## Hylebos Waterway Fish Injury Studies Individual Data and Quality Assurance Results CASE NARRATIVE

### **Juvenile Salmon Injury**

# Cytochrome P4501A - measured as aryl hydrocarbon hydroxylase (AHH) <u>Activity (</u>Tables 1-3)

## **Microsomal Protein Standard Curve Calibration**

Both criteria of the QAP were met for data used in this calibration. Standard curves for each sample set had regression coefficient ( $r^2$ ) of  $\exists 0.990$  and each protein standard run in triplicate had RSD < 20%.

#### **Method Blank**

Each sample set had a mean blank (calculated from 2 sets of duplicates) of 50 dpm <sup>14</sup>C or less meeting the QAP guidelines.

#### **Performance Evaluation**

As stated in the QAP in order to meet an RSD of less than 30% for triplicate analyses of each sample, any outlying replicate was omitted from calculations of final AHH activity. In addition all values were greater than 50 pmol/mg microsomal protein/minute.

### **Juvenile Salmon Injury**

# Analyses of Cytochrome P4501A [measured as aryl hydrocarbon hydroxylase(AHH) Activity]

#### Table 1 Notes

### **Microsomal Protein Standard Curve Calibration**

The Microsome # (column heading abbr. <u>usome #)</u>. Protein sample # (column heading abbr. <u>prot. sample#)</u> and <u>Sample #</u> are internal lab use numbers and for identification only. Column <u>Final Liver Comp#</u> is the official number assigned to each composited sample.

The number under Plate # column is the number assigned to each assay plate.

The concentrations of total protein were calculated using Bovine Albumin Serum (BSA) as the standard.

Column for Protein Standard concentrations (column abbr. <u>Prot. Std. Conc.</u> #) identify each standard concentration dilution used : 2.0, 4.0, 8.0, 16.0, 20.0, and 30.0 mg/ml, with all standards loaded on plate as triplicates (e.g. 2.0-1, 2.0-2, 2.0-3 etc).

The <u>Prot. Std OD value</u> column is the absorbance of the sample measured as Optical Density (OD) at a wavelength of 620nm.

Percent coefficient of variation (column <u>OD %CV</u> is standard deviation (SD) of each triplicate analyses (reported in <u>Prot, Std OD value)</u> divided by mean of each triplicate, multiplied by 100.

Correlation Coefficient (column abbr. <u>Corr. Coeff.</u>) is generated from linear standard curve fit of mean CID values of protein standards.

TABLE 1: Protein Standard Curve Calibration

						Drotoin C	Drotein Standard Curve	NA.	
# emosn	Sample #	Final		Prot.	mg Prot.	Prot. Std.	Prot.Std	00	Corr. Coeff.
		Liver Comp#	Plate #	Sample #	/m1	Conc.#	OD value	%CV (<20%)	(>0.95)
95012401	94 Chum PTH-C1	-	950124	A01	17.1	2.0-1	0.230		0.993
95012402	94 Chum PTH-C2	8		A02	15.0	2.0-2	0.230		
95012403	94 Chum PTH-C3	ო		A03	16.1	2.0-3	0.230	0.731	
95012404	94 Chum PTH-C4	4		A04	17.7	4.0-1	0.242		
95012405	94 Chum SKO-C1	S.		A05	12.5	4.0-2	0.230		
95012406	94 Chum SKO-C2	9		A06	14.3	4.0-3	0.245	3.317	
95012407	94 Chum HYL(5/25)-C1	7		A07	12.1	8.0-1	0.284		
95012408	94 Chum HYL(5/25)-C2	<b>©</b>		A08	16.3	8.0-2	0.282		
95012409	94 Chum HYL(5/25)-C3	o		A09	14.4	8.0-3	0.278	1.086	
95012410	94 Chum HYL(6/1)-C1	10		A10	11.4	16.0-1	0.420		
95012411	94 Chum HYL(6/1)-C2	-		A11	13.8	16.0-2	0.431		
95012412	94 Chinook HYL(6/1)-C3	12		A12	11.9	16.0-3	0.435	1.81	
95012413	94 Chum HYL(6/2)-C4	13		A13	10.9	20.0-1	0.477		
95012414	94 Chum HYL(6/2)-C5	14		A14	11.0	20.0-2	0.473		
95012415	94 Chinook HYL(6/2)-C6	1.5		A15	14.1	20.0-3	0.448	3.37	
95012416	94 Chum HYL(6/8)-C1	10		A16	12.1	30.0-1	0.546		
95012417	94 Chum HYL(6/8)-C2	17		A17	13.8	30.0-2	0.580		
95012418	94 Chinook HYL(6/8)-C3	8 -		A18	9.8	30.0-3	0.604	5.051	
95012419	94 Chum HYL(6/9)-C4	6 1		A19	13.9				
95012420	94 Chum HYL(6/9)-C5	20		A20	14.1				
95012421	94 Chinook PSH(6/13)-C1	21		A21	13.6				Assay date
95012422	94 Chinook PSH(6/13)-C2	22		A22	14.4				1/26/95
95012423	94 Chinook PSH(6/13)-C3	23	े. .व	A23	14.8				
95012424	94 Chinook PSH(6/13)-C4	24		A24	12.5			33	
95012501	94 Chinook PSH(6/13)-C5	25	950125	A01	13.7	2.0-1	0.236		0.991
95012502	94 Chum HYL(6/15)-C1	56		A02	18.9	2.0-2	0.236		
95012503	94 Chinook HYL(6/15)-C2	27		A03	13.3	2.0-3	0.239	0.731	
95012504	94 Chinook HYL(6/16)-C3	28		A04	13.4	4.0-1	0.242		
95012505	94 Chinook HYL(6/22)-C1	29		A05	18.4	4.0-2	0.245		
95012506	94 Chinook HYL(6/23)-C2	30		90V	15.1	4.0-3	0.244	0.628	
95012507	94 Chinook HYL(6/29)-C1	31		A07	14.4	8.0-1	0.293		

Hylebos/NRDA Juvenile Flatfish Survey

TABLE 1: Protein Standard Curve Calibration

						Protein S	Protein Standard Curve	Ve	. 22
# emosn	Sample #	Final		Prot.	mg Prot.	mg Prot. Prot. Std.	Prot.Std	00	Corr. Coeff.
		Liver Comp#	Plate #	Sample #	m/	Conc.#	OD value	%CV (<20%)	(>0.95)
95012508	95012508 94 Chinook NisH(5/9)-C1	32		A08	18.2	8.28.0-2	0.295		
95012509	95012509 94 Chinook NisH(5/13)-C2	33		A09	18.5	8.58.0-3	0.289	1.045	•
95012510	94 Chinook NisH(5/13)-C3	34		A10	17.8	7.8 16.0-1	0.419	•	
95012511	94 Chinook NisE(5/18)-C1	35		A11	18.8	18.8 16.0-2	0.437		
95012512	94 Chinook NisE(5/20)-C2	36		A12	21.1	16.0-3	0.445	3.073	
95012513	94 Chinook NisE(5/31)-C3	37		A13	16.3	16.3 20.0-1	0.461		
						20.0-2	0.463		
						20.0-3	0.482	2.474	
				-		30.0-1	0.533		
					-	30.0-2	0.569		Assay date
						30.0-3	0.581	4.455	1/26/95

### **Juvenile Salmon Injury**

# Analyses of Cytochrome P4501A [measured as aryl hydrocarbon hydroxylase(AHH) Activity]

#### **Table 2 Notes**

## **Method Blank - AHH Assay**

The numbers under columns Microsome # (column abbr. <u>usome #)</u>, <u>AHH # and Sample #</u> are internal lab use numbers and for identification only. Column <u>Final Liver Comp#</u> is the official number assigned to each composited sample.

The <u>AHH Assay Date</u> column is the day sample set was run. The column <u>AHH Set Blk (µome #)</u> is the sample run as the blank for each set.

<u>Substrate blank</u> is reaction mixture without substrate ( <sup>14</sup>C-Benzo[a]Pyrene ) added. <u>Boiled blank</u> is reaction mixture with an aliquot of boiled microsome sample. Both blank sets run as duplicates (i.e. DPM1 and DPM2). All 4 blanks are then averaged to give <u>AHH Assay Set Blk Avg DPM</u>.

Table 2 : Method Blank - Aryl hydrocarbon hydroxlase (AHH) Assay

АНН									
Assay	# emosn	AHH #	Sample #	Final	AHH Set BIK	Substrate	Substrate	<b>Boiled BIK</b>	<b>Boiled BIK</b>
Date				Liver Comp#	(#ewosn)	BIK (DPM1)	BIK (DPM2)	(DPM1)	(DPM2)
1/31/95	95012401 1	1/31/95-1	94 Chum PTH-C1	1	95012401	47	48	41	50
	95012402 1	1/31/95-2	94 Chum PTH-C2	7					
	950124031	1/31/95-3	94 Chum PTH-C3	ო					
	95012404 1	1/31/95-4	94 Chum PTH-C4	4					,
	95012405 1	1/31/95-5	94 Chum SKO-C1	Ŋ		AHH As	<b>AHH Assay Set BIk</b>		
	95012406 1	1/31/95-6	94 Chum SKO-C2	ဖ		Avg DPM	46		
	95012407 1	1/31/95-7	94 Chum HYL(5/25)-C1	_					
	95012408 1	1/31/95-8	94 Chum HYL(5/25)-C2	<b>&amp;</b>					
	95012409 1	/31/95-9	94 Chum HYL(5/25)-C3	6					
	95012410 1/	/31/95-10	94 Chum HYL(6/1)-C1	10					
	95012411 1/	/31/95-11	94 Chum HYL(6/1)-C2	-					
	95012412 1	/31/95-12	94 Chinook HYL(6/1)-C3	12					
	95012413 1/	/31/95-13	94 Chum HYL(6/2)-C4	13					
	95012414 1/	/31/95-14	94 Chum HYL(6/2)-C5	4 +					
	151	/31/95-15	94 Chinook HYL(6/2)-C6	T 22					
	16 1	/31/95-16	94 Chum HYL(6/8)-C1	16					
	171		94 Chum HYL(6/8)-C2	17					
	4181	/31/95-18	94 Chinook HYL(6/8)-C3	18					
	_	/31/95-19	94 Chum HYL(6/9)-C4	9					
	95012420 1/	/31/95-20	94 Chum HYL(6/9)-C5	20					
	_	/31/95-21	94 Chinook PSH(6/13)-C1	21					
	┰.	/31/95-22		22					
	95012423 1/	/31/95-23	94 Chinook PSH(6/13)-C3	23					
	95012424 1/	1/31/95-24	94 Chinook PSH(6/13)-C4	2.4					
2/1/95	95012501 2/01/95-1	/01/95-1	94 Chinook PSH(6/13)-C5	25	95012501	45	48	49	53
	95012502 2/	2/01/95-2	94 Chum HYL(6/15)-C1	26					
	95012503 2/	2/01/95-3	94 Chinook HYL(6/15)-C2	27					
	95012504 2/	2/01/95-4	94 Chinook HYL(6/16)-C3	28					
	95012505 2/	2/01/95-5	94 Chinook HYL(6/22)-C1	29	•	AHH AS	<b>AHH Assay Set BIk</b>		
	95012506 2/	2/01/95-6	94 Chinook HYL(6/23)-C2	30		<b>Avg DPM</b>	49		
	95012507 2/	2/01/95-7	94 Chinook HYL(6/29)-C1	31					

Table 2: Method Blank - Aryl hydrocarbon hydroxlase (AHH) Assay Hylebos/NRDA Juvenile Salmon Survey

Boiled BIK (DPM2)						
Boiled Blk (DPM1)						
Substrate Bik (DPM2)						
Substrate Substrate BIk (DPM1) BIk (DPM2)						
AHH Set BIK Substrate (µsome#) BIk (DPM1)						
Final Liver Comp#	32	33	34	35	36	37
Sample #	94 Chinook NisH(5/9)-C1	94 Chinook NisH(5/13)-C2	94 Chinook NisH(5/13)-C3	94 Chinook NisE(5/18)-C1	94 Chinook NisE(5/20)-C2	94 Chinook NisE(5/31)-C3
<b>АНН</b> #	/01/95-8	/01/95-9	/01/95-10	/01/95-11	/01/95-12	/01/95-13
msome #	95012508 2/01/95-8	95012509 2/01/95-9	95012510 2/01/95-10	95012511 2/01/95-11	95012512 2/01/95-12	95012513 2/01/95-13
AHH Assay Date						

## **Juvenile Salmon Injury**

# Analyses of Cytochrome P4501A [measured as aryl hydrocarbon hydroxylase(AHH) Activity]

#### Table 3 Notes

## **Performance Evaluation Final - AHH Activity**

Microsome # (column abbr. <u>usome #)</u>, <u>AHH # and Sample # are internal lab use numbers and for identification only. Column <u>Final Liver Comp#</u> is the official number assigned to each composited sample.</u>

The <u>AHH Assay Date</u> column is the day sample set was run. The <u>AHH Set Blk (µsome #)</u> is the sample run as the blank for each set.

Samples are run in triplicate. The percent coefficient of variation (%CV of DPM column) is the standard deviation (value reported in column **DPM SD**) of each triplicate (reported in columns :AHH DPM1, AHH PM2, AHH DPM3) divided by mean of each triplicate (column AHH DPM Avg) multiplied by 100.

Triplicate sample outlier indicated by •, was not used in final calculation to meet criteria of %CV < 30%.

The column <u>Avg AHH minus set Blk</u>, is <u>AHH DPM Avg</u>, minus the value calculated in **Table 2** <u>AHH set Blk</u>) for each set.

Final AHH Activity is normalized for total microsomal prote in i.e. "Avg AHH minus set Blk" divided by "mg prot/ ml' (from **Table 1**).

Final units for AHH activity are picomoles BaP metabolized per minute per mg microsomal protein.

Table 3: Performance Evaluation Final - Aryl hydrocarbon hydroxiase (AHH) Activity Calculations Hylebos/NRDA Juvenile Salmon Survey

			Final		-				AHH		%CV of	Avg AHH	Final
Assay	usome # AHH #	Sample #	Liver	AHH Set mg prot. AHH	mg prot.	AHH	AHH	AHH	DPM	DPM	DPM	minus	AHH
Date			Comp#	BK	lm/	DPM1	DPM2	<b>DPM3</b>	Avg	S	(<30%)	Set Bik	Activity
1/31/95	95012401 1/31/95-1	94 Chum PTH-C1	-	46	17.1	66	104	86	96	6	10	50	20
	95012402 1/31/95-2	94 Chum PTH-C2	N		15.0	113	121	124	119	9	ιΩ	73	33
	95012403 1/31/95-3	94 Chum PTH-C3	က		16.1	160	150	136	149	12	•	103	43
•	95012404 1/31/95-4	94 Chum PTH-C4	4		17.7	115	137	124	125	Ξ	6	4	30
	95012405 1/31/95-5	94 Chum SKO-C1	r0		12.5	83	94	78	85	œ	10	39	21
	95012406 1/31/95-6	94 Chum SKO-C2	9		14.3	87	108	104	100	=	11	54	26
	95012407 1/31/95-7	94 Chum HYL(5/25)-C1	7		12.1	178	175	152	168	14	80	122	69
	95012408 1/31/95-8	94 Chum HYL(5/25)-C2	80	*	16.3	343	351	309	334	22	7	288	120
	95012409 1/31/95-9	94 Chum HYL(5/25)-C3	o		14.4	202	226	246	225	22	10	179	84
		94 Chum HYL(6/1)-C1	10		11.4	167	165	147	160	=	7	114	68
	95012411 1/31/95-11	94 Chum HYL(6/1)-C2	1		13.8	172	222	221	205	50	14	159	7.8
	95012412 1/31/95-12	94 Chinook HYL(6/1)-C3	12		11.9	213	222	180	205	22	-	159	91
	95012413 1/31/95-13	94 Chum HYL(6/2)-C4	13		10.9	148	154	120	141	18	13	92	29
	95012414 1/31/95-14	94 Chum HYL(6/2)-C5	14		11.0	91	102	88	94	7	80	48	29
	95012415 1/31/95-15	94 Chinook HYL(6/2)-C6	15		14.1	222	229	265	239	23	10	193	8
	95012416 1/31/95-16	94 Chum HYL(6/8)-C1	16		12.1	129	150	- 35	140	15	11	94	53
	95012417 1/31/95-17	94 Chum HYL(6/8)-C2	17		13.8	225	246	231	234	=	S	188	93
	95012418 1/31/95-18	94 Chinook HYL(6/8)-C3	18		9.8	107	113	06	103	12	12	57	40
	95012419 1/31/95-19	94 Chum HYL(6/9)-C4	6		13.9	256	287	299	281	22	<b>c</b>	235	115
	95012420 1/31/95-20		20		14.1	306	285	269	287	19	9	241	116
	95012421 1/31/95-21		21		13.6	94	92	101	97	4	4	51	25
	95012422 1/31/95-22	94 Chinook PSH(6/13)-C2	22		14.4	•164	92	75	84	12	14	38	18
	95012423 1/31/95-23	94 Chinook PSH(6/13)-C3	23		14.8	112	113	111	112	-	-	99	30
	95012424 1/31/95-24	94 Chinook PSH(6/13)-C4	24		12.5	87	85	81	84	က	4	38	21
2/1/95	95012501 2/01/95-1	94 Chinook PSH(6/13)-C5	25	49	13.7	66	112	104	105	7	9	59	29
	95012502 2/01/95-2	94 Chum HYL(6/15)-C1	26		18.9	573	528	488	530	43	80	484	174
	95012503 2/01/95-3	94 Chinook HYL(6/15)-C2	27		13.3	257	232	221	237	18	80	191	86
	95012504 2/01/95-4	94 Chinook HYL(6/16)-C3	28		13.4	324	318	277	306	56	80	260	132
	95012505 2/01/95-5	94 Chinook HYL(6/22)-C1	29		18.4	527	521	424	491	28	12	445	164
	95012506 2/01/95-6	94 Chinook HYL(6/23)-C2	30		15.1	341	288	324	318	27	6	272	122
					•	) Oulti	er of t	Oultier of triplicate	•				

Table 3: Performance Evaluation Final - Aryl hydrocarbon hydroxlase (AHH) Activity Calculations Hylebos/NRDA Juvenile Salmon Survey

				Final						AHH		%CV of	Avg AHH	Final
Assay	msome #	AHH *	Sample #	Liver	<b>AHH Set</b>	AHH Set mg prot. AHH	AHH	AHH	AHH	DPM	DPM	DPM	minus	AHH
Date				Comp#	器	lm/	DPM1	DPM2	<b>DPM3</b>	Avg	S	(%06>)	Set BIK	Activity
	95012507 2/01/95-7	7-26/10/	94 Chinook HYL(6/29)-C1	31	1	14.4	141	141 116 124	124	127	13	10	81	38
	95012508 2/01/95-8	1/01/95-8	94 Chinook NisH(5/9)-C1	32		18.2	415	409	380	401	19	S	355	133
	95012509 2/01/95-9	101/95-9	94 Chinook NisH(5/13)-C2	33		18.5	343	342	320	335	13	4	286	105
	95012510 2/01/95-10	//01/95-10	94 Chinook NisH(5/13)-C3	34		17.8	415		361	399	33	00	350	134
	95012511 2/01/95-11	/01/95-11	94 Chinook NisE(5/18)-C1	3		18.8	277		230	265	30	11	216	78
	95012512 2/01/95-12	/01/95-12	94 Chinook NisE(5/20)-C2	36		21.1	307		261	293	27	<b>o</b>	244	79
	95012513 2/01/95-13	/01/95-13	94 Chinook NisE(5/31)-C3	37		16.3	181	185	181	182	8	-	133	26
							• Oulti	er of t	<b>Jultier of triplicate</b>	m				