

# Attachment 1: Chain of Custody Forms

## SWAMP REQUEST FOR ANALYSIS AND CHAIN OF CUSTODY (COC) RECORD

<b>Fiscal Year:</b> 11	<b>Project ID:</b> 11SWSBG1	<b>Contact Person:</b> Autumn Bonnema	
<b>Region:</b>	<b>Season:</b> summer	<b>Phone:</b> 831-771-4175	
<b>Field Crew:</b>	<b>Date:</b>	<b>email:</b> <a href="mailto:bonnema@miml.calstate.edu">bonnema@miml.calstate.edu</a>	
		<b>Mailing Address:</b> 7544 Sandholdt Rd. Moss Landing, CA 95039	

StationCode	Station Name	LabID	Sample Date	Tissue THg	Tissue Archive			# of Containers Plastic Bag	Preservation Frozen
			<b>TOTAL</b>	0	0	0	0	0	0

**Comments:**

<b>Samples Relinquished by:</b>		<b>Samples Received by:</b>	
Name (Print and Sign)	Date	Name (Print and Sign)	Date


# Chain of Custody Record



Forest and Rangeland Ecosystem Science Center  
 Contaminant Ecology Program  
 Collin Eagles-Smith  
 3200 SW Jefferson  
 Corvallis, OR 97330

Transferring to: Name: Address:  Phone: Email:	Project:  Persons performing analysis:	
Sample ID(s)	Sample Description	Analysis Requested
Relinquished By: _____ <small>Print</small> <small>Sign</small>		Date / Time: _____
Received By: _____ <small>Print</small> <small>Sign</small>		Date / Time: _____
Relinquished By: _____ <small>Print</small> <small>Sign</small>		Date / Time: _____
Received By: _____ <small>Print</small> <small>Sign</small>		Date / Time: _____

## Attachment 2: Field Data Sheets

SWAMP Tissue Sampling - Non-Trawl (Event Type = T1) SWB WildLk 2012				Entered in d-base (initial/date)		Pg of Pgs	
*StationCode: _____		*StationName: _____		*Purpose Failure Code: _____		Agency	
*FundingCode: 1 1 S W B G 0 1		*Date (mm/dd/yyyy): / /		WIND DIRECTION (from):		PHOTOS (RB & LB assigned when facing downstream; RENAME to StationCode_yyyy_mm_dd_uniquecode)	
*Sampling Crew:		Arrival Time:	BEAUFORT SCALE (see attachment):			1: (RB / LB / BB / US / DS / ##)	
		Departure Time:					
DOMINANT SUBSTRATE: Concrete, Cobble, Gravel, Sand, Mud, Other _____, unk			WATERCOLOR: Colorless, Green, Yellow, Brown			2: (RB / LB / BB / US / DS / ##)	
WATER CLARITY: Clear (see bottom), Cloudy (>4" vis), Murky (<4" vis)			OTHER PRESENCE: Vascular, Nonvascular, Oily Sheen, Foam, Trash, O				
Comments:						3: (RB / LB / BB / US / DS / ##)	
<b>Tissue Collection</b>							
COLLECTION DEVICE: RV _____ Masta-Blasta, Big E, Sparky _____, Backpack Model _____, Net (length & mesh) _____							
Target:	Lat (dd.dddd)		Long (dd.dddd)		-		
GPS Model:			Datum: NAD83 WGS84 Other _____		*GPS / DGPS		Elevation (ft):
Location	*Depth (m):	Distance from Bank (m):		Accuracy (ft / m)	Latitude (dd.dddd)	Longitude (-ddd.dddd)	Depth (m)
COLLECTION METHOD:	E-boat, Backpack shocker, Fyke net, gill net, seine, hook & line		Start Time	Coord. 1			
SAMPLE LOCATION:	Bank, Thalweg, Midchannel, Open Water, NA			Coord. 2			
HYDROMODIFICATION:	None, Bridge, Pipes, Concrete Channel, Grade Control, Culvert,		End Time	Coord. 3			
HYDROMODLOC(to sample):	US / DS / NA / WI	Other _____	Geoshape: Line Poly Point		Coord. 4		
Location	*Depth (m):	Distance from Bank (m):			Latitude (dd.dddd)	Longitude (-ddd.dddd)	Depth (m)
COLLECTION METHOD:	E-boat, Backpack shocker, Fyke net, gill net, seine, hook & line		Start Time	Coord. 1			
SAMPLE LOCATION:	Bank, Thalweg, Midchannel, Open Water, NA			Coord. 2			
HYDROMODIFICATION:	None, Bridge, Pipes, Concrete Channel, Grade Control, Culvert,		End Time	Coord. 3			
HYDROMODLOC(to sample):	US / DS / NA / WI	Other _____	Geoshape: Line Poly Point		Coord. 4		
Location	*Depth (m):	Distance from Bank (m):			Latitude (dd.dddd)	Longitude (-ddd.dddd)	Depth (m)
COLLECTION METHOD:	E-boat, Backpack shocker, Fyke net, gill net, seine, hook & line		Start Time	Coord. 1			
SAMPLE LOCATION:	Bank, Thalweg, Midchannel, Open Water, NA			Coord. 2			
HYDROMODIFICATION:	None, Bridge, Pipes, Concrete Channel, Grade Control, Culvert,		End Time	Coord. 3			
HYDROMODLOC(to sample):	US / DS / NA / WI	Other _____	Geoshape: Line Poly Point		Coord. 4		
Failure Codes: Dry (no water), Instrument Failure, No Access, Non-sampleable, Pre-abandoned, Other							
Comments:							

SWAMP Tissue Sampling - Non-Trawl (Event Type = TI) SWB_WildLk_2012					Entered in d-base (initial/date)		Pg	of	Pgs
*StationCode: _____			*StationName: _____				*Purpose Failure Code:	Agency	
*FundingCode: 1 1 S W B G 0 1			*Date (mm/dd/yyyy): / /						
<b>Tissue Collection</b>									
Location	*Depth (m):	Distance from Bank (m):		Accuracy (ft / m)	Latitude (dd.ddddd)	Longitude (-ddd.ddddd)	Depth (m)		
COLLECTION METHOD:	E-boat, Backpack shocker, Fyke net, gill net, seine, hook & line			Start Time	Coord. 1				
SAMPLE LOCATION:	Bank, Thalweg, Midchannel, Open Water, NA				Coord. 2				
HYDROMODIFICATION:	None, Bridge, Pipes, Concrete Channel, Grade Control, Culvert,			End Time	Coord. 3				
HYDROMODLOC(to sample):	US / DS / NA / WI	Other _____ Geoshape: Line Poly Point			Coord. 4				
Location	*Depth (m):	Distance from Bank (m):			Latitude (dd.ddddd)	Longitude (-ddd.ddddd)	Depth (m)		
COLLECTION METHOD:	E-boat, Backpack shocker, Fyke net, gill net, seine, hook & line			Start Time	Coord. 1				
SAMPLE LOCATION:	Bank, Thalweg, Midchannel, Open Water, NA				Coord. 2				
HYDROMODIFICATION:	None, Bridge, Pipes, Concrete Channel, Grade Control, Culvert,			End Time	Coord. 3				
HYDROMODLOC(to sample):	US / DS / NA / WI	Other _____ Geoshape: Line Poly Point			Coord. 4				
Location	*Depth (m):	Distance from Bank (m):			Latitude (dd.ddddd)	Longitude (-ddd.ddddd)	Depth (m)		
COLLECTION METHOD:	E-boat, Backpack shocker, Fyke net, gill net, seine, hook & line			Start Time	Coord. 1				
SAMPLE LOCATION:	Bank, Thalweg, Midchannel, Open Water, NA				Coord. 2				
HYDROMODIFICATION:	None, Bridge, Pipes, Concrete Channel, Grade Control, Culvert,			End Time	Coord. 3				
HYDROMODLOC(to sample):	US / DS / NA / WI	Other _____ Geoshape: Line Poly Point			Coord. 4				
Location	*Depth (m):	Distance from Bank (m):			Latitude (dd.ddddd)	Longitude (-ddd.ddddd)	Depth (m)		
COLLECTION METHOD:	E-boat, Backpack shocker, Fyke net, gill net, seine, hook & line			Start Time	Coord. 1				
SAMPLE LOCATION:	Bank, Thalweg, Midchannel, Open Water, NA				Coord. 2				
HYDROMODIFICATION:	None, Bridge, Pipes, Concrete Channel, Grade Control, Culvert,			End Time	Coord. 3				
HYDROMODLOC(to sample):	US / DS / NA / WI	Other _____ Geoshape: Line Poly Point			Coord. 4				
Location	*Depth (m):	Distance from Bank (m):			Latitude (dd.ddddd)	Longitude (-ddd.ddddd)	Depth (m)		
COLLECTION METHOD:	E-boat, Backpack shocker, Fyke net, gill net, seine, hook & line			Start Time	Coord. 1				
SAMPLE LOCATION:	Bank, Thalweg, Midchannel, Open Water, NA				Coord. 2				
HYDROMODIFICATION:	None, Bridge, Pipes, Concrete Channel, Grade Control, Culvert,			End Time	Coord. 3				
HYDROMODLOC(to sample):	US / DS / NA / WI	Other _____ Geoshape: Line Poly Point			Coord. 4				
Location	*Depth (m):	Distance from Bank (m):			Latitude (dd.ddddd)	Longitude (-ddd.ddddd)	Depth (m)		
COLLECTION METHOD:	E-boat, Backpack shocker, Fyke net, gill net, seine, hook & line			Start Time	Coord. 1				
SAMPLE LOCATION:	Bank, Thalweg, Midchannel, Open Water, NA				Coord. 2				
HYDROMODIFICATION:	None, Bridge, Pipes, Concrete Channel, Grade Control, Culvert,			End Time	Coord. 3				
HYDROMODLOC(to sample):	US / DS / NA / WI	Other _____ Geoshape: Line Poly Point			Coord. 4				
Location	*Depth (m):	Distance from Bank (m):			Latitude (dd.ddddd)	Longitude (-ddd.ddddd)	Depth (m)		
COLLECTION METHOD:	E-boat, Backpack shocker, Fyke net, gill net, seine, hook & line			Start Time	Coord. 1				
SAMPLE LOCATION:	Bank, Thalweg, Midchannel, Open Water, NA				Coord. 2				
HYDROMODIFICATION:	None, Bridge, Pipes, Concrete Channel, Grade Control, Culvert,			End Time	Coord. 3				
HYDROMODLOC(to sample):	US / DS / NA / WI	Other _____ Geoshape: Line Poly Point			Coord. 4				
Failure Codes: Dry (no water), Instrument Failure, No Access, Non-sampleable, Pre-abandoned, Other									
Comments:									

SWAMP Tissue Sampling - Fish Abundance SWB_WildLk_2012				Entered in d-base (initial/date)				Pg:      of      Pgs				
*StationCode: _____			StationName: _____			Date (mm/dd/yyyy):     /     /						
Location #	Organism ID	Tag #	Species Name/Code	FL (mm)	TL (mm)	StdL (mm)	Weight (g)	Count	Count Est.	Sex	Anomaly	Condition
										M F U L		
										M F U L		
										M F U L		
										M F U L		
										M F U L		
										M F U L		
										M F U L		
										M F U L		
										M F U L		
										M F U L		
										M F U L		
										M F U L		
										M F U L		
										M F U L		
										M F U L		
										M F U L		
										M F U L		
<b>Location #:</b> Match fish with Location # from Tissue Collection sheet				<b>Organism ID:</b> Combine composite # and fish # (e.g., fish 1 of composite WC01 is WC01-01) to be un( Tag #: Use if applicable								
<b>Species Code:</b> Largemouth Bass (LMB), Smallmouth Bass (SMB), Spotted Bass (SPB), Sacramento Pike Minnow (SPM), Rainbow Trout (RT), Brown Trout (BT), Brook Trout (BKT), White Catfish (WC), Carp (CAR), Channel Catfish (CC), Brown Bullhead (BRB), Sacramento Sucker (SS), Redear (RES), Black Crappie (CRP), Bluegill (BG), Tilapia (TIL), Green Sunfish (GRS), Kokanee (KOK)												
<b>Stage:</b> Adult (A), Juvenile (J), Subadult (SA), Not Recorded (NR)				<b>Count Est:</b> If appropriate, add < or > if count is €								
<b>Anomalies:</b> Ambicoloration (A), Albinism (B), Cloudiness (CL), Deformity-skeletal (D), Discoloration (DC), Depression (DS), Fin Erosion (F), Gill Erosion (T), Hemorrhage (H), Lesion (L), Parasite (P), Popeye (PE), Tumor (T), Ulceration (U), White Spots (W), and any combination												
				<b>Sex:</b> unk(U), taken at Lab(L) <b>BodyLocation:</b> Branchial Chamber(BRC), Buccal Cavity(BC), Eyes(E), Musculoskeleton(M), Skin/Fins(SF)								
<b>Comments:</b> Mark fish requiring further ID; SEPARATE FISH BY LOCATION AND INDICATE LOCATION # ON LABEL												

<u>Fish Species</u>	<u>Species Code</u>	<u>Fish Species</u>	<u>Species Code</u>
American shad	AMS	Sacramento sucker	SAS
black crappie	BCR	Sacramento blackfish	SBF
bluegill	BGL	sculpin ssp	SCP
black bullhead	BLB	shiner perch	SHP
blue catfish	BLC	smallmouth bass	SMB
brown trout	BNT	spotted bass	SPB
brown bullhead	BRB	Sacramento perch	SPH
brook trout	BRT	Sacramento pike minnow	SPM
carp, common	CAR	Sacramento splittail	SST
channel catfish	CHC	striped bass	STB
chinook salmon	CHS	steelhead	STH
coho salmon	COH	striped mullet	STM
California roach	CRH	sturgeon, white	WST
delta smelt	DTS	threadfin shad	TFS
flathead catfish	FHC	tilapia ssp	TIL
fathead minnows	FHM	topsmelt	TPS
goldfish	GLF	threespine stickleback	TSS
golden trout	GLT	tui chub	TUC
green sunfish	GRS	tule perch	TUP
hitch	HIT	warmouth	WAR
hardhead	HRH	white crappie	WCR
inland silverside	ISS	white catfish	WHC
killifish	KIL		
kokanee salmon	KOK		
lamprey	LAM		
longfin smelt	LFS		
lake trout	LKT		
largemouth bass	LMB		
mosquitofish	MQF		
pumpkinseed sunfish	PKS		
rainbow trout	RBT		
redestye bass	REB		
redestye sunfish	RES		
red shiner	RSR		

2010 USGS SBSP Waterbird Hg Project

**Egg Collection/Salvage Sheet**

ID Code	Date (M/D/Y)	Pond	Nest Number	UTM Easting	UTM Northing	Species	Life Stage	Approx. age	condition when found	Collection Status	Notes

**Egg Collection Status**

NS	Dead	Rolled
FTH	Cracked	Abandoned (Researcher)
FTH-drilled	Flooded	Abandoned (partial dep)

**Life Stage**

Egg	Adult
Chick	
Juvenile	

Abandoned (Natural)
Abandoned (foam)

# Attachment 3: Analysis Authorization Forms

Analysis Authorization  
 Fiscal Year: 1112  
 Region:

Project ID: SWB\_WildLk\_2012  
 Season:  
 Date:

Contact Person: Autumn Bonnema  
 Phone: 831-771-4175  
 email: bonnema@mml.calstate.edu  
 Mailing Address:

Station	StationName	OrganismName	CompositeIDText	Dissect and Analyze					Archive
				Tissue	Flesh	Ind Hg	Comp Hg	%Moisture	
Total				0	0	0	0	0	0



## Attachment 4: Laboratory Data Sheets

SWAMP Lab Data Sheet - FISH											
ProjectID: SWB_WildLk_2012				PrepPres: Skin ON; Scales ON				LabID:		Pg: 1 of 2 Pgs	
StationCode:				Tissue: Whole Body				Entered d-base (initial/date)			
StationName:				Homog. Method: USGS SOP Mortar/Pestle				Staff. Diss. Homog.			
Species Name:				Date Diss. (mm/dd/yyyy): / /				Date Homog. (mm/dd/yyyy): / /			
#	Tissue/Bag ID	Fish #	Organism ID	Composite / Individual ID	empty weigh boat	Part Wt w/ Boat (FreezerWet Wt)	Dry Weight w/ Boat	Sex	Part	Anomaly	Body Location
1								NR	Whole		
2								NR	Whole		
3								NR	Whole		
4								NR	Whole		
5								NR	Whole		
6								NR	Whole		
7								NR	Whole		
8								NR	Whole		
9								NR	Whole		
10								NR	Whole		
OrganismID: xxxxxxxxLXX##YYzz-ZZ; unique code - StationCode (xxxxxxx), Location (LL), Project (XX), ProjectYear (##), OrganismCode (YYY), Bag # (zz), Fish # (ZZ); ex. 203SRF101L1SW04CAR01-01											
TissueID: Differentiates different parts from same fish or differentiates composited vs. individual fish											
Part: Tissue (T), Liver (L), Other (O) - list in Comments											
Comp/IndID: Unique code; include Agency code in the ID; e.g., 2003-1823-MLML or C031501-MLML											
Anomalies: Ambicoloration (A), Albinism (B), Cloudiness (CL), Deformity-skeletal (D), Discoloration (DC), Depression (DS), Fin Erosion (F), Gill Erosion (T), Hemorrhage (H), Lesion (L), Parasite (P),											
Body Locations: Branchial Chamber (BRC), Buccal Cavity (BC), Eyes (E), Musculoskeleton (M), Skin/Fins (SF)								Popeye (PE), Tumor (T), Ulceration (U), White Spots (W), and any combination			
Comments: Measure length to nearest 1 mm; Measure weight to nearest 0.01 g; Keep archive tissue if possible; If a duplicate is made, use DupID as identification for analysis											
ModEed 06/09/07											

<b>SWAMP Lab Data Sheet - FISH</b>		ProjectID: SWB_WildLk_2012	PrepPres: Skin ON; Scales ON	LabID:	Pg: 1 of 2 Pgs
StationCode:	Tissue: Whole Body		Entered d-base (initial/date)		
StationName:	Homog. Method: USGS SOP Mortar/Pestle		Staff: Diss.	Homog.	
Species Name:	Date Diss. (mm/dd/yyyy): / /		Date Homog. (mm/dd/yyyy): / /		
<b>CHEMISTRY JARS</b>					
Individual ID: _____		Individual ID: _____		Individual ID: _____	
Analysis: Mercury		Analysis: Mercury		Analysis: Mercury	
Jar Weight Full (g): _____		Jar Weight Full (g): _____		Jar Weight Full (g): _____	
Jar Weight Empty (g): _____		Jar Weight Empty (g): _____		Jar Weight Empty (g): _____	
Comp Tissue Wt (Jar Full - Empty; g): _____		Comp Tissue Wt (Jar Full - Empty; g): _____		Comp Tissue Wt (Jar Full - Empty; g): _____	
Individual ID: _____		Individual ID: _____		Individual ID: _____	
Analysis: Mercury		Analysis: Mercury		Analysis: Mercury	
Jar Weight Full (g): _____		Jar Weight Full (g): _____		Jar Weight Full (g): _____	
Jar Weight Empty (g): _____		Jar Weight Empty (g): _____		Jar Weight Empty (g): _____	
Comp Tissue Wt (Jar Full - Empty; g): _____		Comp Tissue Wt (Jar Full - Empty; g): _____		Comp Tissue Wt (Jar Full - Empty; g): _____	
Individual ID: _____		Individual ID: _____		Individual ID: _____	
Analysis: Mercury		Analysis: Mercury		Analysis: Mercury	
Jar Weight Full (g): _____		Jar Weight Full (g): _____		Jar Weight Full (g): _____	
Jar Weight Empty (g): _____		Jar Weight Empty (g): _____		Jar Weight Empty (g): _____	
Comp Tissue Wt (Jar Full - Empty; g): _____		Comp Tissue Wt (Jar Full - Empty; g): _____		Comp Tissue Wt (Jar Full - Empty; g): _____	
Individual ID: _____		Individual ID: _____		Individual ID: _____	
Analysis: Mercury		Analysis: Mercury		Analysis: Mercury	
Jar Weight Full (g): _____		Jar Weight Full (g): _____		Jar Weight Full (g): _____	
Jar Weight Empty (g): _____		Jar Weight Empty (g): _____		Jar Weight Empty (g): _____	
Comp Tissue Wt (Jar Full - Empty; g): _____		Comp Tissue Wt (Jar Full - Empty; g): _____		Comp Tissue Wt (Jar Full - Empty; g): _____	
Individual ID: _____ Analysis: Mercury Jar Weight Full (g): _____ Jar Weight Empty (g): _____ Comp Tissue Wt (Jar Full - Empty; g): _____					
Comments: _____ _____ _____					