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Tracking of marine predators to protect Southern Ocean ecosystems

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101

102 **Southern Ocean ecosystems are under pressure from resource exploitation and climate**
103 **change^{1,2}. Mitigation requires identification and protection of Areas of Ecological Significance**
104 **(AES), which have eluded identification at the ocean-basin scale. For this globally significant**
105 **region, we identify AES using assemblage-level tracking of marine predators and assess current**
106 **threats and protection levels. Integration of >4000 tracks from 17 bird and mammal species**
107 **reveal AES around sub-Antarctic islands in the Atlantic and Indian Oceans and over the**
108 **Antarctic continental shelf. Fishing pressure is disproportionately concentrated inside AES, and**
109 **climate change over the next century is predicted to impose pressure on these areas,**
110 **particularly around the Antarctic continent. Currently, 7.1% of the ocean south of 40°S is under**
111 **formal protection, including 29% of the total AES area. The establishment and regular revision of**
112 **networks of protection encompassing AES is needed to provide long-term mitigation of growing**
113 **pressures on Southern Ocean ecosystems.**

114

115 The Southern Ocean, defined here as the circumpolar waters south of 40°S, is home to a unique fauna
116 and plays an important role in biogeochemical cycles and the global climate system¹. Past industrial
117 sealing, whaling and demersal fishing caused significant perturbations from which some Southern
118 Ocean ecosystems are only now starting to recover³. Squid and toothfish harvesting continue^{4,5} and
119 interest in the expansion of Antarctic krill fisheries is growing⁶. These target species are important prey
120 for upper trophic organisms with krill a key component of the Southern Ocean food web, raising
121 substantial concerns about impacts on Southern Ocean ecosystems². Anthropogenic greenhouse gas
122 emissions are simultaneously causing large changes to the Southern Ocean⁷. Strong interest has
123 therefore developed in long-term conservation of the Southern Ocean, but management authorities face
124 the considerable challenge of implementing conservation goals within existing management
125 frameworks².

126

127 A first step in meeting this challenge is to identify regions that should be considered for protection, for
128 example because of their high biodiversity, biological productivity, or particular importance for life-history
129 stages of species^{8,9}. The distribution and demography of marine predators provides a viable basis for
130 this¹⁰, particularly in the vast and remote Southern Ocean where integrated ecosystem measures are
131 difficult to obtain at management-relevant, ocean-basin scales¹¹. Indeed, on-shore measures of
132 Southern Ocean marine predators have been used as regional indicators of ecosystem status for
133 several decades¹². Spatial aggregations of predators at sea identify not only areas that are important to
134 the predator species themselves, which depend on lower trophic levels¹³, but also areas of broader
135 ecosystem importance such as regions of elevated productivity and biomass at lower trophic levels¹⁴.
136 Combining information across predator species with diverse diets and life histories is essential for an

137 ecosystem-wide approach that is less susceptible to factors affecting individual species¹². Recognition
138 of the value of tracking data for broad-scale conservation decision-making is growing¹⁵.

139

140 **Using predator tracking data to identify Areas of Ecological Significance**

141 In the Southern Ocean, many predator species with differing diets and movement patterns have been
142 tracked¹⁶. We synthesised tracking data from 4060 individuals of 17 species (Fig. 1a) to provide a
143 circumpolar assessment of regions of ecological importance in the Southern Ocean. We identified
144 regions preferred by multiple predator species as indicators of high levels of lower trophic biomass and
145 biodiversity, and refer to these as Areas of Ecological Significance¹⁷ (AES). Our definition of AES is not
146 the same as Ecologically and Biologically Significant Marine Areas (EBSAs), or Key Biodiversity Areas
147 (KBAs). However, it is consistent with several of the criteria used for defining EBSAs or KBAs,
148 particularly biological productivity and diversity⁸, and so provides a similar qualitative, integrated
149 assessment of biodiversity patterns.

150

151 We assembled tracking data from 17 species of seabirds (12 species) and marine mammals (5
152 species), collected between 1991 and 2016¹⁶. We used habitat selection models (Methods and
153 Supplement, Extended Data 1–3) of individual predator species and then combined their spatial
154 predictions to identify regions important to our full suite of species (Fig. 1b). This enabled us to account
155 for incomplete tracking coverage (i.e., colonies from which no animals were tracked) and predict habitat
156 importance for each species across the entire Southern Ocean. Combined, these provided an
157 integrated and spatially explicit assessment of areas of high biodiversity and biomass at multiple trophic
158 levels. Sea surface temperature (SST) and wind strength were most often the best predictors of habitat
159 selectivity in these species-specific models (Extended Data 4). SST has been linked to global patterns
160 of marine biodiversity¹⁸; in the Southern Ocean it acts as an indicator of water masses with different
161 ecological properties¹⁹. Wind exerts several influences including driving ocean currents and mixing,
162 transport of iron, sea-ice dynamics, and determining primary production²⁰ and has been linked, for
163 example, to the global distribution of albatrosses and petrels²¹. The importance of other predictor
164 variables differed among species (Extended Data 4). The relationship between habitat selectivity and
165 environmental predictors differed across species showing how species used their environments in
166 different ways (Extended Data 5).

167

168 **Distribution of Areas of Ecological Significance**

169 Regions with the highest overall habitat importance scores were identified as Areas of Ecological
170 Significance (AES; calculated as the upper decile of those scores). These were located over the
171 Antarctic continental shelf (89% of AES pixels south of 60°S were over or within 200 km of the shelf)
172 and in two northerly aggregations: one encompassing much of the Scotia Sea and surrounding waters,
173 and the second covering the chain of sub-Antarctic islands from the Prince Edward Islands through to

174 parts of the Kerguelen Plateau (Fig. 1c). Regions of lower importance were identified in the southern
175 Pacific and Indian Oceans. The distribution of AES is associated with the availability of suitable
176 breeding/resting habitat, as well as regional oceanography and sea-ice dynamics that affect biological
177 production (Fig. 1c). The AES were based on a combination of island-breeding and wholly pelagic
178 species, and therefore reflect broad-scale patterns of importance. These patterns are supported by: (i)
179 broad-scale patterns of primary production — Southern Ocean land masses provide iron fertilization that
180 stimulates downstream production in this otherwise iron-limited ecosystem²²; (ii) historical whaling
181 catches north of 60°S, which show that fewer whales were taken in the southern Indian or Pacific
182 Oceans, and that the region identified as an AES in the south Atlantic corresponds with high whale
183 catches²³; and (iii) recent and historical estimates of Antarctic krill distribution, with high concentrations
184 in the south Atlantic and lower concentrations in the south Pacific and southern Indian Ocean²⁴. The
185 AES in the south Atlantic corresponds to the area of elevated krill biomass, whereas the AES in the
186 Indian Ocean partially corresponds to a region dominated by myctophid fish and other euphausiids²⁵.

187

188 **Exposure of Areas of Ecological Significance to potential stressors in the Southern Ocean**

189 The Southern Ocean is subject to several stressors that influence its ecosystems, including expanding
190 resource extraction and rapid climate change²⁶. We note that both temperature and wind, which were
191 important parameters in many of our species-specific habitat models, are changing and projected to
192 continue to do so²⁷.

193

194 Fishing has both direct effects, through incidental bycatch, and indirect effects through resource
195 competition²⁸. Many demersal finfish were exploited during the latter part of the 20th century, leading to
196 the decimation of some stocks in the Antarctic and sub-Antarctic⁵. Finfish fishing in the Antarctic is now
197 regulated and focused on toothfish species caught with longlines. Fisheries for Antarctic krill began in
198 the 1960s and are now concentrated in the south Atlantic sector, most notably at the Antarctic
199 Peninsula and South Shetland Islands, the South Orkney Islands and South Georgia⁶. Krill is managed
200 with a low, precautionary catch limit, taking account of the key role it plays in the Antarctic food web. By
201 global standards, fishing pressure in the Southern Ocean is low²⁹, but indications are that pressure on
202 its marine resources will grow^{2,5,6}. Fishing effort (Fig. 2a) was significantly different inside and outside of
203 the AES (Fig. 2b), with a disproportionate amount of moderate-to-high effort (≥ 100 total hours of fishing)
204 occurring inside AES. Of cells with moderate-to-high fishing effort, 37.9% were AES, despite AES only
205 representing 10% of the study area. Areas of conspicuous fishing effort around southern South
206 America, New Zealand, and Australia should be treated with caution, since our study does not include
207 temperate predator species likely to figure prominently in these ecosystems (Fig. 2a). Nonetheless,
208 relatively high intensity fishing areas directly relevant to the Southern Ocean occurred around the
209 Falkland Islands/Islas Malvinas, where squid and some finfish are targeted, South Georgia (ice fish, krill

210 and toothfish), at the West Antarctic Peninsula (krill), and over the Kerguelen (toothfish and ice fish) and
211 Campbell (squid and finfish) plateaux⁴⁻⁶. Relatively important fisheries for toothfish also occur within the
212 Ross Sea³⁰.

213
214 The physical attributes of the Southern Ocean are changing. Sea-ice is a critical component of high-
215 latitude ecosystems, playing central roles in oceanographic, biogeochemical, and ecological processes.
216 The biological consequences of sea-ice changes in the Southern Ocean include changes in breeding
217 site availability or access, prey availability, and changes to ecosystem structure and function³¹. The
218 pattern of change in sea-ice in the Antarctic displays considerable regional and temporal variation. In
219 the West Antarctic Peninsula, sea-ice extent has declined markedly in recent decades, but has
220 increased in other areas³². Most climate projections indicate that overall sea-ice will decline over the
221 next century²⁷. Given the broad influence of both SST and wind on ecosystems, these components can
222 also influence aspects of an animal's biology, including breeding phenology, foraging success, survival
223 and reproductive performance²⁶. However, when we contrasted the rates of change of sea-ice duration,
224 SST and wind patterns inside and outside of the AES there were only slight differences, and
225 considerable regional variation (Extended Data 6). The subtle nature of the differences in environmental
226 change inside AES versus outside them does not negate the fact that the study area, overall, is
227 undergoing marked changes in physical environmental processes, and that ecologically important areas
228 are not being spared from these changes.

229

230 **Assessment of current and proposed spatial management**

231 Management of marine systems is complex, especially in areas beyond national jurisdiction³³ where
232 international effort is required, particularly for species that move between national and international
233 waters³⁴. Relevant management includes traditional process-oriented tools, such as individual species
234 protection, stock assessments, decision rules and catch limits, as well as spatial tools such as marine
235 protected areas (MPAs)³⁵, but also altered fishing practices for mitigating bycatch³⁶. In the high-latitude
236 Southern Ocean, the Commission for the Conservation of Antarctic Marine Living Resources (CCAMLR)
237 employs an ecosystem-based management framework intended to ensure that there are no long-term
238 effects from fisheries on marine ecosystems³⁷. This includes setting precautionary, spatially explicit
239 catch quotas and a call for the establishment of a network of MPAs, the design considerations of which
240 can include the potential to provide climate change refugia and the inclusion of reference areas to help
241 separate the effects of fishing from climate related environmental change. Both approaches will benefit
242 from better understanding of the locations of AES. Outside the CCAMLR framework, MPAs have also
243 been established by sovereign management authorities around some sub-Antarctic islands (Fig. 3a).
244 Several other MPAs are currently under development, including within CCAMLR and by national
245 authorities (Fig. 3a). However, the level of protection afforded by any individual MPA depends upon its
246 governance structure and the type and level of permitted activities (e.g., fishing)^{9,38}

247

248 An appropriately designed protected-area network can help buffer the effects of climate change and
249 reduce the effect of stressors such as bycatch or competition from fisheries³⁹. We therefore quantified
250 the coverage and placement of individual MPAs with reference to identified AES. Overall, 7.1% of the
251 ocean south of 40°S is currently protected by MPAs, and this would increase to 11.2% if all currently
252 proposed MPAs were implemented (Fig. 3b). This already meets, in a regional setting, the 10% global
253 Aichi Target 11 for 2020. The Southern Ocean's level of protection is high by global standards, as only
254 3.6% of the world's oceans fall within MPAs, increasing to 7.3% with the addition of planned and
255 announced MPAs³⁸. However, protection needs to be targeted at areas of high conservation value,
256 including those important for the persistence of biodiversity⁹. Existing MPAs cover 27% of the AES
257 identified (Fig. 3b). Southern Ocean MPAs are predominantly in sub-Antarctic regions, and here they
258 show high levels of congruence with AES (Fig. 3a). Of note is the Davis Bank region, south of the
259 Falkland Islands/Islas Malvinas, where there are high levels of fishing inside AES (Fig. 1, 2a, b). This
260 area is now part of a recently implemented MPA by Argentina (Fig. 3a). Adoption of proposed MPAs for
261 the Antarctic continental margins would raise MPA coverage of AES to 39% (Fig. 3b), including areas in
262 East Antarctica, the Weddell Sea, and the Antarctic Peninsula. The largest total AES (4.0 million km²,
263 56% of AES) are under CCAMLR jurisdiction (Fig. 3a,c), followed by 1.9 million km² (27% of AES) in
264 national waters (EEZs) and only 1.2 million km² (16% of AES) are outside the CCAMLR Convention
265 Area and national waters (Fig. 3c). Implementation of MPA proposals would benefit Southern Ocean
266 ecosystems, especially those in the Antarctic Peninsula, East Antarctic and Weddell Sea.

267

268 **Likely effects of future climate change**

269 We estimated the likely effects of future climate change on the distribution of AES under two
270 Representative Concentration Pathway (RCP) simulations: a medium-forcing scenario (RCP4.5) and a
271 more extreme, high-forcing scenario (RCP8.5)⁴⁰. For each scenario, eight global climate models,
272 considered to be most suitable for Southern Ocean studies by reliably reproducing extant sea-ice
273 conditions, were used to predict locations of AES-like habitat in 2100. Here we discuss only the RCP8.5
274 results, since current CO₂ emissions are in line with this scenario⁴¹. Results for the moderate RCP4.5
275 scenario are presented in Extended Data 7. There was an overall reduction in AES-like area (-3.3%),
276 partitioned into an increase in sub-Antarctic AES-like cells (+5.7%), but outweighed by a decrease in
277 Antarctic AES-like cells (-10.2%).

278

279 In the sub-Antarctic, AES-like areas generally moved south (Fig. 4a), resulting in an overall growth in
280 sub-Antarctic AES area (Fig. 4b). This general southward migration of important habitat is consistent
281 with projections for individual predator species (e.g.,⁴²) as well as for other species including krill and
282 salps^{43,44}. The advantages for predators from the overall increase in area of sub-Antarctic AES may be
283 offset by increased cost of travel to more distant foraging grounds, at least for diving central place

284 foragers (penguins and fur seals), while volant species (albatrosses and petrels) or those unconstrained
285 by terrestrial breeding sites (whales) may benefit from increased sub-Antarctic foraging opportunities
286 (e.g., ⁴⁵). Changes in the future distribution of AES-like areas along the Antarctic margin are more
287 spatially heterogeneous, with areas of AES loss interspersed with areas of AES gain or retention (Fig.
288 4a). However, there will be a net loss (-10.2%) of AES-like cells in the CCAMLR Convention area (Fig.
289 4b). The heterogeneity of this pattern is due in part to the dynamic nature of the high-latitude Antarctic
290 marine environment and the uncertainty across a number of climate-model variables in this region. This
291 uncertainty is due to the variability in skill of models in reproducing current climate and the large
292 envelope of projected responses from those models. Our projections are based on unchanged future
293 availability (i.e., colony locations and sizes) and species-environment relationships. However, as
294 species adapt to future pressures and changes to available breeding habitat, populations are likely to
295 change both their preferred colony locations and habitat usage. Sub-Antarctic-breeding species have
296 limited availability of alternative breeding sites, but colony sizes might change. Ice-breeding species
297 may be able to relocate, whilst land-breeding species that require ice-free terrain may be able to occupy
298 previously vacant areas, or some may move to regions that become ice-free due to changing local
299 conditions (e.g., ⁴⁶). Our projected loss of AES-like habitat on the Antarctic margin suggests that these
300 populations will be under pressure as the climate continues to change, and therefore continued
301 monitoring of these species, and on-going assessment of the effectiveness of management actions (e.g.
302 MPAs), will be important. Monitoring of colonies will need to detect local colonisations, particularly when
303 populations are small⁴⁷. As part of the designation of MPAs within CCAMLR, research and monitoring
304 plans are necessary and required; *inter alia*, these plans should consider changes to species-
305 environment relationships and other dynamic processes within and adjacent to the protected area, given
306 the pressures of ongoing climate change.

307
308 There was a mixed response across the eight climate models, with changes in the number of AES-like
309 cells included in current MPAs ranging from -8.7% to +8.4% (Fig 4b). When the proposed MPAs are
310 included (current + proposed), all climate models indicated a decrease (between -16.9% to -0.9%) in the
311 number of AES-like cells within MPAs. This suggests that proposed MPAs are in areas projected to
312 become less-similar to existing AES by 2100. Any protection afforded by MPAs in such areas could
313 provide better medium-term opportunities for populations to adapt as they will not have to cope with
314 both climate change and other stressors during that period.

315 316 **Conclusion**

317 Our work provides strong evidence in support of the ecological importance of existing and proposed
318 Southern Ocean MPAs. By integrating tracking data from a suite of predators we identified regions likely
319 to have high biodiversity and biomass of the prey (and concomitant ecosystems) of the animals that
320 were tracked. Our AES are clearly candidates for protection; as such, the implementation of the

321 proposed MPAs within the CCAMLR region would greatly increase the protection of important Southern
322 Ocean habitat. Several MPA proposals have failed to reach consensus within the CCAMLR process
323 and, even when adopted result in MPAs with varying degrees of protection. Many sources of input are
324 needed to establish MPAs, but the AES described here will play a key role in making the scientific case
325 in this multi-faceted process^{2,48}, by providing ecosystem-level analysis of areas most warranting
326 protection. MPA design should also consider future conditions. Pressures on AES due to climate
327 change will affect all parts of the Southern Ocean, but their effects are likely to be strongest along the
328 Antarctic margin. Species responses to these pressures are currently difficult to predict, highlighting the
329 importance of continued monitoring as part of ongoing management actions. Because only 16% of all
330 Southern Ocean AES lie outside the CCAMLR Convention Area or national waters, the responsibilities
331 for these future actions fall mostly to CCAMLR members and those nations with sovereign territory in
332 the sub-Antarctic. Adaptive management approaches to conservation measures (including MPAs) will
333 be necessary to deal with these future changes in a timely way. The Southern Ocean can be an
334 exemplar of how science, policy and management can interact to meet the challenges of a changing
335 planet. In the Southern Ocean, these challenges will be considerable, including increased fishing
336 pressure as the global demand for marine resources grows⁴⁹. Our results highlight where future
337 science-informed policy efforts might best be directed, including both adaptive spatial protection and
338 improved robust fisheries management. Similar synthetic approaches should capitalise on the
339 increasing amount of tracking data being collected through large-scale initiatives (e.g.,⁵⁰) to indicate
340 regions in need of protection globally.

341

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458
459

460 METHODS

461 Analytical overview

462 We assembled tracking data from 17 species of seabirds and marine mammals, collected between
463 1991 and 2016, from across the Antarctic predator research community¹⁶. Birds and mammals comprise
464 the majority of top predator species in the Southern Ocean, which has few other large, highly mobile
465 marine predator taxa (bony and cartilaginous fishes). These include toothfish, southern bluefin tuna
466 (*Thunnus maccoyii*, which occur in the northernmost part of our study area), and a small number of
467 shark species. Very few of these fish and shark species have been tracked, with very few tracking data
468 available south of 40°S⁵¹. However, this does not detract from the underlying logic of our approach: that

469 by using the at-sea distributions of an ecologically diverse suite of predators we can identify areas of
470 ecological importance. This benchmark dataset represents 4060 individual tracks and more than 2.9
471 million location estimates (Fig. 1a). After filtering and quality control, we retained 2823 tracks comprising
472 2.3 million locations¹⁶. The ~30% of tracks that were excluded were those with poor quality location fixes
473 that could not be properly filtered, tracks from individuals that did not actually depart the colony, or
474 tracks other problems detected during the rigorous quality control process that we implemented. The full
475 process is described in our companion data paper¹⁶, which makes available all of the data for use by the
476 broader community, without providing further analytical investigation to consider the matters raised
477 here. The environmental covariate values along each of these tracks (the *used* habitat) were compared
478 statistically with the habitat *available* to each animal, thereby allowing each species' habitat selection to
479 be determined (e.g.,^{52,53}; see Extended Data 1 and 2). We fitted habitat selection models for different
480 life-history stages within a species. Despite the considerable size of the data set, it is not an exhaustive
481 representation of animals from all known colonies (for central-place foragers) or geographic regions (for
482 non-central-place foragers). To account for incomplete tracking coverage, we used the fitted habitat
483 selection models to map habitat importance for each life-history stage of each species across the entire
484 Southern Ocean, including areas around colonies without tracking deployments (Extended Data 3). For
485 each species we calculated the average habitat importance across life-history stages. For colony-
486 breeding species, colony sizes were used to weight the habitat importance values, upweighting areas
487 important to large colonies (Extended Data 8). Southern Ocean predator species can be clustered into
488 Antarctic and sub-Antarctic species (Extended Data 9). We mapped assemblage-level habitat
489 importance (Extended Data 10) for each of these two groups (hereafter "overall habitat importance"
490 maps) by averaging across species-level maps. To calculate the overall map, we took the maximum of
491 the two assemblage-level importance values in each cell. Areas with high values of overall habitat
492 importance (in the top decile of values) indicate areas that are attractive to many species; these
493 represent Areas of Ecological Significance¹⁷. We then compared the overall habitat importance values
494 inside and outside Areas of Ecological Significance in the context of fishing effort and changes in
495 physical environmental conditions (duration of sea-ice cover, sea surface temperature (SST) and wind
496 speed). We finally quantified the spatial protection afforded to Areas of Ecological Significance under
497 current and proposed spatial management plans.

498

499 We describe the methods in more detail in the Supplementary Information. We conducted all the
500 analyses in R⁵⁴.

501

502 **Tracking data**

503 The data represent output from a variety of tracking tag types, providing location estimates at different
504 spatio-temporal resolution and accuracy. We applied a state-space model⁵⁵ to estimate most probable
505 locations at regular temporal intervals while accounting for potential errors in the location estimates with

506 automatic and manual quality control before and after filtering¹⁶. While this procedure does not make the
507 track from a light-based tag as accurate as one from GPS device, it does provide a consistent
508 characterization of the positional accuracy across different tag types, allowing the uncertainty in position
509 to propagate into the uncertainty in the parameters of the fitted movement model and in the track
510 simulation step (see below). We further note that the light-based tag deployments were made almost
511 exclusively on sub-Antarctic animals (albatrosses and fur seals). The spatial scale of our results (Areas
512 of Ecological Significance) in the sub-Antarctic zone (~5 million km²) is considerably larger than the
513 likely scale of positional error of light-based tags (~100 km) and so we do not believe that using a
514 mixture of tag types has adversely affected our results.

515

516 **Life-history stages**

517 Most of the species in the study are central-place foragers (*i.e.*, they return periodically to land or sea-
518 ice to breed, moult or rest). The constraints faced by these predators at different stages in their life-
519 history cycle mean that their movements differ markedly across these stages. We therefore fitted
520 models separately for up to five predefined life-history stages in each species' breeding cycle. We
521 automatically assigned tracks to these stages based on calendar date, with manual reassignment where
522 necessary following examination of individual movement patterns. This resulted in 40 data subsets (17
523 species * 1–4 life-history stages) with sufficient data for habitat selection modelling (Supplementary
524 Information Table S1).

525

526 **Simulating tracks to estimate available space**

527 The observed locations only provide information about where animals occur, not about where they could
528 have gone. To estimate the geographic space potentially available to animals, we simulated sets of
529 tracks for each observed track. For each observed track, we simulated 50 tracks using the movement
530 model described above⁵⁵. This yielded simulated tracks with movement characteristics (distributions of
531 step length and turning angle) that are the same as the observed track, but they are random and
532 independent of environmental effects. Thus, the simulated tracks offer an estimate of the geographic
533 space that each animal could have occupied (given its movement characteristics and track length) if it
534 had no habitat preferences. The environmental differences between the available geographic space and
535 the utilized geographic space allow the habitat selection of the organisms to be estimated, as detailed
536 below. Locations at the animal's home colony, and locations at known terrestrial resting sites, were fixed
537 at the corresponding time and date in the simulated tracks in order to accurately simulate central place
538 foraging behaviour (Supplementary Information).

539

540 **Environmental data**

541 To characterize the biophysical environment at observed and simulated locations, we compiled a suite
542 of 19 environmental covariates (Extended Data 2, Supplementary Information Table S2) and extracted

543 the value of these at each location. The covariates were remotely-sensed, measured *in-situ* or model-
544 estimated and represent biophysical features that influence the movement, distribution and density of
545 marine predators (e.g.,^{52,53}). It was not computationally feasible to temporally match environmental data
546 to each location estimate. Rather, we created a climatology spanning each tracking data subset
547 (species by life-history stage combination), using the predefined stage dates. We took the mean (or
548 standard deviation) of the environmental data that fell on these days of the year (stage dates) over the
549 whole study period (November 1991 to June 2016). Some covariates (e.g., salinity difference) were only
550 available as monthly climatologies, and we used the months corresponding with the stage dates to
551 calculate the mean (or standard deviation). All covariates were resampled to a 0.1° x 0.1° grid; hereafter
552 we refer to the pixels of this grid as 'cells'. We checked the covariates for each data subset for missing
553 values and if >10% of values were missing we excluded the covariate from that model. This influenced
554 mainly chlorophyll-a concentration, which was excluded from 17 of the 40 habitat models
555 (Supplementary Information Table S1). This affected life-history stages with a large proportion of winter
556 days, since chlorophyll-a data has poor winter satellite coverage due to being obscured by high cloud
557 cover. However, chlorophyll-a was rarely an important predictor in the models in which it was included;
558 thus, excluding chlorophyll-a from models probably had only a negligible effect.

559

560 **Habitat selection models**

561 We used a habitat selection modelling framework⁵⁶ to model and predict the space use of marine birds
562 and mammals of the Southern Ocean. These models use the observed locations of each individual
563 animal and an estimate of the geographic space available to each individual, along with covariates that
564 characterize their environment. The environmental differences between the habitat that was utilized and
565 the habitat that was available allow the habitat selection of the organisms to be estimated. To fit the
566 models, we used boosted regression trees, a machine-learning algorithm that produces an ensemble of
567 regression trees that have been iteratively fitted in a boosting process to improve accuracy⁵⁷. We tested
568 several other algorithms but boosted regression trees showed the best predictive performance in
569 another study⁵³ and in our tests. For a given location, the response variable was whether the location
570 was an observed or simulated (available) location, and the explanatory covariates were the associated
571 environmental covariates. Boosted regression trees have four parameters that must be set: the number
572 of trees (boosting iterations), the maximum tree depth, the learning rate (shrinkage) and the minimum
573 number of observations in a node. We chose these values as the combination that minimised the area
574 under the receiver operating characteristic curve (a measure of model predictive performance) during
575 10-fold cross-validation. We also used this metric to evaluate the final fitted models. We used the fitted
576 model to generate spatial predictions for the entire study region and we estimated the uncertainty
577 associated with these predictions using a bootstrap approach (Supplementary Information)

578

579 *Accessibility model*

580 The modelling procedure described above does not account for the accessibility of a given location to
581 an individual animal (in effect, it estimates the habitat selection of a given location in terms of its
582 environmental characteristics, but without considering whether or not the animal could actually reach
583 that location). For central-place foragers in particular, this is an important consideration. We therefore
584 used a second set of models to account for this⁵³. We modelled accessibility in terms of the number of
585 observed + simulated locations in a given cell as a function of that cell's distance to the deployment
586 colony. We fitted binomial models with a smooth, monotonic decreasing constraint⁵⁸, under the
587 assumption that the accessibility of cells should decrease with geographic distance. To estimate
588 uncertainty, we sampled curves from the posterior distribution of each fitted accessibility model to use in
589 a bootstrap approach (Supplementary Information).

590
591 We used these models to predict the accessibility of each cell over the study region to each species
592 during each life-history stage (that is, given the distance of a cell from a colony, the fitted accessibility
593 model provides an estimate of the probability that animals from that colony would be able to visit that
594 cell). For colony-breeding species (those other than humpback whales, crabeater and Weddell seals),
595 colony sizes were used to weight this accessibility estimate: for a given cell, the accessibility from all
596 known colonies of that species was calculated. A weighted mean of these accessibilities was then
597 taken, using colony sizes as weights. Thus, this weighted accessibility represents the probability that a
598 randomly-selected individual from the global population would be able to visit that cell, effectively
599 upweighting cells in the vicinity of large colonies.

600
601 For the non-colony breeding, ice-associated seals (crabeater and Weddell seals), we modelled
602 accessibility as a function of distance beyond the ice edge (15% ice concentration contour), rather than
603 distance to the colony. For humpback whales, we assumed that the whole study area was equally
604 accessible.

605

606 **Habitat Importance**

607 *Transforming output and combining models*

608 The habitat selection models predict the value of the habitat at a location given that the animals could
609 access that location. The predictions of the habitat selection models were therefore multiplied by the
610 predictions of the accessibility models to yield an index that reflects both the habitat selection of each
611 cell and its accessibility to the animals. This is not an estimate of the probability of a species using a
612 given cell, because that probability also depends on the prevalence of the species⁵⁹. Since prevalence
613 varies between species, our habitat selection estimates cannot be compared directly between species.
614 We therefore partitioned the cells into decreasing percentiles based on area⁵² in order to obtain a
615 habitat importance map expressed in terms of area (e.g., cells with values of 90 or higher represent the
616 top 10% most important habitat by area for that species). We refer to this as habitat importance, and

617 these maps can be compared among species. To create a single habitat importance layer for each
618 species, we averaged the stage-specific habitat importance layers.

619

620 *Species grouping*

621 We calculated community-level habitat importance by averaging species-specific habitat importance
622 maps. Sub-Antarctic regions are naturally more species-diverse than those of the Antarctic, and so a
623 simple average of all species together tended to strongly favour sub-Antarctic areas simply because of
624 their greater species diversity. To account for the differences in species richness between the Antarctic
625 and sub-Antarctic, we first defined two species groups using an Unweighted Pair Group Method with
626 Arithmetic Mean hierarchical clustering with Manhattan distance, applied to habitat importance scores
627 (Extended Data 9). This produced two clear groups: an Antarctic species group (emperor penguin,
628 crabeater seal, Antarctic petrel, Adélie penguin, Weddell seal) and a sub-Antarctic species group
629 (Antarctic fur seals, black-browed albatross, wandering albatross, sooty albatross, grey-headed
630 albatross, king penguin, macaroni/royal penguin, light-mantled albatross, white-chinned petrel). The
631 wide-ranging humpback whales and elephant seals did not clearly fall into either cluster, and so were
632 treated as belonging to both groups. The mean habitat importance was calculated for each of these
633 groups separately and then combined (Extended Data 10) by taking the maximum of the two values
634 (Antarctic and sub-Antarctic) in each pixel. We refer to this final layer as the overall habitat importance.

635

636 **Areas of Ecological Significance**

637 To identify the most important areas, we calculated the 90th percentile (top decile) of the overall habitat
638 importance values. Cells with overall habitat importance values above this threshold together comprised
639 Areas of Ecological Significance.

640

641 **Environmental pressures**

642 To assess past environmental stressors on the Southern Ocean ecosystem, we calculated change in
643 SST, wind speed and sea-ice duration. We selected SST and wind because they were frequently the
644 most important predictor variables in the habitat models (Extended Data 4), and sea-ice concentration
645 since this was an important predictor for Antarctic species. Moreover, these variables are considered
646 important drivers of ocean and ecosystem dynamics (e.g.,^{18,60}) and key axes on which environmental
647 change in the Southern Ocean has been detected (e.g.,²⁶). For each cell, we calculated change in SST
648 (°C) or wind speed (m/s) as the difference between mean SST or wind speed in 1987–1999 and 2007–
649 2017. For sea-ice duration, we calculated the difference in the mean number of days per year that each
650 pixel had a sea-ice concentration >15%, for the same periods. These periods represent the decades at
651 the beginning and end of a 30-year period covering our study period. Thirty years is also the
652 recommended period for climate assessments⁶¹. We also obtained data on fishing effort, which is
653 considered to be a major environmental stressor in many regions of the Southern Ocean (e.g.,^{29,62})

654 from the Global Fishing Watch dataset, covering the period 2012–2016²⁹. We compared the values of
655 these four stressors in the Areas of Ecological Significance and outside cells using random permutation
656 tests with 10,000 permutations. The null hypothesis is that stressor values inside and outside Areas of
657 Ecological Significance are from the same distribution.

658

659 **Future projections of Areas of Ecological Significance**

660 Our predicted AES (under current environmental conditions) are determined by both the oceanographic
661 and climatic conditions of an area, as well as the accessibility of that area to each of our species of
662 interest. In principle it would be possible to use future projections of environmental data and
663 accessibility along with our fitted models in order to obtain future projections of AES. However, some
664 predictor variables are not available from the climate models used for the future projections, and while
665 other variables might appear to be available, they have different properties due to factors such as
666 different temporal and spatial resolution in the output, or the ability of the climate model to resolve the
667 relevant processes. For example, sea surface height from satellite altimetry gives information about
668 frontal and mesoscale features. Yet, while sea surface height is available as an output from many
669 CMIP5 models, those models do not explicitly resolve mesoscale features⁶³ and so the model-output
670 sea surface height data will not be acting as a proxy for the same oceanographic properties that
671 satellite-derived altimetry does.

672

673 To assess future distributions of AES-like habitat, we therefore used a k-nearest neighbour classifier
674 approach, conceptually similar to climate analogues (e.g.,⁶⁴). For each grid cell we compiled current
675 (end of 20th century) environmental conditions, as well as projected conditions at the end of the 21st
676 century from climate models (see below). In terms of accessibility, most of our study species breed in
677 colonies, and “accessibility” for these species is determined by both the geographic distribution of their
678 colonies as well as the colony sizes. Currently, future projections of colony location and size do not exist
679 for our study species, although initial work has begun for some species (e.g.,⁴⁶). Colony locations and
680 sizes were therefore assumed to remain constant, and so the accessibility of each grid cell to each
681 species was assumed to remain unchanged. For each grid cell, we compared its projected future
682 environmental and accessibility conditions to every cell in the current (20th century) grid and selected
683 the most similar five cells. If the majority of those cells were from current AES areas, the projected cell
684 was labelled as “AES-like”, otherwise “not AES-like”. These projections therefore provide an indication
685 of the future distribution of AES-like environmental conditions, under the assumptions that colonies do
686 not move or change in size, and that the animals do not change their habitat preferences. These
687 assumptions are unlikely to hold in reality; however, examining the changes in AES-like habitat under
688 these assumptions allows us to isolate the effects of environmental change from colony or habitat-
689 usage changes. As environmental change occurs, species are likely to adapt by changing their colony

690 distributions and habitat usage. The AES projections offer insights into the likely distribution of
691 environmental pressures, and thus where adaptation by species might be important.

692
693 Climate data were compiled from eight global climate models (ACCESS1.0, BCC-CSM1.1, CanESM2,
694 CMCC-CM, EC-EARTH, GISS-E2-H-CC, MIROC-ESM, and NorESM-M) considered to be most suitable
695 for Southern Ocean studies, by virtue of reliably reproducing extant sea-ice conditions⁶⁵. These models
696 were from phase five of the World Climate Research Programme's Coupled Model Intercomparison
697 Project (CMIP5). For each model, we extracted data for a 30-year period concomitant with our tracking
698 data (1976–2005), and for a thirty-year end-of-21st-century (2071–2100) period. We extracted future
699 (2071–2100) climate data from projections under two Representative Concentration Pathway (RCP)
700 simulations: a medium-forcing scenario (RCP4.5, which assumes that society implements changes to
701 limit future CO₂ emissions in the near future, with peak emissions occurring in 2040) and a more
702 extreme, high-forcing scenario (RCP8.5, which assumes little curbing of emissions and retains a strong
703 reliance on fossil fuels into the foreseeable future)⁴⁰. Reference (1976-2005) data were extracted from
704 hindcast model runs that attempt to simulate historical conditions, and consequently use observed CO₂
705 concentrations over the past 160 years to guide the models.

706
707 A maximum of eight variables were extracted for each model, depending on the available data (not all
708 models provide all variables), at monthly time resolution. The variables used were sea-ice
709 concentration, SST, sea surface salinity, sea surface height, the spatial gradient of sea surface height,
710 near-surface current speed, near-surface wind speed, and surface downward heat flux. The 30-year
711 mean and standard deviation of each variable was calculated over summer (December to February)
712 and winter (July to September) months. All variables were normalized to the range 0–1 prior to further
713 analysis.

714
715 The resulting set of up to 48 predictors (mean and SD of up to 8 environmental variables, each for
716 summer and winter, plus accessibility layers for 16 species) naturally showed high correlation between
717 many of the variables. We used a principal components analysis to reduce the dimensionality of this
718 data set, choosing the lowest number of principal components required to explain at least 95% of the
719 variance in the original data; this number ranged from 14–17 components, depending on the model and
720 scenario. For each projected-climate cell, the nearest neighbours in the historical-climate grid were
721 calculated using Euclidean distance on these normalized and dimension-reduced data.

722
723 Animal Ethics Statement.

724 All work was conducted under the appropriate National or Institutional Ethics approvals. There were:
725 Argentina (Dirección Nacional del Antártico), Australia (Australian Antarctic program; the University of
726 Tasmania), Belgium (Belgian Science Policy Office), Brazil (Brazilian Antarctic Programme; National

727 Council for Scientific and Technological Development - CNPq; Ministry of Science, Technology,
728 Innovation and Communications – MCTIC; Ministry of the Environment; CAPES), France (Agence
729 Nationale de la Recherche; Terres Australes et Antarctiques Françaises), Germany (Deutsche
730 Forschungsgemeinschaft, Hanse-Wissenschaftskolleg - Institute for Advanced Study), Italy (Programma
731 Nazionale di Ricerche in Antartide, PNRA), Japan (Japanese Antarctic Research Expedition; JSPS
732 Kakenhi grant), Monaco (Fondation Prince Albert II de Monaco), New Zealand (Ministry for Primary
733 Industries - BRAG; Pew Charitable Trusts), Norway (Norwegian Antarctic Research Expeditions;
734 Norwegian Research Council), Portugal (Foundation for Science and Technology), South Africa
735 (Department of Environmental Affairs; National Research Foundation; South African National Antarctic
736 Programme), UK British Antarctic Survey; Natural Environment Research Council, and USA U.S. AMLR
737 Program of NOAA Fisheries; US Office of Polar Programs.

738

739 **Data availability**

740 The tracking data are available in¹⁶. Computer code is available at <https://github.com/SCAR/RAATD>

741

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771

772

773 SUPPLEMENTARY INFORMATION

774 Supplementary Information is linked to the online version of the paper at www.nature.com/nature.

775

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815

816 **AUTHOR CONTRIBUTIONS**

817 MAH conceived and led the project.

818 RA, BA, GB, JB, MNB, LB, HB, C-AB, PB, J-BC, RC, DPC, RJMC, LDR, PJNdB, KD, SD, MD, KD, LE,
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821 ATak, ATar, LGT, PNT, WT, EW, HW, BW and JX collected and contributed data.

822 VA-G, HB, J-BC, SC, BD, MAH, LAH, KJ, AK, IJ, MAL, DN, BR, RRR, YRC, DT, LGT, PNT, AVdP and
823 SW processed and analysed the data.

824 MAH, HB, J-BC, SC, BD, LAH, IJ, MAL, BR, RRR, YRC, DT, LGT, PNT, AVdP, SW and SLC drafted
825 the paper.

826 All authors edited and proofread the paper.

827

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830 requests for materials should be addressed to mark.hindell@utas.edu.au.

831

832 **COMPETING INTERESTS**

833 HB, J-BC, BD, MAH, LAH, IJ, MAL, BR, RRR, YRC, DT, LGT, PNT, AVdP and SW are members of the
834 Scientific Committee on Antarctic Research (SCAR) Expert Group on Birds and Marine Mammals. SLC
835 is President of SCAR.

836 **FIGURE CAPTIONS**

837

838 **Figure 1. Areas of Ecological Significance in the Southern Ocean.** Tracking data from 17 predator
839 species (a) were used to model the habitat importance for each species. Combining these model
840 outputs gives overall habitat importance (b) and the upper decile of overall habitat importance delimits
841 Areas of Ecological Significance (white contours). (c) shows these Areas of Ecological Significance
842 (blue) in context. In (a), black points indicate tracking data and yellow points indicate tagging locations
843 (location names in ¹⁶). In (b), black points indicate colony locations for the 14 colony-breeding species.
844 In (c), major oceanographic fronts (grey lines) are: Sub-Antarctic Front (SAF), Polar Front (PF) and
845 Southern Antarctic Circumpolar Current Front (SACCF).

846 [DOUBLE COLUMN]

847

848 **Figure 2. Fishing effort in the Southern Ocean.** (a) Map showing fishing effort (total fishing hours
849 between 2012 and 2016²⁹). Contour lines (white) indicate the Areas of Ecological Significance. (b)
850 Kernel density plot showing the distribution of values of fishing effort (zero values not shown) inside
851 (red) and outside (grey) Areas of Ecological Significance (AES). Two-tailed permutation tests (n =
852 1,098,226 grid cells) indicate a significant difference. (c) Proportion of cells inside and outside AES that
853 had some (>0 hours, yellow) or no (0 hours, purple) fishing effort.

854 [SINGLE COLUMN]

855

856 **Figure 3. Spatial protection of Southern Ocean Areas of Ecological Significance.** (a) Current
857 (orange polygons) and proposed (magenta polygons) Marine Protected Areas (MPAs) superimposed on
858 overall habitat importance. White contours denote Areas of Ecological Significance (AES), black lines
859 show national Exclusive Economic Zones, and the blue line shows the CCAMLR Convention Area. (b)
860 Area (million km²) in current (orange) and proposed (magenta) MPAs, and outside MPAs (blue). (c)
861 Area (million km²) inside and outside AES in national Exclusive Economic Zones, the CCAMLR
862 Convention Area and the international waters outside these two areas.

863 [DOUBLE COLUMN]

864

865 **Figure 4. Projected change in distribution of Areas of Ecological Significance (AES) under**
866 **RCP8.5.** a) Cells that were AES in the original results are shown as blue (remain as AES) or orange
867 (become non-AES in the future). The gradation from orange to blue shows the proportion of climate
868 models that indicate loss (orange) or retention (blue) of AES. Similarly, the gradation from white to
869 green shows the proportion of models that indicate non-AES cells will remain as non-AES (white) or
870 become AES (green). Orange and magenta polygons show current and proposed Marine Protected
871 Areas, respectively. b) Shows the percentage change in AES area according to the 8 different climate
872 models (black points), and the mean of these (red points). In boxplots, the box is from the 25th – 75th

873 percentile, and the whiskers extend to the smallest/largest value that is not further than 1.5 times the
874 interquartile range from the 25th/75th percentile.
875 [SINGLE COLUMN]

876 **EXTENDED DATA CAPTIONS**

877

878 **Extended Data 1. Overview of the modelling process.** a) shows how habitat importance for a given
879 life-history stage (e.g., chick-rearing) of a given species (e.g., king penguin [*Aptenodytes patagonicus*])
880 is calculated using two models (grey boxes): the habitat selection model and the accessibility model. b)
881 shows how these stage-specific, species-specific habitat importance predictions are combined to
882 calculate mean habitat importance for multiple species (e.g., king penguin and Antarctic fur seal
883 [*Arctocephalus gazella*]). Note that in the habitat accessibility model—Box 2 in (a)—distance to colony
884 can be weighted by relative colony size or not. The unweighted version is shown in this figure.

885

886 **Extended Data 2. Maps showing the 19 environmental covariates used to model marine predator**
887 **habitat selection in the Southern Ocean.** Grey lines indicate major oceanographic fronts.
888 Abbreviations, sources and units of measurement are defined in the Supplementary Information Table
889 S2.

890

891 **Extended Data 3. Habitat importance scores for 16 marine predator species in the Southern**
892 **Ocean.** The maps show predicted habitat importance for each species. Predictions for macaroni
893 (*Eudyptes chrysocome*) and royal penguins (*E. schlegeli*) are combined. Black circles show all known
894 colony locations for the 14 colony-breeding species, which we used to predict the models across the
895 whole Southern Ocean.

896

897 **Extended Data 4. Covariate importance.** Relative importance of 19 environmental variables used as
898 predictors in 40 boosted regression tree models of the habitat selection of Southern Ocean marine
899 predators. Higher variable relative importance values indicate that the variable has higher predictive
900 power. Points show the values for each model and boxplots (in grey, behind) show the distribution of
901 values. Variables are ordered (top to bottom) by decreasing median importance. Panels show the
902 results for three different species groups that were identified by hierarchical cluster analysis (see text
903 and Extended Data 7). Full covariate names are given in Supplementary Information Table S2. In
904 boxplots, the median is shown, the box is from the 25th – 75th percentile, and the whiskers extend to the
905 smallest/largest value that is not further than 1.5 times the interquartile range from the 25th/75th
906 percentile.

907

908 **Extended Data 5. Varied relationships between covariates and habitat selection across species.**
909 Scatterplot smooths (black lines) of the relationship between predictions of the species habitat selection
910 models (boosted regression trees) (vertical axis) and the values of covariates used as predictors in our
911 boosted regression tree models (horizontal axis). The smooths were drawn by fitting generalized
912 additive models for large datasets with a thin plate regression spline basis, since loess smoothing was
913 not computationally feasible. Full covariate names and units are given in Supplementary Information

914 Table S2. Higher habitat selection values indicate higher probabilities of use, irrespective of availability
915 in this case. A smooth is shown for each species. Since each species had 1–5 predictions, for different
916 life-history stages, we took the mean habitat selection estimate per cell for each species. Rug marks on
917 the horizontal axis indicate the distributions of the data points.

918

919 **Extended Data 6. Potential environmental stressors in the Southern Ocean.** Maps (a-c) showing
920 the change (mean in 1987–1998 compared to mean in 2007–2017) in (a) sea-ice duration (days), (b)
921 sea surface temperature (SST, °C), and (c) wind speed (m/s). Contour lines (black) indicate the Areas
922 of Ecological Significance. Kernel density plots (d-f) show the distribution of values of each of a-c inside
923 (red) and outside (grey) Areas of Ecological Significance (AES). Horizontal lines represent zero change.
924 Two-tailed permutation tests indicate significant differences in each case; the number of grid cells
925 included in the test is given in each case (n).

926

927 **Extended Data 7. Change in AES distribution under RCP4.5.** a) Cells that were AES in the original
928 results are shown as blue (remain as AES) or orange (become non-AES in the future). The gradation
929 from orange to blue shows the proportion of climate models that indicate loss (orange) or retention
930 (blue) of AES. Similarly, the gradation from white to green shows the proportion of models that indicate
931 non-AES cells will remain as non-AES (white) or become AES (green). Orange and magenta polygons
932 show current and proposed Marine Protected Areas, respectively. b) Shows the percentage change in
933 AES area according to the 8 different climate models (black points), and the mean of these (red points).

934

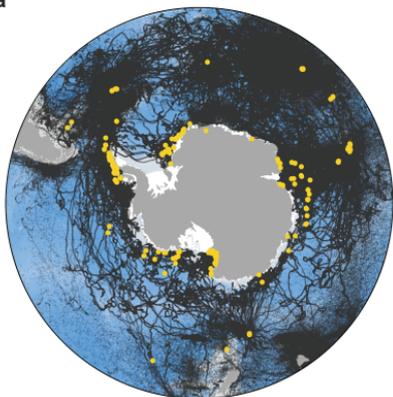
935 **Extended Data 8. Comparison of a) unweighted and b) weighted overall habitat importance.** a)
936 shows overall habitat importance calculated without accounting for colony sizes, while b) shows overall
937 habitat importance if colony sizes are taken into account. See Methods and Supplementary Information
938 for details. Black points indicate colony locations for the 14 colony-breeding species, and the white
939 contours indicate the Areas of Ecological Significance.

940

941 **Extended Data 9. Dendrogram of hierarchical cluster analysis showing species groups in the**
942 **dataset.** We performed UPGMA (Unweighted Pair Group Method with Arithmetic Mean) hierarchical
943 cluster analysis on the Manhattan distance among species, calculated from the habitat importance
944 scores. The results show two clear species groups (Antarctic - blue, and sub-Antarctic - magenta).
945 Humpback whales and southern elephant seals (orange) did not fall into either group and we assigned
946 them to both groups for subsequent analyses. The cophenetic correlation coefficient between the
947 distance matrix and the dendrogram was 0.86, which means the dendrogram is a good representation
948 of the Manhattan distance values among the species. Values can range from 0 (no correlation) to 1
949 (perfect correlation).

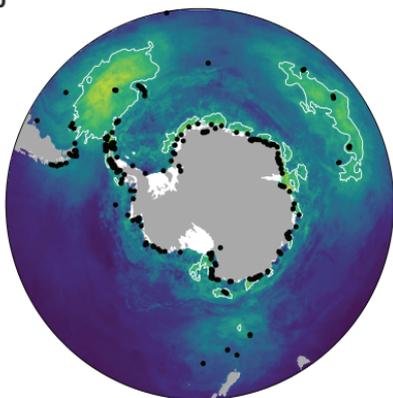
950

951 **Extended Data 10. Mean habitat importance of a) Antarctic and b) sub-Antarctic species.** To
952 account for regional differences in species richness we defined two species groups (see Methods and
953 Extended Data 5) and calculated mean habitat importance for these two groups separately. These two
954 mean habitat importance layers—(a) and (b)—were then combined into a single overall habitat
955 importance layer by choosing the maximum value in each cell. Black points indicate the colony locations
956 of colony-breeding species in each species group.

a

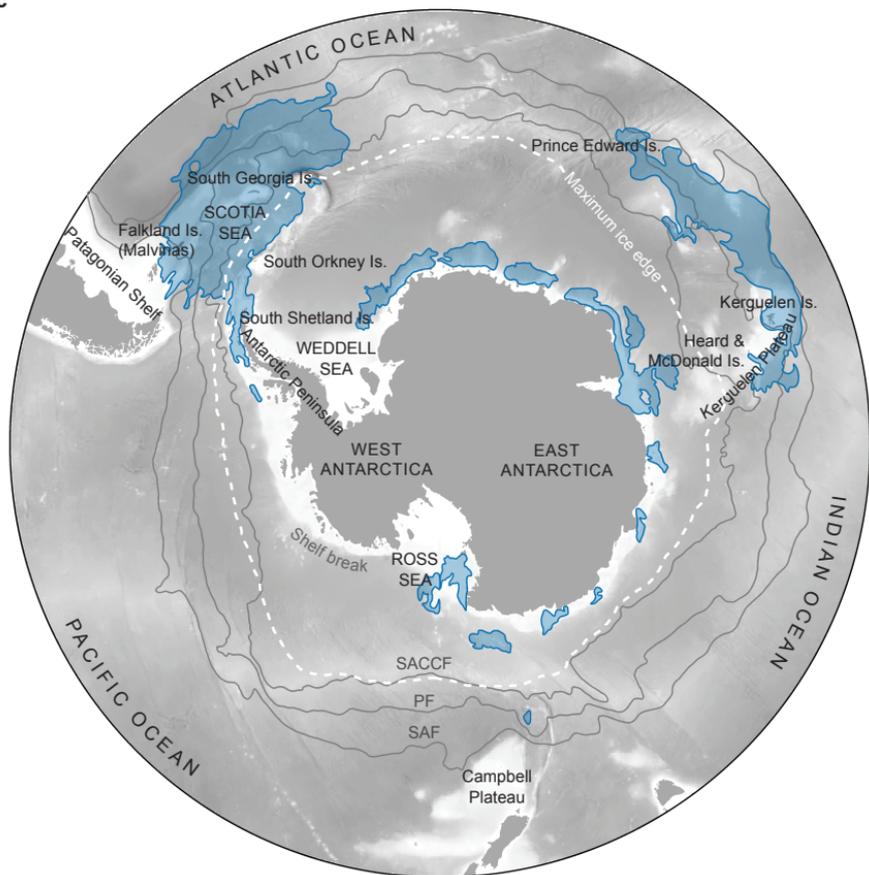
Depth (m)

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b

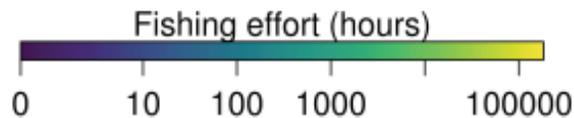
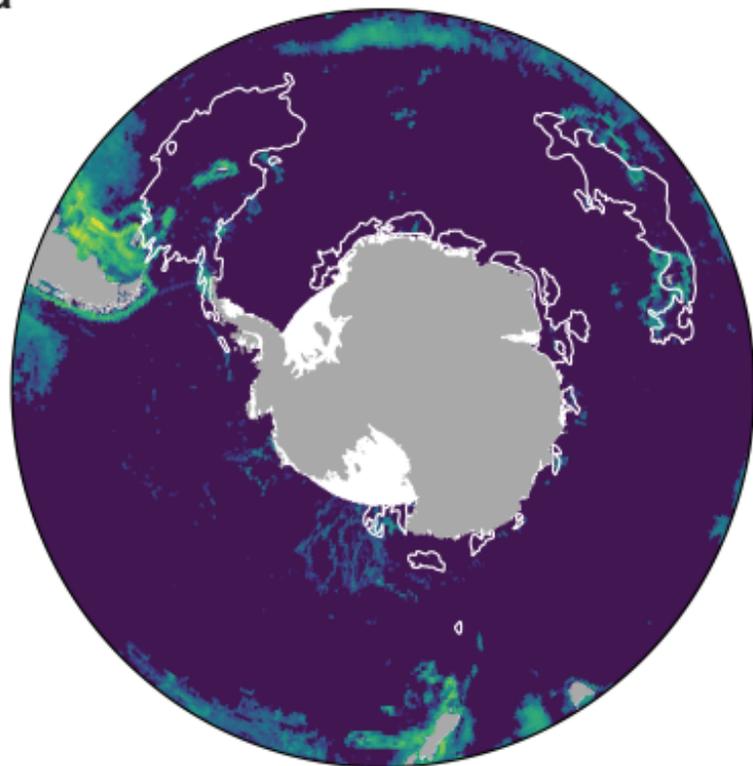
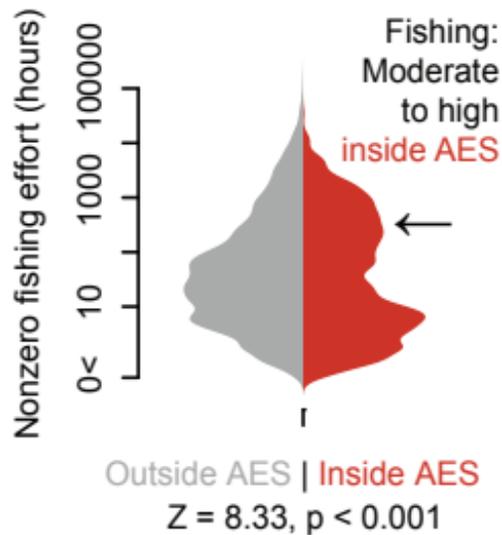
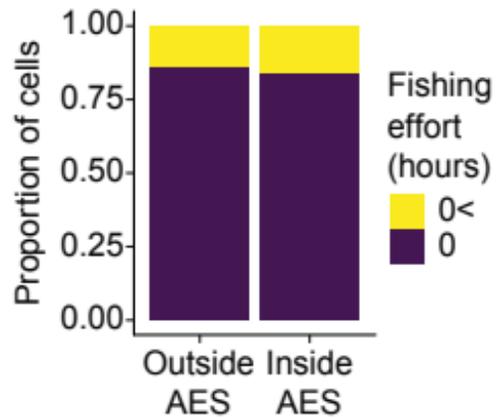
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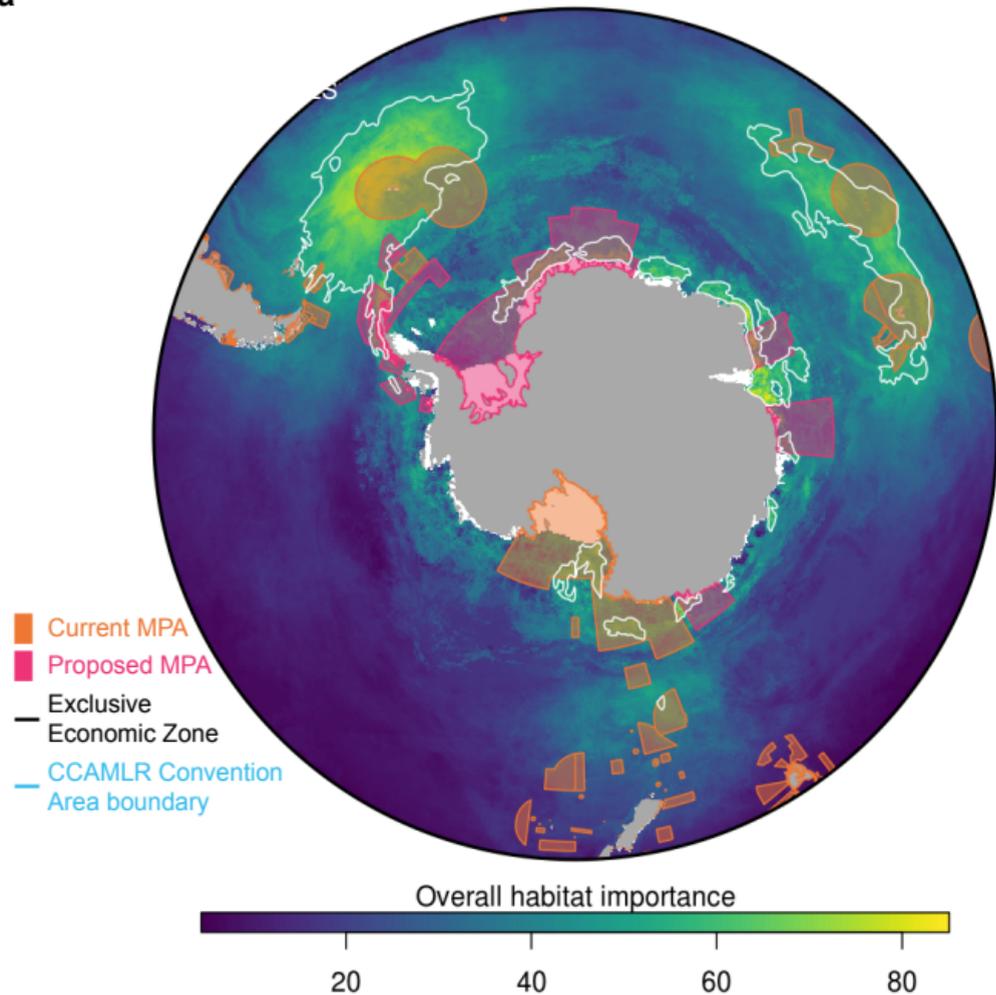
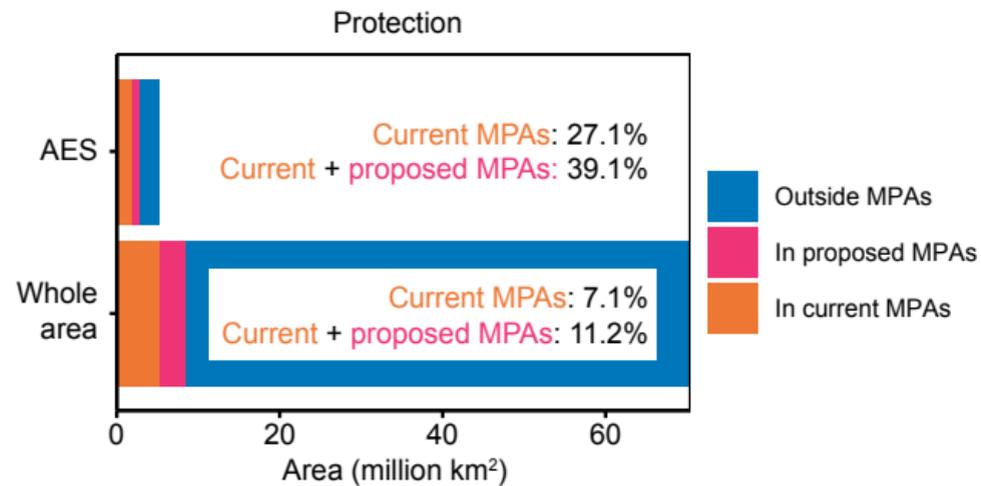
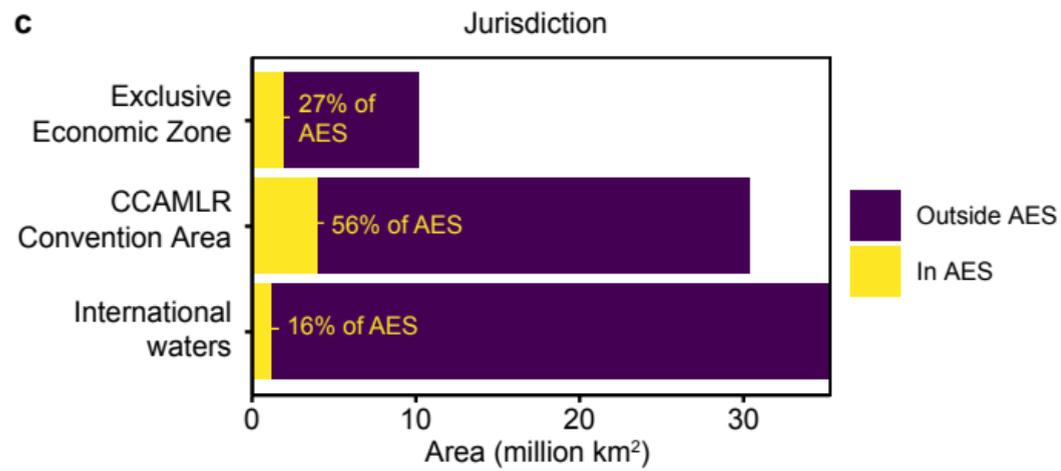
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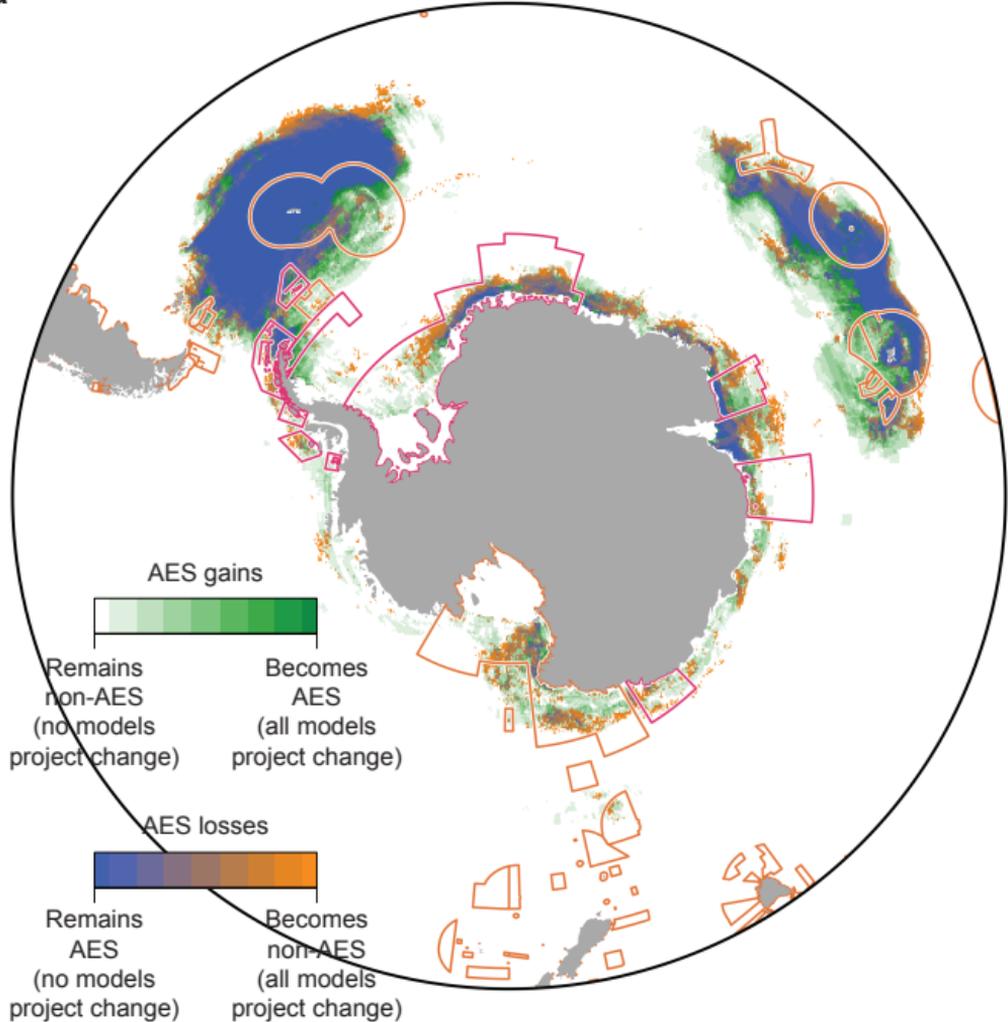
c

Depth (m)

-6000 -4000 -2000 0

a**b****c**

a**b****c**

a**b**