

## **APPENDIX S2 | Details for models, priors, MCMC sampling, and results for all parameters from mark-recapture models**

### *RJMCMC selection of CJS model*

The *multimark* CJS framework uses a probit link for capture probability and apparent annual survival. We used the default uninformative priors for all parameters (Table S2.1; McClintock 2015). Each model was fitted with three MCMC chains of 500,000 iterations each, including a 25,000-sample burn-in, thinned to every 250 samples. We used the same number of iterations for the RJMCMC as the final sample size for each model, i.e., three chains of 1,900 iterations each for a total of 5,700 samples.

### *Estimation of survival*

We fitted the final CJS model with three parallel MCMC chains of 1,000,000 iterations each thinned to every 50 samples, including a 25,000-iteration burn-in, resulting in 58,500 samples.

### *Estimation of abundance*

We used the default uninformative priors for all parameters (Table S2.2; McClintock 2015 for more details). We fitted the model with three parallel MCMC chains of 1,000,000 iterations each thinned to every 100 samples, including a 250,000-iteration burn-in, resulting in 22,500 samples.

### *Estimation of annual rate of change through multiple imputation*

For each fixed value of  $\phi$ , we fitted the model with three MCMC chains, each including an adaptation phase of 500 samples, burn-in of 10,000 samples, and 20,000 iterations thinned at 100-sample intervals, resulting in 600 samples. This was repeated over 250 samples of  $\phi$ , resulting in 150,000 samples.

**TABLE S2.1** Summaries of priors and results for parameters estimated by final CJS model in *multimark*:  $\beta_\phi$  is probit-link apparent annual survival;  $\beta_{p,0}$  is the probit-link intercept of annual capture probability  $p$ ;  $\beta_{p,e}$  is the probit-link slope of capture probability with standardized Beaufort- and season-adjusted effort index (see Methods);  $\sigma_{p,z}^2$

is the variance of probit-link individual random effects in capture probability;  $\alpha$  is probability of simultaneous left- and right-sided detections, given both types encountered;  $\tau$  is the sum of the conditional probabilities of left- and right-sided encounters, respectively, given detection (we assumed equal probabilities for both sides);  $\psi$  is probability that a randomly selected capture history belongs to one of the unique individuals encountered at least once;  $p_t$  is capture probability for occasion  $t$ , and  $\phi$  is apparent annual survival. See McClintock (2015) for more details on model parameterization. Additional estimated parameters not reported here are the latent state matrix (dead or alive) and the vector of individual random effects for capture probability.

Parameter	Prior	Mean	SD	Mode	P <sub>5</sub>	P <sub>25</sub>	P <sub>50</sub>	P <sub>75</sub>	P <sub>95</sub>	<i>n</i>
$\beta_{p,0}$	(0, 1)	-1.256	0.154	-1.222	-1.52	-1.355	-1.249	-1.148	-1.015	11,577
$\beta_{p,e}$	(0, 1)	0.189	0.079	0.185	0.060	0.135	0.188	0.242	0.321	38,293
$\beta_\phi$	(0, 1)	1.702	0.282	1.634	1.278	1.505	1.682	1.876	2.195	9,100
$\sigma_{p,z}^2$	Inv- $\gamma(1,0.01)$	0.230	0.188	0.023	0.01	0.083	0.195	0.330	0.582	12,487
$\alpha$	Unif(0,1)	0.926	0.048	0.959	0.835	0.900	0.936	0.962	0.986	51,068
$\tau$	Beta(1,1)	0.564	0.060	0.564	0.464	0.524	0.564	0.604	0.660	53,919
$\psi$	Beta(1,1)	0.915	0.030	0.923	0.863	0.896	0.917	0.936	0.959	48,365
$p_2$	N/A (derived)	0.155	0.027	0.145	0.114	0.135	0.152	0.171	0.204	17,297
$p_3$	N/A (derived)	0.099	0.022	0.096	0.066	0.083	0.097	0.113	0.138	44,732
$p_4$	N/A (derived)	0.125	0.022	0.120	0.092	0.110	0.123	0.139	0.163	24,466
$p_5$	N/A (derived)	0.085	0.023	0.080	0.050	0.068	0.082	0.099	0.127	52,296
$p_6$	N/A (derived)	0.121	0.021	0.116	0.088	0.106	0.119	0.134	0.158	26,548
$p_7$	N/A (derived)	0.101	0.022	0.098	0.068	0.086	0.100	0.115	0.139	43,013
$p_8$	N/A (derived)	0.168	0.032	0.158	0.121	0.146	0.165	0.187	0.225	16,335
$p_9$	N/A (derived)	0.117	0.021	0.111	0.084	0.102	0.115	0.130	0.154	29,258
$p_{10}$	N/A (derived)	0.170	0.033	0.160	0.123	0.147	0.167	0.190	0.229	16,301
$p_{11}$	N/A (derived)	0.210	0.049	0.201	0.139	0.175	0.205	0.239	0.299	17,139
$\phi$	N/A (derived)	0.950	0.027	0.962	0.899	0.934	0.954	0.970	0.986	9,811

**TABLE S2.2** Summaries of priors and results for parameters estimated by closed population model in *multimark*:  $\beta_{p,0}$  is the logit-link intercept of annual capture probability  $p$ ;  $\beta_{p,e}$  is the logit-link slope of capture probability with standardized Beaufort- and season-adjusted effort index (see Methods);  $\sigma_{p,z}$  is the standard deviation of logit-link individual random effects in capture probability;  $\alpha$  is probability of simultaneous left- and right-sided detections, given both types encountered;  $\tau$  is the sum of the conditional probabilities of left- and right-sided encounters, respectively, given detection (we assumed equal probabilities for both sides);  $\psi$  is probability that an individual in the data-augmented capture history matrix was encountered at least once;  $p_t$  is capture probability for occasion  $t$ ;  $N_{nc}$  is noncalf abundance, including a correction factor of 1.052;  $N$  is total abundance, including a correction factor of 1.107. See McClintock (2015) for more details on model parameterization. Not reported here are the latent capture history matrix and the vector of individual random effects for capture probability.

Parameter	Prior	Mean	SD	Mode	P <sub>5</sub>	P <sub>25</sub>	P <sub>50</sub>	P <sub>75</sub>	P <sub>95</sub>	<i>n</i>
$\beta_{p,0}$	(0, 1.75)	-1.774	0.695	-1.401	-3.108	-2.154	-1.643	-1.269	-0.876	3,961
$\beta_{p,e}$	(0, 1.75)	0.759	0.219	0.76	0.406	0.609	0.756	0.902	1.126	22,279
$\sigma_{p,z}$	half-Cauchy(25)	0.848	0.493	0.319	0.11	0.459	0.819	1.187	1.711	3,481
$\alpha$	Unif(0,1)	0.9	0.06	0.926	0.786	0.865	0.909	0.945	0.978	24,194
$\tau$	Beta(1,1)	0.556	0.066	0.556	0.446	0.51	0.556	0.6	0.662	21,871
$\psi$	Beta(1,1)	0.972	0.026	0.994	0.921	0.96	0.979	0.991	0.998	21,586
$p_9$	N/A (derived)	0.104	0.042	0.089	0.045	0.073	0.099	0.13	0.18	7,610
$p_{10}$	N/A (derived)	0.21	0.07	0.19	0.1	0.159	0.207	0.257	0.329	5,289
$p_{11}$	N/A (derived)	0.303	0.1	0.305	0.145	0.229	0.3	0.37	0.474	5,338
$N_{nc}$	N/A (derived)	115	49	84	67	83	101	131	208	5,207
$N$	N/A (derived)	121	52	89	71	87	106	138	219	5,207

**TABLE S2.3** Summaries of priors and results for parameters estimated by Pradel-lambda model:  $\beta_{p,0}$  is the logit-link intercept for annual capture probability  $p$ ;  $\beta_{p,e}$  is the logit-link slope for capture probability with standardized Beaufort- and season-adjusted effort index (see Methods);  $\sigma_{p,t}$  is the standard deviation of logit-link temporal random effects in capture probability,  $\beta_\lambda$  is the log-link intercept for population growth rate, and  $\lambda$  is population growth rate.

Parameter	Prior	Mean	SD	Mode	P <sub>5</sub>	P <sub>25</sub>	P <sub>50</sub>	P <sub>75</sub>	P <sub>95</sub>	<i>n</i>
$\beta_{p,0}$	(0, 3.16)	-2.378	0.227	-2.38	-2.749	-2.53	-2.379	-2.228	-2.003	1,001
$\beta_{p,e}$	(0, 10)	0.402	0.133	0.399	0.187	0.316	0.401	0.487	0.617	44,705
$\sigma_{p,t}$	Unif(0,2)	0.187	0.146	0.037	0.015	0.076	0.157	0.263	0.463	12,1827
$\beta_\lambda$	(0, 10)	-0.009	0.03	-0.01	-0.057	-0.029	-0.009	0.011	0.04	38,731
$p_1$	NA (derived)	0.086	0.025	0.076	0.053	0.068	0.081	0.098	0.133	4,822
$p_2$	NA (derived)	0.104	0.025	0.099	0.067	0.087	0.102	0.119	0.149	1,704
$p_3$	NA (derived)	0.055	0.017	0.052	0.031	0.044	0.054	0.065	0.085	5,171
$p_4$	NA (derived)	0.084	0.021	0.079	0.054	0.069	0.082	0.096	0.122	2,041
$p_5$	NA (derived)	0.056	0.018	0.048	0.032	0.043	0.053	0.065	0.089	7,440
$p_6$	NA (derived)	0.076	0.019	0.071	0.049	0.063	0.075	0.088	0.11	1,912
$p_7$	NA (derived)	0.065	0.017	0.059	0.04	0.052	0.062	0.075	0.096	2,885
$p_8$	NA (derived)	0.127	0.029	0.12	0.086	0.107	0.124	0.144	0.18	938
$p_9$	NA (derived)	0.069	0.018	0.067	0.042	0.057	0.068	0.08	0.101	1,991
$p_{10}$	NA (derived)	0.126	0.029	0.117	0.085	0.106	0.123	0.142	0.177	803
$p_{11}$	NA (derived)	0.169	0.04	0.16	0.111	0.14	0.164	0.192	0.241	795
$\lambda$	NA (derived)	0.992	0.029	0.989	0.944	0.972	0.991	1.011	1.041	39,021