Region-specific magnetic fields structure sea turtle populations

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Abstract

Philopatry and long distance migrations are common in the animal kingdom, of which sea
turtles are flagship examples. Recent studies have suggested sea turtles use the Earth's
magnetic field to navigate across ocean basins to return to their natal area, yet the
mechanisms underlying this process remain unknown. If true though, the genetic structure at
nesting sites should positively correlate with differences in location-specific magnetic vectors
within nesting regions. Here, we confirm this working hypothesis but only in certain regions
of the world and for all sea turtle species nesting in those regions. Reversely, where no
correlations were found between genetic differentiation and geomagnetic vectors, this was
the case for all nesting sea turtle species. Our approach hence reveals parallel but not
universal use of geomagnetic cues in sea turtles. We describe magneto-sensing regions as
characterized by sharp clines of total and vertical field intensity vectors offering the
navigation cues that increase philopatric accuracy and promote genetic structuring among
sea turtle populations.

Introduction

Philopatry is a well-documented trait of many species, including insects, birds, mammals and
reptiles (Greenwood 1980; FitzSimmons *et al.* 1997). With this behavior, after long-distance
migration, individuals return to their natal area upon sexual maturity to mate, sometimes
decades after leaving at an early life stages. While philopatry can be a learned trait from
older individuals (e.g. sheep, moose (Jesmer *et al.* 2018), whales (Baker *et al.* 2013)), in
others, it is an evolved behaviour associated with long-distance navigation and imprinting,
such as butterflies, salmon, and sea turtles (Brower 1995; Dittman & Quinn 1996; Lohmann
et al. 1999). Sea turtles provide perhaps the best illustration of philopatry in the marine
environment since they return to their natal area after ~25-30 years and thousands of kms of
juvenile migration (Wyneken *et al.* 2013). Philopatry can be as accurate as 50 km (Hatase *et*
al. 2002; Bowen & Karl 2007) and shapes population structure (Stiebens *et al.* 2013b). Yet,
its underlying mechanisms remain partly speculative.

Several studies have demonstrated that the Earth's magnetic field provides sufficient cues to

guide animal migration (Wiltschko & Wiltschko 2005; Lohmann *et al.* 2008b). The Earth

magnetism is driven by the motion of the molten iron in Earth's core and is defined by seven

vectors (Campbell 2003). The shifting of liquid iron also means the magnetic field is in a

constant state of flux over various time-scales. While it is acknowledged that imprinting

occurs during the early stages of development or birth (Lohmann *et al.* 2008b), the

underlying mechanisms of geomagnetic navigation remain unknown, particularly in sea

turtles. Nevertheless, if, globally, geomagnetic imprinting determines philopatric behaviour

(Brothers & Lohmann 2015), it should leave a signature of selection on the population

genetic structure of sea turtles. This is because, within a region, if a location's (nesting site

or natal area) geomagnetic profile differs extensively from neighbouring locations, the

navigation accuracy of sea turtles should be high and genetic structure should increase, i.e.

gene flow among populations should decrease.

By examining the possible determinants of genetic structure across the nesting aggregations

(regions) of sea turtles, and across species with overlapping nesting distributions, we should

be able to disentangle species-specific features of magneto-sensing from region-specific

geomagnetic characteristics. This approach compensates for the lack of known underlying

molecular mechanisms and the associated cline analysis of allele frequency change

(Günther & Coop 2013). Positive correlations between geomagnetic field vectors and

population genetic structure would confirm that sea turtles use vectors of the earth magnetic

field for their philopatric migration. Finding no such pattern would challenge the imprinting

hypothesis of early life stages as a potentially universal mechanism of navigation.

Here, we combine a thorough literature search with sampling and mitochondrial control

region sequencing. We were able to obtain population structure data from five of the seven

sea turtle species from 144 locations (nesting sites and natal areas) in nine regions globally,

for ~ 17,470 sequences (Fig. 1). We obtained data from nesting regions where several

species co-exist allowing us to disentangle species from region-specific effects and explore

whether all species are equally responsive to geomagnetism. Specifically, we tested the

correlation between genetic structure, estimated from Wright fixation indices, and

geomagnetic vectors at natal areas. F_{ST} fixation indices from mitochondrial genes are widely

used as markers of philopatric behaviours because of their maternal mode of inheritance

(Bowen & Karl 2007).

Methods

<u>Literature search and data collection</u>

We took advantage of the well-characterized population genetic structure of sea turtle across
the globe. Data were collected from published literature following a two-step process. Firstly,
we used SWOT (The State of the World's Sea Turtles), IUCN (International Union for
Conservation of Nature) and the grey literature to identify all the world nesting aggregations
(Electronic supplementary material, Table S1). Secondly, we screened the primary and
secondary literature for all studies with indices of population structure expressed as a
pairwise Wright's fixation index (F_{ST}, γ_{ST} or φ_{ST}). To find this literature, we searched for "Sea
turtle AND" either "Phylogeography", "Genetic structure", "Population structure" or
"Mitochondrial DNA" together with the names of the nesting aggregations. We also
performed individual searches for each of the seven sea turtle species substituting "Sea
turtle" for: "Loggerhead", "Green turtle", "Hawksbill", "Leatherback", "Flatback", "Kemp's
Ridley" and "Olive Ridley".

As our goal was to specifically compare genetic and geomagnetic information within sea
turtle rookeries, obtaining exact nesting beach locations to derive accurate magnetic field
information was important. Therefore, if the literature contained only one nesting site from a
specific nesting region, this data point was removed (5 nesting sites removed). For literature
that contained both nesting site and feeding ground genetic information, feeding ground data
points were ignored. Data points in which genetic structure information was pooled from
multiple geographic locations into a single general natal area were removed (18 nesting sites
removed). When the exact nesting site sampled for a location could not be identified, data
points were also removed (3 nesting sites removed). This resulted in the use of 144
locations (Electronic supplementary material, Table S2).

Nesting sites and areas among sea turtle species were grouped into what we define as
"nesting regions" based on geographic location. We did not use regional management units
(RMUs) due to the lack of consistency of RMUs among species, which would prevent direct
comparison. It resulted into a dataset composed of 144 locations, from 9 different nesting
regions worldwide for 5 of the 7 sea turtle species. Noteworthy, as gene flow across regions
does not occur, data points between nesting regions did not enter the analyses to avoid
biasing for high values of genetic structure indices.

<u>Population structure in the largest North East Atlantic loggerhead turtle rookery</u>

To complement the existing information, we also added data from the Cape Verde
Archipelago, the 3rd largest aggregation of loggerhead turtles in the world. Field surveys took
place in the Cape Verde archipelago in 2011, 2012 and 2013 during the nesting season from
June to October. Nine different islands were sampled: Boavista, Fogo, Maio, Sal, Santa
Luzia, Santiago, Santo Antão, São Nicolau and São Vicente. A 3mm piece of non-

keratinized tissue was removed from the right front flipper using a single-use disposable

scalpel and samples were stored in ethanol (Stiebens *et al.* 2013a; Stiebens *et al.* 2013b).

Turtles were tagged with metal tags and/or pit tags directly after egg deposition to track

nesting behaviour and to avoid multiple sampling. In the laboratory, each sample was

washed in distilled water for 20 seconds and cut into smaller pieces. DNA from tissues was

extracted using the DNeasy® 96 Blood & Tissue Kit (QIAGEN, Hilden, Germany). Elution

was conducted in twice 75 µl of AE Buffer. All other steps followed the manufacturer's

protocol.

The control region of the mitochondrial DNA was amplified using the Primers LCM15382 (5'-

GCTTAACCCTAAAGCATTGG-3') and H950 (5'-GTCTCGGATTTAGGGGTTTG-3')(Abreu-

Grobois *et al.* 2006) following (Stiebens *et al.* 2013a). PCR products were cleaned with

ExoSAP-IT® following the manufacturer's protocol. Cycle sequencing reactions were

performed with Big Dye® Terminator v3.1 Cycle Sequencing Kit (Applied Biosystems,

Darmstadt, Germany). Sequences were obtained from the forward direction. In cases where

a sufficiently long fragment could not be obtained, fragments were additionally sequenced

from the reverse direction to build a longer contig. Sequencing was performed in an ABI

3730 Genetic Analyzer (Applied Biosystems, Darmstadt, Germany). Sequences were

assembled in Codon Code Aligner v5.0 (CodonCode Corporation, Dedham, Massachusetts)

and ambiguities were corrected by hand. All the amplified mitochondrial sequences were

classified accordingly to the standardized nomenclature of the Archie Carr Centre for Sea

turtle Research (http://accstr.ufl.edu). Wright's fixation index (F_{ST}) were computed with

10.000 permutations using Arlequin v3.5.1[28] (Electronic supplementary material, Table S3).

<u>Defining the geomagnetic profiles</u>

To evaluate whether the global nesting regions have significantly different geomagnetic

profiles and to test for the stability of the difference in the last century, for each nesting site,

we collected geomagnetic profiles at 3 time-points (the present day, standardized as

01/08/2015, 50 and 100 years prior to the present day). This confirmation step is important

since it serves as a pre-requisite for the geomagnetic field to act as a reliable source of

navigation cues everywhere and on an ecologically relevant timeframe. We used the

National Oceanic and Atmospheric Administration (NOAA) Magnetic Field Calculator

(http://www.ngdc.noaa.gov/geomag-web/#igrfwmm) and the International Geomagnetic

Reference Field (IGRF) model, by entering the GPS coordinate of each natal area. The

resulting geomagnetic field vectors included: declination and inclination in degrees (°),

horizontal intensity in nano-teslas (nT) as well as the northern component (nT), eastern

component (nT), vertical component (nT) and total field intensity (nT) at each natal area.

<u>Estimating migration error rate</u>

To estimate sea turtle navigational error (error rate), we focused on distances between
isolines (bands of equal magnetic intensity) within a region. We first established the most
northern and southern natal areas and their coordinates within each region. Next, we
obtained geographic coordinates at each change of 1000nT (i.e. a change in 1 isoline)
between these coordinates using the NOAA Magnetic Field Calculator
(https://www.ngdc.noaa.gov/geomag-web/#igrfgrid) with the IGRF. Each region was given a
standardized longitude comprised of the mean longitude of its most northern and southern
locations. We then calculated the distance in kilometres between the geographic coordinates
of each isoline using the NOAA Latitude/Longitude Distance Calculator. A region's
navigational error rate is therefore given by the mean of the distances between its isolines.

<u>Magnetic variation across regions</u>

To quantify the selection acting on turtle philopatric accuracy, we also assessed how much
error a turtle might accrue if it mistook adjacent vertical and total field intensity isolines.
Vertical field intensity and total field intensity at each 1-degree change in latitude across
regions were collected using the NOAA Magnetic Field Calculator with the IGRF. We defined
each region by their most northern and most southern natal areas and longitude for a region
was standardized as the mean of the longitudes of the natal areas within the region. This
particular data collection was done in 2-year intervals across 24 years till the present day
(01/08/1991 – 01/08/2015), based on the time it would take a turtle to sexually mature from
hatching. Natal locations were categorized into northern or southern hemisphere based on
their latitude.

<u>Data analyses</u>

Statistical analyses were performed in R Studio version 3.2.5 [(C) 2016 The R Foundation
for Statistical Computing]. Three between-class analyses (BCA from the *ade4* package
(Dray & Dufour 2007)) were performed (for the present day, 50 and 100 years prior to the
present day) to establish whether geomagnetic field vectors differ among sea turtle nesting
regions worldwide and whether these differences were stable across time.

To test for the possible correlation between geomagnetism and population structure (indices
of population genetics differentiation at specific locations), the individual geomagnetic
vectors were converted into distance matrices using the maximum distance (supremum
norm) between any two points of a breeding region. We used the maximum distance as this
accounts for potential land-barriers and Euclidean or ordinary-line distance would
oversimplify distances among natal locations. As the geomagnetic field is intrinsically linked

to geographic location (confirmed here; Pearson's product-moment correlation, t = 39.658, df
= 1287, p < 0. 001), geomagnetism in our models was used as the residuals of the
correlation between the magnetic field and geographic distance. Our global model included
sea turtle fixation indices as the response variable with geomagnetic residuals, geographic
distance between locations, sea turtle region and the two way interactions between the
geomagnetic residuals and the other variables. This model was backward selected using the
Akaike's Information Criterion (Burnham & Anderson 2004)(stepAIC). Further independent
linear models were performed to explore both the relationship between sea turtle population
genetic structure and the geomagnetic field as well as the relationship between the rate of
error in "magneto-sensing" and "non-magneto sensing" regions in the northern and southern
hemispheres.

Results and Discussion

Sea turtles require region-specific geomagnetic vectors that are stable over at least one
generation if they are to use the earth's magnetic field to conduct their philopatric migration.
Hence, we studied the geomagnetic signatures over the last century (present, 50 and 100
years before present) from all regions for the five sea turtle species for which we obtained
genetic data. We confirmed that each region has had a specific geomagnetic signature at
any time point over the last century (Randomization Test - Present: Obs = 0.049, p = 0.001;
50 years before present (BP): Obs = 0.827, p = 0.001; 100 years BP: Obs = 0.821, p =
0.001, Fig. 2). This result is a mandatory pre-requisite for the geomagnetic field to provide
informative navigational cues across nesting regions and over time.

A principal component analysis, for the present day, shows that PC1 explains 89.4% of total
variance, with PC2 explaining 6.42% (50-years before present (BP) PC1 = 90.23%, PC2 =
4.91%; 100-years BP PC1 = 89.55 %, PC2 = 5.9%). For PC1, 78.6% was explained by the
vertical field component (Z) followed by total field intensity (F) with 21.3%. The pattern
remained the same for both 50 (Z = 76.9%, F = 23.1% of PC1) and 100 years BP (Z =
77.2%, F = 22.8% of PC1). Thus, over the last century, this vector has been stable enough
for turtles to differentiate between nesting regions. Previous studies showed that sea turtles
from Florida can perceive magnetic total field intensity (Lohmann & Lohmann 1996;
Lohmann *et al.* 2001; Lohmann *et al.* 2008b; Putman *et al.* 2011), an element of which is the
vertical field component. Ultimately, this finding shows that geomagnetism is a basis acting
as a navigational cue that underlies philopatric behaviours in sea turtle species at a time-
scale relevant with both ecological (navigation) and evolutionary (population structure)
processes.

As a second step, we investigated the correlation between geomagnetic vectors and indices
of population structure for the five sea turtle species and 144 locations (Table S1). We found
that within nesting regions, the stronger the geomagnetic difference between nesting groups
is, the stronger the population structure ($F_{1,\ 694}$ = 52.845, p < 0.001). We also found
differences across species ($F_{4,\ 715}$ = 17.37, p < 0.001) with generally higher genetic structure
for the Hawksbill turtle (mean $\pm$ sd F_{ST} = 0.451 $\pm$ 0.283) and lowest for the Olive Ridley (F_{ST}
0.009 $\pm$ 0.044). Strikingly, no interaction between species and geomagnetic vectors was
detected ($F_{4,\ 694}$ = 0.997, p = 0.409). Instead, an interaction between regions and
geomagnetic vectors was a better predictor of the observed genetic structure ($F_{7,\ 694}$ =
11.620, p < 0.001, Fig. 3a). Specifically, we found that the Northern/Western Caribbean,
Northeast Pacific and Southwest Indian Ocean showed the expected positive correlation
between genetic structure and geomagnetic field following the geomagnetic imprinting
hypothesis (Northern/Western Caribbean: $F_{1,\ 256}$ = 12.75, p < 0.001, Northeast Pacific: $F_{1,\ 21}$
= 6.318, p = 0.020, Southwest Indian Ocean: $F_{1,\ 43}$ = 3.772, p = 0.059). Crucially, this pattern
was observed for all turtle species nesting within those regions, i.e. Green turtle, Hawksbill,
Leatherback, Loggerhead and the Olive Ridley (Electronic supplementary material, Table
S1). Thus, even though the absolute level of genetic differentiation may vary among sea
turtle species, all turtles nesting in those three regions have retained parallel use of
geomagnetic profile as a cue for their philopatric migration.

Conversely, in the six other nesting regions, we did not detect the expected relationship
between population genetic structure and region-specific magnetic field. Here as well, this is
true for all species nesting within those regions (All F < 1.199, all p > 0.281, Fig. 3b) and
suggests that, in these regions, the magnetic field might not provide sufficient information to
locate natal areas (i.e. area around their nesting site) with sufficiently high accuracy.
Noteworthy, as expected, across the globe and across species, geographic distance was a
good predictor of indices of population structure ($F_{1,\ 694}$ = 64.163, p < 0.001). Overall, we
demonstrate that geomagnetism-mediated philopatric behaviour is region-specific and not
species-specific. This conclusion is reinforced by the parallel patterns observed across
species within a region. But what are the characteristics of the earth magnetic field in some
regions compared to others that would explain geomagnetic imprinting?

For each geomagnetic vector, the Earth is stratified by lines of constant magnetic values,
known as isolines (Lohmann *et al.* 2008b)(Fig. 1). As coastlines are marked by isolines, sea
turtles are thought to follow magnetism of equal value back to the area near their nesting
sites (Lohmann *et al.* 2008b; Endres *et al.* 2016). Aggregated magnetic isolines across a
geographic area highlight intense clines of the specific geomagnetic vector, whereas more

dispersed isolines over the same distance represent weaker gradients of the same vector.
We tested whether the different groups of regions varied in the vertical and total field
intensity vectors, which we show best characterize the specific rookeries. We also assessed
whether there was any consistency in the variation across both hemispheres. We found
magnetic-regions, i.e. those with correlations between population structure and the
geomagnetic vectors, to differ significantly from non-magnetic regions for vertical field
intensity in both the northern and the southern hemisphere (Northern Hemisphere: $F_{1, 1376}$ =
383.5, p < 0.001, Southern Hemisphere: $F_{1, 713}$ = 219.5, p < 0.001, Fig. 4a). Magnetic
regions, therefore, are those that typically have increased vertical field intensity compared to
non-magnetic regions, regardless of hemisphere. Magnetic and non-magnetic regions also
differ in their total field intensity in the northern hemisphere ($F_{1, 1376}$ = 120.7, p < 0.001),
however no difference was found between regions in the southern hemisphere ($F_{1, 713}$ =
2.087, p = 0.149, Fig. 4b).
256

To quantify the selection acting on turtle philopatric accuracy, we determined how far a turtle
might stray if it mistakenly followed adjacent vertical and total field intensity isolines. We
show that sea turtles from geomagnetic regions have significantly lower rates of philopatric
error (223km ± 93km) based on total field intensity compared to those in non-geomagnetic
regions (390km ± 184km, $F_{1, 217}$ = 8.366, p < 0.001, Fig. 5). In contrast, no such difference
was found between geomagnetic regions (120km ± 21km) and non-geomagnetic regions
(124km ± 46km) for the error-rate if sea turtles were to use vertical field isolines alone ($F_{1, 227}$
= -0.889, p = 0.375). This result is consistent with research based on hatchling geomagnetic
orientation, which follow bi-coordinate geomagnetic map from which both longitudinal and
latitudinal information can be derived (Lohmann *et al.* 2008b; Putman *et al.* 2011). As the
vertical field only gives information towards the magnetic equator (i.e. equator = 0), it is
unlikely sea turtles would use it alone. Instead, turtles likely derive information from the
combined use of vertical and total field intensity.
270

While our findings are consistent with previous studies (Lohmann *et al.* 2008b; Endres *et al.*
2016; Brothers & Lohmann 2018), we demonstrate that geomagnetic imprinting is regional,
not universal. This result concurs with secular variation of the Earth's magnetic field, which
causes female sea turtles to change natal locations as magnetic signatures drift along
coastlines (Brothers & Lohmann 2015). In general, our result demonstrates that slight
inaccuracy in the detection of the geomagnetic field vector may lead turtles to fail to return to
breeding sites (Lohmann *et al.* 2008b) creating extreme selection pressure to increase
nesting site fidelity.
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Overall, our study reveals the selective pressure exerted by the geomagnetic field on sea
turtle population structure and how this contributes to sea turtle evolution. In geomagnetic-
regions, isolines are close enough to one another to provide sufficiently accurate
navigational information for turtles to reach their natal area and reproduce. This pattern of
philopatry will ultimately restrict gene flow among populations within regions and among
regions. In regions with distant isolines, navigation mistakes could result in the best case in
increased gene flow among natal areas, or in the worst case in failure to reproduce at all. In
these regions, philopatric journeys will likely require sea turtles to utilize multi-modal cues,
e.g. olfaction and geomagnetism (Lohmann *et al.* 1999; Lohmann *et al.* 2008a; Endres *et al.*
2016). Previous genetic analyses also show that the accuracy of sea turtle philopatric
journeys vary between different populations (Meylan *et al.* 1990; Bowen & Karl 2007; Lee *et*
al. 2007) and our results could suggest these differences in accuracy may be determined by
local cue. It would also explain why turtles with disrupted magneto-sensing capacity by the
application of magnet on their head, despite longer journey, find their natal area (Luschi *et*
al. 2007).

In our study, we cannot account for all the other parameters that also influence the
distribution of sea turtles. For instance, modern and historical climates (Poloczanska *et al.*
2009; Pike 2013), ocean currents (Putman *et al.* 2010) or plate tectonics (Bowen *et al.* 1989)
have all been implicated in the distribution of sea turtles. Our results suggest that rookery-
specific environmental pressures will define which parameter is the most important.
Interestingly, the framework we have detailed in this study should facilitate testing the
different possible drivers of population genetics and ultimately determine their relative
contribution in other species that have evolved philopatric behaviours.

In conclusion, while our results demonstrate that there is sufficient differentiation among the
geomagnetic fields of sea turtle rookeries, location-specific use of given vectors suggests the
parallel (across species) and local (to regions) adaptation of navigational cue recognition.
Whilst our analyses focus specifically on sea turtles, its relevance extends to other taxa
conducting long-distance migrations to and from their natal areas. Many other marine
organisms e.g. elephant seals, European eels or salmon have been hypothesized to imprint
on a natal geomagnetic field to complete their philopatric migrations (Matsumura *et al.* 2011;
Putman *et al.* 2013; Putman *et al.* 2014; Baltazar-Soares & Eizaguirre 2017; Naisbett-Jones
et al. 2017). Our results suggest that underlying mechanisms for geomagnetic navigation are
locally adapted and future research into such mechanisms should focus on evolved region-
specific and coupled systems rather than a universal molecular/physiological mechanism.

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Author Contributions

CE and SJKC designed the study, CE and SJKC conducted sampling in Cape Verde. CE
and MBS conducted the molecular analyses. SJKC performed all the statistics of the meta-
analysis and drafted the manuscript. All authors edited the manuscript and contributed to its
final version.

Declaration of Interests

The authors declare no competing interests.

Figures

Fig. 1: **World Magnetic Model (Epoch 2015.0) of total field intensity.** Turtle species, locations split by nesting regions and field intensity isolines ($\square$=1000nT) are shown. Abbreviated nesting regions: EC = Eastern Caribbean, I-P = Indo-Pacific, MED = Mediterranean, NEP = Northeast Pacific, N/WC = Northern/Western Caribbean, NWP = Northwest Pacific, SEA = Southeast Atlantic, SWA = Southwest Atlantic, SWIO = Southwest Indian Ocean.

Fig. 2: **Geomagnetic fields' representation.** Principle component analysis including eigenvectors (D = Declination, I = Inclination, H = Horizontal Component, F = Total Field Intensity, X = North Component, Y = East Component, Z = Vertical Component) illustrating the geomagnetic vectors for each region from all sea turtle species. (a) The present day (01/08/2015), (b) 50 years and (c) 100 years prior to the present day.

Fig. 3a & b: **Relationship between sea turtle population structure and the geomagnetic field.** The geomagnetic field is expressed as residuals of the correlation between the geomagnetic field and geographic distance. (a) Shows the overall relationship for all regions and (b) these relationships for geomagnetic regions, that offer low rates of navigational error, and for non-geomagnetic (high rates of navigational error) regions.

Fig. 4a & b: **Magnetic field vector intensity for geomagnetic and non-geomagnetic regions split by hemisphere.** Means and standard deviation depicted. Data collected at each change of 1 degree of latitude, across each region from the most northern sea turtle natal areas to the most southern. Each data point (isoline) was sampled over 24 years from 01/08/1991 to 01/08/2015 at 2-year intervals, i.e. the average time a sea turtle would take to mature. (a) Shows vertical field intensity and (b) shows total field intensity.

Fig. 5: **Estimated error rate in geomagnetic and non-geomagnetic regions**. Means and confidence intervals are depicted. Values are obtained for "geomagnetic" and "non-geomagnetic" regions from the estimation of distance if turtles confused two adjacent isolines for magnetic total field intensity and vertical field intensity.

| Loggerhead | Green | Hawksbill | Leatherback | Olive Ridley |

(a) Total field intensity error rate (km)

(b) Vertical field intensity error rate (km)

Regions grouped based on their relationship between
sea turtle population structure and geomagnetic difference