

## EUNIS habitat type F9.2, predicted habitat suitability - version 1, June 2016

The modelled suitability for the EUNIS habitat type is an indication of where conditions are favourable for the habitat type based on sample plot data (Braun-Blanquet database) and the Maxent software package. The modelled suitability map may be used as a proxy for the geographical distribution of the habitat type. Note however that it is not representing the actual distribution of the habitat type.

Also note that predictions are less reliable due to data deficiency in the eastern part of Europe, and to a lesser extent to the Scandinavian countries.

Geographic restriction for plot observations: n/a

Remarks: Prediction in eastern part of Europe uncertain due to lack of data for that area.

### Simple

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<b>Point of contact</b>	<b>Organisation name</b>	<b>Individual name</b>	<b>Electronic mail address</b>	<b>Website Role</b>
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### Point of contact

No information provided.

<b>Maintenance and update frequency</b>	Unknown
<b>GEMET - INSPIRE themes, version 1.0</b>	<ul style="list-style-type: none"> <li>Habitats and biotopes</li> </ul>
<b>GEMET</b>	<ul style="list-style-type: none"> <li>natural area</li> <li>heathland</li> <li>terrestrial ecosystem</li> <li>tundra</li> </ul>
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<b>Place</b>	<ul style="list-style-type: none"> <li>Europe</li> </ul>
<b>EEA topics</b>	<ul style="list-style-type: none"> <li>Biodiversity</li> </ul>
<b>Use limitation</b>	EEA standard re-use policy: unless otherwise indicated, re-use of content on the EEA website for commercial or non-commercial purposes is permitted free of charge, provided that the source is acknowledged ( <a href="http://www.eea.europa.eu/legal/copyright">http://www.eea.europa.eu/legal/copyright</a> ). Copyright holder: European Environment Agency (EEA).

<b>Access constraints</b>	Other restrictions
<b>Other constraints</b>	<a href="#">no limitations to public access</a>
<b>Spatial representation type</b>	Grid
<b>Distance</b>	1 1 km
<b>Language of dataset</b>	English
<b>Character set</b>	UTF8
<b>Topic category</b>	<ul style="list-style-type: none"><li>• Biota</li></ul>

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<b>Begin date</b>	1940-01-01		
<b>End date</b>	2011-12-31		
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<b>Distribution format</b>	• GeoTIFF ( )		
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<b>Hierarchy level</b>	Dataset		

## Conformance result

<b>Date (Publication)</b>	2010-12-08
<b>Explanation</b>	See the referenced specification

<b>Statement</b>	<p>The database compiled for the Braun-Blanquet project is a compilation of various national and regional vegetation databases. The maintenance of these databases is in principle in the hands of the custodians. However, before uploading the databases into Braun-Blanquet database a quality check is performed by Alterra and Masaryk University. If possible, detected errors are corrected and reported back to the data provider. For the modelling of the habitat suitability map the Maxent software is used ( <a href="http://www.cs.princeton.edu/~schapire/maxent/">http://www.cs.princeton.edu/~schapire/maxent/</a> ). The grid values in the map represent the probability (ranging from 0-1) that the cell is suitable for the habitat.</p> <p>The grid file represents the habitat suitability of the EUNIS type. For the modelling the widely used software Maxent for maximum entropy modelling of species' geographic distributions was used. Maxent is a general-purpose machine-learning method with a simple and precise mathematical formulation, and has a number of aspects that make it well-suited for species distribution modelling when only presence (occurrence) data but not absence data are available (Phillips et al. 2006). Because EUNIS habitats have a particular species composition, they are assumed to respond to specific ecological requirements, allowing us to generate correlative estimates of geographic distributions. Modelling habitats that have been floristically defined is a well-known procedure for ecological modelling at local scales, and a promising technique to be applied also at the continental level.</p> <p>The Maxent method considers presence data (known observations of a given entity) and the so-called background data. Background data comprise a set of points used to describe the environmental variation of the study area according to the available environmental layers. It is assumed that these layers represent well the most important ecological gradients on a European scale. As layers the following environmental parameters have been used: Potential Evapotranspiration, Topsoil pH, Solar radiation, Temperature Seasonality (standard deviation *100), Mean Temperature of Wettest Quarter, Annual Precipitation, Precipitation Seasonality (Coefficient of Variation), Precipitation of Warmest Quarter &amp; Distance to water (rivers, lakes, sea).</p> <p>Statistical output of the model:</p> <p>#Training samples: 1234</p> <p>Regularized training gain: 0.472</p>
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Unregularized training gain: 0.5897

Iterations: 500

Training AUC: 0.7945

#Test samples: 137

Test gain: 0.4925

Test AUC: 0.7679

AUC Standard Deviation: 0.0168

#Background points: 5798

bio\_12\_etr2\_ras contribution: 2.8352

bio\_15\_etr2\_ras contribution: 3.6013

bio\_18\_etr2\_ras contribution: 11.8177

bio\_4\_etr2\_ras contribution: 1.0261

bio\_8\_etr2\_ras contribution: 1.021

bld\_m\_sd1\_1km\_eu\_ll contribution: 1.6898

cecum\_m\_sd1\_1km\_eu\_ll contribution: 0.2901

clyppt\_m\_sd1\_1km\_eu\_ll contribution: 1.8138

crvol\_m\_sd1\_1km\_eu\_ll contribution: 31.0597

dist2water1km contribution: 1.0777

orcdrc\_m\_sd1\_1km\_eu\_ll contribution: 1.153

pet\_he\_yr contribution: 5.1577

sltppt\_m\_sd1\_1km\_eu\_ll contribution: 2.4878

sndppt\_m\_sd1\_1km\_eu\_ll contribution: 2.8443

solar\_1km contribution: 32.1247

bio\_12\_etr2\_ras permutation importance: 4.558

bio\_15\_etr2\_ras permutation importance: 5.6519

bio\_18\_etr2\_ras permutation importance: 11.506

bio\_4\_etr2\_ras permutation importance: 3.1255

bio\_8\_etr2\_ras permutation importance: 2.0508

bld\_m\_sd1\_1km\_eu\_ll permutation importance: 5.028

cecum\_m\_sd1\_1km\_eu\_ll permutation importance: 0.8506

clyppt\_m\_sd1\_1km\_eu\_ll permutation importance: 5.5312

crvol\_m\_sd1\_1km\_eu\_ll permutation importance: 23.9301

dist2water1km permutation importance: 1.1233

orcdrc\_m\_sd1\_1km\_eu\_ll permutation importance: 1.8734

pet\_he\_yr permutation importance: 7.7424

sltppt\_m\_sd1\_1km\_eu\_ll permutation importance: 2.8432

sndppt\_m\_sd1\_1km\_eu\_ll permutation importance: 3.1127

solar\_1km permutation importance: 21.0731

Training gain without bio\_12\_etr2\_ras: 0.466

Training gain without bio\_15\_etr2\_ras: 0.4594

Training gain without bio\_18\_etr2\_ras: 0.4605

Training gain without bio\_4\_etr2\_ras: 0.468

Training gain without bio\_8\_etr2\_ras: 0.4678

Training gain without bld\_m\_sd1\_1km\_eu\_ll: 0.4665

Training gain without cecsum\_m\_sd1\_1km\_eu\_ll: 0.4705

Training gain without clyppt\_m\_sd1\_1km\_eu\_ll: 0.4667

Training gain without crvol\_m\_sd1\_1km\_eu\_ll: 0.4411

Training gain without dist2water1km: 0.4678

Training gain without orcdrc\_m\_sd1\_1km\_eu\_ll: 0.4681

Training gain without pet\_he\_yr: 0.4585

Training gain without sltppt\_m\_sd1\_1km\_eu\_ll: 0.4694

Training gain without sndppt\_m\_sd1\_1km\_eu\_ll: 0.4687

Training gain without solar\_1km: 0.445

Training gain with only bio\_12\_etr2\_ras: 0.1094

Training gain with only bio\_15\_etr2\_ras: 0.033

Training gain with only bio\_18\_etr2\_ras: 0.1928

Training gain with only bio\_4\_etr2\_ras: 0.0593

Training gain with only bio\_8\_etr2\_ras: 0.0847

Training gain with only bld\_m\_sd1\_1km\_eu\_ll: 0.0798

Training gain with only cecsum\_m\_sd1\_1km\_eu\_ll: 0.1247

Training gain with only clyppt\_m\_sd1\_1km\_eu\_ll: 0.0173

Training gain with only crvol\_m\_sd1\_1km\_eu\_ll: 0.2713

Training gain with only dist2water1km: 0.0341

Training gain with only orcdrc\_m\_sd1\_1km\_eu\_ll: 0.0866

Training gain with only pet\_he\_yr: 0.1019

Training gain with only sltppt\_m\_sd1\_1km\_eu\_ll: 0.1146

Training gain with only sndppt\_m\_sd1\_1km\_eu\_ll: 0.0766

Training gain with only solar\_1km: 0.2602

Test gain without bio\_12\_etr2\_ras: 0.4698

Test gain without bio\_15\_etr2\_ras: 0.4778

Test gain without bio\_18\_etr2\_ras: 0.4733

Test gain without bio\_4\_etr2\_ras: 0.4828

Test gain without bio\_8\_etr2\_ras: 0.482

Test gain without bld\_m\_sd1\_1km\_eu\_ll: 0.511

Test gain without cecsum\_m\_sd1\_1km\_eu\_ll: 0.495

Test gain without clyppt\_m\_sd1\_1km\_eu\_ll: 0.4921

Test gain without crvol\_m\_sd1\_1km\_eu\_ll: 0.4835

Test gain without dist2water1km: 0.476

Test gain without orcdrc\_m\_sd1\_1km\_eu\_ll: 0.4875

Test gain without pet\_he\_yr: 0.4828

Test gain without sltppt\_m\_sd1\_1km\_eu\_ll: 0.485

Test gain without sndppt\_m\_sd1\_1km\_eu\_ll: 0.4792

Test gain without solar\_1km: 0.5002

Test gain with only bio\_12\_etr2\_ras: 0.1721

Test gain with only bio\_15\_etr2\_ras: 0.0342

Test gain with only bio\_18\_etr2\_ras: 0.229  
Test gain with only bio\_4\_etr2\_ras: 0.0821  
Test gain with only bio\_8\_etr2\_ras: 0.1543  
Test gain with only bld\_m\_sd1\_1km\_eu\_ll: 0.0971  
Test gain with only cecsum\_m\_sd1\_1km\_eu\_ll: 0.1399  
Test gain with only clyppt\_m\_sd1\_1km\_eu\_ll: 0.0329  
Test gain with only crvol\_m\_sd1\_1km\_eu\_ll: 0.2595  
Test gain with only dist2water1km: 0.0712  
Test gain with only orcdrc\_m\_sd1\_1km\_eu\_ll: 0.0989  
Test gain with only pet\_he\_yr: 0.1181  
Test gain with only sltppt\_m\_sd1\_1km\_eu\_ll: 0.1319  
Test gain with only sndppt\_m\_sd1\_1km\_eu\_ll: 0.116  
Test gain with only solar\_1km: 0.2465  
AUC without bio\_12\_etr2\_ras: 0.7616  
AUC without bio\_15\_etr2\_ras: 0.7637  
AUC without bio\_18\_etr2\_ras: 0.7635  
AUC without bio\_4\_etr2\_ras: 0.7654  
AUC without bio\_8\_etr2\_ras: 0.7633  
AUC without bld\_m\_sd1\_1km\_eu\_ll: 0.7743  
AUC without cecsum\_m\_sd1\_1km\_eu\_ll: 0.7686  
AUC without clyppt\_m\_sd1\_1km\_eu\_ll: 0.768  
AUC without crvol\_m\_sd1\_1km\_eu\_ll: 0.7692  
AUC without dist2water1km: 0.762  
AUC without orcdrc\_m\_sd1\_1km\_eu\_ll: 0.7658  
AUC without pet\_he\_yr: 0.767  
AUC without sltppt\_m\_sd1\_1km\_eu\_ll: 0.7641  
AUC without sndppt\_m\_sd1\_1km\_eu\_ll: 0.7642  
AUC without solar\_1km: 0.7702  
AUC with only bio\_12\_etr2\_ras: 0.6604  
AUC with only bio\_15\_etr2\_ras: 0.5555  
AUC with only bio\_18\_etr2\_ras: 0.6796  
AUC with only bio\_4\_etr2\_ras: 0.6188  
AUC with only bio\_8\_etr2\_ras: 0.6718  
AUC with only bld\_m\_sd1\_1km\_eu\_ll: 0.5966  
AUC with only cecsum\_m\_sd1\_1km\_eu\_ll: 0.6361  
AUC with only clyppt\_m\_sd1\_1km\_eu\_ll: 0.5725  
AUC with only crvol\_m\_sd1\_1km\_eu\_ll: 0.6815  
AUC with only dist2water1km: 0.6108  
AUC with only orcdrc\_m\_sd1\_1km\_eu\_ll: 0.6134  
AUC with only pet\_he\_yr: 0.6106  
AUC with only sltppt\_m\_sd1\_1km\_eu\_ll: 0.6365  
AUC with only sndppt\_m\_sd1\_1km\_eu\_ll: 0.634  
AUC with only solar\_1km: 0.6787

Entropy: 8.1975

Prevalence (average of logistic output over background sites): 0.3091

Fixed cumulative value 1 cumulative threshold: 1

Fixed cumulative value 1 logistic threshold: 0.0747

Fixed cumulative value 1 area: 0.7585

Fixed cumulative value 1 training omission: 0.0049

Fixed cumulative value 1 test omission: 0.0146

Fixed cumulative value 1 binomial probability: 2.74E-10

Fixed cumulative value 5 cumulative threshold: 5

Fixed cumulative value 5 logistic threshold: 0.2263

Fixed cumulative value 5 area: 0.609

Fixed cumulative value 5 training omission: 0.0284

Fixed cumulative value 5 test omission: 0.073

Fixed cumulative value 5 binomial probability: 1.19E-14

Fixed cumulative value 10 cumulative threshold: 10

Fixed cumulative value 10 logistic threshold: 0.3038

Fixed cumulative value 10 area: 0.5233

Fixed cumulative value 10 training omission: 0.0583

Fixed cumulative value 10 test omission: 0.0949

Fixed cumulative value 10 binomial probability: 1.81E-19

Minimum training presence cumulative threshold: 0.0973

Minimum training presence logistic threshold: 0.0137

Minimum training presence area: 0.8974

Minimum training presence training omission: 0

Minimum training presence test omission: 0

Minimum training presence binomial probability: 3.78E-05

10 percentile training presence cumulative threshold: 15.8761

10 percentile training presence logistic threshold: 0.3701

10 percentile training presence area: 0.4514

10 percentile training presence training omission: 0.0997

10 percentile training presence test omission: 0.146

10 percentile training presence binomial probability: 1.39E-21

Equal training sensitivity and specificity cumulative threshold: 36.1342

Equal training sensitivity and specificity logistic threshold: 0.481

Equal training sensitivity and specificity area: 0.2832

Equal training sensitivity and specificity training omission: 0.2828

Equal training sensitivity and specificity test omission: 0.2847

Equal training sensitivity and specificity binomial probability: 1.52E-29

Maximum training sensitivity plus specificity cumulative threshold: 26.821

Maximum training sensitivity plus specificity logistic threshold: 0.4385

Maximum training sensitivity plus specificity area: 0.3513

Maximum training sensitivity plus specificity training omission: 0.1904

Maximum training sensitivity plus specificity test omission: 0.2482

Maximum training sensitivity plus specificity binomial probability: 4.64E-23

Equal test sensitivity and specificity cumulative threshold: 35.9049

Equal test sensitivity and specificity logistic threshold: 0.48

Equal test sensitivity and specificity area: 0.2848

Equal test sensitivity and specificity training omission: 0.2796

Equal test sensitivity and specificity test omission: 0.2847

Equal test sensitivity and specificity binomial probability: 2.95E-29

Maximum test sensitivity plus specificity cumulative threshold: 36.7815

Maximum test sensitivity plus specificity logistic threshold: 0.4832

Maximum test sensitivity plus specificity area: 0.2787

Maximum test sensitivity plus specificity training omission: 0.2869

Maximum test sensitivity plus specificity test omission: 0.2847

Maximum test sensitivity plus specificity binomial probability: 2.14E-30

Balance training omission, predicted area and threshold value cumulative threshold: 2.47

Balance training omission, predicted area and threshold value logistic threshold: 0.1364

Balance training omission, predicted area and threshold value area: 0.6787

Balance training omission, predicted area and threshold value training omission: 0.0089

Balance training omission, predicted area and threshold value test omission: 0.0219

Balance training omission, predicted area and threshold value binomial probability: 3.08E-14

Equate entropy of thresholded and original distributions cumulative threshold: 4.2318

Equate entropy of thresholded and original distributions logistic threshold: 0.2101

Equate entropy of thresholded and original distributions area: 0.6263

Equate entropy of thresholded and original distributions training omission: 0.0243

Equate entropy of thresholded and original distributions test omission: 0.0584

Equate entropy of thresholded and original distributions binomial probability: 1.18E-14

Source

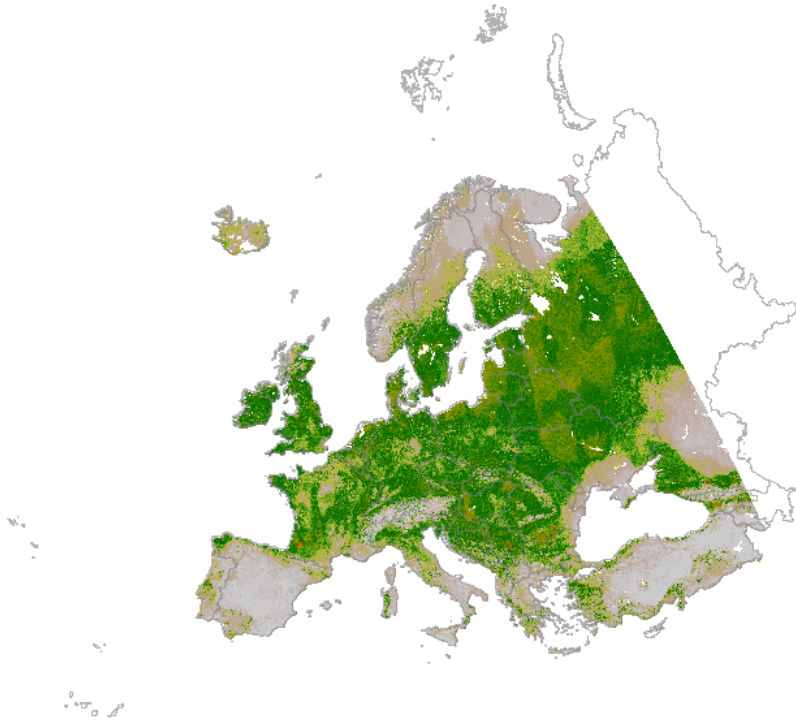
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## Overviews



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