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# Dataset for the Incorporation of Climate Change into a Multiple Stressor Risk Assessment for the Chinook Salmon (Oncorhynchus tshawytscha) Population in the Yakima River, Washington USA

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#### **Supplemental Information**

**Table S1.** Descriptions and derivation of conditional probability tables (CPTs). CPTs were generated using equations, case learning, and pegging the corners, defined below.

Equations, such as derived from regressions on exposure-response data, can be used to derive conditional probability tables. The use of equations in Netica is described here [\(https://www.norsys.com/WebHelp/NETICA/X\\_Equations.htm\)](https://www.norsys.com/WebHelp/NETICA/X_Equations.htm).

Case learning: Case learning is a machine learning tool used to derive relationships between variables. In the BN-RRM we are using parameter case learning since the causal relationships have already been identified. Details on the methodology in Netica can be found at [\(https://www.norsys.com/WebHelp/NETICA/X\\_Learning\\_from\\_Cases.htm\)](https://www.norsys.com/WebHelp/NETICA/X_Learning_from_Cases.htm).

Pegging the corners. When an interaction is indicated and data do not exist to quantify the interaction a 'pegging the corners" technique can be employed (Marcot et al. 2006, Marcot 2017). Pegging the corners sets the extreme cases to the maximum or minimum response possible and then back interpolates the other entries.











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## **Summary of methods used in the BN from Mitchell et al (2021).**

Mitchell et al. (2021) constructed four age-based matrix population models representing each spring Chinook subpopulation in the Yakima River Basin, WA. The study focused on direct effects, and not the indirect effects that may be caused by changes to benthic communities and water flows. The transition matrices were based on stream-type Chinook salmon life history graphs and had a maximum age of 5 years. Matrix transition values (survival and reproduction) and dispersal rates were estimated from subpopulation-specific life history information described in detail in the Supplemental Data of Mitchell et al. (2021). The matrices used baseline models representing metapopulation dynamics prior to impact from pesticides and ecological stressors. An equation for diazinon-malathion synergistic mixture toxicity was derived using the drc package in R (Ritz et al. 2015) to model a log logistic 3 parameter model of the pesticide toxicity data from Laetz et al. (2013). Diazinon and malathion concentrations were converted to toxic units (TU) in the equation for this model by normalizing measured concentrations with EC50s calculated from individual diazinon and malathion dose-response curves (Laetz et al. 2009, 2013).

Measured organophosphate concentrations (malathion and diazinon in μg/L), water temperature 7-day average of the daily maxima (°C), and dissolved oxygen (DO) data (mg/L) were downloaded from the Washington Department of Ecology's Environmental Integrated Management (EIM) database (WDOE 2020) using search filters "Lower Yakima" and "fresh/surface waters" for the monitoring period of 2006 to 2016. The resulting frequency distributions of the data were discretized into bins and incorporated into conditional probability tables (CPTs) dependent on the season. For pesticide data, non-detects ("U" ‐qualified) and data detected below the reporting limit ("J"‐qualified) were captured in the lowest pesticide concentration bins in the Bayesian network. The lowest pesticide concentration bins were parameterized as 0 μg/L to the US Environmental Protection Agency (USEPA) aquatic life criteria for diazinon (0.17 μg/L) or malathion (0.1 μg/L) (USEPA 2020) and were associated with no toxic effect (no effect on AChE activity) in the model. Additional details can be found in Mitchell et al. (2021) and the associated Supplemental Information.

The Puget Sound Partnership's (PSP's) recovery goal for Chinook salmon is "no net loss" of population abundance (PSP 2020). This metric was used to define risk to Chinook in the present study and has previously been used to define risk by Landis et al. (2020) and Mitchell et al. (2021).The initial abundance for all subpopulation models was set to 500 000 fish, which approximates a typical salmon population's spawner abundance under a stable age distribution for the region. Risk was defined as the probability that the resultant metapopulation size is below the initial abundance of 500 000 fish.

The sensitivity of the Bayesian network Chinook population size endpoint to various nodes was calculated using Netica's "Sensitivity to findings" function to calculate percent mutual information (Norsys 2014).

## **Supplemental Information Tables**

SI Table 1. Modifications (additions) to the temperature and dissolved oxygen distributions for seasons and climate change scenarios. All modification (mod) distributions are normal distributions with the mean and standard deviation indicated below. Modifications adapted from Ficklin et al. (2013) percentile and standard deviation ensemble model statistics across the Sierra Nevada for the 2050s and 2080s for spring and summer. Low=25<sup>th</sup> percentile; medium =  $50<sup>th</sup>$  percentile; high =  $75<sup>th</sup>$  percentile of SWAT model simulations forced with 16 general circulation models.





SI Table 2. Scenarios explored in the amended Bayesian network model from Mitchell et al (2021) for the lower Yakima metapopulation. Present = original distributions from Mitchell et al. (2021) model. See Tables 1 and 2 and text for additional explanation of scenarios.



#### **Supplemental Information Figures**



SI Figure 1. Conceptual flow for modifying monitoring concentration distribution nodes from Mitchell et al. (2021) for dissolved oxygen. (a) "Present" state selected in *Scenario* with no resulting change to the original *Dissolved Oxygen (mg/L)* distribution so both *Dissolved Oxygen (mg/L)* nodes contain the same distributions, (b) "Future" state selected in *Scenario* so the distribution in the parent *Dissolved Oxygen (mg/L)* is subtracted from the distribution in the modification node to create a modified distribution in the child *Dissolved Oxygen (mg/L)*. A similar modification was made for the Water *Temperature 7-DADMax* distribution. Pesticide distributions were modified through percent additions or subtractions.



SI Figure 2. Adjustment network for the malathion, diazinon, water temperature, and dissolved oxygen monitoring nodes. Nodes are grouped into pesticides (green); scenarios (bright blue); and water quality (dark blue). Mod=modification factor; DO= dissolved oxygen; temp = temperature. Pesticide concentration distributions are reduced or increased by percentages based on hypothetical futures in the *Spray scenario* node. Water quality nodes are adjusted by adding their corresponding "mod" nodes based on scenario projections.



SI Figure 3. Scenario example for summertime monitoring data in the Lower Yakima with a high 2080 projected adjustment to water quality (temperature and dissolved oxygen) and a high spray scenario. Note the changes in the values of the nodes compared to SI Figure 2.



SI Figure S4. Sensitivity analysis to population node for the present, 2050, and 2080 year scenarios with none, present, and high pesticide loads using mutual information (%). Note the differences in the contributions as measured by mutual information for the spring (yellow) compared to summer (blue). Note the importance of the metapopulation node is in summer compared to the spring.