HMMoce: An R package for improved geolocation of archival-tagged fishes using a hidden Markov method

³ Camrin D. Braun^{1,2*}, Benjamin Galuardi^{3,4}, Simon R. Thorrold²

4 1. Massachusetts Institute of Technology-Woods Hole Oceanographic Institution Joint Program in Oceanog-

⁵ raphy/Applied Ocean Science and Engineering, Cambridge, MA 02139

6 2. Biology Department, Woods Hole Oceanographic Institution, Woods Hole, MA 02543

⁷ 3. School of Marine Science and Technology, University of Massachusetts Dartmouth, Fairhaven, MA
 02719

Greater Atlantic Regional Fisheries Office, National Marine Fisheries Service, National Oceanic and
 Atmospheric Administration, Gloucester, MA 01930

¹¹ 1 Summary

1. Electronic tagging of marine fishes is commonly achieved with archival tags that rely on light levels 12 and sea surface temperatures to retrospectively estimate movements. However, methodological issues 13 associated with light-level geolocation have constrained meaningful inference to species where it is 14 possible to accurately estimate time of sunrise and sunset. Most studies have largely ignored the 15 oceanographic profiles collected by the tag as a potential way to refine light-level geolocation estimates. 16 2. Open-source oceanographic measurements and outputs from high-resolution models are increasingly 17 available and accessible. Temperature and depth profiles recorded by electronic tags can be integrated 18 with these empirical data and model outputs to construct likelihoods and improve geolocation estimates. 19 3. The R package HMMoce leverages available tag and oceanographic data to improve position estimates 20 derived from electronic tags using a hidden Markov approach. We illustrate the use of the model and 21 test its performance using example blue and make shark archival tag data. Model results were validated 22 using independent, known tracks and compared to results from other geolocation approaches. 23

4. HMMoce exhibited as much as 6-fold improvement in pointwise error as compared to traditional light-level
 geolocation approaches. The results demonstrated the general applicability of HMMoce to marine animals,
 particularly those that do not frequent surface waters during crepuscular periods.

1

27 Key words: satellite telemetry; movement ecology; oceanography; state-space model; behavioral state

²⁸ * Correspondence author. Email: cbraun@whoi.edu

²⁹ 2 Introduction

Electronic archival tags have been widely adopted by ecologists to track movements of wide-ranging species 30 that are difficult to monitor using other techniques. In ocean environments, implanted archival and pop-up 31 satellite archival transmitting (PSAT) tags have proved particularly valuable in the study of life history 32 patterns (e.g. Thorrold et al., 2014), biophysical interactions and habitat use (e.g. Braun et al., 2015b; Lam 33 et al., 2014), horizontal and vertical movements (e.g. Braun et al., 2014; Lam et al., 2016; Werry et al., 2014), 34 and the spatial structure of populations (Skomal et al., 2009; Galuardi et al., 2010; Galuardi and Lam, 2014) 35 in a number of commercially important fishes (Block et al., 2011) and species of conservation concern (Braun 36 et al., 2015a). Yet, tracks provided by electronic tags that rely on light-level geolocation often exhibit large 37 error in daily position estimates (Musyl et al., 2011; Braun et al., 2015b) that may hinder inferences drawn 38 from the tag data. Approaches that provide more certainty in movement estimates derived from light level 39 data (Galuardi and Lam, 2014; Luo et al., 2015) would increase the power of ecological hypotheses tested 40 using tag data. 41

Electronic archival tags typically use light levels to estimate position when it is not possible for the tag to 42 interrogate geo-location satellites (Sibert et al., 2003; Nielsen and Sibert, 2007). Accuracy of geolocation using 43 light levels, however, is limited (\pm 100-200 km; ~10,000 km²) even for surface-oriented species where good light data is available (Wilson et al., 2007; Braun et al., 2015b). While several studies have incorporated ancillary 45 data, including sea surface temperature (Smith and Goodman, 1986; Lam et al., 2010), tidal fluctuation 46 (Pedersen et al., 2008) or ocean heat content (Luo et al., 2015) to help improve geolocation estimates, only 47 one used all data collected from archival tags within a rigorous statistical framework to improve geolocation 48 estimates (Summer et al., 2009). Although excursions from the photic zone, including diel vertical migration 49 (Neilson et al., 2009) and extended mesopelagic occupation (Skomal et al., 2009) may render light geolocation 50 impossible, the depth-temperature profiles recorded by the tags provide diagnostic oceanographic signatures 51 that can be leveraged to help constrain position (Skomal et al., 2009; Aarestrup et al., 2009). 52

⁵³ Hidden Markov Models (HMMs) have gained popularity in recent years as a tool for analyzing animal
⁵⁴ movement data and have been applied to understand movements of a number of organisms (Holzmann
⁵⁵ et al., 2006; Thygesen et al., 2009; Pedersen et al., 2011). Much of the progress in ocean environments

stems from a HMM used to track cod in the North Sea using tidal information (Pedersen et al., 2008). The 56 approach combined a number of desirable features, including inference about the underlying behavioral 57 state of the animal, mobilization of oceanographic data in a spatial likelihood framework (Nielsen et al., 58 2006), and later incorporated formal treatment of barriers to movement (Pedersen et al., 2011). Generally, 59 the Bayesian HMM approach uses a model of animal movements (e.g. Brownian motion) and a model or 60 observations of the environment (e.g. in situ oceanography) to estimate the posterior distribution of the 61 state (e.g. animal position and behavior). Several R packages exist for analyzing movement data with HMMs, 62 including ctmm (Calabrese et al., 2016) and moveHMM (Michelot et al., 2016), but none are designed for 63 geolocating marine fishes with archival tag data. An electronic tag manufacturer (Wildlife Computers, Inc.) 64 recently updated their proprietary software (GPE3) to geolocate archival tag data based on light levels and 65 sea surface temperature (SST) in a HMM framework following Pedersen et al. (2008). However, GPE3 is 66 limited by a lack of behavior state-switching dynamics and does not include functionality for non-surface 67 oriented species. GPE3 is also proprietary software that cannot be modified by the user and is limited to 68 tags built by Wildlife Computers. 69

Our primary objective was to build an analysis toolkit to improve geolocation estimates from electronic 70 archival tags deployed on marine animals that alleviates many of the limitations imposed by previous 71 approaches. The new R package HMMoce uses available electronic tag data and oceanographic data mined 72 from ocean observing system portals to estimate animal movements, behavior, and residency from uncertain 73 and temporally correlated movement data. We modify and expand a hidden Markov approach (Thygesen 74 et al., 2009; Pedersen et al., 2008, 2011) that, in addition to estimating animal movements, allows behavior 75 state estimation and provides information about the posterior distribution of the modeled states that can 76 be used as a residency metric (Pedersen et al., 2011). The modeling framework we developed is sufficiently 77 flexible to accommodate other tag types and animal movement questions, can be applied in any geographic 78 location, and benefits from the transparency of a widely-used open source platform. Here we describe the 79 model framework and demonstrate its applicability using example data. For specific details on package use 80 and functions and a full tutorial with an example dataset, please refer to the package and its accompanying 81 vignette, available on CRAN. 82

3 Overview of HMMoce

⁸⁴ 3.1 Model formulation

We present a process-based approach to estimate residency and behavior from movement data collected with 85 electronic archival tags. The logic of this approach involves calculating gridded observation likelihoods at 86 each time point based on tag and environmental data, forming the state-space model, estimating model 87 parameters and model selection and interpretation. The application of grids to explicitly resolve space is a 88 key component that allows state estimation (location and behavior, in this case) to be supplemented by or 89 based entirely on environmental data (e.g. temperature at depth). The details of the discretized grid HMM 90 approach are thoroughly explained elsewhere (e.g. Thygesen et al., 2009; Pedersen et al., 2011). A detailed 91 methodology for our approach can be found in the supplement. 92

Briefly, observation-based likelihoods (Eq. S1) were derived from in situ SST (Eq. S2), light-based longitude 93 and depth-temperature profile data (Eqs. S3, S4, S5) collected by the tags using five separate likelihood 94 calculations: 1) An SST likelihood (Eq. S2) was generated for tag-based SST values integrated according 95 to an error term $(\pm 1\%)$, based on tag sensor accuracy) and compared to remotely-sensed SST from daily, 96 optimally-interpolated SST fields (OI-SST, 0.25° resolution; Banzon et al., 2016). 2) Light-based longitude 97 likelihood was derived using estimates of longitude from GPE2 software (Wildlife Computers, Inc.), which 98 facilitated visual checking of light curves. Depth-temperature profiles recorded by the tag were compared 99 to 3) monthly climatological mean depth-temperature data from the World Ocean Atlas 2013 (WOA, 0.25° 100 resolution; Locarnini et al., 2013) and 4) daily reanalysis model depth-temperature products from the HYbrid 101 Coordinate Ocean Model (HYCOM, 0.08° resolution; Chassignet et al., 2007) at standard depth levels 102 available in these products (Eq. S5). Individual likelihood surfaces for each depth level were then combined 103 for an overall profile likelihood at that time point (Eq. S6). 5) Ocean Heat Content (OHC, Eq. S3) was 104 obtained by integrating the heat content of the water column above the minimum daily temperature recorded 105 by the tag for both the tag profiles and HYCOM fields (Eq. S4; Luo et al., 2015). Start and end locations 106 were considered known in all cases and model runs. 107

The resulting observation likelihoods (in various combinations; Eq. S1) were used in a two-step Bayesian state-space approach to estimate the posterior distribution of the state (in this case, a joint probability distribution of location and behavior at each time point). Probability distributions were first calculated forward in time using alternating time and data updates of the current state estimate using a HMM filter (for a detailed methodology of the HMM filter see Appendix 2 in Pedersen et al., 2011). The filter recursions ¹¹³ also returned a likelihood measure indicating how well the model fit the data, which facilitated calculating ¹¹⁴ model parameters (e.g. behavior state-switching probabilities). In Bayesian statistics, the maximum a priori ¹¹⁵ (MAP) estimate of the model parameters is typically used to calculate the posteriors; however, in practice, ¹¹⁶ ample a priori information is rarely available and maximum likelihood (ML) estimates are often very similar ¹¹⁷ to MAP estimates (Jonsen et al., 2005). Thus, we implemented recent advances by Woillez et al. (2016) that ¹¹⁸ further exploited the discretization of space in this model by employing a joint ML estimation of all model ¹¹⁹ parameters using an iterative Expectation-Maximization framework (Supp. 1.4.1).

Model selection in this context would typically use Bayesian Information criterion (BIC), but this approach requires approximation that imposes restrictions on the priors. Instead, we used Akaike's Information criterion (AIC) for model selection following Pedersen et al. (2011). The HMM smoother recursion was the final step that worked backwards in time using filtered state estimates and all available observation data to determine smoothed state estimates. This step provided the time marginal of the probability distributions based on observations (posterior distributions).

Results from the final smoothing step represent the posterior distribution of each state over time. Distributions 126 are summed for each behavior state and time step to determine the most likely behavior state through time. 127 HMMoce can calculate the mean or mode of the posterior distribution grid, at each time step, to estimate the 128 animal's position. The posteriors can be further analyzed for additional inference including uncertainty and 129 residency. A residency distribution (RD) is conceptually similar to the utilization distribution (UD), but 130 the concept of UD (and other space-use metrics) is often vaguely defined (Royle and Dorazio, 2008). In this 131 case, RD is interpreted as the estimate of the time spent in a given space within a time interval (see Eq. 5 in 132 Pedersen et al., 2011). 133

¹³⁴ 3.2 Computational improvements and requirements

While the basic framework of HMMoce was based on previous work (Pedersen et al., 2008; Thygesen et al., 135 2009; Pedersen et al., 2011), several improvements were made to accommodate user needs. We focused several 136 enhancements on improving computation efficiency, which was a limitation of previous techniques (SPHMM 137 code for R; Pedersen et al., 2011). Image processing routines replaced sparse matrix convolution yielding 138 orders of magnitude improvements in computation time, particularly for large, high-resolution grids that 139 characterize geolocation approaches for highly migratory species. In addition, all likelihood routines (except 140 simple light-based likelihood calculations) were parallelized, yielding marked performance improvements, 141 particularly for likelihoods comparing 3D depth-temperature profiles to high-resolution 3D HYCOM grids. 142

Despite these improvements, HMMoce remains relatively computationally intensive; however, cloud computing is becoming more inexpensive and accessible to a broad user group. The HMMoce package includes a vignette demonstrating simple plug and play functionality for the model using Amazon Web Services's computational resources and an associated machine image containing RStudio Server and all the required dependencies for running HMMoce with user-provided tag data.

¹⁴⁸ 4 Case study: pelagic shark movements

To illustrate the application of HMMoce, we analyzed tag data from three blue sharks (*Prionace glauca*) and 149 one shortfin mako (Isurus oxyrinchus) that were double-tagged with satellite-linked radio telemetry tags 150 (Wildlife Computers finmount SPOT5 tags) and PSAT tags (Table 1). Full tagging methods are provided in 151 the supplement. We considered the resulting Argos-based tracks as "known" because errors on geolocation 152 estimates from the SPOT tags are much lower (typically < 10 km; Witt et al., 2010; Patterson et al., 2010) 153 than PSAT-based outputs (> 50 km; Winship et al., 2012). The PSAT tags were deployed for an average of 154 150 days (range 107-180) in the northwest Atlantic with overall movements of 5403-12122 km. The PSAT 155 data contained depth-temperature profiles for 53-72% of days at liberty and SPOT locations were recorded 156 for 72-96% of deployment days (Table 1). 157

Blue sharks made frequent dives to the mesopelagic zone (~600-800m, max 680-1688m) but also frequented 158 the surface-air interface where the PSAT tags collected good quality light and SST data (94-100% deployment 159 days with light, 82-92% SST)(Fig. 1). The make occupied a restricted area (~200 km latitudinal distance) 160 near Cape Hatteras during the winter months and spent relatively little far from the edge of the continental 161 shelf compared to the more nomadic blue sharks. The make also had high quality light and SST data (96%162 and 69%, respectively) while regularly diving shallower than the blue sharks (~400m). Consistent exposure of 163 the dorsal fin allowed the SPOT tag to acquire Argos positions throughout the duration of each deployment 164 (Table 1). 165

We calculated movements of the sharks from PSAT tag data using three modeling approaches that are currently available (Ukfsst, Trackit, GPE3) and HMMoce (Supp. 1.6). Results for the mako are shown in the main text (Fig. 2), and blue shark figures are provided in the supplement (Figs. S2, S3, S4). In general, HMMoce and GPE3 produced the most accurate tracks while those from Ukfsst and Trackit were often unrealistic with errors as high as >1300 km (Table 2). For 3 of 4 individuals, HMMoce tracks had the lowest pointwise error and correspondingly lowest root-mean-square error (RMSE) values. For the fourth individual (blue shark 141259), the mean error and RMSE in latitude for GPE3 ouput was lower than HMMoce, which had

a lower RMSE in longitude. The traditional approaches (light only, Trackit; light and SST, Ukfsst) vielded 173 much larger error than HMMoce in all cases and only one Trackit output (blue shark 141254 without SST) 174 exhibited marginally smaller error than GPE3 (with SST). In 3 of 4 cases, HMMoce demonstrated the best 175 fitting model by leveraging either OHC (n=1) or HYCOM profiles (n=2) (Table 2) in addition to light-based 176 longitude and SST data used in the other geolocation approaches. The movements of blue shark 141259, in 177 which the HMMoce model did not use profile-based observations to build the final estimated track, included 178 time in both dynamic Gulf Stream water and the more homogenous Sargasso Sea. It proved difficult in 179 both areas to match water column profiles recorded by the tag (or integrated OHC) with an accurate and 180 constrained position in the climatological (WOA) or model-based (HYCOM) oceanographic data (Fig. S5). 181

While HMMoce was designed to improve geolocation estimates for all tagged marine organisms, the main 182 impetus for the work was to fulfill a need for improving track estimates in cases where light and SST data 183 were lacking due to minimal surface occupation. We tested the ability of HMMoce to recover accurate tracks 184 with only limited light-level data by randomly removing (using sample in base R, without replacement) 75% 185 and 50% of deployment days with adequate light and SST data, respectively, from the shark PSAT data while 186 keeping the depth-temperature profile data for these days. The removals approximated PSAT data quality 187 typical of swordfish tag deployments in the Atlantic Ocean due to crepuscular diving behavior and light 188 avoidance (Braun et al., 2015a; Neilson et al., 2009). The data removal experiment (Fig. 1) demonstrated 189 the power of incorporating the depth dimension in likelihood calculations when light and/or SST data is 190 poor. In all 4 cases, HMMoce estimated tracks with lower mean error than corresponding GPE3 results (Table 191 2), but model selection favored including depth-temperature profile information in only 2 of 4 final models. 192 Error in the removal experiment for HMMoce was only marginally higher as compared to the full dataset for 3 193 of 4 individuals (Table 2). 194

Both GPE3 and HMMoce provide estimated residency distributions (RD; a form of utilization distribution) 195 (Pedersen et al., 2011). However, only HMMoce incorporates a state-switching component that facilitates 196 explicit modeling of distinct animal behaviors (Fig. 3). The state-switching is governed by movement kernels 197 (based on speed) and probability of switching between states is calculated by the EM algorithm (Table S1). 198 For the make, the RDs indicated areas of core use focused largely where resident behavior was most probable. 199 The RD for the migratory state showed the offshore movement to the SE into the Gulf Stream region before 200 the fish returned to the shelf break and moved SW toward Cape Hatteras. The most notable features of the 201 migratory RD are the overlap areas where the fish transitioned from migratory to resident behaviors (Fig. 4). 202

203 5 Conclusions

We present a flexible, customizable and transparent HMM framework that may be applied to nearly any marine species utilizing electronic archival tags through a novel use of oceanographic data. Tests of the model demonstrated that HMMoce is a valuable tool for estimating movements from low quality PSAT data through consideration of the vertical structure of the water column in the state estimation. This can be especially beneficial for tag data that is lacking adequate light-level data or other measurements.

For further development, we anticipate several improvements to the HMMoce package. Current priorities include support for other tag types, direct versus derived use of light data, and additional algorithms (e.g. Viterbi) to calculate the most probable track (Pedersen et al., 2011). Behavior state estimation could be expanded to include advection or modified to update probability with respect to environmental data (Patterson et al., 2009).

We anticipate user feedback will help prioritize further improvements, and we welcome bug reports, feedback, and suggestions for the development of HMMoce via our Github repository https://github.com/camrinbraun/HMMoce. Additional usage information, including an example dataset and a tutorial for using HMMoce on Amazon Web Services, can be found by installing HMMoce in R (install.packages("HMMoce")) and accessing the package vignette.

²¹⁹ 6 Acknowledgements

We thank P. Gaube, M. Kaplan, D. McGillicuddy and A. Solow for helpful discussions during model development. We thank M. Pedersen for inspiration and previous work on model development. This work was funded by awards to C. Braun from the Martin Family Society of Fellows for Sustainability Fellowship at the Massachusetts Institute of Technology, the Grassle Fellowship and Ocean Venture Fund at the Woods Hole Oceanographic Institution, and the NASA Earth and Space Science Fellowship. Computational support was provided by the AWS Cloud Credits for Research program.

Funding for the development of HYCOM has been provided by the National Ocean Partnership Program and the Office of Naval Research. Data assimilative products using HYCOM are funded by the U.S. Navy. Computer time was made available by the DoD High Performance Computing Modernization Program. The output is publicly available at http://hycom.org.

²³⁰ All tagging protocols were performed in accordance with the Woods Hole Oceanographic Institution's Animal

²³¹ Care and Use Committee (IACUC) protocol #BI23112.

²³² 7 Data accessibility

All code mentioned here is available in the HMMoce package for R hosted on CRAN. The development version of the package is available on GitHub at https://github.com/camrinbraun/HMMoce. Supporting data (e.g. shark satellite tag data) is distributed with the package from both sources.

236 8 Author contributions

²³⁷ CDB and BG conceived the project and developed the package. CDB and SRT collected the data. All
 ²³⁸ co-authors wrote the manuscript, assisted with edits and approve publication.

²³⁹ 9 Figure captions

Figure 1. Example blue shark data demonstrating the deployment days with [A] light, [B] sea surface temperature and [C] depth-temperature profile data used as the observation portion of the HMM. Full (F) and removal (R) datasets for light and SST are shown [A,B].

Figure 2. Calculated tracks for make shark 141257 using the 4 different geolocation approaches (Ukfsst, purple; Trackit, blue; GPE3, green; HMMoce, yellow) compared to the "known" Argos-based track (red, black crosses). Latitudinal and longitudinal estimates through time are shown in panels B and C, respectively. Lines appear broken when a resulting track is missing daily data.

Figure 3. Movements (A) and behavior (B) calculated using HMMoce for mako 141257. The tagged individual is considered resident where probability of residency is greater than 0.5 (grey points and bars in panels A and B, respectively). Green and red points indicate tag and pop-up location respectively. Black line follows predicted daily locations of tagged shark.

Figure 4. Residency distributions for the overall HMMoce modeled movements (A) and for individual behavior states (B, C). Shaded circles indicate residency behavior, white circles indicate migratory behavior, green triangle is tagging location and red triangle is pop-up location. Residency distributions show the expected proportion of time spent in various grid cells over the course of tag deployment.

Table 1: Tagging summary for double-tagged blue (BSH) and shortfin mako (MKO) sharks used in this study. PDT, Light, SST and SPOT = percent of deployment period with depth-temperature profile (PDT), light and sea surface temperature (SST) data from the PSAT tag and percent of deployment period with Argos-based positions (SPOT), respectively.

Species	Tag ID	Start Date	End Date	Duration (d)	PDT (%)	Light $(\%)$	SST $(\%)$	SPOT $(\%)$
BSH	141254	2015-10-21	2016-02-05	107	72	100	92	96
BSH	141256	2015 - 10 - 13	2016-02-24	134	66	94	88	87
BSH	141259	2015 - 10 - 13	2016-04-10	180	53	94	82	85
MKO	141257	2015 - 10 - 15	2016-04-12	180	58	96	69	72

Table 2: Validation metrics for double-tagged blue (BSH) and shortfin mako (MKO) shark tracks estimated using HMMoce, GPE3, Trackit (TI) and Ukfsst. Reported error values (mean, sd, median, range) are pointwise distance calculations (mean great circle distance) from known positions (km). Root-mean-square errors (RMSE) are Lat, Long (degrees). HMMoce.r and GPE3.r indicate fit metrics for data removal experiments in which 75% of daily light and 50% of daily SST data was randomly removed. Input indicates input data type: light (L), SST (S), ocean heat content (O), World Ocean Atlas profiles (W) and HYCOM profiles (H). All runs were performed on a 0.08° grid with fixed migratory speed of 2 m/s (except 141259 used 4 m/s).

Species	Tag ID	Type	Mean~(SD)	Median	Range	RMSE	Input
BSH	141254	HMMoce	117.4(96.7)	92.4	0.5-443.6	1.21, 0.81	LSO
		GPE3	175.8(117.1)	164.3	3.2 - 424.7	1.4, 1.64	LS
		TI	162.3(71.6)	158.2	1 - 328.2	0.97, 1.65	\mathbf{L}
		KF	179.5(99.5)	178.5	1 - 435.2	1.29, 1.24	\mathbf{L}
		HMMoce.r	131.2(96.2)	101.9	0.5 - 440.5	1.23, 1.01	LS
		GPE3.r	157.6(100.6)	143.5	1.4-408.9	1.25, 1.44	LS
BSH	141256	HMMoce	83.8(63)	63.7	4.9-297.4	0.52, 0.93	LSH
		GPE3	84.9(68.8)	66.9	5.9 - 345	0.66, 0.89	LS
		TI	474.2(244.1)	459.9	0-854.3	1.98, 4.84	L
		KF	192.7(152.4)	172.6	0-699.8	1.35, 0.65	L
		HMMoce.r	93.4(57.8)	79.1	4.2-286	0.59, 0.92	LSH
		GPE3.r	423.5(432)	197.8	2.1 - 1394	4.25, 3.96	LS
BSH	141259	HMMoce	179.4(126)	150.3	4.4 - 575.2	1.79, 1	LS
		GPE3	158.1(109.6)	139.5	4.9 - 434.5	1.44, 1.17	LS
		TI	367.5(239.1)	291.4	2.4 - 861.5	3.3, 2.36	L
		\mathbf{KF}	245.8(225.5)	194.5	1.7 - 1078.7	2.31, 0.88	L
		HMMoce.r	183.3(132.2)	140.5	4.4 - 560.5	1.9, 0.88	LS
		GPE3.r	198(129.5)	162.0	6.1 - 625.8	1.61, 1.77	LS
MKO	141257	HMMoce	99.8(90.7)	66.8	3.8 - 426.9	0.92,0.99	LSH
		GPE3	151.1(161.1)	93.0	6.8 - 675.2	0.65, 2.38	LS
		TI	462.6(347.7)	320.5	0-1332.7	4.6, 2.79	L
		\mathbf{KF}	220.4(151.2)	173.7	0-614.6	1.3, 1.32	L
		HMMoce.r	157.9(128.2)	119.1	3.8 - 494.4	1.05, 1.92	LSH
		GPE3.r	182.3(171.8)	136.4	0.3-711.2	0.88, 2.62	LS

255 **References**

- ²⁵⁶ Aarestrup, K., Okland, F., Hansen, M. M., Righton, D., Gargan, P., Castonguay, M., Bernatchez, L., Howey,
- ²⁵⁷ P., Sparholt, H., Pedersen, M. I., and McKinley, R. S. (2009). Oceanic Spawning Migration of the European
- ²⁵⁸ Eel (Anguilla anguilla). *Science*, 325(5948):1660–1660.
- ²⁵⁹ Banzon, V., Smith, T. M., Mike Chin, T., Liu, C., and Hankins, W. (2016). A long-term record of blended
- satellite and in situ sea-surface temperature for climate monitoring, modeling and environmental studies.
- Earth System Science Data, 8(1):165-176.
- ²⁶² Block, B. A., Jonsen, I. D., Jorgensen, S. J., Winship, A. J., Shaffer, S. A., Bograd, S. J., Hazen, E. L., Foley,
 ²⁶³ D. G., Breed, G. A., and Harrison, A. L. (2011). Tracking apex marine predator movements in a dynamic
 ²⁶⁴ ocean. Nature, 475(7354):86–90.
- Braun, C. D., Kaplan, M. B., Horodysky, A. Z., and Llopiz, J. K. (2015a). Satellite telemetry reveals physical
 processes driving billfish behavior. *Animal Biotelemetry*, 3(1):2.
- Braun, C. D., Skomal, G. B., Thorrold, S. R., and Berumen, M. L. (2014). Diving Behavior of the Reef
 Manta Ray Links Coral Reefs with Adjacent Deep Pelagic Habitats. *PLoS One*, 9(2):e88170.
- Braun, C. D., Skomal, G. B., Thorrold, S. R., and Berumen, M. L. (2015b). Movements of the reef manta ray
 (Manta alfredi) in the Red Sea using satellite and acoustic telemetry. *Marine Biology*, 162(12):2351–2362.
- ²⁷¹ Calabrese, J. M., Fleming, C. H., Gurarie, E., and Freckleton, R. (2016). Ctmm: an R Package for Analyzing
- Animal Relocation Data As a Continuous-Time Stochastic Process. Methods in Ecology and Evolution,
 7(9):1124–1132.
- Chassignet, E. P., Hurlburt, H. E., Smedstad, O. M., Halliwell, G. R., Hogan, P. J., Wallcraft, A. J., Baraille,
 R., and Bleck, R. (2007). The HYCOM (HYbrid Coordinate Ocean Model) data assimilative system. *Journal of Marine Systems*, 65(1-4 SPEC. ISS.):60–83.
- Galuardi, B. and Lam, C. H. T. (2014). Chapter Nineteen Telemetry Analysis of Highly Migratory Species.
 pages 447–476.
- Galuardi, B., Royer, F., Golet, W., Logan, J., Neilson, J., and Lutcavage, M. (2010). Complex migration routes
 of Atlantic bluefin tuna (Thunnus thynnus) question current population structure paradigm. *Canadian*
- Journal of Fisheries and Aquatic Sciences, 67(6):966–976.
- Holzmann, H., Munk, A., Suster, M., and Zucchini, W. (2006). Hidden Markov models for circular and
 linear-circular time series. *Environmental and Ecological Statistics*, 13(3):325–347.

- Jonsen, I. D., Flemmings, J. M., and Myers, R. a. (2005). Robust State Space Modeling of Animal Movement Data. *Ecology*, 86(11):2874–2880.
- Lam, C. H., Galuardi, B., and Lutcavage, M. E. (2014). Movements and oceanographic associations of bigeye
 tuna (Thunnus obesus) in the Northwest Atlantic. *Canadian Journal of Fisheries and Aquatic Sciences*,
 1543(September 2013):1–15.
- Lam, C. H., Galuardi, B., Mendillo, A., Chandler, E., and Lutcavage, M. E. (2016). Sailfish migrations
- ²⁹⁰ connect productive coastal areas in the West Atlantic Ocean. *Scientific Reports*, 6(August):38163.
- Lam, C. H., Nielsen, A., and Sibert, J. R. (2010). Incorporating sea-surface temperature to the light-based
 geolocation model TrackIt. *Marine Ecology Progress Series*, 419:71–84.
- Locarnini, R. A., Mishonov, A. V., Antonov, J. I., Boyer, T. P., Garcia, H. E., Baranova, O. K., Zweng,
 M. M., Paver, C. R., Reagan, J. R., and Johnson, D. R. (2013). World Ocean Atlas 2013, Volume 1:
 Temperature. NOAA Atlas NESDIS, 73:40.
- Luo, J., Ault, J. S., Shay, L. K., Hoolihan, J. P., Prince, E. D., Brown, C. a., and Rooker, J. R. (2015). Ocean
 Heat Content Reveals Secrets of Fish Migrations. *Plos One*, 10(10):e0141101.
- Michelot, T., Langrock, R., and Patterson, T. A. (2016). moveHMM: An R package for the statistical
 modelling of animal movement data using hidden Markov models. *Methods in Ecology and Evolution*,
 7(11):1308–1315.
- Musyl, M. K., Domeier, M. L., Nasby-Lucas, N., Brill, R. W., McNaughton, L. M., Swimmer, J. Y., Lutcavage,
 M. S., Wilson, S. G., Galuardi, B., and Liddle, J. B. (2011). Performance of pop-up satellite archival tags.
 Marine Ecology Progress Series.
- Neilson, J. D., Smith, S., Royer, F., Paul, S. D., Porter, J. M., and Lutcavage, M. (2009). Investigations of
 horizontal movements of Atlantic swordfish using pop-up satellite archival tags. In *Tagging and tracking of marine animals with electronic devices*, pages 145–159. Springer.
- Nielsen, A., Bigelow, K. A., Musyl, M. K., and Sibert, J. R. (2006). Improving light-based geolocation by
 including sea surface temperature. *Fisheries Oceanography*, 15(4):314–325.
- ³⁰⁹ Nielsen, A. and Sibert, J. R. (2007). State–space model for light-based tracking of marine animals. *Canadian*
- Journal of Fisheries and Aquatic Sciences, 64(8):1055–1068.

321

- Patterson, T. A., Basson, M., Bravington, M. V., and Gunn, J. S. (2009). Classifying movement behaviour in 311
- relation to environmental conditions using hidden Markov models. Journal of Animal Ecology, 78(6):1113-312 1123.313
- Patterson, T. A., McConnell, B. J., Fedak, M. A., Bravington, M. V., and Hindell, M. A. (2010). Using 314
- GPS data to evaluate the accuracy of state-space methods for correction of Argos satellite telemetry error. 315

Ecology, 91(1):273-285.316

- Pedersen, M., Berg, C., Thygesen, U., Nielsen, a., and Madsen, H. (2011). Estimation methods for nonlinear 317 state-space models in ecology. Ecological Modelling, 222(8):1394–1400. 318
- Pedersen, M. W., Righton, D., Thygesen, U. H., Andersen, K. H., and Madsen, H. (2008). Geolocation of 319 North Sea cod (Gadus morhua) using hidden Markov models and behavioural switching. Canadian Journal 320 of Fisheries and Aquatic Sciences, 65(11):2367-2377.
- Royle, J. A. and Dorazio, R. M. (2008). Hierarchical modeling and inference in ecology: the analysis of data 322 from populations, metapopulations and communities. Academic Press. 323
- Sibert, J. R., Musyl, M. K., and Brill, R. W. (2003). Horizontal movements of bigeve tuna (Thunnus 324 obesus) near Hawaii determined by Kalman filter analysis of archival tagging data. Fisheries Oceanography, 325 12(3):141-151.326
- Skomal, G. B., Zeeman, S. I., Chisholm, J. H., Summers, E. L., Walsh, H. J., McMahon, K. W., and Thorrold, 327 S. R. (2009). Transequatorial migrations by basking sharks in the western Atlantic Ocean. Current Biology, 328 19(12):1019-1022. 329
- Smith, P. and Goodman, D. (1986). Determining fish movements from an" archival" tag: precision of 330 geographical positions made from a time series of swimming temperature and depth. US Department 331 of Commerce, National Oceanic and Atmospheric Administration, National Marine Fisheries Service, 332 Southwest Fisheries Center. 333
- Sumner, M. D., Wotherspoon, S. J., and Hindell, M. A. (2009). Bayesian estimation of animal movement 334 from archival and satellite tags. PLoS One, 4(10):e7324. 335
- Thorrold, S. R., Afonso, P., Fontes, J., Braun, C. D., Skomal, G. B., and Berumen, M. L. (2014). Extreme 336 diving behavior in devil rays link surface waters and the deep ocean. Nature Communications, 5(4274). 337
- Thygesen, U. H., Pedersen, M. W., and Madsen, H. (2009). Geolocating Fish Using Hidden Markov Models 338
- and Data Storage Tags. Tagging and Tracking of Marine Animals with Electronic Devices, 9:23-34. 339

- ³⁴⁰ Werry, J. M., Planes, S., Berumen, M. L., Lee, K. A., Braun, C. D., and Clua, E. (2014). Reef-Fidelity and
- Migration of Tiger Sharks, Galeocerdo cuvier, across the Coral Sea. *PLoS One*, 9(1):e83249.
- ³⁴² Wilson, S. G., Stewart, B. S., Polovina, J. J., Meekan, M. G., Stevens, J. D., and Galuardi, B. (2007). Accuracy
- and precision of archival tag data: a multiple-tagging study conducted on a whale shark (Rhincodon typus)

in the Indian Ocean. Fisheries Oceanography, 16(6):547-554.

- ³⁴⁵ Winship, A. J., Jorgensen, S. J., Shaffer, S. a., Jonsen, I. D., Robinson, P. W., Costa, D. P., and Block,
- B. a. (2012). State-space framework for estimating measurement error from double-tagging telemetry
- experiments. Methods in Ecology and Evolution, 3(2):291–302.
- Witt, M. J., Åkesson, S., Broderick, A. C., Coyne, M. S., Ellick, J., Formia, A., Hays, G. C., Luschi, P., Stroud,
- ³⁴⁹ S., and Godley, B. J. (2010). Assessing accuracy and utility of satellite-tracking data using Argos-linked
- ³⁵⁰ Fastloc-GPS. Animal behaviour, 80(3):571.
- ³⁵¹ Woillez, M., Fablet, R., Ngo, T. T., Lalire, M., Lazure, P., and de Pontual, H. (2016). A HMM-based model
- to geolocate pelagic fish from high-resolution individual temperature and depth histories: European sea
- bass as a case study. *Ecological Modelling*, 321:10–22.





Methods in Ecology and Evolution



Probability of resident

