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## Mechanisms driving phenological and range change in migratory species

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*Does your article include research that required ethical approval or permits?:* This article does not present research with ethical considerations

*Statement (if applicable):* CUST\_IF\_YES\_ETHICS :No data available.

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Statement (if applicable):

All data are presented and accessible in the figures

#### Conflict of interest

I/We declare we have no competing interests

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## Authors' contributions

This paper has multiple authors and our individual contributions were as below

## Statement (if applicable):

All three authors conceived and designed the study and generated the data. JAG conducted the analyses and all three authors contributed to the writing of the manuscript.

1 2		
3	1	Mechanisms driving phenological and range change in migratory species
5	2	
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22	12	
23 24 25	13	Summary
26 27 28	14 15 16	Many migratory systems are changing rapidly in space and time, and these changes present challenges for conservation. Changes in local abundance and site occupancy across species' ranges have raised concerns over the efficacy of existing protected area networks, while
29 30 31 32 34 35 36 38 39	17 18 19	changes in phenology can potentially create mismatches in the timing of annual events with the availability of key resources. These changes could arise either through individuals shifting in space and time or through generational shifts in the frequency of individuals
	20 21 22	using different locations or on differing migratory schedules. Using a long-term study of a migratory shorebird in which individuals have been tracked through a period of range expansion and phenological change, we show that these changes occur through
	23 24 25	generational shifts in spatial and phenological distributions, and that individuals are highly consistent in space and time. Predictions of future rates of changes in range size and phenology, and their implications for species conservation, will require an understanding of
41 42 43 44	26 27 28	the processes that can drive generational shifts. We therefore explore the developmental, demographic and environmental processes that could influence generational shifts in phenology and distribution, and the studies that will be needed to distinguish among these
45	29	mechanisms of change.
47 48	30	
49 50	31	Keywords: avian, climate change, migration, mismatch
51 52	32	
53 54	33	Introduction
55 56	34	Migratory populations throughout the world are changing rapidly, with declines in
57 58	35 36	abundance being reported on all major flyways [1-3] and driving international calls for action [4]. Identifying the causes of population changes in migratory species is inherently
<b>5</b> 0	37	complex because of the range of sites and conditions experienced by individuals on their

annual journeys. Changing conditions in any or all parts of migratory ranges could drive changes in abundance and distribution. For example, changes in local conditions could influence local demography and thus alter relative abundances and site occupancy anywhere within a migratory range. However, the effects of local changes in one part of a range can also cascade through to influence abundance and distribution across a range [5,6]. This potential for interactions between local environmental conditions, demography, individual development and range-wide distribution and phenology make migratory systems complex, and designing conservation actions to halt and reverse declines in migratory species is therefore a major challenge. Recent changes in abundance of migratory species have been linked to their distribution and phenology. For example, phenological change is most commonly observed as shifts in the timing of migration to breeding grounds, and the magnitude of these shifts can varv greatly among species. Among European breeding birds, declines have been more frequent in species for which advances in spring arrival dates have not occurred [7], and species with non-overlapping breeding and wintering ranges are both more likely to be declining and less  $\frac{22}{23}$ likely to have shown advances in spring arrival [2]. A lack of advance in spring arrival dates can potentially increase the impact of any trophic mismatch resulting from climate-driven changes in the timing of availability of key food resources for breeding [8,9]. Declining species with little or no phenological change are often assumed to be constrained from responding to changing climatic conditions in breeding areas (for example because they migrate to more distant non-breeding locations), but the nature of any such constraint is unknown. Identifying the mechanisms through which phenological change occurs, and thus the factors constraining or facilitating these changes, may therefore be key to designing and targetting conservation actions to mitigate the effects of trophic mismatch. 35 36 Changes in the distribution of breeding and non-breeding ranges have also occurred in many migratory species, with polewards range shifts being particularly prevalent [10,11]. Concerns have consequently arisen over the efficacy of existing protected area networks [5,12]. Range change is often interpreted as a response to changes in the suitability of environmental conditions (eg colonisation of areas that were previously unsuitable and/or contraction from areas of declining suitability). However, the mechanisms driving changes in the distribution of individuals across a range are rarely known. Both range change and phenological change could arise through individual plasticity in use of space and time. For example, range change may occur through individual movement to locate suitable conditions, while changes in timing of migration could arise through individual decisions on departure timings or time spent on migratory journeys varying between years. By contrast, these changes could also result from generational shifts in the frequency of individuals that use different locations or migrate at specific times. Generational shifts would not require individual plasticity but would require changes in the conditions determining the frequency of individuals within a population with different phenologies or probabilities of occupying different parts of a range, such that the spatial and temporal distribution of recruits to the population would differ from their predecessors. For example, changes in distribution could arise through shifts in the conditions influencing 60

the probability of occupancy of different locations by new recruits, and/or by shifts in the survival rates achieved within different locations. Similarly, shifts in timing of migration could occur through changes in the conditions influencing adoption of migration schedules by new recruits (or survival rates associated with different schedules) altering the frequency of individuals on different schedules within a population [13]. Individual plasticity in spatial distribution and migratory timings would facilitate relatively rapid responses to changing environmental conditions. However, generational shifts would likely result in slower responses to environmental changes, particularly in long-lived species, as the direction and magnitude of the changes would depend on the proportion of annual recruits within a population and the proportion of those recruits experiencing changing drivers of use of space and time. If individual variation in migratory destination or timing has a genetic component, then generational shifts could drive microevolutionary change. However, genetic change is not an inevitable consequence of generational shifts, as individual destinations or timings could also be determined by environmental or social cues. Changes in range size or migration phenology of populations as a result of individual 25 plasticity in use of space and time has not been demonstrated, and a growing number of individual tracking studies are reporting high levels of repeatability in individual migratory destinations and timings [13-20]. If generational shifts are the more likely driver of population changes in space and time, then our focus should be on understanding drivers of settlement and phenology of recruits, as well as their subsequent survival and recruitment. Identifying the relative contributions of generational shifts and individual plasticity requires model systems in which individuals can be tracked across space and time, through periods of shifts in range and phenology. Such large-scale, long-term tracking data are rare but one system that provides all of these elements is the Icelandic black-tailed godwit, Limosa *limosa islandica*, which has been the subject of intensive individual and population studies since the mid-1990s [21-23]. In common with many migratory bird species at temperate and subarctic latitudes, advances in the phenology of spring migration have occurred in the Icelandic godwit population in recent decades [24]. Iceland supports very large breeding populations of several shorebird populations which migrate south to locations ranging from temperate Europe to sub-Saharan Africa [25]. The first recorded arrival dates of these species into south Iceland in spring have advanced over the last three decades, with godwits showing one of the most rapid advances (~2 weeks earlier now than in the 1990s; [24]). In addition, this godwit population has shown sustained increases in number for over a century, likely as a consequence of warming conditions in Iceland facilitating earlier and more successful breeding and recruitment [6,26,27]. This population growth has been accompanied by range expansion in both the breeding and non-breeding ranges; in Iceland, godwits have expanded from a breeding range that was confined to the south-west corner of the country around 1900, to occupy progressively more northerly and easterly locations [27]. In the non-breeding range, which spans coastal areas of north-west Europe from Britain and Ireland to Iberia and northern Morocco, colonisation and population increases have primarily occurred in the northern part of the range (east and north-west England, Scotland and east Ireland) since the 1970s, when surveys began [21,28]. 60

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3	122	In the mid-1990s, a programme of population-wide and life-long tracking of individual
4 5	123	godwits was initiated [21,29]. Across the breeding and non-breeding ranges, godwits have
6	124	been caught and marked with unique combinations of coloured leg-rings, and ~1-2% of the
8	125	population(which numbers ~ 50,000 individuals; [30]) is marked. A citizen science network
9	126 127	of >2000 observers across Europe regularly report marked individuals, allowing the locations of individuals to be repeatedly tracked within and across years [31]. Here, we use this range-
10	128	wide tracking of an expanding (in space) and advancing (in time) migratory shorebird, to
12	129	explore whether individual plasticity or generational shifts are likely to have caused these
14	130	changes in space and time, and to consider the evidence for potential developmental,
15 16	131	environmental and demographic drivers of such changes.
17	132	
18 19 20	133	Methods
21 22	134	Phenological change: repeated measures of individual spring arrival dates
23 24	135	Black-tailed godwits return to Iceland between mid-April and mid-May and, on arrival, flocks
25	136	congregate on a small number of arrival sites [32]. Since 1999, standardised surveys of
26	137	arrival sites in south and west Iceland have taken place from mid-April to early May, with
27 28 29	138 139	days [13,32]. Between 1999 and 2018, arrival dates were recorded in at least three years for
30	140	85 individuals (Fig. 1). In order to quantify (a) the trends in arrival dates of individuals, we
31 22	141	variable, and vear and individual as fixed effects (an extension of the model reported in Gill
33 24	143	et al. 2014 for a smaller sample). In order to then quantify (b) whether individual trends in
34 35	144	arrival dates have changed in magnitude or (c) whether the frequency of individuals with
36	145	differing arrival dates has changed over the 16-year survey period, we then constructed two
38	146	further GLMs with the (b) trend in arrival date and (c) mean arrival date for each individual
39 ₄0	147	individual was observed on arrival in Iceland) observation as a fixed effect.
41 42	140	,
43	149	
44 45	150	Range change: non-breeding locations of marked individuals
46	151	Locations of individually-marked godwits across the migratory range have been recorded by
47 48	152	anetwork of citizen scientists since the mid-1990s. Here we use all recorded non-breeding
49	153	locations (ie excluding records within Iceland) of 631 individuals marked at the main post-
50 E4	154	(mid-Octiomid-Eeb) records of the 419 of these individuals observed during that period. To
52 52	156	assess the role of individual movement in driving the northward range expansion we use
53 54	157	these sightings to quantify the total number of non-breeding locations (individual estuaries
52	158	and wetlands), regions and countries (Table 1) used by individuals tracked for differing
57	159	numbers of years. To quantify the contribution of generational shifts in the frequency of
58	160	individuals using different locations to the northwards range expansion, we then use these
59 60	וסו	signungs to assess whether individuals marked in more recent years (which will be younger,

on average, than previously marked individuals) are more likely to use recently colonised locations. We constructed generalised linear models with a binomial structure and a logit link, with the number of individuals marked in consecutive 5-year time periods since 1995 that winter in sites (a) colonised before or after the 1960s and (b) north and south of 52°N (most of the recently colonised sites are north of 52°N; [31,33]) as the dependent variable, and time period as a fixed effect. Results Mechanisms driving phenological change: individual plasticity or generational shifts Although first spring observations of godwits in South Iceland have advanced by more than two weeks in the last two decades [24], trends in arrival dates of 85 marked individuals over the last 16 years vary significantly among individuals but show no consistent advances in <u> 2</u>9 individual arrival date (Table 2a), and no change in individual arrival trends over the survey 74 period (Fig. 2a, Table 2b). However, the distribution of arrival dates of marked individuals has changed over time, with individuals first observed in more recent years tending to arrive earlier than individuals first recorded in the late 1990s/early 2000s (Fig. 2b, Table 2c). Thus, 28 while the arrival dates of individuals have not advanced, the frequency of early-arriving individuals has increased over this time period. 30 32 Mechanisms driving range change: individual plasticity or generational shifts Repeated tracking of individual godwits across the migratory range for up to 23 years 36 indicates a remarkably low level of individual plasticity in site use throughout their lives. On average, individuals are recorded on a total of only ~4 (±2.5 SE) non-breeding locations, regardless of the number of years for which they have been tracked (Fig. 3). In addition, these few locations are spread across, on average,  $2.5(\pm 1.5)$  regions in  $1.5(\pm 0.8)$  countries. Individuals are therefore highly restricted and consistent in their use of a small number of passage and winter locations, and these can be spread across the migratory range. However, over the two decades of continuous marking and tracking, with newly marked individuals being added to the population each year, the relative frequency of individuals using 'new' (colonised since the 1960s, Table 1) winter sites has increased (Fig. 4a), and this is primarily through increased numbers of individuals using more northerly sites (Fig. 4b, Table 1). 52 Discussion The spring arrival dates of godwits in Iceland has advanced [24] and the breeding population has expanded northwards in recent decades, with rapid population increases in more northerly non-breeding locations [21]. Repeated tracking of individuals in space and time over this period has shown that these expansions and advances are driven by generational shifts in the frequency of individuals occupying different locations and migrating at different 

times, and not by individual plasticity. Individuals show life-long consistency in use of a small number of non-breeding locations but the proportion of individuals occupying recently colonised sites is greater for more recently-ringed birds. As recently colonised sites are primarily (but not exclusively) in the north of the range, and as the network of observers recording individual locations has been in operation throughout the 20 year survey period. the recorded shift in distribution is unlikely to be influenced by changes in reporting rates across the range. The widely reported changes in phenology and distribution of many migratory species in 15 recent decades may therefore be likely to result from generational shifts in the frequency of individuals with differing phenologies and using different locations. Generational shifts have been shown to drive shifts in distribution in other migratory systems. For example, a shift in use of spring passage sites by a migratory wader population (continental black-tailed godwits, *Limosa limosa limosa*) was driven by new recruits to the population, while adults in 22 the population continued to use the site they had always previously occupied [33]. Quantifying the role of generational shifts requires long-term tracking of individuals, and relatively few studies have this information during periods of change. However, tracking studies are increasingly being conducted, and studies tracking individuals over multiple years are typically reporting high levels of repeatability of individual timings and destinations [13-20], suggesting that the benefits of philopatry in use of space and time are very strong [34,35]. Thus generational shifts may be the primary mechanism through which phenological shifts and range change occur. 31 32 Identifying the environmental and demographic processes that drive generational shifts in space and time is therefore likely to be a critical step in understanding population-level responses to environmental change and the associated implications for conservation. Generational shifts in distribution or phenology could potentially arise through processes occurring at the following points in early life: 39 1. Developmental drivers of generational shifts Conditions experienced at the natal stage, such as timing of hatching and/or conditions for growth and development, could influence the probability of those individuals undertaking different subsequent migratory routes and timings. For example, individuals hatched late in the season and/or with insufficient resources to fuel rapid growth are likely to migrate later, on average, and may thus have less time to locate more distant non-breeding locations. In such a case, an increase in the number of later-fledging individuals could drive recruitment into non-breeding locations that are closer to the breeding grounds. Similarly, natal conditions could potentially influence subsequent migration phenology, either directly through impacts on individual condition, or indirectly through impacts on the conditions and potential flockmates encountered during the non-breeding season. An important aspect of natal conditions that could potentially influence juvenile distribution and migratory timings is the changes in breeding phenology that are widely reported in many migratory populations [9,36]. Advances in nesting dates have been reported in many species and individual plasticity in nesting dates is common [37,38]. Thus, current evidence 60

1		
2 3	2/2	suggests that individual timings of arrival of migrants on breeding grounds tend to be
4	243	suggests that individual timings of an valid migrants on breeding grounds tend to be
5	244 245	consistent, but subsequent limings of breeding can vary greatly, and often vary in response to local weather conditions [6, 39, 40]. This changing phenology of natal conditions could
6 7	246	potentially drive changes in subsequent phenology and distribution of new recruits to the
8	247	population if timing of fledging influences subsequent migratory routes and timings. A key
9 10	248	mechanism that could drive such links is the potential for timing of fledging to influence the
11	249	likely flockmates on migratory journeys, and the destinations to which they are travelling.
12	250	For example, tracking of adult and juvenile Lesser Spotted Eagles, <i>Clanga pomarina</i> , on their
14	251	migratory journeys has shown that juveniles that departed at the same time as adults were
15	252	significantly more likely to take the same routes as adults, and to have higher subsequent
19	253	survival rates, than juveniles who departed without adults [41]. Timing-driven access to
18	254	social cues in migratory flocks could therefore be an important driver of the migratory
19	255	routes and destinations located by juveniles, and changes in timing of fledging could
20 21	256	therefore drive changes in the non-breeding distribution of migratory species.
22	257	2. Environmental effects on generational shifts
25 74	200	2. Environmental energies on generational shifts
25	209	
26	260	migrating at specific times or locating specific non-breeding locations. For example, changing weather conditions (e.g. windspeeds or directions) could alter the proportions of
27 28	262	recruits migrating at different time or taking different routes. However, while wind
29 20	263	conditions can have important effects on migrating birds [42] individual consistency in
30 31	264	migratory destinations and timings would mean that such effects could only operate in early
32	265	life (ie during settlement/recruitment).
33 34	266	
35	267	3. Demographic effects on generational shifts
$\frac{36}{37}$	268	Disproportionate changes in survival rates of recruits that differ in distribution or timing
38	269	could lead to generational shifts. For example, while the numbers of individuals recruiting
39	270	into more northerly winter locations or arriving early on the breeding grounds may not be
40	271	changing, those individuals could be increasingly likely to survive, for example as a
<b>41</b>	272	consequence of ameliorating weather conditions in northerly areas or on arrival in the
43 11	273 274	range expansion into areas in which weather conditions are changing as a consequence of
77 15	275	climatic change [43] and these effects could operate alongside developmental or
46 ⊿7	276	environmental effects
47	277	
49	278	Drivers of generational shifts in Icelandic godwits
50 51	279	In Icelandic black-tailed godwits, rapid warming has occurred on the breeding grounds in
51 52	280	recent decades, and nesting dates are earlier [6] and productivity is higher [26] in warmer
53	281	years. This warming-driven increase in productivity is likely to have fuelled the colonisation
54 55	282	of colder breeding areas in the north, where nest-laying and hatching dates are, on average.
56	283	later than in more southerly breeding areas [6]. Individuals from these colder and more
58	284	recently colonised breeding areas are more likely to also winter in the more recently
59	285	colonised non-breeding areas [31], and these seasonal links could thus result from regional-
60		

1 2		
3	286	scale differences in timing of fledging and subsequent social cues available to juveniles
4 5	287	undertaking their firstmigration.
6 7	288	Were in a driven advence in pacting dates as uld also have driven the advencing environ
8	289	warming-driven advances in nesting dates could also have driven the advancing spring
9 10	290 291	in more southerly areas of Europe [29] and breeding in the warmer areas of south and west
10	292	Iceland [32] arrive first, and (b) more recently hatched individuals tend to have earlier spring
13	293	$arrival dates than individuals hatched in the 1990s \cite{13}. This suggests that the generational$
14 15	294	shifts driving the phenological advance in this system (Fig. 2) could potentially result from
15	295	Increased numbers of early-flatched individuals from the traditionally occupied areas of
17 18	290	and social cues to both locate traditionally occupied winter areas and return early in spring
19	298	Increased survival rates of early-arrivers and northerly winterers could also be contributing
<del>3</del> 9	299	to these changes in space and time.
22	300	
23 74	301	Climate change and generational shifts in space and time
25	30Z	In climate-driven smits in breeding prenology can alter the frequency of juvernies
26 27	303 304	route through which climate-associated shifts in range and phenology are manifest. A
28	305	common pattern among migratory species at present is those migrating longer distances are
30	306	less likely to show shifts in migration phenology [44,45]. As longer-distance migrants
31	307 308	typically arrive later on the breeding grounds and have a smaller gap between arrival and laving than shorter-distance migrants [13], they may have a more limited capacity to alter
33	309	breeding phenology (and thus generational shifts resulting from shifts in breeding
34 35	310	phenology are less likely to occur). The effects of climate warming on breeding phenology
36	311	can thus have potentially far-reaching consequences for migratory populations.
38	312	<i>_ ,</i>
39 ⊿0	313 314	<i>Future research</i>
40	315	generational shifts and the conditions in which each might be most relevant will require
42 43	316	studies in which the effects of natal conditions, environmental conditions experienced by
44	317	juvenile individuals undertaking different migratory routes and timings, and the
45	318	demographic consequences of conditions experienced at destinations can be measured.
47 48	319 320	I racking individuals from fledging to recruitment into adulthood is difficult because survival rates at this life stage are typically low, and retrieval of tags can be challenging because the
40 49	321	subsequent breeding locations of these individuals is typically unknown, but technological
50 51	322	advances will hopefully make these issues more tractable in the near future. Long-term
52	323	studies of seasonal patterns of nest loss, timing of replacement clutches and juvenile
53 54	324	fledging will also be particularly valuable in identifying the potential role of breeding
55 54	325 326	phenology in driving change in migratory systems. Quantifying the developmental, environmental and demographic processes that influence individual migratory routes
50 57	327	destinations and timings will be key to understanding future rates and directions of spatial
58 59	328	and phenological change in migratory species, and the associated implications for designing
60	329	effective protected area networks and conservation actions for these species.

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45 67 8 9 10 1 <u>3</u> 14	331	Ackr	nowledgements			
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15 16 17	338	Com	peting interests			
17	339	Weł	nave no competing interests			
19 20	340	Ethics				
21 22	341	All re	All required approval and licenses were obtained			
23 24 25	342	Auth	Author contributions			
25 26 27 28	343 344	All three authors conceived and designed the study and generated the data. JAG conducted the analyses and all three authors contributed to the writing of the manuscript.				
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529	Table 1. The 121 winter locations across 26 regions and nine countries	sused by the
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530 individual godwits shown in Figure 3. Regions in bold are colonised since the 1960s.

0				
7 531	Country	Region	Lat-Long	No. of locations
8	N Ireland	East	54°N, 05°W	1
10	Ireland	West	53°N, 08°W	1
11		Central	52°N, 08°W	2
12		East	53°N, 06°W	6
14		South	51°N, 08°W	11
15	Wales	North	53°N. 03°W	2
16 17		West	52°N. 04°W	1
17	England	North-west	53°N, 03°W	9
19	5	Central	52°N 01°W	2
20		East	52°N 01°F	27
21		South	50°N 01°W	10
23		South-east	51°N 01°E	3
24		South-west	50°N 03°W	7
25 26	Netherlands	North	53°N 06°E	1
27	Nethenands	Contral	53 N, 00 L	2
28		Weet	52 N, UO E	2
29	<b>F</b>	VVest	52 N, 04 E	3
30	France	North	48°N, 01°W	3
32		North-west	47°N, 02°W	1
33		West	46°N, 01°W	10
34	Portugal	South	37°N, 08°W	4
35		West	38°N, 09°W	3
30 37	Spain	North	43°N. 03°W	1
38	·	North-west	42°N. 08°W	1
39		West	38°N 06°W	1
40 41		South	37°N 06°W	2
42	Morocco	West	30°N 00°N/	- 1
12			00 14, 00 14	•

#### Table 2. Results of GLMs of (a) annual and individual variation in arrival dates of 85 god wits(3-12 years between 1999 and 2018) and variation in (b) annual trends in arrival dates and (c) mean arrival dates, in relation to the year in which those 85 individuals were first

observed on arrival in Iceland. 

	d.f.	F	p	Slope (±SE)
(a) Year Individual Error	1 84 300	0.99 2.06	0.32 <b>0.001</b>	0.062 ±0.06
(b) First observation year Error	1 83	0.02	0.89	0.004 ±0.03
(c) First observation year Error	1 83	8.83	0.004	-0.21 ± 0.07

3

# 544 Table 3. Results of binomial models of variation through consecutive time intervals in the

545 frequency of individually-marked godwits recorded in winter locations (a) occupied since 546 the 1960s (old, **0**) or colonised since then (new, 1), and (b) north (0) and south (1) of 52°N.

-/ ~	Estimat	e SE	р	Odds ratio
(a)				
Intercept	0.4	0.25	0.001	0.46
rear	-0.77	0.13	0.001	0.40
(b)				
tercept	-0.008	0.23		
ear	-0.29	0.11	0.006	0.74

2 3 4	552	Figure Legends
5 67 8 9 10 11 12 13 14 15	553	
	554 555	Figure 1. Number of years on which 85 individually-marked Icelandic godwits have been recorded on spring arrival in Iceland, between 1999 and 2018
	556	
	557 558 559	Figure 2. Changes in the mean (a) annual change in arrival date and (b) arrival date (days since 1 April) in Iceland of 85 individual godwits with the year in which they were first observed on arrival in Iceland (see Table 2c for model details).
17	560	
18 19 20 21 22 23 24	561 562 563 564	Figure 3. The total number of (a) countries, (b) regions and (c) locations on which individual godwits have ever been recorded in the total number of years over which each has been tracked. The number of individuals tracked for each total number of years is shown above the bars in (a), see Table 1 for numbers of locations, regions and countries.
25 26	565	
26   27   28   29   30   31   32   33   34   35   36   37   38   39   40   41   42   43   44   45   46	566 567	Figure 4. Changes through consecutive 5-year time intervals in the (a) proportion of individual godwits wintering in locations that have been occupied since the 1960s (old) or
	568 569	colonised since then (new), and (b) latitude of those winter locations. Numbers observed in each time period are shown in (a).







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