

A Bayesian state-space mark-recapture model to estimate exploitation rates in mixed stock fisheries

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1 **Abstract**

2 A Bayesian state-space mark-recapture model is developed to estimate the exploitation rates
3 of fish stocks caught in mixed stock fisheries. Expert knowledge and published results on
4 biological parameters, reporting rates of tags and other key parameters are incorporated into
5 the mark-recapture analysis through elaborations in model structure and the use of
6 informative prior probability distributions for model parameters. Information on related
7 stocks is incorporated through the use of hierarchical structures and parameters that represent
8 differences between the stock in question and related stocks. Fishing mortality rates are
9 modelled using fishing effort data as covariates. A state-space formulation is adopted to
10 account for uncertainties in system dynamics and the observation process. The methodology
11 is applied to wild Atlantic salmon (*Salmo salar*) stocks from rivers located in the north-
12 eastern Baltic Sea that are exploited by a sequence of mixed and single stock fisheries.
13 Estimated fishing mortality rates for wild salmon are influenced by prior knowledge about tag
14 reporting rates and salmon biology and to a limited extent by prior assumptions about
15 exploitation rates.

16

17 Keywords: mark-recapture, mixed fishery, fishing mortality rates, Bayesian, Atlantic salmon

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19

20

20 **Introduction**

21 Fish stocks in mixed-stock fisheries often exhibit different migration patterns, life
22 histories, productivity rates and susceptibilities to natural and fishing mortality rates. This
23 makes it difficult to assess the impact of mixed stock fisheries on the individual stocks. Catch
24 data, catch-per-unit-effort data and research indices of abundance, often applied in stock
25 assessments, may contain sufficient information to assess exploitation rates of the combined
26 population, but due to the level of aggregation, these data are unlikely to provide sufficient
27 information about exploitation rates on the individual stocks (Hilborn 1990; Hampton and
28 Fournier 2001).

29 In contrast, mark and recapture data, provided that the tagging design is adequate and
30 the reported recapture rates are sufficiently high, can be among the most informative types of
31 data available for fish stock assessment (Punt et al. 2000; Martell and Walters 2002). In this
32 paper we demonstrate how mark-recapture data can be used for the assessment of the
33 exploitation rates of stocks within a mixed stock fishery. Even though the amount of tagging
34 data for small or heavily depleted stocks may be limited, there may exist additional biological
35 information on these stocks, not necessarily expressed quantitatively or as data series. By
36 analysing the mark-recapture data within a Bayesian framework, this additional information
37 can be incorporated to improve assessments of individual fish stocks.

38 This paper proposes a Bayesian state-space mark-recapture model to estimate the
39 fishing mortality rates of stocks caught within a mixed stock fishery, particularly when data
40 for some of the stocks are sparse. The next, i.e., second, section outlines some of the general
41 aspects of the methodology. The third section provides a background on Atlantic salmon
42 (*Salmo salar*) stocks from the north-eastern rim of the Baltic Sea. This section contains
43 subsections on the data, information about the population biology, the population dynamics
44 model, the observation model and the estimation of exploitation rates. The fourth and fifth
45 sections contain the results, and discussion and conclusions.

46 **Methods**

47 Several review papers have been written related to capture, recapture and removal
48 statistics for the estimation of population size and demographic parameters (Seber 1982;
49 Pollock 1991; Schwarz and Seber 1999). The early mark-recapture models applied in fisheries
50 used explicit maximum likelihood estimates when fitting a model to tagging data (Seber
51 1982). The use of mark-recapture methods within a Bayesian context is more recent (Gazey
52 and Staley 1986; Newman 2000; Mäntyniemi and Romakkaniemi 2002). Bayesian methods
53 allow researchers to use both quantitative data and qualitative information that may be
54 obtained from experts (expert opinions) or from previous studies (prior knowledge) (Malakoff
55 1999). This makes the Bayesian approach particularly useful in situations when data are
56 sparse and the associated uncertainty in population parameters is large (Ludwig et al. 2001).
57 The problem of sparse tagging data series is common in fisheries since most mark-recapture
58 experiments are done on an opportunistic basis, rather than based on careful tagging
59 experiments designed to facilitate ongoing fisheries stock assessments (Martell and Walters
60 2002). Within this paper a Bayesian mark-recapture model is proposed to estimate the fishing
61 mortality rates of stocks (including data-poor ones) within a mixed fishery. The Bayesian
62 approach explicitly deals with uncertainty, i.e., uncertainty in the parameter values and
63 uncertainty in the observations or data (Punt and Hilborn 1997). Natural variability in the
64 population dynamics is accounted for by applying a state-space formulation of the mark-
65 recapture model (Rivot et al. 2004).

66 The mark-recapture model consists of a population dynamics model, describing the
67 dynamics of the population of interest, and an observation model, describing how the
68 recapture data have been obtained (McAllister et al. 1999). The mark-recapture model uses
69 prior probability density functions (pdfs) for model parameters such as population parameters
70 (natural mortality rates, maturation rates and/or migration rates), fisheries related parameters
71 (catchability coefficients), data collection parameters (tag reporting rates) and for parameters

72 that convey process error in system dynamics and observation error in the data. These prior
73 pdfs reflect the prior beliefs about the values for these parameters and can be used together
74 with the data in Bayes' theorem to compute the joint posterior density function of the state-
75 space model parameters or the Bayesian probability that the parameter values are true given
76 the recovered tags (Gelman et al. 1995). The joint posterior density function therefore states
77 the degree of belief in values of state-space model parameters given the mark-recapture data.
78 In order to avoid possible confounding of the reporting rates with other parameters, the tag
79 reporting rate priors are not updated by the tagging data.

80 The Bayesian mark-recapture analysis is run using WinBUGS (Bayesian inference
81 Using Gibbs sampling) software (<http://www.mrc-bsu.cam.ac.uk/bugs>), version 1.4.
82 WinBUGS uses MCMC (Markov Chain Monte Carlo) methods to sample from the posterior
83 probability density function (Thomas et al. 1992). All of the modelling results described in
84 this paper have undergone tests to remove the 'burn-in' associated with the use of MCMC
85 methods and to assess convergence (Best et al. 1995). It is therefore assumed that the reported
86 distributions are representative of the underlying stationary distributions. In addition, the fit of
87 the model to the data has been assessed by comparing the data to the posterior predictive
88 distribution of the model i.e. the distribution of data simulated from the model (Gelman et al.
89 1995; Michielsens and McAllister 2004). The result of this comparison can be expressed in
90 terms of a Bayesian p-value (Meng 1994). Bayesian posterior predictive p-values indicate the
91 probability that the replicated data could be as extreme or more extreme than the observed
92 data. Alternative model structures have been compared using the Deviance Information
93 Criterion (DIC), which is a Bayesian measure of model complexity and fit (Spiegelhalter et al.
94 2002). The model that best replicates the data will have the smallest DIC.

95 **Application: estimation of the exploitation rates of Atlantic salmon**
96 **(*Salmo salar*) stocks in the Baltic Sea**

97 Wild Atlantic salmon stocks are exploited in the Baltic Sea by a sequence of six
98 salmon fisheries. While foraging in the Baltic Main Basin, salmon are captured by offshore
99 driftnet and offshore longline fisheries. Upon maturing, they migrate back to the rivers to
100 spawn. During their migration they may be captured by coastal driftnet, coastal trapnet or
101 coastal gillnet fisheries or by the river fishery, consisting predominantly of rod fishing.
102 Trends in wild salmon abundance cannot be ascertained from catch and fishing effort data
103 because of the large numbers of hatchery-reared salmon stocked annually in the Baltic Sea
104 and the paucity of records on the fraction of wild fish in annual catch records (Karlsson and
105 Karlstöm 1994; Romakkaniemi et al. 2003).

106 **Data**

107 Between 1987 and 2003, about 27000 wild salmon smolts have been tagged from the
108 rivers Tornionjoki and Simojoki, which are located at the north-eastern rim of the Baltic Sea
109 (Figure 1) (ICES 2005). Recapture records have been obtained for each of the main fisheries
110 in the Baltic Sea area. With a tag recovery rate of 5.6 %, the annual number of wild salmon
111 tag recoveries is relatively low, especially given the lack of tagging in 1989-90 and 1995-97.

Figure 1

112 The mark-recapture model therefore relies also on tagging data from related hatchery-reared
113 salmon stocks of the neighbouring rivers Iijoki and Kemijoki to facilitate parameter
114 estimation for wild salmon stocks (ICES 2005). It is assumed that these hatchery-reared
115 salmon stocks have similar sea migration patterns, i.e., to the feeding grounds at the Main
116 Basin and back to the spawning rivers (Romakkaniemi et al., 2003), and similar biological
117 characteristics as wild salmon stocks, with the exceptions specified in the next section. For
118 simplicity it is assumed that the wild stocks of the two rivers do not differ from each other nor
119 do the two hatchery-reared stocks differ from each other in any respect relevant for the

120 modelling. A total of around 170000 hatchery-reared salmon smolts have been tagged and
121 released in these rivers between 1987-2003 (ICES 2005).

122 In addition to the tagging data, the mark-recapture model also uses fishing effort data
123 as a covariate for fishing mortality rates (Figure 2). This reduces the number of estimated
124 parameters and may increase efficiency in parameter estimation (Seber and Schwarz 2002).

Figure 2

125 The unit of fishing effort is in gear-days (number of units of gear deployed (e.g. long line
126 hooks) x number of days deployed per year) (ICES 2005). Since the mid-nineties, fishing
127 effort on salmon in the Baltic Sea has decreased markedly. Since the mid-eighties about 10%
128 of the annual catch has been taken by river anglers. In absence of annual river fishing effort
129 data, only the long-term average fishing mortality rate in rivers can be estimated using a
130 single constant value for river fishing effort. Tables containing the data can be found in the
131 ICES working group reports for the assessment of Baltic salmon and trout (ICES 2002; 2005).

132 **Information on the population biology of wild and hatchery-reared stocks**

133 In addition to the mark-recapture data, there exists tagging independent information
134 about the life history characteristics of wild salmon stocks, especially in relation to their
135 hatchery-reared counterparts, e.g., the differences in age of maturation and in natural post-
136 smolt mortality (Salminen et al. 1995; Kallio-Nyberg and Koljonen 1997; Jutila et al. 2003).
137 It is of interest to find methods to incorporate such basic biological knowledge into stock
138 assessments (Ulltang 1996). This information has played a key role in the specification of the
139 model structure and the prior pdfs of model parameters.

140 Tagging data for both wild and hatchery-reared salmon are analysed together whereby
141 the model structure describes the relationship between certain parameters for wild and
142 hatchery-reared salmon. The maturation rate for wild grilse is assumed to be lower than for
143 hatchery-reared grilse due to the lower growth rate (Kallio-Nyberg and Koljonen 1997; Jutila
144 et al. 2003). This is implemented by multiplying a mean maturation rate for grilse respectively
145 with a yearly maturation effect for wild or hatchery-reared salmon. Maturation rates for wild

146 grilse are thereby allowed to be the same or smaller than maturation rates for hatchery-reared
147 grilse. In addition, the post-smolt mortality rate of hatchery-reared salmon is assumed to be
148 higher than that of wild salmon (Olla et al. 1998; Brown and Laland 2001). This is
149 implemented similarly as for the maturation rates. The post-smolt mortality rates are allowed
150 to differ from year to year (Salminen et al. 1995), and it assumed that these annual changes
151 are the same for both wild and hatchery-reared salmon. Unlike the post-smolt mortality rate,
152 the instantaneous natural mortality rate for adult salmon is assumed to be constant over the
153 years and the same for wild and hatchery-reared salmon.

154 Existing information about the salmon stocks is also incorporated by assigning prior
155 pdfs to biological model parameters. The prior pdfs for these parameters are obtained through
156 the use of expert knowledge about biological parameters for Atlantic salmon in general or
157 Baltic salmon in particular. When depending on expert judgment it is better to depend on a
158 group of experts (Punt and Hilborn 1997) and keep the methods to elicit prior information as
159 simple as possible (Chaloner 1996). For the analysis in this paper, twelve experts have been
160 asked to provide the most likely value and a minimum and maximum value for the biological
161 model parameters based on previous studies and relevant literature. Care has been taken to
162 ensure that the expert judgment is not based on data used within the mark-recapture model to
163 avoid using the same data twice and thus rendering the results too informative. The use of
164 multiple experts resulted in multiple priors for the different biological parameters. Each
165 expert was given the same weight when combining the priors from the different experts
166 through arithmetic pooling (Genest and Zidek 1986; Spiegelhalter et al. 2004). An overview
167 of the different model parameters and their prior pdfs is provided (Figures 3 and 4) as well as
168 a list of all the symbols used for the different model parameters (Table 1).

Figure 3

Figure 4

Population dynamics model

Table 1

170 The population dynamics model used within the mark-recapture analysis is age-
171 structured and assumes that all salmon are the same smolt age when tagged and released. The

172 offshore driftnet fishery (DF), offshore longline fishery (LF), coastal driftnet fishery (CDF),
 173 coastal fishery (CF) and river fishery (RF), are assumed to take place during sequential points
 174 in time (Figure 5). The population dynamics equations are of the following general form

175

Figure 5

176 (1)
$$N_{r,t_2,a} = N_{r,t_1,a} e^{-F_{f,y,a} - \Delta t M_{y,a}/12} \epsilon_{y,t}$$

177

178 whereby $N_{r,t,a}$ is the abundance of tagged salmon in month t during their a^{th} year at liberty
 179 after release in year r and $F_{f,y,a}$ is the instantaneous fishing mortality rate by fishery f in year
 180 y whereby $y = r + a - 1$. $M_{y,a}$ is the instantaneous natural mortality rate in year y . During their
 181 first year at liberty, salmon experience high natural mortality rates when migrating from the
 182 freshwater environment to sea (Salminen et al. 1995). The natural mortality rate during the
 183 first year at liberty i.e. the post-smolt mortality ($M_{y,1}$), is therefore different from the adult
 184 mortality rate which is assumed to be the same for different age groups ($a = 2$ to 4) and across
 185 years. Because equation 1 covers $t_2 - t_1 = \Delta t$ months, the yearly instantaneous natural
 186 mortality rate is adjusted to cover the same period ($\Delta t \cdot M_{y,a}/12$). In the coastal areas, it is
 187 assumed that the percentage of salmon mauled by seals has increased annually by 5.5%
 188 between 1995 and 2001, following the increase in the seal population (ICES 2002). In coastal
 189 areas, an additional seal related mortality factor ζ_y is used to increase the instantaneous
 190 natural mortality rate above the average rate.

191 The population dynamics model includes four different life history types that spend
 192 from one to four winters at the sea before returning to the rivers to spawn. Assuming tagged
 193 smolts migrate to sea immediately after being released, the number of years at liberty
 194 corresponds to the number of years at sea. Each year, a fraction of the salmon population will
 195 mature (L_a) and start migrating back to the river,

196

197 (2)
$$N_{r,t_4,a+1} = L_a N_{r,t_3,a} e^{-F_{f,y,a} - \Delta t \cdot M_{y,a}/12} \epsilon_{y,t}$$

198

199 while the immature salmon will remain another year at sea

200

201 (3)
$$N_{r,t_5,a+1} = (1 - L_a) N_{r,t_3,a} e^{-F_{f,y,a} - \Delta t \cdot M_{y,a}/12} \epsilon_{y,t} .$$

202

203 Salmon that return after 1 winter at sea are called 1-Sea-Winter (1SW) salmon or grilse. In
204 case they remain several years at sea before spawning, they are named Multi-Sea-Winter
205 (MSW) spawners. It is assumed that all salmon die after spawning. Wild and hatchery-reared
206 salmon are modelled as separate fish stocks without causal dependencies, even though some
207 of their life history parameters are linked in the sense that knowledge from the hatchery-
208 reared population is assumed to help in the assessment of wild salmon.

209 Deviations from the population dynamics model predictions are modelled within each
210 survival process by including a process error term $\epsilon_{y,t}$. In the absence of data, a symmetrical
211 uniform distribution around 1 is proposed for the process error in survival rates, whereby the
212 variance of the process error is made dependent on the size of the time step and on the
213 mortality rate Z . In general, state-space models use yearly time steps when modelling a
214 population. Because of the importance of within-year detail when modelling migratory
215 species exploited by several different fisheries, smaller than yearly time steps may be
216 required. Assuming that variance components are additive, the variance of the yearly process
217 error is divided by 12 and multiplied by the number of months over which the process error is
218 applied. The smaller the time steps, the smaller the variance.

219 In addition to the time-steps, the process error is also dependent on the total mortality
 220 rate Z , based on the assumption that without recruitment, the product of the total survival rate
 221 e^{-Z} and the process error ε_t should be smaller than 1 ($0 < N_{t+1} = N_t e^{-Z} \varepsilon_t < N_t$ and $e^{-Z} \varepsilon_t < 1$).
 222 By definition, the process error will therefore be larger than 0 and smaller than e^Z . Because
 223 the uniform prior pdf is assumed to be symmetrical around 1, the value closest to 1 will
 224 determine the minimum and maximum value of the uniform distribution. At each point during
 225 the life history, the process error is assumed to be the same for wild and hatchery-reared
 226 salmon. In case wild and hatchery-reared salmon have different survival rates, the smallest
 227 resulting process error is applied to both.

228 This proposed process error term differs significantly from earlier process error terms
 229 such as the one of Schnute and Richards (1995),

230

231 (4)
$$\varepsilon_t = \frac{e^{\sigma\delta}}{1 - e^{-M} + e^{-M} e^{\sigma\delta}}$$

232

233 which is derived from independent standard normal variates δ , natural mortality M , and a
 234 parameter σ related to the variance of ε_t . This process error term can be adjusted to be
 235 dependent on the total mortality rate. The process error of Schnute and Richards (1995) has a
 236 more pronounced peak around one compared to the formulation proposed in this paper. A
 237 flatter process error term accounts for more uncertainty in system dynamics and allows data to
 238 update the process error term more readily. When using the new process error term within the
 239 mark-recapture model, the WinBUGS program runs more than four times faster than when
 240 using the process error term of Schnute and Richards (1995), due to faster mixing MCMC
 241 chains. The new process error formulation thus appears to provide advantages over existing
 242 alternatives.

243 **Observation model**

244 The number of reported tags under-represents the total number of tagged salmon
245 caught because a proportion of the recaptured tags remains unreported. Several analyses have
246 been undertaken to try to estimate the tag reporting rates in the different salmon fisheries of
247 the Baltic Sea (ICES 2003). In general, these analyses are based on certain fishermen or
248 certain fleets and only give a first indication of the possible reporting rates in the different
249 fisheries. Expert judgment is used to extrapolate the resulting reporting rates to the entire
250 fisheries or to other fisheries, as experts are believed to have information about fisheries
251 which is not directly available from existing data sets.

252 Expert judgment about the tag reporting rates by fishermen of their national fishing
253 fleet was elicited from twelve experts. The experts based their judgments on data obtained
254 from these studies, information from literature and their own experiences and observations.
255 The expert information was obtained in the same way as for biological parameters but was
256 combined by weighting the pdfs for each nation by the nation's contribution to salmon
257 catches (Table 2). This reflects the assumption that reporting rates of tags are dependent on
258 the country of origin of the fishermen and that experts only have knowledge of reporting rates
259 by fishermen from their own country. Reporting rates and tag shedding rates for the offshore
Table 2 driftnet fishery and the coastal driftnet fishery are assumed to be the same.
260

261 The expected number of tags caught and reported by the different fisheries during the
262 salmon's a^{th} year at liberty is therefore given by the following equation

263

264 (5)
$$C_{f,r,a} = \gamma_f \lambda_f \varphi_f N_{r,t,a} e^{-F_{f,y,a} - M_{y,a}/24}$$

265

266 whereby $C_{f,r,a}$ is the expected number of caught and reported tags from salmon released in
267 year r and recaptured after a years at liberty by fishery f in the middle of the month ($M/12/2$),

268 γ_f is the probability that the fishermen will report the tags when recaptured by fishery f and
 269 λ_f is the probability that the salmon will retain the tags when caught by fishery f . Tag
 270 retention problems are assumed to occur only when catching the tagged salmon with driftnets.
 271 In order to account for tagged salmon removed from traps or nets by seals, a factor φ_f is used
 272 to adjust the reporting rate of tags recaptured by the coastal fishery. This adjustment factor
 273 should decrease the reporting rate of the coastal fishery annually by 5.5% between 1995 and
 274 2001.

275 It is assumed that the reported tags are distributed according to a negative binomial
 276 distribution, taking into account the schooling behaviour of the salmon (Christensen and
 277 Larsson 1979) and the somewhat patchy distribution of the total fishing effort. The following
 278 version of the negative binomial probability density function (pdf) was used for the
 279 probability that the number of recaptured and reported tagged fish equals c , given a particular
 280 set of model parameters (θ),

281

282 (6)
$$p(c|\theta) = \frac{\Gamma(k+c)}{\Gamma(k)c!} \left(\frac{k}{k+m}\right)^k \left(\frac{m}{m+k}\right)^c.$$

283

284 In this equation m represents the sample mean, i.e., the model-predicted number of tagged fish
 285 reported in a given fishery in a given year, and k represents the overdispersion parameter
 286 (Hilborn and Mangel 1997). It is important to note, that the overdispersion parameter has a
 287 direct biological explanation: it represents the propensity for schooling behaviour in salmon.
 288 Parameter k is therefore assumed to be the same across the years but to differ across fish
 289 stocks and fisheries. The variance in c given θ is determined through the equation

290

291 (7)
$$\text{Var}(c) = m + \frac{m^2}{k}.$$

292

293 The larger the value of the overdispersion parameter, the closer the approximation to a
294 Poisson distribution. Both the recaptured and reported hatchery-reared tagged salmon as well
295 as the recaptured and reported wild tagged salmon are assumed to follow a negative binomial
296 distribution.

297 **Estimation of the exploitation rates**

298 The main outputs of the model are the fishing mortality rates or exploitation rates
299 which are dependent on the fishing effort ($E_{f,y}$) and the catchability coefficients ($q_{f,a}$) of the
300 different fisheries, according to the following equation

301

302 (8)
$$F_{f,y,a} = q_{f,a} E_{f,y}.$$

303

304 The uncertainty over values for catchability plays a key role within assessments when
305 effort based management systems are applied. The catchability coefficients, $q_{f,a}$ or q , have
306 been estimated independently for different age groups in case the fisheries have different
307 efficiencies to catch different age groups as assumed to be the case for the driftnet, trapnet and
308 gillnet fisheries. It has been assumed that q can vary between different fisheries, and between
309 different age groups of wild and reared salmon and that there is an underlying distribution for
310 q across these groups. This has been implemented within the model by assuming a
311 hierarchical model structure defined through a mean catchability coefficient of fishery f for
312 fish of age a for the combined set of stocks ($\mu_{q,f,a}$) and a cross-stock variance of the

313 catchability coefficients ($\sigma_{q,f,a}^2$) (Gelman et al. 1995; Millar and Methot 2002). An overview
314 of the estimated catchability coefficients for the different fisheries can be found in Figure 4.

315 Selecting appropriate prior pdfs for these parameters can be difficult. Non-informative
316 prior pdfs for the catchability coefficients may result in bimodal distributions for the
317 corresponding harvest rates with peaks at 0 and 1. Therefore priors are placed on the harvest
318 rates for each fishery in the first year of the data series, i.e., 1987 ($H_{f,y=1987,a}$), and based on
319 the fishing efforts during that year, the prior pdfs for the corresponding catchability
320 coefficients are calculated.

321

322 (9)
$$q_{f,a} = \frac{-\log(1 - H_{f,1987,a})}{E_{f,1987}}$$

323

324 The prior pdf's for the harvest rates in 1987 are given by uniform distributions
325 between 0 and 1, Unif(0, 1). Some tagged salmon are caught incidentally immediately after
326 their release. Even though the associated fishing mortality is assumed to be small, these
327 tagged salmon need to be accounted for in order not to overestimate the survival rate. The
328 prior pdf of the incidental harvest rate during feeding migration is therefore given by Beta(1,
329 20). In combination with the fishing effort in subsequent years, the prior pdfs for the
330 catchability coefficients determine the prior pdfs for the harvest rates in subsequent years.

331 For each year the model estimates different harvest rates or fishing mortality rates
332 depending on the fishery, depending on the age of the fish and depending on whether it is a
333 wild or hatchery-reared fish. To present these values at this detailed level of disaggregation
334 would be confusing and not necessarily useful for management purposes. Instead the total
335 cumulative fishing mortality rates are reported for wild or hatchery-reared fish depending on
336 the number of winters (a-1) the salmon stay at sea. This total cumulative fishing mortality rate

337 relates to the total fishing pressure a fish is subjected to during its life history. For example a
 338 2-Sea-Winter (2SW) salmon can be caught by the river and coastal fishery when migrating to
 339 the feeding grounds (as a non-target species of miscellaneous types of fisheries), by the
 340 driftnet and longline fishery during its first and second winter at sea, by the coastal driftnet,
 341 trapnet and gillnet fishery during the migration to the spawning grounds and by the river
 342 fishery. The general formula for the total cumulative fishing mortality rate for 1SW to 4SW
 343 salmon is given by the following equation

344

345 (10)
$$F_{(a-1)SW,y} = \sum_{f=RF,CF} F_{f,y-a+1,1} + \sum_{f=DF,LF} \sum_a F_{f,y-a+1,a-1} + \sum_{f=CDF,CF,RF} F_{f,y,a}$$
, with a = 2 to 5.

346

347 The total cumulative fishing mortality rate thus spans the entire life history of the
 348 salmon and it shows directly how large the impact of fishing is on the spawning capacity
 349 compared to an unexploited situation. The total cumulative fishing mortality rates only reflect
 350 the mortality due to fishing and do not include any natural mortality. To simplify the
 351 interpretation of the results, cumulative fishing mortality rates are expressed as total harvest
 352 rates.

353 **Results**

354 The prior pdfs of model parameters for hatchery-reared salmon have been updated
 355 considerably due to utilizing informative tagging data for reared salmon (Figures 3 and 4).
 356 For wild salmon, the prior pdfs for the maturation rates and natural mortality rates, have been
 357 updated to a lesser extent because the priors had already been quite informative and the
 358 information available in the tagging data for wild salmon was limited. The annual posterior
 359 estimates for the post-smolt mortality rate of wild and reared salmon are shown in Figure 6.

Figure 6

360 There is a trend in the results indicating that post-smolt mortality rates have been higher in

361 recent years. The reasons for this shift in post-smolt mortality rates are still unclear but may
362 be linked to environmental factors (Kallio-Nyberg et al. 2004).

Figure 7
363 The main outputs of the model are the marginal posterior pdfs for total harvest rates
364 for the different life history types (Figure 7) based on the catchability coefficients of the
365 different fisheries for different age groups. The priors for the catchability coefficients and the
366 derived priors for the harvest rates have been updated considerably due to the information
367 contained in the tagging data and the fishing effort data. The longer the salmon stay out at sea,
368 the higher the chance that they will be captured by the fishery and the higher the
369 corresponding harvest rate. The total harvest rates for 2SW wild salmon have been compared
370 to the preliminary precautionary reference point (ICES 2002). The total harvest rate of 2SW
371 wild salmon is higher than the precautionary reference point for 2SW salmon, indicating that
372 from a fisheries management perspective exploitation rates are too high.

373 Several diagnostic measures have been calculated. The posterior predictive
374 distribution of model quantities indicates whether the model's predictions are plausible given
375 the observed data. About 1.8% of the observed data points were located outside the 95%
376 probability intervals of the posterior predictive distributions. This indicates that the data could
377 have been obtained from the model and the posterior pdfs of model parameters. When
378 calculating Bayesian p-values, data points with a value of 0 have been excluded. These data
379 points would result in Bayesian p-values of 1 when assuming a negative binomial likelihood
380 function. For the remaining data points, only 2.6% of the posterior predictive p-values are
381 larger than 0.975 and 2.1% are smaller than 0.025.

382 One of the main concerns when using tagging data in stock-assessments is the
383 uncertainty regarding the tag return rates (Hilborn and Walters 1992). In this paper, the tag
384 return rate parameters (i.e., tag reporting and retention rates) have been based on partial data,
385 extrapolated through expert judgement. In order to assess the impact of the choice of the prior
386 pdfs for the tag reporting and retention rates on the resulting estimates for the exploitation

387 rates, three different scenarios have been compared against the base case. First it has been
388 assumed that less information was available regarding the tag return rates resulting in less
389 informative prior pdfs. This has been implemented by doubling the CV of the prior pdfs for
390 the tag return rate parameters. Secondly more informative prior pdfs with half the original
391 CV's have been specified in order to assess the benefit of investing in studies to provide more
392 precise estimates of parameters determining the tag return rates. Thirdly, the parameters
393 determining the tag return rates have been given prior pdfs with different means, assuming the
394 means of the pdfs for the tag return rate parameters are one third lower than under the base
395 case scenario.

Figure 8

396 The resulting probability density functions for the cumulative harvest rate of 2SW fish
397 returning in 2004 are presented (Figure 8, panel a). The four posterior pdfs overlap but there
398 are distinct differences between them. As could be expected, less informative prior pdfs for
399 tag return rates result in a wider posterior pdf for the total harvest rate with a higher
400 probability of very high total harvest rates. More informative prior pdfs for tag return rates
401 result in slightly more informative estimates of the exploitation rate. When tag return rates are
402 assumed to be lower than was the case for the base case scenario, the total harvest rate needs
403 to be higher in order to obtain the same number of reported tagged salmon in the catch. In
404 general it can be concluded that the amount of uncertainty in the prior pdfs of parameters
405 determining the tag return rates has a clear impact on the uncertainty in the estimates of the
406 total fishing mortality rate. Greater uncertainty in these prior pdfs will result in higher
407 estimates of the total cumulative exploitation rates. This would lead to more restrictive
408 fisheries management advice. These arguments underline both the importance of including
409 uncertainty in the values used for tag return rates and the importance of investing in studies to
410 reduce as much as possible the uncertainty in the prior pdfs for the tag return rates. Such
411 studies could include scientific observer programs and high reward tagging studies. Similar
412 sensitivity analyses for the maturation rates and natural mortality rates (Figure 8, panel b and

413 c) indicate the limited impact of the priors pdfs for the maturation rates and natural mortality
414 rates on the posterior probability distributions for the harvest rates.

415 The use of a negative binomial likelihood function was compared to alternative model
416 structures that use Poisson, binomial and beta-binomial pdfs for the reported tags. The
417 Poisson distribution is a special case of the negative binomial distribution, obtained by
418 assuming a random spatial and temporal distribution of fish and reported tags instead of a
419 clustered distribution. The beta-binomial pdf is more realistic than the negative binomial
420 distribution because it assigns zero probability to catches higher than the number of tagged
421 fish. When tag return probabilities are low, the beta-binomial pdf approximates the negative
422 binomial distribution. The binomial distribution is a special case of the beta-binomial
423 distribution that assumes a random spatial and temporal distribution of fish and tags. The DIC
424 for the scenario based on negative binomial likelihood functions was 4315 while the DIC for
425 the models using Poisson, binomial or beta-binomial pdfs for the likelihood functions were
426 6287, 6641 and 4468 respectively. This indicates that models assuming a clustered
427 distribution of fish obtain a better fit to the mark-recapture data than models assuming a
428 random distribution. The model using the negative binomial distribution gives a slightly better
429 fit to the mark-recapture data than the model assuming a beta-binomial distribution. When
430 calculating the DIC, no negative values have been obtained for the effective number of
431 parameters, suggesting there is no indication of conflicts between the priors and the data or of
432 problems related to the parameterisation of the model.

433 **Discussion**

434 We have demonstrated that when reported recapture rates are sufficiently high (e.g., >
435 5%), as in the case of salmon fisheries, mark-recapture analyses can allow for the estimation
436 of annual fishing mortality rates. Yet, few fishery stock assessments actually use the tagging
437 data in mark-recapture analyses and take advantage of the fact that mark-recapture analyses
438 do not necessarily need catch or catch-per-unit-effort data. Many fisheries stock assessments

439 however use tagging data to examine stock structure (Kohler and Turner 2001), movement or
440 migration patterns (Sibert et al. 1999; McGarvey and Feenstra 2002) and the allocation of
441 quotas (Caron et al. 2002). Tagging data are currently also used in a number of instances to
442 help estimate harvest rates on fish stocks. For example, the incorporation of tagging data in
443 MULTIFAN-CL (Hampton and Fournier 2001), a methodology which analyses length-
444 frequency distributions of catches, facilitates the estimation of harvest rates. Mark-recapture
445 data are also analysed using conventional mark-recapture methods (e.g., Jolly-Seber) to help
446 estimate harvest rates for some stocks, for example, individual Canadian Atlantic cod stocks
447 (Cadigan and Bratley 2002) and north-east Atlantic mackerel (Skagen 2003). In contrast, this
448 paper has presented a Bayesian state-space methodology to estimate harvest rates on
449 individual stocks imposed by a sequence of mixed-stock and single stock fisheries.

450 One of the main factors constraining the use of tagging data for stock assessment
451 purposes is the uncertainty over reporting rates of tags by fishermen (Hilborn and Walters
452 1992). By analysing tagging data within a Bayesian setting, the uncertainty regarding the
453 reporting of tagged fish can be taken into account. The information for the probability
454 distributions of the reporting rates can be obtained from studies designed to estimate the
455 reporting rates, from expert opinion or a combination of the two. Using this approach, mark-
456 recapture analyses can provide estimates of exploitation rates independent from catch data. By
457 incorporating mark-recapture analyses in conventional fisheries stock-assessments (Patterson
458 1999; Punt et al. 2000), the results become more robust (Martell and Walters 2002).

459 In contrast to many other Bayesian state-space formulations, this paper follows a
460 different convention in which the prior is not placed on the state variable, e.g., abundance
461 (Millar and Meyer 2000), but instead on the process error term itself. This latter convention
462 has already been implemented in a number of papers (McAllister et al. 1994; McAllister and
463 Ianelli 1997). By placing priors on the process errors terms, $\varepsilon_t \sim \text{Norm}(0, s^2)$, rather than on the
464 state variable, $N_t \sim \text{logNorm}(\text{fn}(N_{t-1}), s^2)$, the computational efficiency and convergence

465 properties improve due to removal of the highly correlated state variables from the set of
466 parameters estimated by the MCMC algorithm (Cunningham 2002).

467 The survival rate process error model also offers an alternative functional form for the
468 survival rate process error that efficiently keeps to the constraint that $N_{t+1} \leq N_t$. Schnute and
469 Richards' (1995) process error formulation keeps to the same constraint but is
470 computationally less efficient. Lewy and Nielsen (2003) offer a lognormal process error term
471 for survival rate processes. This density function however needs to be truncated to prevent
472 $N_{t+1} > N_t$ and produces more positively skewed density functions than the other two.

473 For the assessment of mixed-stock fisheries, conventional datasets such as catch data
474 and catch-per-unit effort data offer little information on the individual stocks. Mark-recapture
475 data can provide this kind of information. However, tagging data are often more sparse for
476 overexploited stocks than for more abundant stocks. The Bayesian approach taken facilitates
477 the estimation of exploitation rates of stocks within mixed stock fisheries, even when data are
478 sparse. The methodology utilizes knowledge about key differences and similarities between
479 the fish stocks of interest and biological information about the individual fish stocks based on
480 published literature, expert judgment. This prior knowledge is combined with tagging and
481 effort data to help estimate exploitation rates of each of the fish stocks.

482 The use of data from similar or related populations is a well-known concept within
483 ecology where data from similar populations are combined within meta-population analyses
484 in order to estimate their population parameters and predict the parameters for unsampled
485 populations (Gurevitch et al. 2001). The dependence among parameters that vary predictably
486 among populations can be modelled though the use of a hierarchical model structure (Gelman
487 et al. 1995). Within this paper, hierarchical structures have been used to model stock
488 dependent catchability coefficients (q). The use of an hierarchical structure however requires
489 that the individual stocks are exchangeable i.e. that the differences between the individual
490 stocks should be unpredictable (Gelman et al. 1995). For the example presented within this

491 paper, this might be the case for the efficiency of the fisheries to catch different stocks. This is
492 not the case for other model parameters where prior information about the stocks already
493 dictates that the model parameter of one stock cannot be larger than the model parameters for
494 another stock. In such cases, it is more appropriate to use an additional parameter, indicating
495 the difference between the parameter values for both stocks.

496 The use of prior information is an integral part of Bayes' theorem (Gelman et al.
497 1995). However, some scientists believe that the use of subjective priors, e.g., those derived
498 from experts, is inappropriate and unscientific because these priors may be incorrect or biased
499 (Cox and Hinkley 1982). To reduce the impact of subjective beliefs, the use of uninformative
500 (e.g., flat or low gradient) priors has often been recommended (Walters and Ludwig 1994).
501 An uninformative prior for particular parameters may however lead to informative priors for
502 other variables of interest and, when data are relatively uninformative, this may lead to results
503 that are inconsistent with biologists' understanding about the population (Punt and Hilborn
504 1997). Uninformative priors for q for example may lead to informative and spurious priors for
505 the associated harvest rates. In contrast, and as demonstrated in this paper and others (e.g.,
506 McAllister et al. 1994), the use of informative priors that have been carefully constructed
507 using available knowledge and data other than those to which the model has been fitted will
508 lead to more scientifically credible and defensible results. Therefore management actions can
509 more easily be justified in a precautionary context whereby the increases in uncertainty can
510 help to justify more restrictive management advice.

511 When applying the assessment methodology to Baltic salmon stocks, links have been
512 built between the life history parameters of wild and hatchery-reared stocks. Although wild
513 and hatchery-reared salmon have somewhat different life histories (Kallio-Nyberg and
514 Koljonen 1997; Jutila et al. 2003), certain population parameters can be regarded as similar or
515 related while others are assumed to differ. In the current model it has also been assumed that
516 within the group of wild stocks and within the group of reared stocks there are no differences

517 in biological characteristics or in harvesting patterns. In our example, the rivers are located
518 next to each other, and the same stock (Iijoki salmon) has been partly used for releases in both
519 of the dammed rivers. The findings of Järvi (1938; 1948) indicate that the differences in the
520 characteristics of the salmon stocks within the groups in the study area are relatively small. In
521 this case the impact of this simplifying assumption is likely to be small. When expanding the
522 model to include more stocks, an hierarchical modelling approach should be considered since
523 local adaptations can lead to differences in migration patterns for different stocks within the
524 group of wild or reared salmon (Alm 1934; Power 1981; Jonsson et al. 1991).

525 In addition there are several simplifying yet incorrect assumptions about the fisheries
526 and stock characteristics. The number of repeat spawners for example has increased over
527 time. During the last few years, up to 15% repeat spawners have been sampled in the river
528 Tornionjoki (Romakkaniemi et al. 2003). The catchability coefficient of the driftnet fishery
529 may also have changed systematically over time (ICES 2003). In addition, the fishing effort is
530 assumed to be known without error (Pollock 2002) and the process errors are assumed
531 independent. At this stage of the methodological development, these facts have not been
532 accounted for within the model.

533 Within the current example, no catch data have been used because the mark-recapture
534 model only relates to stocks from rivers located in the north-east of the Baltic Sea while the
535 catch data of the mixed stock fishery relate to all Baltic salmon stocks. Because the
536 exploitation rates of salmon stocks differ depending for example on the location of the river
537 where they spawn within the Baltic Sea, it is not possible to extrapolate the results of the
538 exploitation rates to the entire Baltic Sea, and to estimate the salmon abundance based on the
539 total Baltic salmon catch data. Instead abundance estimates could be obtained by linking the
540 current exploitation rates with estimates of wild smolt production, as obtained by Mäntyniemi
541 and Romakkaniemi (2002), in order to estimate the number of salmon returning to the rivers
542 for spawning (ICES 2005). Alternatively, genetic stock identification methods can be used to

543 estimate the proportion of the catch originating from certain groups of stocks (Koljonen and
544 Pella 1997; Koljonen et al. 2004; ICES 2005).

545 Even without the estimation of wild salmon abundances and even though there are
546 several simplifications in the model, the current estimates of harvest rates are useful for
547 management purposes by comparing the estimates with limit or precautionary harvest rate
548 reference points (ICES 2002; Martell and Walters 2002). Since 2002, the current mark-
549 recapture methodology has been used for this purpose within the ICES working group for the
550 assessment of Baltic salmon and trout (ICES 2002).

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Table 1: List of symbols used within the model

Indices	
y	Year
t	Month
r	Release year
a	Years at liberty i.e. years at sea
f	Fishery
Model parameters	
$M_{y,1}$	Instantaneous natural post-smolt mortality rate in year y (year^{-1})
$M_{y,a \neq 1}$	Instantaneous natural adult mortality rate (year^{-1})
L_a	Proportion of salmon that mature after a years at sea
$q_{f,a}$	Catchability coefficient or efficiency of fishery f to catch salmon during their a^{th} year at sea (geardays^{-1})
$\varepsilon_{y,t}$	Process error term
γ_f	Probability that the fishermen will report the tags when recaptured by fishery f
λ_f	Probability that the salmon will retain the tags when caught by the fishery f
φ_f	Adjustment factor for the reporting rate in the coastal fishery f to account for the tagged salmon removed from the traps or nets by seals
ζ_y	Adjustment factor for the instantaneous natural mortality rate to account for increased predation by seals in coastal areas in year y
Model variables	
$N_{r,t,a}$	Abundance of tagged salmon in month t of the a^{th} year at liberty after release in year r
$C_{f,r,a}$	Expected number of tags caught and reported by fishery f during the salmon's a^{th} year at liberty after release in year r
$F_{f,y,a}$	Instantaneous rate of fishing mortality in year y by fishery f on salmon that spend a years at sea (year^{-1})
$H_{f,y,a}$	Harvest rate in year y by fishery f on salmon that spend a years at sea
Data	
$E_{f,y}$	Fishing effort by fishery f in year y (geardays)
$C_{f,r,a}^{\text{obs}}$	Observed number of tags caught and reported by fishery f during the salmon's a^{th} year at liberty after release in year r

Table 2: Summary of tag return rate parameters, their prior probability density functions (pdfs) and the corresponding prior median, coefficient of variation (CV) and 95% probability interval (PI). The pdfs follow the same parameterisation as used within the WinBUGS program. Beta denotes a beta pdf determined by two shape parameters. These pdfs have been truncated (e.g. $I(a,b)$) to indicate the prior belief that the random variable can not be smaller than a or larger than b.

Parameters	Distribution	Median	CV	95% PI
Tag retention rate	Beta(20,8) $I(0.5,1)$	0.72	0.11	0.55 - 0.86
Tag reporting rate in the river fishery	Beta(16,6) $I(0.3,0.95)$	0.73	0.13	0.53 - 0.89
Tag reporting rate in the coastal fishery	Beta(11,9) $I(0.3,0.8)$	0.55	0.19	0.35 - 0.75
Tag reporting rate in the driftnet fishery	Beta(8,4) $I(0.2,0.95)$	0.68	0.20	0.39 - 0.89
Tag reporting rate in the longline fishery	Beta(10,4) $I(0.3,0.95)$	0.72	0.16	0.46 - 0.91

Legends to Figures

Figure 1. Migration route of Atlantic salmon stocks (*Salmo salar*) from the rivers Torne (Tornionjoki), Simojoki, Kemijoki and Iijoki in Sweden and Finland. The driftnet and longline fisheries take place predominantly in the Baltic Main Basin while the trapnet and gillnet fisheries take place in the Gulf of Bothnia. The presence of dams in the rivers Kemijoki and Iijoki, which prevents access to spawning grounds, are indicated by lines across the rivers.

Figure 2. Fishing effort of the driftnet, longline, trapnet and gillnet fisheries on Atlantic salmon stocks (*Salmo salar*) between 1987 and 2004. River fishing effort is assumed to be constant over time. The unit of fishing effort is in gear-days (number of units of gear deployed x number of days per year).

Figure 3. Overview of the prior (dotted lines) and posterior (solid lines) probability density functions for maturation rates and instantaneous natural mortality rates of 1-3 Sea-Winter (SW) wild and hatchery-reared Atlantic salmon (*Salmo salar*) in the Baltic Sea area.

Figure 4. Overview of the prior (dotted line) and posterior probability density functions for the catchability coefficients (10^{-3} gear-days⁻¹ x year⁻¹) of 1-4 Sea-Winter (SW) wild (posterior – solid line) and hatchery-reared (posterior – dashed line) Atlantic salmon (*Salmo salar*) in the Baltic Sea area by the offshore driftnet and longline fishery, the coastal trapnet and gillnet fishery and the river fishery. The catchability coefficients of the coastal driftnet fishery are assumed the same as for the offshore driftnet fishery.

Figure 5. Schematic presentation of the mark-recapture model for Atlantic salmon (*Salmo salar*) in the Baltic Sea area. The offshore driftnet and longline fisheries in the Baltic Main Basin are assumed to take place in October and December respectively. During the migration to the spawning grounds, the salmon can be intercepted by the coastal driftnet fishery in May, the trapnet and gillnet fisheries in June and the river fishery in August.

Figure 6. Medians and 95% probability intervals for the annual estimates of post-smolt mortality rates (% per year) for wild and hatchery-reared Atlantic salmon (*Salmo salar*) in the Baltic Sea area between 1987 and 2004.

Figure 7. Medians and 95% probability intervals for the total cumulative harvest rate for 1-4 Sea-Winter (SW) wild and hatchery-reared Atlantic salmon (*Salmo salar*) in the Baltic Sea area. The total cumulative harvest rate for 2SW wild salmon can be compared against the precautionary harvest rate reference point (ICES 2002).

Figure 8. Posterior probability density functions of the total harvest rate for 2SW wild Atlantic salmon (*Salmo salar*) returning to north-eastern Baltic Sea rivers in 2004 when using different prior probability density functions for model parameters determining (a) tag return rates i.e., tag reporting rates and tag retention rates, (b) maturation rates and (c) natural mortality rates.















