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The political biogeography of migratory marine predators

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The political biogeography of migratory marine predators

48 During their migrations marine predators experience varying levels of protection and face 49 many threats as they travel through multiple countries' jurisdictions and across ocean 50 basins. Some populations are declining rapidly. Contributing to declines is a failure of 51 international agreements to ensure effective cooperation by the stakeholders responsible 52 for managing species throughout their ranges, including in the high seas, a global 53 commons. Here we use biologging data from marine predators to provide quantitative 54 measures with great potential to inform local, national, and international management 55 efforts in the Pacific Ocean. We synthesized a large tracking dataset to show how the 56 movements and migratory phenology of 1,648 individuals representing 14 species—from 57 leatherback turtles to white sharks-relate to geopolitical boundaries of the Pacific Ocean 58 throughout species' annual cycles. Cumulatively, these species visited 86% of Pacific Ocean 59 countries and some spent three quarters of their annual cycles in the high seas. With our 60 results, we offer answers to questions posed when designing international strategies for 61 managing migratory species.

Marine migrations can span ocean basins and are dynamic in space and time¹. Migratory species 62 are thus exposed to a variety of threats² as they travel through multiple countries' jurisdictions 63 64 and the open ocean. As a result, numerous migratory marine species from diverse taxa have 65 experienced recent drastic population declines including leatherback turtles (Dermochelys *coriacea*)³, Pacific bluefin tuna (*Thunnus orientalis*)⁴, and some sharks⁵ and seabirds⁶. Under 66 current management frameworks, migratory species have received varying levels of protection 67 and many gaps remain⁷⁻¹⁰. National rights over marine resources are delineated by Exclusive 68 69 Economic Zones (EEZs) which include waters out to 200 nautical miles from a country's

shoreline¹¹. Areas beyond national jurisdiction, the 'high seas', are legally recognized as a global 70 71 commons. Regional Fisheries Management Organizations (RFMOs) are the primary multijurisdictional mechanism for managing transboundary and high seas fish stocks¹². In a joint 72 73 management structure with member states, conservation and management rules are adopted by 74 the RFMO while enforcement of these measures falls to individual countries. As such, individual 75 nations are responsible for fishing and non-fishing related threats within their EEZs and, through 76 their high seas fleets and flag vessels, share responsibility beyond their EEZs. To recover 77 populations and to prevent declines of healthy populations, improved management and effective international cooperation and governance⁷ are urgently needed. Key information needs at all 78 79 levels include quantitative measures to indicate who has management jurisdiction over migratory 80 species across their range and at different times during their migratory cycle, including for 81 breeding, foraging, and migrating. Here we use biologging data to provide this information. We 82 show how the migratory cycles of populations of 14 species relate to geopolitical boundaries of 83 the Pacific Ocean using a subset of a large tracking dataset collected between 2000-2009 by the Tagging of Pacific Predators project (TOPP)¹ For each species of tuna (Pacific bluefin, 84 85 yellowfin, Thunnus albacares; albacore, Thunnus alalunga); shark (blue, Prionace glauca, 86 shortfin mako, Isurus oxyrinchus; white, Carcharodon carcharias; salmon, Lamna ditropis), 87 pinniped (northern elephant seal, Mirounga angustirostris; California sea lion, Zalophus 88 californianus), seabird (Laysan albatross, *Phoebastria immutabilis*; black-footed albatross, 89 Phoebastria nigripes; sooty shearwater, Puffinus griseus), sea turtle (leatherback), and cetacean 90 (blue whale, Balaenoptera musculus) we asked: 1) Which EEZs were visited? 2) What 91 proportion of time was spent in each EEZ and the high seas? 3) When during their migratory 92 cycle were animals within each EEZ or the high seas?

93 **Results and Discussion**

94 Individual animals (n=1,648) representing 265,881 tracking days (Supplementary Table 1) 95 visited 63 Pacific Ocean EEZs (Fig. 1) under the jurisdiction of 37 countries (some sovereignties 96 are disputed; disjunct EEZs for a given country were treated separately, Supplementary Table 2). 97 Some species (Pacific bluefin tuna, leatherback turtle, sooty shearwater, Laysan albatross) 98 travelled across the Pacific and all species entered numerous jurisdictions. The high seas were 99 visited by 48% (n=797) of individuals. Tag deployments occurred primarily in the eastern Pacific 100 Ocean and over 83% of daily locations were either in Mexico (31%), the high seas (29%), or the 101 United States (23%); 71% of all locations were within the boundaries of an EEZ (Supplementary 102 Table 3). 103 While these simple statistics provide insight into overall occurrence, they may be biased by 104 effects of deployment location and sampling imbalances common to electronic tracking datasets.

105 We addressed biases due to variability in sample size during the year (Supplementary Table 1,

106 Supplementary Fig. 5), deployment dates (Supplementary Fig. 6), and track durations

107 (Supplementary Figs. 7-8) using multinomial generalized additive models^{13,14}. We predicted

108 seasonal patterns of occurrence within specific countries and the high seas for multiple taxa (Fig.

109 2), breeding populations (Fig. 3 and Supplementary Fig. 1), life history stages (Supplementary

Fig. 2), and years (Supplementary Figs. 3-4). California sea lions and Yellowfin tuna were not

111 modeled because greater than 90% of locations were within a single EEZ (Supplementary Table

112 3). From model predictions we also estimated the percentage of an annual cycle spent in EEZs or

113 in the high seas (Table 1). The TOPP project was unprecedented in producing a large multi-

114 species, multi-year simultaneous animal movement dataset at an ocean-basin scale. Nonetheless,

115 many datasets are not fully representative at the species level within the Pacific Ocean. TOPP

116 focused primarily on North American populations and many species datasets are age or sex-117 biased. For example, this study includes results from female salmon sharks in the eastern North 118 Pacific, but not males from the western North Pacific. It includes female Northern elephant seals 119 from Mexican and American rookeries, but not males. Additionally, some species in this study 120 include few individuals relative to population size (for example, Sooty Shearwaters). Our results 121 therefore describe only the specific geographic subsets of populations and life history stages studied by TOPP (see Methods, Supplementary Information and ¹ for full dataset details and 122 123 deployment locations).

Using our results, we offer examples of scientific answers to key questions posed whendesigning international strategies for managing migratory marine species.

126 When during the year are marine predators present within countries' waters?

127 Marine predators cue on shifts of habitats and prey, which in turn concentrate individuals in specific regions during defined time periods¹. Consequently, residency within each EEZ is not 128 129 equally probable throughout the year (Fig. 2). It may be highly punctuated in time, for example the central Pacific island migration corridor of fast-moving sooty shearwaters¹⁵ (Fig. 2). Or, a 130 131 single EEZ may constitute half or more of yearly residency, for example, salmon sharks in 132 Alaska and Pacific bluefin tuna in Mexico (Table 1). Some populations in this study remained 133 almost entirely within the EEZs in which tag deployments occurred, making management more 134 straightforward. For example, California sea lions from the U.S. breeding population remained within U.S. waters except during years of anomalous oceanographic conditions¹⁶ when they 135 136 ventured to the high seas (Supplementary Table 2). Some life history stages not represented in 137 our dataset also remain in one or two EEZs, for example juvenile white sharks in the eastern north Pacific remain in USA and Mexican EEZs¹⁷. 138

139 Among the six taxa of marine predators studied, some co-occurred seasonally within the same 140 EEZs (Fig. 2). Tunas, sharks, and whales occurred within U.S. waters from July to December; 141 female elephant seals, albatrosses, and leatherback turtles ranged throughout the high seas from 142 April to November; and Laysan albatrosses and sooty shearwaters visited Russian waters from 143 July through October. There are examples of similar patterns from other stocks and populations 144 in the Pacific. White shark data modeled here represent individuals migrating between the U.S. 145 EEZ (Central California) and the high seas. A second group of northeastern Pacific white sharks 146 shows near identical phenology in migrations between the high seas and Guadalupe Island, Mexico¹⁸. By identifying seasonal patterns of co-occurrence across guilds, species, and 147 148 populations, our results can help managers maximize their efforts across a range of migratory 149 taxa. For example, dynamic and ecosystem-based management approaches require a synthetic 150 understanding of the migratory cycles of multiple species. Our results also could help identify 151 when and where to focus management efforts focused on human interactions, for example to 152 help maximize bycatch mitigation efforts in places where currently there is a lack of observer 153 coverage and enforcement.

Which countries should be cooperating, either directly, or through established international bodies and frameworks?

We identified the set of countries visited by each species (Fig 1., Supplementary Tables 1-2) and predicted when during the year animals moved among countries or into the high seas according to their cycles of breeding, foraging, and migration (Fig. 2-3, Supplementary Figs. 1-4).
Understanding the political biogeography of leatherback turtles in the Pacific Ocean is especially

160 important as they are highly threatened¹⁹ and their management is jurisdictionally complex.

161 During this study, leatherback turtles in the Pacific Ocean moved through 32 countries and the

162 high seas. Globally, seven leatherback turtle subpopulations are recognized and all are 163 considered vulnerable to extinction. However, the western Pacific and eastern Pacific 164 subpopulations we studied are critically endangered with estimates of a 96% population decline by 2040¹⁹. We compared Eastern (Fig. 2) and Western Pacific leatherbacks (Fig. 3), and breeding 165 166 populations of Western Pacific leatherbacks (Fig. 3). In the Western Pacific, turtles that breed in 167 the austral winter pass through Asian and Central Pacific EEZs; turtles that breed in the austral 168 summer migrate to EEZs of the South Pacific (Fig. 3). We show that political biogeography is 169 linked to population structure and breeding phenology for this species and our results thus 170 provide the ability to link observed locations of human interactions to specific leatherback turtle 171 breeding populations. To save leatherback turtles from extinction in the Pacific Ocean, a multi-172 lateral, cooperative approach is the only way forward, often stemming from private, local, or 173 regional collaborations that provide a first step in cooperative research and conservation. 174 Examples of such international coordination include the Inter-American Tropical Tuna 175 Commission, the Commission for the Conservation of Antarctic Marine Living Resources, the 176 Northern Fur Sea Treaty, and the Agreement on the Conservation of Albatrosses and Petrels. Our results could be a key ingredient in a holistic conservation strategy²⁰ that integrates protections 177 178 throughout the pan-Pacific leatherback life cycle including: optimizing reproductive success on 179 nesting beaches (e.g. beach protection, monitoring, and enforcement, conservation payments to 180 local communities), and preventing deaths due to incidental catch by fisheries within EEZs (e.g. 181 tailored approaches to scale of fishery and socio-economic context, adoption of gear-technology-182 handling standards to reduce incidental catch and increase probability of post-release survival, 183 incentive-based mechanisms, use rights, time-area-closures) and in the high seas (e.g. expanding

pan-Pacific policy actions, increasing and enforcing observer coverage, adoption of gear technology-handling standards, etc.)²¹.

186

187 How important are the high seas to marine predator populations?

The high seas are one of the world's last global commons²² and are among the least protected 188 places on Earth²³. Despite recent progress, many RFMOs have not ensured that all fish stocks 189 under their mandates are fished sustainably⁹ and/or have not suitably protected non-target species 190 such as seabirds, sharks, turtles, and marine mammals¹⁰. Many approaches have been suggested 191 192 or used to improve the sustainability of high seas fisheries, including: rights-based management, 193 adopting and enforcing best practice gear technology standards, increasing observer coverage, 194 time/area restrictions, protected areas, vessel monitoring, increasing and sharing scientific 195 research, market and trade-based mechanisms, and the adoption of a new international legal instrument.^{7,10}. To implement many of these suggestions, quantitative measures of high seas use 196 197 are needed.

198 Our results provide measures of the time multiple populations spend within the high seas at a 199 basin-wide scale (Table 1 and Fig. 2). For example, Pacific bluefin tuna tracked during their 200 trans-Pacific migration (n=12, Supplementary Fig. 2), and seabirds, leatherback turtles, white 201 sharks, and northern elephant seals spent between 45-75% of the year in the high seas (Table 1). 202 Attention to high seas management issues is increasing. The United Nations General Assembly 203 in 2015 resolved to develop an international legally binding instrument for the conservation and sustainable use of marine biodiversity of the high seas²⁴. This process will advance in 2018 to 204 205 full negotiation. The knowledge we present of how and when animals use the high seas is a 206 critical contribution to these next steps to sustain marine biodiversity and is a complement to

207 new satellite services (for example, AIS, maritime Automated Identification System made
208 publicly available through Global Fishing Watch²⁵) that can be used to increase the transparency
209 of high seas fishing.

210 Conclusions

211 Our analysis of tracking data collected from Pacific predators describes seasonal patterns 212 of national and international management jurisdiction over migratory species. Actions to protect 213 marine migratory species are needed throughout their range, including on the high seas. Multiple international conventions and agreements^{11,12,26} seek to promote cooperation within and beyond 214 215 national jurisdictions for managing migratory species. Nevertheless, scaled-up international 216 collaboration and effective governance are essential. While our results demonstrate the 217 jurisdictional complexity of managing some critically endangered highly migratory species like 218 Pacific leatherback turtles, they also demonstrate that for some species or populations, 219 agreements between just a few countries could help reverse declines. Our approach capitalizes on what biologging technologies do best²⁷: provide continuous movement data on individual 220 221 animals who spend most of their lives away from direct scientific observation. This information can^{28-30} and should be used to inform management. 222



232 Fig. 1. Daily locations of marine predators electronically tracked within EEZs and the high seas of the Pacific Ocean. (a) State space modeled daily locations of 14 marine predator species 233 234 electronically tracked 2000-2009 in EEZs (transparent overlay) and the high seas (ocean water 235 falling outside transparent overlay). Modified from¹. (b) Key to visited EEZs. EEZ boundaries 236 from VLIZ Maritime Boundaries Geodatabase (2016); some are disputed. Refer to official record 237 for all claimants and alternative geographies. 1 Alaska (USA); 2 Canada; 3 United States of 238 America; 4 Mexico; 5 Clipperton Island (France); 6 Guatemala; 7 El Salvador; 8 Nicaragua; 9 239 Costa Rica: 10 Galapagos Islands (Ecuador): 11 Panama; 12 Peru; 13 Desventuradas Islands 240 (Chile); 14 Chile (includes Juan Fernandez Islands); 15 Easter Island (Chile); 16 Pitcairn (UK); 17 241 French Polynesia (France); 18 Line Islands Group (Kiribati); 19 Jarvis Island (USA); 20 Palmyra 242 Atoll (USA); 21 Johnston Atoll (USA); 22 Hawaii (USA); 23 Wake Island (USA); 24 Marshall 243 Islands; 25 Nauru; 26 Kiribati; 27 Tuvalu; 28 Howland Island and Baker Island (USA); 29 Phoenix 244 Islands Group (Kiribati); 30 Tokelau (New Zealand); 31 Wallis and Futuna (France); 32 American 245 Samoa (USA); 33 Niue (New Zealand); 34 Cook Islands (New Zealand); 35 Samoa; 36 Tonga; 37

- Fiji; 38 Norfolk Island (AUS); 39 New Zealand; 40 Macquarie Island (AUS); 41 Antarctica; 42
- Australia; 43 New Caledonia (France); 44 Vanuatu; 45 Solomon Islands; 46 Papua New Guinea;
- 248 47 Indonesia; 48 Brunei; 49 Malaysia; 50 Micronesia; 51 Palau; 52 Philippines; 53 Spratly Islands
- 249 (Disputed); 54 Viet Nam; 55 Paracel Islands (Disputed); 56 Taiwan; 57 Northern Mariana Islands
- and Guam (USA); 58 Japan; 59 China; 60 South Korea; 61 Japan-Korea Joint Development Zone;
- 251 62 Southern Kuriles (Disputed); 63 Russia.



253 Fig. 2. Seasonal probability of marine predator occurrence in Pacific Ocean EEZs and the 254 high seas from electronic tracking. Lines represent the estimated effect of day of the year on the 255 probability of a randomly selected individual from the tracked population occurring in each region. 256 Shading represents the inter-quartile range of estimates simulated from the posterior distribution of the 257 model parameters. Total sample size for each population is represented by n; tracking duration varied 258 among individuals (see Methods). Model details: Methods and Supplementary Table 4. Leatherback 259 turtles have a multi-year migratory cycle; estimates begin on January 21 and continue through the first 260 year of this cycle following breeding/tag deployment.





Fig. 3. Population differences in Western Pacific leatherback turtle use of EEZs and the high seas. (A, C, E) Probability of turtle occurrence in EEZs and the high seas during the 7-9 months following tag deployment. Tags were deployed during the breeding period (A-D), or on foraging

267 grounds (E-F). Lines represent the estimated effect of days elapsed after tag deployment on the 268 probability of a randomly selected individual from the tracked population occurring in each region. 269 Shading represents the inter-quartile range of estimates simulated from the posterior distribution of the 270 model parameters. Sample size and longevity of tags dictated the number of days elapsed since tagging 271 (7-9 months) considered in the models. Model details: Methods and Supplementary Table 4. (B, D, F) 272 State space modeled daily locations of individuals electronically tracked from each population (colors 273 correspond to EEZ where tags were deployed) and EEZ boundaries (colors correspond to facing panel). 274 (A-B) Austral summer breeding populations tagged during breeding in Indonesia, Solomon Islands, and 275 Papua New Guinea (C-D) Austral winter breeding population tagged during breeding in Indonesia, (E-276 F) Foraging population in California Current Large Marine Ecosystem.

278 Table 1: Percentage of the year marine predators are estimated to spend within Pacific Ocean 279 EEZs and the high seas. Percentages were estimated from model results presented in Fig. 2 and 280 Supplementary Fig. 2 (PBT, TP) using electronic tagging data. Model details: Methods and 281 Supplementary Table 4. Species codes: PBT, Pacific bluefin tuna (all individuals); PBT (TP), trans-282 Pacific migrants (see Supplementary Fig. 2); AT, albacore tuna; WS, white shark; MS, mako shark; BS, 283 blue shark; SS, salmon shark; NELE, female northern elephant seal; BFAL, black-footed albatross; 284 LAAL, Laysan albatross; SOSH, sooty shearwater; LET(CR), leatherback turtle from Costa Rica; 285 BLWH, blue whale. Percentages may not total 100 due to rounding. Leatherback turtles have a multi-286 year migratory cycle; estimates are for the first year of this cycle following breeding/tag deployment. 287 Uncertainty in these estimates and estimates for additional data subsets are presented in Supplementary 288 Table 5.

	PBT	PBT	AT	WS	MS	BS	SS	NELE	BFAL	LAAL	SOSH	LET	BLWH
		(TP)										(CR)	
High Seas	0.2	44.9	12.2	62.5	3.7	0.8	23.7	66.6	66.7	74.5	65.9	78.2	30.3
USA	28.7	25.7	27.6	37.4	46.8	55.9	3.2	33.1	7.7		<1		55.8
Mexico	71.1	28.3	60.2	<1	49.5	35.7	<1				<1		13.2
Alaska							70.0	0.1	1.0	4.5	<1		
Canada						<1	3.1	0.3	2.7				
Hawaii	<1		<1	<1	<1				21.6	17.5			
Russia										3.2	4.2		
Central Pac. Isl.											3.8		
Japan	<1	1.1									4.0		
New Zealand											20.7		
Costa Rica												17.8	
Galapagos												3.7	

Other	<1	<1		<	1		<1	<1	<1	<1	<1
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292 Methods:

293 *1. Data summary and availability*

From 2000 to 2009, researchers with the Tagging of Pacific Predators (TOPP) project deployed 4,306 electronic tags which provided 1,791 individual animal tracks from populations of 23 species in the Pacific Ocean¹. Animal research was conducted in accordance with institutional animal care and use protocols from Stanford University and the University of California, for endangered species in accordance with guidance from the U.S. Endangered Species Act and for marine mammals in accordance with the U.S. Marine Mammal Protection Act.

A Bayesian state-space model³¹ was fitted to the tag data to derive regular, daily mean estimates of locations at sea while accounting for tag observation error³². The state-space model also provided estimates of the uncertainty in the location estimates. This modeled TOPP dataset is archived in the U.S. Animal Tracking Network Data Assembly Center.

We used a subset of this TOPP dataset; only species tracked over multiple years were included. The dataset we analyzed included 14 species, 1,648 individuals and 265,881 modeled daily locations (Supplementary Table 1). Yearly sampling effort varied (Supplementary Table 1). Tags were deployed within the boundaries of eight EEZs (Supplementary Tables 2-3). For full deployment details:¹. There are multiple populations in the Pacific Ocean of many species considered here---we refer only to the specific populations and life history stages in the TOPP dataset¹.

311 *Variability in deployment date and track duration*: Timing of tag deployments was multi-modal
312 for some species (Supplementary Fig. 6), and track duration varied among individuals

(Supplementary Fig. 7). This variability in a tracking dataset can affect spatio-temporal analyseswhen deployment dates and track duration are unrelated to species life history.

315 Pinnipeds and seabirds: Distributions of deployment dates and track durations reflected 316 these species' life histories. To capture the full annual cycle of land-breeding and moulting 317 species, tags were deployed multiple times in a given year (northern elephant seals, prior to the 318 short post-breeding and long post-moult migrations; seabirds: prior to the short breeding and 319 long post-breeding migrations). Typically, unique sets of individuals were tracked during each 320 migration although some seals were tracked during both migrations in a given year, or during the 321 same migration in multiple years. California sea lions were predominantly tagged while nursing 322 pups to facilitate tag recovery; most individuals were tracked only during the breeding period. In 323 general, tag failure was rare and tags were recovered upon recapture of the animals. Pinnipeds 324 included in this analysis were females and all pinnipeds and seabirds were adults.

325 *Tunas, sharks, whales*: For these species, deployment timing varied among years partially 326 because of reasons unrelated to species' life history (sampling design considerations or cruise 327 availability). Primary tagging months were: Pacific bluefin tuna, January, March, July-328 September, November-December; yellowfin tuna, February, August, October-December; 329 shortfin mako shark June-August, November; blue shark, January-February, June-August, 330 October-December; salmon shark, July-August; and white shark, January and December. 331 Because of a higher frequency of tag failure and the difficulty of targeted recapture, the 332 distributions of track durations for these species (Supplementary Fig. 7) were a function of tag 333 attrition and harvest recapture. In general, these datasets contained a high number of individuals 334 tracked for less than a year. Salmon shark was an exception with 16 individuals tracked for at 335 least two years (Supplementary Fig. 7). Salmon sharks studied were all females; white sharks

were large adults and subadults; Pacific bluefin tuna, albacore tuna, shortfin mako shark, andblue sharks were largely juveniles

338 *Leatherback turtles:* Leatherback turtles have a multi-year migratory cycle and all tags 339 experienced attrition before recording the full multi-year migration. The eastern Pacific and 340 western Pacific subpopulations of leatherback turtles were considered separately in this study. Tags on eastern Pacific leatherbacks were deployed in January during the nesting period³³. 341 342 Tracking of the western Pacific subpopulation included both summer (Indonesia) and winter 343 breeders (Indonesia, Papua New Guinea, Solomon Islands), and animals captured while foraging in the California Current. Tag deployment thus occurred in three different pulses (see ³⁴ for more 344 345 detail on the multi-year migratory cycles of Western Pacific leatherbacks and their population 346 dynamics among breeding rookeries). All turtles included here were breeding adults.

347

348

8 2. Location Classification

349 Global EEZ boundaries were obtained as shapefiles from the VLIZ Maritime Boundaries 350 Geodatabase (v.8, 2014). Some EEZ boundaries between countries are disputed; full details of 351 boundary delineation are available (VLIZ: http://www.vliz.be/vmdcdata/marbound/). Shapefiles 352 were converted to polygon vectors using the MATLAB mapping toolbox (The MathWorks Inc., 353 R2015b). We developed a custom script based upon MATLAB's "inpolygon" function to 354 classify each location as present or absent (binary, ones and zeros) in each EEZ of the Pacific 355 basin. We classified locations on an EEZ boundary as inside the EEZ (and thus, those few 356 locations located exactly on the boundary/edge of two EEZs would be classified as within two 357 EEZs). If a location was neither on land nor in an EEZ, we classified it as a high seas location.

358 Disjunct EEZs for a given country were treated separately. For example, Hawaii and Alaska
359 were each treated as unique to the mainland USA EEZ.

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3. EEZ and High Seas Occurrence and models of seasonal use

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363 For each species we calculated the percentage of individuals (Supplementary Table 2) and the 364 percentage of daily locations (Supplementary Table 3) spent in each EEZ and the high seas. We 365 began our analysis exploring the proportion of time spent by individuals of each species in each EEZ and the high seas, a value most often reported in related literature (for example, see ³⁵). 366 367 However, for our dataset these simple statistical summaries of EEZ use were biased in the 368 following ways: 1) statistical summaries of individual EEZ use calculated from data sets with 369 high tag attrition (and thus a high number of abbreviated tracks: tunas, sharks, whales, and 370 turtles) were biased toward the EEZs in which tags were deployed (Supplementary Figs. 7-8); 2) 371 deployment date affected interpretation of EEZ use (Supplementary Figs. 6 and 8), and 3) yearly 372 summaries were affected by variability in sample size (Supplementary Table 1 and 373 Supplementary Fig. 5) and deployment dates (Supplementary Fig. 6) across years. One solution 374 is to remove from analysis individuals with abbreviated tracking durations (for example <30375 days or <1 year). However, we felt there was value in retaining all available information to elucidate seasonal patterns of EEZ use. 376

We therefore took a modeling approach to better estimate seasonal EEZ and high seas occurrence for the tagged population given the effects of individual variability in track duration and tagging location and date. The presence of an individual from the tagged sample in each EEZ or the high seas was modeled with a generalized additive model¹⁴, specifically a

multinomial logistic regression¹³. For species predicted to have an annual migratory cycle (all 381 382 species except leatherback turtles), a cyclic effect was used for day of the year enforcing 383 continuity in the estimated probabilities from year to year. Individual identity was treated as a 384 random intercept effect to account for differences in behavior and sample size throughout the 385 year among tagged individuals. Both day-of-year and individual effects were allowed to vary 386 across EEZs and the high seas. The multinomial model structure ensured that the probabilities of 387 presence in EEZs and the high seas summed to 1 for any given day of the year for any given individual. Models were fitted using the 'mgcv' package¹⁴ in R (R Development Core Team 388 389 2016). The ordering of categories in the model formulation (i.e., EEZs and the high seas) can 390 somewhat affect the fit of the multinomial models used (see R package mgcv reference manual, GAM multinomial logistic regression, for more detail ³⁶), typically (in our case) by inflating 391 392 uncertainty estimates when EEZs with few observed locations are ordered first. For consistency 393 we ordered the categories for each model from the highest to lowest numbers of 'observed' 394 locations.

For each species, the formulation of the models depended upon species life history and dataset quality (Supplementary Table 4). Models were not developed for species who spent most of their time within a single EEZ (yellowfin tuna; California sea lion). Yearly models were considered for species with balanced datasets over multiple years (Supplementary Tables 1 and 4): female northern elephant seals (Supplementary Fig. 3) and salmon shark (Supplementary Fig. 400 4).

For some species, there was enough information to separately model life history stages or
breeding populations. Separate models were fitted for female northern elephant seals from U.S.
and Mexican breeding populations (Supplementary Fig. 1). Two model groups were also

404 constructed for Pacific bluefin tuna. Group 1 included all PBT individuals. The second PBT
405 model group included only those tuna that undertook trans-Pacific migrations (n=12,
406 Supplementary Fig. 2).

407 For western Pacific leatherbacks, we modeled each breeding population and the 408 California foraging population separately due to the multi-modality of tag deployments and our 409 interest in differences in EEZ use by the different populations. In order to compare EEZ 410 occurrence in the months following breeding (regardless of whether breeding occurred in 411 summer or winter), and in the months following tag deployment in the California foraging 412 grounds, the number of days elapsed following tag deployment was the fixed effect (rather than 413 day of year, Supplementary Table 4). Therefore, the model was related to a life history event, 414 estimating EEZ and high seas occurrence during the migration following this event, rather than 415 to the calendar year.

416

417

4. Model predictions and uncertainty

418 From fitted models we predicted the probability that a randomly selected individual from the 419 tracked population would occur in an EEZ or in the high seas on each day of the year. Use of 420 EEZs by tagged western Pacific leatherback turtles was estimated only for the 7.5-9 months 421 following tag deployment, due to the impacts of tag attrition on data availability after this time. 422 EEZs with few occurrences were grouped together into an "Other" category. For Sooty 423 Shearwaters, island EEZs falling within the Central Pacific were also grouped together, representing their migration corridor¹⁵. Population-level predictions were derived by setting the 424 425 random intercept effect of individual in the model to zero. We estimated the proportion of the

426 year spent in each EEZ or in the high seas for each tracked population by summing the 427 population-level predicted daily probabilities over the course of a year and dividing by 365. 428 Models also provided predictions of the probabilities of specific tagged individuals occurring in 429 EEZs or the high seas on each day of the year. These probabilities were sometimes highly 430 variable among tagged individuals of a species/population (Supplementary Fig. 9). In some 431 cases, non-negligible numbers of individuals of a species visited an EEZ during a year (as an 432 annual summary), but the mean population response on any given day of year may not represent 433 this. For example, 20% of individual white sharks tracked in this study (Supplementary Table 1) 434 visited Hawaii and 6% of all locations were in Hawaii (Supplementary Table 2), but the mean 435 population response on any day of year was near 0 (Fig. 2, Table 1). A similar pattern was 436 observed for elephant seals in Canada (Supplementary Fig. 9). Because the models estimate a 437 population response on a given day of year, a substantial proportion of individuals would need to 438 visit the EEZ at the same time to be represented in the population response. Summaries of 439 individual use of EEZs and the high seas (Supplementary Table 1), therefore provide information 440 that may be of use to managers in addition to model results, keeping in mind caveats due to variability in tracking duration discussed in Methods sections 1 and 5. 441

Uncertainty in model predictions was characterized by simulating a sample of estimates from the posterior distribution of the model parameters¹⁴. The posterior distribution was assumed to be multivariate normal with means equal to the parameter estimates and variances/covariances from the estimated covariance matrix. The estimated uncertainty generally reflected sample size across species/populations and throughout the year for individual species/populations demonstrating the usefulness of the model in accounting for sample size. For example, compare uncertainty and sample sizes in northern elephant seal yearly models to results for the full

449 population. Uncertainty in the proportion of the year spent in each EEZ and in the high seas was 450 estimated by calculating these proportions for each parameter set from the posterior sample and characterizing the distribution of proportions across the sample. 451

452 The estimates of uncertainty presented here are likely underestimates of the true 453 uncertainty in the effect of day of the year on the occurrence of tagged animals in EEZs and the 454 high seas. Although individual identity was included as a model effect, sequential correlation in 455 the model residuals for an individual could have remained, in which case the true uncertainty in 456 the day-of-year effect would be greater. Also, the presence data that the models were fitted to 457 were themselves derived from state-space model location estimates with associated positional uncertainty¹ that was not accounted for here. Nevertheless, the estimates of uncertainty 458 459 presented here provide an upper bound on the confidence that should be placed in the estimated 460 effects of day of year on the occurrence of the tagged populations in EEZs and the high seas.

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5. Additional considerations: Effects of tag deployment location and variability in 463 deployment data and track duration on probability estimates

464 Statistical summaries of time spent in EEZs from electronic tracking data are influenced by the 465 distribution of track durations and deployment dates and locations. Early in a track, individuals 466 have a high likelihood of being located within the deployment EEZ because tags were deployed 467 there. As time passes, individuals have the ability to disperse from the release location and the 468 proportion of time spent within the deployment EEZ should level to a more biologically 469 representative proportion unaffected by the initial tag deployment event. Therefore, an 470 interaction between deployment location and track duration has the potential to bias estimates of 471 EEZ use in favor of the EEZ in which tags were deployed when a large proportion of the tracked472 population has short-duration tracks.

473 We explored the effects of track duration and timing of tag deployment in the TOPP 474 dataset by calculating the running proportion of time spent by each individual within primary 475 EEZs and the high seas according to the relative day along each individual's track (i.e. days 476 elapsed since deployment) and according to the month in which the tag was deployed 477 (Supplementary Fig. 8). For example, most tags were deployed on Pacific bluefin tuna in March, July-August, and November-December within the Mexican EEZ¹. Individuals tracked for less 478 479 than 30 days spent 80-100% of their time within Mexico. Individuals tracked for greater than a 480 year, spent 50% of their time in Mexico (Supplementary Fig. 8) with little change in this 481 proportion as track length increased beyond a year. Tuna released in Mexico in November spent 482 a higher proportion of their time in Mexico in the few months after being tagged than those 483 tagged in July (Supplementary Fig. 8).

484 We attempted to account for this effect of track duration and tag deployment location in 485 the models by exploring the use of a day-of-track term. Ideally such a term would capture the 486 higher probability of being in the tagging EEZ at the beginning of a track and the effect would 487 diminish during a track. Some of the models we explored partially captured the expected day-of-488 track effect, but the predicted effects did not diminish monotonically over time and often 489 exhibited non-intuitive patterns later in a track. For example, the Pacific bluefin tuna model 490 exhibited an expected decrease in the predicted probability of being in the Mexico EEZ (the 491 deployment EEZ) during the early part of a track, but the predicted probability increased later in 492 the track and exhibited non-intuitive patterns for some days of the year (Supplementary Fig. 10).

These results suggested that the combination of data and model structure used was unable tocapture the expected effect of track duration, so these models were not considered further.

Tag deployment location is an experimental design feature of a tracking dataset that can bias interpretation of space use for the tracked population. In this paper, it might have imposed an upward bias on estimates of probability of occurrence in EEZs in which tags were deployed, especially when combined with a dataset subject to a high amount of tag attrition (i.e. "short" tracks). For datasets in which many individuals were tracked for multiple years (i.e. salmon sharks), we expect a minimal effect of this bias for the tracked population.

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584 End Notes

585 **Supplementary Information** is linked to the online version of the paper.

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- and K.C.W. designed experiments and deployed electronic tags on fish and sharks. S.R.B.,

612 P.H.D., G.L.S., and B.A.B., designed experiments and deployed electronic tags on sea turtles.

613 D.P.C., P.W.R., S.E.S., and B.R.M. designed experiments and deployed electronic tags on

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618

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621 **Competing interests:**

622 Kristina Gjerde is an unpaid member of the Sargasso Sea Project Inc. Board, the Global Ocean

- 623 Biodiversity Initiative Scientific Steering Committee, the Deep Ocean Stewardship Initiative
- 624 Executive Board, the High Seas Alliance Steering Committee and the Deep Ocean Observing
- 625 Strategy Scientific Steering Committee.

626 Data Availability Statement: The tracking data used are archived in the U.S. Animal Tracking
627 Network Data Assembly Center. <u>http://oceanview.pfeg.noaa.gov/ATN/</u>

- 628 **Code Availability Statement:** Custom R scripts will be made available via GitHub upon
- 629 publication in the political-biogeography project, and are also available upon request to the
- 630 corresponding author.