

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. Homogeneous physiography is a necessary but not necessarily sufficient condition for a region to be an ecoregion, thus machine delineation of ecophysigraphic regions is the first, important step toward global ecoregionalization. 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). At the finer 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 frame-

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48 work. TEW is a compilation of local regions taken from pre-105
49 existing, independently conducted studies. On one hand, this-106
50 may be viewed as a positive because TEW combines expert-107
51 knowledge of the broad community. On the other hand, there-108
52 are no straightforward means to inspect materials and protocols-109
53 that contributed to the creation of TEW. As there is no under-110
54 lying quantitative framework, there are no quantitative criteria-111
55 to assess the quality of TEW. Therefore, no systematic checks-112
56 modifications or objective updates to TEW are possible. More-113
57 over, although many individual regions in TEW may be well-114
58 delineated, as a whole, TEW lacks overall consistency. A user-115
59 has no means of knowing which regions are well-delineated-116
60 and which are not. TEW legend conveys a short description-117
61 of a region which usually pertains to a combination of region's-118
62 geography, climate, and flora. Because regions in TEW lack-119
63 quantitative description, the inter-regions comparison is limited-120
64 to contrasting their short descriptions in the legend. 121

65 The weight-of-evidence approach (Omernik, 1987; Omernik-122
66 and Griffith, 2014) also lacks quantitative framework, but, it is-123
67 rooted in a clear conceptual framework – “Ecoregions should-124
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 lineated (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 the-133
77 character similar to that in TEW, the inter-regions comparison-134
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' climatic-138
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 using-140
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 variables-143
88 are associated with each pixel (a tract of land corresponding to-144
89 the resolution of the data) and pixels agglomerated into larger-145
90 zones (ecoregions) on the basis of the Euclidean distance be-146
91 tween these vectors. Such automated approach addresses issues-147
92 related to objectivity, consistency, and inter-region comparabil-148
93 ity (see our discussion above), however, its ability to yield a-149
94 useful ecoregionalization is limited by the choice of clustering-150
95 as a technique enabling the automation. Clustering leads to a-151
96 delineation of non-contiguous, highly fragmented zones, with-152
97 the fragments spread over wide areas. Clustering may be well-153
98 suited for classification but it is ill-suited for mapping. Mapping-154
99 needs to be based on characteristics which are macroscopically-155
100 recognizable (Klijn et al., 1995), which environmental variables-156
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 landmass-158
104 capable of producing a useful global map of ecophysiological-159

regions. We call the resultant regions “ecophysiological” be-
cause they are mapped based on physiography but aim at de-
lineating ecosystems as well. This is consistent with the no-
tion that ecoregionalization on larger scales should be based
on physiography (Klijn et al., 1995; Sayre et al., 2014). Fol-
lowing Omernik and Griffith (2014), our mapping is based on
macroscopically recognizable *patterns* of physiographic cate-
gorical variables, but a decision on where to put boundaries be-
tween 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 no-
tion (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 be-
hind 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 frame-
work calls for. However, we find a high level of spatial asso-
ciation 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 quantita-
tive 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, cli-
mate, 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. We also use the newest global land cover
dataset – the European Space Agency (ESA) Climate Change
Initiative (CCI) global land cover map (thereafter referred to as
CCI-LC). 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 informa-
tion about geographical distribution of biodiversity and ecolog-
ical 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 CCI-LC land cover dataset in-
cluding its accuracy can be found in the Land Cover CCI Prod-
uct User Guide V.2 (ESA, 2017).

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)

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 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 ecophysiological 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 ecoregionalizations

Ecoregions should be characterized by homogeneous patterns of physiographic variables (Klijn et al., 1995; Sayre et al., 2014; Omernik and Griffith, 2014). In addition, it is desirable that patterns of physiographic variables in adjacent regions differ from each other. We assess a degree to which these conditions are met by ECORs using statistics of regions homogeneity and isolation metrics with respect to patterns of all physiographic variables. These statistics are calculated over all ECOR’s segments. We compare ECOR-derived statistics with analogous statistics calculated over all land units in BEC, and TEW. Note that in BEC and TEW land units are individual polygons (land units) in their respective shapefiles. The term “ecoregion” in BEC and TEW does not 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.

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

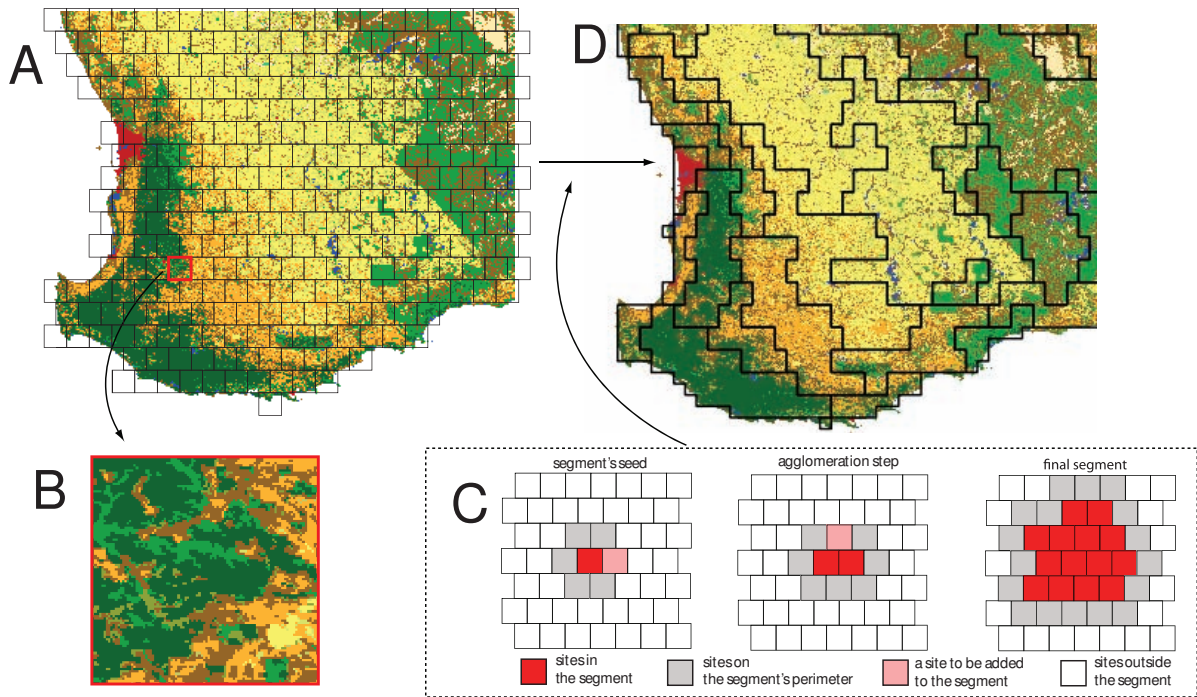


Figure 1: Basic concept of pattern-based segmentation using a fragment of landmass located in the southwestern Australia around the city of Perth. (A) A grid of sites. (B) A zoom-in onto a single $30\text{km} \times 30\text{km}$ site to show its pattern. (C) The concept of seeded region growing algorithm; 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).

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 Supplement S3). Minimum possible value of H is zero and it occurs when a segment is completely within a single climate category (it is completely homogeneous). The larger the value of H the more inhomogeneous the segment is with respect to climate.

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

3. Results

3.1. Associations between physiographic variables

We first estimate a degree of association between our four physiographic variables in order to provide a priori rationale for using land cover patterns as the only input to the segmentation

algorithm. We want to check to what degree categories of different variables co-occur on the scale of our sites. To start we regridded the four variables from their native resolutions (see Table 1) to grids with $9\text{km} \times 9\text{km}$ and $30\text{km} \times 30\text{km}$ cells using the mode values method. Because we deal with categorical variables we use Cramér's V measure of association (Cramér, 2016). Table 2 shows the values of Cramér's V for all combinations of variables.

Table 2: Degree of association between physiographic variables

	LC	BC	LF	S	Mean	St.Dev.
9km × 9km 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
30km × 30km 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.

Our results in Table 2 indicate that mutual associations between land cover, soils and climate are higher (0.3–0.5) than association of these variables with landforms (0.09 – 0.2). According to one interpretation (Corbett and LeRoy, 2003) of Cramér's V values $V < 0.2$ indicates a weak association, $V =$

0.2–0.25 indicates a moderate relationship, $V = 0.25–0.30$ indicates a moderately strong association, and $V > 0.3$ indicates a strong association. Using this interpretation, values in Table 2 indicate three physiographic variables, land cover, soils, and bioclimate to be strongly mutually associated. The landforms variable is only weakly associated with the remaining three variables, but most associated with the land cover. 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 $30\text{km} \times 30\text{km}$ sites, $15\text{km} \times 15\text{km}$ sites, and $9\text{km} \times 9\text{km}$ 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 $1.2 \times 10^7 \text{ km}^2$. 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,

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.

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

The semantic meaning of the code can be deciphered from the legends of the four variables (see Supplement S3). For example, 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 $(1 - \text{inhomogeneity/isolation})$. 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

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.

374 has, on average, better-defined regions with respect to a given 405
375 variable. In general, ECORs regions are more homogeneous 406
376 but less isolated than TEW and BEC. For the best overall char- 407
377 acterization of regionalization, the inhomogeneity and isolation 408
378 metrics need to be considered together; this is achieved by the 409
379 quality metric. According to the unweighted method, ECORs 410
380 are characterized by smaller values of quality than TEW but 411
381 by higher values of quality than BEC. According to the area- 412
382 weighted method, ECORs are characterized by higher values 413
383 of quality than both TEW and BEC. 414

384 For landforms, land cover, and soils, the numbers in Table 4 415
385 could also be compared within a row (for a given regionaliza- 416
386 tion) to indicate, on average, a quality of a region delineation 417
387 with respect to patterns of different physiographic variables. 418
388 As expected, ECORs regions are best delineated with respect 419
389 to the land cover. The value of 0.57 (unweighted quality for 420
390 land cover in ECOR 30) can be interpreted as follows: in an 421
391 average region, the similarity of its constituent sites with re- 422
392 spect to patterns of land cover is 2.3 times higher than an av- 423
393 erage similarity of land cover patterns between this region and 424
394 its neighbors. Following this interpretation for patterns of soils 425
395 and landforms yields the ratios of 2 and 1.67, respectively. This 426
396 result is consistent with our expectations based on associations 427
397 between physiographic variables (section 3.1). 428

398 Homogeneity of regions with respect to bioclimate requires 429
399 a separate discussion because it is measured by the entropy. To 430
400 get some intuition to the meaning of entropy values we give 431
401 few examples. In the region where 90% of the area has climate 432
402 A and 10% of the area has climate B the value of entropy is 433
403 0.47. If the region is divided equally between two climates the 434
404 entropy value is 1. Small regions are covered by a single cli- 435

mate and have entropy values equal to 0. All regionalizations, except the BEC, are, on average, climate-homogeneous. Average values of isolation with respect to bioclimate must be small because most regions are small and are surrounded by regions with the same climate type.

Based on results in Table 4 we conclude that our method yields a very good regionalization of land cover patterns (quality = 0.55/0.69 using unweighted/area-weighted method for ECOR 9). It also yields a reasonable regionalization of the entire physiography with the average quality (calculated from land cover, soils, and landforms) equal to 0.5/0.48 (using unweighted/area-weighted method for ECOR 9). For comparison, the average quality for TEW is 0.61/0.32, and the average quality for BEC is 0.27/0.29. Note a significant difference between the unweighted and area-weighted values of quality for TEW. This is explained by the fact that distribution of region areas in TEW is heavily skewed toward very small regions. In TEW a small number of large regions occupy almost the entire landmass, and a large number of small regions occupy a small fraction of the landmass.

In addition, we have produced maps showing geographical distributions of inhomogeneity, isolation, and quality metrics (see Supplement S1). Locations with high values on the maps of inhomogeneity identify regions where a pattern of a given variable is under-segmented. In ECOR there are no such regions on the maps for land cover, soils, and climate (as measured by entropy), but there are few regions which are under-segmented on the map of landforms. Inhomogeneity maps for TEW and BEC have more under-segmented regions. Under-segmentation is a significant issue because it indicates that physiography varies across a region putting its status as an

ecoregion in doubt. Locations with high values on the maps of isolation identify regions where a pattern of a given variable is over-segmented. Over-segmentation is a problem because it indicates that neighboring regions have similar physiography and a single ecoregion may extent over several segments. ECOR maps are generally over-segmented to a higher degree than TEW and BEC maps. In algorithmic regionalizations there is always a trade-off between minimizing inhomogeneity of segments and maximizing isolation between different segments. This trade-off is set by maximizing the quality metric. Locations with high values on the maps of quality identify regions with relatively low inhomogeneity and relatively high isolation. These are the location where delineation of regions is the most successful. Comparing quality maps in Supplement 1 indicates that ECOR is overall a more successful ecoregionalization than TEW or BEC when using physiography as the criterion for the comparison.

4. Discussion

ECOR is the first attempt to obtain a global map of ecophysiological regions purely by means of an autonomous pattern-based segmentation algorithm. Pixel-based segmentation was previously used by Bisquet 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 variables. 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 segmentation) 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 biotic composition (Kerr et al., 2001; Pearson et al., 2004; Luoto et al., 2007; Coops et al., 2009) because it provides habitat resources for species. For these reasons, land cover is often used to provide the first-order information about geographical distribution 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 association 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 ecoregionalization is our criterion that the regions should, at the minimum, contain cohesive patterns of all physiographic variables – a quality quantitatively measured by the inhomogeneity metric. The analysis presented in section 3.3 shows that although ECOR does not yet fully meet patterns cohesiveness criterion, it meets it to the sufficient degree to be considered a viable ecoregionalization. The argument for that follows from the fact that ECOR meets patterns cohesiveness criterion to a higher degree than BEC and TEW (see Table 4 and Supplement S1), the two

regionalizations of landmass generally accepted as ecoregionalizations.

The higher cohesiveness of patterns in ECOR follows mostly from its design and from the existence of the spatial association between categories of physiographic variables. Isolation of ECOR regions is on average smaller than for regions in BEC and TEW. The overall quality of ECOR regionalization is much higher than the quality of BEC regionalization, and comparable or higher (depending on the type of measurement) to the quality of TEW regionalization.

Fig. 2 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. 2 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 any specific information available only through on the ground inspection.

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

Fig. 3 compares ECOR with the EPA Level III Ecoregions of the U.S. (Omernik, 1987; Omernik and Griffith, 2014) using the state of New Mexico as an example. Both, ECOR and EPA rely on patterns of environment for their delineation, except that ECOR delineation is algorithmic and EPA delineation is manual. Because both regionalizations follow the same underlying concept we expect a higher level of correspondence between ECOR and EPA than between ECOR and TEW.

Indeed, a clear correspondence between the two regionalizations is observed in Fig. 3A. Each EPA ecoregion is dominated by an ECOR region. The Chihuahuan Desert is dominated by a region characterized as (shrub; aridisols/mollisols; scat. 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

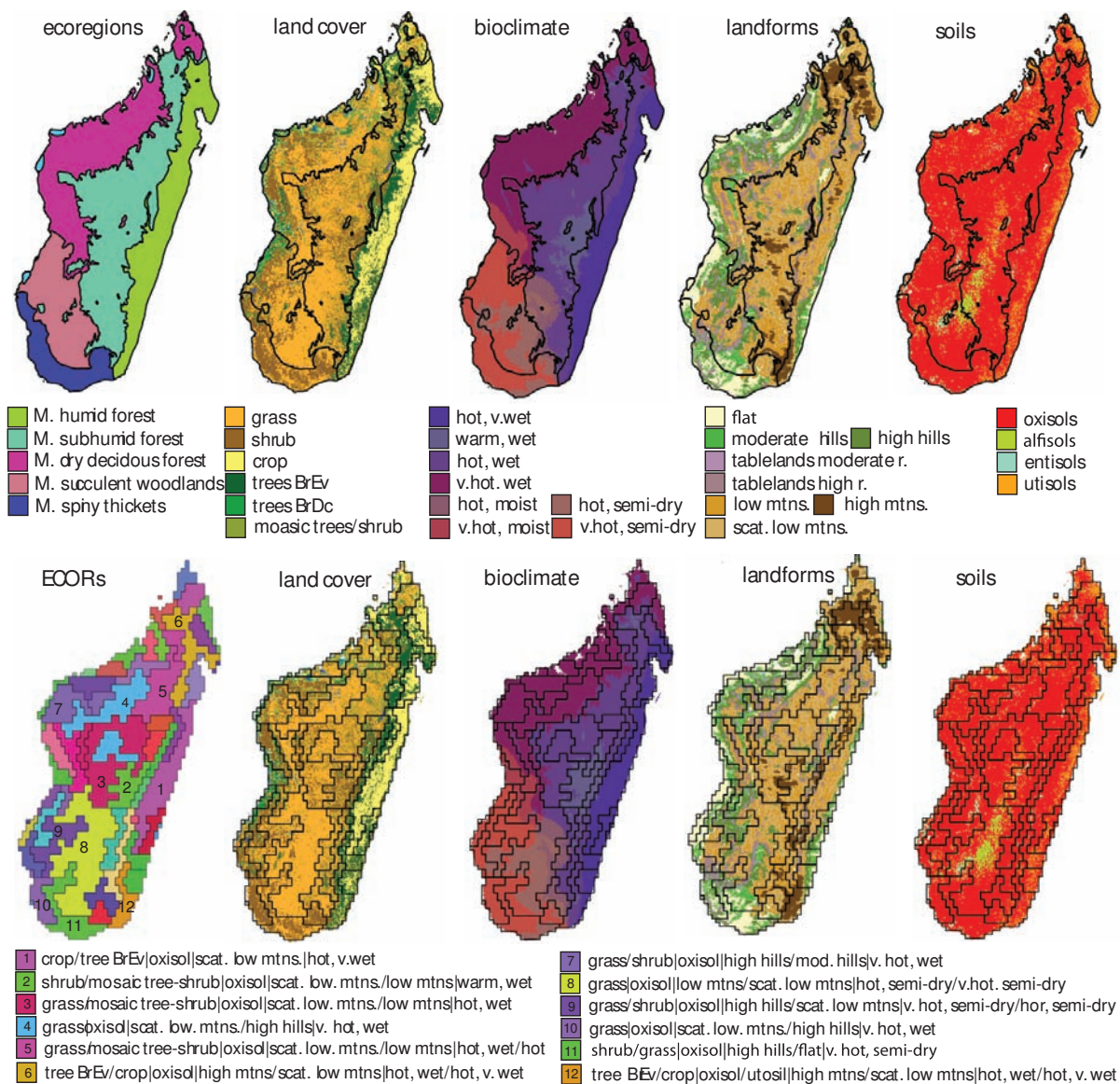


Figure 2: Comparison of ecoregionalizations in TEW and ECOR 30km using the island of Madagascar as an example. The upper row of maps shows TEW regions and how they divide the island's physiography. The lower row of maps shows the same for ECOR. Abbreviations: M. – Madagascar, v. – very, r. – relief, scat. – scattered, BrEv – broadleaf evergreen, mtns. – mountains.

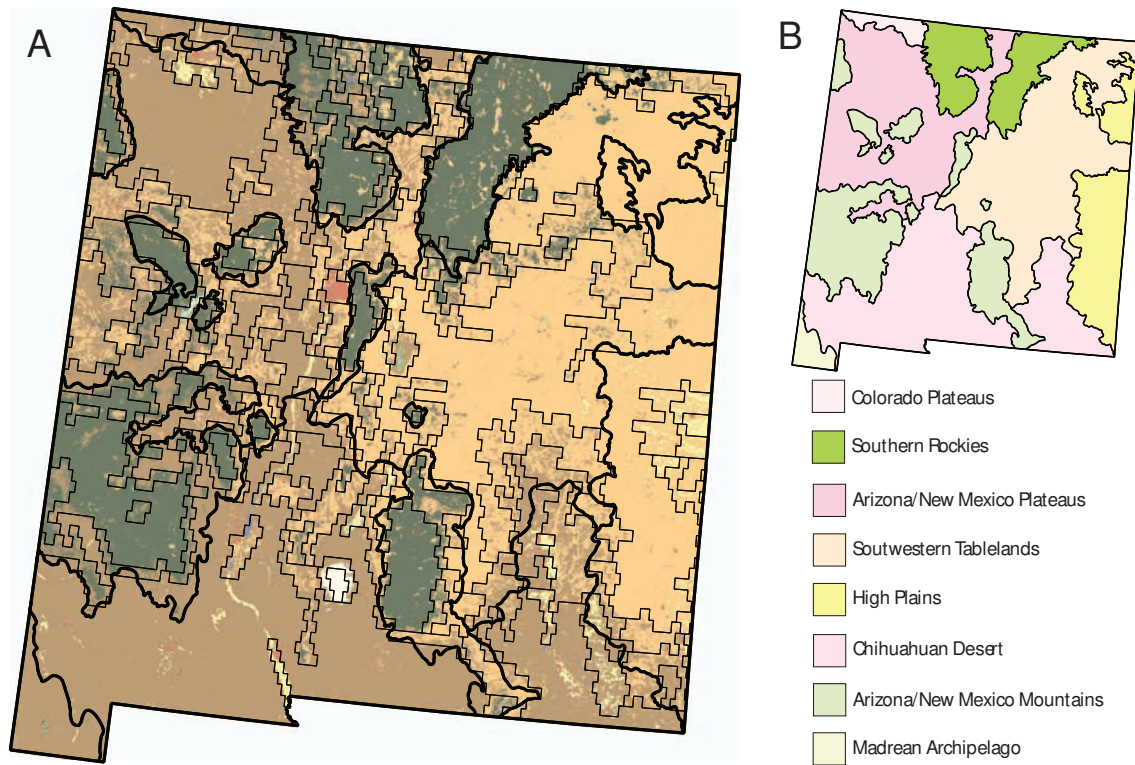


Figure 3: Comparison of ECOR 9km and EPA Level III ecoregionalizations of the state of New Mexico, U.S. (A) EPA ecoregions (thick lines) and ECOR regions (thin lines) overlying the map of land cover. (B) Eight EPA Level III ecoregions in New Mexico.

547 mtns./scat. low mtns.; cool, semi-dry/cold, moist. The two re-573
 548 gions, Southwestern Tablelands and High Plains are dominated574
 549 by the same ECOR region (grass; mollisols/aridisols; moderate575
 550 hills/flat; warm, semi-dry/cool, semi-dry). They differ by pre-576
 551 dominant landforms which the present version of segmentation577
 552 was not able to take into account. 578

553 In addition, ECOR also delineated smaller regions, where579
 554 pattern of land cover departs from surroundings. For example,580
 555 in the Chihuahuan Desert ecoregion, there are several inclu-581
 556 sions, one is the large field of white sand dunes, and another the582
 557 San Andreas mountains just west of the dunes. ECOR delin-583
 558 eated these features as independent regions, whereas they ap-584
 559 pear only at the higher, IV Level of the EPA mapping. 585

560 5. Conclusions

561 A possibility of delineating ecoregions using quantitative589
 562 methodology was discussed (McMahon et al., 2001; Loveland590
 563 and Merchant, 2004) and attempted by Hargrove and Hoffman591
 564 (2005) using multivariate clustering. However, the quantitative592
 565 method presented in this paper is the first to achieve some level593
 566 of success. This is because, instead of relying on clustering, it594
 567 employs a method that attempts to emulate in computer code595
 568 the qualitative, weight-of-evidence approach. The presented596
 569 global delineation of ecophysiological regions (ECOR) is the597
 570 first iteration of this new method. Although, we presented a de-598
 571 lineation based on a specific land cover dataset (CCI-LC), using599
 572 different dataset of comparable resolution would yield a very

similar result due to the fact that all land cover datasets must reflect the same on-the-ground reality. Indeed, we repeated calculations using the 1 km resolution GLC 2000 dataset and obtained very similar regionalization.

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

ECOR will get an update when the pattern-based segmentation technology achieves a multi-layer capability. The challenge of segmenting on the basis of multiple patterns simultaneously is how to incorporate similarities between patterns of individual variables into a similarity of the common, physiographic patterns. We expect that such update will result in improvement of regions’ physiographic homogeneity, but at the cost of an even larger number of regions.

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600 sity of Cincinnati Space Exploration Institute.

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Supplement S1: Global Maps of Regionalization Metrics

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

1 Description

This supplement contains figures each containing a set of maps showing spatial distributions of regions inhomogeneity, isolation, and an overall quality ($1 - \text{inhomogeneity/isolation}$) of regionalization with respect to a given physiographic variable. As there are four variables (land cover, soils, landforms, and bioclimate) and three regionalizations (ECOR, TEW, and BEC), there are twelve figures. Nine of these figures have three panels (inhomogeneity, isolation, and quality), the remaining three figures, corresponding to the bioclimatic variable, have only two panels. This is because, unlike in the case of the remaining three variables, we measure inhomogeneity of regions with respect to bioclimate in terms of entropy so inhomogeneity and isolation do not have the same units, and the quality metric is not defined.

The values of inhomogeneity, isolation, and quality vary from 0 to 1, except for the biodiversity variable where the value of inhomogeneity varies from 0 to $\log_2 37$. For inhomogeneity the smaller values are more desirable, but for isolation and quality, the larger values are more desirable. Legends in the figures are arranged so the gradation from a green color to a red color indicates a decrease in desirability.

For ECOR and TEW metrics are calculated using $9\text{km} \times 9\text{km}$ sites, and for BEC using $30\text{km} \times 30\text{km}$ sites (see section 2.2 of the paper for the explanation).

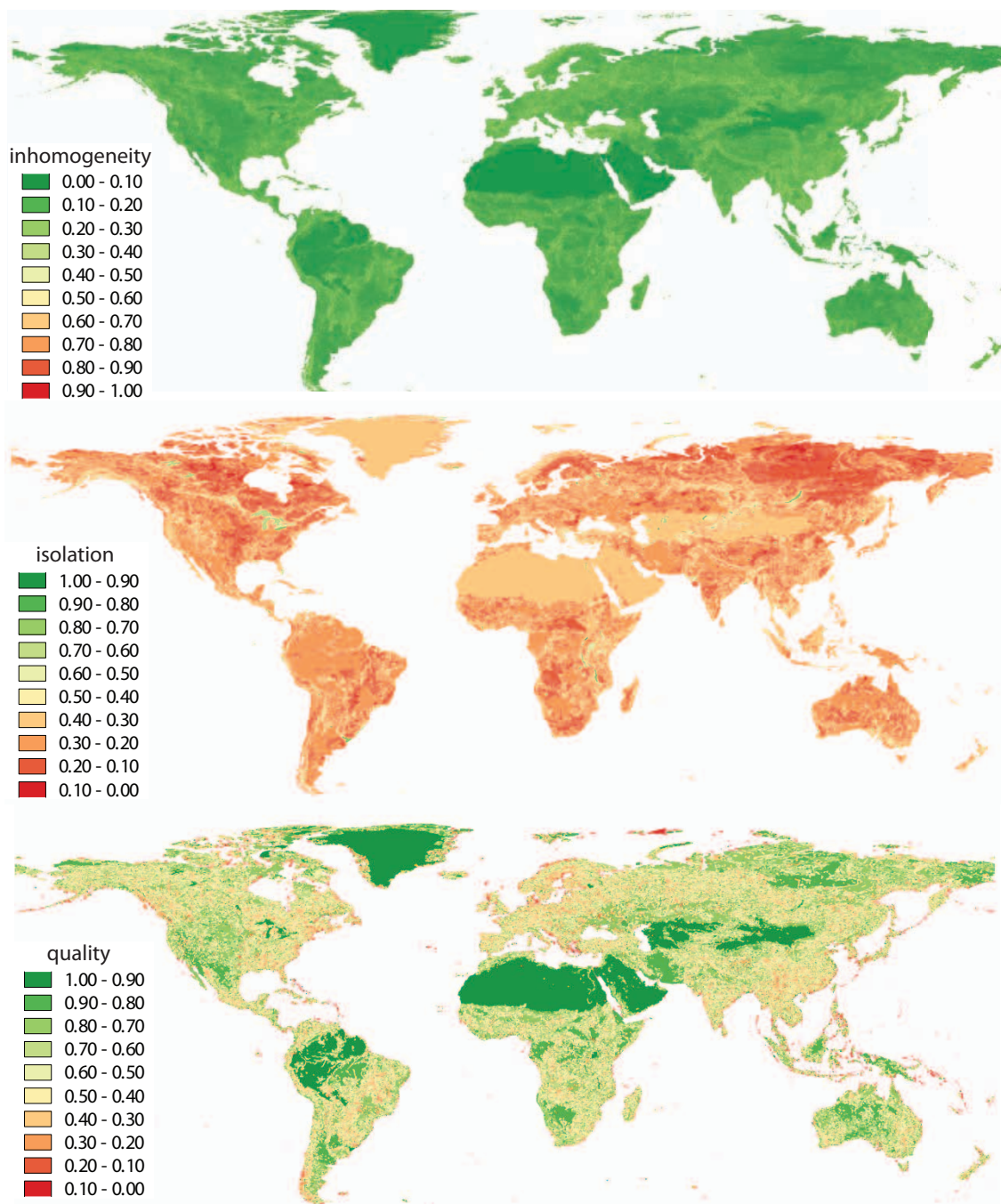


Figure 1: Maps of quality metrics, inhomogeneity (top), isolation (middle), and quality (bottom) with respect to **land cover** for the **ECOR** regionalization.

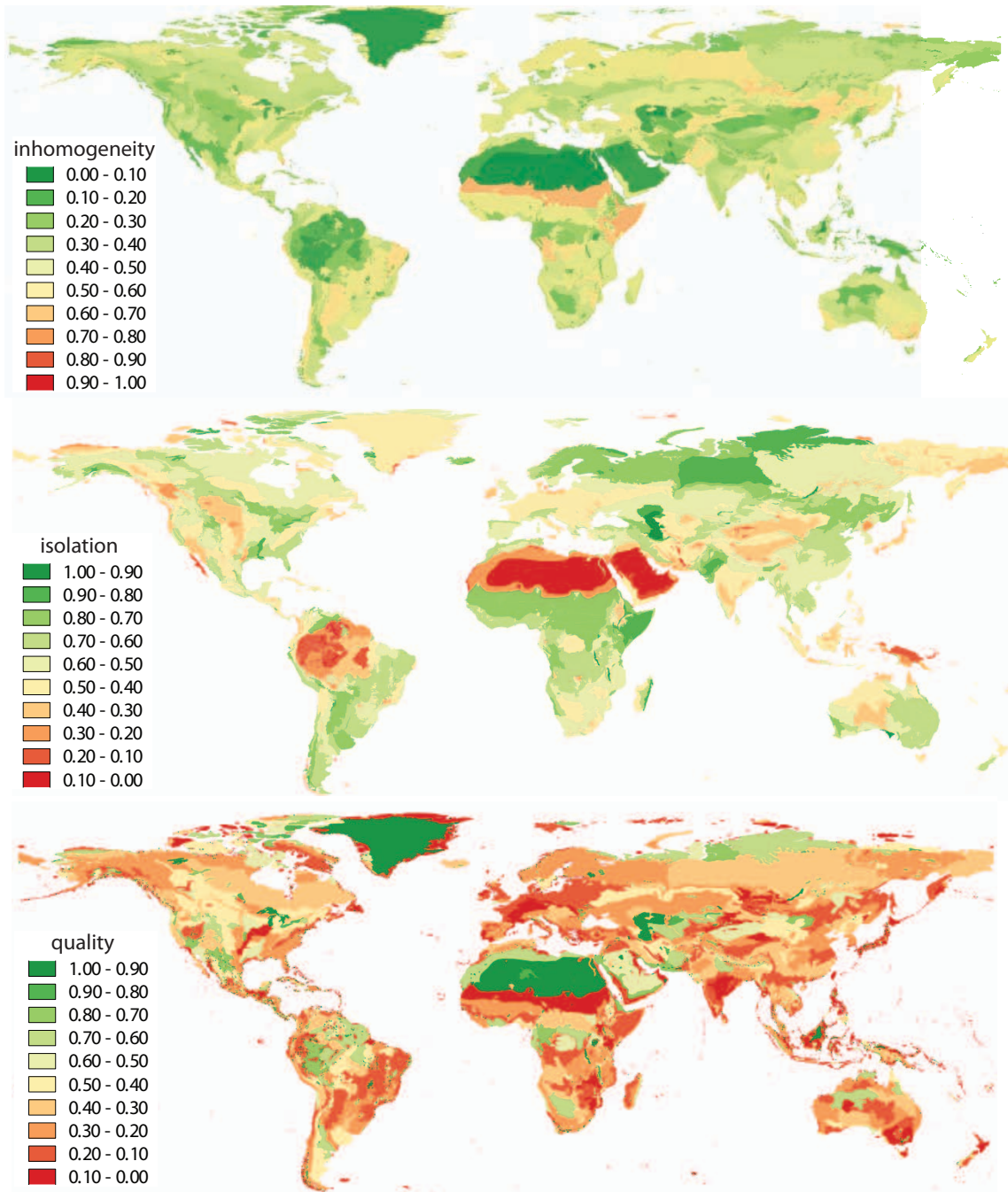


Figure 2: Maps of quality metrics, inhomogeneity (top), isolation (middle), and quality (bottom) with respect to **land cover** for the **TEW** regionalization.

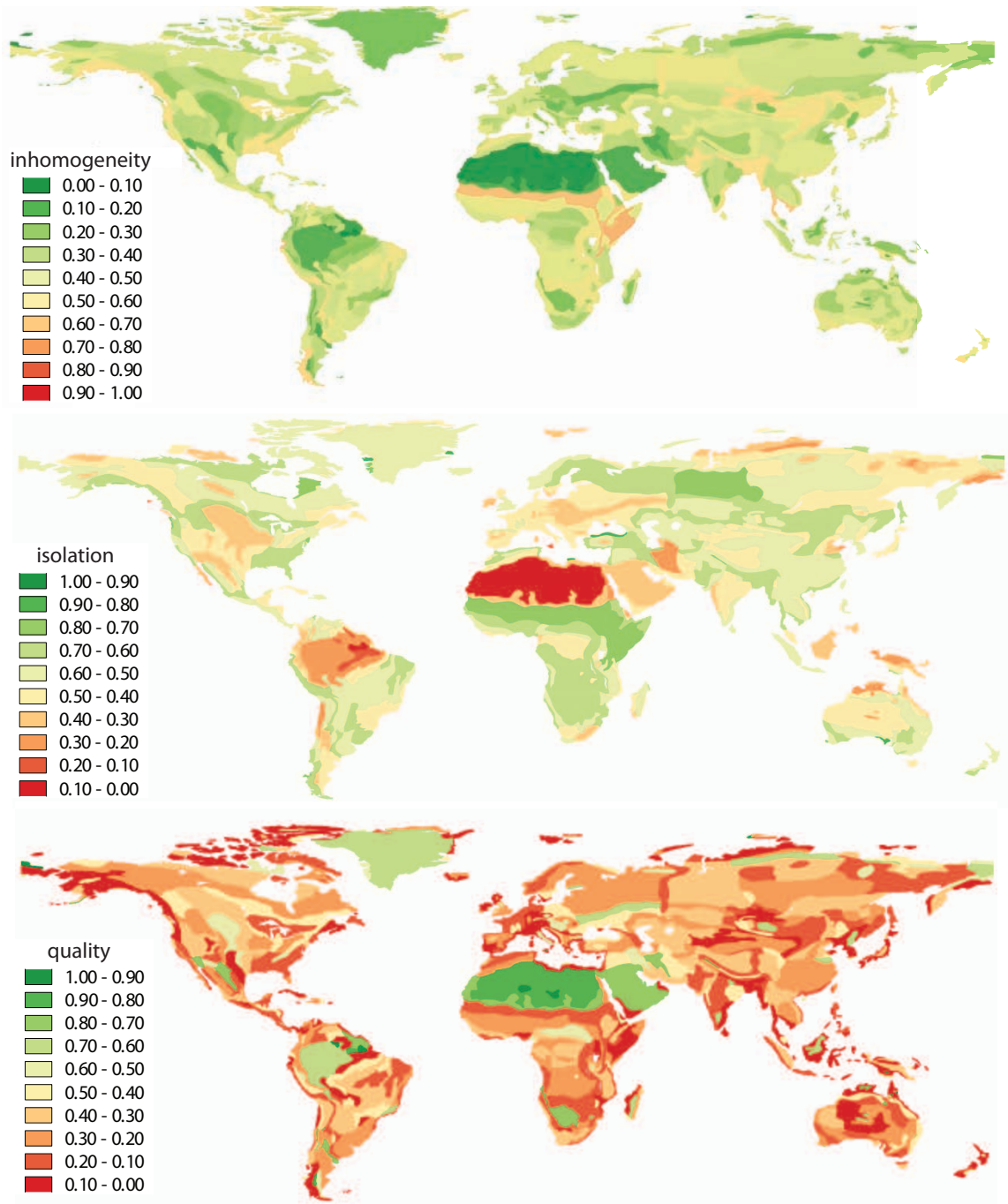


Figure 3: Maps of quality metrics, inhomogeneity (top), isolation (middle), and quality (bottom) with respect to **land cover** for the **BEC** regionalization.

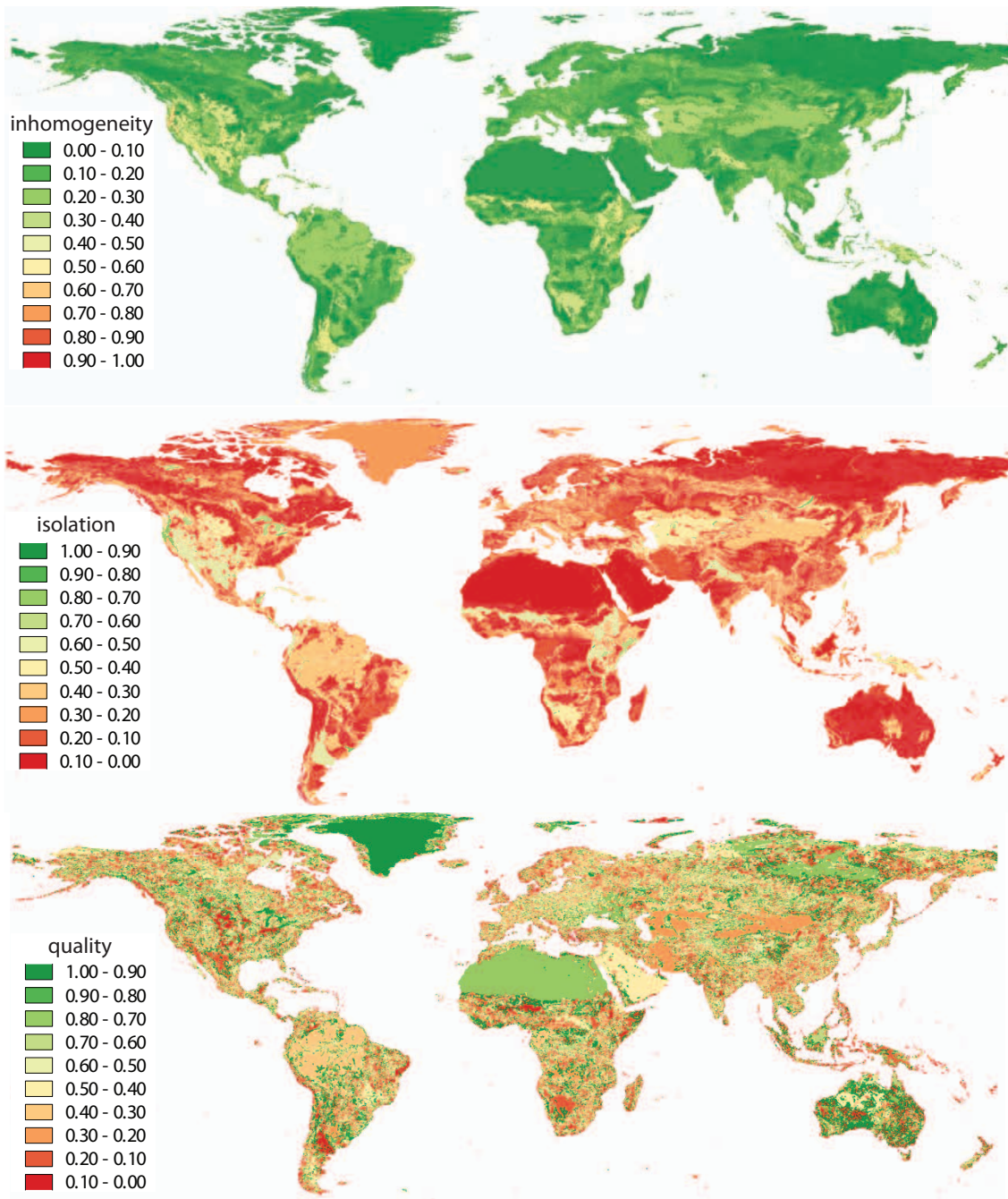


Figure 4: Maps of quality metrics, inhomogeneity (top), isolation (middle), and quality (bottom) with respect to **soils** for the **ECOR** regionalization.

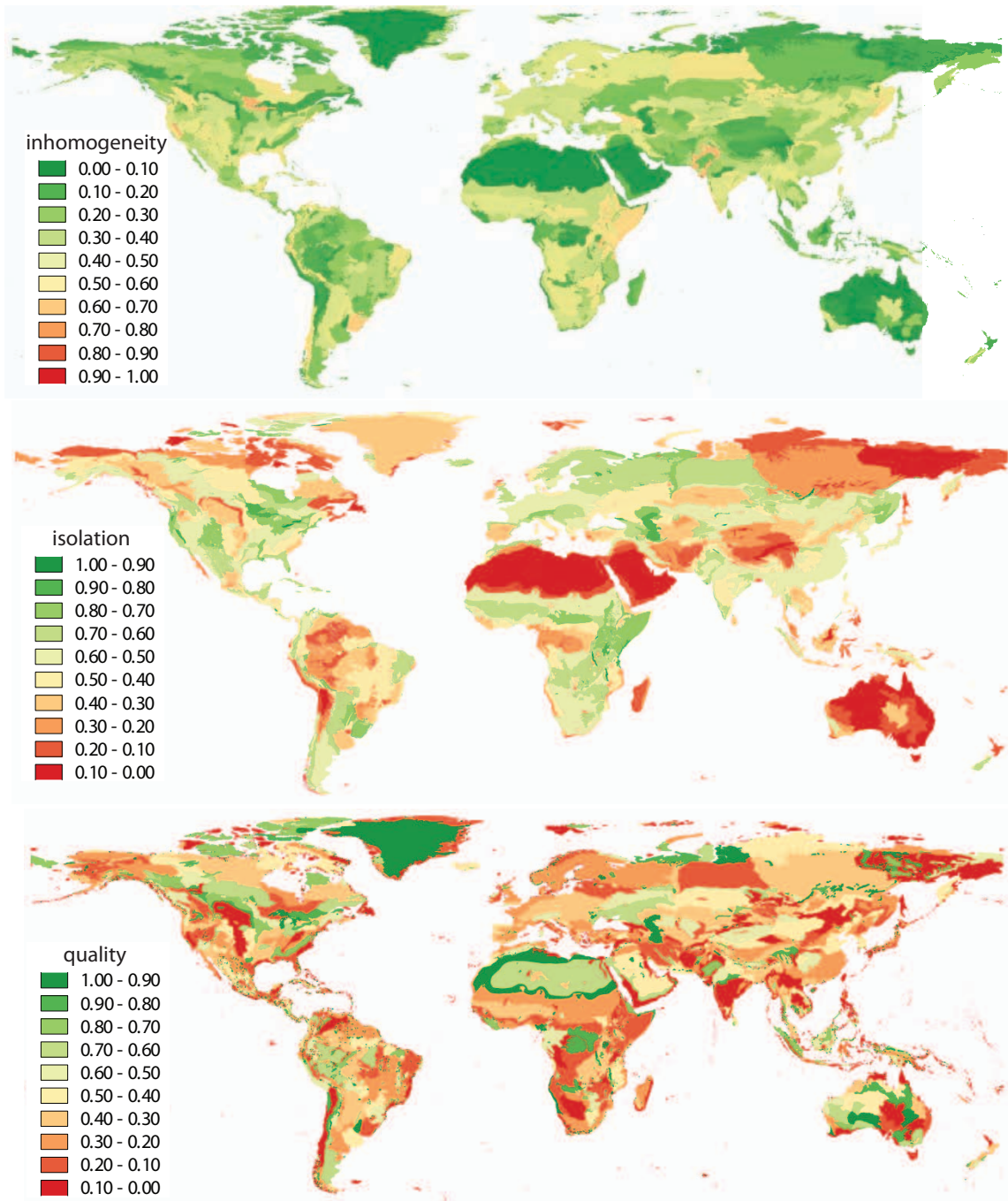


Figure 5: Maps of quality metrics, inhomogeneity (top), isolation (middle), and quality (bottom) with respect to **soils** for the **TEW** regionalization.

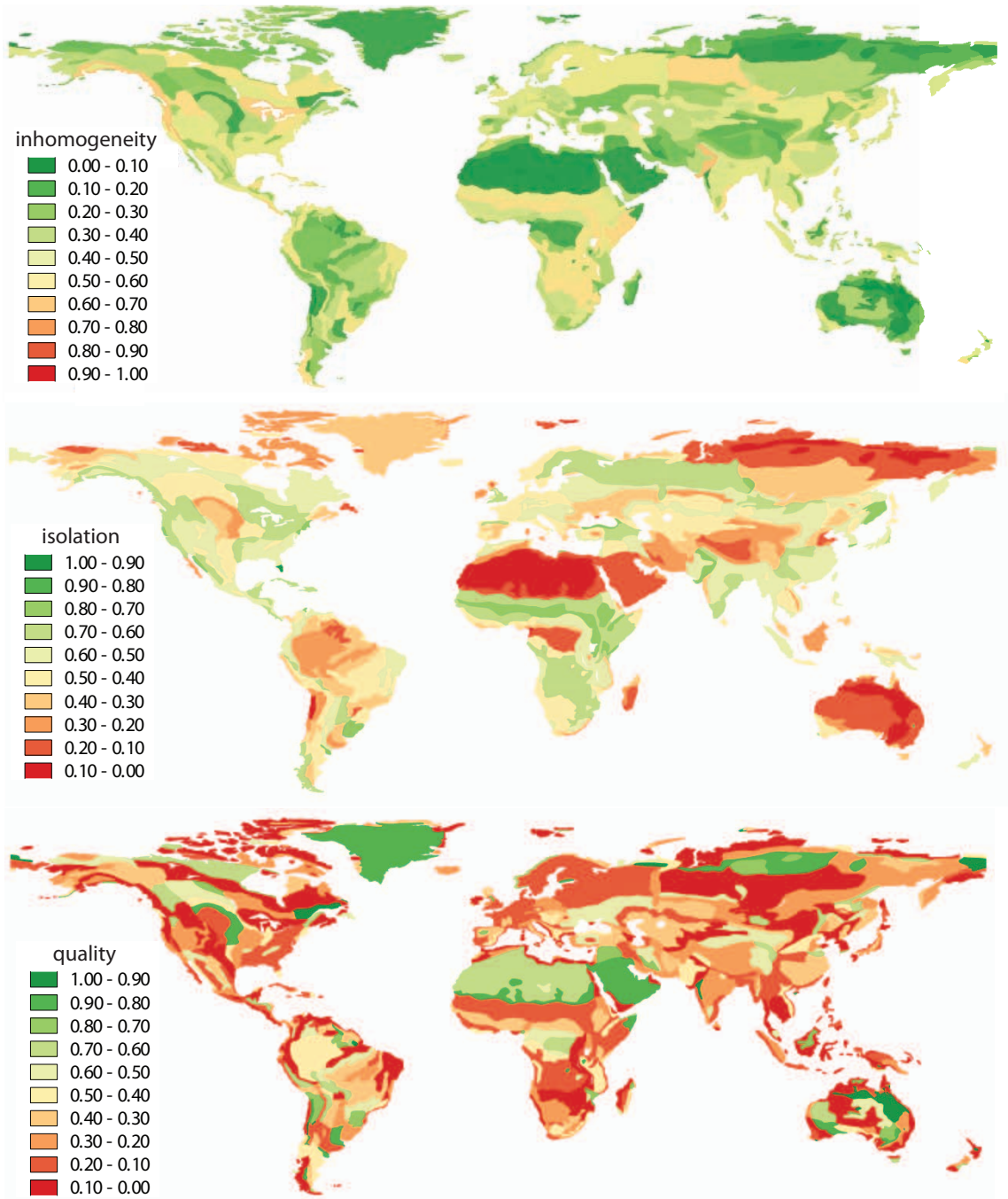


Figure 6: Maps of quality metrics, inhomogeneity (top), isolation (middle), and quality (bottom) with respect to **soils** for the **BEC** regionalization.

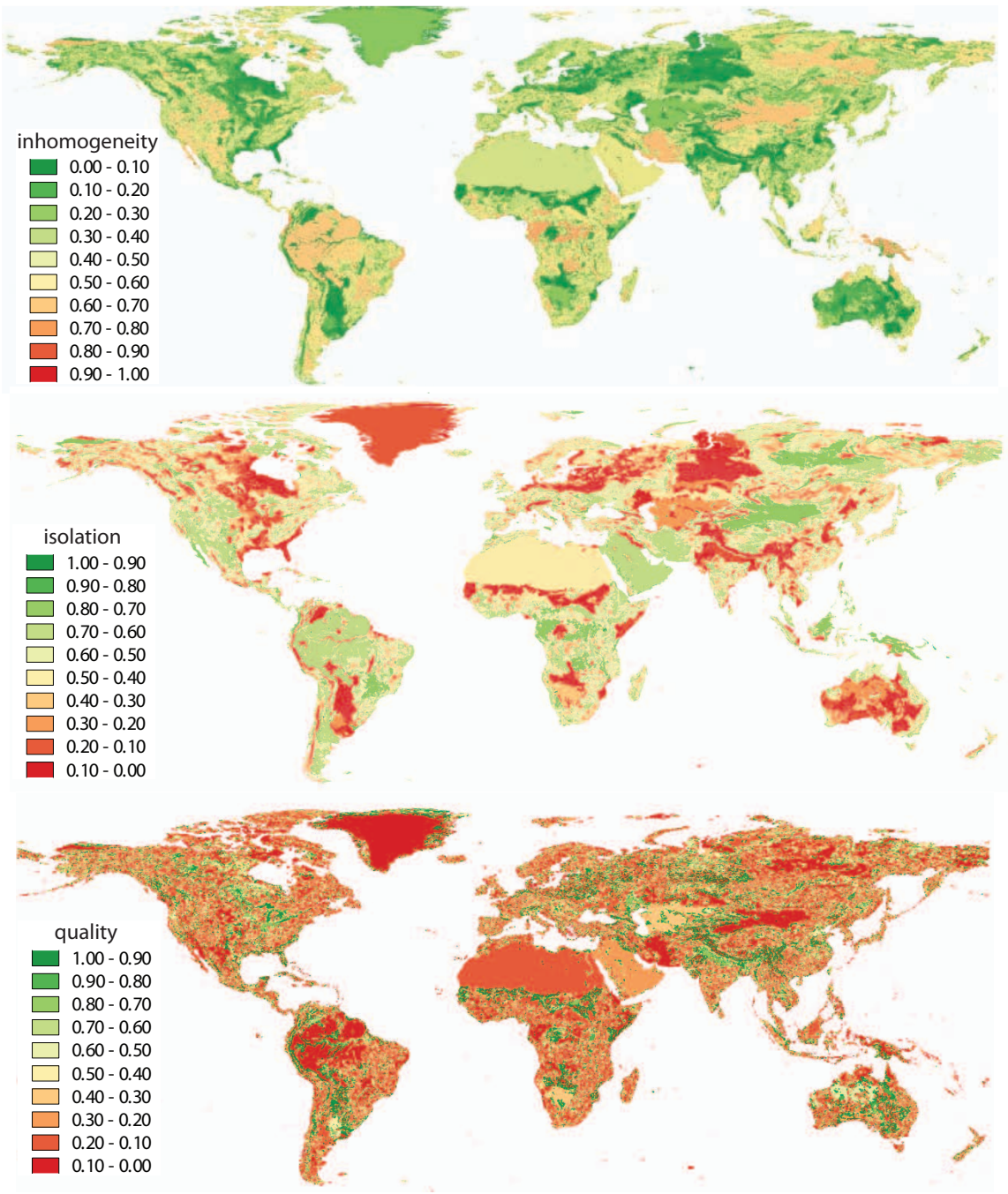


Figure 7: Maps of quality metrics, inhomogeneity (top), isolation (middle), and quality (bottom) with respect to **landforms** for the **ECOR** regionalization.

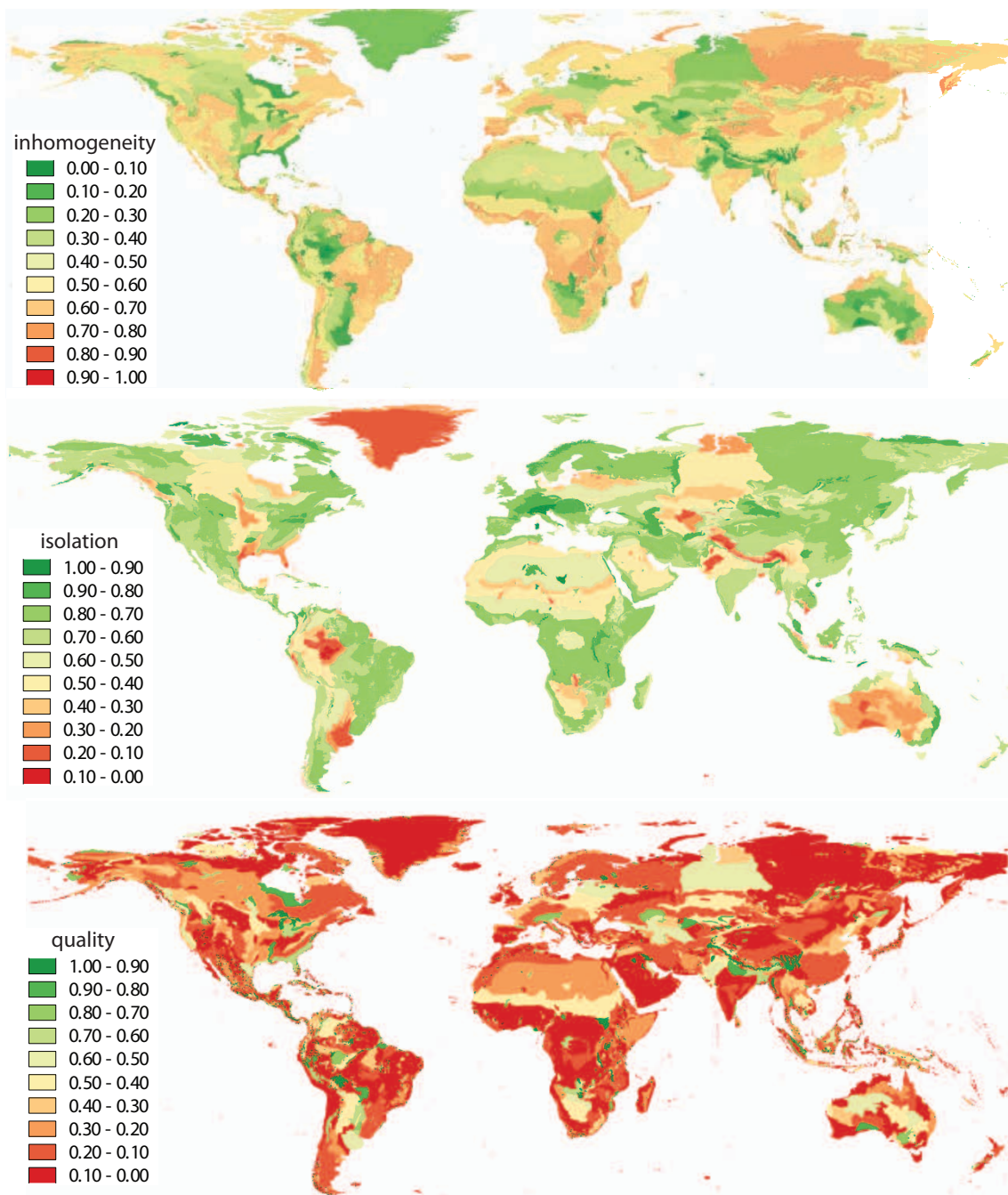


Figure 8: Maps of quality metrics, inhomogeneity (top), isolation (middle), and quality (bottom) with respect to **landforms** for the **TEW** regionalization.

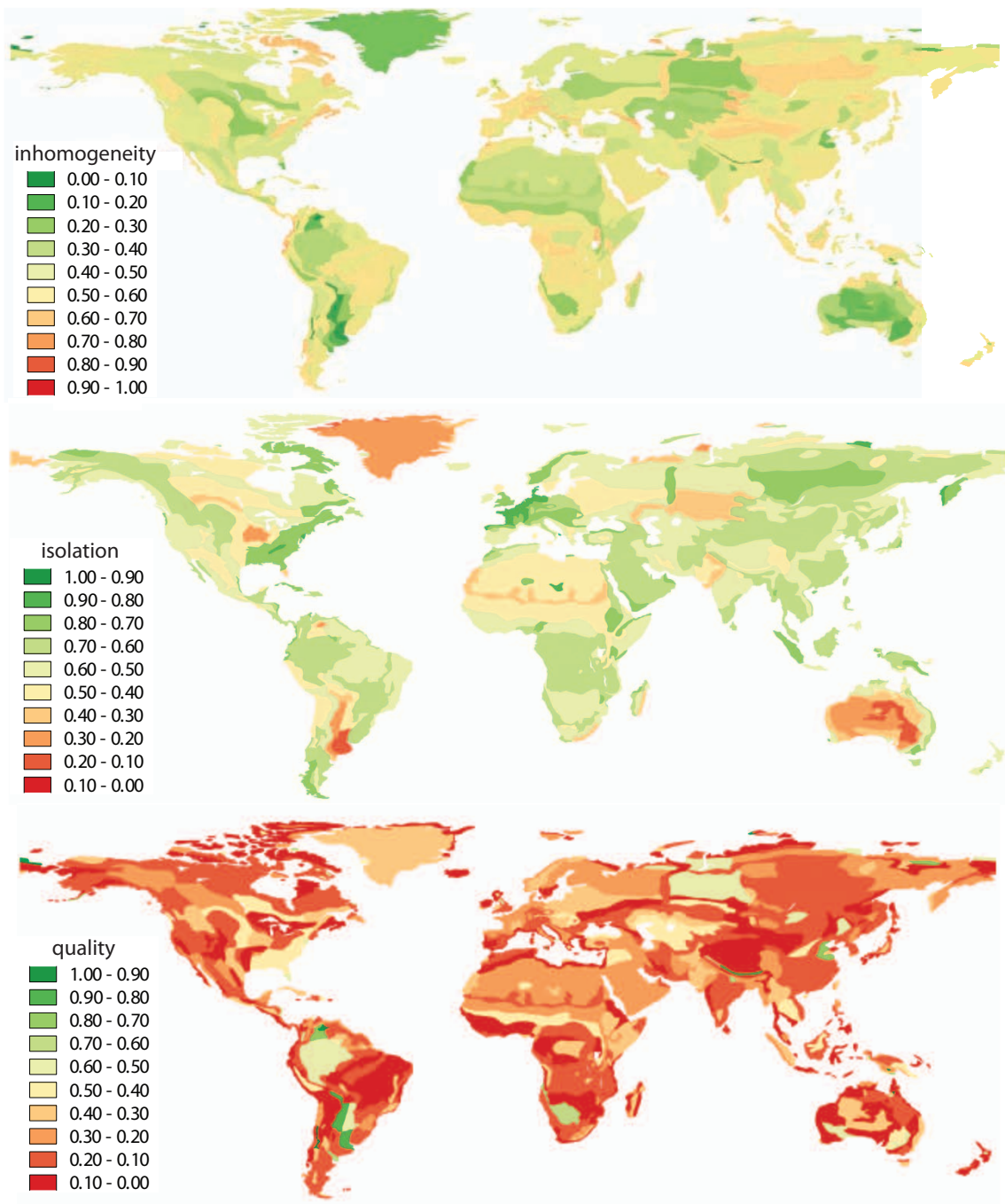


Figure 9: Maps of quality metrics, inhomogeneity (top), isolation (middle), and quality (bottom) with respect to **landforms** for the **BEC** regionalization.

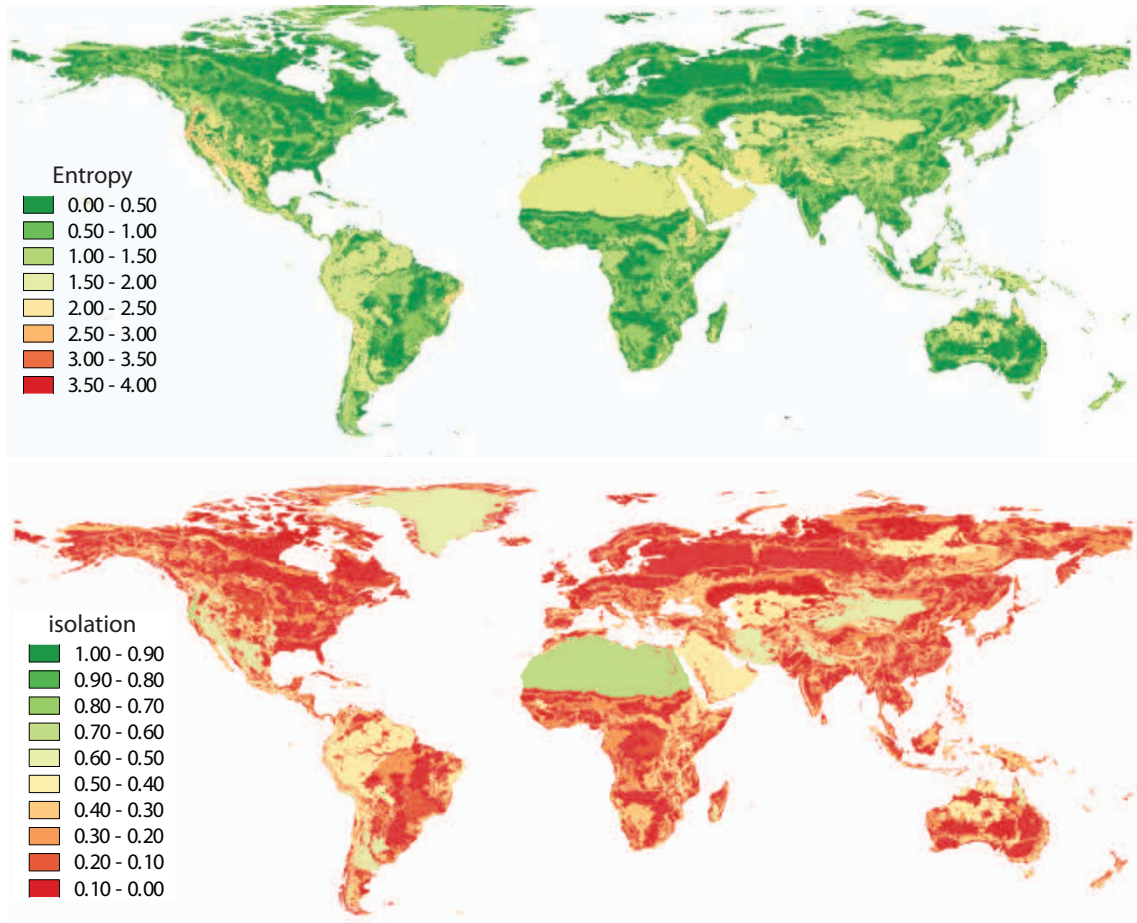


Figure 10: Maps of quality metrics, inhomogeneity (top), isolation (middle), and quality (bottom) with respect to **bioclimate** for the **ECOR** regionalization.

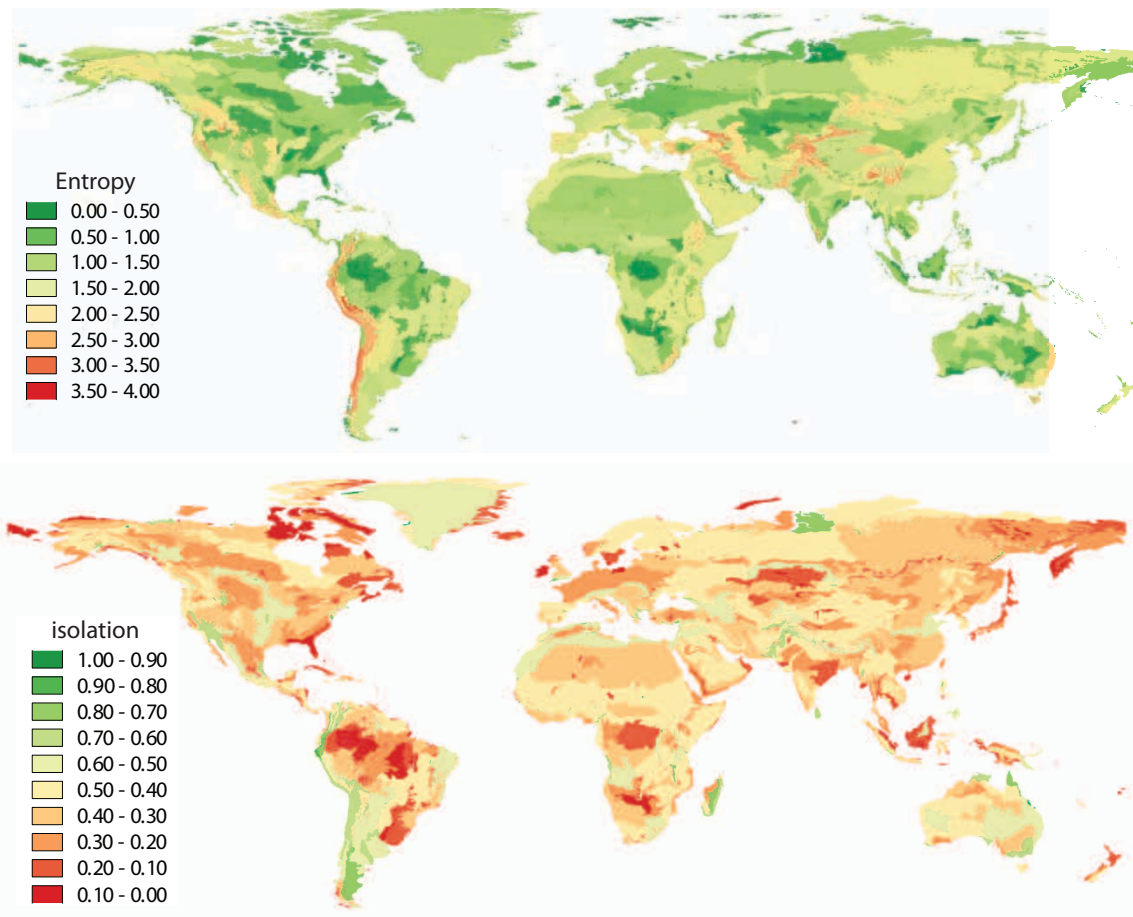


Figure 11: Maps of quality metrics, inhomogeneity (top), isolation (middle), and quality (bottom) with respect to **bioclimate** for the **TEW** regionalization.

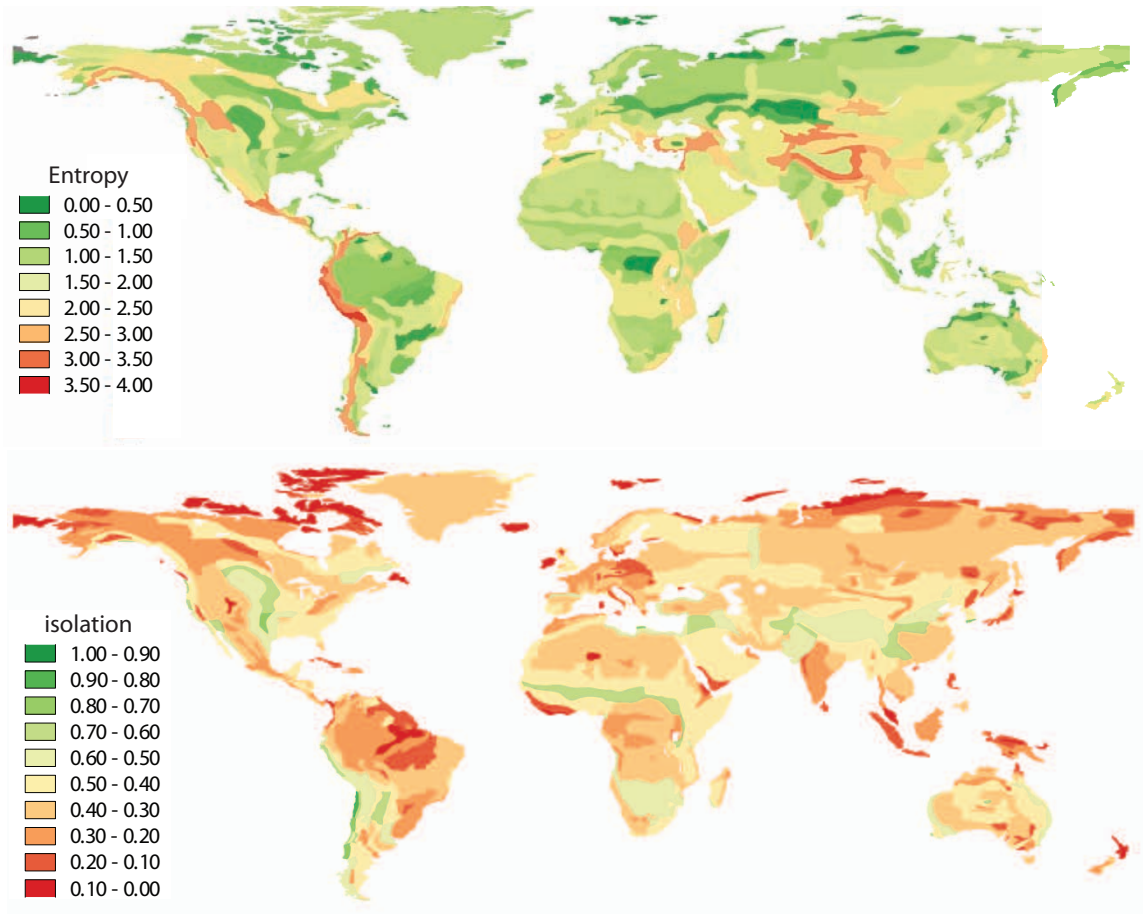


Figure 12: Maps of quality metrics, inhomogeneity (top), isolation (middle), and quality (bottom) with respect to **bioclimate** for the **BEC** regionalization.

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.

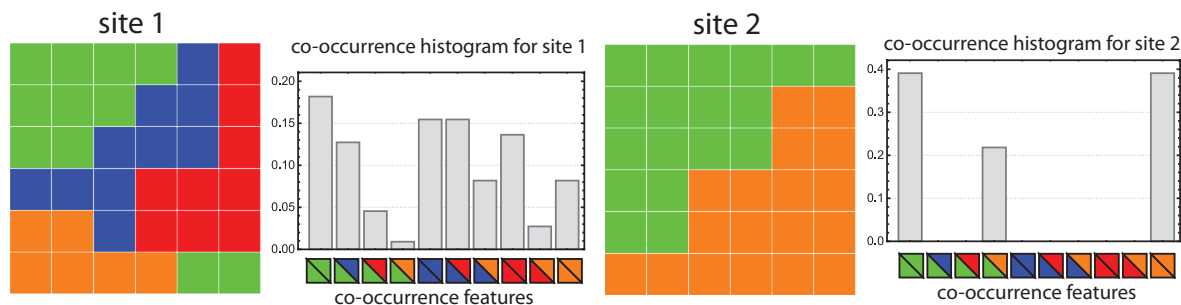


Figure 1: Co-occurrence histograms for two hypothetical sites with different patterns of variable categories. Four colors, red, blue, green, and orange indicate the four categories of the variable.

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.

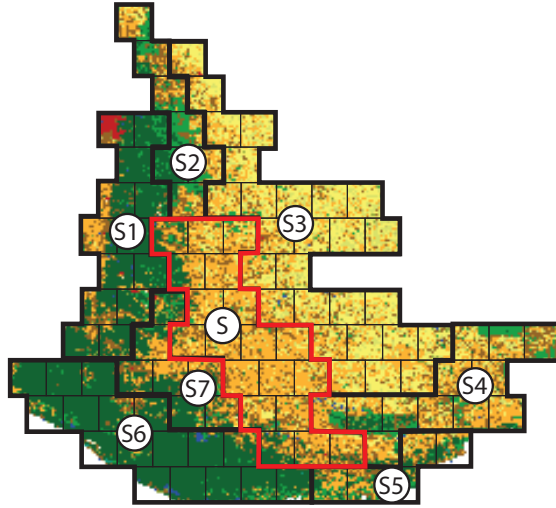


Figure 2: Focus segment S (outlined in red) has seven neighbors labeled as S_1 to S_7 and outlined in black. A linkage D is calculated between S and every neighbor. The seven values of D are averaged using weights which correspond to lengths of borders between S and the neighbors. The value of isolation (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.

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Supplement S3: Legends to categories of physiographic variables

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

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]

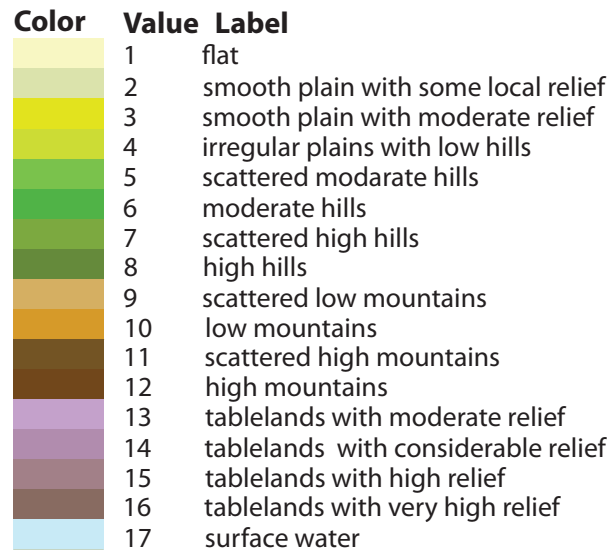


Figure 4: Legend for 17 categories of landforms. See Karagulle et al. [2017]

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