14% of regions have patterns of land cover dominated (on average 89% share of region’s area) by a single category, and the entry 86/79 means that 86% of regions have patterns of land cover dominated by top two categories (on average 79% of such region’s area is occupied by top two categories). Thus, a land cover in a given region can be succinctly described by a four-digit number ABCD, where the first two digits, AB, indicate the top category (one of 22, see Table 1) and the last two digits, CD, indicate the second top category. If a region is monothematic CD=00. This procedure creates 429 unique land cover codes in the 9km sites-base regionalization and 357 unique land cover codes in the 30km site-based regionalization. The same procedure is repeated for remaining variables, and individual four-digit numbers are combined into a single 16-digit number,

$$\text{region's code} = \overbrace{\text{ABCD}}^{\text{land cover}} \underbrace{\text{EFGH}}_{\text{soils}} \overbrace{\text{IJKL}}^{\text{landforms}} \underbrace{\text{MNPR}}_{\text{bioclimate}}$$

The semantic meaning of the code can be deciphered from the legends of the four variables (see Supplement S3). For exam-

Table 3: Statistics of regions category histograms

| | monothematic | bi-thematic | # of codes |
|--|--------------|-------------|------------|
| 9km sites-based regionalization | | | |
| land cover | 14/89 | 86/79 | 429 |
| bioclimate | 74/98 | 26/93 | 307 |
| landforms | 38/96 | 62/80 | 167 |
| soils | 63/96 | 37/91 | 117 |
| 30km sites-base regionalization | | | |
| land cover | 13/90 | 87/77 | 357 |
| bioclimate | 59/96 | 41/89 | 256 |
| landforms | 29/94 | 71/71 | 111 |
| soils | 57/96 | 43/89 | 109 |

See main text for explanation of the entries in the Table.

ple, the code 1207080012001920 has the following meaning: land cover dominated by the mixture of shrubland and needle-leave evergreen forest, soils dominated by mollisols, landform dominated by high mountains, and climate a mixture of warm semi-dry and warm moist. There is only one region with this particular code and it contains Santa Catalina Mountains near Tucson, Arizona, U.S. There are 8251 unique 16-digit codes in the 30km site-based ecoregionalization, and 23,660 unique 16-digit codes in the 9km site-based ecoregionalization. Note that the number of unique existing codes is much smaller than combinatorially possible due to the high correlation between physiographic variables. On the other hand, a large number of unique codes indicates a high diversity of physiographic conditions over the landmass.

ECORs databases, as well as shapefiles for BEC and TEW containing the values of regions’ inhomogeneity and isolations metrics as attributes, are available from <http://sil.uc.edu>.

3.3. Quality of regionalizations

Results of quality of regionalization calculations are summarized in Table 4. This table has three sections showing values of average inhomogeneity, average isolation, and average overall quality, respectively. Averages are calculated over all regions in the regionalization. An overall quality of delineation for a single region is defined as (1 - inhomogeneity/isolation). This metric has a 0 to 1 range with higher numbers indicating better delineation. The quality metric is not applicable to climate because climate’s inhomogeneity and isolation are not measured in the same units. We calculate the standard, unweighted average (the left part of Table 4) and the area-weighted average (the right part of Table 4). Area-weighted average metrics may be better for comparison between different regionalizations due to significant differences between regions area distribution in BEC, TEW, and ECOR.

The numbers in Table 4 should be compared within a single column (for a given variable) to indicate which regionalization has, on average, better-defined regions with respect to a given variable. In general, ECORs regions are more homogeneous but less isolated than TEW and BEC. For the best overall characterization of regionalization, the inhomogeneity and isolation metrics need to be considered together; this is achieved by the quality metric. According to the unweighted method, ECORs

Table 4: Average inhomogeneities and isolations of segments in different regionalizations

| Name | Unweighted | | | | Area-Weighted | | | |
|-------------------------|-------------|-------------|-------------|-------------|---------------|-------------|-------------|-------------|
| | BioClim | Landform | Land Cover | Soils | BioClim | Landform | Land Cover | Soils |
| Average inhomogeneities | | | | | | | | |
| BEC | 1.32 | 0.43 | 0.34 | 0.28 | 1.54 | 0.40 | 0.33 | 0.28 |
| TEW | 0.38 | 0.18 | 0.15 | 0.10 | 1.31 | 0.44 | 0.32 | 0.24 |
| ECOR 9 | 0.37 | 0.22 | 0.13 | 0.07 | 0.81 | 0.31 | 0.08 | 0.10 |
| ECOR 15 | 0.47 | 0.23 | 0.12 | 0.09 | 0.89 | 0.31 | 0.08 | 0.11 |
| ECOR 30 | 0.62 | 0.22 | 0.12 | 0.10 | 1.00 | 0.27 | 0.08 | 0.11 |
| Average isolations | | | | | | | | |
| BEC | 0.32 | 0.56 | 0.49 | 0.41 | 0.38 | 0.51 | 0.46 | 0.40 |
| TEW | 0.29 | 0.51 | 0.41 | 0.32 | 0.37 | 0.55 | 0.48 | 0.36 |
| ECOR 9 | 0.12 | 0.36 | 0.29 | 0.17 | 0.24 | 0.39 | 0.25 | 0.13 |
| ECOR 15 | 0.15 | 0.37 | 0.28 | 0.18 | 0.25 | 0.43 | 0.26 | 0.14 |
| ECOR 30 | 0.20 | 0.36 | 0.28 | 0.21 | 0.28 | 0.37 | 0.25 | 0.19 |
| Average quality | | | | | | | | |
| BEC | n/a | 0.22 | 0.29 | 0.31 | n/a | 0.21 | 0.34 | 0.32 |
| TEW | n/a | 0.61 | 0.60 | 0.63 | n/a | 0.22 | 0.38 | 0.38 |
| ECOR 9 | n/a | 0.44 | 0.55 | 0.51 | n/a | 0.29 | 0.69 | 0.47 |
| ECOR 15 | n/a | 0.41 | 0.56 | 0.49 | n/a | 0.28 | 0.66 | 0.46 |
| ECOR 30 | n/a | 0.40 | 0.57 | 0.50 | n/a | 0.29 | 0.61 | 0.47 |

The best value for each variable is indicated in the bold font. n/a – not applicable. 9, 15, and 30 in ECOR regionalizations refer to the size of a single site in km.

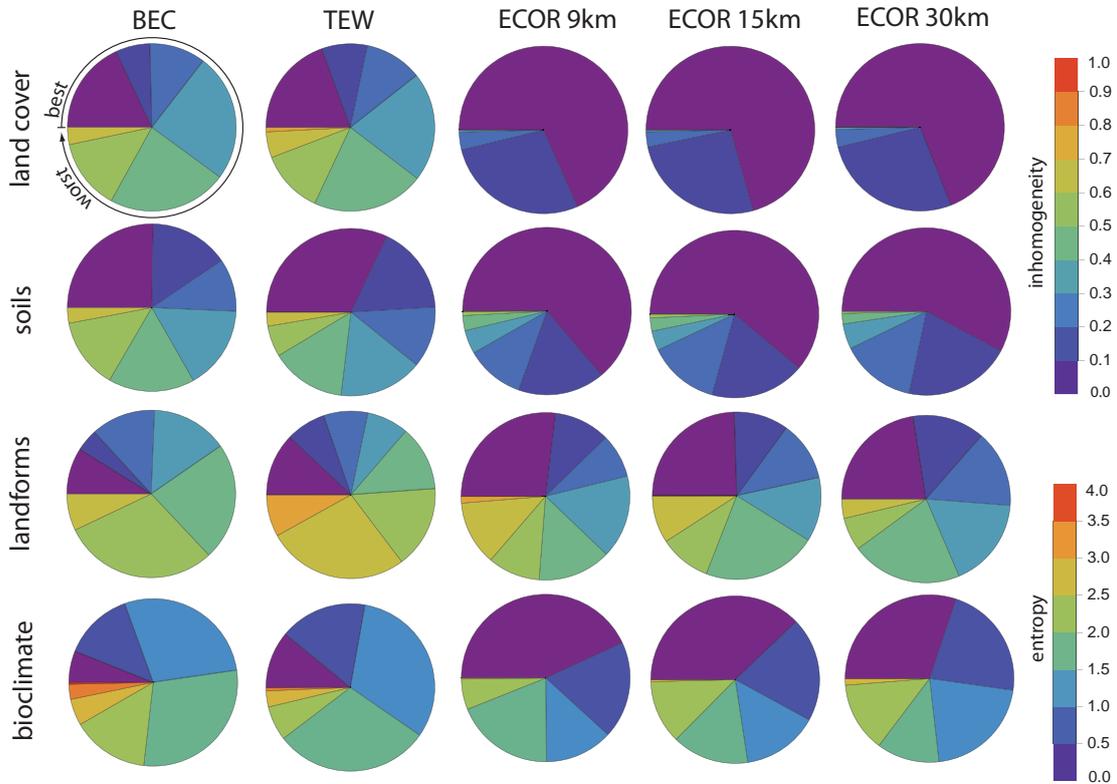


Figure 2: Pie diagrams illustrating division of Earth's landmass into zones of different levels of inhomogeneity. Rows correspond to different physiographic variables and column correspond to different regionalizations. The top legend pertains to land cover, soils, and landforms, and the bottom legend pertains to bioclimate .

374 are characterized by smaller values of quality than TEW but⁴³¹
375 by higher values of quality than BEC. According to the area-⁴³²
376 weighted method, ECORs are characterized by higher values⁴³³
377 of quality than both TEW and BEC. ⁴³⁴

378 For landforms, land cover, and soils, the numbers in Table 4⁴³⁵
379 could also be compared within a row (for a given regionaliza-⁴³⁶
380 tion) to indicate, on average, a quality of a region delineation⁴³⁷
381 with respect to patterns of different physiographic variables.⁴³⁸
382 As expected, ECORs regions are best delineated with respect⁴³⁹
383 to the land cover. The value of 0.57 (unweighted quality for⁴⁴⁰
384 land cover in ECOR 30) can be interpreted as follows: in an⁴⁴¹
385 average region, the similarity of its constituent sites with re-⁴⁴²
386 spect to patterns of land cover is 2.3 times higher than an av-⁴⁴³
387 erage similarity of land cover patterns between this region and⁴⁴⁴
388 its neighbors. Following this interpretation for patterns of soils⁴⁴⁵
389 and landforms yields the ratios of 2 and 1.67, respectively. This⁴⁴⁶
390 result is consistent with our expectations based on associations⁴⁴⁷
391 between physiographic variables (section 3.1). ⁴⁴⁸

392 Homogeneity of regions with respect to bioclimate requires⁴⁴⁹
393 a separate discussion because it is measured by the entropy. To⁴⁵⁰
394 get some intuition to the meaning of entropy values we give⁴⁵¹
395 few examples. In the region where 90% of the area has climate⁴⁵²
396 A and 10% of the area has climate B the value of entropy is⁴⁵³
397 0.47. If the region is divided equally between two climates the⁴⁵⁴
398 entropy value is 1. Small regions are covered by a single cli-
399 mate and have entropy values equal to 0. All regionalizations,
400 except the BEC, are, on average, climate-homogeneous. Aver-⁴⁵⁵
401 age values of isolation with respect to bioclimate must be small
402 because most regions are small and are surrounded by regions⁴⁵⁶
403 with the same climate type. ⁴⁵⁷

404 Based on results in Table 4 we conclude that our method⁴⁵⁸
405 yields a very good regionalization of land cover patterns (qual-⁴⁵⁹
406 ity = 0.55/0.69 using unweighted/area-weighted method for⁴⁶⁰
407 ECOR 9). It also yields a reasonable regionalization of the⁴⁶¹
408 entire physiography with the average quality (calculated from⁴⁶²
409 land cover, soils, and landforms) equal to 0.5/0.48 (using⁴⁶³
410 unweighted/area-weighted method for ECOR 9). For compari-⁴⁶⁴
411 son, the average quality for TEW is 0.61/0.32, and the average⁴⁶⁵
412 quality for BEC is 0.27/0.29. Note a significant difference be-⁴⁶⁶
413 tween the unweighted and area-weighted values of quality for⁴⁶⁷
414 TEW. This is explained by the fact that distribution of region⁴⁶⁸
415 areas in TEW is heavily skewed toward very small regions. In⁴⁶⁹
416 TEW a small number of large regions occupy almost the entire⁴⁷⁰
417 landmass, and a large number of small regions occupy a small⁴⁷¹
418 fraction of the landmass. ⁴⁷²

419 In addition to comparing regionalization on the basis of met-⁴⁷³
420 rics in Table 4, we also compare them on the basis of percent-⁴⁷⁴
421 age of landmass grouped into regions of high homogeneity of⁴⁷⁵
422 a pattern. Fig. 2 shows pie diagrams illustrating a division of⁴⁷⁶
423 landmass into zones characterized by different levels of inho-⁴⁷⁷
424 mogeneity with respect to a pattern of a given physiographic⁴⁷⁸
425 variable. An area of each circle represents the area of an entire⁴⁷⁹
426 terrestrial landmass and slices represent proportions of land-⁴⁸⁰
427 mass area covered by regions with inhomogeneity values as⁴⁸¹
428 encoded by their colors. Comparing pie diagrams in a given⁴⁸²
429 row inform about differences between overall homogeneities⁴⁸³
430 of regions in different regionalizations with respect to a given⁴⁸⁴

variable. ECORs clearly divides the land in a way that maxi-
mizes the percentage of landmass grouped into homogeneous
patterns.

Finally, we have produced maps showing geographical distri-
butions of inhomogeneity values (see Supplement S1). ECORs
maps of inhomogeneity with respect to bioclimate reveals that
its relatively higher overall inhomogeneity value stems mostly
from a few large segments in arid areas (like, for example, the
Sahara Desert). In these places, our algorithm delineates very
large segments because arid areas are large tracts of same land
cover. However, the bioclimatic classification assigns few dif-
ferent arid climate categories to these areas resulting in an in-
creased value of inhomogeneity metric. However, these regions
are still covered in their entirety by the arid climate. Simi-
larly, ECORs maps of inhomogeneity with respect to patterns of
landforms reveals that some regions of uniform land cover (for
example, the Amazonian forest) contain multiple categories of
landforms classification. Overall, the limitation of using only
patterns of land cover for ecoregionalization manifest itself in
cases where topographically different areas are covered by the
same land cover, or where large areas of the same land cover
extend through more than one climatic zone. Even with this
limitations, the maps in Supplement S1 shows that ECOR out-
performs TEW and BEC.

4. Discussion

ECOR is the first attempt to obtain a global map of ecophys-
iographic regions purely by means of an autonomous pattern-
based segmentation algorithm. Pixel-based segmentation was
previously used by Bisquert et al. (2015) for regionalization of
France using MODIS time series imagery, but no attempt was
made to check whether obtained segments are homogeneous in
terms of landscapes, soils, climate, or other physiographic vari-
ables. In section 2.1 we described our overall strategy for such
automatic regionalization as well as an implementation of this
strategy given the present status (the single layer-based segmen-
tation) of the enabling technology. After performing analysis
of associations between four physiographic variables (section
3.1) we determined that patterns of land cover are best suited
for the single layer-based segmentation. Land cover is also a
natural choice because it can be used as a proxy for vegetation
structure. In turn, vegetation can be used as a proxy for bi-
otic composition (Kerr et al., 2001; Pearson et al., 2004; Luoto
et al., 2007; Coops et al., 2009) because it provides habitat re-
sources for species. For these reasons, land cover is often used
to provide the first-order information about geographical distri-
bution of biodiversity and ecological processes (Siriwardena
et al., 2000; Eyre et al., 2004; Heikkinen et al., 2004; Fuller
et al., 2005; Luoto et al., 2006). We also found enough asso-
ciation between all the variables to expect that the land cover-
based regionalization may indeed be a viable ecophysiological
regionalization.

The key to evaluating whether ECOR is a viable ecoregion-
alization is our criterion that the regions should, at the mini-
mum, contain cohesive patterns of all physiographic variables

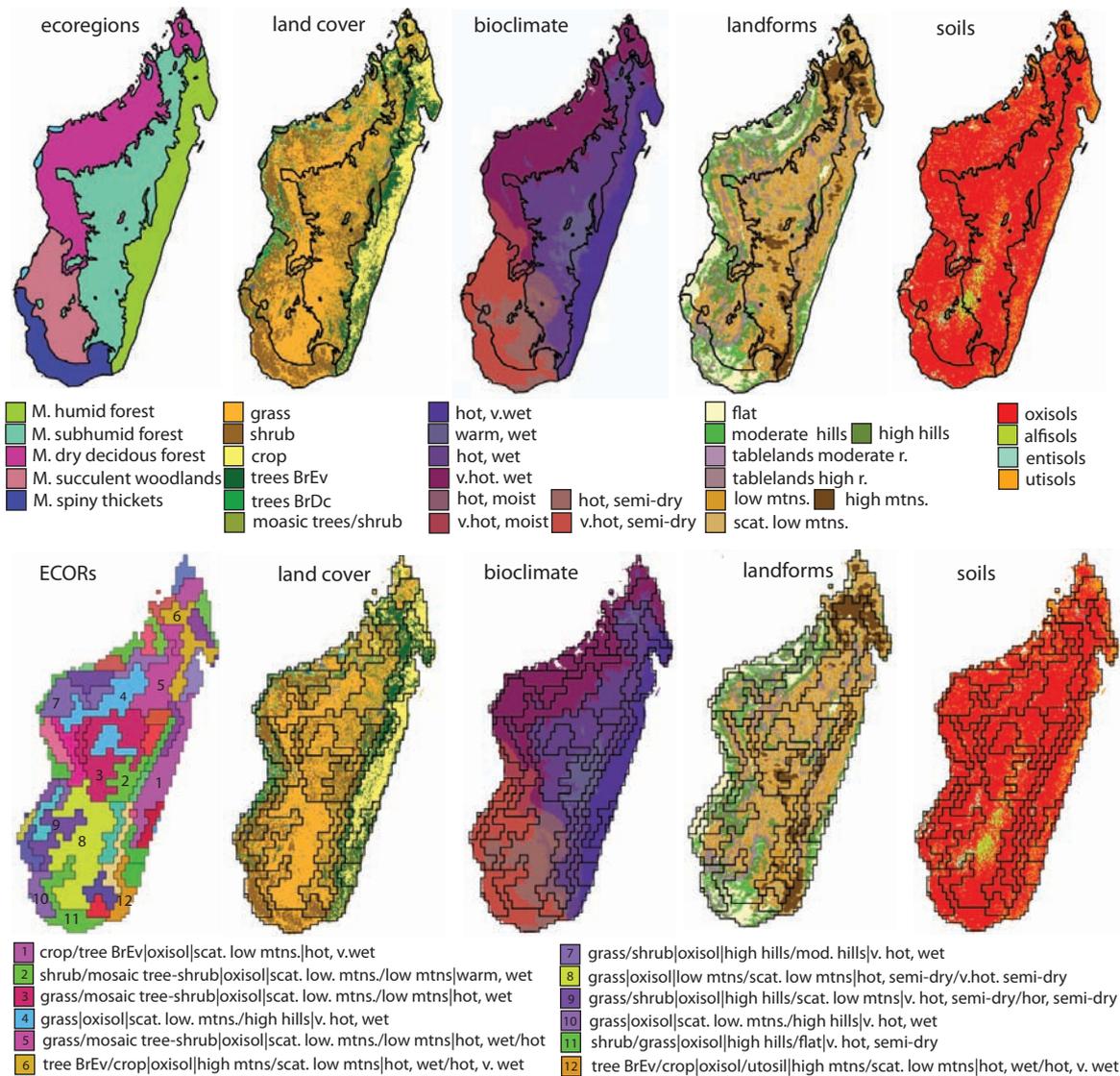


Figure 3: Comparison of ecoregionalizations in TEW and ECOR 30km using the island of Madagascar as an example. The upper row of maps shows TEW regions and how they divide the island's physiography. The lower row of maps shows the same for ECOR. Abbreviations: M. – Madagascar, v. – very, r. – relief, scat. – scattered, BrEv – broadleaf evergreen, mtns. – mountains.

485 – a quality quantitatively measured by the inhomogeneity met-500
 486 ric. The analysis presented in section 3.3 shows that although-501
 487 ECOR does not yet fully meet patterns cohesiveness criterion, 502
 488 it meets it to the sufficient degree to be considered a viable 503
 489 ecoregionalization. The argument for that follows from the fact 504
 490 that ECOR meets patterns cohesiveness criterion to a higher 505
 491 degree than BEC and TEW (see Table 4, Fig. 3, and Supplement 506
 492 S1), the two regionalizations of landmass generally accepted as 507
 493 ecoregionalizations. 508

494 The higher cohesiveness of patterns in ECOR follows mostly 509
 495 from its design and from the existence of the spatial associa-510
 496 tion between categories of physiographic variables. Isolation-511
 497 of ECOR regions is on average smaller than for regions in BEC-512
 498 and TEW. The overall quality of ECOR regionalization is much-513
 499 higher than the quality if BEC regionalization, and comparable-514

or higher (depending on the type of measurement) to the quality of TEW regionalization.

Fig. 3 shows a difference between TEW and ECOR using the island of Madagascar as an example. The most noticeable difference between the two regionalizations is the number of regions, 5 for TEW and 55 for ECOR. A large number of ECOR regions reflects its design – the algorithm painstakingly delineates all variations in the pattern of land cover. Closer inspection reveals that indeed each ECOR region contains a homogeneous pattern of land cover, and to a somewhat lesser degree, a homogeneous pattern of the entire physiography. In Fig. 3 we also included a portion of algorithm-generated legend for 12 out of 55 ECOR regions. Note that this legend is quite specific as it informs on the state of each physiographic variable in the region. However, the auto-generated legend does not contain

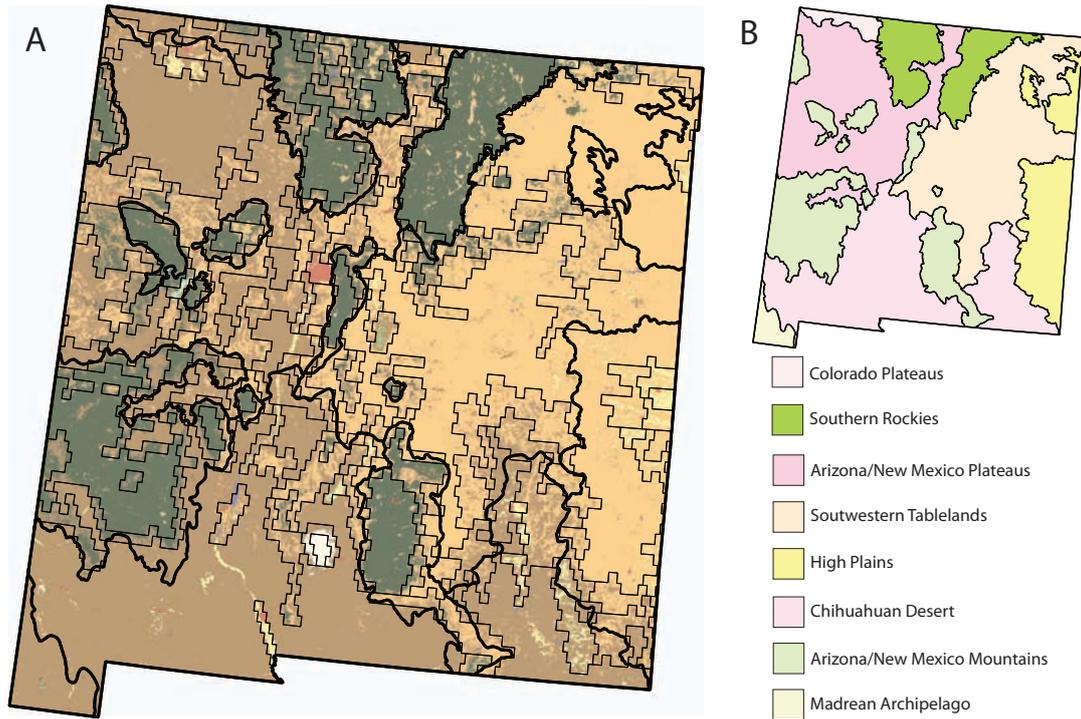


Figure 4: Comparison of ECOR 9km and EPA Level III ecoregionalizations of the state of New Mexico, U.S. (A) EPA ecoregions (thick lines) and ECOR regions (thin lines) overlying the map of land cover. (B) Eight EPA Level III ecoregions in New Mexico.

any specific information available only through on the ground⁵⁴³
inspection.⁵⁴⁴

TEW delineates five ecoregions in Madagascar. Note that⁵⁴⁵
boundaries of TEW regions divide pretty well the climate, and⁵⁴⁶
two of them (humid forest and spiny thickets) are delineating⁵⁴⁷
patterns of land cover (although not to the same precision as⁵⁴⁸
ECOR), but the landforms are definitively not well divided by⁵⁴⁹
TEW ecoregions. The most inaccurate part of the TEW are the⁵⁵⁰
names of ecoregions. Four of them have “forest” or “ wood⁵⁵¹
land” in their names even so Madagascar lost about 80% of its⁵⁵²
original forest, and the forest is presently very scarce across the⁵⁵³
island (see the land cover map). We speculate that these names⁵⁵⁴
originated before the island was deforested. Such dramatic land⁵⁵⁵
change must have change island’s ecosystems, so TEW division⁵⁵⁶
may not be any longer valid for the present day Madagascar.⁵⁵⁷
This goes to the difficulty of updating manual regionalizations.⁵⁵⁸

Fig. 4 compares ECOR with the EPA Level III Ecoregions⁵⁵⁹
of the U.S. (Omernik, 1987; Omernik and Griffith, 2014) using⁵⁶⁰
the state of New Mexico as an example. Both, ECOR and EPA⁵⁶¹
rely on patterns of environment for their delineation, except that⁵⁶²
ECOR delineation is algorithmic and EPA delineation is manual.⁵⁶³
Because both regionalizations follow the same underlying⁵⁶⁴
concept we expect a higher level of correspondence between⁵⁶⁵
ECOR and EPA than between ECOR and TEW.⁵⁶⁶

Indeed, a clear correspondence between the two regionaliza-⁵⁶⁵
tions is observed in Fig. 5A. Each EPA ecoregion is dominated⁵⁶⁶
by an ECOR region. The Chihuahuan Desert is dominated⁵⁶⁷
by a region characterized as (shrub; aridisols/mollisols; scat.⁵⁶⁸

low mtns./low mtns.; warm, semi-dry/cool, semi-dry). Arizona/
New Mexico Mtns. is dominated by (tree NeEv; mollisols;
low mtns./high mtns; cool, semi-dry/cool, moist). Arizona/
New Mexico Plateaus is dominated by (shrub; entisols/aridisols, high hills/scat. low mtns.; cool, semi-dry). Southwestern
Rockies are dominated by (tree NeEv; alfisols/mollisols; high
mtns./scat. low mtns.; cool, semi-dry/cold, moist). The two re-
gions, Southwestern Tablelands and High Plains are dominated
by the same ECOR region (grass; mollisols/aridisols; moderate
hills/flat; warm, semi-dry/cool, semi-dry). They differ by pre-
dominant landforms which the present version of segmentation
was not able to take into account.

In addition, ECOR also delineated smaller regions, where
pattern of land cover departs from surroundings. For example,
in the Chihuahuan Desert ecoregion, there are several inclu-
sions, one is the large field of white sand dunes, and another the
San Andreas mountains just west of the dunes. ECOR delin-
eated these features as independent regions, whereas they ap-
pear only at the higher, IV Level of the EPA mapping.

5. Conclusions

A possibility of delineating ecoregions using quantitative
methodology was discussed (McMahon et al., 2001; Loveland
and Merchant, 2004) and attempted by Hargrove and Hoffman
(2005) using multivariate clustering. However, the quantitative
method presented in this paper is the first to achieve some level
of success. This is because, instead of relying on clustering, it

630 employs a method that attempts to emulate in computer code
631 the qualitative, weight-of-evidence approach. The presented
632 global delineation of ecophysiological regions (ECOR) is the
633 first iteration of this new method.
634

635 In addition to describing the method behind ECOR, we make
636 available the complete, worldwide database of ECOR regions
637 so that the scientific community can evaluate its usefulness for
638 various tasks. We have already identified several areas where
639 ECOR can be useful. At the minimum, it offers a valuable
640 “first draft map” for analysts to manually modify it using their
641 expert knowledge. This would save a lot of time and effort,
642 and expedite updating existing maps, such as TEW. It would,
643 perhaps, make possible a construction of the EPA-style map of
644 ecoregions on the global scale. ECOR makes available detailed
645 quantitative information about physiographic patterns in each
646 region. Moreover, this information is SQL-searchable. As such
647 data was not previously available, we need to start thinking how
648 it could be utilized.
649

650 ECOR will get an update when the pattern-based segmen-
651 tation technology achieves a multi-layer capability. The chal-
652 lenge of segmenting on the basis of multiple patterns simulta-
653 neously is how to incorporate similarities between patterns of
654 individual variables into a similarity of the common, physio-
655 graphic patterns. We expect that such update will result in im-
656 provement of regions’ physiographic homogeneity, but at the
657 cost of an even larger number of regions.
658

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661

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Figure S1: Maps of regions' inhomogeneity values with respect to patterns of land covers in different ecoregionalizations.

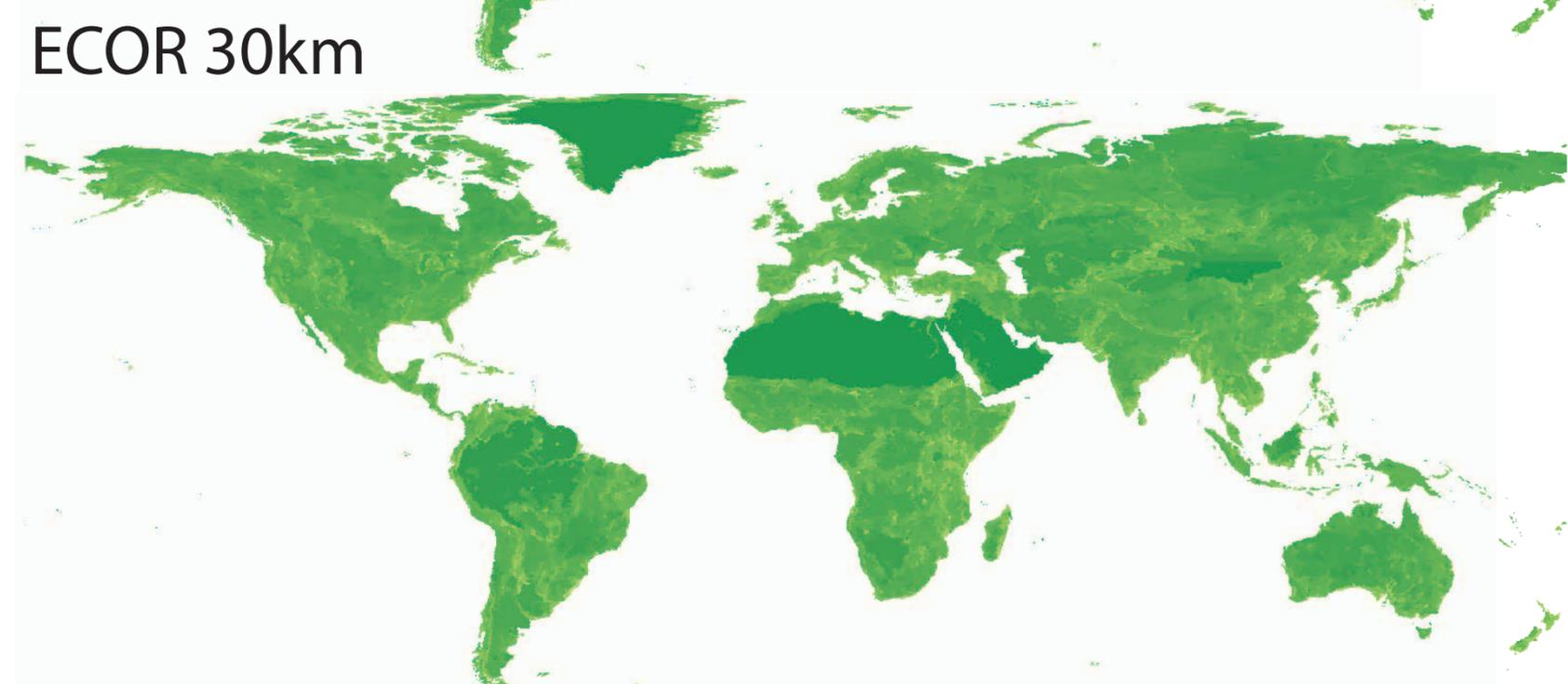
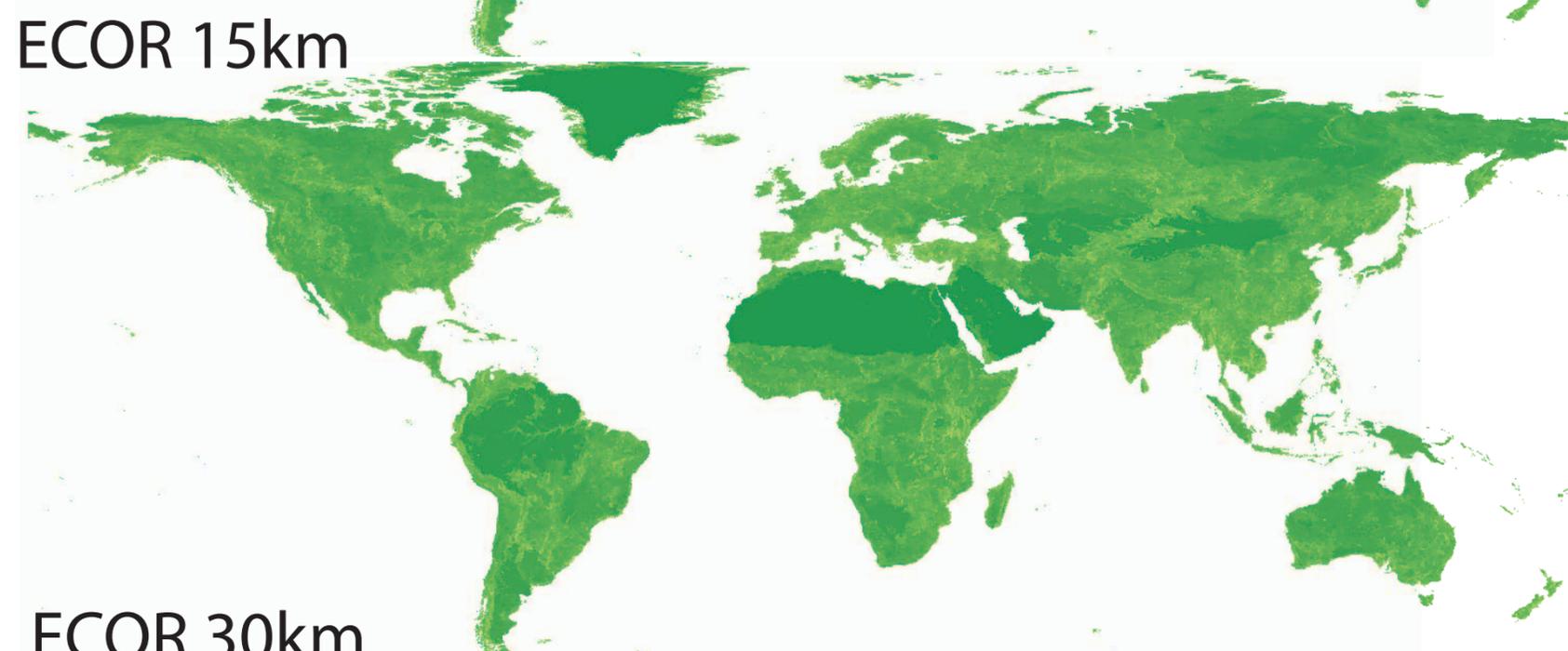
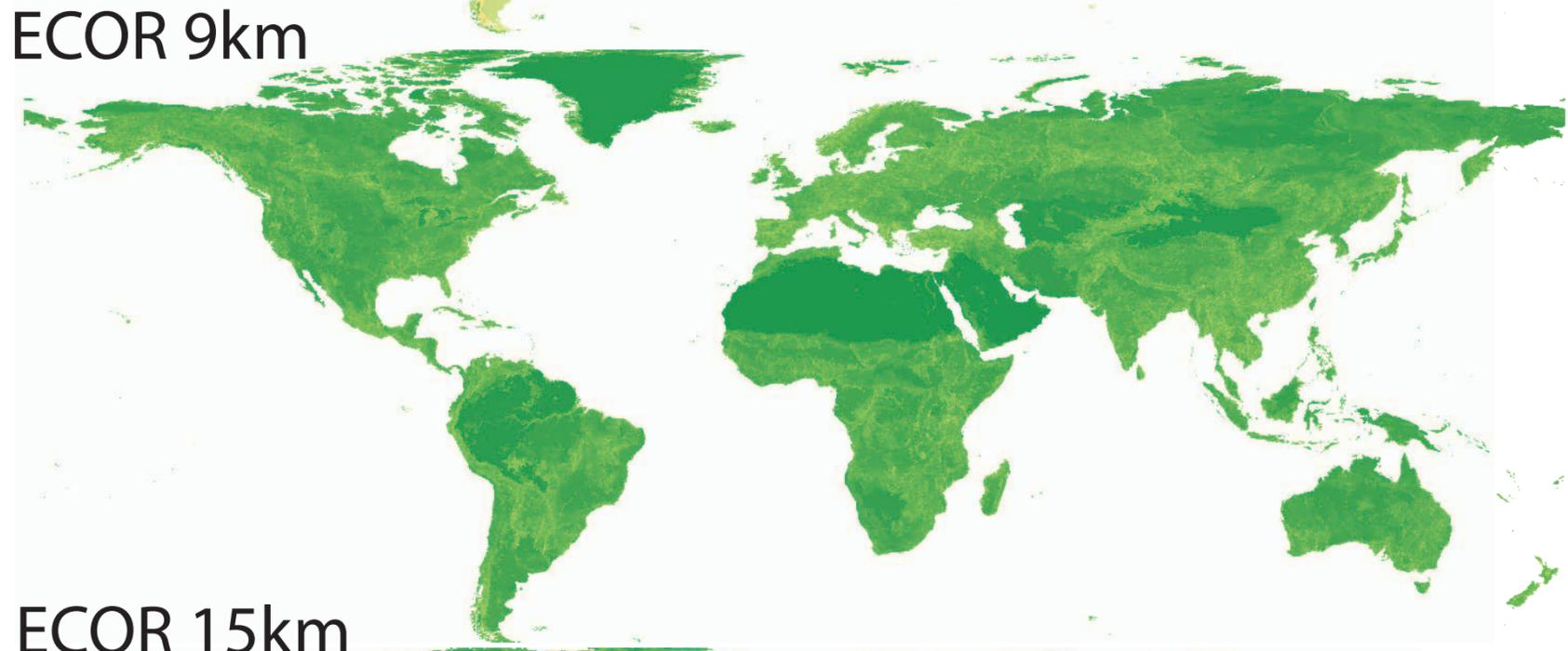
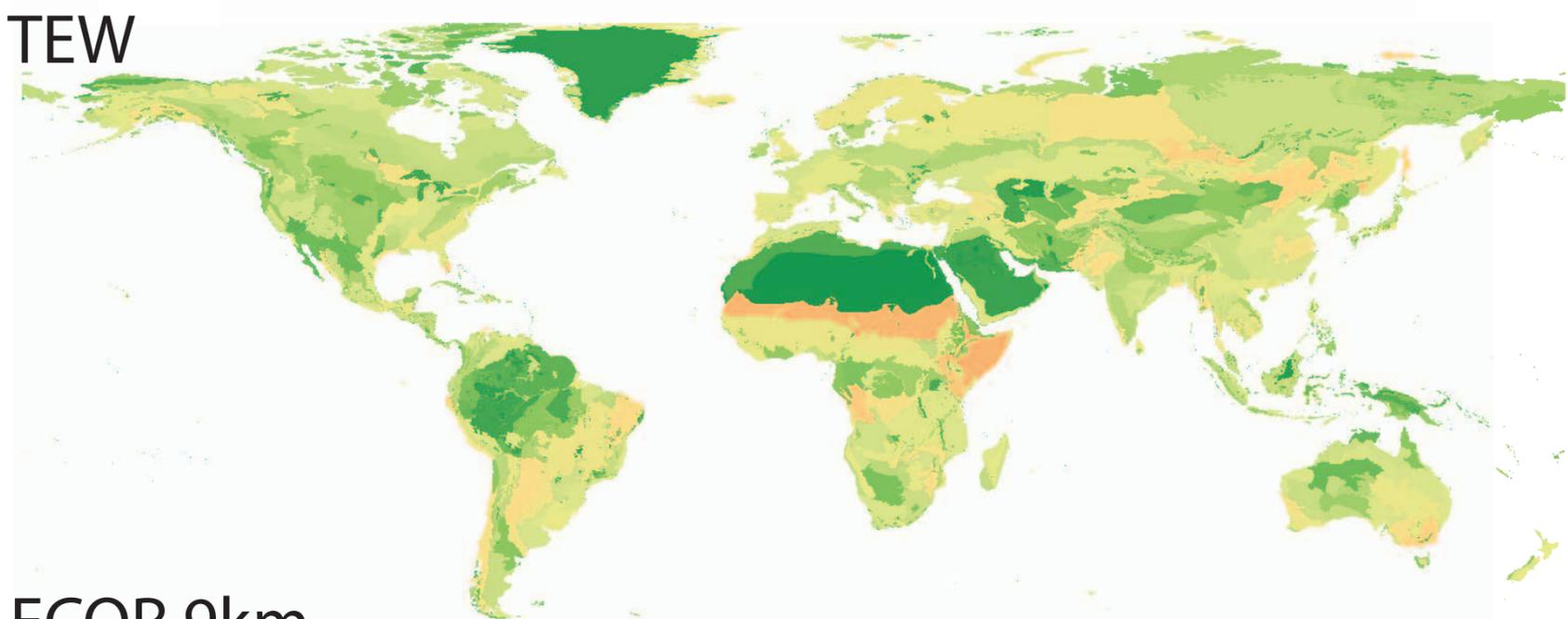
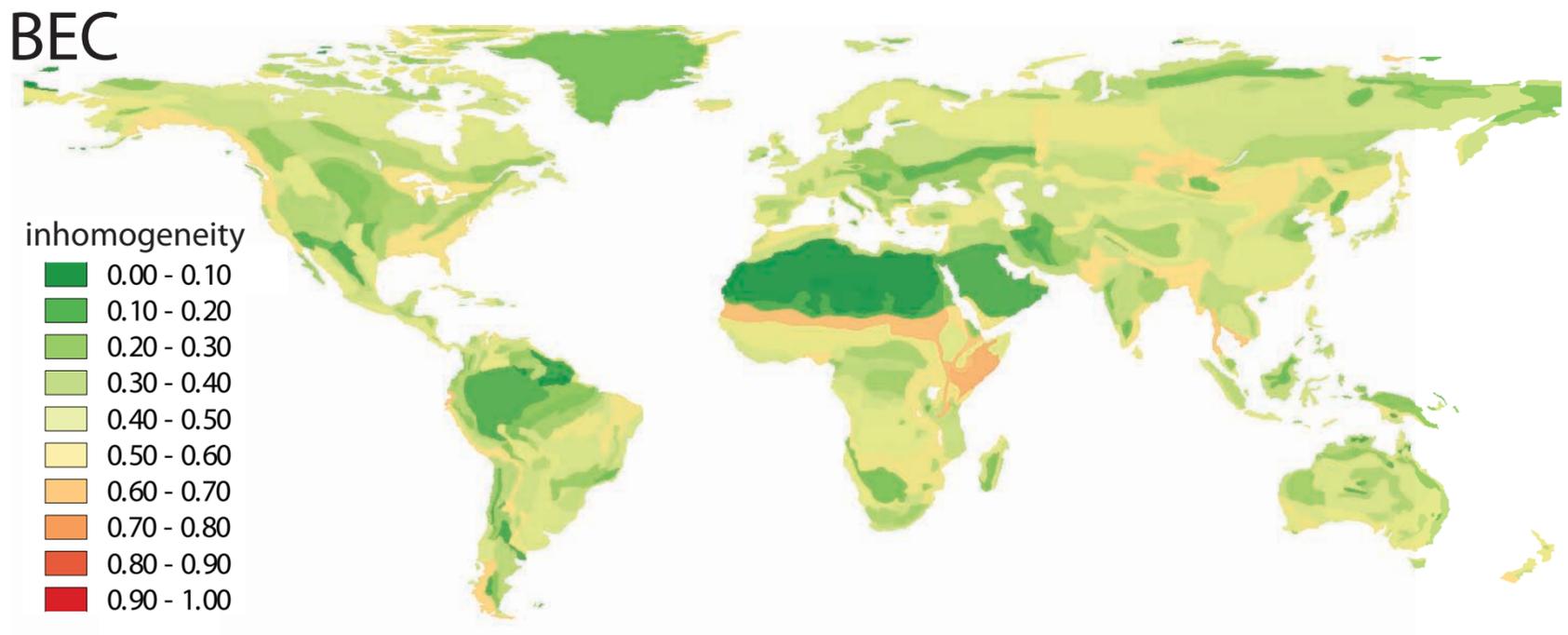


Figure S2: Maps of regions' inhomogeneity values with respect to patterns of soils in different ecoregionalizations

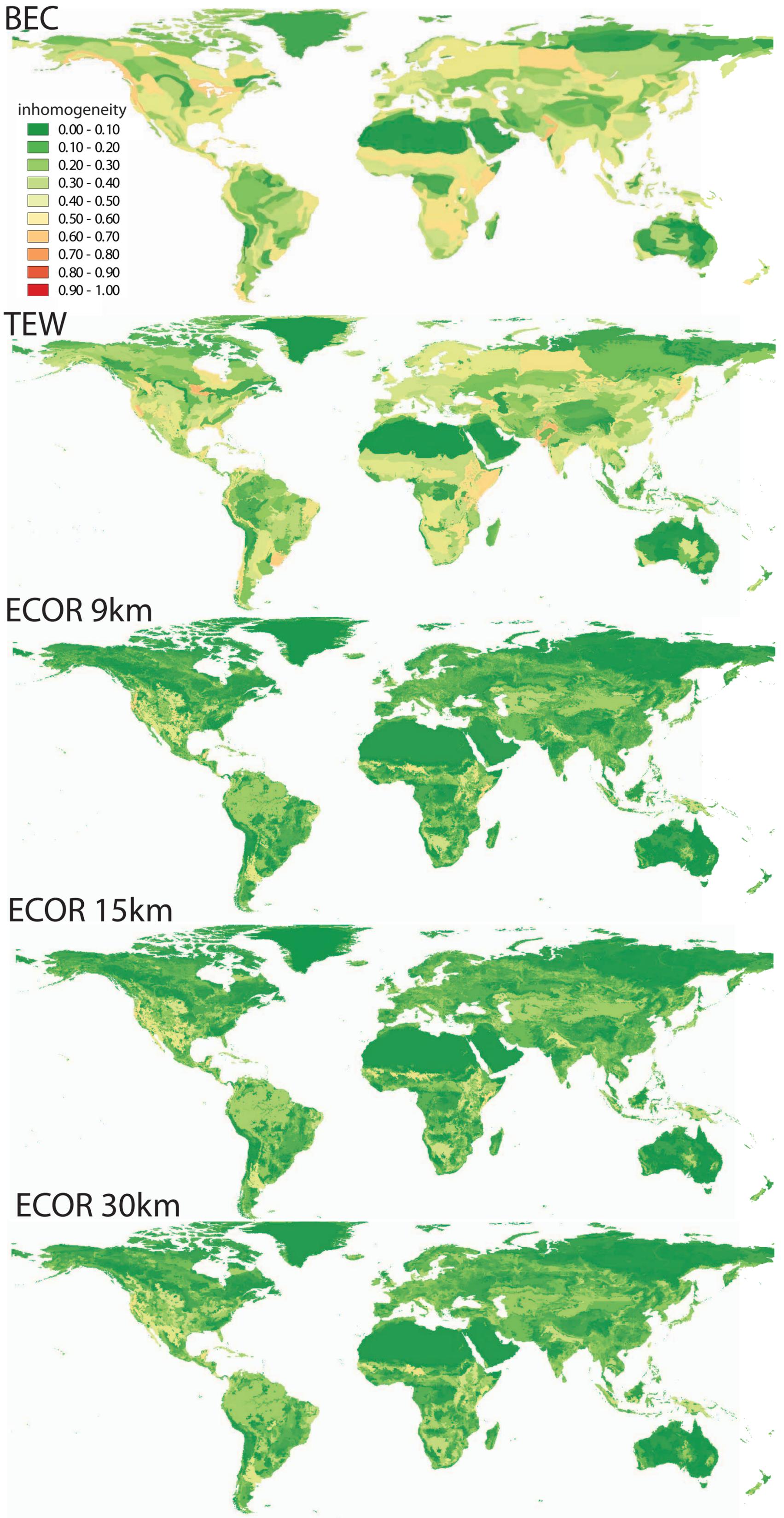


Figure S3: Maps of regions' inhomogeneity values with respect to patterns of landforms in different ecoregionalizations.

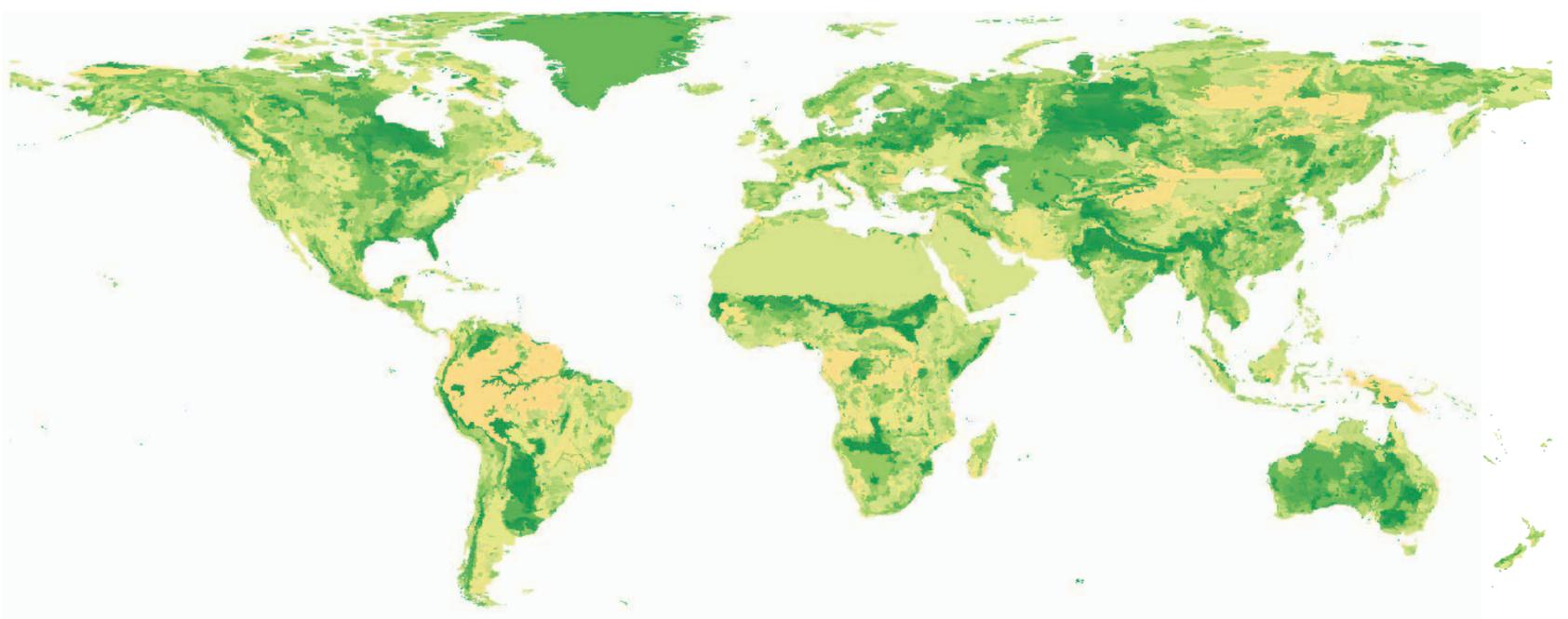
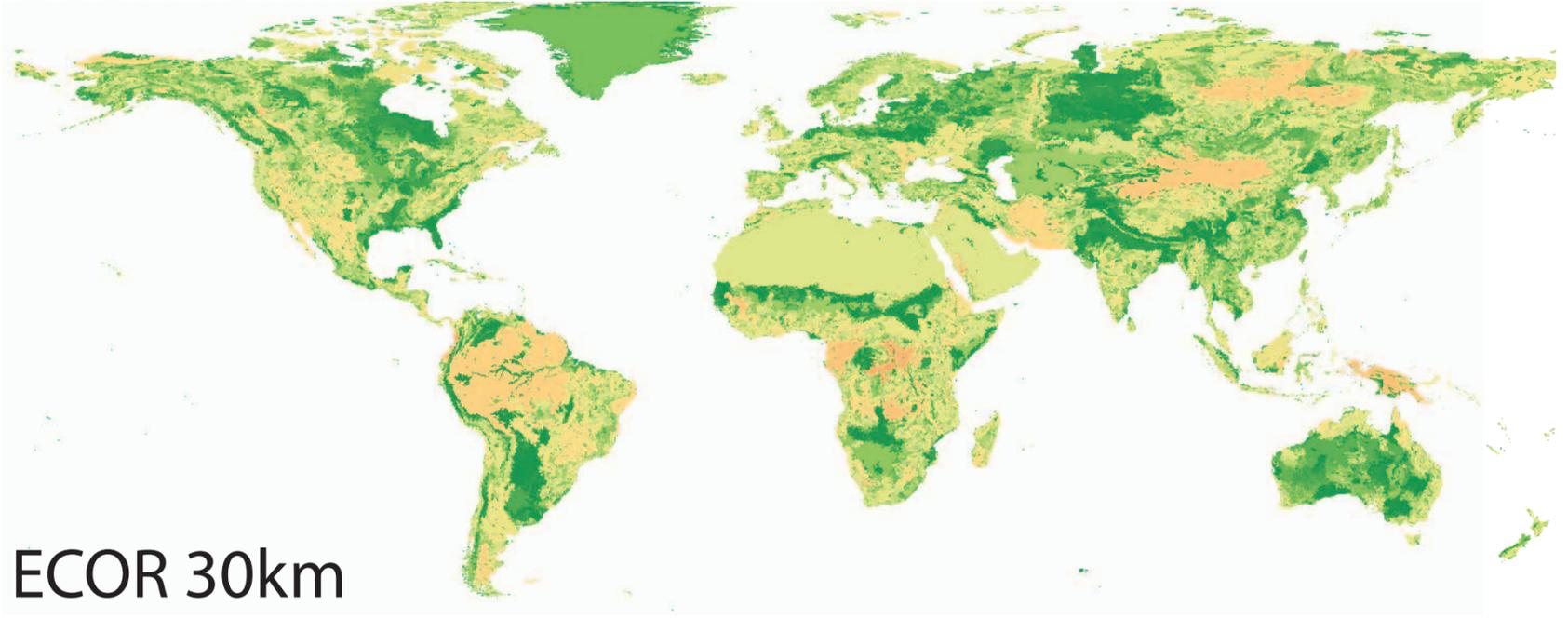
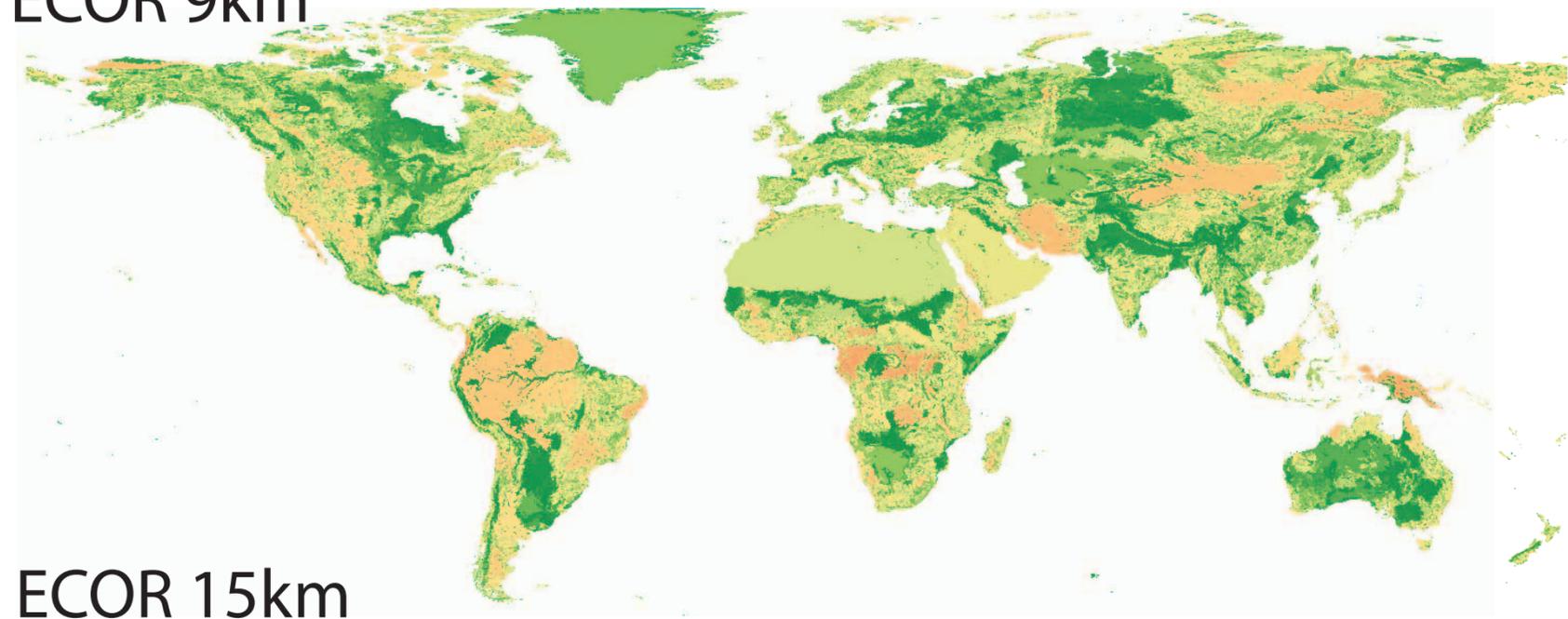
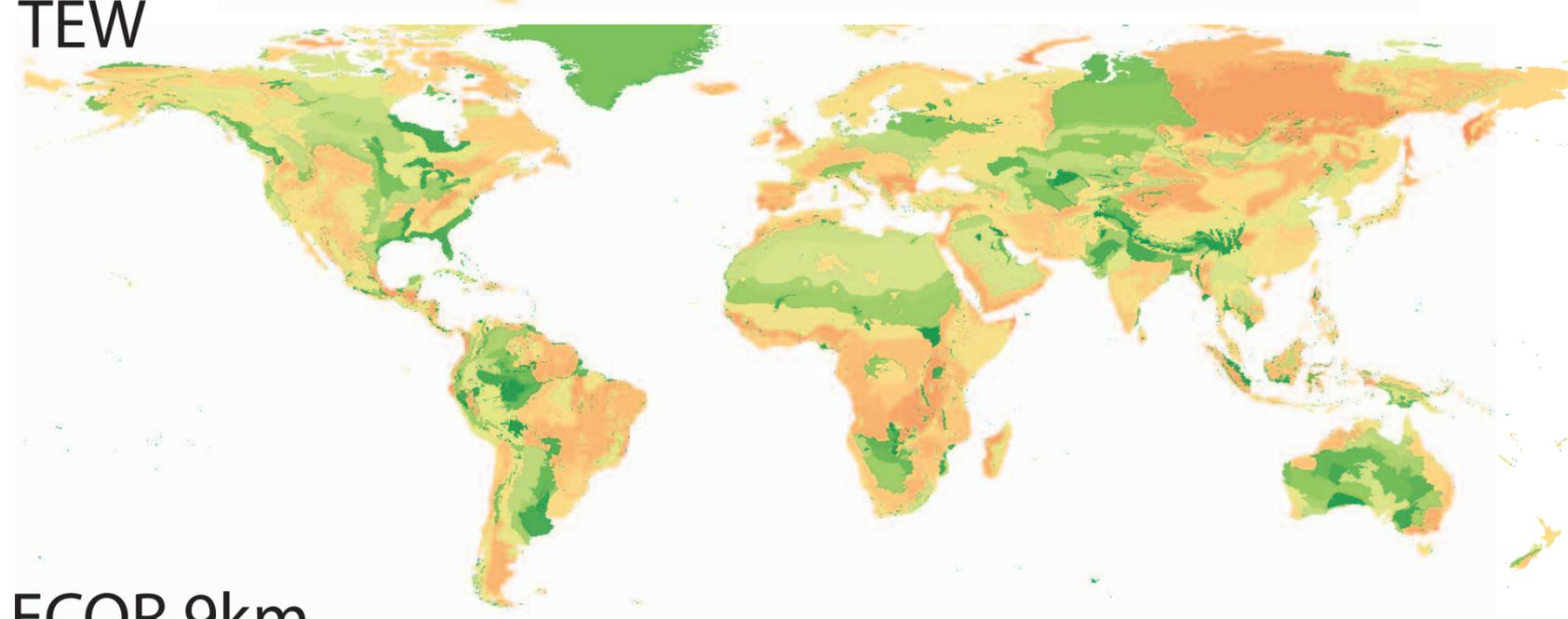
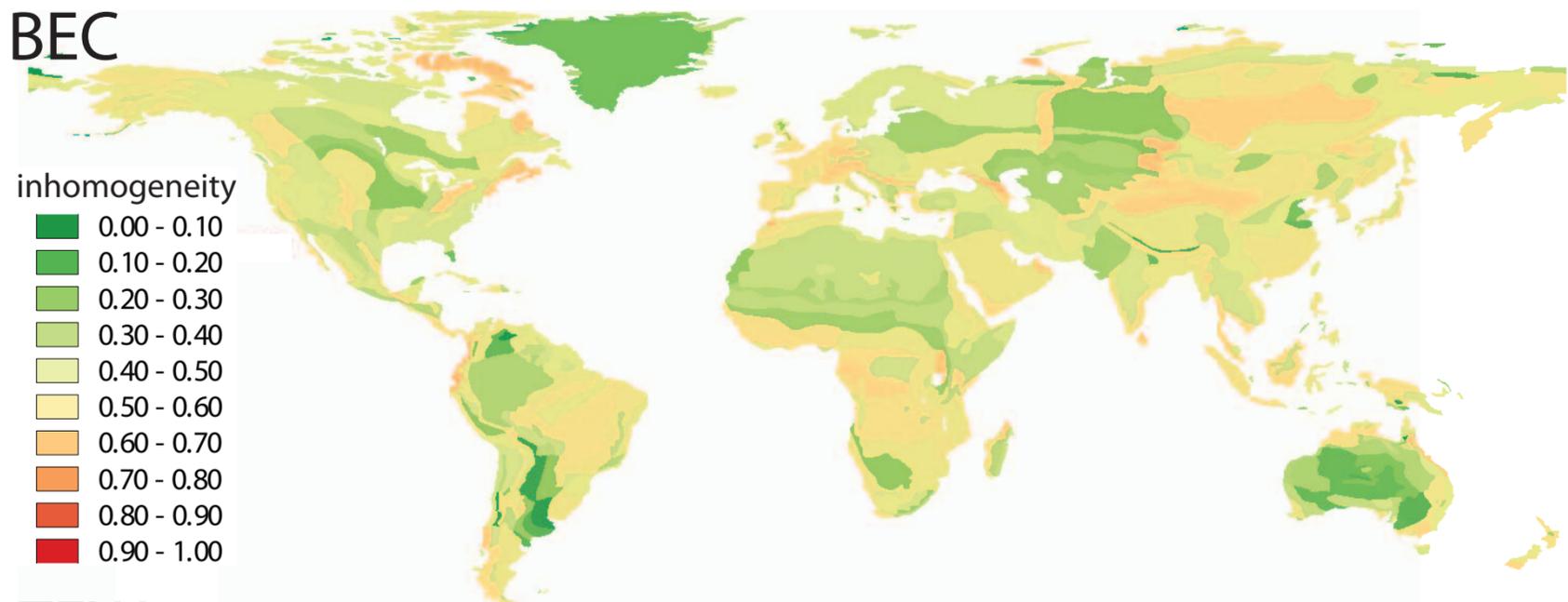
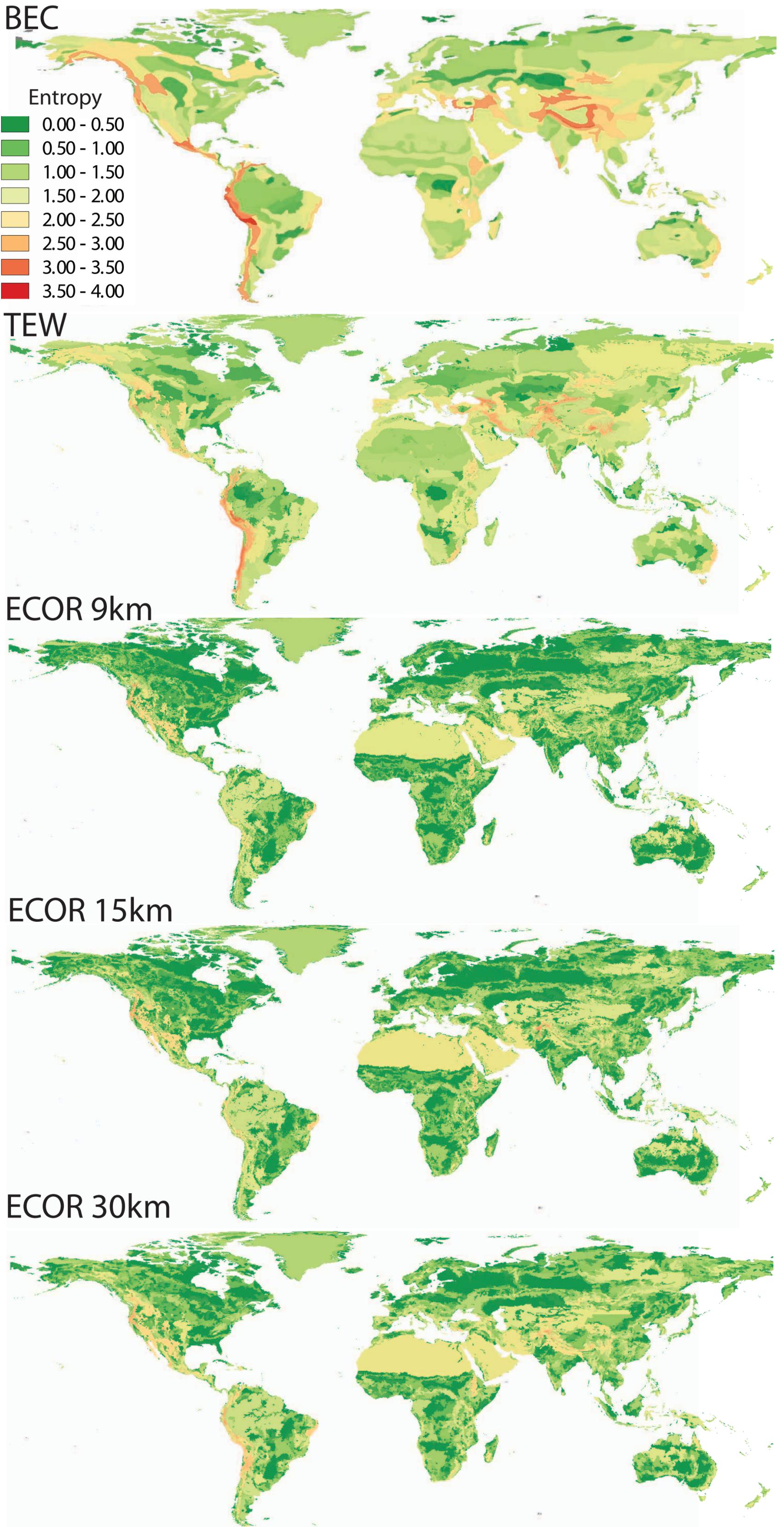


Figure S4: Maps of regions' inhomogeneity values with respect to patterns of bioclimates in different ecoregionalizations.



Supplement S2: Inhomogeneity and Isolation Metrics

Towards machine ecoregionalization of Earth’s landmass using pattern segmentation method

1 Co-occurrence histograms

Recall from section 2.1 that the landmass is tessellated into sites – square blocks of cells in the variable categorical raster. For the numerical description of a pattern of variable’s categories in the site we use a histogram of category co-occurrence pattern features [Barnsley and Barr, 1996; Chang and Krumm, 1999]. A co-occurrence feature is a pair of categories assigned to two neighboring cells. Features are extracted from a site by combining co-occurrence matrices calculated for eight different displacement vectors along principal directions. For a raster with k possible categories, the result is a symmetric matrix which we reduce to a histogram with $d = (k^2 + k)/2$ bins. Fig. 1 show examples of co-occurrence histograms stemming from two different hypothetical sites. In this hypothetical case $k = 4$ resulting in a co-occurrence histograms with 10 bins. In the case of CCI-LC, $k = 22$ and the co-occurrence histogram has 253 bins. A bin in a histogram gives a (normalized; divided by the sum of all bins) number of co-occurrences (either horizontal, vertical or diagonal) between given two categories. The k bins correspond to the co-occurrence of same-category pairs and their values reflect both, the abundance of the category and its spatial arrangement. The remaining $(k^2 - k)/2$ bins correspond to co-occurrences between different-categories pairs and their values reflect a geometric configuration of the pattern.

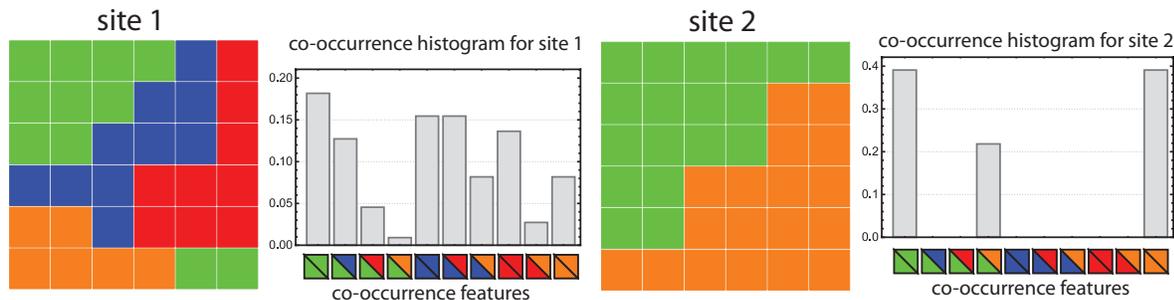


Figure 1: Co-occurrence histograms for two hypothetical sites with different patterns of variable categories. Four colors, red, blue, green, and orange indicate the four categories of the variable.

2 Dissimilarity measure

We use the Jensen-Shannon Divergence (JSD) [Lin, 1991] as a measure of dissimilarity between two sites represented by corresponding normalized co-occurrence histograms M_1 and M_2 . The JSD expresses the informational distance between the two histograms as a deviation between Shannon’s entropy of the conjugate of the two histograms $(M_1 + M_2)/2$ and the mean entropy of individual histograms M_1 and M_2 . The value of JSD, denoted by $d(M_1, M_2)$, is given by the following formula:

$$d(M_1, M_2) = H\left(\frac{M_1 + M_2}{2}\right) - \frac{1}{2}[H(M_1) + H(M_2)], \quad (1)$$

where $H(M)$ indicates a value of the Shannon’s entropy of the histogram M :

$$H(M) = -\sum_{i=1}^{|M|} m_i \log_2 m_i. \quad (2)$$

where m_i is the value of i th bin in the histogram M and $|M|$ is the number of bins (the same for both histograms). For normalized histograms, the JSD dissimilarity always takes values from 0 to 1 with the value of 0 indicating that two motifs are identical, and the value of 1 indicating maximum dissimilarity (none of the classes existing in one motif can be found in the other).

3 Linkage, inhomogeneity, and isolation

The segmentation algorithm not only requires calculating a value of dissimilarity between two sites (eq. 1) but also a value of dissimilarity between two segments (sets of sites), which we refer to as a *linkage*. Consider two segments, $S_1 = \{M_{1,1}, \dots, M_{1,k_1}\}$ consisting of k_1 sites and $S_2 = \{M_{2,1}, \dots, M_{2,k_2}\}$ consisting of k_2 sites. To measure a dissimilarity between these two segments we use the so-called average linkage or Unweighted Pair Group Method with Arithmetic Mean (UPGMA) [Sokal and Michener, 1958] given by

$$D(S_1, S_2) = \frac{1}{k_1 k_2} \sum_{i=1}^{k_1} \sum_{j=1}^{k_2} d(M_{1,i}, M_{2,j}) \quad (3)$$

where function $d(x, y)$ is given by eq.(1). The value of $D(S_1, S_2)$ has a range between 0 and 1 because the values of d are restricted to this range.

Let S be a focus segment and S_1, \dots, S_N be its neighbors. The isolation metrics γ is a weighted average linkage between the focus segment and its N neighbors,

$$\gamma(S) = \frac{1}{N} \sum_{i=1}^N w_i D(S, S_i) \quad (4)$$

where w_i are the weight set to a fraction of focus segment S perimeter shared with segment S_i . Isolation is a property of a single segment, its value has a range between 0 and 1 because the values of D are restricted to this range. Large values of γ indicate that a focus segment is dissimilar to its neighbors. Fig. 2 illustrates the concept of isolation.

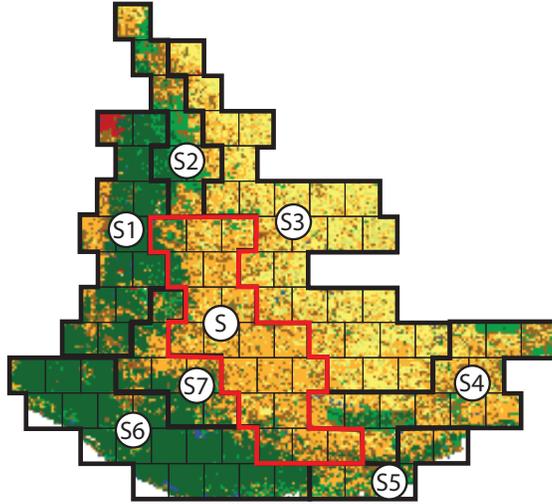


Figure 2: Focus segment S (outlined in red) has seven neighbors labeled as S_1 to S_7 and outlined in black. A linkage D is calculated between S and every neighbor. The seven values of D are averaged using weights which correspond to lengths of borders between S and the neighbors. The value of isolation (with respect to land cover) for S is $\gamma = 0.38$ whereas its inhomogeneity is 0.11.

Inhomogeneity is also a property of a single segment; it measures a degree of mutual dissimilarity between all sites within the segment. As a measure of inhomogeneity, we use an average distance between all distinct pairs of sites in a segment. For a segment $S = \{M_1, \dots, M_{k_1}\}$ with k_1 sites the inhomogeneity is given as:

$$\delta(S) = \frac{1}{k_1(k_1 - 1)} \sum_i \sum_{j \neq i} d(M_i, M_j) \quad (5)$$

as there is $k_1(k_1 - 1)$ distinct pairs of motifs in the segment S . The value of δ has a range between 0 and 1 because values of d are restricted to this range. The small value of δ indicates that all sites in the segment represent consistent patterns so the segment is pattern-homogeneous. Note that segment is considered homogeneous even if its constituent sites represent complex patterns of categories as long as the pattern of this complexity is approximately the same among all sites within a segment. Segment S in Fig. 2 has 19 sites. To calculate $\delta(S)$ we first calculate $19 \times 18 = 324$ values of dissimilarity (eqn. 1) (between every pair of sites in S) and then calculate an unweighted average.

References

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Supplement S3: Legends to categories of physiographic variables

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This supplement contains legends to the four physiographic variables we use in the paper. The colors are as they appear in the rasters of these variables we make available for download from <http://sil.uc.edu>. The value is the number in the raster that corresponds to a given category. It is also a number utilized for auto-generation of 16-digits codes for each region; use these legends to decipher a code. The label is the name of a category.

| Color | Value | Label |
|---|-------|--|
|  | 1 | cropland rainfed |
|  | 2 | cropland irrigated |
|  | 3 | mosaic cropland / natural vegetation |
|  | 4 | mosaic natural vegetation / cropland |
|  | 5 | tree cover broadleaved evergreen |
|  | 6 | tree cover broadleaved deciduous |
|  | 7 | tree cover needleleaved evergreen |
|  | 8 | tree cover needleleaved deciduous |
|  | 9 | tree cover mixed |
|  | 10 | mosaic tree and shrub / herbaceous cover |
|  | 11 | mosaic herbaceous cover / tree and shrub |
|  | 12 | shrubland |
|  | 13 | grassland |
|  | 14 | lichens and mosses |
|  | 15 | sparse vegetation |
|  | 16 | tree cover flooded fresh water |
|  | 17 | tree cover flooded saline water |
|  | 18 | shrub or herbaceous cover flooded water |
|  | 19 | urban areas |
|  | 20 | bare areas |
|  | 21 | water bodies |
|  | 22 | permanent snow and ice |

Figure 1: Legend for 22 CCI-LC land cover categories (<http://maps.elie.ucl.ac.be/CCI/viewer/>)

| Color | Value | Label |
|---|-------|-------------|
|  | 1 | alfisols |
|  | 2 | andisols |
|  | 3 | aridisols |
|  | 4 | entisols |
|  | 5 | gelisols |
|  | 6 | histosols |
|  | 7 | inceptisols |
|  | 8 | mollisols |
|  | 9 | oxisols |
|  | 10 | spodosols |
|  | 11 | ultisols |
|  | 12 | vertisols |

Figure 2: Legend for twelve soil orders. See <https://globalrangelands.org/topics/rangeland-ecology/twelve-soil-orders> for description of the orders.

| Color | Value | Label |
|---|-------|---------------------|
|  | 1 | very cold, wet |
|  | 2 | very cold, very wet |
|  | 3 | very cold, moist |
|  | 4 | very cold, semi-dry |
|  | 5 | arctic |
|  | 6 | cold, very wet |
|  | 7 | cold, wet |
|  | 8 | cold, moist |
|  | 9 | cold, semi-dry |
|  | 10 | cool, very wet |
|  | 11 | cool, wet |
|  | 12 | cool, moist |
|  | 13 | cool, semi-dry |
|  | 14 | warm, wet |
|  | 15 | warm, very wet |
|  | 16 | cool, dry |
|  | 17 | cold, dry |
|  | 18 | warm, dry |
|  | 19 | warm, semi-dry |
|  | 20 | warm, moist |
|  | 21 | cool, very dry |
|  | 22 | warm, very dry |
|  | 23 | hot, wet |
|  | 24 | hot, moist |
|  | 25 | very cold, dry |
|  | 26 | cold, very dry |
|  | 27 | hot, semi-dry |
|  | 28 | hot, very wet |
|  | 29 | High mountains |
|  | 30 | hot, very dry |
|  | 31 | very hot, very dry |
|  | 32 | very hot, semi-dry |
|  | 33 | very hot, wet |
|  | 34 | very hot, moist |
|  | 35 | very hot, dry |
|  | 36 | very hot, very wet |
|  | 37 | very cold, very dry |

Figure 3: Legend for 37 types of bioclimates. See Sayre et al. [2014]

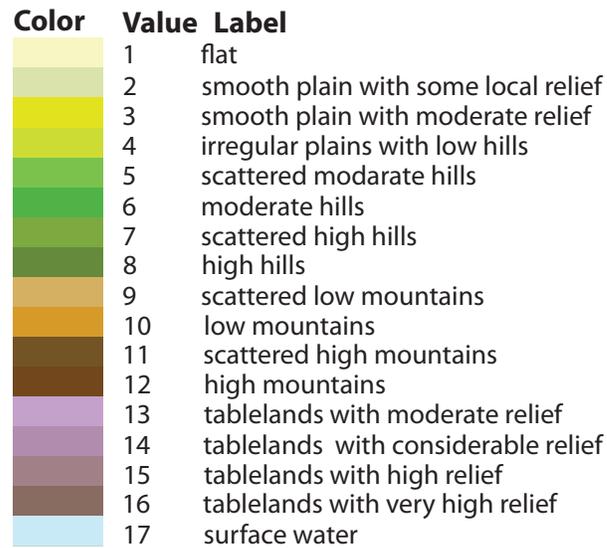


Figure 4: Legend for 17 categories of landforms. See Karagulle et al. [2017]

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