Population diversity in Kuskokwim Chinook: New findings on trade-offs for different harvest strategies

A research team funded by the Arctic-Yukon-Kuskokwim Sustainable Salmon Initiative used estimates of Chinook population diversity to build computer simulations that evaluated how well alternative harvest policies meet Chinook population diversity and fishery objectives in the Kuskokwim.

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Variation among salmon populations in their productivity, carrying capacity and life history characteristics (population diversity) contributes to stable fisheries as a result of portfolio effects, whereby fisheries that harvest multiple populations benefit from the averaging effects of their variable dynamics (**Figure 1**). This diversity also supports fishery resilience because typically at least some populations will do well when others do not and populations that are less productive today might be more productive under future conditions.

As a result population diversity is increasingly viewed as a foundation of sustainable and resilient resource management and the importance of protecting population diversity is recognized in fisheries management policies such as <u>Alaska's</u> <u>Sustainable Salmon Policy</u>.

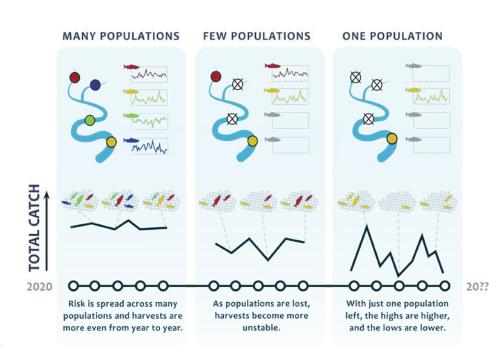
When mixed-stock fisheries for multiple populations overlap in space and time there can be an inherent conflict between harvest rates and population diversity: high harvest rates, which can be sustained by the most productive populations, can come at the cost of increased risk of overfishing less productive ones (**Figure 2**).

Despite the potential importance of these harvestpopulation diversity tradeoffs, Chinook salmon management, including in the large river basins of

Figure 1. Illustration of how population diversity contributes to harvest stability. When diversity is high, individual populations doing very well can compensate for those that are doing poorly, leading to a more stable harvest over time. When diversity is low, harvests are more variable because there are fewer populations to buffer the effects of a variable environment.

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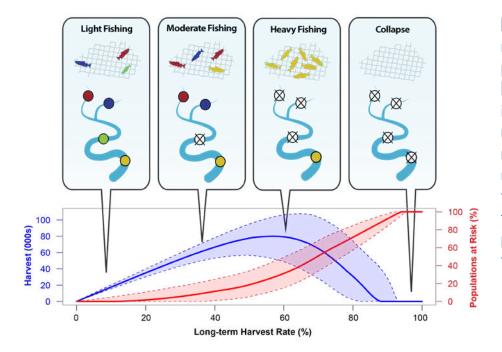


Figure 2. Relationship between long-term mixed-population harvest and risks to individual populations across a range of long-term average harvest rates. Overall harvest from the system is predicted to increase as the average harvest rate increases up to a point, after which it rapidly declines because most populations are overfished and then driven to extinction. This figure illustrates that you have to be willing to accept some risk to the weakest (least productive) populations if you want to maximize total harvest from the system.

Western Alaska and the Yukon, does not often explicitly consider them. To help address this knowledge gap we characterized the degree of Chinook population diversity within the Kuskokwim River basin by fitting spawner-recruitment models to all available tributary level data on Chinook escapement along with estimates of age composition and total harvest (13 populations accounting for approximately half of the total production from the Kuskokwim). We found clear evidence of population diversity in the system where population productivity and size were estimated to vary by as much as 3-fold and 18-fold among populations, respectively (**Figure 3**).

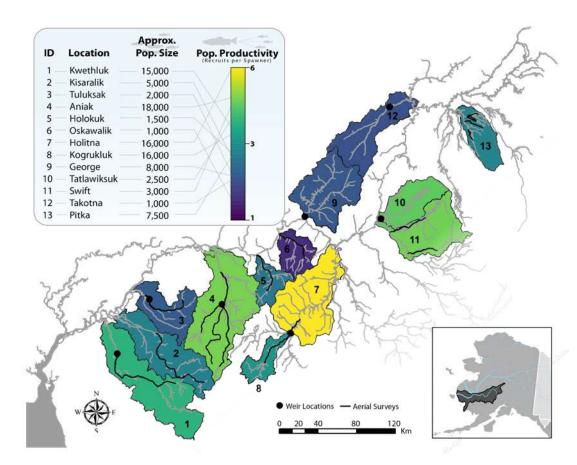


Figure 3. Chinook population diversity in the Kuskokwim River basin. Each polygon depicts the general spawning distribution of the 13 Chinook populations for which there are estimates. of spawner abundance based on either weir (points) or aerial (thick line) surveys. Populations are color coded by their productivity (recruits produced per spawner at small population size) and estimated average population size (predicted in absence of fishing) is listed next to each system's name. This biodiversity emphasizes that large populations are not necessarily the most productivity, and vice-versa.

We used estimates of Chinook population diversity to build computer simulations that evaluated how well alternative harvest policies (defined by basin wide escapement and harvest goals) meet Chinook population diversity and fishery objectives in the Kuskokwim. We found that harvest policies focused on meeting minimum subsistence needs were unlikely to jeopardize long-term prospects for basin-wide sustainable use. However, Chinook population diversity gives rise to asymmetric tradeoffs among fishery and conservation objectives in the Kuskokwim. For example, relative to a harvest policy that seeks to maximize mixed-stock harvests, foregoing relatively small amounts of mixed-stock harvest is predicted to yield relatively large increases in the chances of ensuring equitable access to Chinook (i.e., meeting tributary goals) and to nearly eliminate biological risks to weak populations (Figure 4).

The approach we developed for the Kuskokwim provides a general framework for characterizing salmon population diversity in large river basins and evaluating harvest-population diversity tradeoffs among alternative harvest policies within them. With support from AYK-SSI we are now exploring Chinook salmon population diversity in the Yukon River basin. Ongoing research analyzing Chinook salmon ear stones (otoliths) in the Kuskokwim and Yukon suggests that different parts of these large watersheds are hot spots for salmon production and growth, and that favorable locations change year to year. Together with our efforts to characterize Chinook population diversity, this research emphasizes the importance of protecting and monitoring salmon habitats and populations (both large and small) throughout these large free flowing river basins to maximize their resilience to environmental change and the benefits communities derive from them.

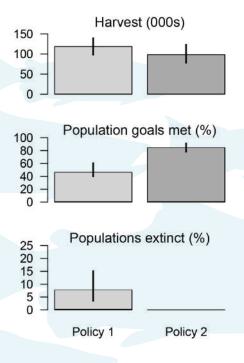


Figure 4. Illustration of the predicted performance of two alternative harvest policies for Kuskokwim Chinook. Policy 1 seeks to maximize total mixed-population harvest while Policy 2 seeks to maximize harvests only once the risk of driving the weakest populations extinct is minimized. Shown for each policy is (1 – top panel) the predicted average harvest, (2 – middle panel) proportion of individual populations whose average spawner abundance is above a population (or tributary) level goal which is also a proxy for equitable access to Chinook, and (3 – lower panel) the proportion of populations predicted to be driven to extinction.

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I. AYK SSI TITLE PAGE:

2019 Arctic-Yukon-Kuskokwim Sustainable Salmon Initiative Project Final Product*

Chinook harvest-population diversity tradeoffs

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II. ABSTRACT:

Variability among fish populations in ecological characteristics and productive capacity (i.e., population diversity) can be critical to maintaining resilience to environmental change and to dampening variability in harvest for fisheries that depend upon them. However, when fisheries for multiple populations overlap in space and time there can be an inherent conflict between mixed-stock harvest and population diversity: high harvest rates, which can be sustained by the most productive populations, can come at the cost of increased risk of overfishing those that are less productive. The importance of these harvest-population diversity trade-offs in salmon management is well-recognized, but they are difficult to characterize due to incomplete information, and they are not often explicitly evaluated in contemporary fisheries management. We developed an integrated age-structured multi-population state-space spawner-recruit model and fitted it to data from 13 Chinook salmon populations from the Kuskokwim drainage in western Alaska. We found clear evidence of population diversity in the system where productivity and carrying capacity can vary by as much as 3-fold and 18-fold among populations, respectively. Simulation testing of the model illustrated that it was largely unbiased with respect to leading parameters, abundance states, and derived biological reference points, whereas more commonly applied regression-based approaches showed substantial bias. We used the state-space model to parameterize closed-loop simulations that evaluated how well alternative harvest policies meet Chinook population diversity and fishery objectives in the Kuskokwim. We found that Chinook population diversity gives rise to asymmetric trade-offs among fishery and conservation objectives whereby foregoing relatively small amounts of mixed-stock harvest resulted in relatively large increases in the chances of ensuring equitable access to Chinook (i.e., meeting tributary goals) and nearly eliminated the risk of weak stock extirpation. We also found that harvest policies focused on meeting minimum subsistence needs were unlikely to jeopardize long-term prospects for basin-wide sustainable use. The fishery and biological performance of alternative harvest policies, and the magnitude of resulting trade-offs, were moderately sensitive to potential future changes in population productivity and capacity and to uncertainty in the underlying drivers of recruitment variation. Our approach provides a general framework for characterizing salmon population diversity in large river basins and evaluating harvest-population diversity trade-offs among alternative harvest policies within them.

III. PROJECT EVALUATION:

Our original proposal had six objectives; a seventh objective was added in consultation with the AYK-SSI Science Coordinator as the project was being carried out. These objectives, and our progress towards meeting them, are described below.

Objective 1: To determine the shape of the relationship between spawner abundance and adult recruitment for each individual Chinook spawning population in the Kuskokwim watershed.

We met this objective by developing an integrated age-structured multi-population state-space spawnerrecruit model modelling framework that is described in detail in the manuscript in Appendix A. See Figure 5 in the manuscript for an illustration of the inferred shape of these relationships across a range of alternative model formulations. Simulation testing of the model illustrated the state-space model was largely unbiased with respect to leading parameters, abundance states, and derived biological reference points, whereas more commonly applied regression-based approaches showed substantial bias.

Objective 2: To determine the predicted average annual fishery yield across all Kuskokwim Chinook populations - the sum of the equilibrium harvest that each population could sustain, and the proportion of populations that are predicted to not be overexploited - in order to evaluate the tradeoff between harvest and conservation of population diversity across a range of mixedpopulation harvest rates.

We met this objective as detailed in the manuscripts in Appendix A and B (see for example Figure 7 in Appendix A and Figure 3 in Appendix B).

Objective 3: To determine the extent to which the erosion of population diversity resulting from overharvest of weak populations may limit long-term fishery yield under changing environmental conditions that might favor what are now weak populations over those that are currently strong.

We met this objective by developing empirically parameterized closed-loop simulations of the Kuskokwim system where we evaluated the fishery and biological consequences of alternative harvest policies under a scenario with large future change in Chinook population diversity (i.e., weak populations became strong, and strong ones became weak). We found that when the environment drives pronounced shifts over time in the size and productivity of populations in the system (e.g., a regime shift) giving up a specific amount of harvest results in larger gains in spatial equity in access to the salmon for subsistence needs relative to scenarios where there is not a regime shift. These findings are described in detail in the manuscript in Appendix B.

Objective 4: To determine both fishery manager and stakeholder perspectives on acceptable conservation constraints (i.e., which point[s] on the curve in Figure 1 is acceptable) and fishery objectives (e.g., minimize frequency of fishery closures) in order to inform the estimation of optimal harvest policies in objective 5.

We met this objective by piggy backing on a broader exercise focused on building capacity among Kuskokwim stakeholders to engage in salmon management through a series of workshops with influential community members from throughout the river basin with a long history of active engagement in fishery management as well as US Fish and Wildlife Service and Alaska Department of Fish and Game biologists and fishery managers. These workshops included discussion of existing and potential Kuskokwim Chinook fishery and biological objectives and alternative management actions associated with them. From these discussions, we identified a suite of long-term objectives against which to quantify the performance of alternative management actions in the system. This is detailed in the manuscript in Appendix B.

Objective 5: To determine optimal harvest policies for Kuskokwim Chinook based upon the magnitude of persistent productivity changes in Kuskokwim Chinook and the conservation constraints and fishery objectives from Objective 4.

We met this objective by quantifying the performance of alternative harvest policies against objectives from Objective 4 using closed-loop simulations across a range of plausible hypotheses representing alternative states of nature. This is detailed in the manuscript in Appendix B. It is important to note however that our analysis do not identify an "optimal" strategy but instead help uncover trade-offs amongst objectives across a range of strategies.

Objective 6: To determine the harvest and population diversity costs of assuming Kuskokwim Chinook recruitment is driven by (1) over-compensatory processes or (2) extrinsic environmental forcing when the alternative hypothesis is true.

We fully met this objective by quantifying the performance of alternative harvest policies against the objectives from Objective 4 using closed-loop simulations for scenarios where recruitment is driven by either over-compensatory processes or extrinsic environmental forcing that gives rise to the appearance of overcompensation. We found that absolute harvest and its inter-annual stability were predicted to be lower, and that more harvest had to be foregone to minimize extirpation risk and increase chances of

ensuring equitable access to Chinook, under the extrinsic environmental forcing scenario relative to the over-compensatory one. These findings are detailed in the manuscript in Appendix B.

Objective 7: To generate infographics that communicate key insights from the proposed research.

We met this objective by developing a web-based tool that allow for interactive visualization of the conservation and fishery outcomes of our closed-loop simulations (see deliverables). In addition we generated numerous figures and visuals for the manuscripts and associated presentations that helped to communicate key insights from the proposed research (see figure in Appendix A, B and C).

IV. DELIVERABLES:

The findings of our project have been, and will continue to be, disseminated via conference presentations, management meetings and peer-reviewed manuscripts. To date we have given six presentations and attended five meetings where we have presented and discussed our research. We anticipate submitting two manuscripts by the fall of 2019 and further refining the web visualization. Copies of presentations are available from the project PIs upon request.

Presentations:

- Staton B., M. Catalano, B.M. Connors, L. Coggins, M. Jones, C. Walters, S. Fleischman and D. Gwinn. 2019. State-space models for estimating population diversity in mixed-stock Pacific salmon fisheries. American Fisheries Society Annual Meeting.
- Staton B., M. Catalano, B.M. Connors, L. Coggins, M. Jones, C. Walters, S. Fleischman and D. Gwinn 2019. Assessment approaches for mixed-stock Pacific salmon fisheries: Empirical and simulation-estimation applications. Auburn University; dissertation seminar
- Connors B. M., L. Coggins, B. Staton, C. Walters, M. Jones, M. Catalano and J. Harding. 2018. Incorporating harvest-population diversity tradeoffs into salmon management in large river basins: insights from Chinook in the Kuskokwim and Yukon River Basins. Western Division American Fisheries Society Meeting.
- Connors B. M., L. Coggins, B. Staton, C. Walters, M. Jones, M. Catalano and J. Harding. 2018. Incorporating harvest-population diversity tradeoffs into salmon management in large river basins: insights from Chinook in the Kuskokwim and Yukon River Basins. Institute of Ocean Sciences.
- Staton B., M. Catalano, B.M. Connors, L. Coggins, M. Jones, C. Walters, S. Fleischman and D. Gwinn 2017. Problems and solutions in the assessment of mixed-stock salmon fisheries. University of British Columbia.
- Connors B. M., L. Coggins, B. Staton, C. Walters, and M. Jones. 2016. Incorporating harvest-population diversity trade-offs into salmon management. Canadian Conference for Fisheries Research.
- Staton B., M. Catalano, B.M. Connors, L. Coggins, M. Jones, C. Walters, S. Fleischman and D. Gwinn 2016. Estimation of stock-specific productivity to assess trade-offs in mixed-stock Pacific salmon fisheries. Alabama-Georgia joint American Fisheries Society Meeting.

Draft manuscripts:

Staton B., M. Catalano, B.M. Connors, L. Coggins, M. Jones, C. Walters, S. Fleischman and D. Gwinn. In preparation. Evaluation of methods for quantifying population diversity in mixed-stock Pacific salmon fisheries.

Connors B. M., B. Staton, L. Coggins, C. Walters, M. Jones, D. Gwinn, M. Catalano and S. Fleischman. In preparation. Incorporating harvest-population diversity trade-offs into salmon management in large river basins: a case study of Kuskokwim River Chinook.

Final manuscripts will be provided to AYK-SSI once published.

Online visualizations:

An interactive online visualization of the results of the harvest policy analysis detailed in Appendix B can be found at:

https://brendanmichaelconnors.shinyapps.io/kuskokwimShinyApp/

These visualizations will continue to be developed until the manuscript they are associated with is published.

Meetings:

2015. National Fish and Wildlife Federation Capacity Building Workshop no. 1. Aniak, Alaska
2016. National Fish and Wildlife Federation Capacity Building Workshop no. 2. Anchorage/Bethel, Alaska
2016. National Fish and Wildlife Federation Capacity Building Workshop no. 3. Bethel, Alaska
2017. National Fish and Wildlife Federation Capacity Building Workshop no. 4. Bethel, Alaska
2017. Salmon Harvest Trade-offs workshop. Anchorage, Alaska

Reports:

Semiannual progress reports July 2015, January and July 2016, 2017, and 2018.

V. PROJECT DATA SUMMARY:

The raw data used in the run-reconstructions and spawner-recruitment analyses described in Appendix A were sourced from the Alaska Department of Fish and Game. These data and source code for the analyses performed in Appendix A can be found in the html supplement provided with this report (Appendix_A_online_supplement_S2.4.html). Source code for the closed-loop simulations performed in Appendix B, can be found at: <u>https://github.com/brendanmichaelconnors/Kusko-harvest-diversity-tradeoffs</u>

Note that the source code for the manuscripts in Appendices A and B will continue to be refined until the manuscripts are published.

VI. APPENDICES:

Appendix A: Evaluation of Methods for Quantifying Population Diversity in Mixed-stock Pacific Salmon Fisheries

Appendix B: Incorporating harvest-population diversity trade-offs into salmon management in large river basins: a case study of Kuskokwim River Chinook

Evaluation of Methods for Quantifying Population Diversity in Mixed-stock Pacific Salmon Fisheries

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Keywords: mixed-stock fisheries, Pacific salmon, state-space models, Bayesian inference

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¹ Abstract

Populations harvested in mixed-stock Pacific salmon fisheries show diversity in genotypic, 2 behavioral, and life history characteristics which can lead to heterogeneity in productivity 3 and population size. Methods to quantify this heterogeneity within mixed-stocks are not 4 well-established but are required for full acknowledgement of trade-offs when setting harvest 5 policies. We developed an integrated age-structured state-space model that allows for more 6 complete use of available data and sharing of information than simpler methods. We developed 7 a suite of state-space models of varying structural complexity for comparison to simpler 8 regression-based approaches and fitted them to data from 13 Chinook Salmon populations in 9 the Kuskokwim drainage in western Alaska. We found biological and policy conclusions were 10 largely consistent between state-space models but differed strongly from regression-based 11 approaches. Simulation trials illustrated our state-space models were largely unbiased with 12 respect to leading parameters, abundance states, and derived biological reference points, 13 whereas the regression-based approaches showed substantial bias. These findings suggest our 14 state-space model shows promise for informing harvest policy evaluations in the context of 15 harvest-biodiversity trade-offs in mixed-stock salmon fisheries. 16

17 **1** Introduction

Many Pacific salmon *Oncorhynchus* spp. populations in large drainage systems are commonly 18 harvested in a relatively small spatial area and are managed as a single stock. However, 19 these "stocks" are instead stock-complexes, in which the aggregate stock is composed of 20 several (and sometimes, many) substocks, hereafter referred to as "populations." These 21 populations can vary in genotypic (Templin et al. 2014), phenotypic (e.g., morphology; 22 Hendry and Quinn 1997), behavioral (e.g., run timing; Clark et al. 2015; Smith and Liller 23 2017a,b), and life history (e.g., age-at-maturation, Blair et al. 1993) characteristics that are 24 the result of adaptations to local environments after many generations of high spawning-site 25 fidelity and reproductive isolation from conspecifics in other tributaries located within the 26 same basin. It has been widely proposed that maintaining this diversity of local adaptation 27 (hereafter, "biodiversity") is favorable both from ecosystem and exploitation perspectives. 28 One argument is that in a system where many parts contribute to the whole, the variability 29 in the aggregate characteristics can be dampened due to asynchrony in the subcomponent 30 dynamics, a phenomenon commonly referred to as the "portfolio effect" (Schindler et al. 2010, 31 2015). 32

Diversity in these population characteristics can ultimately lead to heterogeneity in 33 their productivity (Walters and Martell 2004). Productivity in this context (hereafter, α) 34 is the ability of a population to replace itself after harvesting or some other perturbation, 35 often represented for salmon populations as the maximum number of future returning adults 36 (recruits) produced by one spawner, which is attained at low spawner abundances due 37 to density-dependent survival. Populations j with higher α_i values can sustain greater 38 exploitation rates (U) than those with smaller α_j values; α_j can be expressed in terms of 39 the exploitation rate that maximizes sustained yield from population j ($U_{MSY,j}$; Schnute and 40 Kronlund 2002): 41

(1)
$$\alpha_j = \frac{e^{U_{\text{MSY},j}}}{1 - U_{\text{MSY},j}}.$$

Given that there is likely some level of heterogeneity in α_j and $U_{\text{MSY},j}$ among individual populations, the logical conclusion is that in a mixed-stock fishery where the exploitation rate in year t is common among all populations, some weaker populations must be exploited at $U_t > U_{\text{MSY},j}$ in order to fish the more productive populations at $U_{\text{MSY},j}$. This of course implies a trade-off, such that it may be necessary to over-exploit some populations in order to maximize harvest benefits from others.

Before these trade-offs can be considered by managers in a well-informed way, the 48 shape and magnitude of the trade-off must first be quantified. These trade-offs are typically 49 expressed as the amount of mixed-stock harvest that must be foregone to reduce the number 50 or fraction of populations at risk of overfishing or extirpation (Walters and Martell 2004; 51 Walters et al. 2018). However, to conduct these assessments, the estimated productivity 52 and carrying capacity of all (or a representative sample) of the populations contributing 53 harvest to a mixed-stock fishery. Using a set of assumptions about the underlying population 54 dynamics, these quantities are obtained via spawner-recruit analysis, which involves tracking 55 the number of recruits produced in each brood year by the number of fish that spawned in 56 that same year and fitting a curve to the resulting pattern. Other methods exist for obtaining 57 estimates of population-level parameters, such as habitat-based methods (e.g., Lierman et al. 58 2010; Parken et al. 2006), but spawner-recruit analyses provide other information such as 59 estimates of variability in recruitment anomalies and maturation schedules, which may be 60 useful for forecasting (e.g., Murphy et al. 2017) or parameterizing operating models for use 61 in closed-loop policy evaluations (e.g., Cunningham et al. 2018a; Catalano and Jones 2014; 62 Connors et al. nd). 63

64

The spawner-recruit literature is extensive, but focuses primarily on assessing popu-

lations or stocks as a whole, as opposed to population components (but see the work on 65 Skeena River, British Columbia sockeye salmon O. nerka populations; Korman and English 66 2013; Walters et al. 2008). Population-specific analyses are uncommon because of two fac-67 tors: (1) the data to conduct well-informed population-specific spawner-recruit analyses 68 are often unavailable (20 - 30 years of continuous spawner and harvest counts/estimates 69 and age composition for each population) and (2) management actions in large mixed-stock 70 fisheries may not be dexterous enough to deliberately exert higher exploitation rates on more 71 productive populations even if the nature of the trade-off was known perfectly. Regarding 72 the former reason, there are some cases where the data do exist to perform these kinds 73 of analyses (e.g., the Kuskokwim River Chinook salmon O. tshawytscha stock in western 74 Alaska is used here as case study), however methods to conduct mixed-stock spawner-recruit 75 analyses are not well-developed. Regarding the latter, even in cases where management 76 cannot target particular populations over others, understanding the nature of the trade-offs 77 can be informative for evaluating candidate harvest policies for the mixed-stock in the context 78 of population biodiversity (Walters et al. 2018). 79

Methods to fit spawner-recruit models can be grouped into two broad categories: 80 regression-based approaches (e.g., Clark et al. 2009) and state-space (i.e., time series) models 81 (e.g., Fleischman et al. 2013; Su and Peterman 2012). The regression-based approaches 82 treat spawner-recruit pairs as independent observations, and are thus subject to the pitfalls 83 of dealing with the inherent time-dependent properties and oftentimes large amounts of 84 observation error found in spawner-recruit data sets (Walters and Martell 2004, Ch. 7). 85 The consequence of ignoring the first issue is "time-series bias", which causes positive biases 86 in α and negative biases in β , resulting in the same directional biases in U_{MSY} and S_{MSY} , 87 respectively (i.e., spuriously providing too aggressive harvest policy recommendations; Walters 88 1985). The second is known as "errors-in-variables bias" and is known to cause an apparent 89 scatter which inserts additional variability that commonly used regression estimators do not 90 account for and can also lead to positive biases in α (Ludwig and Walters 1981). Though 91

these methods have been known for their problems for over 30 years, they are still somewhat 92 widely used (e.g., Clark et al. 2009; Korman and English 2013). Unlike the regression-based 93 approaches, the state-space class of models attempts to capture the process of recruitment 94 events leading to future spawners while simultaneously accounting for variability in the 95 biological and measurement processes that gave rise to the observed data (de Valpine and 96 Hastings 2002: Fleischman et al. 2013). For these reasons, state-space spawner-recruit 97 analyses have rapidly gained popularity, particularly in Alaska (Fleischman et al. 2013; 98 Staton et al. 2017; Su and Peterman 2012). This level of additional model complexity comes 99 at computational costs, as these models are well-suited for Bayesian inference with Markov 100 Chain Monte Carlo (MCMC) methods (Newman et al. 2014, Ch. 4), but have been shown to 101 reduce bias in estimates in some circumstances relative to regression-based approaches (Su 102 and Peterman 2012; Walters and Martell 2004). 103

In cases where data are available to perform mixed-stock spawner-recruit analyses on a 104 population-specific basis, it is difficult to know which assessment method is most appropriate. 105 Two prevalent issues arise which may benefit from the development of a mixed-stock state-106 space spawner-recruit model that simultaneously estimates population dynamics parameters, 107 recruitment states, and biological reference points on a single population and aggregate 108 mixed-stock basis. First, reconstructing population-specific recruitment time series is difficult 109 without genetic stock identification (Cunningham et al. 2018b; Michielsens and Cave 2018; 110 Beacham et al. 2004) of the populations contributing to in the mixed-stock harvest, because 111 they are otherwise assessed primarily in aggregate. This situation requires assumptions 112 about the relative exploitation rates experienced by each population, and an approach 113 that integrates the population-specific dynamics into one model can provide flexibility and 114 transparency in how these simplifications are made. Second, available population-specific 115 data are generally escapement time series, whether they are cenuses, indices, or estimates. In 116 cases where reasonably long time series (20 + years) are available for a significant portion 117 (>50%) of the populations in a mixed-stock, it is common to have non-contiguous sampling 118

data. An additional benefit of integrating the analysis in a time series framework may be the ability to share information between populations by exploiting patterns in their recruitment covariance. Inappropriate treatment of either factor may render the final estimates of population parameters inappropriate for management use, thus the sensitivity and performance of a variety of approaches should be assessed, including an integrated state-space model.

In this article, we evaluate the performance of a range of assessment models for mixed-stock salmon fisheries via simulation-estimation and apply them to Chinook salmon populations in the Kuskokwim River in western Alaska as a case study. Our objectives were to:

- (1) develop a set of varyingly complex mixed-stock versions of state-space spawner-recruit
 models,
- (2) determine the sensitivity of biological and trade-off conclusions to assessment model
 complexity (including those obtained using regression-based approaches) using empirical
 data from Kuskokwim River Chinook salmon populations, and

(3) test the performance of the assessment models via simulation-estimation trials.

$_{135}$ 2 Methods

We conducted our analysis in two parts to evaluate the sensitivity and performance of assessment strategies for the mixed-stock Pacific salmon fisheries. First, all assessment methods (two regression-based and four state-space models) were fitted to observed data from the Kuskokwim River populations ($n_j = 13$) to determine the extent to which the choice of assessment model structure influences biological and management conclusions with real data. Next, a hypothetical salmon system composed of several age-structured populations and with known properties, subjected to observation with error and inconsistent sampling frequencies, and then all models were fitted to each simulated data set. Lastly, we quantified model performance based on the bias and precision of quantities with relevance to mixed harvest management (e.g., $U_{\rm MSY}$ and $S_{\rm MSY}$).

¹⁴⁶ 2.1 Mixed-stock spawner-recruit models

147 2.1.1 Regression-based models

Two regression-based approaches for estimating Ricker (1954) spawner-recruit parameters on a population-specific basis were assessed: (a) a single mixed-effect regression model with random intercepts (also referred to as a hierarchical model) and (b) independent regression models. The Ricker spawner-recruit model can be written as:

(2)
$$R_y = \alpha S_y e^{-\beta S_y + \varepsilon_y}$$

where R_y is the total recruitment produced by escapement S_y in brood year y, α is the 152 maximum expected recruits per spawner (RPS), β is the inverse of the escapement that is 153 expected to produce maximum recruitment (S_{MAX}) , and ε_y are mean zero deviations from 154 the expected curve often assumed to be attributed to environmentally driven fluctuations in 155 juvenile survival. Primary interest lies in estimating the population dynamics parameters α 156 and β as they can be used to obtain biological reference points from which sustainable harvest 157 policies can be developed. The Ricker function in (2) is increasing at small escapements and 158 declining at large ones, and it can be linearized: 159

(3)
$$\log_e(\operatorname{RPS}_y) = \log_e(\alpha) - \beta S_y + \varepsilon_y,$$

allowing for estimation of the parameters $\log_e(\alpha)$ and β in a linear regression framework using 160 the least squares method or likelihood methods under the assumption that $\varepsilon_y \sim N(0, \sigma_R^2)$ 161 (Clark et al. 2009; Hilborn and Walters 1992). This relationship is nearly always declining, 162 implying a compensatory effect on juvenile survival (i.e., RPS) with reductions in spawner 163 abundance (Rose et al. 2001). Obtaining the R_y component of RPS_y can be problematic for 164 salmon populations that mature at multiple ages (like Chinook salmon, for which we have 165 incorporated ages four to seven) given the prevalence of non-consecutive sampling years of 166 either age composition or escapement data. For fitting regression models, only brood year 167 recruitment events were used in which all contributing ages were observed. For example, 168 to include brood year recruitment for 2010 in the model, escapement data must have been 169 available for calendar years 2014 - 2017, as well as in 2010 to complete the pair. 170

¹⁷¹ A mixed-stock formulation of this model can be expressed by including population-¹⁷² specific random effects on the intercept $[\log_e(\alpha)]$:

(4)

$$\log_e(\operatorname{RPS}_{y,j}) = \log_e(\alpha_j) - \beta_j S_{y,j} + \varepsilon_y,$$

$$\log_e(\alpha_j) = \log_e(\alpha) + \varepsilon_{\alpha,j},$$

$$\varepsilon_{\alpha,j} \sim \operatorname{N}(0, \sigma_{\alpha}^2).$$

It would be nonsensical to include population-level random effects on the slope, given that β is a capacity parameter related to the compensatory effect of habitat and resource limitation experienced by juveniles, likely in the freshwater environment (i.e., amount of habitat as opposed to quality of habitat). Fitting the individual population models in this hierarchical fashion allows for the sharing of information such that the more intensively assessed populations can help inform those that are more data-poor.

The mixed-effect model may have the benefit of sharing information to make some populations more estimable, but it should also have the tendency to pull the extreme α_i

(those in the tails of the hyperdistribution) toward α . This behavior may not be preferable for 181 policy recommendations, as it should tend to dampen the extent of heterogeneity estimated in 182 α_i . For this reason, independent regression estimates for each population were also obtained 183 for evaluation. In estimating the parameter $\log_e(\alpha)$, a lower bound constraint of zero was 184 used in all regression models. This was necessary to prevent the models from estimating 185 biologically implausible parameters: if $\log_e(\alpha) < 0$, then no amount of spawners would be 186 expected to replace themselves (let alone provide surplus) in their most productive state, in 187 which case the population would not likely be in existence. 188

189 2.1.2 State-space models

We developed four versions of the state-space formulation. As three versions were simplifications of a more full model, the full model is presented completely here and the changes resulting in the other three model structures are described following the description of the full model.

The state-space formulation of the mixed-stock spawner-recruit analysis developed and 194 evaluated here is an extension of various single-stock versions (e.g., Fleischman et al. 2013). 195 Walters et al. (2008) used a similar model using maximum likelihood methods to provide 196 estimates of >50 populations in the Skeena River drainage, British Columbia. The model 197 presented here was fitted with Bayesian methods using program JAGS (Plummer 2017). 198 and allows for relaxation of certain assumptions made by Walters et al. (2008) such as the 199 important notion of all populations having the same recruitment residual time series (i.e., 200 anomalies – deviations from the expected population response). 201

The state-space model is partitioned into two submodels: (a) the process submodel which generates the latent (i.e., true but unobserved) states of $R_{y,j}$ and the resulting calendar year states (e.g., $S_{t,j}$) and (b) the observation submodel which fits the latent states to the observed data (notation summarized in Tables 1, 2). Note that this method does not require excluding brood year recruitment events that were not fully observed as was necessary for
the regression-based models.

The recruitment process operated by producing a mean prediction from the deterministic 208 portion of the Ricker relationship in (2) for n_y brood years for each of the n_j populations. 209 From these deterministic predictions, auto-correlated process variability was added to generate 210 the realized latent recruitment states. To populate the first n_a calendar year abundance 211 states with recruits of each age a, the first a_{max} brood year expected recruitment states 212 were not linked to a spawner abundance through (2) (because the S_y component was not 213 observed), but instead were assumed to have a constant mean equal to the unfished equilibrium 214 recruitment (where non-zero S_j produces $R_j = S_j$ when unexploited and in the absence of 215 process variability): 216

(5)
$$\dot{R}_{y,j} = \frac{\log_e(\alpha_j)}{\beta_j},$$

where $\dot{R}_{y,j}$ is the expected (i.e., deterministic) recruitment in brood year y from population j with Ricker parameters α_j and β_j . The remaining $n_y - a_{max}$ brood years had an explicit time linkage:

(6)
$$\dot{R}_{y,j} = \alpha_j S_{t,j} e^{-\beta_j S_{t,j}}$$

where $t = y - a_{max}$ is the t^{th} calendar year index in which the escapement produced the recruits in the y^{th} brood year index.

From these deterministic predictions of the biological recruitment process, lag-1 autocorrelated process errors were added to produce the latent states:

(7)
$$\log_e(R_{y,1:n_j}) \sim \text{MVN}\left(\log_e(\dot{R}_{y,1:n_j}) + \omega_{y,1:n_j}, \Sigma_R\right),$$

²²⁴ where

(8)
$$\omega_{y,1:n_j} = \phi \left(\log_e(R_{y-1,1:n_j}) - \log_e(\dot{R}_{y-1,1:n_j}) \right),$$

and $R_{y,1:n_j}$ is the vector of latent recruitment states across the n_j stocks in brood year y, 225 $\omega_{y,1:n_j}$ is the portion of the total process error attributable to serial auto-correlation, ϕ is 226 the lag-1 auto-correlation coefficient (constant across populations), and Σ_R is a covariance 227 matrix representing the white noise portion of the total recruitment process variance. In the 228 full model, Σ_R was estimated such that each population was assigned a unique variance and 229 covariance with each other population. This was achieved by using an inverse Wishart prior 230 distribution, with degrees of freedom equal to $n_j + 1$ and the scale matrix populated with 231 zero-value elements along the off-diagonals and each diagonal element equal to one, which 232 inserts little information about the covariance matrix Σ_R (Plummer 2017). The multivariate 233 normal errors were on the logarithmic scale so the variability on $R_{y,j}$ was lognormal, which 234 is the most commonly used distribution for describing recruitment variability (Walters and 235 Martell 2004). Further, the multivariate normal was used as opposed to n_j separate normal 236 distributions so the degree of synchrony in brood year recruitment deviations (i.e., recruitment 237 process errors) among populations could be captured and freely estimated in Σ_R . 238

The maturity schedule is an important component of age-structured spawner-recruit models, as it determines which calendar years the brood year recruits $R_{y,j}$ return to spawn (and be observed). Recent state-space spawner-recruit analyses have accounted for brood year variability in maturity schedules as Dirichlet random vectors drawn from a common hyperdistribution characterized by a mean maturation-at-age probability vector $(\pi_{1:n_a})$ and an inverse dispersion parameter (D) (see Fleischman et al. 2013; Staton et al. 2017, for implementation in JAGS), and the same approach was used for the full model with maturity schedules shared perfectly among populations within a brood year. Brood year-specific maturity schedules were treated as random variables such that:

(9)
$$p_{y,a} \stackrel{\text{iid}}{\sim} \text{Dirichlet}(\pi_{1:n_a} \cdot D).$$

where $p_{y,a}$ is the probability a fish spawned in brood year y will mature and make the spawning migration at age a.

In order to link $R_{y,j}$ with calendar year observations of escapement from each population, $R_{y,j}$ was allocated to calendar year runs-at-age $(N_{t,a,j})$ based on the maturity schedule:

(10)
$$N_{t,a,j} = R_{t+n_a-a,j} p_{t+n_a-a,a},$$

and the total run returning to population j in year t was the sum of mature fish within a calendar year across ages:

(11)
$$N_{t,j} = \sum_{a=1}^{n_a} N_{t,a,j}.$$

The harvest process was modeled using a freely estimated annual exploitation rate (U_t) time series, which was assumed to apply equally to all populations (but see Section 2.2.3 for a relaxation of this assumption):

(12)
$$H_{t,j} = N_{t,j}U_t,$$

²⁵⁷ and escapement was obtained as:

(13)
$$S_{t,j} = N_{t,j}(1 - U_t).$$

The quantity H_t (mixed-stock harvest) was obtained by summing $H_{t,j}$ within a t index across the j indices. The true age composition returning in year t to population j ($q_{t,a,j}$) was obtained as:

(14)
$$q_{t,a,j} = \frac{N_{t,a,j}}{N_{t,j}}.$$

Three data sources were used to fit the state-space model: (a) observed escapement from each population $(S_{obs,t,j})$ with assumed known coefficients of variation (CV), (b) total harvest arising from the aggregate stock $(H_{obs,t})$ with assumed known CV, and (c) the age composition of populations with these data each calendar year $(q_{obs,t,a,j};$ which had associated effective sample size $ESS_{t,j}$ equal to the number of fish successfully aged for population j in year t). The CVs were converted to lognormal variances:

(15)
$$\sigma_{\log}^2 = \log_e(\mathrm{CV}^2 + 1),$$

and used in lognormal likelihoods to fit the time series $S_{t,j}$ to $S_{obs,t,j}$ and H_t to $H_{obs,t}$. Calendar

year age composition was fitted using independent multinomial likelihoods with parameter vectors $q_{t,1:n_a,j}$ and observed vectors of $(q_{obs,t,1:n_a,j} \cdot ESS_{t,j})$.

Three alternative formulations of the state-space model were evaluated, and all were 270 simplifications of the full model described above regarding the structure of (a) the covariance 271 matrix on recruitment residuals (Σ_R) and (b) the maturity process (see Table 3 for a summary 272 of the models fitted in this analysis). The simplest state-space model did not include brood 273 year variability in maturity schedules and Σ_R was constructed by estimating a single σ_R^2 and ρ 274 (common across all populations and population pairs) and populating the diagonal elements 275 with σ_R^2 and off-diagonal elements with $\rho \sigma_R^2$. This simplest model is denoted as SSM-vm (lower 276 case letters indicate simple versions; v = recruitment covariance; m = maturity variability). 277 In one intermediate model (SSM-vM), brood year maturation variability was included but Σ_R 278 was constructed as in the simplest model. In the other intermediate model (SSM-Vm), brood 279 year variability in maturation was not included but Σ_R was fully estimated as in the full 280 model (SSM-VM). These two structural uncertainties (complexity in recruitment covariance 281 and maturity variability) were chosen for evaluation here because they are two key areas 282 where an analyst might question if the available data are adequate for model fitting and 283 inference. In other words, these are two key model components where it may be important 284 to know if the complex versions are reliably estimable with a reasonable amount of data. 285

286 2.2 Kuskokwim empirical analysis

287 2.2.1 Study system

All six assessment models (two regression-based and four state-space models) evaluated were fitted to empirical data from Chinook salmon populations of the Kuskokwim River located in western Alaska (Figure 2). The Kuskokwim River is one of the largest subsistence salmon fisheries in the state, and the largest for Chinook salmon (Fall et al. 2018). Commercial harvests have been historically important to the region, but have not been directed toward

Chinook salmon since 1984. The fishery can very well be described as operating on a mixed-293 stock, both for multiple salmon species (predominately Chinook, chum O. keta, and sockeye 294 salmon) and for multiple populations of the same species. Rigorous monitoring of harvest, 295 escapement, and age composition for Chinook salmon began in the mid-1970s with efforts 296 expanding significantly in the early-1990s and 2000s (Figure 1). Approximately 90% of the 297 subsistence salmon harvest in the drainage occurs in the main-stem Kuskokwim River using 298 primarily drift gillnet gear, downstream of the confluence with the Aniak River (Hamazaki 299 2011). Migrating adult fish originating from and returning to the various tributaries (of 300 which there are at least 20 that support salmon; Figure 2) enter through this bulk of the 301 fishery as a mixed-stock: first Chinook in early June, then chum and sockeye later in June 302 and July but with substantial overlap in entry timing. Chinook salmon stocks migrating to 303 the headwaters have been illustrated to enter the main-stem earliest in the summer migration 304 (Smith and Liller 2017a,b) so a limited ability to direct harvest toward or away from these 305 populations may exist by manipulating the front portion of the fishery. It is acknowledged 306 that the assessment program does not sample all tributaries within the Kuskokwim River 307 where Chinook salmon spawn (Figure 2), but total run size between 1976 and 2017 has been 308 estimated via run reconstruction (Liller et al. 2018) and large-scale mark-recapture studies 309 (Schaberg et al. 2012; Smith and Liller 2017a,b; Stuby 2007). 310

311 2.2.2 Data sources

The data set used included counts of Chinook salmon at many locations throughout the Kuskokwim River drainage (Figure 2). Nearly all data were collected by projects managed by the Alaska Department of Fish and Game (ADF&G) and a complete description of data needs and preparation procedures is provided in Online Supplement S1. The raw escapement data set spanned 20 different escapement monitoring projects (six weirs and 14 aerial surveys) and 42 calendar years from 1976 – 2017; see Head and Smith (2018) for details on tributary escapement monitoring. Some pre-processing was required to convert the aerial survey index counts to estimates of total escapement so that abundance-based reference points could be calculated. Annual estimates of Chinook salmon harvest originating from both subsistence and commercial fisheries in each year was also available, as were the estimated exploitation rates of the aggregate stock. Finally, age composition data were available for the six populations monitored via weir programs (complete details on data processing are provided in Online Supplement S1).

325 2.2.3 Sensitivity Analyses

Two analyses were conducted to test the sensitivity of inference from state-space models fitted 326 to Kuskokwim data. First, the default assumption that all populations have been fished at 327 the same rate each year is tenuous. We therefore included a term in the model that allowed 328 for populations to have differential vulnerability to harvest (v_j) by replacing U_t with $U_t v_j$ 329 in (12) and (13) which allowed acknowledgement of heterogeneous exploitation rates. This 330 alteration changes the interpretation of the parameter vector U_t to be the exploitation rate 331 of fully vulnerable populations. Without additional information on what portion of $H_{obs,t}$ 332 was attributable to each population going back in time, the v_j elements are not estimable. 333 In the absence of this information for Kuskokwim River Chinook salmon, v_i was assigned 334 by calculating the fraction of the fishing households residing along the main-stem of the 335 river that each population must travel past in order to reach their natal spawning grounds. 336 Fishing household data were available from post-season interviews conducted by ADF&G 337 (e.g., Hamazaki 2011; Shelden et al. 2016). Although this method ignores the temporal 338 overlap of the fishery (Hamazaki 2008) with the arrival timing of particular population groups 339 (Smith and Liller 2017a,b), it was intended as a first step at determining how much the 340 conclusions might depend on how the internal harvest accounting was specified. No attempt 341 was made to alter how harvest was apportioned for use in the regression-based models. 342

As a secondary sensitivity analysis, the information content of the age composition data was reduced. In the default case, each annual multinomial age composition vector had sample size equal to the number of fish successfully aged for that population/year combination. For some populations/years, this number was quite high from a multinomial sampling perspective when the number of categories is small (e.g., $\approx 1,200$ samples across four age categories). To assess whether this strength of information had an impact on the inference, the effective sample size was manipulated such that the maximum number of fish sampled for a population was assigned $ESS_{t,j} = 100$, and the other years with data were scaled proportionately.

351 2.2.4 Comparisons of model output

Key population dynamics parameters and biological reference points were compared among the six assessment models wherever possible to determine the extent to which the management conclusions for Kuskokwim River Chinook salmon populations might change based on the model structure used. Where appropriate, quantities were averaged across populations to facilitate comparisons and included indicators of the average population's productivity $(\bar{U}_{MSY,j};$ the "bar" denotes average across populations) and size $(\bar{S}_{eq,j} \text{ and } \bar{S}_{MSY,j})$. Reference points for the aggregate mixed-stock included U_{MSY} and S_{MSY} .

Model fits to the data $(S_{obs,t,s}, H_{obs,t}, and q_{obs,t,a,j})$ were examined for all state-space 359 models and noteworthy differences among model structures were identified. Estimates of 360 synchrony in recruitment anomalies were examined, both between two average populations 361 $(\bar{\rho}_{i,j})$, and among all population pairs. For the simple recruitment variance models (SSM-362 vm and SSM-vM), correlations between each population pair were conducted by applying 363 Pearson's r coefficient to two populations' estimated recruitment anomaly time series; in the 364 complex variance models these correlations were captured in the freely estimated covariance 365 matrix (Σ_R) , and they were extracted and summarized. Although the state-space models 366 were ignorant of spatial relationships among the populations, visual comparisons were made 367 to determine if populations closer in proximity showed higher synchrony than those spaced 368 more distantly, as might be expected. The auto-correlation parameter (ϕ) and characteristics 369 of the maturity schedules (π_a and D) were also compared among state-space models. 370

Harvest-biodiversity trade-offs were assessed visually using estimated equilibrium mixed-371 stock states and metrics of biodiversity. First, the equilibrium escapement and harvest were 372 calculated (Schnute and Kronlund 2002) for each population at each level of an exploitation 373 rate that affected all populations equally, then summed to obtain the mixed-stock expected 374 states at each level. The fraction of populations overfished and trending towards extirpation 375 (defined as the case where equilibrium $S_j \leq 0$) were also calculated. Some populations could 376 not be fitted using the regression approach because they had fewer than three observed pairs 377 of $RPS_{y,j}$. When comparing quantities to the estimates from the state-space models, the 378 populations that could not be fitted with regression approaches were removed from state-space 370 model outputs. 380

381 2.3 Simulation-estimation trials

To test the performance of these models, we simulated 160 data sets designed to mimic the Kuskokwim River empirical data set. Each of the 160 data sets were passed to each of the six assessment models to evaluate which methods return estimates closest to the true parameters.

Given that the state-space model is a much more natural model of this system (which 385 has intrinsic time series properties) than the regression-based versions, it was used as the 386 foundation of the operating model (i.e., state-generating model). The biological submodel 387 was more complex than the most complex estimation model – namely with regards to the 388 maturity schedule, which had a modest level of population variability in mean maturity 389 but with highly correlated brood year variability. In order to serve as the state-generating 390 model for the simulation, the state-space model needed only to be populated with true 391 parameters, initial states, and a harvest control rule. A fixed exploitation rate policy was 392 used (chosen to maximize yield without overfishing more than 30% of the populations) with 393 a modest amount of implementation error to ensure the escapement data time series were 394 generated with contrast. $n_j = 13$ populations were simulated with different parameters $U_{MSY,j}$ 395 and $S_{MSY,j}$ which took on the values of random posterior draws from the most complex 396

state-space model fitted to the Kuskokwim River Chinook salmon population data. All other parameters were chosen to mimic the estimated values from the Kuskokwim analysis, with the exception of Σ_R , which was set to have a modest amount of population recruitment variability $(\bar{\sigma}_{j,j} \approx 0.4); \rho_{i,j}$ for each pair of populations was simulated randomly to be between -1 and 1, but approximately zero when averaged across all populations.

For a given set of simulated true states, a set of observed states $(S_{obs,t,j}, H_{obs,t}, q_{obs,t,a,j})$ 402 was generated by adding sampling error to each year following a fifty year "burn-in" period 403 of the harvest policy into the population dynamics. This was intentionally done to violate 404 the assumption made by the state-space estimation model that sampling was initiated under 405 unfished conditions, as shown in (5). Sampling errors in escapement and harvest estimates 406 were generated with lognormal variability and multinomial sampling for the age composition, 407 as assumed in the state-space estimation model. Frequency of sampling on each population was 408 set to approximately mimic the Kuskokwim River historical monitoring program (Figure 1). 409 The sampling frequency was designed to continue to generate random sampling schedules until 410 one was found that ensured no population had fewer than three observations of $RPS_{y,j}$ which 411 allowed the regression-based models to be fitted to all populations. Aggregate harvest data 412 $(H_{obs,t})$ were assumed to be available every year as were annual estimates of the mixed-stock 413 exploitation rate. 414

Estimation performance in terms of accuracy among assessment models was calculated 415 using the proportional error $\left(\frac{x_{\text{est}}-x_{\text{true}}}{x_{\text{true}}}\right)$. Key quantities of interest for comparison between 416 the regression approaches and the state-space models included: S_{MSY} , U_{MSY} , α_j , $U_{MSY,j}$, and 417 $S_{MSY,j}$, as well as two metrics that incorporate population biodiversity considerations: S_p^* 418 and U_p^* . These quantities represent the equilibrium mixed-stock escapement or exploitation 419 rate, respectively, that would result in no more than $p \cdot 100\%$ of populations expected to 420 be overfished. "Overfished" is defined here as the case where a population would be fished 421 at $U > U_{MSY,j}$. Three levels of p were extracted: 0.1, 0.3, and 0.5. For the state-space 422 models, the ability to accurately estimate the abundance states of $R_{y,j}$, $S_{t,j}$, and H_t was also 423

⁴²⁴ assessed, summarized for early years and all years separately to investigate the influence of ⁴²⁵ the assumption that the observation time series began at unfished equilibrium, as described ⁴²⁶ by (5). Coverage of the 95% equal-tailed credible interval was assessed for these quantities by ⁴²⁷ determining the fraction of obtained credible intervals that captured true values. Additionally, ⁴²⁸ model run times and convergence diagnostics were summarized for all models that were ⁴²⁹ successfully fitted to the simulated data sets.

430 2.4 Computation

All parameter estimation was conducted in the Bayesian framework using the MCMC engine JAGS (Plummer 2017) invoked through R (R Core Team 2018) using the package "jagsUI" (Kellner 2017). All priors for the regression-based methods were uniform and sufficiently non-informative to exclude only highly implausible values (e.g., $\log_e(\alpha_j) \sim U(0, 5)$). Priors were selected with the same intent for state-space models, though other distributional forms were required in some cases (Table 4).

MCMC sampling was conducted using sufficiently long chains (Table 5) to ensure 437 adequate sampling of the posterior parameter space for inference and was assessed using 438 visual inspection of MCMC sampling behavior and the convergence diagnostic proposed 439 by Brooks and Gelman (1998). Adequate sampling was further verified for key estimated 440 quantities using the effective sample size and the Raftery-Lewis diagnostic (Raftery and Lewis 441 1992). All posterior distributions were summarized using the median and 95% equal-tailed 442 credible intervals. In all cases where a quantity was derived from estimated parameters, (e.g., 443 S_p^*), the calculation was conducted for each sample from the joint posterior, then the resulting 444 marginal posterior was summarized, which proved to be a straightforward approach to carry 445 the uncertainty forward from the model estimates to other quantities of interest. All code 446 for simulation, JAGS models, data processing, making use of High Performance Computing 447 resources, and output summarization is documented in Staton (YYYY; a Github repository 448 that stores code and data, will have a DOI associated with it). 449

450 **3** Results

451 3.1 Kuskokwim River empirical analysis

452 3.1.1 State-space models fit to data

The four state-space models produced generally similar latent escapement state time series. 453 especially in years with observed escapement data (Figure 3). For several populations, there 454 were no escapement data prior to the mid-1990s, and this is one area the various state-space 455 models produced different escapement estimates. In this early portion of the time series, 456 models with brood year variability in maturity (SSM-vM and SSM-VM) tended to estimate 457 higher escapement abundance than the models with time-constant maturity (SSM-vm and 458 SSM-Vm). For example, populations spawning in the Kwethluk, George, Holokuk, and 459 Takotna rivers all showed this pattern (Figure 3). 460

There were several cases where extremely high (and seemingly unrealistic) escapement 461 states were estimated by the state-space model, though these only occurred in models with 462 simple maturation schedules (Figure 3). The period in the late-1980s and the early 1990s had 463 much (i.e., 5-10 times) higher estimated escapement than ever observed for the Holitna, 464 Pitka, and Tatlawiksuk populations under either models SSM-vm or SSM-Vm. The George 465 River population had abnormally large escapements in the mid-1990s, and was again most 466 exaggerated for the SSM-vm and SSM-Vm versions. All of these cases occurred when no 467 escapement data were available; in years with data all state-space models fitted the escapement 468 data quite well (Figure 3). 469

In general, the fit to the aggregate harvest data was good, though the time-varying maturity schedule versions (SSM-vM and SSM-VM) fitted the data nearly perfectly. Constant maturity schedule models resulted in a harvest state that was greater than twice as large as the observed state in 1976 (the first observed year) but never showed discrepancies nearly this large for the rest of the time series. Additionally, the state-space models generally fitted the age composition data well. The only differences among model structures came in the distinction of how maturity was treated. Complex maturity models tended to have better fit than simple maturity models for most populations and ages. Far more detailed model output and model fits from each of the state-space models can be found in the Online Supplements S2 - S5.

480 3.1.2 Comparisons of estimated spawner-recruit dynamics

There were large discrepancies in estimates of population productivity between the regression 481 models and the state-space models. Both regression approaches suggested the maximum 482 productivity of the average population $(\bar{\alpha}_i)$ was far higher than any of the state-space models. 483 The independent regression approach provided an estimate of $\bar{\alpha}_j = 7.74 \ (4.47 - 20.22; 95\%)$ 484 credible limits) and the mixed-effect regression approach suggested $\bar{\alpha}_j = 4.63 \ (3.16 - 7.45).$ 485 Most state-space models suggested $\bar{\alpha}_j < 3$, with the highest upper 95% credible limit being 486 5.01, obtained by SSM-vM. These differences in estimated productivity translated directly to 487 the maximum sustainable exploitation rate for the average population ($\bar{U}_{MSY,j}$; Figure 4a) 488 and even more so for the mixed-stock (U_{MSY} ; Figure 4b). Among state-space models, those 480 that included complex maturity variability tended to suggest the populations were more 490 productive (Figure 4a,b). In comparing state-space model estimates from the 10 substocks 491 able to be fitted with all methods to the complete set of 13 substocks, the average maximum 492 sustainable exploitation rates were quite similar, indicating the stocks with insufficient data 493 for regression were missing at random in this regard (Figure 4a,b; compare squares and 494 circles). 495

One metric of substock size is the spawner abundance expected to exactly replace itself under unfished equilibrium conditions $(S_{eq,j})$. When averaged across substocks $(\bar{S}_{eq,j})$, the regression approaches suggested the Kuskokwim River substocks were approximately 2,500 fish (~25%) smaller than the state space models. $\bar{S}_{MSY,j}$ followed a similar pattern: smaller values for the regression approaches than the state-space models (Figure 4c). In

terms of the mixed-stock escapement expected to produce maximum sustained yield (S_{MSY}) , 501 regression approaches suggested much smaller escapements were necessary than did state-502 space models; by a margin of 12,000 - 15,000 fish ($\sim 25 - 50\%$; Figure 4d). State-space models 503 with time-constant maturity had much greater uncertainty in $\bar{S}_{MSY,j}$ and S_{MSY} than the 504 complex maturity models (Figure 4c,d). In comparing state-space model estimates from 505 the 10 substocks able to be fitted with all methods to the complete set of 13 substocks, the 506 average substock-specific escapement values were quite similar, indicating the stocks with 507 insufficient data for regression were missing at random in this regard (Figure 4c; compare 508 squares and circles). However, the mixed-stock S_{MSY} estimates were quite different between 500 the 10 and 13 substock outputs (Figure 4d; compare squares and circles). It seems the three 510 additional substocks (Holitna, Kisaralik, and Oskawalik) should make up approximately 25% 511 of the escapement among the substocks included in this analysis if the management objective 512 is to maximize long-term yield. 513

These similaries and differences in population size and productivity between approaches 514 are well-illustrated at the population-level by visualizing the expected recruitment at each 515 spawner abundance suggested by each model (Figure 5). It is evident that the four state-space 516 models behaved similarly near the origin (which is governed by α_j), whereas in many cases 517 the regression models suggested steeper slopes near the origin (corresponding to higher 518 values of α_i). State-space models tended to disagree in expected recruitment more at larger 519 spawner abundances (Figure 5), suggesting that inferences about stock size and the strength 520 of compensation are dependent on the details of how recruitment variance and maturity 521 are modeled for some substocks. The mixed-effect regression approach differed from the 522 independent regression estimates the most for substocks with fewer observations due to 523 shrinkage (e.g., Pitka versus Kogrukluk; Figure 5). 524

In terms of recruitment variability, the regression-based approaches suggested that the average standard deviation of the lognormal distribution that describes randomness in the recruitment process $(\bar{\sigma}_{R,j})$ was 0.78 and 0.52 for the independent and mixed-effects versions, respectively (Figure 4e). The state-space models estimated that the average amount of recruitment process variability by substock was higher: point estimates ranged from 0.85 – 1.11 (Figure 4e). The models with time-varying maturity had estimates on the lower end of this range, which could be explained by the inclusion of additional process variability in maturity that could describe variability in the data.

Unlike the regression models, the state-space models estimated the degree of covariance 533 between substock recruitment residuals. All state-space models estimated a moderate amount 534 of correlation in recruitment variance (i.e., synchrony) between two average substocks $(\bar{\rho}_{i,j})$: 535 point estimates ranged between 0.18 and 0.28, and none of the models suggested 95% credible 536 limits that encompassed zero. The simple variance models (SSM-vm and SSM-vM) estimated 537 a single correlation parameter, whereas the complex versions (SSM-Vm and SSM-VM) 538 estimated a unique value for each substock combination. One might expect that substock 539 pairs belonging in the same region to show higher degrees of synchrony than substock pairs 540 in different regions, though this analysis suggested this was not necessarily the case (the 541 state-space models did not take into account spatial relations among substocks). Large 542 correlations (e.g., >0.5) were found between pairs of lower versus upper river substocks, lower 543 versus lower, and upper versus upper substocks (Figure 6). Relatively few large correlations 544 were found among middle river substocks with other substocks, though both SSM-Vm and 545 SSM-VM suggested that the Holokuk and Oskawalik substocks have highly synchronous 546 recruitment dynamics, which is interesting given their close proximity (Figure 2; substocks #5547 and #6, respectively). Conversely, all models suggested the Holitna and Kogrukluk substocks 548 have little synchrony, and they fall within the same subdrainage (Figure 2; substocks #7549 and #8, respectively). Most correlations were positive, especially those that were large 550 in magnitude (Figure 6). A notable exception was the correlation between the Kisaralik 551 substock and the Holokuk and Oskawalik substocks: their dynamics were suggested to be 552 largely opposite. 553

554 3.1.3 Trade-off comparisons

With increasing mixed-stock exploitation rates, the equilibrium mixed-stock escapement 555 declined, but did so more rapidly for the state-space models than the regression models (Figure 556 7). State-space models that included time-varying maturity suggested higher equilibrium 557 escapement and harvest would be available at most exploitation rates than models with 558 time-constant maturity. Regression models suggested that MSY was much larger and occurred 559 at a higher exploitation rate than for the state-space models, as would be expected based on 560 comparing the estimated biological reference points between these methods (Figure 4). In 561 terms of substock diversity, the time-varying maturity models suggested fewer substocks would 562 be overfished or trending towards extirpation at low exploitation rates than the time-constant 563 maturity models, and the regression approaches suggested even more biologically optimistic 564 conclusions (Figure 7). 565

566 3.1.4 Sensitivity analyses

The alternative vulnerability schedule (v_i) based on the spatial distribution of fishing house-567 holds resulted in lower river substocks (those spawning in the Kwethluk, Kisaralik, and 568 Tuluksak rivers) having the lowest v_j ranging between 0.7 - 0.8, the middle river substocks 569 ranging between 0.9 and 0.95, and those in the upper river between 0.95 and 1. When these 570 v_i terms were incorporated in the state-space models, most changes in substock specific 571 $U_{\text{MSY},j}$ and $S_{\text{eq},j}$ were small (±10%, range: -20% - 30%), and most (eight of 13) substocks 572 showed increases in $U_{MSY,j}$. Changes in $U_{MSY,j}$ occurred randomly with respect to changes 573 in vulnerability: some substocks in the three regions showed both increases and decreases. 574 Changes with respect to substock size $(S_{eq,j})$ showed more of a pattern: lower river substocks 575 (i.e., those that became less vulnerable in this sensitivity analysis) tended to become smaller 576 by 5-15%, whereas upper river substocks showed increases of 5-25%. Middle river substocks 577 showed a mix of increases and decreases in $S_{eq,j}$. Despite these changes in substock-specific 578

estimates, derived biological reference points for the aggregate mixed-stock showed a high degree of similarity: escapement-related quantities were ~3,000 fish smaller (<5% change) and harvest-related quantities were ~3,000 fish (~10-15%) larger for the alternative vulnerability assumption. Additionally, the conclusions about substock diversity at MSY were nearly identical: approximately 60% of substocks would be overfished and 23% would be trending towards extirpation.

The alternative age composition weighting resulted in more substantial differences from 585 the default case. Twelve of the 13 substocks showed increases in $U_{MSY,j}$, with two substocks 586 showing increases of approximately 100%. Most substock-specific $S_{eq,j}$ were estimated to be 587 smaller when the alternative scheme was used, with eight of 13 substocks showing decreases 588 between 10 and 35%. Derived biological reference points for the aggregate mixed-stock 589 substantially differed as well: escapement-related quantities were $\sim 6,000 - 10,000$ fish smaller 590 $(\sim 10-15\%$ change) and harvest-related quantities were $\sim 6,000 - 12,000$ fish larger $(\sim 35\%)$ 591 change) for the alternative age composition weighting scheme, however the conclusions about 592 substock biodiversity at MSY were nearly identical, just as for the alternative vulnerability 593 analysis. 594

⁵⁹⁵ 3.2 Simulation-estimation trials

State-space models took between 1.3 and 2.9 days to fit on average (range: 0.8 - 5.1 days), 596 with longer run times associated with more complex models; regression-based models took 597 less than an hour in all cases. The state-space models fitted successfully to the majority of 598 simulated data sets (136 out of 160) and the regression-based models were fitted successfully 590 in all cases. No attempts were made to try different initial values for the relatively few 600 data sets that failed in fitting for the state-space models. MCMC diagnostics suggested 601 that sampling was adequate in the vast majority of cases: >99% of all data sets met the 602 convergence criterion for all parameters in all models and >95% met the criterion for adequate 603 number of samples. 604

Regression models were found to systematically overestimate the population-specific 605 quantities of α_j and $U_{MSY,j}$ and in some cases they produced wildly erroneous estimates. The 606 mixed-effect model was more accurate than the models that fitted independent regressions to 607 each population (Figure 8). State-space models far more accurately and precisely estimated 608 these productivity-based quantities than the regression-approaches, though there was still 609 a slight positive bias (median proportional error $\sim 5\%$; Figure 8). Additionally, regression 610 approaches tended to underestimate $S_{MSY,j}$ more than the state-space models. All state-space 611 models tended to overestimate $\sigma_{R,j}$ by approximately 5% regardless of the assumed covariance 612 structure, though the degree of serial auto-correlation (ϕ) was accurately estimated. The 613 individual state-space models showed essentially no differences in bias for $\sigma_{R,j}$ or ϕ (Figure 614 8). However, models with simple covariance structure tended to overestimate the correlation 615 among populations and vice versa for the complex covariance structure. 616

All state-space models returned unbiased estimates of abundance-related states $(R_{y,j}, S_{t,j}, \text{ and } H_t)$ when including all years. The early portion of the time series had a slight tendency to be overestimated by all models (primarily for $R_{y,j}$) by approximately 5%. This was likely a result of the assumption that data collection began when the populations were unfished.

Just as for the population-specific quantities, the regression-based methods provided 622 generally poorer estimates of mixed-stock biological reference points than state-space models 623 and there was no loss in performance with state-space model complexity (Figure 9). The 624 mixed-effect regression produced more positively biased estimates of $U_{0.1}^*$ and $U_{0.3}^*$ than 625 the independent regression model approach, but this pattern switched for $U_{0.5}^*$ and U_{MSY} , 626 likely as a result of the estimated shape of the distribution of population productivity. The 627 mixed-effect version was less dispersed, meaning that productivities in the lower tail would 628 have been closer to the mean (i.e., larger) than for the independent regression approach. 629 State-space models tended to produce slight underestimates of $U_{0.1}^*$ and $U_{0.3}^*$ and slight over 630 estimates of $U_{0.5}^*$ and U_{MSY} (Figure 9). 631

Credible interval coverage was better for the state-space models than for regression 632 approaches as well. For population-specific parameters, the regression approaches had lower 633 coverage than the state-space models and the models that had the complex recruitment 634 variance structure had more parameters close to the optimal level of 95% (Table 6). All 635 state-space models had low coverage for π which resulted from highly narrow credible intervals, 636 not from inaccurate estimates, though complex maturity models did have slightly better 637 coverage (16% versus 11%; Table 6). In terms of mixed-stock biological reference points, the 638 state-space model provided much better coverage than the regression approaches, particularly 639 for exploitation rate-based points (Table 6). All state-space models exhibited poor coverage 640 for abundance-related states (49 - 66%; Table 6). 641

642 4 Discussion

We presented a novel extension of the age-structured state-space spawner-recruit analytical 643 framework, which has increasingly been applied to single stocks (e.g., Su and Peterman 2012; 644 Hamazaki et al. 2012; Fleischman et al. 2013; Staton et al. 2017; DeFilippo et al. 2018), 645 to the mixed-stock realm. This state-space model was shown to (1) have much less bias 646 and better coverage in key management quantities than regression-based approaches, (2) be 647 robust to structural uncertainty in assumed recruitment covariance and maturity variability, 648 (3) provide good fits to the data, and (4) make more full use of the available data for policy 649 and ecological conclusions. Though it was developed and simulation-tested in the context of 650 Kuskokwim River Chinook salmon, we expect that the mixed-stock state-space framework 651 presented here is general enough to be applied to other systems with similar (and possibly 652 dissimilar) properties and data availability. 653

The simulation-trials illustrated that the state-space model performed superiorly to both evaluated regression-based approaches, regardless of the assumptions made regarding recruitment covariance structure and variability in maturity. Furthermore, the directionality

of the regression biases were consistent with expectations from time-series bias (Walters 657 1985): positive biases in $U_{MSY,j}$ and downward biases in $S_{MSY,j}$, as well as for the respective 658 aggregate mixed-stock quantities $(U_{MSY,j})$ and $S_{MSY,j}$. These biases all but disappeared in 659 comparison for the state-space models for $U_{MSY,j}$ and to a lesser degree for $S_{MSY,j}$. These 660 results speak strongly in favor of the use of the state-space model over the regression-based 661 approaches assessed here. The superior performance of state-space models was likely a result 662 of its ability to (1) explicitly account for the time-series properties in the data, (2) parse 663 observation from process uncertainty, and (3) make more full use of the available data. 664

With respect to the fuller use of the available data, there were 35 brood years for 665 each population in which it was possible to jointly observe spawners and recruits across 666 all four ages (if all calendar years between 1976 and 2017 were monitored). The average 667 Kuskokwim River population fitted using the regression approaches had 17% of the possible 668 observations because partially observed recruitment events were not considered. Conversely, 669 the average population fitted using the state-space model had 31% of possibly observable 670 pairs where the recruitment was observed for three out of four ages, 39% with two ages out 671 of four, and 42% for one age out of the possible four observed. Ignoring whether recruitment 672 and escapement were observed jointly, the average population fitted using the state-space 673 model had 48%, 65%, and 76% of possibly observable recruitment events observed for the 674 same ages combinations, respectively. Clearly, the use of the regression approaches resulted 675 in a severe loss of information. It could be argued that the rule we employed to only use 676 completely observed recruitment pairs for fitting the regression approaches was too strict and 677 that alternative approaches to impute missing observations could be devised. While this may 678 be true, the state-space model provides a comprehensive, rational, and rigorous method to 679 completely reconstruct the brood tables with latent states informed with partial information 680 by fitting to solely observed data. 681

Although the simulation results suggested the state-space model is an unbiased estimator,
 to reduce the dimensionality of our analyses it was assessed under relatively limited conditions.

Su and Peterman (2012) illustrated that state-space models can still show bias under some 684 combinations of measurement error, intrinsic productivity, and fishing intensity. We attempted 685 to evaluate performance across a range of true parameters by randomly sampling leading 686 parameters from the joint posterior from one of the empirical model fits. Though it is possible 687 the model could perform more poorly if (1) population size and productivity were more 688 or less heterogeneous than assumed, (2) fewer populations were used, (3) available data 689 time series were much shorter or sparser, or (4) the magnitude of observation error was 690 incorrectly assumed. All of these scenarios remain exciting avenues for future research, but 691 were beyond the scope of this study. As it currently stands, our analysis suggests that it 692 is reasonable to conclude that the state-space model developed here can be an appropriate 693 estimator for mixed-stock fishery data, though future applications to specific cases should be 694 simulation-tested if the properties of the system and available data differ significantly from 695 the Kuskokwim and the operating model and simulated sampling scheme designed roughly 696 off of it. 697

The policy and trade-off conclusions from the state-space model were generally robust 698 to an alternative assumption regarding relative vulnerability of the populations to harvest, 699 although population-specific estimates did change moderately (primarily $U_{MSY,j}$). The 700 alternative vulnerability vector used was a first attempt at assessing sensitivity to the 701 assumption that all populations have been equally vulnerable to harvest; other more complex 702 approaches to determining this vector could also be assessed in the future. For example, the 703 fishery has historically been focused in on the early portion of the Chinook salmon run in 704 the Kuskokwim River (Hamazaki 2008), and upper river populations have been illustrated 705 to arrive earliest in the summer migration (Smith and Liller 2017a.b). This indicates that 706 upper river populations may be even more vulnerable than what was captured by the vector 707 I used, which was based solely on the spatial distribution of harvesters. Although no data 708 sources were available to directly estimate the vulnerability vector for the Kuskokwim River 709 data, other systems may have these data that could be incorporated, particularly those with 710

⁷¹¹ precise genetic stock identification programs. One would expect larger discrepancies in policy ⁷¹² conclusions to arise when vulnerability covaries more strongly with either population size or ⁷¹³ population productivity; coupled with methods to incorporate information on population-⁷¹⁴ specific harvest data, this provides an provides an interesting avenue for future research.

The conclusions from the state-space models were less robust to alterations in the 715 assumed weighting of age composition data. Specifically, policy recommendations became 716 more aggressive when age composition data received less weight: higher estimates of U_{MSY} 717 and lower S_{MSY} . It is unsurprising that the estimates changed, as this often happens in 718 stock assessment models when the assumed data weighting structure is altered (Hulson et al. 719 2011), but it is unclear as to why weakening the confidence in the age data had the effect of 720 increasing perceived population productivity. Regardless of the cause, this finding suggests 721 that policy conclusions may be conditional on the weighting of the data, and that careful 722 thought should be given to the appropriate weighting scheme. It is likely that the optimal 723 weighting scheme falls somewhere between the two schemes assessed here, given that effective 724 sample size is nearly always less than the true sample size due to violations to the multinomial 725 sampling distribution (e.g., sampled individuals show similarities that result in clustering, 726 non-independence, and overdispersion; Maunder 2011). 727

Regarding state-space model complexity, the important findings were (1) there was 728 no loss in estimation performance with increasing model complexity, (2) credible interval 729 coverage of the complex recruitment covariance models was better than that of the models 730 with simple structures for population-level parameters of interest, and (3) the time-varying 731 maturity models never resulted in wildly large estimates of escapement or harvest as the 732 time-constant maturity models sometimes did for the Kuskokwim data. Based on these 733 findings, it seems that the most complex model is most appropriate, however, as previously 734 stated, the simulation-trials were limited in the scope of biological and sampling scenarios 735 considered, and as such the appropriate model may change if applied to other systems with 736 differing characteristics. 737

The conclusion that the most complex model is most appropriate is contrary to the 738 traditional dogma of assessment model complexity and management performance. Walters 739 and Martell (2004) advise that more complex models may provide more accurate estimates of 740 management quantities, but that their uncertainty will be much greater rendering them less 741 useful for setting harvest policies (see Figure 5.2 and the corresponding discussion therein). 742 These claims have been supported through closed-loop evaluations that have shown simple 743 models known to be wrong but that give conservative advice can provide better management 744 outcomes than complex models that better approximate the true model (e.g., Hilborn 1979; 745 Ludwig and Walters 1985). The simplest model (fewest freely estimated parameters) evaluated 746 here was the mixed-effect regression approach. Although it did provide more conservative 747 and more confident advice than the independent regression approach, the simulation-trials 748 showed that it was biased with respect to U_{MSY} and S_{MSY} in the direction that would lead to 749 more aggressive than optimal harvest policies relative to the state-space models. 750

In some cases, harvest on strong and highly profitable fisheries has been severely 751 curtailed in the name of conserving a few small and unproductive populations. Walters et al. 752 (2018) discuss an example of British Columbia commercial salmon fisheries, were large declines 753 in harvest beginning in the early-1990s and continuing to the present were (in part) a result 754 of intentional reductions in exploitation rates intended to minimize the risk of extinction of 755 a few small and unproductive stocks. The authors argue that fisheries managers have not 756 adequately considered the harvest-biodiversity trade-off in their decision-making processes, 757 and have instead focused on managing for the weakest stocks in the portfolio. Among the 758 authors' four (mostly controversial) recommendations to address this situation, first on their 759 list is for managers to conduct trade-off analyses so that costs and benefits to both fishery and 760 conservation interests can more fully inform decision-making. Our state-space approach shows 761 promise for informing these policy analyses. Particularly for those that involve closed-loop 762 stochastic simulation (e.g., Catalano and Jones 2014), the state-space model provides rich 763 biological estimates to populate the operating models and understand the strength of the 764

portfolio effect (which is inversely related to the magnitude of shared recruitment trends; 765 Schindler et al. 2010, 2015). The model not only provides estimates of leading parameters (α_i 766 and β_j), but also estimates of the extent to which recruitment anomalies are shared among 767 populations and the strength of serial auto-correlation in these time series, all of which would 768 be valuable in populating operating models for policy evaluation. Furthermore, it is possible 769 to use the estimated states of recruitment and spawner abundance and recruitment anomalies 770 at the end of the time series to populate forward simulations from the present to determine 771 which policies might be most likely to achieve short-term objectives in addition to those more 772 focused on the long-term (Connors et al. nd). 773

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TABLE 1. Description of the various index notation used in the description of state-space models. n_t is the number of years observed for the most data-rich stock.

Index	Meaning	Dimensions
y	Brood year index; year in which fish were spawned	$n_y = n_t + n_a - 1$
t	Calendar year index; year in which observations are made	n_t
j	Population index	n_j
a	Age index; $a = 1$ is the first age; $a = n_a$ is the last age	n_a

Symbol	Description
Dimension	nal Constants
n_y	Number of brood years
n_t	Number of calendar years for the population with the longest data time series
n_j	Number of populations
n_a	Number of possible ages of maturation
a_{min}	The first age recruits can mature
a_{max}	The last age recruits can mature
Parameter	rs
$\alpha_j{}^{\mathrm{a}}$	Maximum recruits per spawner for population j
$egin{aligned} eta_j^{\mathbf{a}}\ \sigma^2_{R,j} \end{aligned}$	Capacity parameter for population j ; inverse of $S_{MAX,j}$
$\sigma_{R,j}^2$	Recruitment white noise process variance for population j
$ ho_{i,j}$	Correlation in process variance between subtocks i and j
Σ_R	Recruitment white noise process covariance matrix
ϕ	Lag-1 serial autocorrelation coefficient
$\omega_{y,j}$	Serially autocorrelated portion of recruitment process anomalies (residuals)
π_a	Mean probability a juvenile matures at age a
D^{b}	Dirichlet dispersion parameter for brood year-specific maturity schedules
$p_{y,a}{}^{\mathrm{b}}$	Probability a juvenile belonging to brood year y matures at age a
$U_t{}^{\mathbf{c}}$	Exploitation rate experienced by fully vulnerable populations in calendar year t
$v_j{}^{\mathbf{c}}$	Relative vulnerability term for population j
-	Reference Points
$U_{\mathrm{MSY},j}$	Exploitation rate expected to produce MSY for population j
$U_{\rm MSY}$	Exploitation rate expected to produce MSY for the mixed-stock aggregate
$S_{\mathrm{MSY},j}$	Spawner abundance expected to produce MSY for population j
$S_{\rm MSY}$	Spawner abundance expected to produce MSY for the mixed-stock aggregate
$S_{\mathrm{MAX},j}$	Spawner abundance expected to produce maximum recruitment for population
~	j G
$S_{\mathrm{eq},j}$	Spawner abundance expected to produce the same number of recruits for
C*d	population j
$S_p^{*\mathrm{d}}$	Mixed-stock escapement expected to result in no greater than $p \cdot 100\%$ of
TT+d	populations overfished
$\begin{array}{c} H_p^{*\mathrm{d}} \\ U_p^{*\mathrm{d}} \end{array}$	Same as S_p^* , but for mixed-stock harvest
U_p^{+u}	Same as S_p^* , but for mixed-stock exploitation rate
$p_{OF,MSY}^{d}$	Fraction of populations expected to be overfished at MSY
$p_{EX,MSY}^{e}$	Fraction of populations expected to be trending towards extinction at MSY
States	
$\dot{R}_{y,j}$	Deterministic (expected) recruitment in brood year y for population j
$R_{y,j}$	Realized latent (true) recruitment in brood year y for population j
$N_{t,j}$	Run abundance returning to spawn in calendar year t for population j

TABLE 2. Symbology used in the presentation of the state-space models.

Symbol	Description
$S_{t,j}$	Spawner abundance in calendar year t for population j
H_t	Mixed-stock aggregate harvest in calendar year t
$q_{t,a,j}$	Fraction of the run mature at age a in year t for population j
^a In the st	ate-space models, $U_{\text{MSY},i}$ and $S_{\text{MSY},i}$ were estimated as leading parameters, α_i and

TABLE 2. Symbology used in the presentation of the state-space models. (continued)

^a In the state-space models, $U_{MSY,j}$ and $S_{MSY,j}$ were estimated as leading parameters, α_j and β_j were derived from them using equations found in Schnute and Kronlund (2002).

^b Used only in complex maturity models: SSM-vM and SSM-VM. For simple maturity models, $p_{y,a}$ took the value π_a .

^c In the default case, all populations were assumed to be fully vulnerable, v_j was used in a sensitivity analysis to this assumption.

^d Overfished is defined here as the case where the mixed-stock or any given population is fished with exploitation rate greater than is expected to produce MSY.

^e Trending towards extirpation is defined here as the case where expected equilibrium escapement is less than or equal to zero.

TABLE 3. Summary of evaluated models in this analysis. Regression models are described in Section 2.1.1 and state-space models are described in Section 2.1.2.

Model	n_j	Unique σ_j	AR(1)	Recruitment Covariance	Time-Varying Maturity		
Regress	Regression-Based Models						
LM	10	Yes	No	None	Yes		
LME	10	No	No	None	Yes		
State-S	State-Space Models						
vm	13	No	Yes	Single ρ bounded by $[-0.05 - 1)$	No		
Vm	13	Yes	Yes	Unique $\rho_{i,j}$	No		
vM	13	No	Yes	Same as vm	Yes		
VM	13	Yes	Yes	Same as Vm	Yes		

TABLE 4. Prior distributions used for all parameters in the state-space models for both empirical and simulation analyses. In all cases, priors were selected to be minimally informative while also preventing the sampler from exploring highly unlikely areas of the parameter space. Differences among versions of the state-space model (e.g., SSM-vm and SSM-Vm; Table 3) are described by footnotes.

Parameter	Prior	Description
$U_{\mathrm{MSY},j}$	Uniform(0.01, 0.99)	Exploitation rate that produces MSY
$S_{\text{MSY},j}$	Lognormal(0, 0.001)	Spawner abundance that produces MSY
ϕ	Uniform(-0.99, 0.99)	Lag-1 auto-correlation coefficient
Σ_R^{-1a}	$Wishart(R, n_j + 1)$	Inverse covariance matrix for white-noise
		recruitment process variability
$\sigma_R{}^{\mathrm{b}}$	Uniform(0, 2)	White-noise recruitment process standard
		deviation
$ ho^{ m b}$	Uniform(-0.05, 1)	Correlation in recruitment process variability
		among populations
π	$Dirichlet(\alpha = [1, 1, 1, 1])$	Average probability of maturing at each age
$D^{-0.5c}$	Uniform(0.03, 1)	Dispersion of brood year-specific maturity
		schedules
$p_y{}^{\mathbf{c}}$	$\text{Dirichlet}(\alpha = \pi \cdot D)$	Brood year-specific probability of maturing at
		each age
U_t	Beta(1,1)	Annual exploitation rate of fully vulnerable
		populations

^a Only for SSM-Vm and SSM-VM

^b Only for SSM-vm and SSM-vM, Σ_R was constructed using σ_R and ρ as described at the end of Section 2.1.2.

 $^{\rm c}$ Only for SSM-vM and SSM-VM, all p_y took on π for SSM-vm and SSM-Vm.

TABLE 5. Dimensions for the Markov Chain Monte Carlo algorithms used in this analysis. State-space models were sampled much more intensively than the regression models. Fewer chains were used for the simulation analysis to maximize High Performance Computing efficiency. Oversampling of models fitted to empirical data was intentional to ensure adequate inference.

	Regressio	on Models	State-Space Models		
	Empirical	Simulation	Empirical	Simulation	
Burn-in	20,000	20,000	50,000	50,000	
Post Burn-in	200,000	200,000	800,000	600,000	
Thin Interval	50	50	400	100	
Chains	10	5	10	5	
Total	$2,\!200,\!000$	1,100,000	8,500,000	$3,\!250,\!000$	
Retained	40,000	$20,\!000$	$20,\!000$	30,000	

	Regr	ression	State-space				
Quantity	$\mathbf{L}\mathbf{M}$	LME	vm	Vm	$\mathbf{v}\mathbf{M}$	$\mathbf{V}\mathbf{M}$	
Parameters							
α_j	84	68	95	94	95	95	
β_j	83	73	90	88	90	89	
$U_{\mathrm{MSY},j}$	84	68	95	94	95	95	
$S_{\mathrm{MSY},j}$	85	76	85	84	85	85	
$\sigma_{R,j}$			44	93	45	95	
ϕ			87	97	88	97	
π			12	11	16	16	
$ar{ ho}_{i,j}$			86	93	88	93	
Mixed-sto	ck ref	ference	point	s			
$S_{0.1}^{*}$	99	94	93	92	93	93	
$S_{0.3}^{*}$	99	94	96	96	96	96	
$S_{0.5}^{*}$	94	91	91	90	91	90	
$S_{\rm MSY}$	100	88	94	90	94	91	
$U_{0.1}^{*}$	79	38	96	97	96	96	
$U_{0.3}^{*}$	66	32	99	99	99	99	
$U_{0.5}^{*}$	21	38	95	95	96	96	
$U_{\rm MSY}$	75	80	95	91	95	93	
Abundance states							
U_t			64	64	66	66	
$R_{y,j}$		—	59	58	60	59	
$S_{t,j}$			50	49	51	51	
H_t			51	51	55	55	

TABLE 6. Posterior coverage for key quantities in the simulation-estimation trials. Coverage was calculated as the percentage of all estimated 95% credible intervals across simulated data sets that contained the true value. Bold numbers are those that fall greater than 5 percentage points from the optimal coverage.

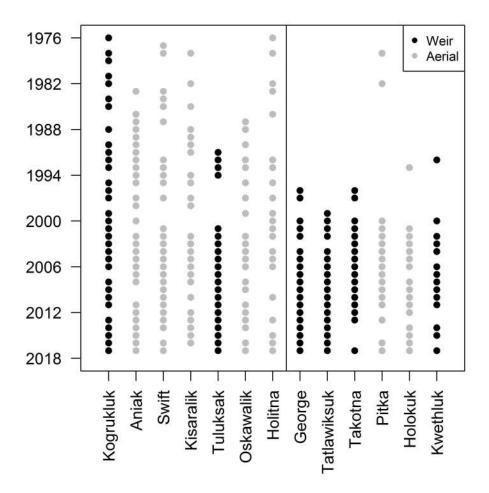
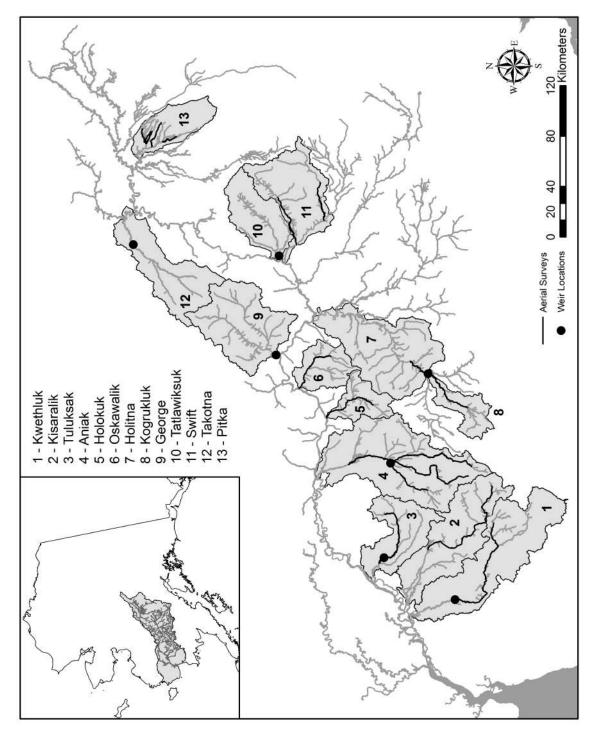


FIGURE 1. The frequency of escapement sampling for each Chinook salmon population monitored in the Kuskokwim River. Black points indicate years that were sampled for populations monitored with a weir and grey points indicate years sampled for populations monitored with aerial surveys. The vertical black line shows a break where > 50% of the years were monitored for a population



this analysis. Black points show the location of weir projects, black sections of river indicate the reaches flown as part of aerial surveys. FIGURE 2. Map of the Kuskokwim River drainage, with the 13 drainage basins representing unique spawning units (populations) used in Drainages monitored via both aerial survey and weir used the weir counts to inform escapement estimates in this analysis, with the exception of the Aniak drainage (#4), for which aerial survey data were much more abundant than available weir data.

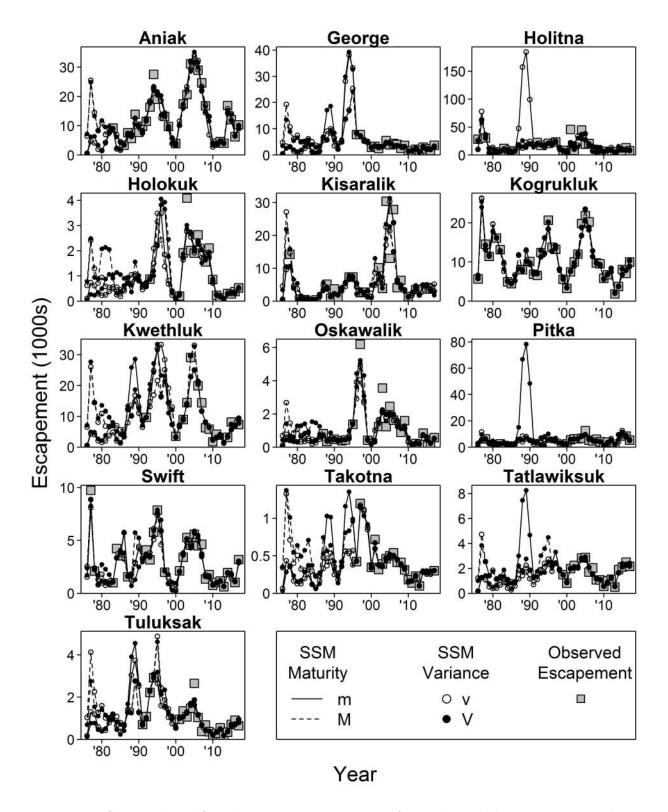


FIGURE 3. Observed and fitted escapement time series for each Kuskokwim River population. Line/symbol types denote the particular state-space model and grey squares denote observed weir counts or expanded aerial survey indices.

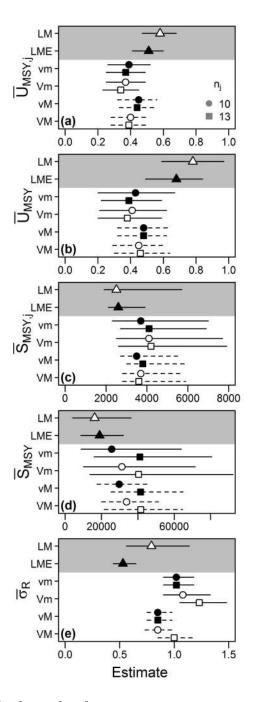


FIGURE 4. Estimates of biological reference points serving as indicators of productivity $(\bar{U}_{MSY,j} - \text{exploitation rate producing MSY}$ from the average population; U_{MSY} – exploitation rate producing mixed-stock MSY) and size ($\bar{S}_{MSY,j}$ and S_{MSY} – same symbology but for escapement), as well as the variability of recruitment anomalies for the average population ($\bar{\sigma}_R$). Six models are shown: two regression models (LM and LME – triangles) and four state-space models (vm, Vm, vM, and VM – circles/squares; models summarized in Table 3). For state-space models, circles represent the values calculated with only the 10 populations able to be fitted with regression approaches included; squares represent the values calculated with all 13 populations.

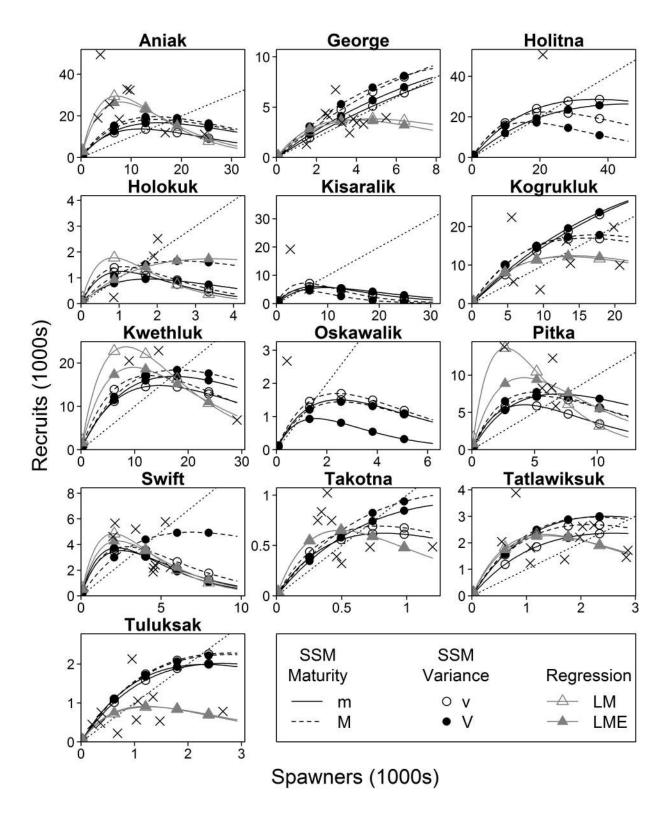


FIGURE 5. Fitted spawner-recruit relationships for the 13 populations monitored in the Kuskokwim River subdrainage included in this analysis. Line and point types denote to different models; crosses are completely observed spawner-recruit pairs. Note that the regression approaches (grey lines/triangles) fitted only to these data, the state-space models (black lines/circles) fitted to all observations of population-specific escapement, aggregate harvest, and age composition.

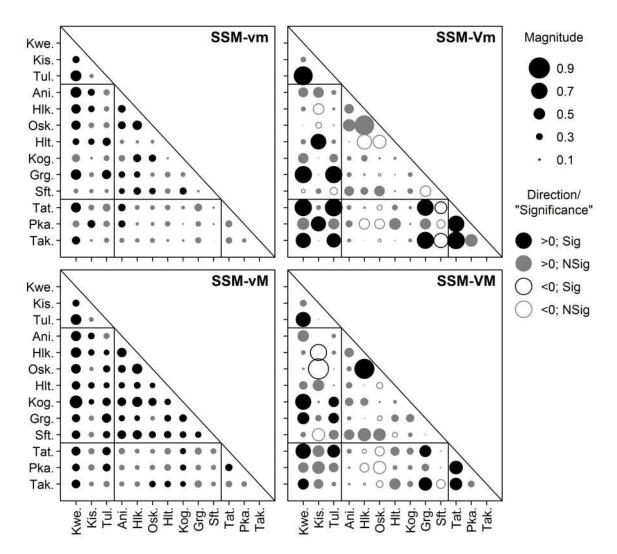


FIGURE 6. Correlation coefficients for recruitment residuals for each pair of populations. The size of each circle represents the magnitude of the correlation, the shade represents significance (whether 95% credible interval included 0), and the fill represents directionality as described in the legend. Populations are ordered from downriver to upriver on both axes, and vertical/horizontal lines denote the boundaries between what are generally considered to be lower, middle, and upper river populations.

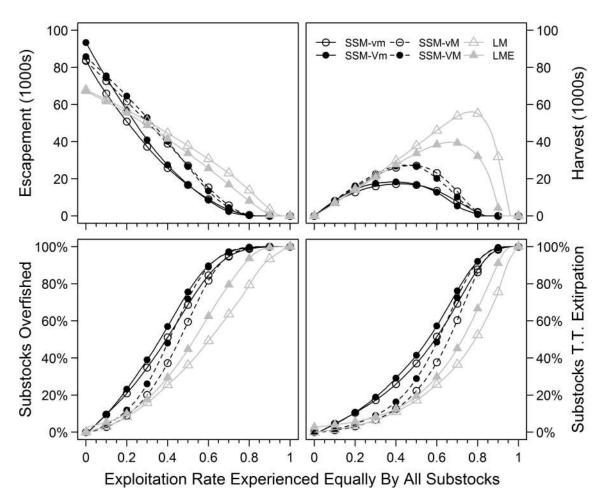


FIGURE 7. Harvest-biodiversity trade-offs based on equilibrium states (escapement and harvest) of the mixed-stock and the percentage of populations expected to be in an undesirable state as a function of the exploitation rate under the assumption that all populations are fished at the same rate. "Overfished" is defined here as $U > U_{MSY,j}$. "T.T." stands for "trending toward", and represents the case where equilibrium escapement would be ≤ 0 . To facilitate comparisons with the regression approaches (grey lines/triangles), the three populations with insufficient data for fitting regression models were excluded from summaries of the state-space models (black lines/circles).

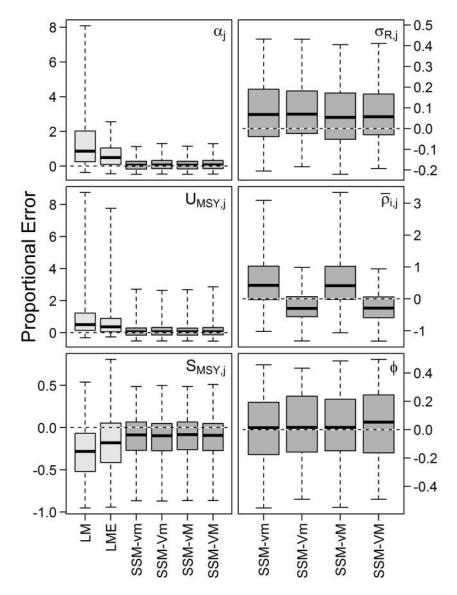


FIGURE 8. Proportional error for key parameters in the multi-stock spawner-recruit models from the simulation-estimation trials. Point estimates used were posterior medians.

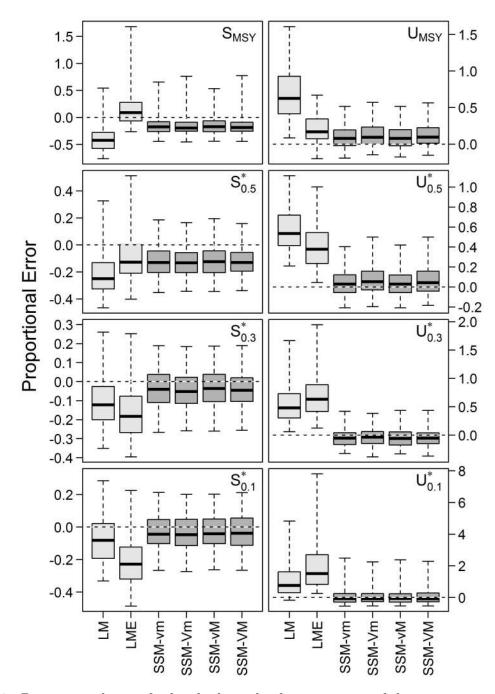


FIGURE 9. Proportional error for key biological reference points of the aggregate mixed stock. S_p^* and U_p^* are the aggregate escapement and fully vulnerable exploitation rate that would ensure no more than $p \cdot 100\%$ of populations are overfished, respectively. Point estimates used were posterior medians.

- 1 Incorporating harvest-population diversity trade-offs into salmon management in large river basins: a
- 2 case study of Kuskokwim River Chinook
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13 Abstract

Variability among fish populations in ecological characteristics and productive capacity (i.e., population 14 diversity) can be critical to maintaining resilience to environmental change and to dampening variability 15 in harvest for fisheries that depend upon them. However, when fisheries for multiple populations overlap 16 in space and time there can be an inherent conflict between harvest and population diversity: high harvest 17 rates, which can be sustained by the most productive populations, can come at the cost of increased risk 18 of overfishing those that are less productive. While the importance of these harvest-population diversity 19 trade-offs in salmon management is well-recognized, they are not often explicitly evaluated in 20 contemporary fisheries management. We used recently obtained estimates of Chinook salmon population 21 diversity within the Kuskokwim River basin of western Alaska, which supports one of the largest 22 subsistence salmon fisheries in the world, to parameterize closed-loop simulations that evaluated how 23 well alternative harvest policies meet population diversity and fishery objectives. We found clear evidence 24 of Chinook population diversity that gives rise to asymmetric trade-offs among fishery and conservation 25 26 objectives. For example, relative to policies that sought to maximize mixed-stock harvest, policies that sought to minimize the risk of driving weak stocks to extirpation did so at the cost of a 20% reduction in 27 harvests and 16% reduction in inter-annual stability in harvests, but with the benefit of increasing the 28 chances of ensuring equitable access to Chinook (i.e., meeting tributary goals) by 84% and nearly 29 eliminating the risk of weak stock extirpation. We also found that harvest policies focused on meeting 30 minimum subsistence needs were unlikely to jeopardize long-term prospects for basin-wide sustainable 31 use. The fishery and biological performance of alternative harvest policies, and the magnitude of resulting 32 trade-offs, were moderately sensitive to potential future changes in population productivity and capacity 33 34 and to uncertainty in the underlying drivers of recruitment variation. Our approach provides a general framework for characterizing salmon population diversity in large river basins and evaluating harvest-35 population diversity trade-offs among alternative harvest policies within them. 36

37 Introduction

Maintaining variation in life history characteristics among species, populations and individuals is 38 increasingly recognized as a hallmark of sustainable and resilient fisheries management. In addition to the 39 inherent importance of protecting such diversity, fisheries that integrate across species and population 40 diversity are often more stable (Sethi 2010; Schindler et al. 2015; Anderson et al. 2017), provide increased 41 food security (Nesbitt and Moore 2016) and are more resilient to environmental change (Anderson et al. 42 2015; Cline et al. 2017). Considerable progress has been made uncovering and quantifying the benefits of 43 species and population diversity, and recognition of the importance of protecting population diversity is 44 now common in national and regional fisheries management policies such as Canada's Wild Salmon Policy 45 (DFO 2005) and Alaska's Sustainable Salmon Policy. However, the tactical incorporation of these 46 considerations into contemporary fisheries management has lagged behind (Walters et al. 2018). 47

Pacific salmon Oncorhynchus spp. provide a classic example of the many dimensions and benefits of 48 species and population diversity. Pacific salmon are broadly distributed across the North Pacific, exhibit a 49 diversity of ecological characteristics and life histories, spawn in thousands of locations across their range 50 and are uniquely adapted to the environmental conditions they experience at fine spatial and temporal 51 scales (Quinn 2018a). This diversity helps to stabilize the aggregate benefits derived from salmon by 52 humans (e.g., Nesbitt & Moore, 2016; Schindler et al., 2010) and wildlife (Schindler et al. 2013; Deacy et 53 54 al. 2019; Service et al. 2019). For example, sockeye salmon fisheries in Bristol Bay, which integrate across hundreds of discrete spawning populations, are two times more stable than if the system consisted of a 55 single homogenous population (Schindler et al. 2010). 56

In situations where fisheries for individual populations within a biodiverse system overlap in space and time, there can be a conflict between harvest and the protection of population diversity because unproductive populations will be unable to sustainably withstand the harvest rates that maximize longterm yield from productive stocks. This gives rise to a trade-off between harvests and the protection of

diversity, known as the "weak" stock problem in fisheries management (Hilborn and Walters 1992; Hilborn
 et al. 2015; Link 2017). Harvest-diversity trade-offs can be acute in large river basins with low levels of
 management precision and control, where fisheries for multiple species and stocks in the marine
 environment or lower river overlap both spatially and temporally (Pestes et al. 2008; Walters et al. 2008).

Harvest-diversity trade-offs may be exacerbated by a changing climate. Climate variation, filtered through 65 spatially variable and nonlinear ecological processes, can give rise to changes in salmon abundance and 66 productivity over inter-decadal and centennial scales (Rogers et al. 2013; Malick et al. 2017). These 67 changes are unlikely to be synchronous across populations and so populations that are relatively 68 69 unproductive and contribute little to fisheries today may become more productive and important contributors to the fisheries of the future (Hilborn et al. 2003; Anderson et al. 2015). It follows that if these 70 weak stocks are traded off for higher harvests today, then their ability to make a greater contribution to 71 harvest in the future may be compromised. 72

73 Harvest-diversity trade-offs may also be influenced by non-stationarity in population productivity which 74 may also obscure the drivers of variation in recruitment in salmon systems. Many salmon spawnerrecruitment relationships are assumed to exhibit over-compensation with declining total recruitment at 75 high spawner abundances (i.e., a Ricker-type spawner-recruitment relationship)(Peterman and Dorner 76 2012; Fleischman et al. 2013; Dorner et al. 2017). Such over-compensation may occur because, for 77 example, at high spawner densities spawning success is reduced due to competition on the spawning 78 grounds. However, cyclical variation in environmental forcing can also give rise to the appearance of 79 overcompensation in spawner-recruit data sets (Parma and Deriso 1990). Such cyclical environmental 80 forcing may also exacerbate harvest-diversity trade-offs because weak stocks are more susceptible to 81 overharvest in years of low productivity than they would be if the primary drivers of recruitment variation 82 were intrinsic. These alternative hypotheses, which are not easily distinguished from each other based on 83

typical spawner-recruit data sets, give rise to structural uncertainty which may impact the evaluation of
 management strategies and perceptions of their ability to meet both harvest and diversity objectives.

Despite the increasing recognition of the importance of salmon biodiversity, there are few published examinations of the performance of alternative harvest policies designed to meet a range of fishery and conservation objectives that incorporate harvest-diversity considerations (but see Hawkshaw & Walters, 2015; Walters et al., 2008). Additionally, quantitative evaluations of the performance of alternative harvest policies in the face of large uncertainty in the drivers, magnitudes and trajectories of recruitment dynamics, have been limited to date (but see Collie, Peterman, & Zuehlke, 2012).

These knowledge gaps were recently highlighted by an Arctic-Yukon-Kuskokwim Sustainable Salmon Initiative independent expert panel that identified the need for empirically grounded closed-loop simulation studies that quantify (1) the predicted consequences of alternative harvest policies in large river basins for both sustainable production over the long-term as well as preserving biocomplexity, and (2) trade-offs among fishery and biological objectives in an attempt to understand the consequences of regime shifts and weak mechanistic understanding of drivers of recruitment (Schindler et al. 2019).

To help close this knowledge gap we characterized salmon population diversity and trade-offs with mixed-98 99 stock harvest in Kuskokwim River Chinook salmon (O. tshawytscha), which support one of the largest subsistence salmon fisheries in the world. We then developed and empirically parameterized a multi-100 stock closed-loop simulation model of the system to evaluate the ability of alternative harvest policies to 101 meet both fishery and conservation objectives. We do this across a range of alternative hypotheses about 102 drivers of recruitment dynamics and non-stationarity in population diversity to quantify how robust 103 alternative harvest policies, and how sensitive harvest-diversity trade-offs, are to a variable environment. 104 105 We find that the inherent Chinook population diversity present in the system gives rise to an asymmetric trade-off between harvest and conserving diversity and that the fishery and biological performance of 106

harvest policies, and the magnitude of resulting trade-offs, are sensitive to directional changes in
 population diversity and uncertainty in the underlying drivers of recruitment variation. Our work provides
 a general framework for characterizing salmon population diversity and evaluating the ability of
 alternative harvest policies to meet multiple biological and fishery objectives in large salmon-producing
 river systems.

112 Methods

Our approach consisted of four steps: (1) establish fishery and population diversity objectives as well as 113 alternative harvest policies through workshops with regional stakeholders, subsistence users and 114 management agencies (US Fish and Wildlife Service and Alaska Department of Fish and Game); (2) 115 characterize population diversity by fitting an age-structured, multi-stock, state-space spawner-116 recruitment model to available data on spawners, harvests, and age composition at a population level¹; 117 (3) quantify the performance of the alternative harvest policies against objectives using closed-loop 118 simulations across a range of plausible hypotheses representing alternative states of nature; and (4) 119 characterize the trade-offs between harvest and diversity that emerge from step 4, and illustrate how 120 alternative harvest policies perform relative to them. 121

122 1.1 Study Area

123 Chinook populations across Western Alaska have declined in abundance and productivity over the past 124 decade (Ohlberger et al. 2016; Dorner et al. 2017). These declines have been particularly pronounced in 125 the Kuskokwim River Basin (Figure 1) where some recent years (e.g., 2012-2014) have seen record low 126 escapement to Chinook spawning grounds, though returns have increased to stable but well below

¹ We use the term population to denote spawning populations that have been the subject of separate assessment related activities within the system (See Figure 1), they are neither a purely biologically or management based unit.

average in the years since then. The Kuskokwim historically supported one of the world's largest Chinook
 subsistence fisheries, which is of importance to dozens of communities along the river that have some of
 the highest levels of subsistence dependence in the state of Alaska. Declines in Kuskokwim Chinook
 abundance have sharply reduced subsistence harvest, closed commercial fisheries, prompted widespread
 stakeholder concern about the future of the subsistence fishery, and led to disagreement among
 managers and stakeholders about the most appropriate way to manage Kuskokwim Chinook.

133 Chinook spawn across at least two-dozen tributaries of the main-stem Kuskokwim, and monitoring of 134 harvest, escapement, and age composition for Chinook has occurred since the mid-1970s with a focus on 135 13 sub-basins monitored by weir or aerial surveys. These populations account for approximately half of 136 annual escapement to the system, and total run size from 1976 and 2017 has been estimated via run 137 reconstruction (Liller and Hamazaki 2016) scaled to drainage-wide mark-recapture estimates of total 138 abundance (e.g., Smith and Liller 2017).

139 1.2 Fishery objectives and alternative management actions

As part of a broader exercise focused on building capacity among Kuskokwim stakeholders to engage in 140 salmon management we held a series of workshops that included influential community members from 141 142 throughout the river basin with a long history of active engagement in fishery management as well as US Fish and Wildlife Service and Alaska Department of Fish and Game biologists and fishery managers. These 143 workshops included discussion of existing and potential Kuskokwim Chinook fishery and biological 144 objectives and alternative management actions associated with them. From these discussions, we 145 identified a suite of long-term objectives against which to quantify the performance of alternative 146 management actions in the system (Table 1). 147

148 1.3 Multi-stock state-space spawner-recruit model

We used a multi-stock state-space spawner-recruit model to characterize Chinook population diversity and dynamics in the Kuskokwim Basin (Staton et al. n.d.). This model is an extension of various singlestock models (Fleischman et al. 2013; Staton et al. 2017) that simultaneously fits separate recruitment curves to incomplete population-specific escapement time series and mixed-stock harvest. We fit the model to time series of spawner abundance, harvest, and age composition in a Bayesian estimation framework using Markov Chain Monte Carlo methods to sample from the joint posterior distribution of all unknown quantities (implemented in JAGS; Plummer, 2017).

The key quantities estimated by the state-space model included population productivity (i.e., maximum 156 recruits-per-spawner), carrying capacity, latent recruitment states, lag-one correlation in recruitment, 157 variance and covariance in recruitment within and among populations, and time-varying maturity 158 schedules. Full details of data processing and the structure of the state-space model, including model code 159 and detailed outputs, are provided in (Staton et al. n.d.) and associated supplements. Our base 160 formulation of the state-space model assumed that all population are equally vulnerable to harvest, but 161 in sensitivity analyses we relaxed this assumption. We chose to use posterior draws obtained from the 162 most complex state-space model evaluated by Staton et al. (n.b). (SSM-VM) with a maximum effective 163 sample size of 100 for the age composition data because (1) effective sample size is nearly always less 164 than the true sample size due to violations to the multinomial sampling distribution (2) their simulations 165 showed no loss in estimation performance with the additional complexity, and (3) it afforded us the ability 166 to parameterize the model more fully with respect to recruitment variance. 167

We used the posterior samples of population productivity and carrying capacity to quantify the range of predicted equilibrium trade-offs between aggregate harvest and conservation of population diversity across a range of mixed-stock harvest rates (Walters and Martell 2004; Walters et al. 2008).

171 1.4 Closed loop forward simulations

To prospectively evaluate the performance of alternative harvest policies with respect to both fishery and 172 conservation objectives, we developed a closed-loop simulation that consisted of three components: (1) 173 an empirically parameterized multi-stock operating model that simulated the dynamics of Kuskokwim 174 Chinook populations over time, (2) a management procedure model that assessed the state (i.e., total 175 176 returning abundance) of the system each year, and applied a given harvest policy (defined by subsistence needs and basin-wide target harvest and spawner abundances), and (3) a performance model that tracked 177 the outcomes of the management procedure for quantitative performance measures related to each 178 fishery and conservation objective. 179

180 *Operating model*

We simulated future population trajectories for the 13 Kuskokwim populations for which we had data by 181 projecting stock dynamics forward over 50 years (approximately eight generations starting in 2017), 182 thereby generating a posterior predictive distribution of future states conditioned on the historical data. 183 By simulating Kuskokwim Chinook dynamics in this manner, we ensured that predicted future spawner 184 185 abundance and age structure were conditioned on the incomplete cohorts at the end of the data series (i.e., those cohorts from which one or more older age classes have not yet returned to spawn) and that 186 uncertainties in the spawner-recruit relationships were propagated (i.e., by drawing from the posterior 187 distributions of each estimated parameter and abundance state in each iteration of the simulation). We 188 chose to use posterior draws obtained from the most complex state-space model evaluated by Staton et 189 al. (n.b)., because their simulations showed no loss in estimation performance with the additional 190 complexity, and it afforded us the ability to parameterize the model more fully with respect to recruitment 191 variance. 192

The simulated population complex was comprised of *n* populations whose dynamics were governed by Ricker type stock recruitment relationships (Ricker 1954):

$$R_{y,j} = \alpha_j S_{y,j} e^{-\beta_j S_{y,j} + \phi \log(R_{y-1,j}/\hat{R}_{y-1,j}) + r_{y,j}}$$
(1)

where $R_{y,n}$ is recruitment from brood year y for population j, α_n and β_j are population specific intrinsic rate of growth (productivity) and within population density dependence, respectively, $S_{y,j}$ is spawner abundance, ϕ is the degree of temporal correlation in recruitment from one year to the next, $\hat{R}_{y-1,j}$ is the expected recruitment in the previous year, and $r_{y,j}$ is white noise variation in recruitment that is correlated among populations according to a common correlation parameter (ρ) following a multivariate normal distribution:

$$r_{y,j} \sim \text{MVN}(0, \mathbf{\Sigma}),$$

$$\mathbf{\Sigma} = \begin{bmatrix} \sigma_1 \sigma_1 & \cdots & \sigma_1 \sigma_j \rho \\ \vdots & \ddots & \vdots \\ \sigma_j \sigma_1 \rho & \cdots & \sigma_j \sigma_j \end{bmatrix}$$
(2)

Returns in year *t*, $N_{t,j}$, were then a function of the proportion of individuals that mature and return to spawn at each age:

$$N_{t,n} = \sum_{a=4}^{7} R_{t-a,n} \pi_{a-3}$$
(3)

where π is a maturity schedule composed of four age classes (4 through 7 for Kuskokwim Chinook). To incorporate the effects of small population size on reproductive success (e.g., allele effects and depensation), we set a quasi-extinction threshold at 50 spawners such that if spawner abundance fell below this threshold recruitment from that brood year was assumed to be zero. For simplicity, we assumed that straying among populations did not occur.

In each year of the forward simulation, harvest $(H_{t,n})$ was subtracted from the return $(N_{t,n})$ according to the harvest control rules described in the following section (Management Procedure model). To simulate the dynamics for the entire Kuskokwim system in each iteration of our forward simulations we expanded spawner abundance, harvest and recruitment by 1/D, where D is normally distributed with a mean of 0.56 and standard deviation of 0.05 and represents an estimate of the proportional contribution of the 13 monitored populations to total Kuskokwim Chinook production (Staton et al. n.d.). This assumes that the monitored populations are a representative sample of all Kuskokwim Chinook populations with respect to population characteristics and time series properties, but was necessary so that the harvest policies we evaluated were on a scale salient to management entities.

217 Management procedure model

The harvest policies we explored were defined by two key management objectives: a basin-wide escapement goal (\tilde{E}) and a harvest goal. Due to the priority placed on subsistence fishing, the harvest goal can be further defined as minimum harvests required to meet subsistence needs (H_{sub}) and a commercial harvest goal (H_{com}) that specifies desired harvest once subsistence needs have been met. These goals, along with the predicted run size in a given year (N_t) determine the harvest rate (U_t) experienced by each population in the forward simulations:

$$U_{t,j} = \begin{cases} (\dot{N}_t - \tilde{E}) / \dot{N}_t, & \hat{N}_t - \tilde{E} \le H_{sub} \\ [H_{sub} + (\dot{N}_t - \tilde{E} - H_{sub})] / \dot{N}_t, & \hat{N}_t - \tilde{E} > H_{sub} \land \hat{N}_t - \tilde{E} \le H_{sub} + H_{com} \\ (H_{sub} + H_{com}) / \dot{N}_t, & \hat{N}_t - \tilde{E} > H_{sub} \land \hat{N}_t - \tilde{E} > H_{sub} + H_{com} \end{cases}$$
(4)

where $\dot{N}_t = \sum_j N_{t,j}$. The forecasted run-size is $\hat{N}_t = \dot{N}_t \varepsilon_N$, where ε_N is forecast error which is assumed to be lognormally distributed with a standard deviation equal to 0.27 (based on retrospective evaluation of forecast error in the Kuskokwim) (Staton and Catalano 2018). Total harvest by population, $H_{t,i}$, is then

$$H_{t,j} = U_t N_{t,j} (1 + \varepsilon_H) \tag{5}$$

where ε_{H} reflects incomplete management control over the harvesting process and outcome uncertainty (Holt and Peterman 2006) and was assumed to be normally distributed with mean zero and standard deviation equal to 0.1. This harvest control rule results in exploitation rates that increase from zero at run sizes less than the escapement goal to a maximum when run size is equal to the escapement goal and target harvest, with the exploitation rate declining thereafter at larger run sizes (Figure 2). Further, it assumed that execution of the fishery was non-selection: the exploitation rate applied to all sub-stocks in the harvest mixture was equal each year.

234 Harvest policies

235 We quantified the performance of harvest policies across a broad range of basin wide escapement and harvest goals and then considered three contrasting harvest policies (Table 2) in more detail to illustrate 236 how policies that prioritize different objectives are predicted to perform across a range of biological and 237 fishery performance measures. The first policy sought to maximize potential harvest from the system by 238 setting a basin wide escapement goal equal to that predicted to produce maximum sustained yield under 239 240 equilibrium conditions. The second was a fixed harvest policy without any escapement goal where the harvest goal is equal to that required to meet minimum subsistence needs. This policy is meant to reflect 241 a situation where fishery managers decide to solely focus on meeting subsistence needs, and minimize 242 inter-annual variation in harvest, regardless of forecasted returns, with the assumption that harvests at 243 the level of subsistence needs will not jeopardize long-term prospects for basin-wide sustainable use. The 244 third policy sought to maximize yield from the system but only once biological risks of extirpation to the 245 least productive populations are minimized. Under this policy the harvest goal was set to equal that in 246 policy 1 but the escapement goal was increased to a level that is predicted to result in no risk of driving 247 the weakest populations towards extinction (see performance measures below). Policy 1 is similar to the 248 basin wide escapement goal approach to management that is currently in place in the Kuskokwim where 249 fishery managers seek to ensure that the aggregate number of Chinook that make it to the spawning 250

grounds fall within the range predicted to provide expected yields greater than 100,000 while also meeting subsistence needs (recommended sustainable escapement goal range of 65,000-120,000)(Hamazaki et al. 2012). However, policy 1 differs from current management because harvesting capacity is constrained in the Kuskokwim due to household processing and consumption constraints.

255 Uncertain states of nature

We simulated the biological and fishery dynamics of the Kuskokwim system across three alternative states of nature. This allowed us to evaluate how robust a given harvest policy is to key structural uncertainties in the system, something that is recognized as a best-practice in the application of closed-loop simulation models to inform decision making (Punt et al. 2016). The first state of nature assumed stationary spawnerrecruitment dynamics that follow a Ricker-type relationship with overcompensation at high spawner abundances (equations 1-3). This is the state of nature that is currently assumed by fishery managers in the Kuskokwim (Hamazaki et al. 2012) and widely across other salmon fisheries.

The second state of nature we considered assumed there were long-term directional changes in 263 population productivity and carrying capacity such that half way through the simulations the least 264 productive stocks became more productive and the most productive stocks became less productive 265 (Figure 5). In this scenario the productivity (α_n) and carrying capacity (α_n/β_n) of individual stocks was 266 assumed to be time-varying and to change over the course of a decade (from t=20 to t=30 in the 267 simulations). These changes were meant to reflect a situation involving large changes in the productive 268 capacity in some stocks and commensurate declines in others (e.g., due to changes in hydrology, 269 geomorphology or life history characteristics) such that the overall production of the system remains 270 unchanged but relative population contributions to production does. The magnitude of change we 271 272 considered under this scenario is biologically plausible; it is within the ranges of changes in productivity

and capacity seen in other salmon systems, albeit at coarser spatial scales (Hilborn et al. 2003; Peterman
and Dorner 2012; Dorner et al. 2017).

The third state of nature we considered assumed that low frequency regime shifts occur that gave rise to the appearance of overcompensation, when in fact none was present. Under this scenario we assumed that individual spawner-recruitment relationships were governed by Beverton-Holt dynamics with cyclical variation in population productivity. To do this we transformed the Ricker spawner-recruitment relationship in equation 1 to a Beverton-Holt form (Hilborn and Walters 1992):

$$\overline{R}_{y,j} = \frac{\alpha_{y,j}}{1 + \frac{\alpha_{y,j}}{\beta_j}} S_{y,j}$$

$$\alpha_{y,j} = \sin(2\pi \left(\frac{y}{f}\right)) \left((\alpha_j \delta + (\alpha_j \delta A)) - \alpha_j \delta \right) + \alpha_j \delta$$
(6)

where population productivity (α_{v_i}) is time-varying following a cycle with a period equal to f and 280 amplitude equal to A. We fixed f and A at 12 and 0.4, respectively based on exploratory simulations, to 281 generate spawner-recruitment relationships with apparent overcompensation that approximates that 282 observed in the Kuskokwim. The term δ is a scalar that adjusts productivity such that long-term population 283 equilibrium abundance in the absence of fishing is approximately the same as those assuming a Ricker 284 spawner-recruitment relationship. We fixed δ at 0.76 based on further exploratory simulations that 285 showed that long-term basin-wide equilibrium spawner abundances were the same under the alternative 286 spawner-recruitment formulations when average productivity in the Beverton-Holt formulation is scaled 287 by 0.76. Note that this scalar may depend on the spawner-recruit relationships used, and should therefore 288 be obtained separately for analyses of different populations in other systems. 289

290 *Performance measures*

Biological and fishery objectives were identified through workshops with fishery managers and stakeholders (Table 2), and we quantified the relative ability of the alternative harvest policies to attain

them according to a set of performance measures. Performance measures related to harvest objectives 293 included average catch and the inter-annual coefficient of variation (CV) in catch, both over the last 20 294 years of the 50 year simulations. We quantified performance relative to the objective of maximizing 295 spatial equity in access to salmon across the river basin as the proportion of populations whose average 296 297 spawner abundance over the last 20 years of each Monte Carlo trial was greater than or equal to the population specific estimate of S_{MSY} . Lastly, we quantified biological performance as the proportion of 298 populations whose average spawner abundance over the last 20 years of the Monte Carlo trial dropped 299 below a guasi-extinction threshold of 5% of equilibrium population size. Each performance measure was 300 summarized across 500 Monte Carlo trials, each parameterized by a unique draw from the posterior 301 distribution of the state-space model fitted to Kuskokwim data (Staton et al. n.b). 302

303 Results

304 2.1 Population diversity

We found clear evidence of heterogeneity in productivity and carrying capacity across the Kuskokwim 305 Chinook populations for which there were data (Figure 3a). Kuskokwim Chinook populations ranged in 306 equilibrium size from approximately 1000 to 18,000 spawners and in productivity from 1.65 to 5 recruits-307 per-spawner (median posterior estimates). There was a weak tendency for larger populations to be more 308 productive, however, no strong spatial patterns in productivity were found (Figure 1 and 3a). We used 309 this observed heterogeneity to quantify equilibrium trade-off between harvest and protection of 310 population diversity in the system (Figure 3b). This illustrates that the relatively high harvest rates that 311 can be sustained by the most productive populations come at the cost of increased risk of over-312 exploitation for those that were found to be less productive. The large uncertainty in our estimates of 313 productivity and carrying capacity result in large uncertainty in these predicted trade-offs. 314

315 2.2 Performance of alternative harvest policies and trade-offs among objectives

Our closed-loop simulations indicated that mixed-stock harvest was maximized at intermediate basin 316 scale escapement goals (~60,000-120,000) (Figure 4a, which is consistent with independent yield analyses 317 from the stock aggregate (Hamazaki et al. 2012). At escapement goals greater than 120,000, harvests are 318 reduced due to more frequent fishery closures, and at escapement goals lower than 60,000 harvests are 319 320 reduced due to the loss of production from the weakest (i.e., least productive) populations in the system. Harvest stability was predicted to be greatest at both low escapement goals and target harvests (Figure 321 4b) and the chances of meeting tributary spawner goals and minimizing risks of driving individual 322 populations to extirpation were greatest when target harvests were low and/or basin-wide escapement 323 goals were high (Figure 4c and d). 324

There was strong asymmetry in trade-offs between harvest and biological or social objectives, with the 325 strength of the asymmetry in trade-offs dependent on the harvest policy that was applied. The harvest 326 327 policy that sought to maximize mixed-stock harvest (policy 1) did so at the costs of increased inter-annual variation in harvest, reduced chances of meeting tributary spawner goals and elevated risk of weak stock 328 extirpation (Figures 4 and 5). In contrast, relative to policy 1, the fixed harvest approach (policy 2) 329 increased inter-annual stability in harvest by 22%, increased the chances of meeting tributary spawner 330 goals by 83%, and reduced median extirpation risk (Figure 3 and 4). This came at the cost, on average, of 331 a 46% reduction in realized overall harvest. The harvest policy that sought to minimize the risk of driving 332 weak stocks to extirpation (policy 3) did so at the cost of a 20% reduction in harvests and 16% reduction 333 in inter-annual stability in harvests, but with the benefit of increasing the chances of ensuring equitable 334 access to Chinook (i.e., meeting tributary goals) by 84% from 46% to 85%. 335

Though there was large uncertainty in the predicted performance of alternative policies, by propagating uncertainty through the closed-loop simulations we can interpret fishery and biological outcomes probabilistically. For example, we found that there was a 41% chance that at least 90% of tributary-level spawner goals would be met if policy 3 was chosen compared to a 1% or 35% chance if policy 1 or 2 were

chosen, respectively. Or from a protection of biocomplexity perspective, there was less than a 20% chance
 of causing at least one population to become extirpated under policy 3, but a 70% or 24% chance if policy
 1, or 2 were chosen, respectively.

343 2.3 Effects of alternative states of nature

We found that the strength of trade-offs between harvest and stability, equity, or risk to diversity were mediated by non-stationarity in the system. When there were large shifts in population diversity over time (Figure 6), we found that similar magnitudes of reductions in harvest must be sacrificed to protect diversity (~30%) but that commensurate increases in equity in access to Chinook were larger than when there were not regime shifts in the system (55% vs 45 %; Figure 7; compare grey and blue bars). In addition, we found that regardless of the harvest policy overall, extirpation risk was elevated were there were pronounced regime shifts in the system.

The performance of harvest policies, and their resulting trade-offs, were also sensitive to the underlying 351 drivers of recruitment variation. When recruitment dynamics were driven by a time-varying Beverton-352 Holt type relationship, absolute harvest and its inter-annual variation were predicted to be lower relative 353 to the alternative spawner-recruitment dynamic scenarios (Figure 7; compare green bars to grey and blue 354 355 bars), and the magnitude of reduction in harvest between harvest policies was greater (50% vs 27%). However, the chances of meeting tributary goals, and extirpation risk, were generally similar across all 356 three spawner-recruitment dynamic scenarios and asymmetry in harvest-diversity trade-offs was 357 dampened: a 50% reduction in predicted harvest from the maximize yield to minimize risk policies was 358 predicted to result in a 45% increase in the chances of meeting tributary goals, compared to 26% and 45%, 359 respectively, under the stationary Ricker scenario. 360

361 **Discussion**

Four key findings emerge from our analyses of Chinook population diversity and performance of 362 alternative harvest policies in the Kuskokwim system. First, we found clear evidence of population 363 diversity where productivity and carrying capacity can vary by as much as 3-fold and 18-fold among 364 populations, respectively. This population diversity gave rise gives to a clear trade-off between mixed-365 stock harvest and conserving population diversity: high harvest rates, which can be sustained by the most 366 productive populations, can come at the cost of increased risk of overfishing those that are less 367 productive. Second, this trade-off was strongly asymmetric: our analysis showed that giving up relatively 368 small amounts of harvest resulted in both large reductions in risk to weak stocks and large gains in spatial 369 equity among subsistence communities in access to the resource. For example, a 20% reduction in average 370 annual mixed-stock harvest by shifting from a policy focused on mixed-stock yield to one that also 371 considers risk to population diversity resulted in an 84% increase in the chances of ensuring equitable 372 access to Chinook across the communities in the river basin and a near complete elimination of risk of 373 374 driving weak populations to extinction.

Third, we found that it is possible for the strength of these harvest-population diversity trade-offs to be 375 mediated by external changes in the environment. When the environment drives pronounced shifts over 376 time in the size and productivity of populations in the system (e.g., a regime shift) giving up a specific 377 378 amount of harvest results in larger gains in spatial equity in access to the salmon for subsistence needs relative to scenarios where there is not a regime shift. These predicted benefits are dependent on the 379 magnitude of change in population size and productivity over time. While the magnitude of change we 380 considered was within the ranges of changes in productivity and capacity seen in other salmon systems 381 (Hilborn et al. 2003; Peterman and Dorner 2012; Dorner et al. 2017), they were of a large magnitude and 382 occurred over a short (10-year) time period (Figure 5). As a result, our findings might be appropriately 383

considered an upper bound to the potential benefits of protecting population diversity within the system
 from a harvest-diversity trade-off perspective.

Fourth, we found that the performance of harvest policies, and their resulting trade-offs were sensitive 386 to the true underlying drivers of recruitment variation. Many salmon systems, including the Kuskokwim 387 are assumed to exhibit strong overcompensation where survival declines at high spawner abundance (i.e., 388 389 a Ricker type spawner-recruitment relationship). However, cyclical environmental forcing can lead to apparent overcompensation in observed spawner-recruitment relationships because years of high 390 recruitment due to favourable environmental conditions are followed by years of low recruitment due to 391 unfavourable conditions (Parma and Deriso 1990). The true drivers of recruitment in a system are rarely 392 known; this uncertainty can fuel debate about the fishery and ecological consequences of alternative 393 assumptions about the structural form of spawner-recruitment relationships when providing harvest 394 advice (Schindler et al. 2019). It is well known that fishery reference points are more biologically 395 conservative (e.g., S_{MSY} is higher) when one assumes a Ricker type spawner-recruitment relationship 396 rather than a Beverton-Holt one (Fleischman et al. 2013); however, the performance of alternative harvest 397 policies across these uncertain states of nature are rarely fully evaluated. We found that absolute harvest 398 and its inter-annual stability were predicted to be lower, and that more harvest had to be foregone to 399 minimize extirpation risk and increase chances of ensuring equitable access to Chinook, under the time-400 401 varying Beverton-Holt scenario relative to the Ricker ones. As a result, the asymmetry in harvest-diversity trade-offs was weaker under the Beverton-Holt scenario. 402

As is inevitable with any analysis of a complex socio-ecological system, we made a number of simplifying assumptions. First, due to incomplete monitoring coverage we were only able to model the dynamics of Chinook populations that comprise approximately half of the production from the system (Figure 1). This implicitly assumes that monitored stocks are a representative sample of all populations with respect to population characteristics and time series properties. If this assumption is severely violated then our

inference about trade-offs and aggregate yield may be biased. For example, if only the most productive populations are monitored then we would have underestimated risks to population diversity. Nonetheless, given that the populations with data are distributed across the basin (Figure 1) and range widely in both their productivity and size we suspect that we have captured the general bounds of the system. It should also be noted that the weakest (least productive) spawning populations in the Kuskokwim may have been extirpated early in the development of the fishery for this river, and are thus absent from our dataset.

Second, our simulations treated each population as a closed spawning population because we did not 415 consider the effects of straying between and within basins. Straying is fundamental characteristic of 416 salmon and clearly an important contributor to adaptation and fitness. Failing to account for straying 417 could lead to overestimating risk of extirpation because populations at low abundance cannot be 418 "rescued" by neighbouring populations. In general, it is hypothesized that the relative frequency of 419 straying in wild salmon is related to stability of habitat quality, extent of specialization for freshwater 420 habitat, and variation in age at maturity (Quinn 2018b). As a result, straying is considered to occur at lower 421 levels in Chinook (and particularly stream-type Chinook like those in the Kuskokwim) than in other Pacific 422 salmon like pink and chum. Empirical estimates of Chinook stray rates are available for hatchery fish and 423 range from 0.1 to 10% for stream-type Chinook in the Columbia Basin (Westley et al. 2013). In light of the 424 above magnitude of straying, and the fact that the populations we considered typically had spawning 425 locations that were greater than 100 km away from the nearest other populations, we considered the 426 magnitude of straying to likely be negligible but nonetheless consider this an area that warrants future 427 study. Strontium isotopes, which have recently been used to uncover fine-scale natal origins and 428 migration histories of Chinook in western Alaska (Brennan et al. 2019), offer a potentially useful tool to 429 quantify the magnitude of straying in the large river basins. 430

Lastly, our closed-loops simulations are empirically parameterized from a multi-stock spawner-recruit 431 analysis (Staton et al. n.d.) that assumes all populations in the system have historically been equally 432 vulnerable to harvest, which was also made in our analysis. While the vast majority of harvest has 433 historically occurred in the lower river near the community of Bethel, differences in run-timing and the 434 435 timing of the fishery (typically front-loaded due to drying weather), are likely to have led to headwater fish being more vulnerable to harvest than lower river populations (Hamazaki 2008). The key findings from 436 our closed-loop simulations are robust to this assumption (Figure S1) but modelling population variation 437 in run-timing and the timing of fisheries are logical extensions to the work we present here. 438

439 We chose three simple and contrasting harvest policies to illustrate their predicted performance against Chinook fishery and population diversity objectives in the Kuskokwim. While the contrasting policies were 440 not intended to exactly match the current approach to management, there are some general insights that 441 emerge from our analysis that are of relevance for management of the Chinook fishery moving forward. 442 We find that a relatively low overall harvest rate is more important than a strict, high escapement goal, 443 with respect to minimizing extirpation risk. This is because our simulations suggest that subsistence needs, 444 and spatial equity in access to fish, can be met with relatively low risk to population diversity even when 445 the escapement goal is very low. However, as harvest goals increase above those required to meet 446 subsistence needs the importance of an escapement goal becomes more pronounced to the point where 447 if target harvest is at or near its historic maximum (~150,000) managing for the upper end of the basin 448 scale escapement goal derived from optimal yield profiles (Hamazaki et al. 2012) is necessary and 449 adequate to minimize extirpation risk. An obvious next step in the Kuskokwim would be to implement a 450 Management Strategy Evaluation that incorporates short-term within-season dynamics and decision 451 making, population differences in run-timing (Smith and Liller 2017b) and harvest vulnerability (Hamazaki 452 2008), with the formal engagement of decision makers and stakeholders in the process (e.g., Cunningham 453 et al. 2018). 454

The performance, and merits, of alternative harvest policies in salmon management have been debated 455 for years. Many salmon systems are managed with escapement goals based on basin scale yield 456 predictions, as in the case in many parts of Alaska and British Columbia. Time varying policies have been 457 shown through simulations to yield improved performance against fishery and conservation objectives 458 459 when there is low frequency changes in the productivity of salmon stocks over time (Collie et al. 2012). However, these types of harvest policies can lead to relatively high variability in harvest, and require 460 precise information on run size to inform annual harvest rates. In instances where stability in harvest and 461 large errors in forecasts dominate, fixed harvest policies (with caps based on conservation constraints) 462 can perform well against fishery and conservation objectives (Hawkshaw and Walters 2015). 463

Maintaining population diversity is increasingly recognized as a hallmark of sustainable and resilient fisheries management. However, in large river basins (e.g., Kuskokwim, Yukon, Skeena, and Fraser) with relatively little fine spatial and temporal management control doing so can be challenging. Our work provides a blueprint for characterizing salmon population diversity in large river basins with relatively limited management control and evaluating harvest-population diversity trade-offs among alternative harvest policies within them.

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- **Table 1.** Fishery and conservation objectives, along with alternative management actions, identified by
- 597 Kuskokwim River stakeholders.

Objectives	Alternatives
 Maintain population and species diversity Maximize equity among villages in access to salmon resource Ensure and protect food security for subsistence users Maximize commercial fishing opportunities 	 Fixed harvest (i.e., minimum required to meet subsistence needs) Manage for basin wide maximum sustainable yield Manage for escapements greater than those predicted to maximize yield so as to protect less productive stocks and ensure equity among villages in access to salmon Manage for population specific escapement goals

Table 2. Alternative harvest policies considered in the closed loop simulations.

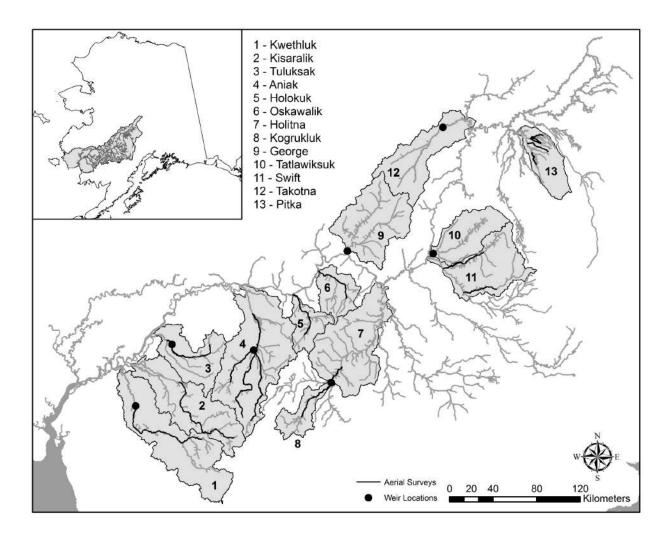
Policy	Basin wide escapement goal (\widetilde{E})	Subsistence needs (H _{sub})	Commercial harvest goal (H _{com})
Maximize basin wide harvest	75,000	65,000	105,000
Fixed harvest at levels needed for subsistence and maximize harvest stability	5,000	65,000	0
Maximize harvest while minimizing biological risk	135,000	65,000	105,000

Table 3. States, biological and policy parameters and associated values. Description of states and

parameters in the close loop simulation model described in the main text along with their associated

values where appropriate.

Parameter	Description (with median values for base scenario where appropriate)	Equation at first use
$S_{y,n}$	Spawner abundance in brood year y from population n	1
$R_{y,n}$	Recruitment in brood year y from population n	1
α_n	Productivity (maximum recruits-per-spawner at small population size) for population n	1
β_n	Magnitude of within brood-year density dependent effects on survival for population <i>n</i>	1
ϕ	Strength of lag-one temporal correlation in survival [0.23]	1
σ_n	Recruitment variation (SD units) for population n [0.69]	2
$ ho_{i,j}$	Correlation in recruitment variation between population i and j [0.18-0.28]	
$\pi_{3:6}$	Age-at-maturity proportions [4yrs=0.25, 5yr=0.36, 6yr=0.35, 7yr=0.4]	3
N _{t,n}	Adult salmon returning to spawn prior to any harvest in fisheries in calendar year t from sub- stock n	3
\dot{N}_t	Total aggregate run-size in calendar year t	4
\widehat{N}_t \widetilde{E}	Forecasted aggregate run-size in calendar year t	4
\tilde{E}	Basin-wide escapement goal	4
H _{sub}	Basin-wide minimum harvest required to meet subsistence needs	4
H _{com}	Basin-wide commercial harvest target	4
$U_{n,t}$	Harvest rate experienced by population n in calendar year t	4
\mathcal{E}_N	Forecast error [~In(1, 0.27)]	4
\mathcal{E}_H	Outcome uncertainty [~N(0,0.1)]	5
f	Period of time-varying productivity in Beverton-Holt spawner- recruitment model formulation [12]	6
Α	Amplitude of time-varying productivity in Beverton-Holt spawner- recruitment model formulation [0.4]	6
δ	Scalar that adjust productivity in Beverton-Holt formulation so that long-term equilibrium abundance is equal to that under the Ricker spawner-recruitment [0.67]	6



- **Figure 1. The Kuskokwim River basin**. General spawning distribution of the 13 Chinook populations for
- ⁶⁰⁷ which there are estimates of spawner abundance based on either weir or aerial surveys.

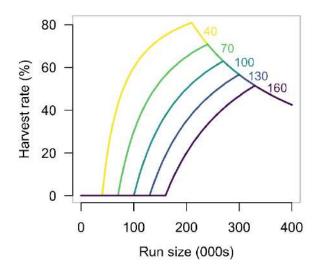


Figure 2. Kuskokwim Chinook harvest control rule. Illustration of mixed-stock harvest rate as a function of true run-size across a range of escapement goals (colored lines) assuming a relatively high (175,000 fish) harvest goal (equation 4). The deterministic harvest control rule is shown but in the closed loop simulations the realized harvest rate in any given year deviate from this relationship as a function of forecast error and outcome uncertainty.

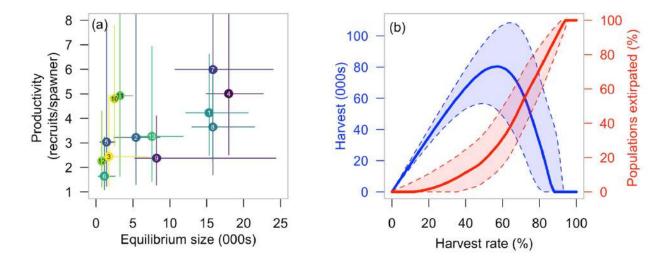
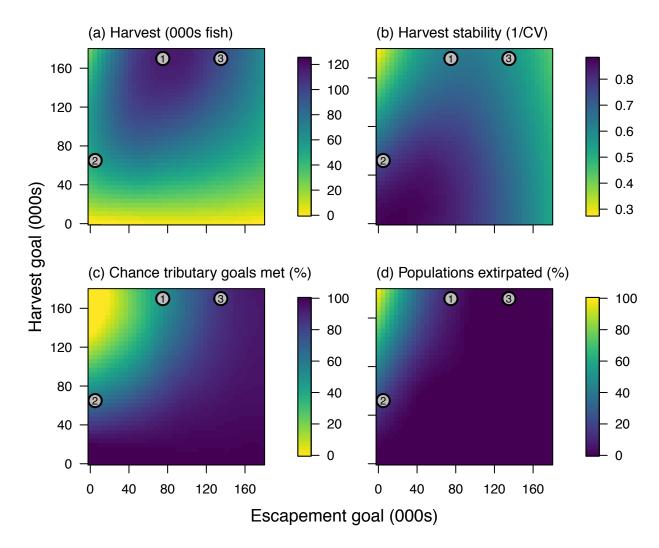
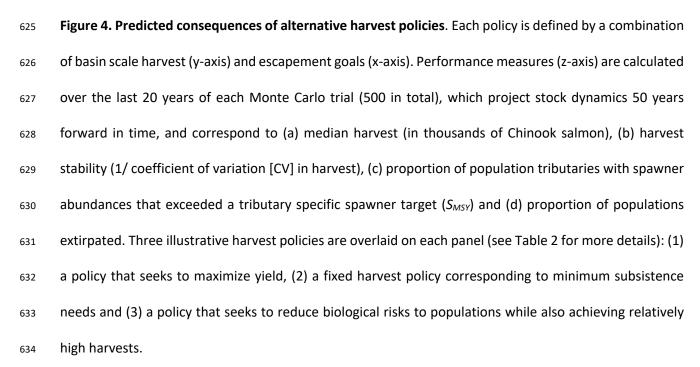




Figure 3. Kuskokwim Chinook population diversity and trade-offs with harvest. (a) Posterior distributions of intrinsic productivity (alpha – recruits per spawner at small population size) for individual population along with corresponding estimates of equilibrium population size. Points are colored to increase visual contrast and numbers correspond to the populations in Figure 1; (b) Predicted basin wide equilibrium yield across a range of fixed harvest rates and corresponding risks to population diversity. Overall yield from the system is predicted to be maximized at a harvest rate of ~ 50%, but this comes at the cost of putting ~20% of the populations at risk of extirpation.







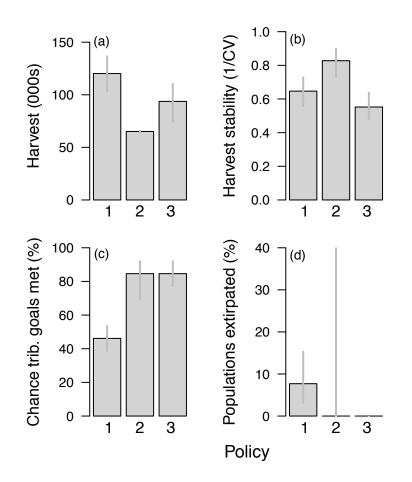


Figure 5. Trade-offs between harvest policies. Predicted ability of three alternative harvest policies (see 637 Figure 4 and Table 2) to meet fishery (a: harvest and b: harvest stability), equity (c: proportion of 638 population tributaries with spawner abundances that exceeded a tributary specific spawner goals) and 639 conservation (d: proportion of populations extirpated) objectives. Each bar is the median (and 25th and 640 75th percentiles) performance of a given policy as calculated over the last 20 years of each Monte Carlo 641 trial (500 in total). Contrasting policies within and among panels illustrates trade-offs in the system 642 between harvest, equity and conservation, as well as asymmetry in them – but it is important to recognize 643 that these three policies are only three points in the policy space as defined by this analysis. 644

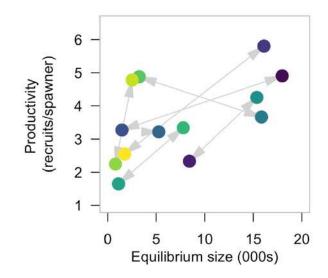




Figure 6. Time varying population diversity. Example of simulated changes (arrows) in the productivity
 and size of individual Chinook populations beginning 20 years into the 50 year forward simulations. Each
 point is the median estimated productivity and equilibrium size for a given population as estimated from
 the multi-stock state-space spawner-recruit model presented in Staton et al (n.b).

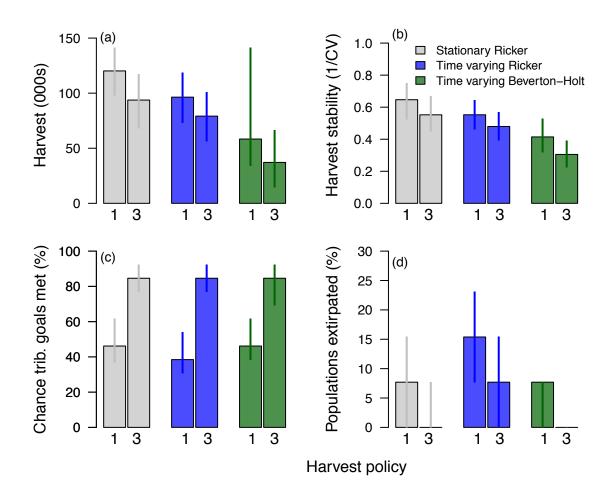


Figure 7. Trade-offs between harvest policies across alternative states of nature. Performance of alternative harvest policies (1 and 3 from Figure 4) across three alternative forms of underlying stock recruitment dynamics (grey bars: stationary Ricker type spawner-recruitment relationship; blue bars: time varying Ricker spawner-recruitment relationship (Figure 6); green bars: time varying Beverton-Holt spawner-recruitment relationship). These plots illustrate that while the general trade-offs are robust to structural uncertainty in the underlying form of the spawner-recruitment dynamics, the magnitude and asymmetry of the trade-offs are sensitive to the underlying drivers of variation in recruitment.

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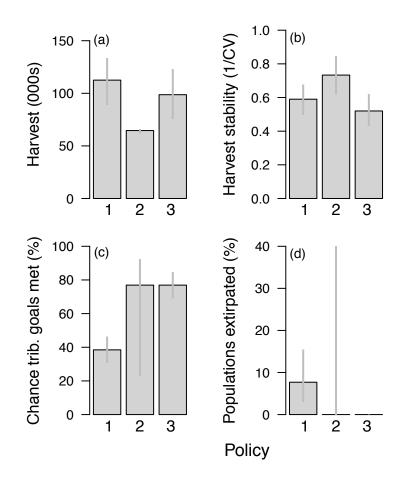


Figure S1. Sensitivity of trade-offs between harvest policies to alternative harvest vulnerability 664 assumptions. Predicted ability of three alternative harvest policies to meet fishery (a: harvest and b: 665 harvest stability), equity (c: proportion of population tributaries with spawner abundances that exceeded 666 a tributary specific spawner goals) and conservation (d: proportion of populations extirpated) objectives 667 based on simulation parameterized from a multi-population spawner-recruit model that assumes all 668 populations in the system have historically been unequally vulnerable to harvest (in contrast to base 669 model that assumes all populations are equally vulnerable; see Figure 5). Each bar is the median (and 25th 670 and 75th percentiles) performance of a given policy as calculated over the last 20 years of each Monte 671 Carlo trial (500 in total). 672