

Estimating Natural Mortality for Atlantic Sea Scallops
(*Placopecten magellenicus*) Using a Size-Based Stock
Assessment Model

Deborah R. Hart*, Jui-Han Chang

Northeast Fisheries Science Center
166 Water St., Woods Hole, MA 02543, USA

*Email: deborah.hart@noaa.gov

June 26, 2022

Abstract

We estimate temporally varying natural mortality within a size-based stock assessment model for U.S. sea scallops in three different regions, and compare the results to corresponding models where natural mortality is not estimated, and (in one case) where only the mean natural mortality is estimated. In one model, for the Georges Bank Closed Areas, natural mortality was estimated by year for all sizes, whereas in the other two models (for the Georges Bank Open Areas and the Mid-Atlantic Bight), annual variation in natural mortality was estimated for juveniles only. Estimating natural mortality by year improved the performance over fixed natural mortality in all three models, as measured by improved model fit, Akaike's information criterion (AIC), and reduced retrospective patterns. There was evidence that natural mortality is density-dependent. Large recruitment events were associated with substantial increases in juvenile natural mortality, and natural mortality in the Georges Bank Closed Areas tended to increase at high biomass. Estimation of natural mortality was facilitated by survey data that gave estimates of absolute abundance together with low fishing mortality that minimized the confounding between natural and fishing mortality. Natural mortality is an important process that should be estimated when there is sufficient information to do so.

Keywords: Natural mortality, Density dependence, Stock assessment models, Length-based models, Invertebrate fisheries, *Placopecten magellanicus*

1 Introduction

Natural mortality (M) is one of the most important parameters in fisheries stock assessments. It is typically directly used to estimate fishing mortality from total mortality, and its value strongly influences estimates of abundance and reference points (Maunder and Piner 2015, Punt et al. 2021). Despite this, natural mortality is frequently not estimated in stock assessment models, and is often assumed constant in time and age or size, even though this is unlikely to be true. The problem is that M is difficult to estimate since it can be confounded by errors in catch estimates, or assumptions regarding fishery and survey selectivity, steepness or stock-recruit relationships, age or growth, length to weight conversions, and survey efficiency (Punt et al. 2021). Erroneous estimates of any of these factors can bias estimates of M (Lee et al. 2011, Johnson et al. 2015), possibly as much or more than a fixed estimate.

The natural mortality of Atlantic sea scallops (*Placopecten magellanicus*) has been estimated by a number of methods. Merrill and Posgay (1964) used the “clapper ratio” method, $M = \frac{1}{S} \frac{C}{L}$, where C is the density of “clappers”, i.e., dead shells with the hinge ligament attached, L is the density of live scallops, and S is the (mean) time for a clapper to separate into two valves after death, to estimate $M = 0.1$ for the Georges Bank sea scallops. There are a number of uncertainties surrounding this method, however, including its equilibrium assumption, the estimate of S , which is likely size specific, and the assumption of equal catchability of clappers and live scallops in survey gear. Additionally, as a ratio estimator, the clapper ratio is biased, which is of particular concern given the high uncertainty in the S term in the denominator (Hart 2013). Applying a bias-correction factor gave a revised estimate of $M = 0.12$ for Georges Bank sea scallops. Estimates of M for Mid-Atlantic sea scallops can be obtained from those from Georges Bank using the idea that M/K , where K is the Brody growth coefficient, should be invariant among populations of the same species (Beverton and Holt 1959, Hart 2013).

In this article, we present a method to estimate size-specific and temporally variable natural mortality in stock assessment models. This method is applied to the U.S. stock assessments of Atlantic sea scallops, and the results are compared these to models where

48 M is fixed. Sea scallops offer an ideal opportunity to estimate M for a number of reasons.
49 First, there have been areas that have been closed to scallop fishing for a number of years,
50 thereby reducing the confounding of natural and fishing mortality. Similarly, estimation of
51 juvenile M is facilitated because surveys reliably detect juvenile scallops two years before
52 they are currently vulnerable to the commercial fishery. Finally, sea scallops are tracked by a
53 number of dedicated, relatively precise surveys, each of which has its own external estimate
54 of gear efficiency. Such surveys can be used to directly estimate absolute abundance which
55 in turn can be combined with estimates of catch and total mortality to estimate natural
56 mortality.

57 **2 Methods**

58 A size-based stock assessment model (CASA [Catch At Size Analysis], Sullivan et al. 1990,
59 Hart et al. 2013) was used that tracked sea scallop numbers at shell height, i.e., the distance
60 from the base (umbo) to the tip of the shell. Growth was modeled using stochastic growth
61 matrices based on growth increment data from shell ring analysis (Hart and Chute 2009);
62 these growth matrices were fixed and not estimated within the stock assessment model.
63 Three separate models for different regions were used: Mid-Atlantic, Georges Bank Open
64 and Georges Bank Closed (Figure 1). The Georges Bank region was split into two models
65 because of the different management history of the two areas; the Georges Bank Closed
66 Areas have experienced much lower fishing mortality than the Georges Bank Open Areas
67 since 1994, including a number of years that they were completely closed to scallop fishing
68 (Hart and Rago 2006). Splitting the region into two models gives more stable and precise
69 model results than using a single model with domed selectivity for the periods where portions
70 of the region were closed (Hart et al. 2013). Technical details regarding the stock assessment
71 model can be found in Jacobson and Chang (2018).

72 The models were fitted by maximum likelihood to abundance trends and size data ob-
73 tained from sea scallop dredge surveys during 1975–2019 (Serchuk and Wigley, 1986, Hart
74 and Rago, 2006), drop camera surveys conducted in most years since 2003 (Stokesbury et

75 al., 2004, Bethoney and Stokesbury 2018), towed camera (Habcam) surveys during 2011-
76 2019 (Howland et al. 2006, Chang et al. 2017) as well as commercial landings and shell
77 height data from port samples and at-sea observers. Model estimates of survey efficiency
78 were constrained by priors (likelihood penalties); the priors from the two optical surveys
79 had an expected efficiency of one, whereas the expected efficiency of the survey dredge was
80 based on paired tow experiments with Habcam (Miller et al. 2019). Standard deviations
81 of the priors were set at 0.1. The drop camera survey was separated into two time series:
82 one prior to 2015, when standard definition video cameras were used, and since 2015, when
83 high definition digital cameras were used. In 2019, neither the drop camera nor the Habcam
84 surveys were complete on Georges Bank, but at least one of these optical surveys covered all
85 portions of this region. These surveys were therefore combined on Georges Bank in 2019; the
86 combined survey was considered part of the Habcam time series for Georges Bank Closed
87 and the drop camera survey for Georges Bank Open.

88 In the Georges Bank Closed model, natural mortality was assumed to not vary with size
89 since there was no evidence otherwise. For that model, we estimated mean natural mortality
90 (over time), \bar{M} , as well as the annual deviations of M from this mean $\gamma(y)$ (except the first
91 year):

$$M(y) = \bar{M} + \gamma(y) \quad (1)$$

92 In the other two models, we varied M by shell height h , using a weighted average of juvenile
93 and adult natural mortality, M_{juv} and M_{ad} :

$$M(h) = \alpha(h)M_{\text{juv}} + [1 - \alpha(h)]M_{\text{ad}} \quad (2)$$

94 where $\alpha(h) = 1 - \frac{1}{1 + \exp(-h_0[h - a])}$ is a decreasing logistic function, with left asymptote
95 equal to one and right asymptote equal to zero, h_0 is fixed at the shell height at about
96 age 2.5 (70mm for Georges Bank Open and 65mm for Mid-Atlantic), and a is fixed at 0.1
97 (Figure 2). As in equation (1), juvenile and adult natural mortality can be decomposed
98 into their mean values and deviations from the mean: $M_{\text{juv}}(y) = \bar{M}_{\text{juv}} + \theta(y)$ and $M_{\text{ad}}(y) =$
99 $\bar{M}_{\text{ad}} + \omega(y)$. However, the values of the adult natural mortality parameters \bar{M}_{ad} and $\omega(y)$
100 were not estimable in these two models due to confounding with fishing mortality. They

101 were fixed at a value of $\bar{M}_{\text{ad}} = 0.2$ and $\omega(y) = 0$ in the Georges Bank Open model, based
102 on a similar estimate in Georges Bank Closed. The Brody growth coefficient K for Mid-
103 Atlantic scallops is about 25% higher than for those on Georges Bank (Hart and Chute
104 2009), and hence life history theory suggests that M_{ad} in the Mid-Atlantic should also be
105 25% higher (Beverton and Holt 1959). Thus, we fixed $\bar{M}_{\text{ad}} = 0.25$ for Mid-Atlantic sea
106 scallops and $\omega(y) = 0$. Mean juvenile natural mortality \bar{M}_{juv} was also set to those values,
107 but the deviations in juvenile M , $\theta(y)$, were estimated for each year except for the first year.

108 We evaluated model performance using likelihood and Akaike’s information criterion
109 (AIC) as a measure of model fit to the data. We also used retrospective analysis (6 peels)
110 to measure model stability, with Mohn’s ρ as a summary statistic (Mohn 1999). It is of
111 interest to examine whether density dependence plays a role in natural mortality. To this
112 end, for each year in the Mid-Atlantic and Georges Bank Open models, we related the model
113 estimates of two year old recruits to juvenile natural mortality. For the Georges Bank Closed
114 Model, estimates of overall natural mortality were compared to biomass for each year. In
115 all three cases, the relationships between natural mortality and $\log(\text{recruitment})$ or biomass
116 were fit to GAM models, using the `mgcv` R package with a Gaussian distribution and for
117 the Georges Bank Open and Mid-Atlantic models, a log link.

118 Results

119 Estimation of natural mortality improved fits for all three models and decreased retrospective
120 patterns (Table 1; Figures 3-4). Mean standard errors for natural mortality ranged from
121 0.044 to 0.049 in the models that temporally estimated M and 0.009 in the Georges Bank
122 closed model that estimated the mean M only (Table 1). In the Georges Bank Closed model,
123 temporal estimation of M was substantially superior in terms of both AIC and Mohn’s ρ than
124 just estimating the mean M , which in turn decreased AIC and slightly decreased Mohn’s ρ
125 compared to the fixed M run. In the terminal year in Georges Bank Closed, both the fixed
126 M and estimated mean M runs considerably overestimated biomass compared to surveys,
127 whereas the run that estimated M for all years was able to match the declines in survey

128 biomass observed in 2018 and 2019 (Figure 3a). For Georges Bank Open and the Mid-
129 Atlantic models, estimating juvenile M improved AIC and reduced the retrospective pattern
130 (especially in Georges Bank Open) compared to using a fixed M , although the differences in
131 estimated biomass between these runs were modest.

132 In the Georges Bank Closed Areas, the three years with the highest natural mortalities all
133 occurred at high biomasses (> 75000 t, Figure 5ab). The one other year with substantially
134 increased natural mortality ($M > 0.3$) occurred at a moderate biomass. In both Georges
135 Bank Open and the Mid-Atlantic, high juvenile natural mortality only occurred after strong
136 recruitment events; there was no evidence of elevated juvenile natural mortality at low or
137 moderate levels of recruitment (< 1500 million, Figure 5c-f). The GAM fits predicted
138 substantial increases in M at high recruitment in these areas.

139 Discussion

140 Like many marine fish and invertebrates, natural mortality in sea scallops can be charac-
141 terized by periods where natural mortality is near constant, punctuated by spikes in M .
142 These increases in natural mortality may be due to high temperatures, disease and/or pre-
143 dation (Dickie and Metcof 1963, Gulka et al., 1983, Belvin et al., 2008, Hart and Shank
144 2011, Levesque et al. 2016, Siemann et al. 2019). Episodic natural mortality has been noted
145 elsewhere for sea scallops such as in Narragansett Bay (Gulka et al., 1983) and the Bay of
146 Fundy (Smith and Lundy 2002), and is commonly associated with high densities.

147 Estimation of natural mortality by year is one method to account for variable natural
148 mortality, and in all three models, it substantially improved model performance, as measured
149 both by model fits and reduced retrospective patterns. However, in doing so, we estimated
150 a parameter for every year (45 extra parameters in our case), most of which were similar to
151 the baseline M and contributed little to model fit. For this reason, AIC values for the Mid-
152 Atlantic model with M estimated for each year were only slightly less than using a fixed value
153 for natural mortality. Future work should consider ways to reduce the number of estimated
154 parameters (e.g., using smoothing splines or likelihood penalties on the M deviations), while

155 still allowing estimates of high natural mortality for the years that it occurs.

156 Estimation of natural mortality in the Georges Bank Closed model was facilitated by the
157 zero or low fishing mortality in many years ($F < 0.01$ in eight years, and $F < 0.1$ in 19
158 of the 45 years of the time series). Indeed, we could not obtain a reliable estimate of the
159 mean M in either of the other two models. Even in the Georges Bank Closed model, some
160 uncertainties remain. In particular, some illegal fishing did occur in these areas, as evidenced
161 by a number of enforcement actions. However, vessel monitoring systems that were required
162 on most scallops boats since 1997 show very limited incursions into these areas, so that the
163 levels of illegal fishing were likely much lower than natural mortality, and any bias introduced
164 by illegal fishing was therefore small.

165 Stokesbury et al. (2007) claimed that there was a “mass mortality”, i.e., very high natural
166 mortality, in Nantucket Lightship Closed Area, a portion of the Georges Bank Closed Areas,
167 between 2004 and 2005, based on a large decline observed in their drop camera survey
168 between those years. Our Georges Bank Closed model does not show evidence of increased
169 natural mortality during that time, and there was only a modest decline observed in the
170 scallop dredge survey. The drop camera survey employed a fixed grid design, where the
171 station locations in 2005 repeated those from 2004. Maps showing densities from the 2004
172 drop camera survey were distributed to fishermen, and the area was reopened to limited
173 fishing in late 2004. It is likely that the large decline in the drop camera survey in 2005 was
174 an artifact due to heavy fishing around the drop camera stations that showed high density
175 in the 2004 survey (as evidenced by vessel monitoring system data), rather than a large
176 increase in natural mortality.

177 Elevated natural mortality in the Georges Bank Closed Areas tended to occur when its
178 biomass was high. However, in many of the years with high biomass, natural mortality
179 was near its baseline level. The increases in natural mortality in the Georges Bank Closed
180 Areas during 2010-12 were associated with elevated levels of “gray meat” disease (Levesque
181 et al. 2016, Siemann et al. 2019). Diseases are more likely to spread at high densities
182 or biomass. Similarly, predation by *Asterias* spp. sea stars are also contributors to adult
183 natural mortality, and high scallop densities may tend to attract these predators. However, it

184 appears that while high (adult) biomass may be conducive to disease or predation processes,
185 these processes are not always elevated when the biomass is high.

186 By contrast, the three large recruitment events in the Georges Bank open and Mid-
187 Atlantic areas were all associated with very high juvenile natural mortality rates. This
188 phenomenon has also been frequently observed in seeding and tank experiments, where
189 juvenile scallops at high densities have experienced high mortality (e.g., Barbeau et al. 1998,
190 Wong et al. 2005, Fr chet te et al. 2013). In most cases, the increased mortality was ascribed
191 to predation by *Cancer* spp. crabs, which appear to act as agents of density dependence
192 (Barbeau et al. 1998, Wong and Barbeau 2005, 2006, Hart and Shank 2011), although
193 Fr chet te et al. (2013) reported self-thinning due to interference or competition between the
194 scallops themselves. These crabs are not predators of larger scallops (Elner and Jamieson
195 1979), which may explain why density-dependent mortality of adults is not as strong.

196 High juvenile mortality on large year classes induced a retrospective pattern in models
197 with fixed M , where biomass (and abundance) estimates declined as more years are added.
198 The conflict between the initial estimate of the large year class and subsequent reduced
199 estimates is eventually resolved in the model by decreasing the estimated size of the large year
200 class, so that the initial high recruitment estimate is (incorrectly) attributed to observation
201 error. Even the variable juvenile M models display a similar retrospective pattern, since
202 it is only after several years of observations of the large year class that the decline can be
203 attributed in the model to increased natural mortality rather than observation error. After a
204 sufficient number of years since the recruitment event, abundance and biomass estimates in
205 the fixed and estimated M models will be similar, although the models arrive at the estimate
206 in different ways. The fixed M models reduce the estimated size of the large year classes
207 to well below that indicated by the surveys, whereas when M is estimated, the size of these
208 year classes is reduced in subsequent years by increasing juvenile natural mortality. This
209 distinction becomes important when projecting into the future the fate of a large year class;
210 a fixed M model may substantially overestimate future biomass and landings. A variable
211 M projection model can be constructed by building in density-dependence, based on the
212 relationships between biomass or recruitment and natural mortality (Fig. 4).

213 Like many other U.S. fisheries, fishing mortality has declined considerably in the sea
214 scallop fishery since the mid-1990s (Hart and Rago 2006). During the years near the peak
215 of fishing mortality, the effects of natural mortality on population dynamics was limited
216 since fishing mortality was many times greater than natural mortality. Thus, for fisheries
217 where such substantial overfishing is occurring, it would be very difficult to estimate (adult)
218 natural mortality, but the consequences of a misspecified M would be limited. On the
219 other hand, when fishing mortality is reduced below F_{MSY} , fishing and natural mortality
220 are commensurate quantities, so obtaining good estimates of natural mortality is important
221 for understanding stock dynamics. The Georges Bank Closed Areas represent an extreme
222 example of this, since natural mortality has been greater than fishing mortality (and often
223 much greater) for most years since 1994.

224 We have presented a case study where both overall and juvenile natural mortality can
225 be estimated by year for a data rich stock with surveys of absolute abundance, and where
226 potentially confounding fishing mortality is relatively low. Estimation of natural mortality
227 not only improved model performance, but it gave important biological insights into the role
228 of density-dependence in regulating sea scallop populations.

229 References

- 230 Barbeau, M.A., Hatcher, B.G., Scheibling, R.E., Hennigar, A.W., Taylor, L.H. and Risk,
231 A.C., 1998. Dynamics of juvenile sea scallop (*Placopecten magellanicus*) and their predators
232 in bottom seeding trials in Lunenburg Bay, Nova Scotia. Can. J. Fish. Aquat. Sci., 53,
233 2494-2512.
- 234 Barbeau, M.A., Scheibling, R.E., Hatcher, B.G., 1998. Behavioural responses of predatory
235 crabs and sea stars to varying density of juvenile sea scallops, Aquaculture, 169, 87-98.
- 236 Barbeau, M.A., Scheibling, R.E., Hatcher, B.G., Taylor, L.H., Hennigar, A.W., 1994. Sur-
237 vival analysis of tethered juvenile sea scallops *Placopecten magellanicus* in field experiments:
238 effects of predators, scallop size and density, site and season. Mar. Ecol. Progr. Ser., 243-256.
- 239 Belvin, S., Tremblay, R., Roussy, M. and McGladdery, S.E., 2008. Inoculation experiments
240 to understand mass mortalities in sea scallop, *Placopecten magellanicus*. J. Shellfish Res.,
241 27, 251-260.
- 242 Bethoney, N.D., Stokesbury, K.D., 2018. Methods for image-based surveys of benthic
243 macroinvertebrates and their habitat exemplified by the drop camera survey for the Atlantic
244 sea scallop. Journal of Visualized Experiments: JoVE, 137.
- 245 Beverton, R.J.H., Holt, S.J., 1959. January. A review of the lifespans and mortality rates
246 of fish in nature, and their relation to growth and other physiological characteristics. In
247 Ciba Foundation Symposium-The Lifespan of Animals (Colloquia on Ageing) (Vol. 5, pp.
248 142-180). Chichester, UK: John Wiley & Sons, Ltd.
- 249 Chang, J.H., Shank, B.V., Hart, D.R., 2017. A comparison of methods to estimate abundance
250 and biomass from belt transect surveys. Limnology and Oceanography: Methods, 15, 480-
251 494.
- 252 Dickie, L.M., Medcof, J.C., 1963. Causes of mass mortalities of scallops (*Placopecten magel-*
253 *lanicus*) in the southwestern Gulf of St. Lawrence. J. Fish. Bd. Can, 20, 451-482.

254 Elner R.W., Jamieson G.S., 1979. Predation of sea scallops, *Placopecten magellanicus*, by
255 the rock crab, *Cancer irroratus*, and the American lobster, *Homarus americanus*. J. Fish.
256 Res. Bd. Can. 36, 537-543

257 Fréchette, M., Urquiza, J.M., Daigle, G., Maheux, D., Dumais, J.F., 2013. Self-thinning
258 dynamics in experimental scallop populations. Aquaculture International, 21, 539-551.

259 Gulka, G., Chang, P.W. and Marti, K.A., 1983. Prokaryotic infection associated with a mass
260 mortality of the sea scallop, *Placopecten magellanicus*. J. Fish Diseases, 6, 355-364.

261 Hart, D.R., 2013. Quantifying the tradeoff between precaution and yield in fishery reference
262 points. ICES J. Mar. Sci., 70, 591-603.

263 Hart, D.R., Chute, A.S., 2009. Estimating von Bertalanffy growth parameters from growth
264 increment data using a linear mixed effects model, with an application to the sea scallop
265 *Placopecten magellanicus*. ICES J. Mar. Sci., 66, 2165-2175.

266 Hart, D.R., Jacobson, L.D., Tang, J., 2013. To split or not to split: assessment of Georges
267 Bank sea scallops in the presence of marine protected areas. Fish. Res., 144, 74-83.

268 Hart, D.R. and Shank, B.V., 2011. Mortality of sea scallops *Placopecten magellanicus* in
269 the Mid-Atlantic Bight: Comment on Stokesbury et al.(2011). Mar. Ecol. Progr. Ser., 443,
270 293-297.

271 Hart, D.R. and Rago, P.J., 2006. Long-term dynamics of US Atlantic sea scallop *Placopecten*
272 *magellanicus* populations. N. Am. J. Fish. Manage., 26, 490-501.

273 Howland, J., Gallager, S., Singh, H., Girard, A., Abrams, L., Griner, C., Taylor, R. and Vine,
274 N., 2006. Development of a towed survey system for deployment by the fishing industry. In
275 OCEANS 2006 (pp. 1-5). IEEE.

276 Jacobson, L., Chang, J.H., 2018. Technical documentation for the CASA length structured
277 stock assessment model used in the SARC-59 sea scallop stock assessment, Appendix A6
278 in: 65th Northeast Regional Stock Assessment Workshop (65th SAW) Assessment Report.
279 Northeast Fisheries Science Center Ref. Doc. 18-11.

280 Johnson, K.F., Monnahan, C.C., McGilliard, C.R., Vert-Pre, K.A., Anderson, S.C., Cun-
281 ningham, C.J., Hurtado-Ferro, F., Licandeo, R.R., Muradian, M.L., Ono, K. and Szuwalski,
282 C.S., 2015. Time-varying natural mortality in fisheries stock assessment models: identifying
283 a default approach. ICES J. Mar. Sci., 72, 137-150.

284 Lee, H.H., Maunder, M.N., Piner, K.R., Methot, R.D., 2011. Estimating natural mortality
285 within a fisheries stock assessment model: an evaluation using simulation analysis based on
286 twelve stock assessments. Fish. Res., 109, 89-94.

287 Levesque, M.M., Inglis, S.D., Shumway, S.E., Stokesbury, K.D., 2016. Mortality assessment
288 of Atlantic sea scallops (*Placopecten magellanicus*) from gray-meat disease. J. Shellfish Res.,
289 35, 295-305.

290 Maunder, M.N., Piner, K.R., 2015. Contemporary fisheries stock assessment: many issues
291 still remain. ICES J. Mar. Sci., 72, 7-18.

292 Merrill, A.S., Posgay, J.A., 1964. Estimating the natural mortality rate of the sea scallop
293 (*Placopecten magellanicus*). I.C.N.A.F. Res. Bull., 1, 88-106.

294 Miller, T.J., Hart, D.R., Hopkins, K., Vine, N.H., Taylor, R., York, A.D., Gallagher, S.M.,
295 2019. Estimation of the capture efficiency and abundance of Atlantic sea scallops (*Placopecten*
296 *magellanicus*) from paired photographic–dredge tows using hierarchical models. Can. J. Fish.
297 Aquat. Sci., 76, 847-855.

298 Mohn, R., 1999. The retrospective problem in sequential population analysis: an investiga-
299 tion using cod fishery and simulated data. ICES J. Mar. Sci., 56, 473-488.

300 Punt, A.E., Castillo-Jordán, C., Hamel, O.S., Cope, J.M., Maunder, M.N., Ianelli, J.M.,
301 2021. Consequences of error in natural mortality and its estimation in stock assessment
302 models, Fish. Res., 233, 105759.

303 Serchuk, F.M., and Wigley, S.E., 1986., Evaluation of USA and Canadian research vessel
304 surveys for sea scallops (*Placopecten magellanicus*) on Georges Bank. J. Northwest Atl. Fish.
305 Sci., 7, 1-13.

306 Siemann, L.A., Garcia, L.M., Huntsberger, C.J. and Smolowitz, R.J., 2019. Investigating the
307 impact of multiple factors on gray meats in Atlantic sea scallops (*Placopecten magellanicus*).
308 J. Shellfish Res., 38, 233-243.

309 Smith, S.J. and Lundy, M.J., 2002. Scallop production area 4 in the Bay of Fundy: stock
310 status and forecast. Canadian Science Advisory Secretariat Research Document 2002/018.

311 Stokesbury, K.D., Harris, B.P., Marino, M.C., Nogueira, J.I., 2004. Estimation of sea scallop
312 abundance using a video survey in off-shore US waters. J. Shellfish Res., 23, 33-41.

313 Stokesbury, K.D., Harris, B.P., Marino II, M.C. and Nogueira, J.I., 2007. Sea scallop mass
314 mortality in a Marine Protected Area. Mar. Ecol. Prog. Ser., 349, 151-158.

315 Sullivan, P.J., Lai, H.L., Gallucci, V.F., 1990. A catch-at-length analysis that incorporates
316 a stochastic model of growth. Can. J. Fish. Aquat. Sci., 47, 84-198.

317 Wong M.C., Barbeau M.A., 2005. Prey selection and the functional response of sea stars
318 (*Asterias vulgaris* Verrill) and rock crabs (*Cancer irroratus* Say) preying on juvenile sea
319 scallops (*Placopecten magellanicus* (Gmelin)), and blue mussels (*Mytilus edulis* Linnaeus).
320 J. Exp. Mar. Biol. Ecol. 327, 1-21.

321 Wong M.C., Barbeau M.A., Hennigar A.W., Robinson S.M.C., 2005. Protective refuges for
322 seeded juvenile scallops (*Placopecten magellanicus*) from sea star (*Asterias* spp.) and crab
323 (*Cancer irroratus* and *Carcinus maenas*) predation. Can. J. Fish. Aquat. Sci. 62, 1766-1781.

324 Wong, M.C., Barbeau, M.A., 2006. Rock crab predation of juvenile sea scallops: the func-
325 tional response and its implications for bottom culture. Aquaculture International, 14, 355-
326 376.

Table 1: Summary of model runs: Number of parameters, log likelihood, AIC, Mohn’s ρ , and the mean standard error of the M estimates. In the “Est M ” runs, natural mortality is being estimated by year, either for all sizes (GB Closed) or juvenile M only (GB Open and MA). In the Mean M run (GB Closed only), the mean M is being estimated, but M is assumed constant over time. In the other runs, M is assumed fixed at the specified value.

Region	Model	NParam	ln(Likelihood)	AIC	ρ	Mean M SE
GBCl	Est M	145	8735	17760	0.22	0.044
GBCl	Mean M	100	8818	17836	0.34	0.009
GBCl	$M = 0.16$	99	8852	17902	0.37	NA
GBOp	Est M	144	9208	18704	0.33	0.049
GBOp	$M = 0.2$	100	9300	18800	0.56	NA
MA	Est M	150	-7208	-14114	0.33	0.048
MA	$M = 0.25$	106	-7116	-14020	0.41	NA

328 **Figures**

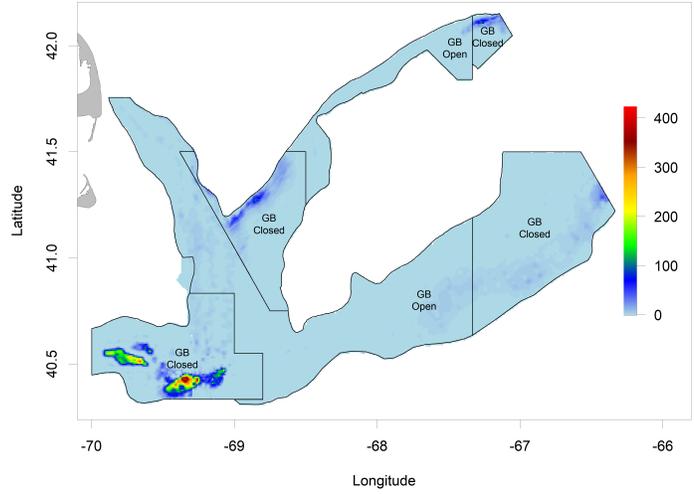
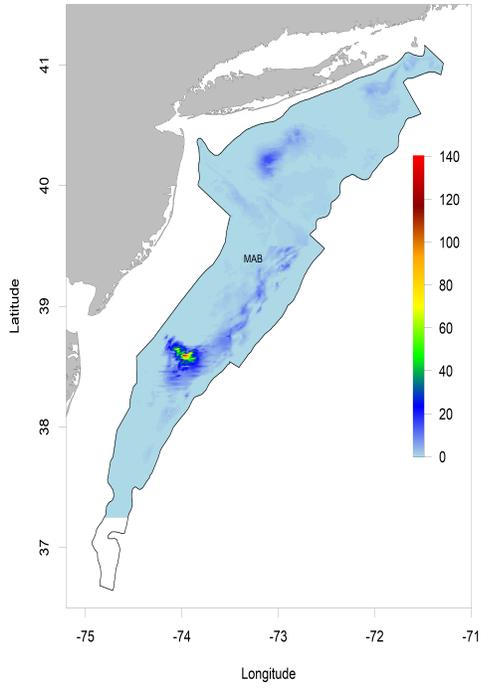


Figure 1

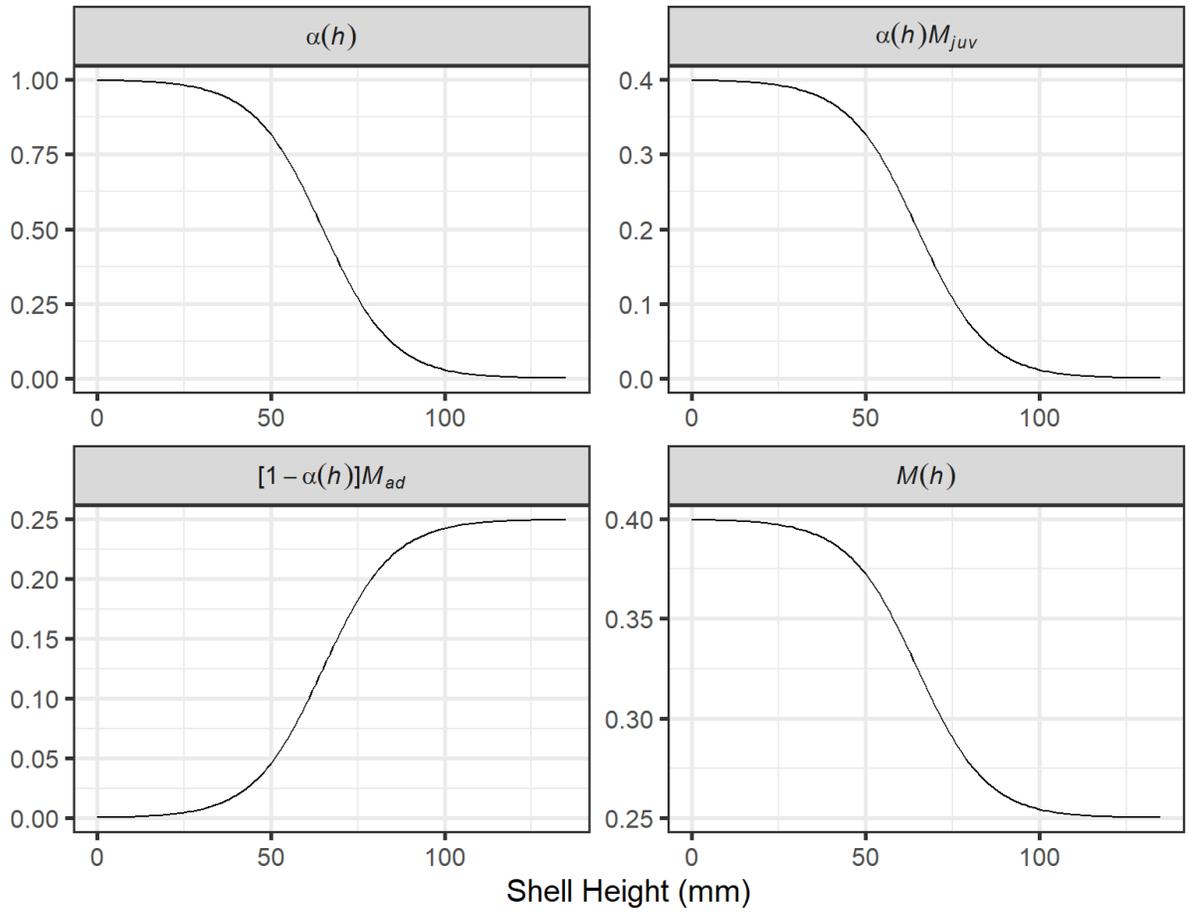


Figure 2

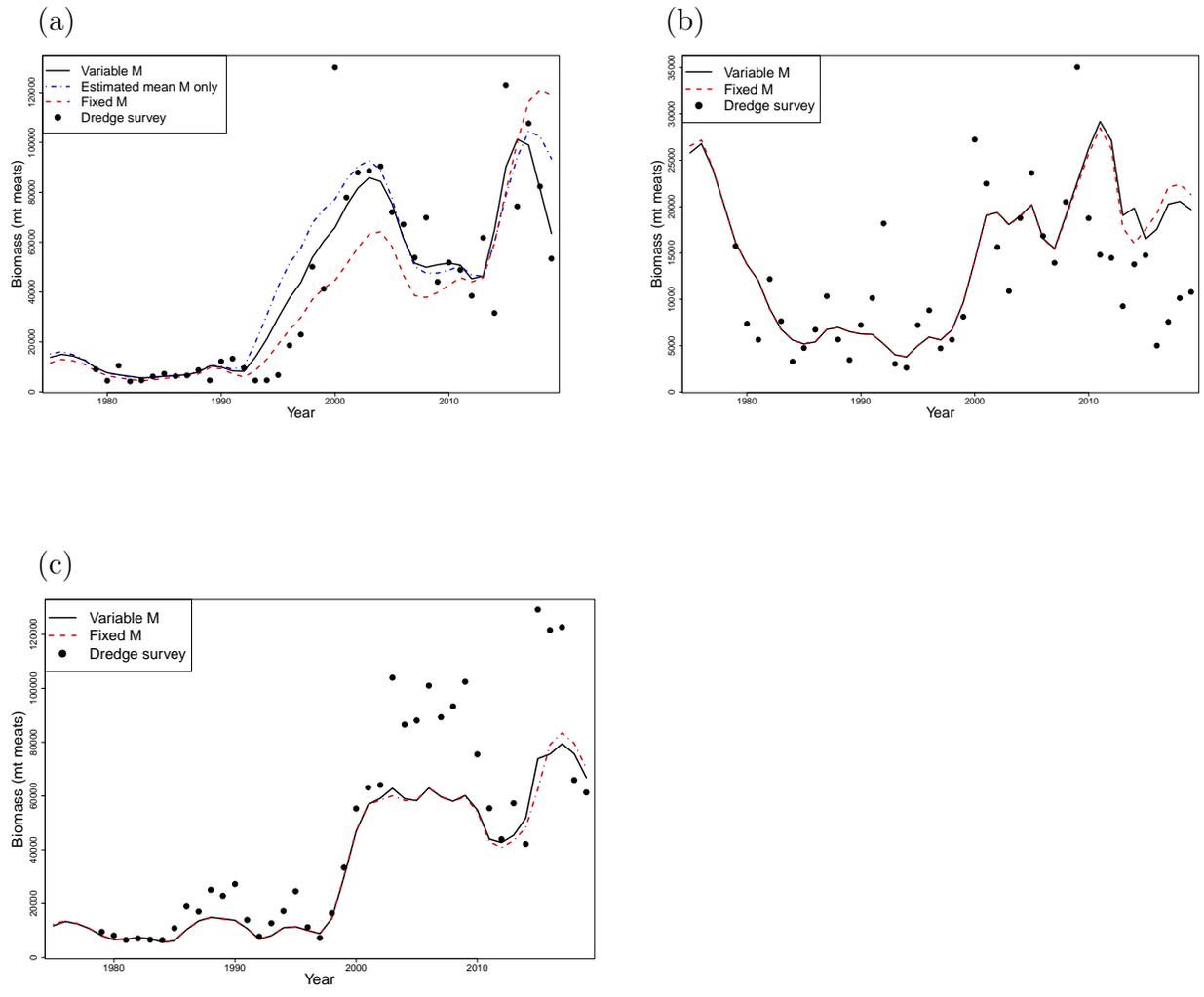


Figure 3

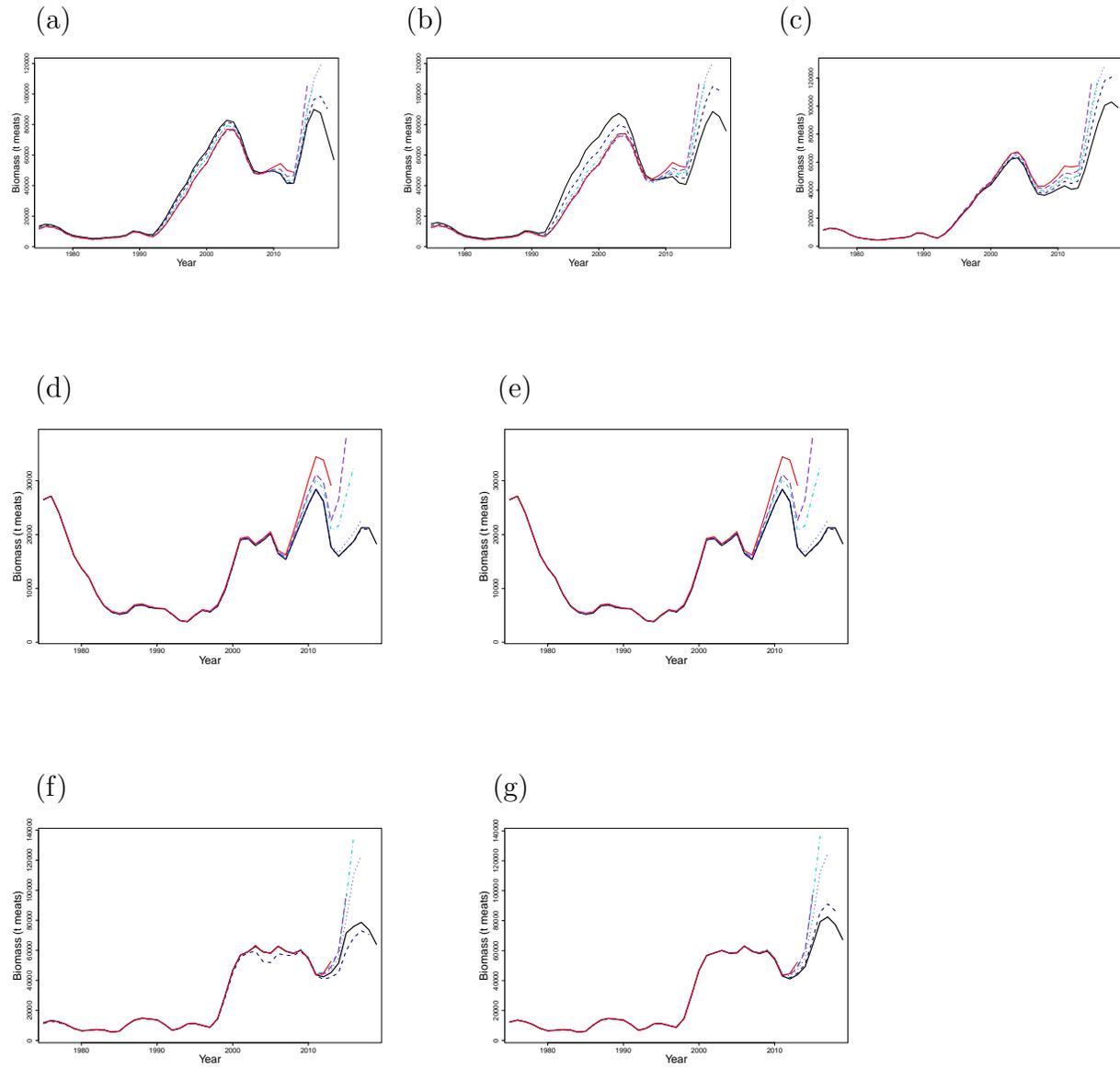


Figure 4

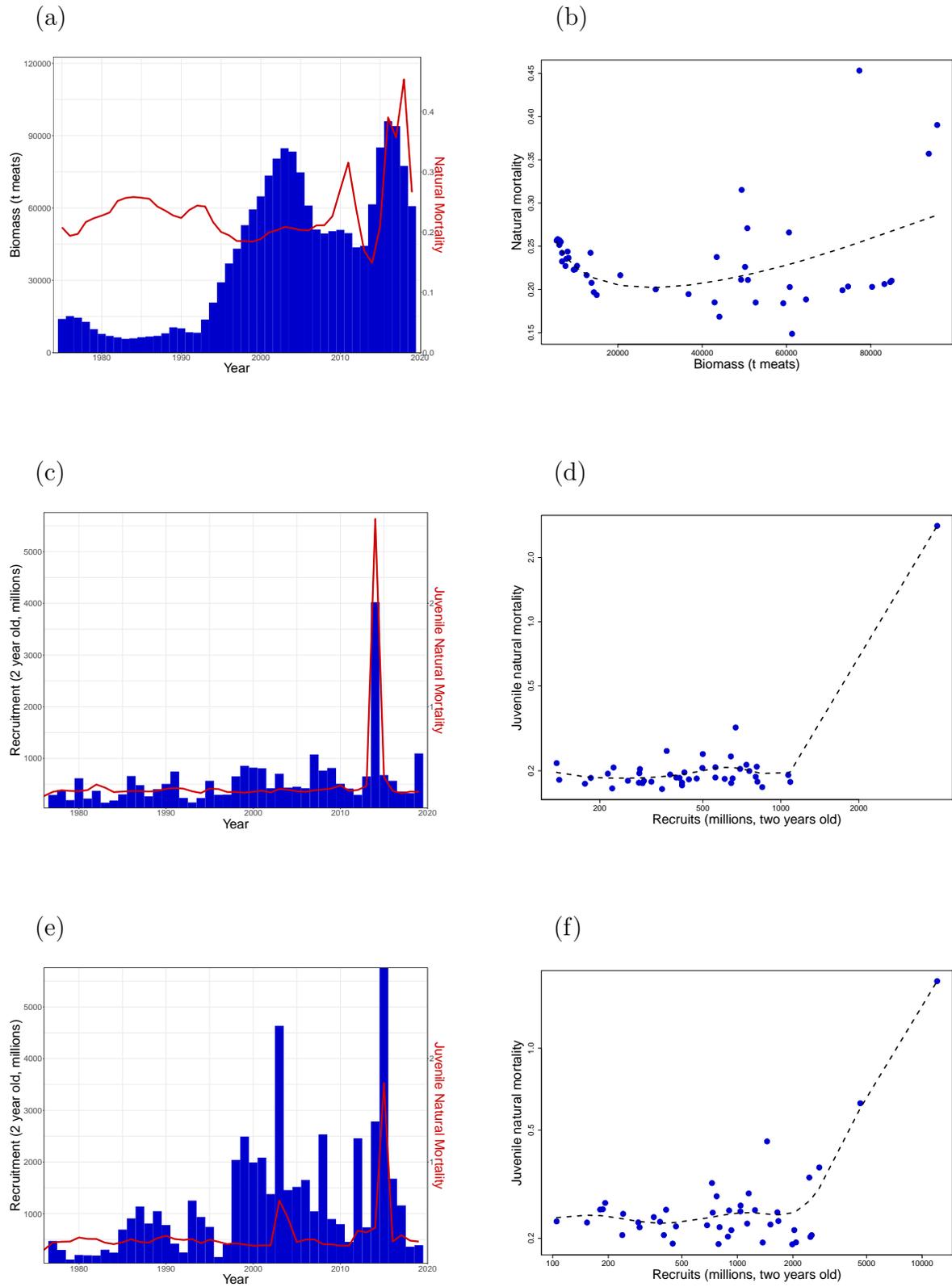


Figure 5

329 Figure Legends

330 **Fig. 1.** Charts of U.S. sea scallop grounds, showing the Mid-Atlantic Bight (MAB, left) and
331 Georges Bank (GB, right), split into open and closed portions, with densities as estimated
332 by the 2018 Habcam survey. The eastern portions of Georges Bank is in Canadian waters
333 and was not modeled.

334 **Fig. 2.** Example plots showing the calculation of natural mortality by size in the Mid-
335 Atlantic, assuming $M_{\text{juv}} = 0.4$ and $M_{\text{ad}} = 0.25$. Clockwise, from the upper left, the figures
336 show the logistic curve $\alpha(h)$, the contribution at shell height to natural mortality by the
337 juvenile component of natural mortality ($\alpha(h)M_{\text{juv}}$), the contribution at shell height to nat-
338 ural mortality by the adult component of natural mortality ($[1 - \alpha(h)]M_{\text{ad}}$), and natural
339 mortality $M(h)$ at shell height.

340 **Fig. 3.** Comparison of model estimates and the dredge survey for (a) Georges Bank Closed,
341 (b) Georges Bank Open, and (c) the Mid-Atlantic Bight. The solid black line shows estimates
342 for models that estimate M by year, whereas the red dashed line are from models where
343 M is fixed. The black dots show estimates directly from the dredge survey, and the blue
344 dashed-dotted line (Georges Bank Closed model only) shows output from the model that
345 estimates mean M only.

346 **Fig. 4.** Six peel retrospectives for the Georges Bank Closed model with (a) M estimated by
347 year, (b) only the mean M estimated and (c) M fixed; for the Georges Bank Open model
348 with (d) juvenile M estimated by year and (e) M fixed; for the Mid-Atlantic model with (f)
349 juvenile M estimated by year and (g) M fixed.

350 **Fig. 5.** Natural mortality (red line) and recruitment estimates in the variable M runs for (a)
351 Georges Bank Closed, (c) Georges Bank Open and (e) Mid-Atlantic. (b) Plot of estimated
352 M vs. biomass for Georges Bank Closed. Plots of estimated M vs. (2 year old) recruitment
353 in (d) Georges Bank Open and (f) Mid-Atlantic. The dashed lines in (b), (d) and (f) show
354 GAM fits.