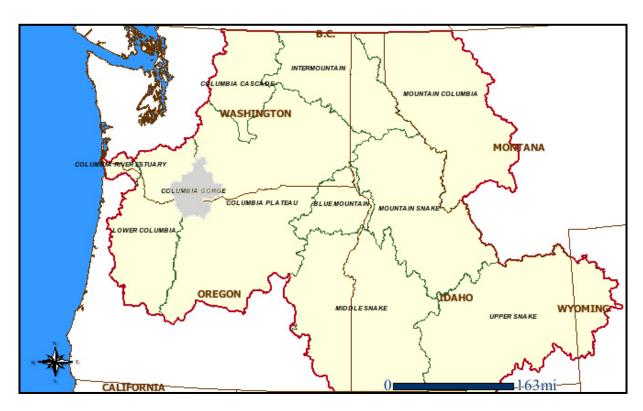


U.S. Fish and Wildlife Service - Pacific Region

Columbia River Basin Hatchery Review Team

Columbia River Basin, Columbia Gorge Province Little White Salmon, Big White Salmon, and Wind River Watersheds



Carson, Spring Creek, Little White Salmon, and Willard National Fish Hatcheries

Assessments and Recommendations

Final Report, Appendix A:

All-H Analyzer (AHA) Output for Salmon and Steelhead Stocks

December 2007

Appendix A: All-H Analyzer (AHA) Output for Salmon and Steelhead Stocks Carson, Spring Creek, and Little White Salmon National Fish Hatcheries, including natural populations in the Wind, Big White Salmon, and Little White Salmon Rivers.

What is AHA?

AHA is an *Excel*-based spreadsheet simulation model that quantifies the mean number of adults returning to a watershed after many generations (years) of reproduction and migration based on equilibrium, or near equilibrium, conditions. Recent versions of AHA (Versions 6.x and 7.x) for the Columbia River allocate returning adults to six physical geographic locations: (1) a hatchery and other recapture facilities within the watershed under consideration; (2) the natural habitat within the watershed where adults spawn; (3) marine harvest areas; (4) the lower Columbia River mainstem downstream from Bonneville Dam; (5) the upper Columbia River mainstem upstream from Bonneville Dam, and (6) a terminal harvest area in the watershed where adults return to spawn. The model was developed primarily by Lars Mobrand (Mobrand Biometrics), in collaboration with the Washington Department of Fish and Wildlife (WDFW) and the Northwest Indian Fisheries Commission (NWIFC), as part of the HSRG review¹ of salmon and steelhead hatcheries in Puget Sound and coastal Washington state.

AHA is based on the Beverton-Holt spawner-recruit model where habitat *capacity* represents the maximum number of adult recruits (asymptote of the Beverton-Holt curve) that the habitat can produce and return to a watershed, and *productivity* represents the slope of the spawnerrecruit curve at the origin (i.e., the number of adult recruits per adult spawner [R/S] when the number of adult spawners is very low and density-dependent factors or competition can be ignored). The actual model (spreadsheet) consists of several sheets (e.g. natural component sheet, hatchery component sheet, genetic fitness sheet, etc.) where estimated mean values of biological and population dynamic parameters are provided by the user (e.g. mean fecundity of females, estimated egg-to-smolt survival, etc.). The genetic fitness function is based on the model of Ford (2002)² and allows the mean fitness of a population (productivity) to decrease incrementally over time depending on (a) the mean proportion of natural spawners composed of hatchery-origin adults (pHOS) relative to the mean proportion of the hatchery broodstock composed of natural-origin adults (pNOB), and (b) the number of generations that hatcheryorigin fish spawn naturally in the watershed. The model is currently being used by fishery comanagers in the Pacific Northwest as a "planning tool" to (a) document assumptions and goals (e.g. current and future habitat conditions, respectively) and (b) assess the likelihood that harvest and conservation goals can be achieved given the aforementioned assumptions and desired future conditions. Only those scenarios achieving realistic or desired outcomes are

¹ www.hatcheryreform.org

² Ford, M.J. 2002. Selection in captivity during supportive breeding may reduce fitness in the wild. Conservation Biology 16: 815-825.

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considered valid. For example, any scenario that results in extirpation of a stock is considered invalid, where any or all of the four H's³ can contribute to stock extirpation.

For more detail on AHA, see AHA Technical Discussion Paper on the Publications page of www.hatcheryreform.org. An AHA user's guide and all AHA analyses are available from the AHA section of the prototype Managing for Success web site at www.mobrand.com/mfs (log in with user name and password "public"). The AHA User's Guide is also available at www.fws.gov/pacific/fisheries/hatcheryreview/reports.html/.

Explanation of Tables A1-A74

Information in the following tables (Tables A1-A7) is intended to provide a summary "snapshot" of the predicted future outcomes associated with current hatchery programs and alternatives considered by the Review Team. Seven sets of simulations are presented: (1) spring Chinook in the Wind River including Carson NFH spring Chinook; (2) summer steelhead in the Wind River including two hatchery program "alternatives"; (3) fall Chinook in the Wind River (current conditions only); (4) tule fall Chinook at Spring Creek NFH and in the Big White Salmon River after removal of Condit Dam; (5) spring Chinook at Little White Salmon NFH and in the Little White Salmon River; and (6) spring Chinook at Little White Salmon NFH and in the Little White Salmon River.

Output of AHA is displayed in a series of colored bar graphs representing adult fish (recruits). Solid green represents natural-origin fish; solid pink represents hatchery-origin fish. Pink diagonal hash bars represent hatchery-origin fish in excess of comanager goals, and gray vertical bars represent hatchery-origin fish that have been selected at least one generation in the natural environment (e.g. as occurs in a genetically-integrated hatchery program).

The graph in the lower-left portion of each table shows the realized mean values of pNOB and pHOS (mean proportions of the hatchery broodstock and natural spawners composed of natural-origin adults and hatchery-origin adults, respectively). The diagonal lines represent combination values of pNOB and pHOS that yield the same value of the parameter, PNI, which stands for proportional natural influence, where PNI = pNOB/(pNOB + pHOS). PNI varies from 0.0 to 1.0 and represents the relative degree to which the genetic constitutions of hatchery-origin fish and/or natural-origin fish are influenced by the natural environment versus the hatchery environment. When pHOS = 0.0 and pNOB > 0.0, then PNI = 1.0 and the genetic constitution of natural-origin fish will be determined by the natural environment only. (Note: In practice, pNOB must be greater than 0.1 to overcome random genetic effects and single-generation selection effects of the hatchery; otherwise, hatchery fish will essentially represent a "hatchery stock" genetically regardless of the value of pHOS.) When pHOS = pNOB, then the hatchery and natural environments will have equal influence on the genetic

Appendix A - AHA Output

³ Habitat, Harvest, Hydropower effects, Hatchery program effects.

⁴ Parameter estimates used to generate the following tables have not all been verified and should be considered preliminary. However, their values are based on the best information available, and the general results presented in the following tables are not expected to change significantly as the parameter estimates are verified and updated.

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constitutions of both hatchery and wild fish, and PNI = 0.5. For integrated hatchery populations and natural stocks, the goal is for PNI to be greater than 0.5 and as close to 1.0 as possible (e.g., > 0.67) as a long-term goal. Symbols on the PNI graph correspond with each of the current and alternative scenarios shown as bar graphs (see also the *Components of this Report* section for definitions of biological significance, population viability, habitat, and harvest ratings).

The outputs presented in Tables A1-A7 are intended to be viewed as part of an electronic report via a desktop computer where portions of the tables can be zoomed in and out for clarity. The Review Team acknowledges that they are difficult to read as printed paper copies.

The data files and specific version of AHA (V.7.2.4) used to generate the outputs in Tables A1-A7 are available upon request form the Chair of the Hatchery Review Team. Some familiarity with AHA is required, and expected, before implementing the software.

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Table A1. AHA output for spring Chinook in Wind River, including Carson NFH spring Chinook. See report for descriptions of Alternatives.

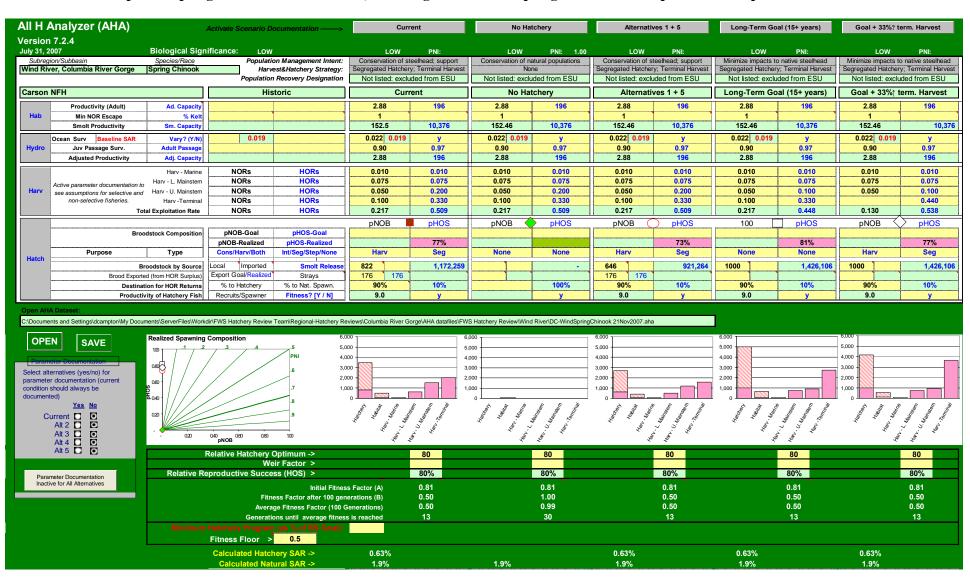
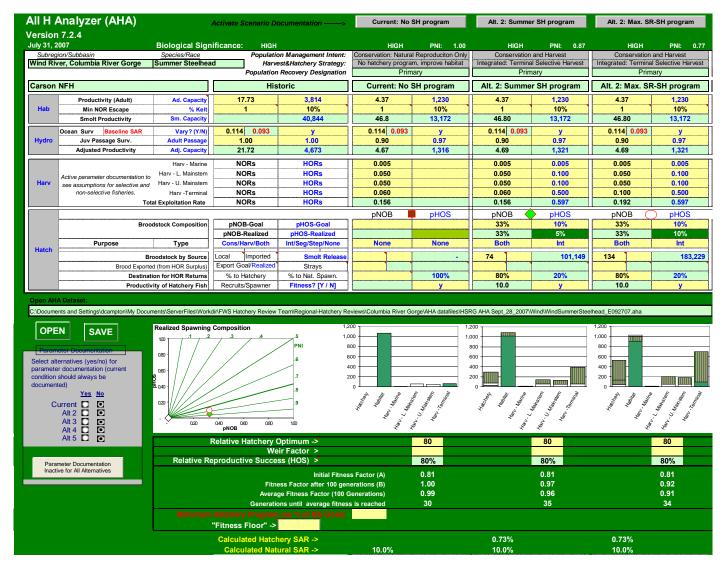
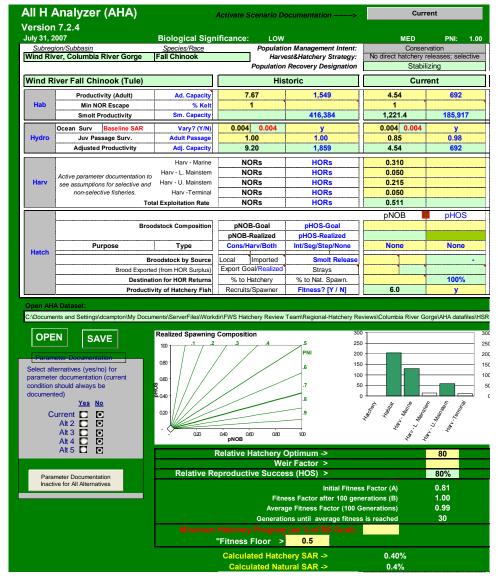


Table A2. AHA output for summer steelhead in Wind River, including an integrated hatchery program at Carson NFH. Two hatchery programs of different size (Alternative 2) are compared to the current "no program" scenario: (1) 100,000 smolt release that maintains pHOS \leq 5%, and (2) the maximum-sized program (180,000) that maintains pHOS \leq 10% and PNI > 0.67. The larger program reduces the number of natural-origin recruits.



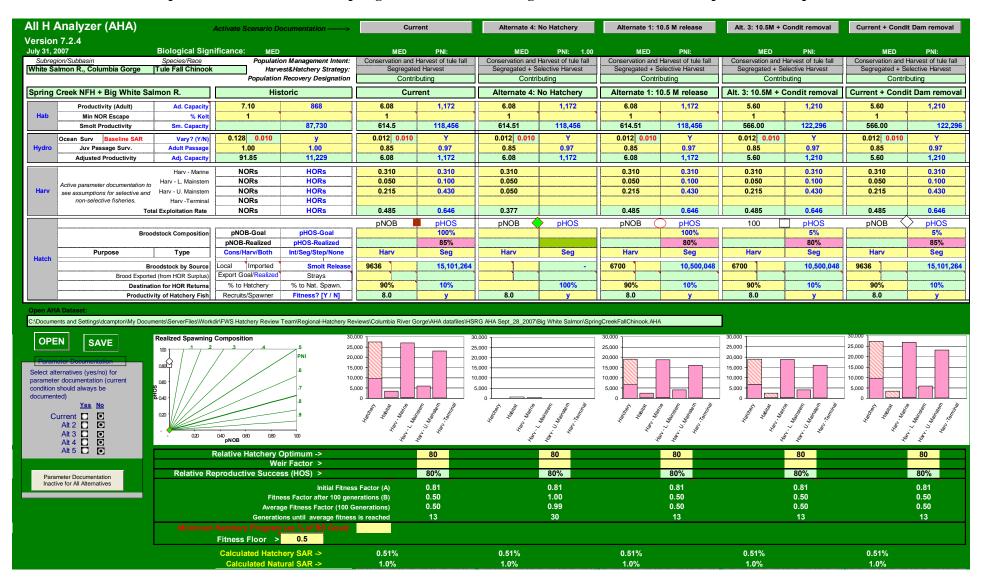
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Table A3. AHA output for fall Chinook in Wind River. The Review Team did not evaluate alternatives for the current management of fall Chinook in the Wind River.



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Table A4. AHA output for tule fall Chinook at Spring Creek NFH and the Big White Salmon River. See report for descriptions of alternatives.



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Table A5. AHA output for spring Chinook at Little White Salmon NFH. The current hatchery program transfers 250,000 yearlings to Walla Walla River, but these transfers are to be assumed by Carson NFH in 2008. Under Alternative 2, the current Carson-strain broodstock is replaced by the Klickitat River stock via a "stepping stone" program where approximately 20% of the broodstock is derived from the integrated Klickitat River stock each generation.

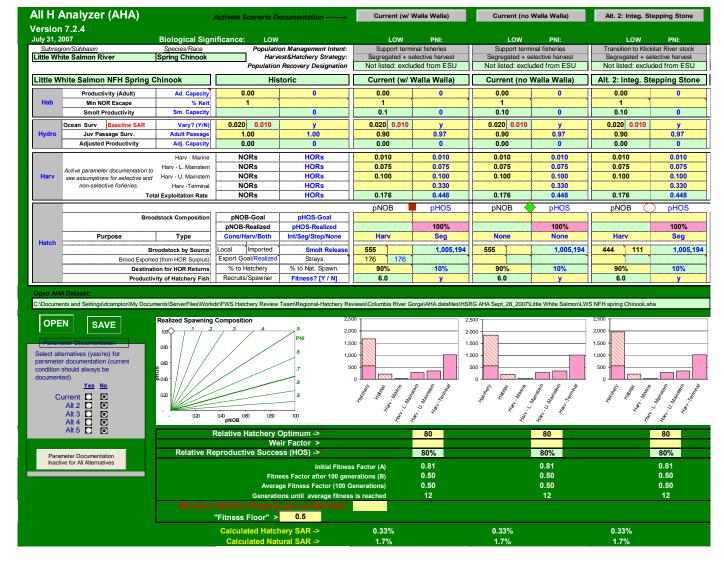
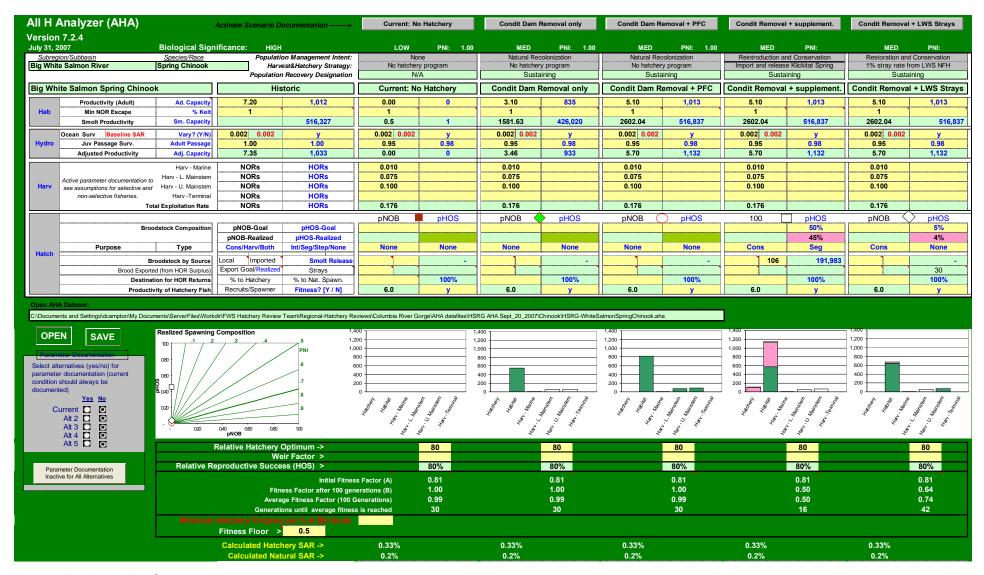


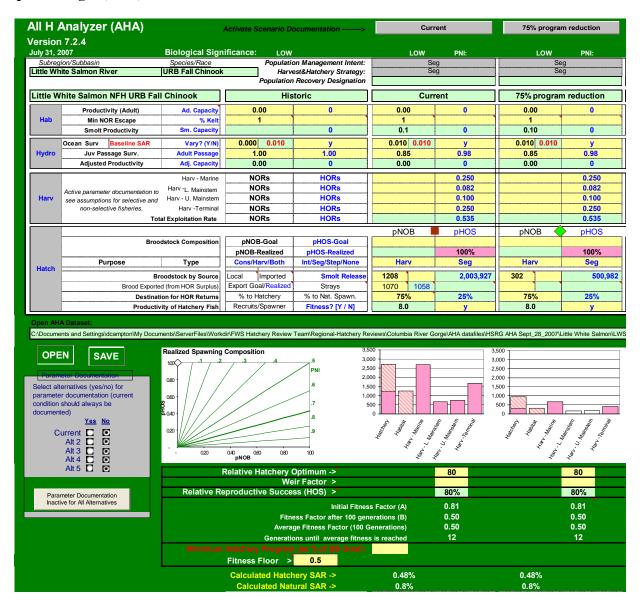
Table A6. AHA output for spring Chinook in the Big White Salmon River. The second and third scenarios after Condit Dam removal show habitat productivities and capacities before and after restoration of "properly functioning conditions" (PFC). The fourth scenario models supplementation outplanting of the Klickitat River stock, while the last (fifth) scenario models strays (1% stray rate assumption) from Little White Salmon NFH.



Appendix A - AHA Output

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Table A7. AHA output for upriver bright (URB) fall Chinook at Little White Salmon NFH and Little White Salmon River.



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For Columbia River Basin Hatchery Review Information www.fws.gov/pacific/Fisheries/Hatcheryreview/

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December 2007

