

**Title:** Defining relative reproductive success: which fish count?  
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1

## Abstract

2 How the Alaska Hatchery Research Group defines reproductive success will have implications  
3 on how the fitness effects of hatchery fish on natural populations are estimated. Here we  
4 present: 1) a proposed mathematical definition for relative reproductive success from the  
5 literature, 2) different definitions of “potential parent” and “adult offspring” used to estimate  
6 reproductive success and how these definitions affect the determination of relative reproductive  
7 success, and 3) concerns about the necessary changes to field sampling in 2014 to accommodate  
8 different definitions of reproductive success. We recommend that reproductive success be based  
9 in the context of management by escapement by including as many  $F_0$  potential parents and  $F_1$   
10 potential offspring as possible, including those that produce no adult offspring such as pre-spawn  
11 mortalities in order to determine productivity in relation to escapement.

12

## Background of AHRG

13 Extensive ocean-ranching salmon aquaculture is practiced in Alaska by private non-profit  
14 corporations (PNP) to enhance common property fisheries. Most of the approximately 1.7B  
15 juvenile salmon PNP hatcheries release annually are pink salmon in Prince William Sound  
16 (PWS) and chum salmon in Southeast Alaska (SEAK; Vercesi 2013). The large scale of these  
17 hatchery programs has raised concerns among some that hatchery fish may have a detrimental  
18 impact on the productivity and sustainability of natural stocks. Others maintain that the potential  
19 for positive effects exists. ADF&G convened a Science Panel (Alaska Hatchery Research  
20 Group; AHRG) whose members have broad experience in salmon enhancement, management,  
21 and natural and hatchery fish interactions. The AHRG was tasked with answering three priority  
22 questions:

23 I. *What is the genetic stock structure of pink and chum salmon in each region (PWS and*  
24 *SEAK)?*

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<sup>1</sup> This document serves as a record of communication between the Alaska Department of Fish and Game Commercial Fisheries Division and other members of the Alaska Hatchery Research Group. As such, these documents serve diverse *ad hoc* information purposes and may contain basic, uninterpreted data. The contents of this document have not been subjected to review and should not be cited or distributed without the permission of the authors or the Commercial Fisheries Division.

- 25 II. *What is the extent and annual variability in straying of hatchery pink salmon in PWS and*  
26 *chum salmon in PWS and SEAK?*
- 27 III. *What is the impact on fitness (productivity) of natural pink and chum salmon stocks due*  
28 *to straying of hatchery pink and chum salmon?*

29 **Introduction**

30 *Measuring the Impact on Fitness*

31 To answer the third question, we need to know the origin and pedigree of each fish captured in  
32 select streams across multiple generations. **Origin** refers to the type of early life-history habitat  
33 (hatchery or natural) that a fish experienced. **Pedigree** refers to the family relationship among  
34 parents and offspring. ‘**Ancestral origin**’ refers to the origin of an individual’s ancestors (e.g.,  
35 two parents of a single origin [hatchery/hatchery or natural/natural] or two parents of mixed  
36 origin [hatchery/natural]). These ancestral origins can be determined by combining information  
37 from three sources: natural or hatchery origin from otolith marks, pedigree from genetic data,  
38 and age from scales (for chum salmon from SEAK). By pairing these data within fish and across  
39 generations, we can estimate **reproductive success (RS)** among cross types (i.e. hatchery-  
40 hatchery, hatchery-natural, and natural-natural origin crosses). The AHRG is using the **relative**  
41 **reproductive success (RRS)** of hatchery-origin fish to natural-origin fish as the measure of  
42 *fitness in this study.*

43 *Problem: What is Relative Reproductive Success?*

44 The RRS of hatchery-origin to natural-origin fish can be calculated in multiple ways depending  
45 upon the definition of RS. *The **average RS, or fitness, of a group is defined in the AHRG***  
46 *priority questions as productivity.* **Productivity** of a system is typically defined as the average  
47 number of adult offspring ( $F_1$  recruits) per potential parent ( $F_0$  spawners). The exact definition  
48 of both ‘**potential parent**’ and ‘**adult offspring**’ is not clear in this study and can be defined as  
49 *one of five categories of adults observed in or near freshwater streams* (Figure 1). Different  
50 definitions of ‘potential parent’ and ‘adult offspring’ produce different measures of RS and  
51 require different field sampling procedures in order to accurately estimate  $F_0$  potential parents  
52 and  $F_1$  adult offspring. The AHRG needs to decide upon a definition of potential parent and  
53 adult offspring prior to the 2014 summer field season to guarantee that the data collected will be  
54 able to adequately answer the proposed research questions.

55 *Goals of Technical Document*

56 Four goals of this technical document are to:

- 57 1) Propose a mathematical definition of RRS from the literature (Araki and Blouin 2005);  
58 2) Explain how the definition of potential parent and adult offspring affects RS and the  
59 estimation of RRS;  
60 3) Describe how different definitions of RS require different field sampling methods; and  
61 4) Request a decision by the AHRG on the definition of RS prior to the 2014 field season.

$$RS = \frac{F_1 \text{ returns}}{F_0 \text{ spawners}} = \frac{\text{Diagram 1}}{\text{Diagram 2}}$$

62

63 **Figure 1:** Equation for RS depends on which category of “adult offspring” and “potential parent” observed in streams is included  
 64 in the numerator and denominator, respectively. Five definitions of adults and offspring can be visualized as concentric circles.  
 65 The orange (bold) and yellow (dashed) shaded circles are different definitions of “potential parent” and “adult offspring” that  
 66 require different field sampling methods regarding the inclusion of pre-spawn mortalities. The largest, most inclusive circle is  
 67 what managers’ observe during aerial surveys of escapement.

68

### Mathematical Definition of RRS

69 While the intuitive definition of RS (i.e. number of  $F_1$  returns per number of  $F_0$  spawners) works  
 70 well in situations where there is little to no error in parentage assignment and all parents are  
 71 sampled, research has shown that both parentage assignment error and failure to sample all  
 72 parents can lead to biases in RS and RRS. Given that some level of parentage assignment error  
 73 is unavoidable and that in this study not all parents will be sampled, it is important to use an  
 74 unbiased estimate of RS and RRS. Below are the formulas and definitions used to calculate an  
 75 unbiased estimate of RRS ( $RRS_{[unbiased]}$ ) as published by Araki & Blouin (2005). Observed  
 76 mean fitness ( $\hat{W}_x$ ) of a group ‘x’ is a commonly used measure of RS for that group. However,  
 77 there are five potential definitions of RS for individuals in a given group. We will expand on  
 78 these five definitions and their implications in the following section.

79  $RS_{x[unbiased]} = W_x = \left(\frac{1}{1-\hat{a}}\right) \left[\hat{W}_x - \left(\frac{N_{offspring} - N_{assigned}}{N_{parent}}\right) \left(\frac{\hat{b}}{1-\hat{b}}\right)\right]$  (Eq. 13; Araki & Blouin 2005)

$$80 \quad RRS_{x/y[\text{unbiased}]} = \frac{\widehat{W}_x - \left( \frac{N_{\text{offspring}} - N_{\text{assigned}}}{N_{\text{parent}}} \right) \left( \frac{\widehat{b}}{1 - \widehat{b}} \right)}{\widehat{W}_y - \left( \frac{N_{\text{offspring}} - N_{\text{assigned}}}{N_{\text{parent}}} \right) \left( \frac{\widehat{b}}{1 - \widehat{b}} \right)} \quad (\text{Eq. 14; Araki \& Blouin 2005})$$

81 where

82  $\widehat{W}_x$  = observed mean (absolute) fitness of individuals in group x;

83  $\widehat{W}_y$  = observed mean (absolute) fitness of individuals in group y;

84  $N_{\text{offspring}}$  = number of offspring sampled;

85  $N_{\text{assigned}}$  = number of offspring assigned to a parent;

86  $N_{\text{parent}}$  = number of potential parents sampled from groups x and y;

87  $a$  = rate of *failed* parentage assignments (*exclusion*) when *true* parent is sampled; and

88  $b$  = rate of incorrect assignments (i.e. *assignment* to an *untrue* parent).

89 The assignment error rates  $a$  and  $b$  are calculated as

$$90 \quad \widehat{a} = 1 - \frac{(1 - \alpha)P_{\text{assigned}}}{P_{\text{offspring}}} \quad (\text{Eq. 30; Araki \& Blouin 2005})$$

$$91 \quad \widehat{b} = \frac{\alpha P_{\text{assigned}}}{1 - (1 - \alpha)P_{\text{assigned}}} \quad (\text{Eq. 31; Araki \& Blouin 2005})$$

92 where

93  $\alpha$  = threshold probability of incorrect assignments (i.e. *assignment* to an *untrue* parent) used in  
 94 likelihood-based or Bayesian-based parentage assignment to accept/reject a parent-offspring  
 95 pairing

96  $x$  = number of potential parents from group x

97  $y$  = number of potential parents from group y

$$98 \quad P_{\text{assigned}} = \frac{N_{\text{assigned}}}{N_{\text{offspring}}} \quad (\text{Appendix I; Araki \& Blouin 2005})$$

$$99 \quad P_{\text{offspring}} = \frac{xW_x + yW_y}{N_{\text{offspring}}} \quad (\text{Eq. 23; Araki \& Blouin 2005})$$

100 When both  $a = 0$  and  $b = 0$ , then  $RS_{[\text{unbiased}]} = \widehat{W}_x$  and  $RSS_{[\text{unbiased}]} = \frac{RS_x}{RS_y} = \frac{\widehat{W}_x}{\widehat{W}_y}$ . In addition,

101 while  $a$  is usually not known (i.e. it is dependent on  $P_{\text{offspring}}$  which is usually not known),

102 estimation of  $RRS_{[\text{unbiased}]}$  only requires  $b$  (which is independent of  $P_{\text{offspring}}$ ).

103

## Defining RS

104

### *Potential Definitions of RS*

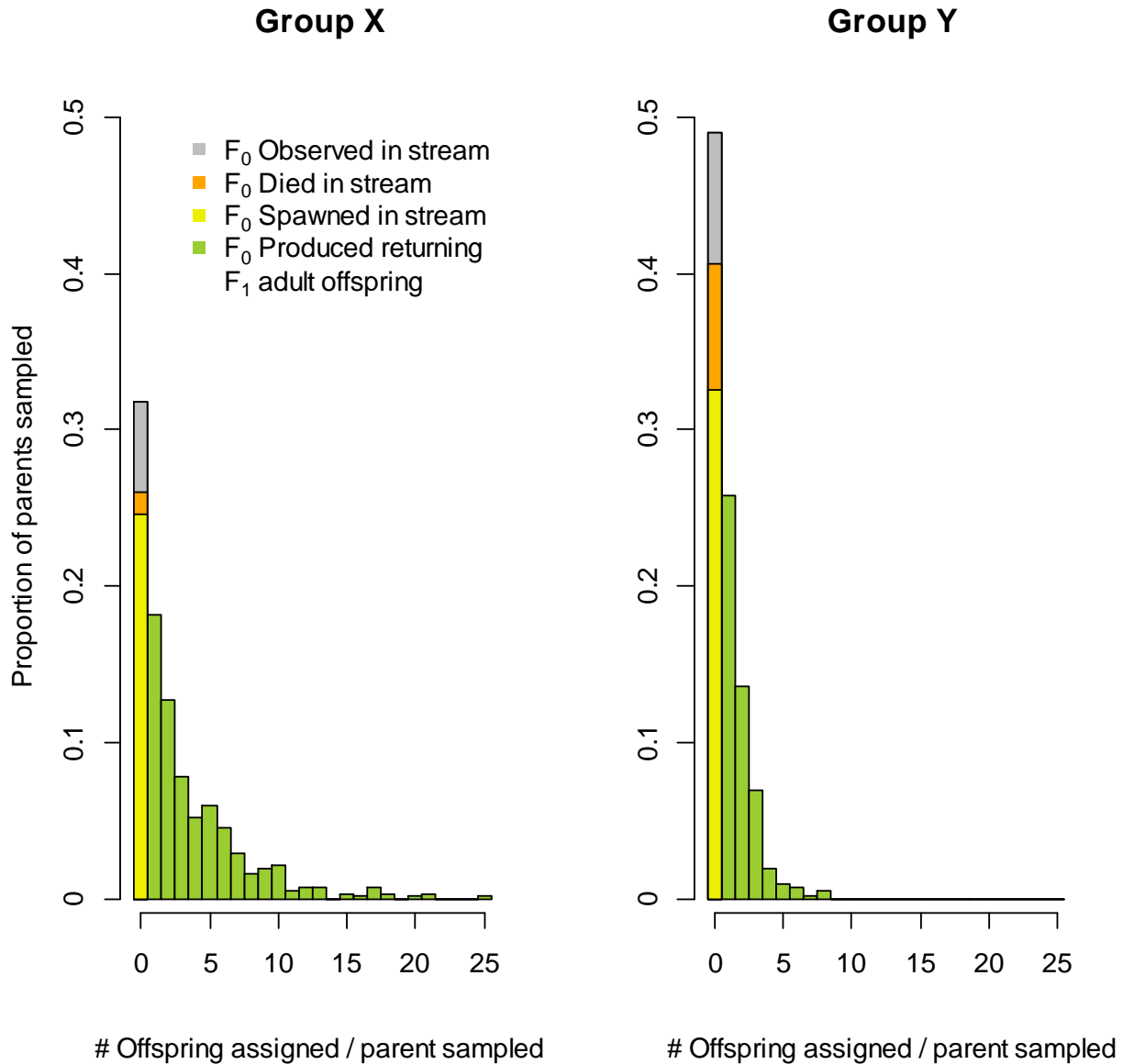
105 The previously described mathematical methods of estimating of  $RRS_{[unbiased]}$  are dependent  
106 upon the definition of both “potential parents” in the denominator and “adult offspring” in the  
107 numerator of RS (Figure 1), where RS is analogous to the observed population mean fitness ( $\widehat{W}_x$   
108 and  $\widehat{W}_y$ ) in the equations from Araki & Blouin (2005). We use a hypothetical example to  
109 illustrate how the definition of “potential parents” in RS affects RRS by comparing recruits-per-  
110 spawner data from two hypothetical groups (Group X and Group Y) under different definitions  
111 of “potential parents” (Figure 2). Figure 2 shows distributions of the proportion of parents that  
112 have a given number of offspring assigned to them after incorporating variation in family size  
113 (note the colors are the same as in Figure 1 with the exception that  $F_0$ 's near the stream mouth  
114 [blue] are not shown). Variation in family sizes of salmon populations can be quite large and  
115 should be considered when assessing genetic variation and short-term population dynamics,  
116 especially given that many families produce zero returning adults (Geiger et al. 2007). To  
117 realistically incorporate variance in family size, we randomly sampled from a negative binomial  
118 distribution based upon the simulation work of Anderson et al. (2011).

119 When “potential parents” are defined as all  $F_0$ 's observed in the stream (Figure 1 & 2), the  
120 proportion of parents that produced zero offspring is represented by the gray bar (including all  
121 colors, except blue in Figure 1). Using this measure of  $F_0$ , the RRS of Group X to Group Y is  
122 calculated as:

123 
$$RS_x = 3.0 \text{ and } RS_y = 1.0, \text{ giving an estimated } RRS = 3.0.$$

124 However, when “potential parents” are defined as only  $F_0$ 's that produced returning  $F_1$  adult  
125 offspring as has been proposed for the AHRG, the proportion of parents that produced zero  
126 offspring is not represented (only green bars). Using this measure of  $F_0$ , the RRS of Group X to  
127 Group Y is calculated as:

128 
$$RS_x = 4.3 \text{ and } RS_y = 1.9, \text{ giving an estimated } RRS = 2.3.$$



129

130 **Figure 2:** Hypothetical data from two groups, X and Y. Note the large difference in the proportion of parents that produced zero  
 131 offspring under different definitions of “potential parents” and the implications this has for calculating RS and estimating RRS.

132 *Example from Literature*

133 Recently Hess et al. (2012) investigated the fitness impacts of a supportive breeding hatchery  
 134 program on a natural population of Chinook salmon in the South Fork Salmon River of the  
 135 Columbia River Basin. This study estimated RSS using the equations from Araki & Blouin  
 136 (2005) based on two definitions of RS (method [i] and [ii]) that differed in how they defined  
 137 “potential parents” (denominator of RS; Figure 1). Method (i), which they reported in  
 138 supplemental information, includes all F<sub>0</sub> parents observed in the stream (e.g., gray in Figure 1 &  
 139 2) regardless of whether they produced any adult progeny in the denominator thus allowing RS  
 140 to vary from  $0 \rightarrow \infty$ ,

141  $RS = \frac{\# F_1 \text{ Returns}}{\# F_0 \text{ Observed in stream}}$  Method (i)

142 Method (ii), which was used in the body of the paper, includes only parents that are known to  
143 have produced adult progeny (i.e. parents that have at least 1 offspring assigned; green in Figure  
144 1 & 2) in the denominator thus restricting RS to vary from  $1 \rightarrow \infty$ ,

145  $RS = \frac{\# F_1 \text{ Returns}}{\# F_0 \text{ Produced returning } F_1 \text{ adult offspring}}$  Method (ii)

146 When Method (ii) was used to calculate RS over two brood years (1998 & 2000) the average  
147  $RRS_{\text{♀}} = 1.11$  and  $RRS_{\text{♂}} = 0.89$  (Table 2 in Hess et al. 2012). However, Method (ii) does *not*  
148 capture differences in the proportion of parents that produce no adult offspring among ancestral  
149 groups (hatchery vs. natural). Method (i) of calculating RS *does* incorporate the difference in  
150 proportion of parents that produce no adult offspring among ancestral origins and resulted in the  
151 average  $RRS_{\text{♀}} = 1.00$  and  $RRS_{\text{♂}} = 0.64$  for same period (Table S3 in Hess et al. 2012). This is  
152 an important consideration given that differences in the mating success among ancestral origins  
153 have been documented in other salmonids (Chinook, Anderson et al. 2013; coho, Thériault et al.  
154 2011). It is also important to note that since the authors had data from every fish that passed a  
155 weir, they were able to present their results in different ways, rather than being forced into a  
156 definition of RS by limitations to the data.

### 157 *Incorporating Relative Reproductive Success into Spawner-Recruit Curves*

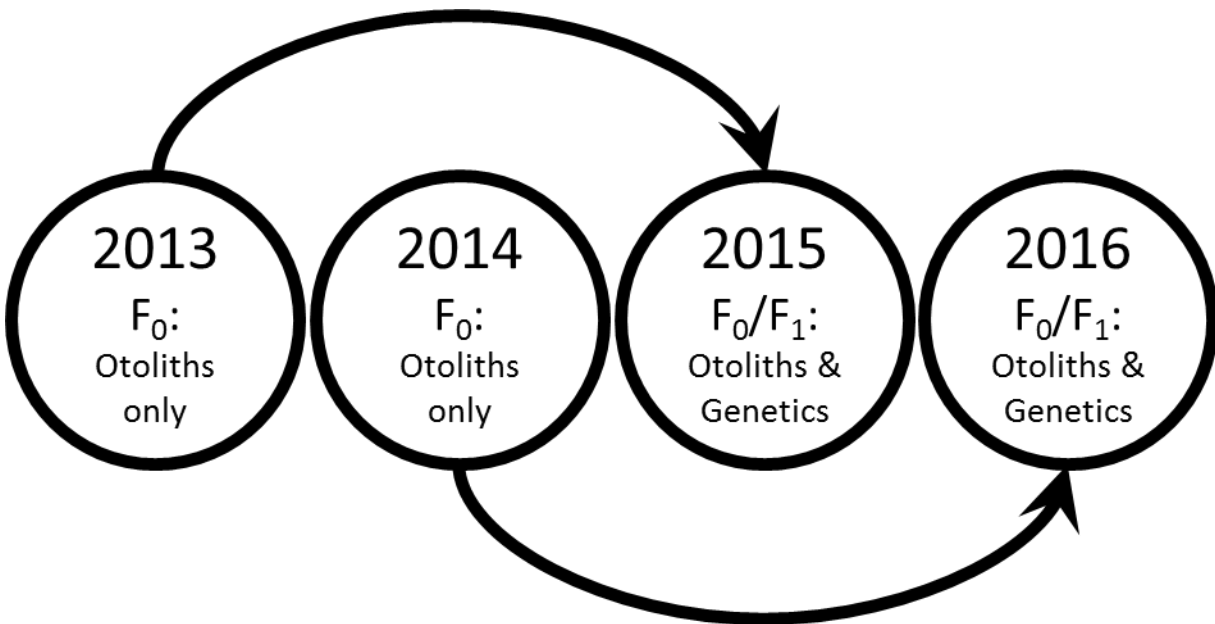
158 Differences in RRS among ancestral origins will influence spawner-recruit curves (and therefore  
159 setting of escapement goals) depending on whether ancestral origin affects reproductive success  
160 and when density-dependent forces occur. For example, if density-dependent forces take place  
161 during spawning (i.e. redd super-imposition) and ancestral origin does not affect redd digging  
162 location, then the number of fish entering the stream is the best measure of that force. On the  
163 other hand, if density-dependent forces occur after hatch (i.e. oxygen levels in the gravel) and  
164 fish of different ancestral origins have different hatch rates, then the best measure of force might  
165 be the number of successfully spawning fish. This is a complex issue that will need to be  
166 addressed by the AHRG in a future technical document.

167 Spawner-recruit relationships are traditionally calculated based on escapement data, because no  
168 information is available on the number of fish successfully contributing to the next generation.  
169 Escapement for pink salmon in PWS is often estimated from aerial stream surveys in which  
170 managers count all fish in-river (alive and dead) and/or milling in the inter-tidal zone at the  
171 mouth of stream. Inclusion of either all live fish in stream or at least all fish that died in a stream  
172 (gray and orange in Figure 1 & 2) is likely more relevant to managers as it most closely puts RS  
173 into the context of escapement by including the largest number of potential parents as counted in  
174 stream surveys.

175 **Implications of Definitions of “Parents” and “Offspring” on Field Sampling**

176 If the inclusion of all fish observed in the stream (gray in Figure 1 & 2) is chosen as the measure  
177 of  $F_0$  for estimating RS (as opposed to other more restrictive measures), this would result in a  
178 tradeoff between sampling a higher proportion of adults that do not contribute to the next  
179 generation (i.e. fish that are ‘nosing in’) and increasing the probability of sampling all  
180 contributing adults.

181 If the inclusion of all fish that died in a stream (orange in Figure 1 & 2) is chosen as the measure  
182 of  $F_0$  for estimating RS, then it is important to include pre-spawn mortalities in the field  
183 sampling, because it will affect the number of potential parents (denominator of RS; Figure 1).  
184 We recommend that pre-spawn mortalities be sampled for otoliths (to determine ancestral origin)  
185 but not genetic tissues for the initial years of the study where only  $F_0$  is being measured (2013-  
186 2014; Figure 3).



187  
188 **Figure 3:** Example sampling strategy for pre-spawn mortalities for pink salmon in PWS.

189 If potential parents are defined as all fish that died in the stream, then knowledge of the ancestral  
190 origin of pre-spawn mortalities is important because they are included as potential parents in the  
191 calculation of RS (Figure 1 & 2). However, to maximize our ability to use genetic data to infer  
192 pedigrees it is not necessary to take a genetic sample from pre-spawn  $F_0$  fish, as they are not  
193 likely to produce any  $F_1$  adult offspring. Yet for years of this study in which  $F_1$ 's are being  
194 sampled (2015-2016), it is important to obtain both otoliths and a genetic sample from pre-spawn  
195 mortalities, as pre-spawn mortalities may include  $F_1$  recruits (Figure 3). It is important to note,  
196 however, that sampling only dead fish will inherently result in a higher probability of missing  
197 contributing adults since it will not be possible to sample every dead fish (i.e. they may wash out,  
198 etc.).



199 Failure to sample pre-spawn mortalities in the field may provide inaccurate estimates of RS in  
 200 the context of escapement data, because managers implicitly assumed them to be potential  
 201 parents when counted in aerial stream surveys of escapement. It is important to recognize that  
 202 pre-spawn mortalities can always be excluded from future analyses if we have the field data, but  
 203 that they will not be able to be included if we do not have the field data. Obtaining field data on  
 204 the greatest number of  $F_0$ 's possible will allow for both the ability to answer a greater range of  
 205 questions **and** increased statistical power to answer those questions. Due to changes in the  
 206 experimental design regarding a reduction in the number of “alevin streams” being sampled,  
 207 there are adequate resources to genotype additional adults.

208 **Summary Table**

<b>Definitions of “potential parent”</b>				
	<i>Present in stream</i>	<i>Died in stream</i>	<i>Spawned in stream</i>	<i>Contributed <math>F_1</math></i>
Pros	Escapement puts RS in a management context. Maximizes probability of sampling contributing fish.	Similar to escapement, incorporates differences in spawning success by origin (H vs. N)	Maximizes efficiency of field sampling for pedigree	Measures family size of spawners that contribute to future generations
Cons	Costs in sampling and genotyping non-contributing parents in the spring of 2014	Reduces probability of sampling contributing fish. Differential field sampling of pre-spawn mortalities (i.e. otoliths only in $F_0$ years, otoliths and genetics in $F_1$ years).	Reduces probability of sampling contributing fish. Does not incorporate differences in spawning success by origin (H vs. N)	Depending on sampling methods, may reduce probability of sampling contributing fish. Does not incorporate differences in spawning success by origin (H vs. N).

209

210 **Other Issues Mentioned, but Not Discussed**

- 211 • Differences in measuring RRS for individual parents vs. cross types (parent pairs)
- 212 • Measuring density-dependent forces
- 213 • Sampling of  $F_1$  recruits in 2015/2016

214 **Questions for the AHRG**

- 215 1. Should we use RRS as defined by Equation 14 of Araki & Blouin (2005)?
- 216 2. How should we define both “potential parents” and “potential offspring” in the equation
- 217 of RS (Figure 1)? Should our calculation of RS attempt to mirror escapement as closely
- 218 as possible?
- 219 3. Should pre-spawn mortalities be included in field sampling, as recommended by the
- 220 ADF&G Gene Conservation Lab?

221

## AHRG Review and Comments

222 *This technical document was discussed at the December 12, 2014 meeting of the AHRG. In*  
223 *addition it was reviewed by email exchange prior to the meeting.*

224 The AHRG did not specifically answer questions 1 and 2, electing to address them at a later date.  
225 It is recommended to sample pre-spawn mortalities.

226 This document is acceptable to the AHRG.

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