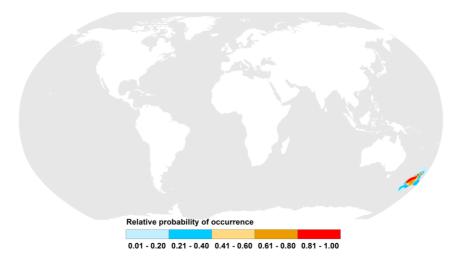
Global Distribution of Hector's Dolphins (2013)



Description: This dataset shows the modelled distribution of Hector's dolphins

(Cephalorhynchus hectori). AquaMaps (www.aquamaps.org) is a species

distribution modelling approach that provides standardised range maps for marine species using available information on species occurrence. IUCN status: Endangered

(Red List of Threatened Species).

Citation(s): Kaschner K, Rius-Barile J, Kesner-Reyes K, Garilao C, Kullander SO, Rees T, Froese R

(2013). AquaMaps: Predicted range maps for aquatic species. World wide web

electronic publication: www.aquamaps.org, Version August 2013

Temporal range:

2013

Global

Geographical

range:

Supplementary information (eg attribute table):

The dataset contains continuous probabilities of occurrence as a global grid of 0.5 dd resolution. Field information: scientific name (Genus, Species); center latitude of 0.5 dd cell (Center Lat); center longitude of 0.5 dd cell (Center Long); unique cell

identifier following the c-squares code system (C-Square Code; see

http://www.cmar.csiro.au/csquares/about-csquares.htm for more information); total predicted relative environmental suitability based on envelope settings

(Overall Probability).

Detailed dataset-specific information (provided by K. Kaschner) is also available.

Purpose of creation:

AquaMaps is a tool for generating model-based, large-scale predictions of natural occurrences of species. The maps can be used to inform environmental impact assessments.

Creation methodology:

The modelled distribution was generated using the AquaMaps online species distribution model. Methodological notes (provided by K. Kaschner) are also available. Observed occurrence records (< 10 cells; FAO areas: 81) were obtained (August 2013) from the Global Biodiversity Information Facility (GBIF; www.gbif.org). These were supplemented by additional information obtained through online species databases such as FishBase (www.fishbase.org) and SeaLifeBase (www.Sealifebase.org), as well as published information on species-

specific habitat usage and expert knowledge. The distribution model was based on four environmental variables (depth, temperature, salinity, and primary



Dataset ID: Kaschner-004

production), and details of the species envelope are in the dataset-specific information (provided by K. Kaschner).

Lineage (versioning): AquaMaps version used: 01 August 2013. In acknowledgement that predicted distributions reflect the current state of knowledge, AquaMaps predictions are not a permanent, fixed output, but instead will be revised and updated as new data or information become available or additional experts provide new input.

Environmental dataset used: HCAF version 5 (August 2013; www.aquamaps.org/envtdata/main.php).

Category: Species distribution

Keywords: marine, coastal, model, pelagic

Similar datasets: None

Quality, limitation(s), fitness for use: Excluded from the model: species misidentifications, fossil records and outliers. For this species, there are less than 10 "good cells" (i.e. Presence records in cells known to fall within the geographic range of the species) and therefore insufficient data to calculate envelopes from occurrence records. All envelope settings are therefore based on expert knowledge only.

The modelled distribution has been expert-reviewed by Kristin Kaschner (15 December 2013), and the quality of predictions ranks 4 out of 5 (see www.aquamaps.org/rating.html for further details). Expert comment: "Good representation of known occurrence of species, except for false predicted presences around Stewart Island and other offshore shallow banks southwest of New Zealand and off the eastern and southern coast of the North Island, which are not supported by data. Also distribution of North Island subpopulation (Maui's dolphin) along the West coast underestimated and predicted concentrations in South Taranaki Bight more representative of historic range (currently highest numbers between Taranaki & Ninety Mile Beach)".

Maintenance frequency:

Data are updated in intervals that are uneven in duration.

Main access/use constraint:

Creative Commons Attribution-NonCommercial 3.0 Unported (CC BY-NC 3.0). See http://creativecommons.org/licenses/by-nc/3.0/ for details. Free to (1) copy/distribute the work, and (2) adapt the work. The material may not be used for commercial purposes.

constraints:

Other access/use Interested users of the dataset should contact Kristin Kaschner who will identify and provide, where appropriate, the most recent updated data.

For commercial use, please contact business-support@unep-wcmc.org.



Dataset ID: Kaschner-004

Contact Albert-Ludwigs-University of Freiburg

organisation:

Organisation Owner Acronym:

type:

Name: Dr. Kristin Kaschner Position: Research Affiliate

City: Country: Germany

E-mail: Kristin.Kaschner@biologie.uni-freiburg.de

Web site: www.uni-freiburg.de

Main format: Tabular (.csv) Other Vector (point; .shp)

format(s):

Distribution Tabular (.csv) Dataset size 14 Kb

format: (uncompressed):

Webpage and/or http://www.aquamaps.org

download:

Other webpage: http://data.unep-wcmc.org/datasets/32

Web map service:

Resolution, scale: 0.5 dd Reference system: WGS 1984

West bounding: 160.0 East bounding: 180.0

South bounding: -90.0 North bounding: 0.0 Date of metadata:

Factsheet: No Metadata standard: UNEP-WCMC Specific 13/10/2014

Hector's dolphin (Cephalorhynchus hectori)



Figure 1: Expert reviewed AquaMaps predictions: Hector's dolphin (*Cephalorhynchus hectori*)

Expert:	Kristin Kaschner					
Review data:	15/12/2013					
File name	Cehec_AQER_122013.csv					
Quality of Predictions:	4 (out of 5) http://www.aquamaps.org/rating.html					
Recommended threshold settings best describing utilized niche (precautionary setting)	0.0					
Expert comment:	Good representation of known occurrence of species, except for false predicted presences around Stewart Island and other offshore shallow banks southwest of New Zealand and off the eastern and southern coast of the North Island, which are not supported by data. Also distribution of North Island subpopulation (Maui's dolphin) along the West coast underestimated and predicted concentrations in South Taranaki Bight more representative of historic range (currently highest numbers between Taranaki & Ninety Mile Beach)					
AquaMaps version:	01/08/2013					
Environmental data set:	http://www.aquamaps.org/envtdata/main.php HCAF ver.5 (08/2013)					
AquaMaps citation:	Kaschner, K., J. Rius-Barile, K. Kesner-Reyes, C. Garilao, S.O. Kullander, T. Rees, and R. Froese. 2013. AquaMaps: Predicted range maps for aquatic species. World wide web electronic publication, www.aquamaps.org, Version 08/2013.					

Detailed dataset-specific information (provided by K. Kaschner)

Input data



Figure 2: Hector's dolphin (*Cephalorhynchus hectori*)- available point occurrence records (GBIF download August 2013). Note that green records represent species misidentification or museum records and associated points are not included in the envelope calculations. There are less than 10 "good cells" (i.e. presence records in cells known to fall within the geographic range of the species) and therefore insufficient data to calculate envelopes from occurrence records. All envelope settings are therefore based on expert knowledge only.

Mapping parameters for Cephalorhynchus hectori (Hector's dolphin)								
Map type: Expert-reviewed								
Map Option: In FAO and Bounding Box								
FAOAreas: 81								
Bounding Box								
(NSWE):	0	-90	160	180				
No of good cells used								
for envelope	_							
calculation	0							
Pelagic: False								
Layer used to generate pr	robabiliti	es: Surf	face					
Species Envelope (HSPEN):								
			Pref Min	Pref Max				
	Used	Min	(10th)	(90th)	Max			
Depth (m)	1	0	1	50	200			
Temperature (°C)	1	5	10	15	20			
Salinity (psu)	1	33.5	34	35	35.6			
Primary Production	1	250	400	1000	1250			
Sea Ice Concentration	0							

Detailed dataset-specific information (provided by K. Kaschner)

Distance to Land (km)	0		

Raw point occurrence data & associated environmental values: Hec_RawOcc_122013.xls

Methodology for producing maps of known and probable worldwide distributions of marine mammals using the AquaMaps approach

Kristin Kaschner, December 2013

General methodology of AquaMaps approach

Aquamaps is an online species distribution model (www.aquamaps.org) that allows the generation of standardized digital range maps of aquatic species, currently covering more than 17 000 species. Maps are generated using a modified version of the relative environmental suitability model (RES¹) developed by (Kaschner et al. 2006) that uses available information about habitat usage of a given species, projected into geographic space, to help visualize its distribution. Habitat usage is quantitatively described with the help of so-called environmental envelopes defining a species' preference with respect to a set of pre-defined environmental conditions, including depth, sea-ice, temperature, salinity and primary production. By default, envelopes are derived from occurrence records available through GBIF (www.gbif.org) supplemented by additional information obtained through online species databases such as FishBase (www.fishbase.org) and SeaLifeBase (www.Sealifebase.org). Acknowledging the sampling biases of currently available online occurrence data, however, AquaMaps explicitly also allows for experts to review and modify environmental envelopes manually. Map outputs represent annual average predictions of the maximum range extent of species (defined as the maximum area between the known outer-most limits of a species' regular or periodic occurrence) and gradients of relative habitat suitability or species occurrences (ranging from 0.00 – 1.00), predicted for each 0.5 degree latitude by 0.5 degree longitude cells. Predictions represent a visualization of the basic environmental niche of a species, which may often be closer to the historic occurrence of species or its potential niche rather than its realized or currently occupied niche. Binary range maps corresponding more closely to areas of known occurrence may be derived using presence thresholds ideally defined by validation analysis (Kaschner et al. 2011) (see below).

¹ Relative Environmental Suitability

AquaMaps predictions for different species have been validated using independent data sets (Kaschner et al. 2006, 2011, Ready et al. 2010) and generally capture existing knowledge of large-scale, long-term annual average species occurrence reasonably well. However, given the overall paucity of data and the frequently large sampling biases in the marine environment, produced outputs should be regarded as hypotheses of species occurrence, based on a clearly defined set of assumptions that can be tested and further refined as new data become available. Moreover, since marine mammal habitat usage often varies across seasons and ocean basins, global predictions should not be used without further review to describe regional species occurrence (and should ideally be checked against independent data) and the overall limitations of data availability, model biases and assumptions etc. should be kept in mind when using produced outputs for management purposes.

Specific methodology for generating updated annual average AquaMaps predictions (all 10 species)

Expert-review was based on environmental envelopes computed from the most recent AquaMaps harvest of occurrence data from GBIF (www.gbif.org) in August 2013. For each species, point occurrence records and resulting 0.5 degree presence cells were reviewed to exclude false records (species misidentifications, fossil records and outliers) based on a comparison of published information about species distributions including, but not limited to, IUCN species pages. Calculated envelopes based on the final subset were further reviewed to ensure that these matched available information about habitat usages as published in the literature. Predictions about the relative probability of occurrence /habitat suitability were then generated based on these reviewed envelopes. Finally, the resulting predictions were reviewed by comparing them with existing information about the maximum range extent and relative occurrence of species within that range, highlighting both false predicted presences and absences. Quality of predictions is reflected in the assigned 5 rank (1 worst best see http://www.aquamaps.org/rating.html) associated with all outputs. It should be noted that the top two ranks are only assigned if predictions have been successfully and

quantitatively validated using independent effort-corrected survey data throughout the whole range ("5") or for at least part of the species range ("4") and as the time available for this project was insufficient for conducting these types of validation, the top rank assigned was a "3" (with the exception of sperm whales for which a quantitative validation had been carried out using data from Antarctic waters).

Presence threshold to be used for producing binary² range maps

Validation analyses have shown strong correlations between observed relative species occurrence and predicted relative environmental suitability as predicted by RES and AquaMaps for the majority of species and areas with enough data from large-scale, long-term dedicated marine mammal surveys to allow testing (Kaschner et al. 2006, 2011, Ready et al. 2010). Observed species densities tend to be highest in areas of predicted probability > 0.4 to 0.6, and validation analysis indicated that this is the most likely presence threshold that should be used to produce the most likely representation of known and probable occurrence of the species, although this may vary for different species. The threshold recommended in the individual species files are based on a precautionary approach that should be used in light of existing uncertainties and in the context of environmental impact assessment.

Kaschner K, Tittensor DP, Ready J, Gerrodette T, Worm B (2011) Current and future patterns of global marine mammal biodiversity. PLoS One 6:e19653

Kaschner K, Watson R, Trites A, Pauly D (2006) Mapping world-wide distributions of marine mammal species using a relative environmental suitability (RES) model. Mar Ecol progess Ser 316:285–310

Ready J, Kaschner K, South AB, Eastwood PD, Rees T, Rius J, Agbayani E, Kullander S, Froese R (2010) Predicting the distributions of marine organisms at the global scale. Ecol Modell 221:467–478

² i.e. presence/absence