

# APPENDIX A. FOOD WEB MODEL PARAMETERS

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## Table of Contents

A.1	SUMMARY OF PARAMETERS	1
	TABLE A-1-1. SUMMARY OF PARAMETERS	1
	TABLE A-1-2. EQUATIONS FOR THE ARNOT AND GOBAS (2004) MODEL	4
A.2	PARAMETERS DERIVED FROM SITE-SPECIFIC DATA	8
A.2.1	Sediment PCBs and OC <sub>sed</sub>	8
	TABLE A-2-1. MODEL COMPONENTS WITH VALUES DETERMINED USING SITE-SPECIFIC DATA	9
	TABLE A-2-2. TISSUE CHEMISTRY DATASETS USED IN THE PRELIMINARY FWM	12
A.2.2	Water chemistry data	14
A.2.3	Tissue data	15
	TABLE A-2-3. CHARACTERISTICS OF THE MODELED SPECIES	16
	TABLE A-2-4. TOTAL PCB TISSUE CONCENTRATIONS IN FISH AND INVERTEBRATE SPECIES TO BE USED TO EVALUATE MODEL PERFORMANCE	32
	TABLE A-2-5. NUMBER OF COMPOSITE TISSUE SAMPLES AVAILABLE FROM EACH LDW MODELING AREA	37
	TABLE A-2-6. WEIGHT, PERCENT LIPIDS, AND PERCENT SOLIDS DATA USED IN THE FWM FOR WHOLE-BODY FISH AND CRABS	39
A.2.4	Relationship between co-located benthic invertebrate tissue and surface sediment total PCB concentrations	40
A.2.5	Benthic Invertebrate taxonomy data	41
A.2.6	Estimation of log K <sub>OW</sub> for PCBs	44
	TABLE A-2-7. AVERAGE LOG K <sub>OWS</sub> FOR EACH MODELED SPECIES DERIVED USING SITE-SPECIFIC TISSUE DATA FROM THE LDW	44
A.3	PARAMETERS DERIVED FROM THE LITERATURE	45
	TABLE A-3-1. MODEL COMPONENTS WITH VALUES DERIVED FROM THE LITERATURE	46
A.3.1	Fish and crab dietary scenarios	47
	TABLE A-3-2. DIETARY STUDIES USED TO CHARACTERIZE MODELED SPECIES' DIETS	47
	TABLE A-3-3. PERCENT OF PHYTOPLANKTON/ALGAE, BENTHIC INVERTEBRATES, ZOOPLANKTON, AND FISH IN ALL FISH AND CRAB SPECIES' DIETS FOR DIETARY SCENARIO 1 (AND DUNGENESS CRAB FOR DIETARY SCENARIO 4)	49
	TABLE A-3-4. PERCENT OF PHYTOPLANKTON/ALGAE, BENTHIC INVERTEBRATES, ZOOPLANKTON, AND FISH IN ALL FISH AND CRAB SPECIES' DIETS FOR DIETARY SCENARIO 2	50
A.4	DEFAULT PARAMETER VALUES FROM ARNOT AND GOBAS MODEL APPLICATION TO THE GREAT LAKES AND SAN FRANCISCO BAY	50
	TABLE A-4-1. DEFAULT VALUES FROM ARNOT AND GOBAS MODEL APPLICATION TO THE GREAT LAKES AND SAN FRANCISCO BAY	51
A.5	REFERENCES	54

## Appendix A. Food Web Model Parameters

This appendix presents the food web model (FWM) parameter values used for the Lower Duwamish Waterway (LDW)-wide and smaller spatial scale initial model runs. Section A.1 presents a summary of Arnot and Gobas (2004) specific model equations. Section A.2 presents parameter values derived using site-specific data. Section A.3 presents parameter values derived from the literature. Section A.4 presents parameter values used or cited in Arnot and Gobas (2004) or Gobas and Arnot (2005). Section A.5 presents parameter values specific to each fish and invertebrate species modeled.

### A.1 SUMMARY OF PARAMETERS

A summary of parameters in the Arnot and Gobas (2004) model are presented in Table A-1-1. The equations for the Arnot and Gobas (2004) model define environmental, biological, or chemical conditions or processes. The Arnot and Gobas model equations are presented in Table A-1-2. Parameter symbols within equations are defined in Tables A-2-1, A-3-1, and A-4-1.

**Table A-1-1. Summary of Parameters**

PARAMETER	SYMBOL	ORIGIN	TABLE WITH DETAILED INFORMATION
Algae, phytoplankton, and aquatic macrophytes – resistance to chemical uptake through aqueous phase	A	default value from model application to the Great Lakes and San Francisco Bay	A-4-1
Algae, phytoplankton, and aquatic macrophytes – resistance to chemical uptake through organic phase	B	default value from model application to the Great Lakes and San Francisco Bay	A-4-1
Bioavailable solute fraction	$\phi$	calibrated in model	A-1-2
Chemical concentration in prey item <i>i</i>	$C_{D,i}$	calibrated in model	A-1-2
Chemical concentration in the modeled species	$C_B$	calibrated in model	A-1-2
Chemical concentration in the sediment, organic carbon normalized	$C_{S,OC}$	calibrated in model	A-1-2
Concentration of DOC in the water column	$\chi_{DOC}$	site-specific empirical data	A-2-1
Concentration of POC in the water column	$\chi_{POC}$	site-specific empirical data	A-2-1
Concentration of suspended solids in water column	$C_{SS}$	site-specific empirical data	A-2-1
Density of lipids	$\delta_\Lambda$	default value from model application to the Great Lakes and San Francisco Bay	A-4-1
Density of water	$\delta_\Omega$	default value from model application to the Great Lakes and San Francisco Bay	A-4-1
Dietary absorption efficiencies of lipid	$\epsilon_L$	default value from model application to the Great Lakes and San Francisco Bay	A-4-1

PARAMETER	SYMBOL	ORIGIN	TABLE WITH DETAILED INFORMATION
Dietary absorption efficiencies of NLOM/NLOC	$\epsilon_N$	default value from model application to the Great Lakes and San Francisco Bay	A-4-1
Dietary absorption efficiencies of water	$\epsilon_W$	default value from model application to the Great Lakes and San Francisco Bay	A-4-1
Dietary chemical transfer efficiency	$E_D$	calibrated in model	A-1-2
Disequilibrium factor for DOC partitioning	$D_{DOC}$	default value from model application to the Great Lakes and San Francisco Bay	A-4-1
Disequilibrium factor for POC partitioning	$D_{POC}$	default value from model application to the Great Lakes and San Francisco Bay	A-4-1
Dissolved oxygen concentration of water column	$C_{OX}$	site-specific empirical data	A-2-1
Fecal egestion rate	$G_F$	calibrated in model	A-1-2
Feeding rate	$G_D$	calibrated in model	A-1-2
Fraction of overlying water ventilated	$m_O$	derived from literature	A-3-1
Fraction of porewater ventilated	$m_P$	derived from literature	A-3-1
Fraction of the diet consisting of prey item $i$	$P_i$	derived from literature	A-3-1
Freely dissolved chemical concentration in the porewater	$C_{WD,P}$	calibrated in model	A-1-2
Freely dissolved chemical concentration in the water (total PCBs as Aroclors)	$C_{WD}$	calibrated in model	A-1-2
Gill ventilation rate	$G_V$	calibrated in model	A-1-2
Henry's Law Constant	$H$	derived from literature	A-3-1
Lipid content of organism	$V_{LB}$	site-specific empirical data	A-2-1
Lipid content of organism (zooplankton)	$V_{LB}$	derived from literature	A-3-1
Lipid content of phytoplankton/algae	$V_{LP}$	derived from literature	A-3-1
Lipid fraction of gut contents	$V_{LG}$	calibrated in model	A-1-2
Mean water column temperature	$T$	site-specific empirical data	A-2-1
NLOC content of phytoplankton/algae	$V_{OCP}$	derived from literature	A-3-1
NLOC fraction of gut contents	$V_{OCG}$	calibrated in model	A-1-2
NLOM content of organism	$V_{NB}$	site-specific empirical data	A-2-1
NLOM content of organism (zooplankton)	$V_{NB}$	derived from literature	A-3-1
NLOM fraction of gut contents	$V_{NG}$	calibrated in model	A-1-2
Octanol-water partition coefficient (total PCBs)	$K_{OW}$	derived from literature	A-3-1
Organic carbon-water partition coefficient	$K_{OC}$	calibrated in model	A-1-2
Organism-water partition coefficient on a wet weight basis	$K_{BW}$	calibrated in model	A-1-2
Overall lipid content of the diet	$V_{LD}$	calibrated in model	A-1-2
Overall NLOC content of the diet	$V_{OCD}$	calibrated in model	A-1-2
Overall NLOM content of the diet	$V_{ND}$	calibrated in model	A-1-2
Overall water content of the diet	$V_{WD}$	calibrated in model	A-1-2

PARAMETER	SYMBOL	ORIGIN	TABLE WITH DETAILED INFORMATION
Partition coefficient of the chemical between the contents of the gastrointestinal tract and the organism	$K_{GB}$	calibrated in model	A-1-2
Phytoplankton/algae-water partition coefficient on a wet weight basis	$K_{PW}$	calibrated in model	A-1-2
Proportionality constant describing similarity in phase partitioning of DOC relative to that of octanol	$\alpha_{DOC}$	default value from model application to the Great Lakes and San Francisco Bay	A-4-1
Proportionality constant describing similarity in phase partitioning of POC relative to that of octanol	$\alpha_{POC}$	default value from model application to the Great Lakes and San Francisco Bay	A-4-1
Proportionality constant expressing the sorption capacity of NLOC relative to that of octanol	$\beta_{OC}$	default value from model application to the Great Lakes and San Francisco Bay	A-4-1
Proportionality constant expressing the sorption capacity of NLOM relative to that of octanol	$\beta$	default value from model application to the Great Lakes and San Francisco Bay	A-4-1
Rate constant for aqueous uptake	$k_1$	calibrated in model	A-1-2
Rate constant for chemical elimination via excretion into egested feces	$k_E$	calibrated in model	A-1-2
Rate constant for chemical elimination via the respiratory area	$k_2$	calibrated in model	A-1-2
Rate constant for chemical uptake via the diet	$k_D$	calibrated in model	A-1-2
Rate constant for growth of aquatic organisms	$k_G$	calibrated in model	A-1-2
Rate constant for growth of phytoplankton/algae	$k_G$	default value from model application to the Great Lakes and San Francisco Bay	A-4-1
Rate constant for metabolic transformation of the chemical	$k_M$	default value from model application to the Great Lakes and San Francisco Bay	A-4-1
Respiratory surface chemical uptake efficiency	$E_W$	calibrated in model	A-1-2
Scavenging efficiency of particles absorbed from the water	$\sigma$	default value from model application to the Great Lakes and San Francisco Bay	A-4-1
Sediment OC content	$OC_{sed}$	site-specific empirical data	A-2-1
Total chemical concentration in the water column (Total PCBs as Aroclors)	$C_{WT}$	site-specific empirical data	A-2-1
Total chemical concentration in the sediment (Total PCBs as Aroclors)	$C_S$	site-specific empirical data	A-2-1
Water content of organism	$V_{WB}$	site-specific empirical data	A-2-1
Water content of organism (zooplankton)	$V_{WB}$	derived from literature	A-3-1
Water content of phytoplankton/algae	$V_{WP}$	derived from literature	A-3-1
Water fraction of gut contents	$V_{WG}$	calibrated in model	A-1-2
Weight of the organism	$W_B$	site-specific empirical data	A-2-1
Weight of the organism (zooplankton)	$W_B$	derived from literature	A-3-1

DOC – dissolved organic carbon  
NLOC – non-lipid organic carbon  
NLOM – non-lipid organic matter

OC – organic carbon  
PCB – polychlorinated biphenyl  
POC – particulate organic carbon

**Lower Duwamish Waterway Group**

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FWM memorandum 3:  
Appendix A  
April 7, 2006  
Page 3

**Table A-1-2. Equations for the Arnot and Gobas (2004) Model**

PARAMETER	SYMBOL	UNITS	EQUATION	VALUE	NOTES	SOURCE
<b>Biological</b>						
Chemical concentration in the modeled species	$C_B$	µg/kg ww	$C_B = \{k_1 \times (m_O \times C_{WD} + m_P \times C_{WD,P}) + k_D \times \sum P_i \times C_{D,i}\} / (k_2 + k_E + k_G + k_M)$	species-specific model output	See Table A-2-4 for tissue chemistry data to be used to evaluate model performance.	Arnot and Gobas (2004)
Chemical concentration in prey item <i>i</i>	$C_{D,i}$	µg/kg ww	same as above	species-specific model output	See Table A-2-4 for tissue chemistry data to be used to evaluate model performance.	Arnot and Gobas (2004)
Rate constant for aqueous uptake (fish, invertebrates, and zooplankton)	$k_1$	L/kg·day	$k_1 = E_W \times G_V / W_B$	calculated in model using equation at left	For chemical uptake via the respiratory area (i.e., gills)	Gobas (1993); Gobas and MacKay (1987) as cited in Arnot and Gobas (2004)
Rate constant for aqueous uptake (algae, phytoplankton, and aquatic macrophytes)	$k_1$	L/kg·day	$k_1 = (A + (B/K_{OW}))^{-1}$	calculated in model using equation at left	For chemical uptake via the respiratory area (i.e., cell wall)	Arnot and Gobas (2004)
Rate constant for chemical elimination via the respiratory area	$k_2$	day <sup>-1</sup>	$k_2 = k_1 / K_{BW}$	calculated in model using equation at left	Loss through respiratory surface (gills or cell membrane/wall)	Gobas (1993) as cited in Arnot and Gobas (2004)
Rate constant for chemical uptake via the diet	$k_D$	kg food/kg organism·day	$k_D = E_D \times G_D / W_B$	calculated in model using equation at left	For phytoplankton/algae, $k_D$ is zero.	Gobas (1993) as cited in Arnot and Gobas (2004)
Rate constant for chemical elimination via excretion into egested feces	$k_E$	day <sup>-1</sup>	$k_E = G_F \times E_D \times K_{GB} / W_B$	calculated in model using equation at left	For phytoplankton/algae, $k_E$ is zero.	Gobas et al. (1993) as cited in Arnot and Gobas (2004)
Rate constant for growth of aquatic organisms	$k_G$	day <sup>-1</sup>	$k_G = 0.000502 \times W_B^{-0.2}$	calculated in model using equation at left	For temperatures around 10°C.	Thomann et al. (1992) as cited in Arnot and Gobas (2004)
Dietary chemical transfer efficiency	$E_D$	%	$E_D = (3.0 \times 10^{-7} \times K_{OW} + 2.0)^{-1}$	calculated in model using equation at left	Transfer of chemical across gut can be characterized by $K_{OW}$ relationship.	Arnot and Gobas (2004)
Respiratory surface chemical uptake efficiency	$E_W$	%	$E_W = (1.85 + (155 / K_{OW}))^{-1}$	calculated in model using equation at left	Transfer of chemical across respiratory surface can be characterized by $K_{OW}$ relationship.	Gobas (1988) as cited in Arnot and Gobas (2004)
Feeding rate – filter feeders	$G_D$	kg/d	$G_D = G_V \times C_{SS} \times \sigma$	calculated in model using equation at left		Morrison et al. (1996) as cited in Arnot and Gobas (2004)

PARAMETER	SYMBOL	UNITS	EQUATION	VALUE	NOTES	SOURCE
Feeding rate – other species	$G_D$	kg/d	$G_D = 0.022 \times W_B^{0.85} \times e^{(0.06 \times T)}$	calculated from weight of biota	Studies of feeding rates in cold-water fish (being used for zooplankton and aquatic invertebrate species as well).	Weiniger (1978) as cited in Arnot and Gobas (2004)
Fecal egestion rate	$G_F$	kg/d	$G_F = \{(1-\epsilon_L) \times V_{LD}\} + (1-\epsilon_N) \times V_{OCD} + (1-\epsilon_N) \times V_{ND} + (1-\epsilon_W) \times V_{WD}\} \times G_D$	calculated in model using equation at left		Arnot and Gobas (2004)
Gill ventilation rate	$G_V$	L/d	$G_V = 1400 \times W_B^{0.65} / C_{OX}$	calculated in model using equation at left		Arnot and Gobas (2004)
Organism-water partition coefficient on a wet weight basis	$K_{BW}$	L water/kg biota	$K_{BW} = k_1/k_2 = V_{LB} \times K_{OW}/\delta_L + V_{NB} \times \beta \times K_{OW} / (V_{LB} \times K_{OW}/\delta_L + V_{WB}/\delta_W)$	calculated in model using equation at left		Arnot and Gobas (2004)
Phytoplankton/algae-water partition coefficient on a wet weight basis	$K_{PW}$	L water/kg phytoplankton/algae	$K_{PW} = V_{LP} \times K_{OW}/\delta_L + \beta_{OC} \times V_{NP} \times K_{OW} + V_{WP}/\delta_W$	calculated in model using equation at left		Arnot and Gobas (2004)
Partition coefficient of the chemical between the contents of the gastrointestinal tract and the organism	$K_{GB}$	kg biota/kg digesta	$K_{GB} = (V_{LG} \times K_{OW}/\delta_L + V_{OCG} \times \beta_{OC} \times K_{OW} + V_{NG} \times \beta \times K_{OW} + V_{WG}/\delta_W) / (V_{LB} \times K_{OW}/\delta_L + V_{NB} \times \beta \times K_{OW} + V_{WB}/\delta_W)$	calculated in model using equation at left		Arnot and Gobas (2004)
Lipid fraction of gut contents	$V_{LG}$	kg lipid/kg digesta ww	$V_{LG} = (1-\epsilon_L) \times V_{LD} / [(1-\epsilon_L) \times V_{LD} + (1-\epsilon_N) \times V_{OCD} + (1-\epsilon_N) \times V_{ND} + (1-\epsilon_W) \times V_{WD}]$	calculated in model using equation at left		Arnot and Gobas (2004)
NLOC fraction of gut contents	$V_{OCG}$	kg lipid/kg digesta ww	$V_{OCG} = [(1-\epsilon_N) \times V_{OCD}] / [(1-\epsilon_L) \times V_{LD} + (1-\epsilon_N) \times V_{OCD} + (1-\epsilon_N) \times V_{ND} + (1-\epsilon_W) \times V_{WD}]$	calculated in model using equation at left	NLOC was added to the model to account for differential affinity of PCBs for NLOC (higher) as compared to NLOM	January 2006 update to Arnot and Gobas model (Arnot and Gobas 2004). Updated model, AQUAWEB, can be found on Environmental Toxicology Research Group website (Gobas 2006)
NLOM fraction of gut contents	$V_{NG}$	kg NLOM/kg digesta ww	$V_{NG} = (1-\epsilon_N) \times V_{ND} / [(1-\epsilon_L) \times V_{LD} + (1-\epsilon_N) \times V_{OCD} + (1-\epsilon_N) \times V_{ND} + (1-\epsilon_W) \times V_{WD}]$	calculated in model using equation at left		Arnot and Gobas (2004)

PARAMETER	SYMBOL	UNITS	EQUATION	VALUE	NOTES	SOURCE
Water fraction of gut contents	$V_{WG}$	kg water/kg digesta ww	$V_{WG} = (1 - \epsilon_W) \times V_{WD} / [(1 - \epsilon_L) \times V_{LD} + (1 - \epsilon_N) \times V_{OCD} + (1 - \epsilon_N) \times V_{ND} + (1 - \epsilon_W) \times V_{WD}]$	calculated in model using equation at left		Arnot and Gobas (2004)
Overall lipid content of the diet	$V_{LD}$	kg lipid/kg food ww	$V_{LD} = \sum P_i \times V_{LB,i}$	calculated in model using equation at left		Arnot and Gobas model spreadsheet (Gobas 2006)
Overall NLOC content of the diet	$V_{OCD}$	kg NLOC/kg food ww	$V_{OCD} = P_p \times V_{OCP} + P_{sed} \times OC_{sed}$	calculated in model using equation at left	NLOC content of diet is determined by fraction of phytoplankton/algae and sediment consumed. These are the only dietary items with NLOC as a constituent.	January 2006 update to Arnot and Gobas model (Arnot and Gobas 2004). Updated model, AQUAWEB, can be found on Environmental Toxicology Research Group website (Gobas 2006)
Overall NLOM content of the diet	$V_{ND}$	kg NLOM/kg food ww	$V_{ND} = \sum P_i \times V_{NB,i}$	calculated in model using equation at left		Arnot and Gobas model spreadsheet (Gobas 2006)
Overall water content of the diet	$V_{WD}$	kg water/kg food ww	$V_{WD} = \sum P_i \times V_{WB,i}$	calculated in model using equation at left		Arnot and Gobas model spreadsheet (Gobas 2006)
<b>Environmental</b>						
Freely dissolved chemical concentration in the porewater	$C_{WD,P}$	$\mu\text{g/L}$	$C_{WD,P} = C_{S,OC} / K_{OC}$	calculated in model using equation at left	This parameter will be calculated for each spatial scale evaluated using sediment data appropriate for that spatial scale.	Kraaij et al. (2002) as cited in Arnot and Gobas (2004)
Chemical concentration in the sediment, organic carbon normalized	$C_{S,OC}$	$\mu\text{g/kg}$	$C_{S,OC} = C_S / OC_{sed}$	calculated in model using equation at left	This parameter will be calculated for each spatial scale evaluated, using sediment data appropriate for that spatial scale.	Calculated using Phase 1 and Phase 2 sediment data
Freely dissolved chemical concentration in the water (total PCBs as Aroclors)	$C_{WD}$	$\mu\text{g/L}$	$C_{WD} = (C_{WT} \times \phi) / 1000$	calculated in model using equation at left	Simulates sequestering of chemical by DOC and POC in the water.	Arnot and Gobas (2004)
Bioavailable solute fraction	$\phi$	unitless	$\phi = 1 / (1 + \chi_{POC} \cdot D_{POC} \cdot \alpha_{POC} \cdot K_{OW} + \chi_{DOC} \cdot D_{DOC} \cdot \alpha_{DOC} \cdot K_{OW})$	calculated in model using equation at left	Simulates sequestering of chemical by DOC and POC in the water.	Arnot and Gobas (2004)



PARAMETER	SYMBOL	UNITS	EQUATION	VALUE	NOTES	SOURCE
<b>Chemical</b>						
Organic carbon-water partition coefficient	$K_{OC}$	L/kg	$K_{OC} = 0.35 \times K_{ow}$	calculated in model from equation at left	There are many different relationships established between $K_{ow}$ and $K_{OC}$ . This relationship was based on the analysis of a wide range of analytes (including PCB congeners) and soil/sediment matrices. The authors excluded data that may not have represented equilibrium conditions that can be very influential for high molecular weight PCBs. It is consistent with the commonly used approximation of $K_{OC} = 0.4 K_{ow}$ .	Seth et al. (1999)

DOC – dissolved organic carbon  
 NLOC – non-lipid organic carbon  
 NLOM – non-lipid organic matter  
 PCB – polychlorinated biphenyl  
 POC – particulate organic carbon  
 ww – wet weight

## **A.2 PARAMETERS DERIVED FROM SITE-SPECIFIC DATA**

The LDW site-specific data presented in this section were derived from various field sampling events conducted in the LDW. Parameter names, symbols, units, selected values, comments, and source information for the initial set of parameters are presented in Table A-2-1. Parameters for which derivation of values cannot be fully explained within the limited space of a table are further discussed in the following subsections.

### **A.2.1 Sediment PCBs and OC<sub>sed</sub>**

Concentrations of total polychlorinated biphenyls (PCBs) (Aroclor sum) and organic carbon (OC<sub>sed</sub>) in surface sediment data were derived from Phase 2 and historical (Phase 1) datasets according to “baseline” conditions, as described in the draft Technical Memorandum: Criteria for Defining the Baseline Surface Sediment Dataset for Use in the Lower Duwamish Waterway Phase 2 RI/FS (Windward 2006). Any changes to the baseline dataset will be reflected in the Phase 2 RI, where the final FWM results will be presented.

**Table A-2-1. Model components with values determined using site-specific data**

PARAMETER	SYMBOL	UNITS	VALUES – MEAN (range)	NO. OF SAMPLES	NOTES	SOURCE
<b>Biological</b>						
Weight of the organism	$W_B$	kg ww	species-specific	see Table A-2-3	see Table A-2-3	see Table A-2-3
Lipid content of organism	$V_{LB}$	% ww	species-specific	see Table A-2-3	see Table A-2-3	see Table A-2-3
Non Lipid Organic Matter (NLOM) content of organism	$V_{NB}$	% ww	species-specific	see Table A-2-3	See Table A-2-3. NLOM is a secondary site of PCB accumulation.	see Table A-2-3
Water content of organism	$V_{WB}$	% ww	species-specific	see Table A-2-3	See Table A-2-3. Water is not a significant contributor to the storage capacity of PCBs but is the third phase of storage in the body.	see Table A-2-3
<b>Environmental</b>						
Total PCB (as Aroclors) concentration in sediment (all LDW)	$C_S$	µg/kg dw	250	1,294	Spatially weighted average concentration calculated using IDW over the entire LDW	Calculated using baseline surface sediment data
Total PCB (as Aroclors) concentration in sediment (modeling area M1)	$C_S$	µg/kg dw	280	305	LDW-wide spatially weighted average concentration calculated using IDW over the entire LDW, but then clipped to M1.	Calculated using baseline surface sediment data.
Total PCB (as Aroclors) concentration in sediment (modeling area M2)	$C_S$	µg/kg dw	160	198	LDW-wide spatially weighted average concentration calculated using IDW over the entire LDW, but then clipped to M2.	Calculated using baseline surface sediment data.
Total PCB (as Aroclors) concentration in sediment (modeling area M3)	$C_S$	µg/kg dw	470	485	LDW-wide spatially weighted average concentration calculated using IDW over the entire LDW, but then clipped to M3.	Calculated using baseline surface sediment data.
Total PCB (as Aroclors) concentration in sediment (modeling area M4)	$C_S$	µg/kg dw	41	265	LDW-wide spatially weighted average concentration calculated using IDW over the entire LDW, but then clipped to M4.	Calculated using baseline surface sediment data.
Sediment OC content (all LDW)	$OC_{sed}$	% dw	1.93	1,294	Spatially weighted average concentration calculated using IDW over the entire LDW.	Calculated using baseline surface sediment data.

PARAMETER	SYMBOL	UNITS	VALUES – MEAN (range)	NO. OF SAMPLES	NOTES	SOURCE
Sediment OC content (modeling area M1)	OC <sub>sed</sub>	% dw	2.00	305	LDW-wide, spatially weighted average concentration calculated using IDW over the entire LDW, but then clipped to M1.	Calculated using baseline surface sediment data.
Sediment OC content (modeling area M2)	OC <sub>sed</sub>	% dw	2.05	198	LDW-wide spatially weighted average concentration calculated using IDW over the entire LDW, but then clipped to M2.	Calculated using baseline surface sediment data.
Sediment OC content (modeling area M3)	OC <sub>sed</sub>	% dw	1.75	485	LDW-wide spatially weighted average concentration calculated using IDW over the entire LDW, but then clipped to M3.	Calculated using baseline surface sediment data.
Sediment OC content (modeling area M4)	OC <sub>sed</sub>	% dw	1.80	265	LDW-wide spatially weighted average concentration calculated using IDW over the entire LDW, but then clipped to M4.	Calculated using baseline surface sediment data.
<b>Water</b>						
Total chemical concentration in the water column (total PCBs as Aroclors)	C <sub>WT</sub>	ng/L	2 (1.5 – 3.1)	King County samples (5 samples) = two stations, two depths, one event (with one field duplicate)	Model water scenarios will also be run using the following set of values: 1, 3, 5, and 10. PCB water mean has one significant figure due to uncertainty and low sample size.	Data received from King County (unpublished) (Williston 2005) for August 2005 sampling event, sample locations and collection methods are in the sampling and analysis plan (King County 2005)
Dissolved oxygen concentration of water column	C <sub>ox</sub>	mg/L	8.0 (6.4 – 9.6)	11	Average of two stations, two depths from King County data, for each month Jan – Nov 2005.	King County Water Quality Monitoring Program (marine) (Mickelson 2006)
Mean water column temperature	T	° Celsius	11.6 (8.1 – 14.7)	11	Average of two stations, two depths from King County data, for each month Jan – Nov 2005.	King County Water Quality Monitoring Program (marine) (Mickelson 2006)
Concentration of DOC in the water column	χ <sub>DOC</sub>	kg/L	$2.2 \times 10^{-6}$ ( $1.4 \times 10^{-6} - 4.0 \times 10^{-6}$ )	11	Average of two stations, two depths from King County data, for each month Jan – Nov 2005.	King County Water Quality Monitoring Program (marine) (Mickelson 2006)

PARAMETER	SYMBOL	UNITS	VALUES – MEAN (range)	NO. OF SAMPLES	NOTES	SOURCE
<b>Concentration of POC in the water column</b>	$\chi_{\text{POC}}$	kg/L	$2.9 \times 10^{-7}$ ( $9.3 \times 10^{-8}$ – $7.7 \times 10^{-7}$ )	11	<b>Calculated concentration (POC = TOC – DOC) Average of two stations, two depths from King County data, for each month Jan – Nov 2005.</b>	<b>King County Water Quality Monitoring Program (marine) (Mickelson 2006)</b>
Concentration of suspended solids in water column	$C_{\text{SS}}$	kg/L	$4.6 \times 10^{-6}$ ( $1.9 \times 10^{-6}$ – $7.6 \times 10^{-6}$ )	11	Average of two stations, two depths from King County data, for each month Jan – Nov 2005.	King County Water Quality Monitoring Program (marine) (Mickelson 2006)

**Bold** text indicates that the model has been demonstrated to be sensitive to that parameter in the past (Arnot 2005).

DOC – dissolved organic carbon

dw – dry weight

EFDC – Environmental Fluid Dynamics Code

IDW – inverse distance weighting

LDW – Lower Duwamish Waterway

NLOM – non-lipid organic matter

OC – organic carbon

PCB – polychlorinated biphenyl

POC – particulate organic carbon

TOC – total organic carbon

WQA – water quality assessment

ww –wet weight

**Table A-2-2. Tissue chemistry datasets used in the preliminary FWM**

SAMPLING EVENT	YEAR	SPECIES	TISSUE TYPE	NUMBER OF INDIVIDUALS PER COMPOSITE TISSUE SAMPLE	N	PARAMETER	SOURCE
LDW Phase 2	2005	Dungeness crab	edible meat	5	3	weight, percent lipids, percent solids, PCB Aroclors	Windward (2006 in prep)
			hepatopancreas	5	3		
		slender crab	edible meat	5	1		
			hepatopancreas	10	1		
		English sole	whole body	5	11		
			paired skin-on fillet and remainder <sup>a</sup>	5	10		
	shiner surfperch	whole body	10	22			
	Pacific staghorn sculpin	whole body	10	4			
	2004	benthic invertebrates	whole body	> 100	20	weight, percent lipids, percent solids, community structure, PCB Aroclors	Windward (2005a, b)
		Dungeness crab	edible meat	5	7	weight, percent lipids, percent solids, PCB Aroclors, PCB congeners <sup>b</sup>	Windward (2005c, d)
			hepatopancreas	6 – 15	3		
		slender crab	edible meat	5	12		
			hepatopancreas	15 – 18	4		
		English sole	whole body	5	21		
Pacific staghorn sculpin		whole body	7 – 10	24			
shiner surfperch	whole body	9 – 10	24				
starry flounder	whole body	5	3				
National Marine Fisheries Service (NMFS) Duwamish injury assessment project	2000	shiner surfperch	whole body	1	2	PCB Aroclors	NMFS (2002)

SAMPLING EVENT	YEAR	SPECIES	TISSUE TYPE	NUMBER OF INDIVIDUALS PER COMPOSITE TISSUE SAMPLE	N	PARAMETER	SOURCE
King County Combined Sewer Overflow Water Quality Assessment for the Duwamish River and Elliott Bay	1996 - 1997	Dungeness crab	edible meat	3	2	percent lipids, percent solids, PCB Aroclors	King County (1999)
			hepatopancreas	3	1		
		shiner surfperch	whole body	10	3		

<sup>a</sup> The remainder is the portion of fish that remains after removal of the skin-on fillet. These remainder and fillet data were used to estimate whole-body English sole concentrations as specified in the QAPP and the data report (Windward 2005f, 2006 in prep).

<sup>b</sup> The following composite samples were analyzed for PCB congeners: three Dungeness crab edible meat samples, two Dungeness crab hepatopancreas samples, five slender crab edible meat samples, two slender crab hepatopancreas samples, seven English sole whole-body samples, nine shiner surfperch whole-body samples, and eight Pacific staghorn sculpin whole-body samples.

N – Number of composite tissue samples analyzed

The total PCB sediment concentrations and  $OC_{sed}$  percentages used in the preliminary FWM runs were calculated from inverse distance weighting (IDW) interpolations derived from 1,294 surface sediment samples. The IDW parameters (e.g., search radius, weighting factor) were selected to optimize the ability of the IDW interpolation to predict concentrations of total PCBs in sediment where data are available for comparison. The optimized interpolation resulted in a predicted spatially weighted average concentration (SWAC) for total PCBs of 250  $\mu\text{g}/\text{kg dw}$  and a spatially weighted average for total organic carbon (TOC) of 1.93% for the entire LDW (Table A-2-1).

Spatially weighted average concentrations and percentages for smaller spatial scales (modeling areas) were calculated using the same interpolation grids generated for the LDW-wide FWM spatial scale. The 10-ft by 10-ft squares from the IDW grid for each modeling area were selected, and spatially weighted average concentrations and percentages were calculated as the mean of all grid cells within a given modeling area of the LDW (Table A-2-1).

### **A.2.2 Water chemistry data**

Water quality parameters with site-specific data include dissolved oxygen (DO), temperature, total suspended solids (TSS), dissolved organic carbon (DOC) and particulate organic carbon (POC), and total PCB concentrations (PCB congener sum). The particulate organic carbon (POC) was estimated from site-specific values for DOC and total organic carbon (TOC) in the water column. Values for these parameters were derived from 2005 data from the King County Marine Ambient and Outfall Water Column Monitoring Program (Mickelson 2006). In 2005, water samples were collected from two depths (1 m below the surface and 1 m above the sediment surface) at each of two stations in the LDW. The two stations were located just south of Harbor Island and at the 16th Avenue Bridge (King County 2005). Samples were collected for analysis of conventional parameters (temperature, TSS, DOC, and POC) monthly from January through November for a total of 44 samples. Summary statistics for these 44 samples were calculated as follows. Concentrations in the bottom and surface water samples on each collection date were averaged for each of the two sample stations, resulting in 11 average monthly concentrations for each parameter at each station. The average monthly values for the two stations were averaged to estimate the “LDW-wide” average monthly value for each parameter. The mean, minimum, maximum, and standard deviation (SD) concentrations of conventional water quality parameters in Table A-2-1 were calculated directly from the river-wide, monthly average values.

Concentrations of total PCBs in the water column were derived from the 2005 King County Duwamish River, Elliott Bay, and Green River water column PCB congener survey (King County 2005). Water samples were collected from the locations and depths described above. The samples were analyzed for all 209 individual PCB congeners. Four sampling events occurred in 2005, with the goal of capturing two low-flow events (August and September) and two high-flow events (November and



December). Validated data are currently available only from the August sampling event.

### **A.2.3 Tissue data**

All Phase 2 tissue data and historical whole-body data identified as acceptable for use in the Phase 2 RI (Windward 2005g) were used for the FWM (Table A-2-2). Lipid content, water content, and total PCB concentrations were calculated from composite samples. Weight data were derived from all individual specimens collected from the LDW in Phase 2 (2004 and 2005). Weight data for juvenile fish were based on individual shiner surfperch ( $\leq 80$  mm) from both background and LDW locations.

Site-specific tissue data from the LDW were used to determine input parameter values for lipid content ( $V_{LB}$ ), water content ( $V_{WB}$ ), and weight ( $W_B$ ) for crabs and fish (Tables A-2-1 and A-2-3). Site-specific tissue data were also used to evaluate FWM model results at various spatial scales by comparing empirical concentrations of total PCBs in fish and crabs to concentrations predicted by the FWM (Table A-2-4). Methods for estimating an average PCB tissue concentration for benthic invertebrates from site-specific data to be used in model performance evaluation are described in Section A.2.4.

**Table A-2-3. Characteristics of the modeled species**

PARAMETER	VALUE – MEAN (range)	SAMPLE SIZE	LOCALE OF DATA COLLECTION	NOTES	SOURCE
<b>Phytoplankton/algae</b>					
Lipid content (% ww)	0.12 (0.10 – 0.14)	27	False Creek, Burrard Inlet, Vancouver, BC	Three samples each of two species of macroalgae and the contents of a plankton tow at three locations in False Creek. Average of green and brown algae and phytoplankton. “Phytoplankton” tissue analyzed was a combination of phytoplankton and zooplankton (236-µm plankton tow net). <sup>a</sup> The range of values was calculated using the plausible value range approach developed for the sensitivity analysis using standard deviation given in paper.	Mackintosh et al. (2004)
NLOC content (% ww)	4.3 (3.4 – 5.2)	27	False Creek, Burrard Inlet, Vancouver, BC	Three samples each of two species of macroalgae and the contents of a plankton tow at three locations in False Creek. Average of green and brown algae and phytoplankton. Phytoplankton and algae carbon is an important organic chemical storage phase due to low lipid concentrations. Carbon rather than “matter” is used for phytoplankton/algae because it is a better predictor of organic chemical content (Mackintosh et al. 2004). The range of values was calculated using the plausible value range approach developed for the sensitivity analysis using standard deviation given in paper.	Mackintosh et al. (2004)
Water content (% ww)	95.6 (94.7 – 96.5)	27	False Creek, Burrard Inlet, Vancouver, BC	Water content is calculated as 100% – % lipid – % carbon. Not a true measure of water content because there are constituents other than lipid and carbon. The range of values was calculated using the ranges of lipids and NLOC.	Mackintosh et al. (2004)
Fraction of porewater ventilated	0	na	na	Phytoplankton live in water column and are not exposed to porewater. Some benthic algae may be exposed to porewater; however, this biota compartment is primarily representing prey for zooplankton, with a little algae consumed by English sole. Therefore, the algae component of this compartment is not modeled as having exposure to porewater.	
<b>Zooplankton</b>					
Weight (kg ww)	$1.6 \times 10^{-7}$ ( $8.8 \times 10^{-8}$ – $2.3 \times 10^{-7}$ )	126	Puget Sound (Budd Inlet)	Twenty-one samples from six stations (over 12 months). Average dry weight mass of zooplankton with assumed 90% water content (zooplankton was composed primarily of crustaceans, cnidarians, larvaceans, and polychaetes).	Giles and Cordell (1998)
Lipid content (% ww)	1.2 (0.9 – 1.7)	nr	Maizura Bay, Japan	Converted from dry weight assuming 90% water content.	Kuroshima et al. (1987) as cited in Delbare et al. (1996)
NLOM content (% ww)	8.8 (7.1 – 12.1)	na	na	NLOM = 100% - % water - % lipids	calculated from water and lipid content

**Table A-2-3, cont.**

PARAMETER	VALUE – MEAN (range)	SAMPLE SIZE	LOCALE OF DATA COLLECTION	NOTES	SOURCE
Water content (% ww)	90 (87-91.2)	nr	Maizura Bay, Japan	Samples collected over 5 months. Species collected not specified.	Kuroshima et al. (1987) as cited in Delbare et al. (1996)
Fraction of porewater ventilated	0	na	na	Zooplankton live in water column and are not exposed to porewater.	
<b>Dietary Scenario 1 for zooplankton (fraction)</b>					
Phytoplankton/algae	1	na	na	It is assumed that the proportion of carnivorous zooplankton in the LDW is insignificant compared to the proportion of herbivorous zooplankton.	
<b>Benthic invertebrates</b>					
Weight (kg ww) (all LDW)	$5.1 \times 10^{-5}$ ( $5.5 \times 10^{-6}$ – $2.1 \times 10^{-4}$ )	10	LDW	Ten intertidal and ten subtidal samples; weight calculated as average number of individuals divided by sample mass using taxonomy samples from the subtidal zone only (“picked” classification). Weight data were not varied by area because of the uncertainty associated with this calculation (see Section A.2.5).	Phase 2 (2004) benthic invertebrate data
Lipid content (% ww) (all LDW)	0.89 (0.35 – 1.4)	20	LDW	ten intertidal and ten subtidal samples	Phase 2 (2004) benthic invertebrate data
Lipid content (% ww) (modeling area M1)	0.94 (0.69 – 1.3)	6	Area M1	two intertidal and three subtidal samples	Phase 2 (2004) benthic invertebrate data
Lipid content (% ww) (modeling area M2)	1.1 (0.79 – 1.4)	6	Area M2	three intertidal and three subtidal samples	Phase 2 (2004) benthic invertebrate data
Lipid content (% ww) (modeling area M3)	0.66 (0.35 – 1.1)	4	Area M3	two intertidal and two subtidal samples	Phase 2 (2004) benthic invertebrate data
Lipid content (% ww) (modeling area M4)	0.78 (0.62-0.95)	4	Area M4	two intertidal and two subtidal samples	Phase 2 (2004) benthic invertebrate data
Water content (% ww) (all LDW)	88.9 (83.4 – 95.9)	20	LDW	ten intertidal and ten subtidal samples	Phase 2 (2004) benthic invertebrate data
Water content (% ww) (modeling area M1)	87.5 (83.4 – 91.7)	6	Area M1	three intertidal and three subtidal samples	Phase 2 (2004) benthic invertebrate data
Water content (% ww) (modeling area M2)	86.9 (84.3 – 90.6)	6	Area M2	three intertidal and three subtidal samples	Phase 2 (2004) benthic invertebrate data
Water content (% ww) (modeling area M3)	91.9 (86.3 – 95.9)	4	Area M3	two intertidal and two subtidal samples	Phase 2 (2004) benthic invertebrate data
Water content (% ww) (modeling area M4)	90.5(89.3 – 92.4)	4	Area M4	two intertidal and two subtidal samples	Phase 2 (2004) benthic invertebrate data

**Table A-2-3, cont.**

PARAMETER	VALUE – MEAN (range)	SAMPLE SIZE	LOCALE OF DATA COLLECTION	NOTES	SOURCE
Fraction of porewater ventilated/ all exposure areas	0.20 (0.05 – 0.3)	na	na	Benthic invertebrates live on or in sediment and ventilate water just above sediment surface.	Winsor et al. (1990)
<b>Dietary Scenario 1 for benthic invertebrates (fraction)</b>					
Phytoplankton/algae	0.11 (0.01 – 0.16)	na		Many taxa have multiple feeding types; dominant feeding type for each taxa estimated using the literature. Feeding guilds were assigned to each phyla (subtidal samples only), and then percent feeding guild was assigned to each sample based on % weight. Average percent feeding guilds were calculated for all 10 samples. Because the model does not allow modeled species to have a fraction of their diet from their own model compartment, and because only one benthic invertebrate compartment was created, sediment was used as a surrogate for benthic invertebrate prey consumed by carnivores. A "detritus" compartment was not modeled because of a lack of data to generate values for such a compartment. Sediment was used as a surrogate for detritus consumed by deposit feeders. Diets were estimated assuming that carnivores consumed 100% sediment, suspension feeders consumed 30% zooplankton and 70% phytoplankton/algae, and deposit feeders consumed 100% sediment.	Literature review performed for feeding guilds of species identified in Phase 2 taxonomy samples
Zooplankton	0.05 (0.01 – 0.07)	na			
Sediment	0.84 (0.77 – 0.99)	na			
<b>Dietary Scenario 2 for benthic invertebrates (fraction)</b>					
Phytoplankton/algae	0.11 (0.01 – 0.16)	na		Many taxa have multiple feeding types; dominant feeding type for each taxa estimated using the literature. Feeding guilds were assigned to each phyla (subtidal samples only), and then percent feeding guild was assigned to each sample based on % weight. Average percent feeding guilds were calculated for all 10 samples. Because the model does not allow modeled species to eat themselves (i.e., have a fraction of their diet from their own model compartment), and because only one benthic invertebrate compartment was created, sediment was used as a surrogate for benthic invertebrate prey consumed by carnivores because of the similarities between total PCB concentrations in benthic invertebrate tissue and sediment. A "detritus" compartment was not modeled because of a lack of data to generate values for such a compartment. Sediment was used as a surrogate for detritus consumed by deposit feeders because it was assumed to have similar PCB concentrations. Assumed that carnivores consumed 50% sediment and 50% zooplankton, suspension feeders consumed 30% zooplankton and 70% phytoplankton/algae, and deposit feeders consumed 100% sediment.	Literature review performed for feeding guilds of species identified in Phase 2 taxonomy samples
Zooplankton	0.12 (0.02 – 0.17)	na			
Sediment	0.77 (0.67 – 0.97)	na			

**Table A-2-3, cont.**

PARAMETER	VALUE – MEAN (range)	SAMPLE SIZE	LOCALE OF DATA COLLECTION	NOTES	SOURCE
<b>Dungeness crab -- combined edible meat and hepatopancreas</b>					
Weight (kg ww) (all LDW)	0.423 (0.096 – 1.130)	51	LDW	Mean of all individual whole crab specimens in composites.	Phase 2 (2004, 2005) Dungeness crab data
Weight (kg ww) (modeling area M1)	0.570 (0.169 – 1.130)	20	Area T1	Mean of all individual whole crab specimens in composites.	Phase 2 (2004, 2005) Dungeness crab data
Weight (kg ww) (modeling area M2)	na	na	na	No Dungeness crabs were found in Area 2.	na
Weight (kg ww) (modeling area M3)	0.381 (0.100 – 0.780)	20	Area T3	Mean of all individual whole crab specimens in composites.	Phase 2 (2004, 2005) Dungeness crab data
Weight (kg ww) (modeling area M4)	0.231 (0.096 – 0.502)	11	Area T4	Mean of all individual whole crab specimens in composites.	Phase 2 (2004, 2005) Dungeness crab data
Lipid content (% ww) (all LDW)	2.6 (1.4 – 5.4)	7	LDW	Based on average Phase 1 and Phase 2 (2004, 2005) data. Whole body = (0.31 × hepatopancreas total PCB concentration) + (0.69 × edible meat total PCB concentration).	Phase 1 and Phase 2 (2004, 2005) Dungeness crab data
Lipid content (% ww) (modeling area M1)	3.2 (1.7 – 5.4)	3	Area T1	Based on average Phase 1 and Phase 2 (2004, 2005) data. Whole body = (0.31 × hepatopancreas total PCB concentration) + (0.69 × edible meat total PCB concentration).	Phase 1 and Phase 2 (2004, 2005) Dungeness crab data
Lipid content (% ww) (modeling area M2)	na	na	na	No Dungeness crabs were collected from Area 2.	na
Lipid content (% ww) (modeling area M3)	1.8 (1.4 – 2.2)	2	Area T3	Based on average Phase 2 (2004, 2005) data. Whole body = (0.31 × hepatopancreas total PCB concentration) + (0.69 × edible meat total PCB concentration).	Phase 2 (2004, 2005) Dungeness crab data
Lipid content (% ww) (modeling area M4)	2.4 (1.9 – 2.9)	2	Area T4	Based on average Phase 2 (2004, 2005) data. Whole body = (0.31 × hepatopancreas total PCB concentration) + (0.69 × edible meat total PCB concentration).	Phase 2 (2004, 2005) Dungeness crab data
Water content (% ww) (all LDW)	82 (78 – 85)	7	LDW	Based on average Phase 1 and Phase 2 (2004, 2005) data. Whole body = (0.31 × hepatopancreas total PCB concentration) + (0.69 × edible meat total PCB concentration).	Phase 1 and Phase 2 (2004, 2005) Dungeness crab data
Water content (% ww) (modeling area M1)	81 (79 – 84)	3	Area T1	Based on average Phase 1 and Phase 2 (2004, 2005) data. Whole body = (0.31 × hepatopancreas total PCB concentration) + (0.69 × edible meat total PCB concentration).	Phase 1 and Phase 2 (2004, 2005) Dungeness crab data
Water content (% ww) (modeling area M2)	na	na	na	No Dungeness crabs were found in Area 2.	na
Water content (% ww) (modeling area M3)	83.1 (81.3 – 84.8)	2	Area T3	Based on average Phase 2 (2004, 2005) data. Whole body = (0.31 × hepatopancreas total PCB concentration) + (0.69 × edible meat total PCB concentration).	Phase 2 (2004, 2005) Dungeness crab data

**Table A-2-3, cont.**

PARAMETER	VALUE – MEAN (range)	SAMPLE SIZE	LOCALE OF DATA COLLECTION	NOTES	SOURCE
Water content (% ww) (modeling area M4)	81.3 (77.9 – 84.7)	2	Area T4	Based on average Phase 2 (2004, 2005) data. Whole body = (0.31 × hepatopancreas total PCB concentration) + (0.69 × edible meat total PCB concentration).	Phase 2 (2004, 2005) Dungeness crab data
Fraction of porewater ventilated	0.02 (0.01 – 0.03)	na	na	Dungeness crabs live on sediment surface and ventilate some water from just above sediment surface (also stir up sediments when foraging).	Winsor et al. (1990); Gobas and Wilcockson (2003)
<b>Dietary Scenario 1 for Dungeness crab (fraction)</b>					
Benthic invertebrates	0.63 (0.42 – 0.84)	369	Grays Harbor, WA	Average % index of relative importance of identifiable prey in stomach contents classified as either phytoplankton/algae, zooplankton, benthic invertebrates or fish. Scenario 1 classifies crabs and shrimp as benthic invertebrates.	Stevens et al. (1982)
Juvenile fish	0.37 (0.16 – 0.58)	369	Grays Harbor, WA	Average % index of relative importance of identifiable prey in stomach contents classified as either phytoplankton/algae, zooplankton, benthic invertebrates, or fish. Scenario 1 classifies crabs and shrimp as benthic invertebrates.	Stevens et al. (1982)
<b>Dietary Scenario 2 for Dungeness crab (fraction)</b>					
Benthic invertebrates	0.16	369	Grays Harbor, WA	Average % of individual identifiable prey in stomach contents classified as either phytoplankton/algae, zooplankton, benthic invertebrates or fish. Scenario 2 classifies crabs and shrimp as zooplankton because no juvenile crab and shrimp prey model compartments were created and the zooplankton model compartment may provide a more realistic surrogate for the crabs and shrimps consumed as prey than the benthic invertebrate compartment.	Stevens et al. (1982)
Zooplankton	0.48	369	Grays Harbor, WA	Average % of individual identifiable prey in stomach contents classified as either phytoplankton/algae, zooplankton, benthic invertebrates or fish. Scenario 2 classifies crabs and shrimp as zooplankton because no juvenile crab and shrimp prey model compartments were created and the zooplankton model compartment may provide a more realistic surrogate for the crabs and shrimps consumed as prey than the benthic invertebrate compartment.	Stevens et al. (1982)
Juvenile fish	0.36	369	Grays Harbor, WA	Average % of individual identifiable prey in stomach contents classified as either phytoplankton/algae, zooplankton, benthic invertebrates or fish. Scenario 2 classifies crabs and shrimp as zooplankton because no juvenile crab and shrimp prey model compartments were created and the zooplankton model compartment may provide a more realistic surrogate for the crabs and shrimps consumed as prey than the benthic invertebrate compartment.	Stevens et al. (1982); Gotshall (1977)

**Table A-2-3, cont.**

PARAMETER	VALUE – MEAN (range)	SAMPLE SIZE	LOCALE OF DATA COLLECTION	NOTES	SOURCE
<b>Dietary Scenario 3 for Dungeness crab (fraction)</b>					
Benthic invertebrates	0.75	na	na	Synthesis of two open-water studies (1977) considering that, as an estuary, the LDW may have a more benthic-dominated food web (Simenstad 1983).	
Juvenile fish	0.15	na	na	Synthesis of two open-water studies (1982) considering that, as an estuary, the LDW may have a more benthic-dominated food web (Simenstad 1983).	
Sediment	0.10	na	na	Up to 50% sediment has been observed in stomach contents.	Stevens et al. (1982)
<b>Dietary Scenario 4 for Dungeness crab (fraction)</b>					
Benthic invertebrates	0.75	416	Humbolt Bay, CA, and in ocean near mouth of Mad River, CA	Average % of individual identifiable prey in stomach contents classified as phytoplankton/algae, zooplankton, benthic invertebrates and fish. Scenario 4 classifies crabs and shrimp as benthic invertebrates.	Gotshall (1977)
Juvenile fish	0.25	416	Humbolt Bay, CA, and in ocean near mouth of Mad River, CA	Average % of individual identifiable prey in stomach contents classified as phytoplankton/algae, zooplankton, benthic invertebrates and fish. Scenario 4 classifies crabs and shrimp as benthic invertebrates.	Gotshall (1977)
<b>Slender crab – combined edible meat and hepatopancreas</b>					
Weight (kg ww) (all LDW)	0.164 (0.112 – 0.260)	74	LDW	Mean of all individual specimens in composites	Phase 2 (2004, 2005) slender crab data
Weight (kg ww) (modeling area M1)	0.170 (0.120 – 0.260)	16	Area T1	Mean of all individual specimens in composites	Phase 2 (2004, 2005) slender crab data
Weight (kg ww) (modeling area M2)	0.170 (0.120 – 0.230)	40	Area T2	Mean of all individual specimens in composites	Phase 2 (2004, 2005) slender crab data
Weight (kg ww) (modeling area M3)	0.150 (0.110 – 0.210)	18	Area T3	Mean of all individual specimens in composites	Phase 2 (2004, 2005) slender crab data
Weight (kg ww) (modeling area M4)	na	na	Area T4	No slender crabs were collected from T4.	Phase 2 (2004, 2005) slender crab data
Lipid content (% ww) (all LDW)	1.1 (0.98 – 1.4)	5	LDW	Based on average Phase 2 (2004, 2005) data. Whole body = (0.31 × hepatopancreas total PCB concentration) + (0.69 × edible meat total PCB concentration).	Phase 2 (2004, 2005) slender crab data
Lipid content (% ww) (modeling area M1)	0.98	1	Area T1	Based on average Phase 2 (2004, 2005) data. Whole body = (0.31 × hepatopancreas total PCB concentration) + (0.69 × edible meat total PCB concentration).	Phase 2 (2004, 2005) slender crab data
Lipid content (% ww) (modeling area M2)	1.1 (0.98 – 1.4)	3	Area T2	Based on average Phase 2 (2004, 2005) data. Whole body = (0.31 × hepatopancreas total PCB concentration) + (0.69 × edible meat total PCB concentration).	Phase 2 (2004, 2005) slender crab data

**Table A-2-3, cont.**

PARAMETER	VALUE – MEAN (range)	SAMPLE SIZE	LOCALE OF DATA COLLECTION	NOTES	SOURCE
Lipid content (% ww) (modeling area M3)	1.0	1	Area T3	Based on average Phase 2 (2004, 2005) data. Whole body = $(0.31 \times \text{hepatopancreas total PCB concentration}) + (0.69 \times \text{edible meat total PCB concentration})$ .	Phase 2 (2004, 2005) slender crab data
Lipid content (% ww) (modeling area M4)	na	na	Area T4	No slender crabs were collected from T4.	Phase 2 (2004, 2005) slender crab data
Water content (% ww) (all LDW)	83.6 (82.5 -- 85.6)	5	LDW	Based on average Phase 2 (2004, 2005) data. Whole body = $(0.31 \times \text{hepatopancreas total PCB concentration}) + (0.69 \times \text{edible meat total PCB concentration})$ .	Phase 2 (2004, 2005) slender crab data
Water content (% ww) (modeling area M1)	82.9	1	Area T1	Based on average Phase 2 (2004, 2005) data. Whole body = $(0.31 \times \text{hepatopancreas total PCB concentration}) + (0.69 \times \text{edible meat total PCB concentration})$ .	Phase 2 (2004, 2005) slender crab data
Water content (% ww) (modeling area M2)	83.2 (82.5 -- 83.6)	3	Area T2	Based on average Phase 2 (2004, 2005) data. Whole body = $(0.31 \times \text{hepatopancreas total PCB concentration}) + (0.69 \times \text{edible meat total PCB concentration})$ .	Phase 2 (2004, 2005) slender crab data
Water content (% ww) (modeling area M3)	85.6	1	Area T3	Based on average Phase 2 (2004, 2005) data. Whole body = $(0.31 \times \text{hepatopancreas total PCB concentration}) + (0.69 \times \text{edible meat total PCB concentration})$ .	Phase 2 (2004, 2005) slender crab data
Water content (% ww) (modeling area M4)	na	na	Area T4	No slender crabs were collected from T4.	Phase 2 (2004, 2005) slender crab data
Fraction of porewater ventilated	0.02 (0.01 – 0.03)	na	na	Slender crabs live on sediment surface and ventilate some water from just above sediment surface (also stir up sediments when foraging).	Winsor et al. (1990); Gobas and Wilcockson (2003)
<b>Dietary Scenario 1 for slender crab (fraction)</b>					
Benthic invertebrates	0.99	40	Hecate Strait, BC	% of individual identifiable prey in stomach contents classified as phytoplankton/algae, zooplankton, benthic invertebrates and fish. Scenario 1 classifies crabs and shrimp as benthic invertebrates.	Bernard (1979)
Juvenile fish	0.01	40	Hecate Strait, BC	% of individual identifiable prey in stomach contents classified as phytoplankton/algae, zooplankton, benthic invertebrates and fish. Scenario 1 classifies crabs and shrimp as benthic invertebrates.	Bernard (1979)
<b>Dietary Scenario 2 for slender crab (fraction)</b>					
zooplankton	0.12	40	Hecate Strait, BC	% of individual identifiable prey in stomach contents classified as phytoplankton/algae, zooplankton, benthic invertebrates and fish. Scenario 2 classifies crabs and shrimp as zooplankton because no crab and shrimp prey model compartment was created and the zooplankton model compartment may provide a more realistic surrogate for the crabs and shrimps consumed as prey than the benthic invertebrate compartment.	Bernard (1979)



**Table A-2-3, cont.**

PARAMETER	VALUE – MEAN (range)	SAMPLE SIZE	LOCALE OF DATA COLLECTION	NOTES	SOURCE
Benthic invertebrates	0.87	40	Hecate Strait, BC	% of individual identifiable prey in stomach contents classified as phytoplankton/algae, zooplankton, benthic invertebrates and fish. Scenario 2 classifies crabs and shrimp as zooplankton because no crab and shrimp prey model compartment was created and the zooplankton model compartment may provide a more realistic surrogate for the crabs and shrimps consumed as prey than the benthic invertebrate compartment.	Bernard (1979)
Juvenile fish	0.01	40	Hecate Strait, BC	% of individual identifiable prey in stomach contents classified as phytoplankton/algae, zooplankton, benthic invertebrates and fish. Scenario 2 classifies crabs and shrimp as zooplankton because no crab and shrimp prey model compartment was created and the zooplankton model compartment may provide a more realistic surrogate for the crabs and shrimps consumed as prey than the benthic invertebrate compartment..	Bernard (2005)
<b>Dietary Scenario 3 for slender crab (fraction)</b>					
Benthic invertebrates	0.90	na	na	Synthesis of available dietary information	C. Jensen (1979);Bernard (1977)
Sediment	0.10	na	na	Based on their primarily benthic diet	
<b>Juvenile Fish</b>					
Weight (kg ww)	0.006 (0.004 – 0.007)	16	LDW, East Passage, and Blake Island	Mean of all individual ≤ 80 mm shiner surfperch specimens for which weight data were available.	Phase 2 (2004, 2005) shiner surfperch data
Lipid content (% ww)	2.5 (1.4 – 3.6)	49	LDW	Estimated assuming that lipids are lower than adult English sole and adult shiner surfperch lipids with range proportional to range observed in adults (on average +/- 45% of mean). Juvenile chinook 0.6 to 2.8 avg 1.4.	Phase 2 (2004, 2005) shiner surfperch data
Water content (% ww)	73.9 (69.6 – 77.2)	46	LDW	Calculated using Phase 2 shiner surfperch data.	Phase 2 (2004, 2005) shiner surfperch data
Fraction of porewater ventilated	0.01 (0.005-0.02)	na	na	Shiner surfperch live in water column and feed at sediment surface, English sole live on sediment surface and burrow into sediment.	
<b>Dietary Scenario 1 for juvenile fish (fraction)</b>					
Zooplankton	0.07 (0.00 – 0.15)	112	north Puget Sound, central Puget Sound, Nisqually Reach	Average % total weight of identifiable prey in juvenile English sole and adult shiner surfperch stomach contents classified as phytoplankton/algae, zooplankton, or benthic invertebrates. Scenario 1 classifies crabs and shrimp as benthic invertebrates.	Miller et al. (1977); Fresh et al. (1979); Wingert et al. (1979)
Benthic invertebrates	0.93 (0.85 – 1)	112	north Puget Sound, central Puget Sound, Nisqually Reach	Average % total weight of identifiable prey in juvenile English sole and adult shiner surfperch stomach contents classified as phytoplankton/algae, zooplankton, or benthic invertebrates. Scenario 1 classifies crabs and shrimp as benthic invertebrates.	Miller et al. (1977); Fresh et al. (1979);Wingert et al. (1979)

**Table A-2-3, cont.**

PARAMETER	VALUE – MEAN (range)	SAMPLE SIZE	LOCALE OF DATA COLLECTION	NOTES	SOURCE
<b>Dietary Scenario 2 for juvenile fish (fraction)</b>					
Zooplankton	0.17 (0.00 – 0.57)	112	north Puget Sound, central Puget Sound, Nisqually Reach	Average % total weight of identifiable prey in juvenile English sole and adult shiner surfperch stomach contents classified as phytoplankton/algae, zooplankton, or benthic invertebrates. Scenario 2 classifies crabs and shrimp as zooplankton because no crab and shrimp prey model compartment was created and the zooplankton model compartment may provide a more realistic surrogate for the crabs and shrimps consumed as prey than the benthic invertebrate compartment.	Miller et al. (1977); Fresh et al. (1979); Wingert et al. (1979)
Benthic invertebrates	0.83 (0.43 – 1)	112	north Puget Sound, central Puget Sound, Nisqually Reach	Average % total weight of identifiable prey in juvenile English sole and adult shiner surfperch stomach contents classified as phytoplankton/algae, zooplankton, or benthic invertebrates. Scenario 2 classifies crabs and shrimp as zooplankton because no crab and shrimp prey model compartment was created and the zooplankton model compartment may provide a more realistic surrogate for the crabs and shrimps consumed as prey than the benthic invertebrate compartment.	Miller et al. (1977); Fresh et al. (1979); Wingert et al. (1979)
<b>Dietary Scenario 3 for juvenile fish (fraction)</b>					
Zooplankton	0.05	na	na	Synthesis of three open-water studies (Fresh et al. 1979; Miller et al. 1977; Wingert et al. 1979), considering additional general juvenile shiner surfperch dietary information and considering that, as an estuary, the LDW may have a more benthic-dominated food web (Simenstad and Watson 1983).	Gordon (1970); Bane and Robinson (1967); Boothe (1977); Miller et al. (1977); Fresh et al. (1979); Wingert et al. (1979)
Benthic invertebrates	0.85	na	na	Synthesis of three open-water studies (Fresh et al. 1979; Miller et al. 1977; Wingert et al. 1979), considering additional general juvenile shiner surfperch dietary information and considering that, as an estuary, the LDW may have a more benthic-dominated food web (Simenstad and Watson 1983).	Gordon (1970); Bane and Robinson (1967); Boothe (1977); Miller et al. (1977); Fresh et al. (1979); Wingert et al. (1979)
Sediment	0.10	na	na	Best professional judgment in consideration of benthic diet	
<b>Shiner surfperch</b>					
Weight (kg ww) (all LDW)	0.017 (0.002 – 0.047)	458	LDW	Mean of all individual specimens in composite samples	Phase 2 (2004, 2005) shiner surfperch data
Weight (kg ww) (modeling area M1)	0.018 (0.007 – 0.047)	119	Area T1	Mean of all individual specimens in composite samples	Phase 2 (2004, 2005) shiner surfperch data
Weight (kg ww) (modeling area M2)	0.017 (0.007 – 0.040)	119	Area T2	Mean of all individual specimens in composite samples	Phase 2 (2004, 2005) shiner surfperch data

**Table A-2-3, cont.**

PARAMETER	VALUE – MEAN (range)	SAMPLE SIZE	LOCALE OF DATA COLLECTION	NOTES	SOURCE
Weight (kg ww) (modeling area M3)	0.017 (0.011 – 0.041)	120	Area T3	Mean of all individual specimens in composite samples	Phase 2 (2004, 2005) shiner surfperch data
Weight (kg ww) (modeling area M4)	0.016 (0.007 – 0.042)	100	Area T4	Mean of all individual specimens in composite samples	Phase 2 (2004, 2005) fish and crab data
Lipid content (% ww) (all LDW)	4.6 (1.6 – 6.9)	49	LDW	Mean of all composite samples	Phase 1 and Phase 2 (2004, 2005) shiner surfperch data
Lipid content (% ww) (modeling area M1)	4.1 (1.6 – 6.2)	15	Area T1	Mean of all composite samples	Phase 1 and Phase 2 (2004, 2005) shiner surfperch data
Lipid content (% ww) (modeling area M2)	4.7 (2.5 – 6.0)	12	Area T2	Mean of all composite samples	Phase 2 (2004, 2005) shiner surfperch data
Lipid content (% ww) (modeling area M3)	4.9 (3.1 – 6.9)	12	Area T3	Mean of all composite samples	Phase 2 (2004, 2005) shiner surfperch data
Lipid content (% ww) (modeling area M4)	5.0 (3.0 – 6.9)	10	Area T4	Mean of all composite samples	Phase 2 (2004, 2005) shiner surfperch data
Water content (% ww) (all LDW)	73.9 (69.6 – 77.2)	46	LDW	Mean of all composite samples	Phase 2 (2004, 2005) shiner surfperch data
Water content (% ww) (modeling area M1)	74.1 (70.4 – 76.5)	12	Area T1	Mean of all composite samples	Phase 2 (2004, 2005) shiner surfperch data
Water content (% ww) (modeling area M2)	74.4 (72.7 – 77.2)	12	Area T2	Mean of all composite samples	Phase 2 (2004, 2005) shiner surfperch data
Water content (% ww) (modeling area M3)	73.5 (69.6 – 77.0)	12	Area T3	Mean of all composite samples	Phase 2 (2004, 2005) shiner surfperch data
Water content (% ww) (modeling area M4)	73.6 (69.7 – 77.2)	10	Area T4	Mean of all composite samples	Phase 2 (2004, 2005) shiner surfperch data
Fraction of porewater ventilated	0.01 (0.005 – 0.02)	na	na	Shiner surfperch live in water column and feed at sediment surface.	Gobas and Wilcockson (1977)
<b>Dietary Scenario 1 for shiner surfperch (fraction)</b>					
Zooplankton	0.14 0.00 – 0.38)	65	north Puget Sound, central Puget Sound, Nisqually Reach	Average % total weight of identifiable prey in stomach contents classified as phytoplankton/algae, zooplankton, or benthic invertebrates. Scenario 1 classifies crabs and shrimp as benthic invertebrates.	Miller et al. (1977); Fresh et al. (1979); Wingert et al. (1979)

**Table A-2-3, cont.**

PARAMETER	VALUE – MEAN (range)	SAMPLE SIZE	LOCALE OF DATA COLLECTION	NOTES	SOURCE
Benthic invertebrates	0.86 (0.62 – 0.95)	65	north Puget Sound, central Puget Sound, Nisqually Reach	Average % total weight of identifiable prey in stomach contents classified as phytoplankton/algae, zooplankton, or benthic invertebrates. Scenario 1 classifies crabs and shrimp as benthic invertebrates.	Miller et al. (1977); Fresh et al. (1979); Wingert et al. (1979)
<b>Dietary Scenario 2 for shiner surfperch (fraction)</b>					
Zooplankton	0.21 (0.00 – 0.57)	65	north Puget Sound, central Puget Sound, Nisqually Reach	Average % total weight of identifiable prey in stomach contents classified as phytoplankton/algae, zooplankton, or benthic invertebrates. Scenario 2 classifies crabs and shrimp as zooplankton because no crab and shrimp prey model compartment was created and the zooplankton model compartment may provide a more realistic surrogate for the crabs and shrimps consumed as prey than the benthic invertebrate compartment.	Miller et al. (1977); Fresh et al. (1979); Wingert et al. (1979)
Benthic invertebrates	0.79 (0.43 – 1)	65	north Puget Sound, central Puget Sound, Nisqually Reach	Average % total weight of identifiable prey in stomach contents classified as phytoplankton/algae, zooplankton, or benthic invertebrates. Scenario 2 classifies crabs and shrimp as zooplankton because no crab and shrimp prey model compartment was created and the zooplankton model compartment may provide a more realistic surrogate for the crabs and shrimps consumed as prey than the benthic invertebrate compartment.	Miller et al. (1977); Fresh et al. (1979); Wingert et al. (1979)
<b>Dietary Scenario 3 for shiner surfperch (fraction)</b>					
Zooplankton	0.10	na	na	Synthesis of three open-water studies (Fresh et al. 1979; Miller et al. 1977; Wingert et al. 1979), considering additional general juvenile shiner surfperch dietary information and considering that, as an estuary, the LDW may have a more benthic-dominated food web (Simenstad and Watson 1983).	Miller et al. (1977); Fresh et al. (1979); Wingert et al. (1979)
Benthic invertebrates	0.80	na	na	Synthesis of three open-water studies (Fresh et al. 1979; Miller et al. 1977; Wingert et al. 1979), considering additional general juvenile shiner surfperch dietary information and considering that, as an estuary, the LDW may have a more benthic-dominated food web (Simenstad and Watson 1983).	Miller et al. (1977); Fresh et al. (1979); Wingert et al. (1979)
Sediment	0.10	na	na	Best professional judgment in consideration of benthic diet	
<b>English sole</b>					
Weight (kg ww) (all LDW)	0.198 (0.073 – 0.600)	245	LDW	Mean of all individual specimens in whole body composite samples	Phase 2 (2004, 2005) English sole data
Weight (kg ww) (modeling area M1)	0.171 (0.076 – 0.500)	67	Area T1	Mean of all individual specimens in whole body composite samples	Phase 2 (2004, 2005) English sole data
Weight (kg ww) (modeling area M2)	0.189 (0.088 – 0.525)	67	Area T2	Mean of all individual specimens in whole body composite samples	Phase 2 (2004, 2005) English sole data

**Table A-2-3, cont.**

PARAMETER	VALUE – MEAN (range)	SAMPLE SIZE	LOCALE OF DATA COLLECTION	NOTES	SOURCE
Weight (kg ww) (modeling area M3)	0.216 (0.079 – 0.404)	70	Area T3	Mean of all individual specimens in whole body composite samples	Phase 2 (2004, 2005) English sole data
Weight (kg ww) (modeling area M4)	0.236 (0.073 – 0.600)	68	Area T4	Mean of all individual specimens in whole body composite samples	Phase 2 (2004, 2005) English sole data
Lipid content (% ww) (all LDW)	5.5 (2.6 – 8.7)	42	LDW	Mean of all composite samples	Phase 2 (2004, 2005) English sole data
Lipid content (% ww) (modeling area M1)	5.2 (3.1 – 6.8)	12	Area T1	Mean of all composite samples	Phase 2 (2004, 2005) English sole data
Lipid content (% ww) (modeling area M2)	6.4 (4.9 – 8.7)	12	Area T2	Mean of all composite samples	Phase 2 (2004, 2005) English sole data
Lipid content (% ww) (modeling area M3)	5.0 (2.6 – 7.5)	12	Area T3	Mean of all composite samples	Phase 2 (2004, 2005) English sole data
Lipid content (% ww) (modeling area M4)	5.4 (3.9 – 6.3)	6	Area T4	Mean of all composite samples	Phase 2 (2004, 2005) English sole data
Water content (% ww) (all LDW)	75.0 (71.0 – 79.0)	42	LDW	Mean of all composite samples	Phase 2 (2004, 2005) English sole data
Water content (% ww) (modeling area M1)	75.5 (73.4 – 79.0)	12	Area T1	Mean of all composite samples	Phase 2 (2004, 2005) English sole data
Water content (% ww) (modeling area M2)	73.9 (71.4 – 76.9)	12	Area T2	Mean of all composite samples	Phase 2 (2004, 2005) English sole data
Water content (% ww) (modeling area M3)	75.4 (73.4 – 78.8)	12	Area T3	Mean of all composite samples	Phase 2 (2004, 2005) English sole data
Water content (% ww) (modeling area M4)	75.0 (74.0 – 76.2)	6	Area T4	Mean of all composite samples	Phase 2 (2004, 2005) English sole data
Fraction of porewater ventilated	0.1 (0.05 – 0.2)	na	LDW	English sole feed at the sediment surface and burrow into the sediment.	(Gobas and Wilcockson 2003)
<b>Dietary Scenario 1 for English sole (fraction)</b>					
Phytoplankton/algae	0.08 (0.05 – 0.10)	135	central Puget Sound, Nisqually Reach	Average % total weight of identifiable prey in stomach contents classified as either phytoplankton/algae, zooplankton, or benthic invertebrates. Scenario 1 classifies crabs and shrimp as benthic invertebrates.	Fresh et al. (1979); Wingert et al. (1979)
Benthic invertebrates	0.92 (0.90 – 0.95)	135	central Puget Sound, Nisqually Reach	Average % total weight of identifiable prey in stomach contents classified as either phytoplankton/algae, zooplankton, or benthic invertebrates. Scenario 1 classifies crabs and shrimp as benthic invertebrates.	Fresh et al. (1979); Wingert et al. (1979)

**Table A-2-3, cont.**

PARAMETER	VALUE – MEAN (range)	SAMPLE SIZE	LOCALE OF DATA COLLECTION	NOTES	SOURCE
<b>Dietary Scenario 2 for English sole (fraction)</b>					
Phytoplankton/algae	0.07 (0.05 – 0.10)	135	central Puget Sound, Nisqually Reach	Average % total weight of identifiable prey in stomach contents classified as either phytoplankton/algae, zooplankton, or benthic invertebrates. Scenario 2 classifies crabs and shrimp as zooplankton because no crab and shrimp prey model compartment was created and the zooplankton model compartment may provide a more realistic surrogate for the crabs and shrimps consumed as prey than the benthic invertebrate compartment.	Fresh et al. (1979); Wingert et al. (1979)
Zooplankton	0.05 (0.00 – 0.09)	135	central Puget Sound, Nisqually Reach	Average % total weight of identifiable prey in stomach contents classified as phytoplankton/algae, zooplankton, or benthic invertebrates. Scenario 2 classifies crabs and shrimp as zooplankton because no crab and shrimp prey model compartment was created and the zooplankton model compartment may provide a more realistic surrogate for the crabs and shrimps consumed as prey than the benthic invertebrate compartment.	Fresh et al. (1979); Wingert et al. (1979)
Benthic invertebrates	0.88 (0.86 – 0.90)	135	central Puget Sound, Nisqually Reach	Average % total weight of identifiable prey in stomach contents classified as either phytoplankton/algae, zooplankton, or benthic invertebrates. Scenario 2 classifies crabs and shrimp as zooplankton because no crab and shrimp prey model compartment was created and the zooplankton model compartment may provide a more realistic surrogate for the crabs and shrimps consumed as prey than the benthic invertebrate compartment.	Fresh et al. (1979); Wingert et al. (1979)
<b>Dietary Scenario 3 for English sole (fraction)</b>					
Benthic invertebrates	0.90	na	na	Synthesis of two open-water studies (Fresh et al. 1979; Wingert et al. 1979) considering that, as an estuary, the LDW may have a more benthic dominated food web (Simenstad and Watson 1983)	Fresh et al. (1979); Wingert et al. (1979)
Sediment	0.10	na	na	Best professional judgment in consideration of benthic diet	Fresh et al. (1979); Wingert et al. (1979)
<b>Pacific staghorn sculpin</b>					
Weight (kg ww) (all LDW)	0.060 (0.013 – 0.227)	272	LDW	Mean of all individual specimens in composite samples	Phase 2 (2004, 2005) Pacific staghorn sculpin data
Weight (kg ww) (modeling area M1)	0.065 (0.018 – 0.180)	67	Area T1	Mean of all individual specimens in composite samples	Phase 2 (2004, 2005) Pacific staghorn sculpin data
Weight (kg ww) (modeling area M2)	0.067 (0.020 – 0.227)	67	Area T2	Mean of all individual specimens in composite samples	Phase 2 (2004, 2005) Pacific staghorn sculpin data

**Table A-2-3, cont.**

PARAMETER	VALUE – MEAN (range)	SAMPLE SIZE	LOCALE OF DATA COLLECTION	NOTES	SOURCE
Weight (kg ww) (modeling area M3)	0.058 (0.016 – 0.164)	70	Area T3	Mean of all individual specimens in composite samples	Phase 2 (2004, 2005) Pacific staghorn sculpin data
Weight (kg ww) (modeling area M4)	0.050 (0.013 – 0.168)	68	Area T4	Mean of all individual specimens in composite samples	Phase 2 (2004, 2005) Pacific staghorn sculpin data
Lipid content (% ww) (all LDW)	2.1 (1.2 – 2.7)	28	LDW	Mean of all composite samples	Phase 2 (2004, 2005) Pacific staghorn sculpin data
Lipid content (% ww) (modeling area M1)	2.2 (1.8 – 2.4)	7	Area T1	Mean of all composite samples	Phase 2 (2004, 2005) Pacific staghorn sculpin data
Lipid content (% ww) (modeling area M2)	2.2 (1.8 – 2.7)	7	Area T2	Mean of all composite samples	Phase 2 (2004, 2005) Pacific staghorn sculpin data
Lipid content (% ww) (modeling area M3)	1.8 (1.3 – 2.1)	7	Area T3	Mean of all composite samples	Phase 2 (2004, 2005) Pacific staghorn sculpin data
Lipid content (% ww) (modeling area M4)	1.9 (1.2 – 2.5)	7	Area T4	Mean of all composite samples	Phase 2 (2004, 2005) Pacific staghorn sculpin data
Water content (% ww) (all LDW)	79.0 (78.0 – 80.5)	28	LDW	Mean of all composite samples	Phase 2 (2004, 2005) Pacific staghorn sculpin data
Water content (% ww) (modeling area M1)	78.6 (78.0 – 79.6)	7	Area T1	Mean of all composite samples	Phase 2 (2004, 2005) Pacific staghorn sculpin data
Water content (% ww) (modeling area M2)	78.9 (78.0 – 80.3)	7	Area T2	Mean of all composite samples	Phase 2 (2004, 2005) Pacific staghorn sculpin data
Water content (% ww) (modeling area M3)	79.2 (78.9 – 79.7)	7	Area T3	Mean of all composite samples	Phase 2 (2004, 2005) Pacific staghorn sculpin data
Water content (% ww) (modeling area M4)	79.2 (78.3 – 80.5)	7	Area T4	Mean of all composite samples	Phase 2 (2004, 2005) Pacific staghorn sculpin data
Fraction of porewater ventilated	0.05 (0.02 – 0.1)	na	LDW	Pacific staghorn sculpin feed at the sediment surface and burrow into the sediment.	

**Table A-2-3, cont.**

PARAMETER	VALUE – MEAN (range)	SAMPLE SIZE	LOCALE OF DATA COLLECTION	NOTES	SOURCE
<b>Dietary Scenario 1 for Pacific staghorn sculpin (fraction)</b>					
Benthic invertebrates	0.56 (0.32 – 0.83)	133	north Puget Sound, central Puget Sound, Nisqually Reach	Average % total weight of identifiable prey in stomach contents classified as either phytoplankton/algae, zooplankton, benthic invertebrates, or fish. Scenario 1 classifies crabs and shrimp as benthic invertebrates.	Miller et al. (1977); Fresh et al. (1979); Wingert et al. (1979)
Juvenile fish	0.44 (0.17 – 0.68)	133	north Puget Sound, central Puget Sound, Nisqually Reach	Average % total weight of identifiable prey in stomach contents classified as either phytoplankton/algae, zooplankton, benthic invertebrates, or fish. Scenario 1 classifies crabs and shrimp as benthic invertebrates.	Miller et al. (1977); Fresh et al. (1979); Wingert et al. (1979)
<b>Dietary Scenario 2 for Pacific staghorn sculpin (fraction)</b>					
Zooplankton	0.37 (0.29 – 0.50)	133	north Puget Sound, central Puget Sound, Nisqually Reach	Average % total weight of identifiable prey in stomach contents classified as either phytoplankton/algae, zooplankton, benthic invertebrates, or fish. Scenario 2 classifies crabs and shrimp as zooplankton because no crab and shrimp prey model compartment was created and the zooplankton model compartment may provide a more realistic surrogate for the crabs and shrimps consumed as prey than the benthic invertebrate compartment.	Miller et al. (1977); Fresh et al. (1979); Wingert et al. (1979)
Benthic invertebrates	0.19 (0.04 – 0.32)	133	north Puget Sound, central Puget Sound, Nisqually Reach	Average % total weight of identifiable prey in stomach contents classified as either phytoplankton/algae, zooplankton, benthic invertebrates, or fish. Scenario 2 classifies crabs and shrimp as zooplankton because no crab and shrimp prey model compartment was created and the zooplankton model compartment may provide a more realistic surrogate for the crabs and shrimps consumed as prey than the benthic invertebrate compartment.	Miller et al. (1977); Fresh et al. (1979); Wingert et al. (1979)
Juvenile fish	0.44 (0.17 – 0.68)	133	north Puget Sound, central Puget Sound, Nisqually Reach	Average % total weight of identifiable prey in stomach contents classified as either phytoplankton/algae, zooplankton, benthic invertebrates, or fish. Scenario 2 classifies crabs and shrimp as zooplankton because no crab and shrimp prey model compartment was created and the zooplankton model compartment may provide a more realistic surrogate for the crabs and shrimps consumed as prey than the benthic invertebrate compartment.	Miller et al. (1977); Fresh et al. (1979); Wingert et al. (1979)
<b>Dietary Scenario 3 for Pacific staghorn sculpin (fraction)</b>					
Zooplankton	0.25	na	na	Synthesis of three open-water studies (Fresh et al. 1979; Miller et al. 1977; Wingert et al. 1979) considering that, as an estuary, the LDW may have a more benthic dominated food web (Simenstad and Watson 1983); but crabs and shrimp have less sediment exposure than infaunal benthic invertebrates, therefore 25% of diet considered zooplankton.	Miller et al. (1977); Fresh et al. (1979); Wingert et al. (1979)



**Table A-2-3, cont.**

PARAMETER	VALUE – MEAN (range)	SAMPLE SIZE	LOCALE OF DATA COLLECTION	NOTES	SOURCE
Benthic invertebrates	0.50	na	na	Synthesis of three open-water studies (Fresh et al. 1979; Miller et al. 1977; Wingert et al. 1979) considering that, as an estuary, the LDW may have a more benthic-dominated food web (Simenstad and Watson 1983); but crabs and shrimp have less sediment exposure than infaunal benthic invertebrates, therefore 25% of diet considered zooplankton.	Miller et al. (1977); Fresh et al. (1979); Wingert et al. (1979)
Juvenile fish	0.15	na	na	Synthesis of three open-water studies (Fresh et al. 1979; Miller et al. 1977; Wingert et al. 1979) considering that, as an estuary, the LDW may have a more benthic-dominated food web (Simenstad and Watson 1983).	Miller et al. (1977); Fresh et al. (1979); Wingert et al. (1979)
Sediment	0.10	na	na	Best professional judgment in consideration of benthic diet.	Miller et al. (1977); Fresh et al. (1979); Wingert et al. (1979)

LDW – Lower Duwamish Waterway

na – not available

NLOC – non-lipid organic carbon

nr – not reported

tbd – to be determined

ww – wet weight

<sup>a</sup> Based on the size of the mesh size, these samples would have consisted mostly of zooplankton and chain-forming phytoplankton.

**Table A-2-4. Total PCB tissue concentrations in fish and invertebrate species to be used to evaluate model performance**

AREA MODELED	TOTAL PCB TISSUE CONCENTRATION (µg/kg ww)		No. OF COMPOSITE SAMPLES	NOTES	SOURCE
	AVERAGE	RANGE			
<b>Benthic invertebrates</b>					
All LDW	170	1) 60 – 1,400 2) 150 – 200	20	Average was estimated using surface sediment total PCBs SWAC of 250 µg/kg dw for the entire LDW and the tissue-sediment regression derived from 20 co-located benthic invertebrate tissue and surface sediment samples. Range 1 is based on 10 intertidal and 10 subtidal samples collected throughout the LDW. Range 2 is based on the 95% confidence interval on the mean (for the regression-estimated average tissue concentration).	Phase 2 (2004) benthic data (for range 1 data) Average and range 2 calculated from Phase 2 (2004) benthic tissue data and co-located sediment data, and Phase 1 and Phase 2 sediment data
Modeling area M1	180	1) 66 – 310 2) 150 – 210	6	Average was estimated using sediment total PCBs SWAC of 280 µg/kg dw in area M1 and the tissue-sediment regression derived from 20 co-located benthic invertebrate tissue and surface sediment samples. Range 1 is based on 3 intertidal and 3 subtidal samples collected in area M1. Range 2 is based on the 95% confidence interval on the mean (for the regression-estimated average tissue concentration).	Phase 2 (2004) benthic data (for range 1 data) Average and range 2 calculated from Phase 2 (2004) benthic tissue data and co-located sediment data, and Phase 1 and Phase 2 sediment data
Modeling area M2	150	1) 100 – 1,400 2) 130 – 170	6	Average was estimated using sediment total PCBs SWAC of 160 µg/kg dw in area M2 and the tissue-sediment regression derived from 20 co-located benthic invertebrate tissue and surface sediment samples. Range 1 is based on 3 intertidal and 3 subtidal samples collected in area M2. Range 2 is based on the 95% confidence interval on the mean (for the regression-estimated average tissue concentration).	Phase 2 (2004) benthic data (for range 1 data) Average and range 2 calculated from Phase 2 (2004) benthic tissue data and co-located sediment data, and Phase 1 and Phase 2 sediment data

Table A-2-4, cont.

AREA MODELED	TOTAL PCB TISSUE CONCENTRATION (µg/kg ww)		NO. OF COMPOSITE SAMPLES	NOTES	SOURCE
	AVERAGE	RANGE			
Modeling area M3	220	1) 99 – 1,400 2) 180 – 270	4	Average was estimated using sediment total PCBs SWAC of 470 µg/kg dw in area M3 and the tissue-sediment regression derived from 20 co-located benthic invertebrate tissue and surface sediment samples. Range 1 is based on 2 intertidal and 2 subtidal samples collected in area M3. Range 2 is based on the 95% confidence interval on the mean (for the regression-estimated average tissue concentration).	Phase 2 (2004) benthic data (for range 1 data) Average and range 2 calculated from Phase 2 (2004) benthic tissue data and co-located sediment data, and Phase 1 and Phase 2 sediment data
Modeling area M4	92	1) 60 – 110 2) 80 – 100	4	Average was estimated using sediment total PCBs SWAC of 41 µg/kg dw in area M4 and the tissue-sediment regression derived from 20 co-located benthic invertebrate tissue and surface sediment samples. Range 1 is based on 2 intertidal and 2 subtidal samples collected in area M4. Range 2 is based on the 95% confidence interval on the mean (for the regression-estimated average tissue concentration).	Phase 2 (2004) benthic data (for range 1 data) Average and range 2 calculated from Phase 2 (2004) benthic tissue data and co-located sediment data, and Phase 1 and Phase 2 sediment data
<b>Dungeness crab – combined edible meat and hepatopancreas</b>					
All LDW	980	420 – 1,900	7	Based on average Phase 1 and Phase 2 (2004, 2005) data. Whole body = (0.31 × hepatopancreas total PCB concentration) + (0.69 × edible meat total PCB concentration)	Phase 1 and Phase 2 (2004, 2005) Dungeness crab data
Modeling area M1	830	450 – 1,400	3	Based on average Phase 1 and Phase 2 (2004, 2005) data. Whole body = (0.31 × hepatopancreas total PCB concentration) + (0.69 × edible meat total PCB concentration)	Phase 1 and Phase 2 (2004, 2005) Dungeness crab data
Modeling area M3	1,000	420 – 1,600	2	Based on average Phase 2 (2004, 2005) data. Whole body = (0.31 × hepatopancreas total PCB concentration) + (0.69 × edible meat total PCB concentration)	Phase 2 (2004, 2005) Dungeness crab data

Table A-2-4, cont.

AREA MODELED	TOTAL PCB TISSUE CONCENTRATION (µg/kg ww)		NO. OF COMPOSITE SAMPLES	NOTES	SOURCE
	AVERAGE	RANGE			
Modeling area M4	1,200	420 – 1,900	2	Based on average Phase 2 (2004, 2005) data. Whole body = (0.31 × hepatopancreas total PCB concentration) + (0.69 × edible meat total PCB concentration)	Phase 2 (2004, 2005) Dungeness crab data
<b>Slender crab – combined edible meat and hepatopancreas</b>					
All LDW	620	250 – 800	5	Based on average Phase 2 (2004, 2005) data. Whole body = (0.31 × hepatopancreas total PCB concentration) + (0.69 × edible meat total PCB concentration)	Phase 2 (2004, 2005) slender crab data
Modeling area M1	650	na	1	Based on average Phase 2 (2004, 2005) data. Whole body = (0.31 × hepatopancreas total PCB concentration) + (0.69 × edible meat total PCB concentration)	Phase 2 (2004, 2005) slender crab data
Modeling area M2	600	250 – 800	3	Based on average Phase 2 (2004, 2005) data. Whole body = (0.31 × hepatopancreas total PCB concentration) + (0.69 × edible meat total PCB concentration)	Phase 2 (2004, 2005) slender crab data
Modeling area M3	630	na	1	Based on average Phase 2 (2004, 2005) data. Whole body = (0.31 × hepatopancreas total PCB concentration) + (0.69 × edible meat total PCB concentration)	Phase 2 (2004, 2005) slender crab data
<b>Shiner surfperch – whole body</b>					
All LDW	1,800	350 – 18,000	51		Phase 1 and Phase 2 (2004, 2005) shiner surfperch data
Modeling area M1	970	350 – 1,800	15		Phase 1 and Phase 2 (2004, 2005) shiner surfperch data
Modeling area M2	2,800	660 – 18,000	12	Average is 1,400 µg/kg ww if the 2004 T2E 18,000-µg/kg ww sample is excluded	Phase 2 (2004, 2005) shiner surfperch data
Modeling area M3	2,700	700 – 8,800	12		Phase 2 (2004, 2005) shiner surfperch data

**Table A-2-4, cont.**

AREA MODELED	TOTAL PCB TISSUE CONCENTRATION (µg/kg ww)		NO. OF COMPOSITE SAMPLES	NOTES	SOURCE
	AVERAGE	RANGE			
Modeling area M4	840	540 – 2,100	12		Phase 1 and Phase 2 (2004, 2005) shiner surfperch data
<b>English sole – whole body</b>					
All LDW	2,300	610 – 4,700	42		Phase 2 (2004, 2005) English sole data
Modeling area M1	2,600	1,100 – 4,700	12		Phase 2 (2004, 2005) English sole data
Modeling area M2	2,900	1,600 – 4,200	12		Phase 2 (2004, 2005) English sole data
Modeling area M3	2,000	610 – 4,300	12		Phase 2 (2004, 2005) English sole data
Modeling area M4	1,400	910 – 1,800	6		Phase 2 (2004, 2005) English sole data
<b>Pacific staghorn sculpin – whole body</b>					
All LDW	900	430 – 2,800	28		Phase 2 (2004, 2005) sculpin data
Modeling area M1	720	580 – 860	7		Phase 2 (2004, 2005) sculpin data
Modeling area M2	750	620 – 1,300	7		Phase 2 (2004, 2005) sculpin data
Modeling area M3	1,400	590 – 2,800	7		Phase 2 (2004, 2005) sculpin data
Modeling area M4	730	430 – 1,300	7		Phase 2 (2004, 2005) sculpin data

LDW – Lower Duwamish Waterway

na – not applicable

PCB – polychlorinated biphenyl

Paired English sole fillet and remainder samples were used to derive concentrations of parameters in “whole-body” samples. Ten whole-body concentrations of lipid content, water content, and total PCB concentrations were estimated for English sole based on the relative weights and analyte concentrations in corresponding skin-on fillet and remainder tissues collected in 2005. These samples were collected to calculate whole-body PCB concentrations as specified in the QAPP (Windward 2005f) and data report (Windward 2006 in prep).

Estimates of lipid content, water content, and total PCB concentrations were calculated for “whole body” crabs by combining the concentration in each composite hepatopancreas sample with concentrations in the corresponding edible meat composite samples (one or more samples) that were collected from the same crabs. Therefore, a single whole-body crab concentration was calculated for each of the 12 hepatopancreas samples in Table A-2-2.<sup>1</sup> Whole-body concentrations were calculated assuming 69% (by weight) edible meat and 31% hepatopancreas, based on the relative weights of these tissues in a 16.6-cm Dungeness crab dissected by Windward in 2004.

Juvenile fish were modeled using shiner surfperch and English sole data to estimate input parameters. Juvenile shiner surfperch and juvenile starry flounder were the most abundant small fish (< 100 mm) captured in trawls during Phase 2 sampling conducted in late summer (Windward 2005c, 2006 in prep). For example, they represented 54 and 30% of the non-target catch, respectively, in the 2004 sampling event and 40 and 42%, respectively, in the 2005 sampling event. Thus, these species are likely prey for Pacific staghorn sculpin and crabs in the LDW. Because data for juvenile starry flounder and juvenile shiner surfperch were not available (with the exception of limited weight data), data from composite samples of adult shiner surfperch and English sole<sup>2</sup> were used. As noted in section A.2.3 and Table A-2-3, empirical weights for juvenile shiner surf perch were used as the basis for estimates of juvenile fish weight for the model.

All data for a given species were combined to determine the average, standard deviation, and range of total lipids, total solids, and total PCB concentrations. Averaging in this way reasonably represents populations foraging throughout the LDW because, for each species, the number of samples is fairly uniform throughout the LDW (Table A-2-5). The relatively even distribution of the data is demonstrated by dividing the LDW into four sections (M1 to M4), and summing the available tissue composite samples per section.

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<sup>1</sup> A total of 11 moisture content values were calculated because total solids data were not available for the Phase 1 hepatopancreas and edible meat samples.

<sup>2</sup> English sole and starry flounder are closely related (and produce viable offspring) so English sole are a reasonable surrogate for starry flounder. The English sole data includes three starry flounder composite samples from tissue sampling area T4.

**Table A-2-5. Number of composite tissue samples available from each LDW modeling area**

SPECIES	AREA	NUMBER OF COMPOSITE TISSUE SAMPLES
Benthic invertebrates	M1	6
	M2	6
	M3	4
	M4	4
English sole (whole body and estimated whole body <sup>a</sup> )	M1	12
	M2	12
	M3	12
	M4	6
Pacific staghorn sculpin (whole body)	M1	7
	M2	7
	M3	7
	M4	7
Shiner surfperch (whole body)	M1	15
	M2	12
	M3	12
	M4	12(10) <sup>c</sup>
Dungeness crab (estimated whole body <sup>b</sup> )	M1	3
	M2	0
	M3	2
	M4	2
Slender crab (estimated whole body <sup>b</sup> )	M1	1
	M2	3
	M3	1
	M4	0

<sup>a</sup> Concentrations in 3 English sole whole-body composite samples in modeling areas M1, M2, and M3, and one composite sample in modeling area M4 were calculated as the weighted average of fillet and remainder composite samples.

<sup>b</sup> All whole-body crab concentrations were estimated as the weighted average of edible meat and hepatopancreas composite samples from the same crabs.

<sup>c</sup> Twelve composite samples were used to calculate total PCB concentrations; however, percent lipids and percent solids data were not available for the two Phase 1 samples.

Data from Phase 1 and Phase 2 (2004 and 2005) were combined to derive model input values. Percent lipids and percent solids whole-body tissue data from Phase 1 and Phase 2 (2004 and 2005) datasets are presented in Table A-2-6. One-way analysis of variance (alpha = 0.05) revealed statistically significant differences between 2004 and 2005 sampling events for Pacific staghorn sculpin percent lipids, and for shiner surfperch weight and percent solids. Pacific staghorn sculpin lipids were on average 24% higher in 2004 samples than 2005. However, 24 Pacific staghorn sculpin composite samples were collected in 2004 versus four samples collected in 2005.

When the four 2005 samples are compared using a paired t-test to the 2004 samples from the same subareas,<sup>3</sup> no statistically significant differences were observed.

Average shiner surfperch weight was 3 g higher in 2005 than in 2004 and average shiner surfperch total solids were 1.8% higher in 2005 than 2004. Because inter-annual variability in these parameters is expected, the variability in parameters between sampling events was considered to be representative of the variability in the data over the time period for which FWM predictions may apply. Variables that had statistically significant differences among sampling events will be given additional consideration during the calibration process, if needed. For the modeling-area scale, the average, standard deviation, and range for lipids and total solids were determined for each individual modeling area, whereas, weight parameter values were based on LDW-wide data (Table A-2-3).

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<sup>3</sup> Each 2005 composite sample was paired with the 2004 composite sample from the same subarea.



**Table A-2-6. Weight, percent lipids, and percent solids data used in the FWM for whole-body fish and crabs**

SPECIES	ANALYTE (UNITS)	PHASE 1		PHASE 2 (2004)		PHASE 2 (2005)	
		N	AVERAGE (SD)	N	AVERAGE (SD)	N	AVERAGE (SD)
English sole <sup>a</sup>	weight (g) <sup>b</sup>	nd	nd	140 <sup>b</sup>	194 (90.6)	105 <sup>b</sup>	204 (115)
	lipids (%)	nd	nd	21	5.8 (1.4)	21	5.22 (1.17)
	total solids (%)	nd	nd	21	25.0 (1.55)	21	25.1 (2.01)
Pacific staghorn sculpin	weight (g) <sup>b</sup>	nd	nd	232 <sup>b</sup>	60.8 (32.7)	40 <sup>b</sup>	55 (46)
	lipids (%)	nd	nd	24	2.1 (0.32)	4	1.65 (0.469)
	total solids (%)	nd	nd	24	21.1 (0.579)	4	20.6 (0.612)
Shiner surfperch	weight (g) <sup>b</sup>	nd	nd	238 <sup>b</sup>	16 (5.7)	220 <sup>b</sup>	19 (6.8)
	lipids (%)	3	2.8 (1.2)	24	3.9 (1.1)	22	5.74 (0.692)
	total solids (%)	nd	nd	24	24.7 (1.27)	22	27.6 (1.54)
Dungeness crab <sup>c</sup>	weight (g) <sup>b</sup>	nd	nd	36 <sup>b</sup>	470 (280)	15 <sup>b</sup>	302 (175)
	lipids (%)	1	5.4 (na)	3	2.3 (0.65)	3	1.98 (0.646)
	total solids (%)	nd	nd	3	19.0 (2.91)	3	17.0 (2.91)
Slender crab <sup>c</sup>	weight (g) <sup>b</sup>	nd	nd	64 <sup>b</sup>	160 (29)	10 <sup>b</sup>	190 (35.7)
	lipids (%)	nd	nd	4	1.1 (0.17)	1	0.980 (na)
	total solids (%)	nd	nd	4	16.1 (1.16)	1	17.5 (na)

<sup>a</sup> Ten of the 21 Phase 2 English sole composite samples were calculated as the weighted average of fillet and remainder composite samples.

<sup>b</sup> Weights were calculated using data for individual specimens rather than composites.

<sup>c</sup> Each whole-body crab lipid and total solids concentration was estimated by combining the concentration in the composite hepatopancreas sample with concentrations in the corresponding edible meat composite samples (one or more samples) that were collected from the same crabs. Therefore, a single whole-body crab concentration for each parameter was calculated for each composite hepatopancreas sample. Whole-body concentrations were calculated assuming 69% (by weight) edible meat and 31% hepatopancreas, based on the relative weight of these tissues in a 16.6-cm Dungeness crab dissected by Windward in 2004.

nd – no data

na – not applicable

SD – standard deviation

N – number of composite samples

#### A.2.4 Relationship between co-located benthic invertebrate tissue and surface sediment total PCB concentrations

Benthic invertebrate tissue and co-located surface sediment samples were collected from 20 locations in the LDW (10 intertidal locations and 10 subtidal locations). These locations were selected to provide a range of total PCB concentrations and spatial coverage throughout the LDW. These data were generated to evaluate whether there was a relationship between chemical concentrations in benthic invertebrate tissue and co-located sediment that could be applied in the FWM and exposure assessments for the risk assessments.

Linear least-squares regression was used to model the relationship between total PCB concentrations<sup>4</sup> in benthic invertebrate tissue and co-located sediment. The relationship between sediment and tissue was not linear and the residuals from a linear fit increased with the total PCB concentration in sediment. The log-log relationship provided a reasonable linear fit with homogeneous residuals (Figure A-2-1)<sup>5</sup> except for two extreme points (locations B5a-1 and B8a). Location B5a-1 had a low-moderate sediment PCB concentration and a very high tissue concentration. The sediment had very low organic carbon content, so this point was not extreme when the data were normalized. However, the normalized sediment relationship with tissue did not provide a good fit. Location B8a had a very high sediment concentration, but the tissue concentration was higher than would be predicted from a linear (log-log) relationship. This point was exerting undue influence on the regression estimates, and it was far higher than the concentrations for which tissue estimates were produced. The R<sup>2</sup> value with the outliers included was 0.51. Without these two points, the regression provided a good fit to the data in the range for which tissue concentrations will be predicted. The R<sup>2</sup> value with the outliers removed was 0.74. The regression parameters were estimated with full reporting-limit concentrations for the two non-detect samples.<sup>6</sup>

Figure A-2-1 displays the log-log linear relationship between PCB concentrations in co-located benthic invertebrate tissue and sediment. The equation for the line with outliers removed is presented as Equation A-2-1.

$$\log_{10}[\text{tissue}] = 1.40 + 0.35 \times \log_{10}[\text{sediment}] \quad \text{Equation A-2-1}$$

Where:

[tissue] = total PCB concentration (µg/kg ww) in benthic invertebrate tissue  
[sediment] = total PCB concentration (µg/kg dw) in sediment

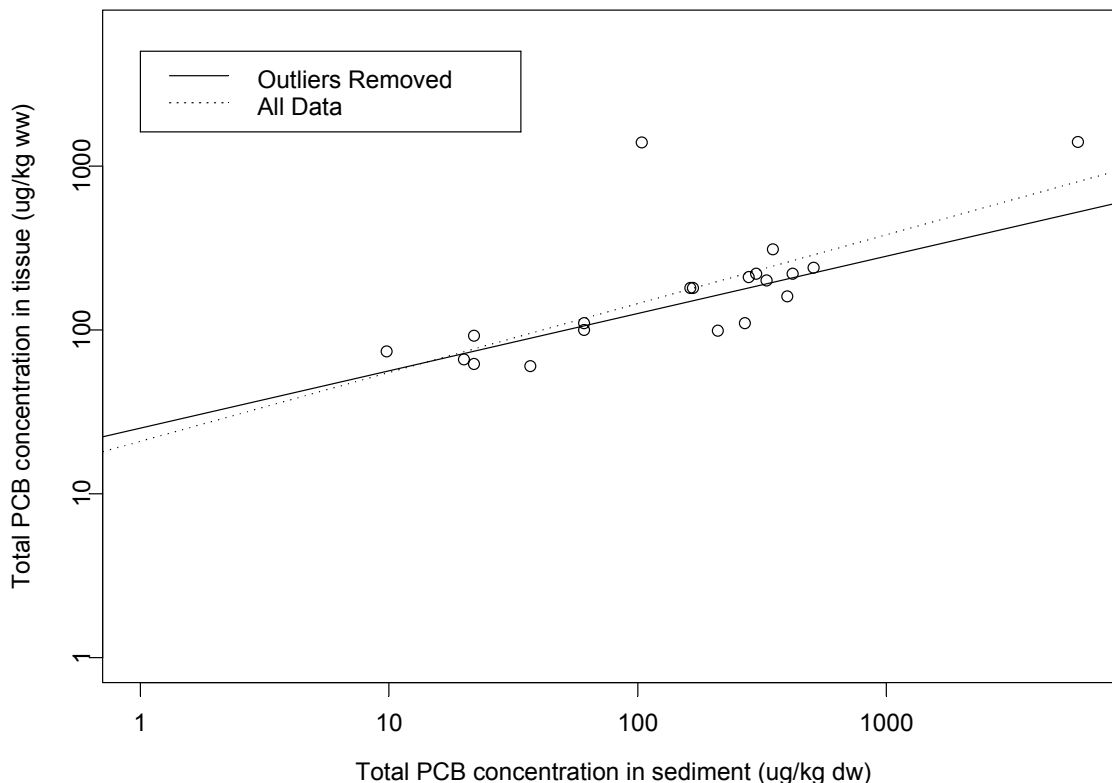
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<sup>4</sup> The relationship between organic-carbon-normalized sediment and lipid-normalized tissue was also tested, but the total PCB relationship without normalization provided a better fit to the data.

<sup>5</sup> The regression analysis was conducted by Alice Shelly of Terrastat Consulting Group.

<sup>6</sup> There was one non-detect sediment concentration (B1a; reporting limit = 20 µg/kg dw) and one non-detect tissue concentration (B4a; reporting limit = 200 µg/kg ww).

Total PCB concentrations in benthic invertebrate tissues for the entire LDW and for each modeling area (Table A-2-4) were estimated from total PCBs in sediment using the equation above. The sediment concentrations used were the SWACs from corresponding areas of the LDW (Table A-2-1).



**Figure A-2-1. Linear least-squares fit to log-transformed total PCB concentration in benthic invertebrate tissue as a function of log-transformed total PCB concentration in sediment**

#### **A.2.5 Benthic Invertebrate taxonomy data**

Benthic community data from both benthic invertebrate taxonomy and tissue samples were used to estimate benthic invertebrate weights and diets (Table A-2-3). Derivations of these parameter values are discussed separately below.

##### ***Benthic invertebrate weights***

Individual weights of benthic invertebrates were not measured as part of the laboratory processing of tissue or taxonomy samples. Therefore, an average individual weight was estimated using Phase 2 taxonomy abundance data and chemistry sample weight data (Windward 2005b, d). An average individual weight

was estimated for each of the 10 subtidal locations based on the major taxonomic groups identified in co-located taxonomy and tissue samples. Several assumptions and data evaluation steps were needed to derive the average and range of weights, as described below.

- ◆ Very small invertebrates identified in the taxonomy samples were not included in the composite tissue samples because taxonomy samples were sorted using a microscope whereas the composite tissue samples were sorted using the naked eye. Based on the size of organisms in the site-specific taxonomic reference collection, invertebrates in each of the composite tissue samples were classified as picked, maybe picked, and not picked. This classification assumed that the majority of invertebrates in the samples were adults because juvenile benthic invertebrates are generally too small to see without a microscope. Those invertebrates classified as “picked” were assumed to be included in the composite tissue samples.
- ◆ The abundances and composition of taxa observed in the three taxonomy samples were assumed to be proportional to the number collected in the 20 composite tissue samples, e.g., if 10% of the invertebrates in a taxonomy sample were *A. salmonis*, then 10% of the invertebrates in the co-located tissue sample were assumed to be *A. salmonis*.
- ◆ The assumption of proportional similarity between taxonomy and composite tissue samples was carried one step further by assuming that if, for example, *A. salmonis* constituted 10% of the crustacean abundance in the taxonomy sample then they also constituted 10% of the crustacean weight in the co-located tissue sample. Implicit in this assumption was that all organisms within a major taxonomic group (i.e., Annelida, Crustacea, Mollusca, and Miscellaneous Phyla) weighed the same.
- ◆ A weight per organism in each major taxonomic group was calculated by relating the number of individuals in each major taxonomic group from taxonomy samples to the weights of each major taxonomic group from co-located tissue samples. Because more sediment grabs were required for tissue samples, the number of organisms in a given taxonomic group was multiplied by the factor difference between the number of taxonomy sample sediment grabs and tissue chemistry sample sediment grabs. For example, for location B-1b, three sediment grabs were included in the taxonomy sample and 11 grabs were included in the co-located tissue sample resulting in a factor difference of 3.67. Therefore, for this location, 188 annelids, assumed picked in the taxonomy sample, resulted in 689 annelids assumed to be present in the co-located tissue sample. The total number of organisms in a major taxonomic group was then divided by the weight data for that group from the tissue sample to determine the weight per organism. Thus, for location B1-b, a total

annelid weight of 7.4 g in the tissue sample was divided by 689 annelids, resulting in 0.011 g per annelid.

- ◆ For a given sample location, the average individual weights of each major taxonomic group were averaged (average of annelids, crustaceans, mollusks, and miscellaneous phyla) to arrive at an average individual weight for that sample location. These averages were then used to generate average, maximum, and minimum individual weights for benthic invertebrates (Table A-2-3).

### ***Benthic invertebrate dietary scenarios***

In order to generate dietary scenarios for benthic invertebrates, the following process was followed.

- ◆ “Picked” invertebrates observed in the subtidal taxonomy samples were assigned a feeding type based on literature review.<sup>7</sup> Feeding types included carnivore, deposit feeder,<sup>8</sup> herbivore, and suspension feeder. Combined feeding types were assigned if different feeding strategies were presented in the literature. Combined feeding types were assumed to participate in each feeding type equally, thus a deposit feeder/carnivore was assumed to be 50% deposit feeder and 50% carnivore.
- ◆ Weight and abundance data (using a “picked” classification of invertebrates) were then used to generate the percent of sample weight comprised of the different feeding types.
- ◆ Average, maximum, and minimum percent of each feeding type over the 10 subtidal sample locations were calculated.
- ◆ Each feeding type was then assigned percentages of available dietary items. For dietary scenario 1, deposit feeders were assumed to ingest 100% sediment, suspension feeders were assumed to ingest 30% zooplankton and 70% phytoplankton/algae, and carnivores were assumed to ingest 100% sediment. Because the model does not allow modeled species to have a fraction of their diet from their own model compartment, and because only one benthic invertebrate compartment was created, sediment was used as a surrogate for benthic invertebrate prey consumed by carnivores. A “detritus” compartment was not modeled because data are unavailable to generate values for such a compartment. Sediment was used as a surrogate for detritus consumed by

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<sup>7</sup> Not all benthic invertebrates were assigned a feeding group because of limited information in the literature.

<sup>8</sup> Detritivore and deposit feeding types were combined into one type (deposit feeder) because of the similarity of their food items, sediment and detritus. Differences in proportions of sediment versus detritus consumed between these two feeding types are insignificant because detritus is represented by sediment in the model.

deposit feeders because it was assumed to have similar PCB concentrations. A limitation of this assumption is that detritus has higher organic matter content than sediment and could potentially have higher PCB concentrations.

Assumptions for dietary scenario 2 were the same except that carnivores were assigned to ingest 50% sediment and 50% zooplankton.

- ◆ Dietary item percentages were then multiplied by the percentage feeding type (by weight) to come up with dietary fractions of sediment, zooplankton, and phytoplankton/algae in the benthic invertebrate diet, for both dietary scenarios.

### A.2.6 Estimation of log K<sub>OW</sub> for PCBs

Estimates of log K<sub>OW</sub> were determined using site-specific tissue data. A concentration-weighted average log K<sub>OW</sub> was calculated using Equation A-2-2 for each tissue sample where all 209 individual PCB congeners were analyzed (Windward 2005a, e). PCB congener-specific K<sub>OWs</sub> were taken from Hawker and Connell (1988). Because there was little variability in concentration-weighted average log K<sub>OWs</sub> among species (Table A-2-7), the average of the species-specific averages (6.62) was used in the FWM. All results will be used for the data distribution in the sensitivity and uncertainty analyses.

$$\text{Average } K_{OW} = \frac{\sum_{i=1}^n C_i \times K_{OWi}}{\sum C_i} \quad \text{Equation A-2-2}$$

Where:

- C<sub>i</sub> – Concentration of PCB congener i
- K<sub>OWi</sub> – K<sub>OW</sub> of PCB congener i
- n – number of detected PCB congeners

**Table A-2-7. Average log K<sub>OWs</sub> for each modeled species derived using site-specific tissue data from the LDW**

SPECIES	N	MINIMUM	MAXIMUM	AVERAGE
Benthic invertebrates	8	6.42	6.87	6.57
Dungeness crab	5 <sup>a</sup>	6.54	6.74	6.64
Slender crab	7 <sup>b</sup>	6.55	6.63	6.58
English sole	7	6.50	6.64	6.56
Shiner surfperch	9	6.42	6.95	6.69
Pacific staghorn sculpin	8	6.63	6.84	6.69
Average of all tissue types				6.62

<sup>a</sup> Three edible meat composite samples and two hepatopancreas composite samples.

<sup>b</sup> Five edible meat composite samples and two hepatopancreas composite samples.

N Number of whole-body composite tissue samples

### **A.3 PARAMETERS DERIVED FROM THE LITERATURE**

The data presented in this section were derived from literature sources investigated by Windward. Parameter names, symbols, units, selected values, comments, and source information for the initial set of parameter values are presented in Table A-3-1. Species-specific diets based on literature data are presented in Table A-2-3. Because the analyses conducted to determine parameter values for modeled fish and crab species' diets cannot be fully described in Table A-3-1, they are further discussed in Section A.3.1.

**Table A-3-1. Model components with values derived from the literature**

MODEL COMPONENT	SYMBOL	UNITS	VALUES – MEAN (range)	NOTES	SOURCE
<b>Biological</b>					
Fraction of the diet consisting of prey item <i>i</i>	$P_i$	fraction (unitless)	species-specific	see A-2-3	see Table A-2-3
Fraction of overlying water ventilated	$m_o$	fraction (unitless)	species-specific	see Table A-2-3	
<b>Fraction of porewater ventilated</b>	<b><math>m_p</math></b>	<b>fraction (unitless)</b>	<b>species-specific</b>	<b>see Table A-2-3</b>	
<b>Lipid content of phytoplankton/algae</b>	<b><math>v_{LP}</math></b>	<b>% lipid ww</b>	<b>see Table A-2-3</b>	<b>see Table A-2-3</b>	<b>see Table A-2-3</b>
<b>NLOC content of phytoplankton/algae</b>	<b><math>v_{OCP}</math></b>	<b>% NLOC ww</b>	<b>see Table A-2-3</b>	<b>See Table A-2-3. NLOC is secondary site of PCB accumulation, for phytoplankton/algae.</b>	<b>see Table A-2-3</b>
Water content of phytoplankton/algae	$v_{WP}$	% water ww	see Table A-2-3	see Table A-2-3	see Table A-2-3
<b>Weight of the organism (zooplankton)</b>	<b><math>W_B</math></b>	<b>kg ww</b>	<b>species-specific</b>	<b>see Table A-2-3</b>	<b>see Table A-2-3</b>
<b>Lipid content of organism (zooplankton)</b>	<b><math>v_{LB}</math></b>	<b>% lipid ww</b>	<b>species-specific</b>	<b>see Table A-2-3</b>	<b>see Table A-2-3</b>
<b>NLOM content of organism (zooplankton)</b>	<b><math>v_{NB}</math></b>	<b>% NLOM ww</b>	<b>species-specific</b>	<b>See Table A-2-3. NLOM is a secondary site of PCB accumulation, for zooplankton.</b>	<b>see Table A-2-3</b>
Water content of organism (zooplankton)	$v_{WB}$	% ww	species-specific	See Table A-2-3. Water is not a significant contributor to the storage capacity of PCBs but is the third phase of storage in the body.	see Table A-2-3
<b>Chemical</b>					
<b>Octanol-water partition coefficient (total PCBs)</b>	<b><math>K_{ow}</math></b>	unitless	<b>6.62 (6.42 – 6.95)</b>	<b>Weighted average of <math>K_{ow}</math> for individual PCB congeners detected in Phase 2 tissue samples (for species in Table A-2-3), weighted by congener concentration (not weighted by species)</b>	<b><math>K_{ow}</math>s for each congener from Hawker and Connell (1988)</b>
Henry's Law Constant	H	(Pa x m <sup>3</sup> )/mol	43.3	This value cancels out in the model calculations.	Mackay et al. (1992)

**Bold** text indicates that the model has been demonstrated as sensitive to this parameter in the past (Arnot 2005).

mol – mole ( $6.022 \times 10^{23}$  entities)

NLOC – non-lipid organic carbon

NLOM – non-lipid organic matter

Pa – Pascals (units of pressure)

PCB – polychlorinated biphenyl

ww – wet weight



### A.3.1 Fish and crab dietary scenarios

One to four dietary scenarios were developed to explore the effect of different dietary assumptions on FWM predictions. The relative proportion of each prey item in modeled species' diets was determined from literature-reported stomach contents analyses. The studies used to characterize fish and crab diets are summarized in Table A-3-2. Dietary data were reported using various metrics. Biomass data were preferred when available. From each study, all prey items constituting at least 1% of diet were assigned to one of the four prey categories used in the FWM:

phytoplankton/algae, benthic invertebrates, zooplankton, and fish. Because some prey were unidentifiable, the percentage of prey biomass<sup>9</sup> assigned to each category was calculated relative to the total biomass of identifiable prey items only. The average fraction of prey biomass in each prey category over all studies was used to determine the relative proportions of prey used for FWM dietary scenarios 1 and 2 (Table A-2-3).<sup>10</sup> Juvenile fish diets were based on those of adult shiner surfperch because quantitative data for juveniles were not available. Juvenile English sole and shiner surfperch diets are similar, and diets are similar for between adult and juvenile life stages (Bane and Robinson 1970; Gordon 1965; Nyberg and Fahey 1988; Toole et al. 1987).

**Table A-3-2. Dietary studies used to characterize modeled species' diets**

STUDY	SPECIES MODELED	LOCATION	HABITAT	GEAR TYPES	SAMPLING FREQUENCY	REPORTED METRICS
Miller et al. (1977)	shiner surfperch, Pacific staghorn sculpin	Multiple North Puget Sound locations (Canada border to Fidalgo Island)	eelgrass, cobble, gravel, kelp bed	tow net, trammel net, beach seine	seasonally	Biomass, %IRI
Fresh et al. (1979)	English sole, shiner surfperch, Pacific staghorn sculpin	Nisqually River estuary, Nisqually Reach	mud, sand, gravel	trawl, beach seine	monthly	Biomass, %IRI
Wingert et al. (1979)	English sole, shiner surfperch, Pacific staghorn sculpin	West Point, Alki Point, Point Pully	eelgrass (only Alki Point reported)	trawl, beach seine	monthly	Biomass, %IRI
Stevens et al. (1982)	Dungeness crab	Grays Harbor, WA	two locations – sand/mud-flat and not reported	trawl	seasonally	%IRI

<sup>9</sup> Percent index of relative importance (%IRI) and % occurrence metrics were used for crab dietary studies.

<sup>10</sup> Prey data from the two Dungeness crab studies were not averaged because they reported different dietary metrics.

STUDY	SPECIES MODELED	LOCATION	HABITAT	GEAR TYPES	SAMPLING FREQUENCY	REPORTED METRICS
Gotshall (1977)	Dungeness crab	Humbolt Bay, CA and nearby ocean	not reported	trawl	Nov to Dec, Aug to Sep	frequency of occurrence, percent of prey
Bernard (1979)	slender crab	Hecate Strait, BC	silt, sand, and gravel	trawl	August	% of individual prey items

%IRI – percent index of relative importance

For all species modeled, dietary scenarios 1 and 2 (and Dungeness crab scenario 4) were statistical estimates of the modeled species diets based solely on stomach contents analyses. Both scenarios used the same prey data, with crabs and shrimp reported as prey assigned to different prey categories. In dietary scenario 1 (and Dungeness crab dietary scenario 4),<sup>11</sup> all crabs and shrimp reported as prey were assigned to the benthic invertebrate category. However, for dietary scenario 2, all crabs and shrimp reported as prey were assigned to the zooplankton category. The percentages of prey in each of the four categories for dietary scenarios 1 (and Dungeness crab dietary scenario 4) are presented in Table A-3-3. The percentages of prey in each of the four categories for dietary scenario 2 are presented in Table A-3-4. Dietary scenario 3 is the only scenario assuming sediment consumption. The relative proportions of prey in dietary scenario 3 represents a synthesis of available information regarding each species' diet, combined with knowledge of the LDW estuarine community. The proportions of prey surrogates assumed for dietary scenario 3 are presented in Table A-2-3 and Table 4-1 of the main document.

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<sup>11</sup> Because the two Dungeness crab studies report different metrics, one dietary scenario was developed for each study rather than averaging the dissimilar metrics to generate an average diet.

**Table A-3-3. Percent of phytoplankton/algae, benthic invertebrates, zooplankton, and fish in all fish and crab species' diets for dietary scenario 1 (and Dungeness crab for dietary scenario 4)**

SPECIES	STUDY	METRIC	N	PHYTO-PLANKTON/ ALGAE	BENTHIC INVERTEBRATES	ZOOPLANKTON	FISH
Dungeness crab <sup>a</sup>	Stevens et al. (1982) <sup>b</sup>	%IRI	410	<b>0</b>	<b>63</b>	<b>0</b>	<b>37</b>
	Gotshall (1977) <sup>c</sup>	% occurrence	337	<b>0</b>	<b>75</b>	<b>0</b>	<b>25</b>
Slender crab <sup>a</sup>	Bernard (1979)	% occurrence	48	<b>0</b>	<b>99</b>	<b>0</b>	<b>1</b>
Shiner surfperch	Fresh et al. (1979)	biomass	10	0	62	38	0
	Miller et al. (1977)	biomass	24	0	100	0	0
	Wingert et al. (1979)	biomass	31	0	95	5	0
Average <sup>b</sup>				<b>0</b>	<b>86</b>	<b>14</b>	<b>0</b>
English sole	Fresh et al. (1979)	biomass	36	5	95	0	0
	Wingert et al. (1979)	biomass	99	10	90	0	0
Average <sup>b</sup>				<b>8</b>	<b>92</b>	<b>0</b>	<b>0</b>
Pacific staghorn sculpin	Fresh et al. (1979)	biomass	57	0	83	0	17
	Miller et al. (1977)	biomass	51	0	52	0	48
	Wingert et al. (1979)	biomass	25	0	32	0	68
Average <sup>b</sup>				<b>0</b>	<b>56</b>	<b>0</b>	<b>44</b>

<sup>a</sup> Crab studies were not averaged because different metrics were used to characterize diets.

<sup>b</sup> Data used in dietary scenario 1.

<sup>c</sup> Data used in dietary scenario 4.

Numbers in **bold** were used in the FWM.

%IRI – percent index of relative importance

N – number of stomachs analyzed

**Table A-3-4. Percent of phytoplankton/algae, benthic invertebrates, zooplankton, and fish in all fish and crab species' diets for dietary scenario 2**

SPECIES	STUDY	METRIC	N	PHYTO- PLANKTON/ ALGAE	BENTHIC INVERTEBRATES	ZOOPLANKTON	FISH
Dungeness crab	Stevens et al. (1982)	% IRI	410	<b>0</b>	<b>16</b>	<b>48</b>	<b>36</b>
Slender crab	Bernard (1979)	% occurrence	48	<b>0</b>	<b>87</b>	<b>12</b>	<b>1</b>
Shiner surfperch	Fresh et al. (1979)	biomass	10	0	43	57	0
	Miller et al. (1977)	biomass	24	0	100	0	0
	Wingert et al. (1979)	biomass	31	0	95	5	0
Average				<b>0</b>	<b>79</b>	<b>21</b>	<b>0</b>
English sole	Fresh et al. (1979)	biomass	36	5	86	9	0
	Wingert et al. (1979)	biomass	99	10	90	0	0
Average				<b>7</b>	<b>88</b>	<b>5</b>	<b>0</b>
Pacific staghorn sculpin	Fresh et al. (1979)	biomass	25	0	32	50	17
	Miller et al. (1977)	biomass	57	0	21	31	48
	Wingert et al. (1979)	biomass	51	0	4	29	68
Average				<b>0</b>	<b>19</b>	<b>37</b>	<b>44</b>

Numbers in **bold** were used in the FWM.

%IRI – percent index of relative importance

N – number of stomachs analyzed

#### **A.4 DEFAULT PARAMETER VALUES FROM ARNOT AND GOBAS MODEL APPLICATION TO THE GREAT LAKES AND SAN FRANCISCO BAY**

The data presented in this section were derived from development of the Arnot and Gobas model and its application to the Great Lakes (Arnot and Gobas 2004) and San Francisco Bay (Gobas and Arnot 2005). Parameter names, symbols, units, selected values, comments, and source information for the initial set of parameter values are presented in Table A-4-1.

**Table A-4-1. Default values from Arnot and Gobas model application to the Great Lakes and San Francisco Bay**

MODEL COMPONENT	SYMBOL	UNITS	VALUES – MEAN (RANGE)	NOTES	SOURCE
<b>Biological</b>					
Density of lipids	$\delta_L$	kg/L	0.9		
Rate constant for growth of phytoplankton/algae	$k_G$	day <sup>-1</sup>	.08	Only phytoplankton/algae has $k_G$ as an input number instead of an equation. This is a mean annual value based on empirical data in which slow-growth conditions (winter) were 0.03 day <sup>-1</sup> and active-growth conditions (summer) were 0.13 day <sup>-1</sup> .	Swackhamer and Skoglund (1993) as cited in Arnot and Gobas (2004)
Scavenging efficiency of particles absorbed from the water	$\sigma$	fraction	1	Used to calculate feeding rate for filter feeders.	Morrison et al. (1996); Reeders et al. (1989); Ten Winkel and Davids (1982) (as cited in Arnot and Gobas (2004))
Algae, phytoplankton, and aquatic macrophytes – resistance to chemical uptake through aqueous phase	A	day <sup>-1</sup>	$6 \times 10^{-5}$ ( $\pm 2.0 \times 10^{-5}$ )	Derived from calibration to phytoplankton field BCF data from the Great Lakes.	Gobas and McLean (2003); Swackhamer and Skoglund (1993) (as cited in Arnot and Gobas (2004))
Algae, phytoplankton, and aquatic macrophytes – resistance to chemical uptake through organic phase	B	day <sup>-1</sup>	5.5 ( $\pm 3.7$ )	Derived by calibration to empirical $k_2$ values from various freshwater phytoplankton, algae, and cyanobacteria species over a range of $K_{OW}$ values.	Koelmans et al. (1993; 1995; 1999); Wang et al. (1996) (as cited in Arnot and Gobas (2004))
Proportionality constant expressing the sorption capacity of NLOM relative to that of octanol	$\beta$	L/kg	0.035	Based on 73-day lab test of HCBP with adult rainbow trout ( <i>Oncorhynchus mykiss</i> ) and a field study that analyzed PCB congener concentrations in tissue and GIT contents of rock bass ( <i>Ambloplites rupestris</i> ).	Gobas et al. (1999) (as cited in Arnot and Gobas (2004))
Proportionality constant expressing the sorption capacity of NLOC relative to that of octanol	$\beta_{oc}$	L/kg	0.35		Seth et al. (1999) (as cited in Arnot and Gobas (2004))

MODEL COMPONENT	SYMBOL	UNITS	VALUES – MEAN (RANGE)	NOTES	SOURCE
Dietary absorption efficiencies of lipid – fish	$\epsilon_L$	fraction	0.92	Based on 73-day lab test with adult rainbow trout ( <i>Oncorhynchus mykiss</i> ) and a field study of rock bass ( <i>Ambloplites rupestris</i> ).	Gobas et al. (1999) (as cited in Arnot and Gobas (2004))
Dietary absorption efficiencies of lipid – invertebrates	$\epsilon_L$	fraction	0.75	Based on studies involving zebra mussels from tidal freshwater section of Hudson River and polychaetes from Cape Cod intertidal flats.	Roditi and Fisher (1999); Berge and Brevik (1996); Gordon (1966); Parkerton (1993) (as cited in Arnot and Gobas (2004))
Dietary absorption efficiencies of lipid – zooplankton	$\epsilon_L$	fraction	0.72	Based on study involving <i>Calanus hyperboreus</i> eating diatoms and flagellates from Gulf of Maine.	Conover (1966) (as cited in Arnot and Gobas (2004))
Dietary absorption efficiencies of NLOM/NLOC – fish	$\epsilon_N$	fraction	0.6	Based on study with tetrachlorobiphenyl and rainbow trout.	Nichols et al. (2001) (as cited in Arnot and Gobas (2004))
Dietary absorption efficiencies of NLOM/NLOC – invertebrates	$\epsilon_N$	fraction	0.75	Based on studies involving zebra mussels from tidal freshwater section of Hudson River and polychaetes from Cape Cod intertidal flats.	Roditi and Fisher (1999); Berge and Brevik (1996); Gordon (1966); Parkerton et al. (1993) (as cited in Arnot and Gobas (2004))
Dietary absorption efficiencies of NLOM/NLOC – zooplankton	$\epsilon_N$	fraction	0.72	<i>Calanus hyperboreus</i> eating diatoms and flagellates from Gulf of Maine.	Conover (1966) (as cited in Arnot and Gobas (2004))
Dietary absorption efficiencies of water – all aquatic animal species	$\epsilon_W$	fraction	0.55	This value has been increased from 25% for the Great Lakes to 55% due to marine conditions (marine organisms retain more water and produce concentrated urine).	Gobas and Arnot (2005)
Rate constant for metabolic transformation of the chemical	$k_M$	day <sup>-1</sup>	0	Assume $k_M$ to be zero for all PCBs. Arnot and Gobas (2003), Fisk et al. (2000), and Van der Linde et al. (2001) identify ways to calculate $k_M$ .	Arnot and Gobas (2000)
<b>Environmental</b>					
Density of water	$\delta_W$	kg/L	1.0		Weast et al. (1985)

MODEL COMPONENT	SYMBOL	UNITS	VALUES – MEAN (RANGE)	NOTES	SOURCE
Proportionality constant describing similarity in phase partitioning of DOC relative to that of octanol	$\alpha_{\text{DOC}}$	unitless	0.08 (0.03 – 0.2)	Used in the bioavailable solute fraction equation for simulating sequestering of chemical by DOC in the water.	Burkhard (1999)
Proportionality constant describing similarity in phase partitioning of POC relative to that of octanol	$\alpha_{\text{POC}}$	unitless	0.35 (0.14 – 0.87)	Used in the bioavailable solute fraction equation for simulating sequestering of chemical by POC in the water.	Seth et al. (2004)
Disequilibrium factor for DOC partitioning	$D_{\text{DOC}}$	unitless	1	Used in the bioavailable solute fraction equation for simulating sequestering of chemical by DOC in the water. Assumes chemicals in the water column are in equilibrium with DOC.	Arnot and Gobas (2004)
Disequilibrium factor for POC partitioning	$D_{\text{POC}}$	unitless	1	Used in the bioavailable solute fraction equation for simulating sequestering of chemical by POC in the water. Assumes chemicals in the water column are in equilibrium with POC.	Arnot and Gobas (2004)

**Bold** text indicates that the model has been demonstrated to be sensitive to this parameter in the past (Arnot 2005).

BCF – bioconcentration factor

DOC – dissolved organic carbon

GIT – gastrointestinal tract

HCPB – PCB 155

NLOM – non-lipid organic matter

PCB – polychlorinated biphenyl

POC – particulate organic carbon

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# APPENDIX B. SENSITIVITY ANALYSES

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## **Appendix B. Sensitivity Analyses**

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This appendix presents the backup information for the sensitivity analyses conducted for the food web model (FWM). The analysis of model sensitivity involves the investigation of how changes in input parameters affect model output and identifies parameters that most influence model predictions. This analysis provides the basis for determining calibration parameters and also for selecting parameters to be evaluated in the uncertainty analysis. As discussed in the main document, two types of sensitivity analyses were conducted: reducing the value of each input parameter by 10% and altering the value of each input parameter based on its plausible range. Both analyses assessed the model's sensitivity to changes in single parameters. However, the first analysis identified the parameters to which the model output was most sensitive only as a result of the mathematical structure of the model. The second analysis helped identify the parameters to which the model output was most sensitive as a result of both its mathematical structure and the potential variability in parameter values.

### **B.1 10 PERCENT CHANGE ANALYSIS**

The 10% change analysis was conducted for all model input parameters identified in Table 5-1 of the FWM Memorandum 2 (Windward 2005) except three: the species-specific diet composition, the scavenging efficiency of particles absorbed from the water for filter feeders, and the concentration of suspended solids in the water column. The sensitivity of the model to differences in dietary composition was evaluated in Section 4.0 of the main document. Scavenging efficiency and suspended solids concentration were not included because they were not needed in the model. These parameters are used to calculate tissue concentrations for filter-feeding organisms, but filter feeders were not included in the current model because benthic invertebrates were modeled as scavengers / detritivores. The complete list of the 29 input parameters evaluated in the 10% change analysis and their initial and 10% adjusted values are presented in Table B-1-1.

All initial parameter values were decreased by 10% with the following exceptions: the octanol-water partition coefficient ( $K_{OW}$ ) and the metabolic transformation rate of PCBs ( $k_M$ ). Because  $K_{OW}$  is a component of many of the equations used in the model, the model is likely to respond differently to changes in  $K_{OW}$  depending upon which direction it is changed. Therefore, the model was run with both a 10% increase and decrease in  $K_{OW}$  to verify that it was not highly sensitive to a change in one direction but not the other. The initial  $k_M$  for PCBs was zero, which was not possible to decrease by 10%. Therefore, as discussed in FWM Memorandum 2 (Windward 2005), the model was run with a  $k_M$  of 0.0001 (Arnot 2005) and again with a 10% lower value (Table B-1-1).

Food ingestion rates ( $G_D$ ), organism growth rates ( $k_G$ ), and the PCB concentration of porewater are not actual input parameters in the model. They are all calculated from empirical equations within the model. For the 10% change analysis, their calculated values were decreased by 10%.



**Table B-1-1. Input parameter values for sensitivity analyses**

PARAMETER	UNITS	VALUE TYPE (% change)	VALUE	SOURCE
<b>Environmental Parameters</b>				
Total concentration of PCBs in the water column	ng/L	baseline	2	King County August water data (2005).
		-10%	1.8	
		upper (55%)	3.1	Full range of empirical data from August 2005 because too few data to calculate 95% confidence interval on the mean.
		lower (-25%)	1.5	Full range of empirical data from August 2005 because too few data to calculate 95% confidence interval on the mean.
Freely dissolved chemical concentration in the porewater ( $C_{WD,P}$ )	$\mu\text{g/kg}$	baseline	$8.88 \times 10^{-5}$	Calculated in model based on the sediment concentration, $OC_{sed}$ , and $K_{OC}$ .
		-10%	$7.99 \times 10^{-5}$	
		upper	na	Calculated in model; not possible to estimate a plausible range.
		lower	na	Calculated in model; not possible to estimate a plausible range.
Concentration of dissolved organic carbon (DOC) in the water column	kg/L	baseline	$2.20 \times 10^{-6}$	Unpublished King County 2005 water data (Mickelson 2006).
		-10%	$1.98 \times 10^{-6}$	
		upper (14%)	$2.50 \times 10^{-6}$	95% confidence interval on the mean from King County 2005 data.
		lower (-18%)	$1.80 \times 10^{-6}$	95% confidence interval on the mean from King County 2005 data.
Concentration of particulate organic carbon (POC) in the water column	kg/L	baseline	$2.90 \times 10^{-7}$	Unpublished King County 2005 water data (Mickelson 2006). Calculated as TOC-DOC in water.
		-10%	$2.61 \times 10^{-7}$	
		upper (41%)	$4.10 \times 10^{-7}$	95% confidence interval on the mean from King County 2005 data.
		lower (-45%)	$1.60 \times 10^{-7}$	95% confidence interval on the mean from King County 2005 data.
Mean water column temperature	$^{\circ}\text{C}$	baseline	11.60	Unpublished King County 2005 water data (Mickelson 2006).
		-10%	10.44	
		upper (11%)	12.90	95% confidence interval on the mean from King County 2005 data.
		lower (-12%)	10.20	95% confidence interval on the mean from King County 2005 data.

**Table B-1-1, cont.**

PARAMETER	UNITS	VALUE TYPE (% change)	VALUE	SOURCE
Dissolved oxygen concentration in the water column	mg/L	baseline	8	Unpublished King County 2005 water data (Mickelson 2006).
		-10%	7.2	
		upper (9%)	8.7	95% confidence interval on the mean from King County 2005 data.
		lower (-9%)	7.3	95% confidence interval on the mean from King County 2005 data.
Concentration of total PCBs in sediment	µg/kg dw	baseline	250	LDW-wide SWAC for baseline surface sediment data.
		-10%	225	
		upper (50%)	375	Based on a given range of results from the sediment groups investigation into different interpolation/SWAC generation methods
		lower (-50%)	125	Based on a given range of results from the sediment groups investigation into different interpolation/SWAC generation methods
Sediment organic carbon content (OC <sub>sed</sub> )	% dw	baseline	1.9	LDWG Phase 1 and 2 data.
		-10%	1.7	
		upper (6%)	2.04	Estimated value from 60th percentile of the distribution of estimates of the mean.
		lower (-8%)	1.78	Estimated value from 40th percentile of the distribution of estimates of the mean.
Proportionality constant describing similarity in phase partitioning of DOC relative to that of octanol ( $\alpha_{DOC}$ )	unitless	baseline	0.08	Arnot and Gobas (2004). Used to calculate bioavailable solute fraction in water in the model; resulting fraction = 0.464.
		-10%	0.072	Resulting bioavailable solute fraction = 0.480.
		upper (150%)	0.2	Arnot and Gobas (2004). Resulting bioavailable solute fraction = 0.307.
		lower (-63%)	0.03	Arnot and Gobas (2004). Resulting bioavailable solute fraction = 0.589.
Proportionality constant describing similarity in phase partitioning of POC relative to that of octanol ( $\alpha_{POC}$ )	unitless	baseline	0.35	Arnot and Gobas (2004). Used to calculate bioavailable solute fraction in water in the model; resulting fraction = 0.464.
		-10%	0.315	Resulting bioavailable solute fraction = 0.473.
		upper (149%)	0.87	Arnot and Gobas (2004). Resulting bioavailable solute fraction = 0.359.
		lower (-60%)	0.14	Arnot and Gobas (2004). Resulting bioavailable solute fraction = 0.526.
Disequilibrium factor for DOC partitioning (D <sub>DOC</sub> )	unitless	baseline	1.0	Arnot and Gobas (2004). Used to calculate bioavailable solute fraction in water in the model; resulting fraction = 0.464.
		-10%	0.9	Resulting bioavailable solute fraction = 0.480.

**Table B-1-1, cont.**

PARAMETER	UNITS	VALUE TYPE (% change)	VALUE	SOURCE
		upper	na	Did not run; no range data available.
		lower	na	Did not run; no range data available.
Disequilibrium factor for POC partitioning ( $D_{POC}$ )	unitless	baseline	1.0	Arnot and Gobas (2004). Used to calculate bioavailable solute fraction in water in the model; resulting fraction = 0.464.
		-10%	0.9	Resulting bioavailable solute fraction = 0.473.
		upper	na	Did not run; no range data available.
		lower	na	Did not run; no range data available.
Density of water ( $\delta_w$ )	kg/L	baseline	1	Weast et al. (1985)
		-10%	0.9	
		upper	1.02	Weast et al. (1985)
		lower	na	
<b>Chemical Parameters</b>				
Octanol-water partition coefficient for PCBs ( $\log K_{ow}$ )	unitless	baseline	6.62	Phase 2 LDWG data and $K_{ow}$ values in Hawker and Connell (1988).
		-10%	6.57	
		+ 10%	6.66	
		upper	6.66	95 % confidence interval on the mean for the LDWG data used above.
		lower	6.58	95 % confidence interval on the mean for the LDWG data used above.
<b>Biological Parameters</b>				
Density of lipids ( $\delta_L$ )	kg/L	baseline	0.9	Arnot (2006)
		-10%	0.81	
		upper	1	
		lower	0.8	
Rate constant for metabolic transformation of PCBs ( $K_m$ )	unitless	separate baseline	$1 \times 10^{-4}$	Arnot (2005). In the initial set of input values $K_m = 0$ .
		-10%	$9 \times 10^{-5}$	
		upper	na	
		lower	na	
Proportionality constant expressing the sorption capacity of NLOM	L/kg	baseline	0.035	Gobas et al. (1999), as cited in Arnot and Gobas (2004).
		-10%	0.0315	

**Table B-1-1, cont.**

PARAMETER	UNITS	VALUE TYPE (% change)	VALUE	SOURCE
relative to that of octanol ( $\beta$ or MAF)		upper (29%)	0.045	Arnot (2005)
		lower (-29%)	0.025	Arnot (2005)
Proportionality constant expressing the sorption capacity of NLOC relative to that of octanol ( $\beta_{OC}$ )	L/kg	baseline	0.35	Seth et al. (1999)
		-10%	0.315	
		upper	na	
		lower	na	
Resistance to chemical uptake through aqueous phase for phytoplankton/algae (A)	day <sup>-1</sup>	baseline	$6.0 \times 10^{-5}$	Arnot and Gobas (2004)
		-10%	$5.4 \times 10^{-5}$	
		upper (33%)	$8.0 \times 10^{-5}$	Gobas and Arnot (2005)
		lower (-33%)	$4.0 \times 10^{-5}$	Gobas and Arnot (2005)
Resistance to chemical uptake through organic phase for phytoplankton/algae (B)	day <sup>-1</sup>	baseline	5.5	Arnot and Gobas (2004)
		-10%	4.95	
		upper (67%)	9.20	Gobas and Arnot (2005)
		lower (-67%)	1.80	Gobas and Arnot (2005)
<b>Growth Rate Constant (<math>k_G</math>)</b>				
Phytoplankton/algae	day <sup>-1</sup>	baseline	$8.00 \times 10^{-2}$	Model default = 0.8 (Arnot and Gobas 2004).
		-10%	$7.20 \times 10^{-2}$	
Zooplankton	day <sup>-1</sup>	baseline	$1.15 \times 10^{-2}$	Calculated in model based on organism weight.
		-10%	$1.03 \times 10^{-2}$	
Benthic invertebrates	day <sup>-1</sup>	baseline	$3.62 \times 10^{-3}$	Calculated in model based on organism weight.
		-10%	$3.26 \times 10^{-3}$	
Juvenile fish	day <sup>-1</sup>	baseline	$1.40 \times 10^{-3}$	Calculated in model based on organism weight.
		-10%	$1.26 \times 10^{-3}$	
Slender crab	day <sup>-1</sup>	baseline	$7.21 \times 10^{-4}$	Calculated in model based on organism weight.
		-10%	$6.49 \times 10^{-4}$	
Dungeness crab	day <sup>-1</sup>	baseline	$5.96 \times 10^{-4}$	Calculated in model based on organism weight.
		-10%	$5.37 \times 10^{-4}$	

**Table B-1-1, cont.**

PARAMETER	UNITS	VALUE TYPE (% change)	VALUE	SOURCE
Pacific staghorn sculpin	day <sup>-1</sup>	baseline	8.81×10 <sup>-4</sup>	Calculated in model based on organism weight.
		-10%	7.93×10 <sup>-4</sup>	
Shiner surfperch	day <sup>-1</sup>	baseline	1.13×10 <sup>-3</sup>	Calculated in model based on organism weight.
		-10%	1.02×10 <sup>-3</sup>	
English sole	day <sup>-1</sup>	baseline	6.94×10 <sup>-4</sup>	Calculated in model based on organism weight.
		-10%	6.25×10 <sup>-4</sup>	
<b>Food Ingestion Rate (G<sub>D</sub>)</b>				
Zooplankton	kg food/ day	baseline	7.38×10 <sup>-8</sup>	Calculated in model based on organism weight and water temperature.
		-10%	6.64×10 <sup>-8</sup>	
Benthic invertebrates	kg food/ day	baseline	9.91×10 <sup>-6</sup>	Calculated in model based on organism weight and water temperature.
		-10%	8.92×10 <sup>-6</sup>	
Juvenile fish	kg food/day	baseline	5.70×10 <sup>-4</sup>	Calculated in model based on organism weight and water temperature.
		-10%	5.13×10 <sup>-4</sup>	
Slender crab	kg food/ day	baseline	9.49×10 <sup>-4</sup>	Calculated in model based on organism weight and water temperature.
		-10%	8.54×10 <sup>-4</sup>	
Dungeness crab	kg food/ day	baseline	2.12×10 <sup>-2</sup>	Calculated in model based on organism weight and water temperature.
		-10%	1.91×10 <sup>-2</sup>	
Pacific staghorn sculpin	kg food/ day	baseline	4.04×10 <sup>-3</sup>	Calculated in model based on organism weight and water temperature.
		-10%	3.63×10 <sup>-3</sup>	
Shiner surfperch	kg food/ day	baseline	1.38×10 <sup>-3</sup>	Calculated in model based on organism weight and water temperature.
		-10%	1.24×10 <sup>-3</sup>	
English sole	kg food/ day	baseline	1.11×10 <sup>-2</sup>	Calculated in model based on organism weight and water temperature.
		-10%	1.00×10 <sup>-2</sup>	
<b>Organism Weight</b>				
Zooplankton	kg	baseline	1.60×10 <sup>-7</sup>	Giles and Cordell (1998)
		-10%	1.44×10 <sup>-7</sup>	
		upper (44%)	2.30×10 <sup>-7</sup>	Range observed in literature.

**Table B-1-1, cont.**

PARAMETER	UNITS	VALUE TYPE (% change)	VALUE	SOURCE
		lower (-45%)	$8.80 \times 10^{-8}$	Range observed in literature.
Benthic invertebrates	kg	baseline	$5.10 \times 10^{-5}$	LDWG Phase 2 data.
		-10%	$4.59 \times 10^{-5}$	
		upper (312%)	$2.10 \times 10^{-4}$	Range observed in LDWG Phase 2 data.
		lower (-89%)	$5.50 \times 10^{-6}$	Range observed in LDWG Phase 2 data.
Juvenile fish	kg	baseline	$6.00 \times 10^{-3}$	LDWG Phase 2 individual shiner surfperch specimens (<80mm) as surrogates for juvenile fish
		-10%	$5.40 \times 10^{-3}$	
		upper (17%)	$7.00 \times 10^{-3}$	95% confidence interval on LDWG Phase 2 data mean.
		lower (0%)	$6.00 \times 10^{-3}$	95% confidence interval on LDWG Phase 2 data mean. Lower bound estimate comes out to be the same as the mean value due to rounding for significant figures.
Slender crab	kg	baseline	0.164	LDWG Phase 2 data.
		-10%	0.148	
		upper (5%)	0.172	95% confidence interval on LDWG Phase 2 data mean.
		lower (4%)	0.157	95% confidence interval on LDWG Phase 2 data mean.
Dungeness crab	kg	baseline	0.423	LDWG Phase 2 data.
		-10%	0.381	
		upper (55%)	0.657	95% confidence interval on LDWG Phase 2 data mean.
		lower (-77%)	0.096	Minimum observed from LDWG Phase 2; not possible to calculate lower confidence interval.
Pacific staghorn sculpin	kg	baseline	0.06	LDWG Phase 2 data.
		-10%	0.054	
		upper (7%)	0.0642	95% confidence interval on LDWG Phase 2 data mean.
		lower (-6%)	0.0562	95% confidence interval on LDWG Phase 2 data mean.
Shiner surfperch	kg	baseline	0.017	LDWG Phase 2 data.
		-10%	0.0153	
		upper (6%)	0.018	95% confidence interval on LDWG Phase 2 data mean.

**Table B-1-1, cont.**

PARAMETER	UNITS	VALUE TYPE (% change)	VALUE	SOURCE
		lower (-2%)	0.0166	95% confidence interval on LDWG Phase 2 data mean.
English sole	kg	baseline	0.198	LDWG Phase 2 data.
		-10%	0.178	
		upper (7%)	0.211	95% confidence interval on LDWG Phase 2 data mean.
		lower (-7%)	0.185	95% confidence interval on LDWG Phase 2 data mean.
<b>Lipid Content</b>				
Phytoplankton/algae	% ww	baseline	0.1%	Mackintosh et al. (2004)
		-10%	0.1%	
		upper (16%)	0.1%	Range observed in literature.
		lower (-16%)	0.1%	Range observed in literature.
Zooplankton	% ww	baseline	1.2%	Kuroshima et al. (1987).
		-10%	1.1%	
		upper (42%)	1.7%	Range observed in literature.
		lower (-25%)	0.9%	Range observed in literature.
Benthic invertebrates	% ww	baseline	0.9%	LDWG Phase 2 data.
		-10%	0.8%	
		upper (12%)	1.0%	95% confidence interval on LDWG Phase 2 data mean.
		lower (-14%)	0.8%	95% confidence interval on LDWG Phase 2 data mean.
Juvenile fish	% ww	baseline	2.5%	LDWG Phase 2 data for English sole and shiner surfperch.
		-10%	2.3%	
		upper (8%)	2.7%	Used the observed variability in means for English sole and shiner surfperch in LDWG Phase 2 data.
		lower (-8%)	2.3%	Used the observed variability in means for English sole and shiner surfperch in LDWG Phase 2 data.
Slender crab	% ww	baseline	1.1%	LDWG Phase 2 data.
		-10%	1.0%	
		upper (21%)	1.3%	95% confidence interval on LDWG Phase 2 data mean.
		lower (-16%)	0.9%	95% confidence interval on LDWG Phase 2 data mean.

**Table B-1-1, cont.**

PARAMETER	UNITS	VALUE TYPE (% change)	VALUE	SOURCE
Dungeness crab	% ww	baseline	2.6%	LDWG Phase 2 data.
		-10%	2.3%	
		upper (38%)	3.6%	95% confidence interval on LDWG Phase 2 data mean.
		lower (-38%)	1.6%	95% confidence interval on LDWG Phase 2 data mean.
Pacific staghorn sculpin	% ww	baseline	2.1%	LDWG Phase 2 data.
		-10%	1.9%	
		upper (4%)	2.2%	95% confidence interval on LDWG Phase 2 data mean.
		lower (-9%)	1.9%	95% confidence interval on LDWG Phase 2 data mean.
Shiner surfperch	% ww	baseline	4.6%	LDWG Phase 2 data.
		-10%	4.1%	
		upper (9%)	5.0%	95% confidence interval on LDWG Phase 2 data mean.
		lower (-7%)	4.3%	95% confidence interval on LDWG Phase 2 data mean.
English sole	% ww	baseline	5.5%	LDWG Phase 2 data.
		-10%	5.0%	
		upper (7%)	5.9%	95% confidence interval on LDWG Phase 2 data mean.
		lower (-7%)	5.1%	95% confidence interval on LDWG Phase 2 data mean.
<b>Water Content</b>				
Phytoplankton/algae	% ww	baseline	95.6%	Mackintosh et al. (2004).
		-10%	86.0%	
		upper (1%)	96.5%	Range observed in literature.
		lower (-1%)	94.7%	Range observed in literature.
Zooplankton	% ww	baseline	90.0%	Kuroshima et al. (1987).
		-10%	81.0%	
		upper (1%)	91.2%	Range observed in literature.
		lower (-3%)	87.0%	Range observed in literature.



**Table B-1-1, cont.**

PARAMETER	UNITS	VALUE TYPE (% change)	VALUE	SOURCE
Benthic invertebrates	% ww	baseline	88.9%	LDWG Phase 2 data.
		-10%	80.0%	
		upper (2%)	90.4%	95% confidence interval on LDWG Phase 2 data mean.
		lower (-2%)	87.3%	95% confidence interval on LDWG Phase 2 data mean.
Juvenile fish	% ww	baseline	73.9%	LDWG Phase 2 data.
		-10%	66.5%	
		upper (4%)	77.2%	Range observed in LDWG Phase 2 data.
		lower (-6%)	69.6%	Range observed in LDWG Phase 2 data.
Slender crab	% ww	baseline	83.6%	LDWG Phase 2 data.
		-10%	75.2%	
		upper (2%)	85.1%	95% confidence interval on LDWG Phase 2 data mean.
		lower (-1%)	82.9%	95% confidence interval on LDWG Phase 2 data mean.
Dungeness crab	% ww	baseline	82.0%	LDWG Phase 2 data.
		-10%	73.8%	
		upper (2%)	83.9%	95% confidence interval on LDWG Phase 2 data mean.
		lower (-3%)	79.5%	95% confidence interval on LDWG Phase 2 data mean.
Pacific staghorn sculpin	% ww	baseline	79.0%	LDWG Phase 2 data.
		-10%	71.1%	
		upper (0.3%)	79.2%	95% confidence interval on LDWG Phase 2 data mean.
		lower (-0.3%)	78.8%	95% confidence interval on LDWG Phase 2 data mean.
Shiner surfperch	% ww	baseline	73.9%	LDWG Phase 2 data.
		-10%	66.5%	
		upper (1%)	74.5%	95% confidence interval on LDWG Phase 2 data mean.
		lower (-1%)	73.3%	95% confidence interval on LDWG Phase 2 data mean.

**Table B-1-1, cont.**

PARAMETER	UNITS	VALUE TYPE (% change)	VALUE	SOURCE
English sole	% ww	baseline	75.0%	LDWG Phase 2 data.
		-10%	67.5%	
		upper (1%)	75.5%	95% confidence interval on LDWG Phase 2 data mean.
		lower (-1%)	74.4%	95% confidence interval on LDWG Phase 2 data mean.
<b>Fraction of Porewater Ventilated</b>				
Benthic invertebrates	fraction	baseline	0.2	Winsor et al. (1990).
		-10%	0.18	
		upper (25%)	0.25	Range observed in literature.
		lower (-75%)	0.05	Range observed in literature.
Juvenile fish	fraction	baseline	0.01	Gobas and Wilcockson (2003)
		-10%	0.009	
		upper (100%)	0.02	Range observed in literature.
		lower (-50%)	0.005	Range observed in literature.
Slender crab	fraction	baseline	0.02	Winsor et al. (1990); Gobas and Wilcockson (2003)
		-10%	0.018	
		upper	0.03	Range observed in literature.
		lower	0.01	Range observed in literature.
Dungeness crab	fraction	baseline	0.02	Winsor et al. (1990); Gobas and Wilcockson (2003)
		-10%	0.018	
		upper (50%)	0.03	Range observed in literature.
		lower (-50%)	0.01	Range observed in literature.
Pacific staghorn sculpin	fraction	baseline	0.05	Value from model components table.
		-10%	0.045	
		upper (100%)	0.1	Range from model components table.
		lower (-60%)	0.02	Range from model components table.
Shiner surfperch	fraction	baseline	0.01	Gobas and Wilcockson (2003)
		-10%	0.009	

**Table B-1-1, cont.**

PARAMETER	UNITS	VALUE TYPE (% change)	VALUE	SOURCE
		upper (100%)	0.02	Range observed in literature.
		lower (-50%)	0.005	Range observed in literature.
English sole	fraction	baseline	0.1	Gobas and Wilcockson (2003)
		-10%	0.09	
		upper (100%)	0.2	Range observed in literature.
		lower (-50%)	0.05	Range observed in literature.
<b>Dietary Absorption Efficiency of Lipids (alpha)</b>				
Zooplankton	fraction	baseline	0.72	Arnot and Gobas (2004)
		-10%	0.65	
		upper (18%)	0.85	Range reported in Arnot and Gobas (2004).
		lower (-42%)	0.55	Range reported in Arnot and Gobas (2004).
Benthic invertebrates (including crabs)	fraction	baseline	0.75	Arnot and Gobas (2004)
		-10%	0.68	
		upper (13%)	0.96	Range reported in Arnot and Gobas (2004).
		lower (-44%)	0.15	Range reported in Arnot and Gobas (2004).
All fish	fraction	baseline	0.92	Arnot and Gobas (2004)
		-10%	0.83	
		upper	na	No range data available.
		lower	na	No range data available.
<b>Dietary Absorption Efficiency of NLOM (beta)</b>				
Zooplankton	fraction	baseline	0.72	Arnot and Gobas (2004)
		-10%	0.65	
		upper (18%)	0.85	Range reported in Arnot and Gobas (2004).
		lower (-42%)	0.55	Range reported in Arnot and Gobas (2004).

**Table B-1-1, cont.**

PARAMETER	UNITS	VALUE TYPE (% change)	VALUE	SOURCE
Benthic invertebrates (including crabs)	fraction	baseline	0.75	Arnot and Gobas (2004)
		-10%	0.68	
		upper (13%)	0.96	Range reported in Arnot and Gobas (2004).
		lower (-44%)	0.15	Range reported in Arnot and Gobas (2004).
All fish	fraction	baseline	0.60	Arnot and Gobas (2004)
		-10%	0.54	
		upper (5%)	0.63	Assumed maximum based on values reported in Arnot and Gobas (2004).
		lower (-5%)	0.57	Reported in Arnot and Gobas (2004).
<b>Dietary Absorption Efficiency of Water</b>				
All organisms	fraction	baseline	0.55	Arnot and Gobas (2004)
		-10%	0.50	
		upper	na	No data were available to calculate a range.
		lower	na	No data were available to calculate a range.

Growth rate constant and food ingestion rate are calculated within the model; it was not possible to incorporate upper and lower range estimates.

na – not applicable

dw – dry weight

ww – wet weight

NLOC– Non-lipid organic carbon

NLOM – Non-lipid organic matter

SWAC – spatially weighted average concentration

## B.2 PLAUSIBLE RANGE SENSITIVITY ANALYSIS

Upper- and lower-bound estimates were developed for all input parameters with significant information to estimate ranges. Plausible ranges were estimated for these parameters using one of two methods. A statistical method was used for parameters where enough data were available to determine the distribution of those data. This approach is described in detail in Section B.2.1. For both site-specific and literature-derived parameters with insufficient data to develop a distribution, range estimates were compiled from the literature. The methods used to determine the plausible range for each parameter are presented in Table B-1-1.

The following parameters were not included in the upper- and lower-bound sensitivity analysis because plausible value ranges were not available:

- ◆ Freely dissolved chemical concentration in the porewater
- ◆ Disequilibrium factor for dissolved organic carbon (DOC) partitioning
- ◆ Disequilibrium factor for particulate organic carbon (POC) partitioning
- ◆ Rate constant for metabolic transformation of PCBs ( $k_M$ )
- ◆ Growth rate constant ( $k_G$ )
- ◆ Food ingestion rate ( $G_D$ )
- ◆ Dietary absorption efficiency of lipids for fish
- ◆ Dietary absorption efficiency of water
- ◆ Concentration of suspended solids in the water column
- ◆ Scavenging efficiency of particles in the water column by filter feeders

### B.2.1 Statistical approach

Mean data were generally used for the initial set of parameter values. The purpose of the plausible-range sensitivity analysis was to evaluate how sensitive the model was to expected or observed variability in the mean parameter values because of uncertainty or natural variability. Therefore, wherever possible, upper and lower estimates of the mean were used in this analysis. For parameters where data distribution information was available, the 95% confidence interval (CI) of the mean (i.e., the 2.5 and 97.5 percentiles of the distribution of estimates of the mean) was used to determine the plausible range. The statistical methods used depended on how the data for each parameter were distributed. The statistical analysis software ProUCL was used to determine how the data for each of these parameters was distributed and the statistical approach for calculation of confidence limits on the mean.

Most of the data were normally distributed. For those parameters, the 95% CI for the mean was approximated with the standard error (SE) using the following equations (Ott 1993):

$$\text{SE} = \text{Standard deviation} / \sqrt{(\text{sample number})}$$
$$95\% \text{ CI of mean} = \text{mean} \pm (1.96 \times \text{SE})$$

The same datasets that were used to determine initial parameter values (Appendix A) were used to estimate the plausible upper and lower ranges.

For data that were not normally distributed, ProUCL recommended another statistical approach to analyze the data. The methods recommended by ProUCL were, therefore, used to calculate the 95% CI of the mean.

For Dungeness crab weight, the data were not normally distributed. ProUCL provided a 97.5 upper confidence limit on the mean, but because larger crabs were selectively taken during sampling, the sample distribution was not expected to reflect population distribution for lower weights. For this reason, the minimum observed value was used as the lower range.

## **B.2.2 Literature-based approach**

For those parameters with insufficient data about their distribution, range estimates were compiled from the literature. For most of these parameters, the ranges presented in the literature were used as the plausible ranges. Many of those ranges were presented in the model documentation (Arnot and Gobas 2004) and are included in the model components table (Appendix A).

As discussed in Appendix A, juvenile fish parameter values were calculated from the Phase 2 data for shiner surfperch and inferences from data in other studies (Gobas and Arnot 2005). The statistical approach described in Section B.2.1 was used to estimate the plausible range for juvenile fish weight and water content. Lipid content range, however, was estimated by applying the measured variability in lipid content from English sole and shiner surfperch (+/- 45% of the mean) to the calculated mean lipid content for juvenile fish.

There are few data directly addressing the dietary absorption efficiencies for lipids, non-lipid organic matter (NLOM), and water for various organisms, and they were not sufficient to provide estimates of the mean. Arnot and Gobas (2004) summarize the available data for all three parameters. Given the uncertainty around these parameter values, the full range of efficiencies for lipids and NLOM for all invertebrates were selected for zooplankton, benthic invertebrates, and crabs. No ranges were presented, however, for water absorption efficiencies for any species. Therefore, no plausible range estimates were used for that parameter.

Ranges were also not presented for the fish dietary absorption efficiencies for lipids and NLOM. Arnot and Gobas (2004) recommend using 60% for the NLOM absorption efficiency for fish, but the study they cite presents an NLOM absorption

efficiency of 57% (Nichols et al. 2001). For the purposes of determining a plausible range in this analysis, 60% was used as the initial value and 57% was used as the lower range value. To be consistent with the lower range estimate and mean, and assuming estimates of the mean are normally distributed, 63% was assumed for an upper range estimate (Table B-1-1).

As discussed in Appendix A, the total PCB concentration in water was estimated using data collected from two stations in the LDW by King County (2005). When the sensitivity analysis was conducted, data were only available from the August sampling event. Consequently, there were insufficient data to determine a 95% CI on the mean for the total PCB concentration in the water column. The observed minimum and maximum concentrations reported in the model components table (Table A-2-1) were, therefore, used for the upper and lower plausible ranges.

As discussed in Appendix A, spatially weighted average concentrations (SWACs) of organic carbon ( $OC_{sed}$ ) and total PCBs in sediment were estimated using an inverse distance weighting (IDW) interpolation model. It was not possible to estimate a meaningful plausible range for  $OC_{sed}$  content because the sample size generated by the interpolation model (187,000) was so large that the standard error estimates were not significant, given the number of significant figures for both  $OC_{sed}$  and total PCBs in sediment. Therefore, the 40th and 60th percentiles of the interpolated data distribution were assumed to represent plausible ranges of the mean  $OC_{sed}$  concentration (Table B-1).

The plausible range for the total PCB SWAC was estimated based on the previous efforts to develop a SWAC for sediment PCBs. Three different methods for determining an organic carbon (OC) normalized PCB SWAC (IDW, Thiessen Polygons, and Kriging) were investigated by LDWG in fall 2005, prior to the calculation of the SWAC using IDW as discussed in Section A.2.1 of Appendix A. The method that resulted in the highest SWAC (Thiessen Polygons) from these previous efforts differed by 50% from the IDW approach. The previous calculations included data through October 2005 and all PCB concentrations were OC normalized. The SWAC estimates were 18mg/kg-OC using Thiessen polygons, 12 mg/kg-OC using IDW, and 10 mg/kg-OC using Kriging. Assuming the ratio between estimates would be the same for OC normalized PCB concentrations as for bulk sediment PCB concentrations, SWAC estimates would be expected to span up to 1.5 times the mean concentration used in the FWM. Based on these analyses, the range around the selected SWAC of 250  $\mu\text{g}/\text{kg dw}$  was estimated to be from 125  $\mu\text{g}/\text{kg dw}$  to 375  $\mu\text{g}/\text{kg dw}$  (Table B-1-1). The IDW interpolation approach and baseline sediment dataset were still being finalized at the time that this memorandum was being completed. Therefore, all SWAC estimates used in this memorandum are preliminary.

### **B.2.3 Evaluation of sensitivity**

The sensitivity of the FWM was evaluated by performing separate model runs with each parameter's adjusted value. For example, the model was run 30 times for the 10% change analysis because it was run separately for each of the 29 parameters, including two separate runs for  $K_{ow}$ . For parameters with species-specific values (e.g., organism weight), all species values were adjusted simultaneously. The output from each model run was compared with the output from the model run with the initial set of values using the species percent difference (SPD) metric described in FWM Memorandum 2 (Windward 2005). The SPD represents the difference in predicted tissue concentration from the initial predicted concentrations; it was calculated for each species for each model run. Parameter sensitivity was evaluated based on the mean and maximum SPDs.

In addition to the SPD, a relative response ratio was calculated for the plausible range analysis. This metric is calculated as the SPD divided by the percent change in parameter value. This metric allows comparison of parameter responses between the 10% sensitivity analysis and the plausible range analysis because it presents the model response relative to the change in parameter value.

For the 10% sensitivity analysis, results were ranked by maximum SPD, and any parameter with a maximum SPD of 8% or more for any species was selected for inclusion in the uncertainty analysis. The threshold of an 8% change in predicted tissue concentration (for any one species) for a 10% change in parameter value was selected to include parameters to which the model is moderately sensitive. A greater than 1:1 response between parameter value change and model prediction change is generally considered highly sensitive (Arnot 2006).

Also identified were parameters that, when run at the upper or lower end of their plausible range, cause a percentage change that is substantial relative to the change caused by other parameters or relative to the magnitude of change in the input value. In order to select parameters for the uncertainty analysis, results were ranked by maximum SPD, and the distribution of results was evaluated to see if any patterns or break points arose from the results. Parameters were also ranked according to a relative response ratio (SPD divided by percent change in parameter value). This metric can be compared to the 10% sensitivity analysis to see if percent changes in model predictions are the same for small or large changes in parameter values.



### B.3 RESULTS OF SENSITIVITY ANALYSES

The results from both sensitivity analyses are discussed in Section 5.0 of the main document. This section presents five results tables. Tables B-3-1 and B-3-2 rank parameters by maximum SPD using results of the 10% sensitivity analysis for target species and for all species, respectively. Tables B-3-3 and B-3-4 rank parameters by maximum SPD using results of the plausible range sensitivity analyses for target species and for all modeled species, respectively. Table B-3-5 ranks parameters by a relative response ratio for the upper- and lower-bound sensitivity analyses for all species. The parameters with a maximum relative response of 0.8 or greater (Table B-5-5) were the same as those with a maximum SPD of 8% or greater presented in Tables B-3-2.

**Table B-3-1. Results of the 10 percent sensitivity analysis for predicted fish and crab total PCB concentrations**

PARAMETER	MAXIMUM SPD (absolute value)	SPECIES WITH MAXIMUM CHANGE	MINIMUM SPD (absolute value)	MEAN SPD (with negatives)
Dietary absorption efficiency of lipids (alpha)	24%	PSS	10%	-14%
Water content	18%	SC	2%	5%
Lipid density	17%	PSS	10%	13%
Food ingestion rate ( $G_D$ )	14%	PS	10%	-12%
Lipid content	14%	PSS	9%	-11%
Dissolved oxygen (DO)	11%	PSS	7%	-9%
Water column temperature	10%	PSS	6%	-8%
Dietary absorption efficiency of NLOM (beta)	9%	DC	6%	-7%
Sediment PCB concentration	8%	SC	8%	-8%
$K_{OW}$	7%	PSS	4%	-5%
Growth rate constant ( $k_G$ )	4%	ES	2%	3%
Sediment organic carbon ( $OC_{sed}$ )	4%	ES	4%	4%
$\beta$ (MAF, proportionality constant for sorption capacity of NLOM)	4%	SC	1%	-2%
PCB concentration in porewater	3%	ES	2%	-3%
Organism weight	3%	PSS	2%	-2%
Porewater, fraction ventilated	2%	ES	2%	-2%
Water PCB concentration	2%	SS	2%	-2%
$\beta_{OC}$ (Proportionality constant for sorption capacity of NLOC)	1.8%	ES	1.2%	1.3%

PARAMETER	MAXIMUM SPD (absolute value)	SPECIES WITH MAXIMUM CHANGE	MINIMUM SPD (absolute value)	MEAN SPD (with negatives)
DOC concentration in water column	0.7%	SS	0.6%	0.6%
D <sub>DOC</sub> (disequilibrium factor for DOC partitioning)	0.7%	SS	0.6%	0.6%
α <sub>DOC</sub> (proportionality constant for DOC)	0.7%	SS	0.6%	0.6%
k <sub>M</sub>	0.5%	ES	0.2%	0.3%
POC concentration in water column	0.41%	SS	0.32%	0.37%
D <sub>POC</sub> (disequilibrium factor for POC partitioning)	0.41%	SS	0.32%	0.37%
α <sub>POC</sub> (proportionality constant for POC)	0.41%	SS	0.32%	0.37%
A (phytoplankton uptake constant)	0.07%	ES	0.04%	0.05%
B (phytoplankton uptake constant)	0.002%	ES	0.001%	0.001%
Dietary absorption efficiency of water (chi)	0.0003%	DC/SC	0.0002%	-0.0003%
Water density	0.000041%	PSS	0.00001%	-0.00001%

DC – Dungeness crab

ES – English sole

PSS – Pacific staghorn sculpin

SC – slender crab

SS – shiner surfperch

**Table B-3-2. Results of the 10 percent sensitivity analysis for all species**

PARAMETER	MAX SPD	SPECIES WITH MAX CHANGE	MIN SPD	MEAN SPD
Water content	102%	P	2%	19%
Dietary absorption efficiency of lipids (alpha)	-24%	PSS	0%	-9%
Lipid density	-17%	PSS	-1%	11%
Food ingestion rate (G <sub>D</sub> )	-14%	PSS	0%	-9%
Lipid content	-14%	PSS	0%	-9%
Dissolved oxygen (DO)	-11%	PSS	0%	-6%
Water PCB concentration	-10%	P/Z	1%	-4%
Water column temperature	-10%	PSS	0%	-6%
Dietary absorption efficiency of NLOM (beta)	-9%	DC	0%	-6%
Sediment PCB concentration	-9%	BI	0%	-6%
K <sub>OW</sub> -10%	-7%	PSS	3%	-5%
K <sub>OW</sub> +10%	6%	PSS	2%	4%

PARAMETER	MAX SPD	SPECIES WITH MAX CHANGE	MIN SPD	MEAN SPD
Growth rate constant (kG)	4.3%	ES	0.6%	2.7%
Sediment organic carbon (OC <sub>sed</sub> )	3.9%	ES	0.0%	2.9%
$\beta$ (MAF, proportionality constant for sorption capacity of NLOM)	-3.7%	SC	0.0%	-1.9%
$\alpha_{DOC}$ (proportionality constant for DOC)	3.5%	P/Z	0.5%	1.3%
DOC concentration in water column	3.5%	P/Z	0.5%	1.3%
D <sub>DOC</sub> (disequilibrium factor for DOC partitioning)	3.5%	P/Z	0.5%	1.3%
Chemical concentration in porewater	-2.7%	ES	0.0%	-2.0%
Weight	-2.5%	PSS	0.0%	-1.4%
A (phytoplankton/algae uptake constant)	2.5%	P	0.0%	0.4%
Porewater, fraction ventilated	-2.4%	ES	0.0%	-1.8%
POC	2.0%	P/Z	0.3%	0.7%
D <sub>POC</sub> (disequilibrium factor for POC partitioning)	2.0%	P/Z	0.3%	0.7%
$\alpha_{POC}$ (proportionality constant for POC)	2.0%	P/Z	0.3%	0.7%
$\beta_{OC}$ (proportionality constant for sorption capacity of NLOC)	1.8%	ES	0.0%	1.1%
K <sub>m</sub>	0.5%	ES	0.0%	0.2%
B (phytoplankton/algae uptake constant)	0.05%	P	0.0%	0.01%
Dietary absorption efficiency of water ( $\chi$ )	-0.0003%	DC/SC	0.0%	-0.0002%
Water density	0.00014%	Z	0.00001%	0.00003%

BI – benthic invertebrate  
 DC – Dungeness crab  
 ES – English sole  
 JF – juvenile fish  
 P – phytoplankton/algae  
 PSS – Pacific staghorn sculpin  
 SC – slender crab  
 SS – Shiner surfperch  
 Z – zooplankton

**Table B-3-3. Results of the plausible range sensitivity analysis for predicted fish and crab total PCB concentrations**

PARAMETER	MAXIMUM SPD (absolute value)	SPECIES WITH MAXIMUM CHANGE	MINIMUM SPD (absolute value)	MEAN SPD (with negatives)
Dietary absorption efficiency of lipids (alpha) (upper)	67%	DC	1%	20%
Dietary absorption efficiency of lipids (alpha) (lower)	54%	DC	3%	-19%
Dietary absorption efficiency of NLOM (beta) (lower)	43%	DC	22%	-29%
Sediment PCB concentration (upper)	42%	SC	40%	41%
Sediment PCB concentration (lower)	42%	SC	40%	-41%
Lipid content (upper)	33%	DC	11%	16%
Lipid content (lower)	31%	DC	11%	-16%
Dietary absorption efficiency of NLOM (beta) (upper)	28%	DC	12%	18%
Weight (lower)	25%	DC	16%	-19%
Lipid density (lower)	20%	PSS	12%	15%
Porewater, fraction ventilated (lower)	17%	ES	16%	-17%
Weight (upper)	17%	DC	13%	15%
Lipid density (upper)	15%	PSS	9%	-12%
Temperature (upper)	12%	PSS	8%	10%
Temperature (lower)	12%	PSS	8%	-9%
Water PCB concentration (upper)	11%	SS	9%	10%
$\beta$ (MAF) upper	11%	SC	3%	6%
$\beta$ (MAF) lower	11%	SC	4%	-6%
DO (lower)	10%	PSS	6%	-8%
DO (upper)	10%	PSS	6%	8%
Porewater, fraction ventilated (upper)	8%	ES	6%	6%
$\alpha$ DOC (proportionality constant for DOC) (upper)	7%	SS	5%	-6%
$K_{ow}$ (lower)	6%	PSS	3%	-5%
$K_{ow}$ (upper)	6%	PSS	3%	4%
$\alpha$ DOC (proportionality constant for DOC) (lower)	6%	SS	4%	5%
Water PCB concentration (lower)	5%	SS	4%	-5%
$\alpha$ POC (proportionality constant for POC) (upper)	5%	SS	4%	-4%
Water content (lower)	4%	JF	0%	2%
Water content (upper)	4%	SC	0%	-2%

PARAMETER	MAXIMUM SPD (absolute value)	SPECIES WITH MAXIMUM CHANGE	MINIMUM SPD (absolute value)	MEAN SPD (with negatives)
OC <sub>sed</sub> (lower)	3%	ES	3%	3%
αPOC (proportionality constant for DOC) (lower)	3%	SS	2%	2%
POC (lower)	2%	SS	2%	2%
OC <sub>sed</sub> (upper)	1.9%	ES	1.8%	-1.8%
POC (upper)	1.5%	SS	1.2%	-1.4%
DOC (lower)	1.4%	SS	1.0%	1.2%
DOC (upper)	0.91%	SS	0.71%	-0.81%
A (lower)	0.26%	ES	0.15%	0.19%
A (upper)	0.22%	ES	0.13%	-0.16%
B (lower)	0.010%	ES	0.006%	0.008%
B (upper)	0.010%	ES	0.006%	-0.008%
Water density (upper) (seawater)	0.000007%	PSS	0.000001%	0.000002%

DC – Dungeness crab

ES – English sole

JF – juvenile fish

PSS – Pacific staghorn sculpin

SC – slender crab

SS – shiner surfperch

**Table B-3-4. Results of the plausible range sensitivity analysis for all species**

PARAMETER	MAX SPD	SPECIES WITH MAX CHANGE	MIN SPD	MEAN SPD
Dietary absorption efficiency of lipids (alpha) (upper)	67%	DC	0%	14%
Water PCB concentration (upper)	55%	P/Z	7%	20%
Dietary absorption efficiency of lipids (alpha) (lower)	-54%	DC	0%	-13%
Sediment PCB concentration (upper)	43%	BI	0%	32%
Sediment PCB concentration (lower)	-43%	BI	0%	-32%
Dietary absorption efficiency of NLOM (beta) (lower)	-43%	DC	0%	-23%
$\alpha_{DOC}$ (proportionality constant for DOC) (upper)	-34%	P/Z	-5%	-12%
Lipid content (upper)	33%	DC	1%	15%
Lipid content (lower)	-31%	DC	-1%	-14%
Dietary absorption efficiency of NLOM (beta) (upper)	28%	DC	0%	14%
$\alpha_{DOC}$ (proportionality constant for DOC) (lower)	27%	P/Z	4%	10%
Water PCB concentration (lower)	-25%	P	-3%	-9%
Weight (lower)	-25%	DC	-0%	-15%
$\alpha_{POC}$ (proportionality constant for POC) (upper)	-23%	P/Z	-3%	-8%
Lipid density (lower)	20%	PSS	1%	12%
Porewater, fraction ventilated (lower)	-17%	ES	0%	-13%
Weight (upper)	17%	DC	0%	11%
Water content (upper)	-15%	P	0%	-4%
Lipid density (upper)	-15%	PSS	-1%	-10%
Water content (lower)	14%	P	0%	4%
$\alpha_{POC}$ (proportionality constant for POC) (lower)	13%	P/Z	2%	5%
Temperature water column (upper)	12%	PSS	0%	7%
Temperature water column (lower)	-12%	PSS	0%	-7%
$\beta$ (MAF, proportionality constant for sorption capacity of NLOM) (upper)	11%	SC	0%	5%
$\beta$ (MAF, proportionality constant for sorption capacity of NLOM) (lower)	-11%	SC	0%	-5%
Dissolved oxygen (lower)	-10%	PSS	0%	-6%

PARAMETER	MAX SPD	SPECIES WITH MAX CHANGE	MIN SPD	MEAN SPD
Dissolved oxygen (upper)	10%	PSS	0%	6%
POC (lower)	10%	P/Z	1%	3%
A (phytoplankton/algae uptake constant) (lower)	9%	P/Z	0%	1%
Porewater, fraction ventilated (upper)	8%	ES	0%	5%
A (phytoplankton/algae uptake constant) (upper)	-8%	P/Z	0%	-1%
POC (upper)	-8%	P/Z	-1%	-3%
DOC (lower)	7%	P/Z	1%	2%
K <sub>OW</sub> (lower)	-6%	PSS	-2%	-4%
K <sub>OW</sub> (upper)	6%	PSS	2%	4%
DOC (upper)	-4%	P/Z	-1%	-2%
OC <sub>sed</sub> (lower)	3%	ES	0%	2%
OC <sub>sed</sub> (upper)	-2%	ES	0%	-1%
B (phytoplankton/algae uptake constant) (lower)	0.36%	P	0.01%	0.06%
B (phytoplankton/algae uptake constant) (upper)	-0.36%	P	-0.01%	-0.06%
Water density (upper) (seawater)	-0.00003%	Z	-0.00000%	-0.00001%

**Table B-3-5. Relative response ratio for upper and lower bound sensitivity analyses for all species**

PARAMETER	RELATIVE RESPONSE		RESPONSE TO CHANGES IN INPUT VALUES			% CHANGE IN PARAMETER INPUT VALUES	
	MAXIMUM <sup>a</sup>	MEAN <sup>b</sup>	MAXIMUM SPD	SPECIES WITH MAXIMUM SPD	MEAN SPD	MEAN	MAXIMUM <sup>c</sup>
Water content (lower)	-15.6	-2.1	14%	P	4%	-2%	-0.9%
Water content (upper)	-15.0	-2.5	-15%	P	-4%	2%	1.0%
Dietary absorption efficiency of Lipids (alpha) (upper)	2.4	0.6	67%	DC	14%	23%	28%
Lipid density (lower)	-1.8	-1.1	20%	PSS	12%	-11%	
Lipid density (upper)	-1.4	-0.9	-15%	PSS	-10%	11%	
Dissolved Oxygen (DO) (upper)	1.1	0.7	10%	PSS	6%	9%	
Dissolved Oxygen (DO) (lower)	1.1	0.7	-10%	PSS	-6%	-9%	
Water column temperature (upper)	1.1	0.6	12%	PSS	7%	11%	
Water PCB concentration (upper)	1.0	0.4	55%	P/Z	20%	55%	
Dietary absorption efficiency of NLOM (beta) (upper)	1.0	0.8	28%	DC	14%	17%	28%

PARAMETER	RELATIVE RESPONSE		RESPONSE TO CHANGES IN INPUT VALUES			% CHANGE IN PARAMETER INPUT VALUES	
	MAXIMUM <sup>a</sup>	MEAN <sup>b</sup>	MAXIMUM SPD	SPECIES WITH MAXIMUM SPD	MEAN SPD	MEAN	MAXIMUM <sup>c</sup>
Water PCB concentration (lower)	1.0	0.4	-25%	P	-9%	-25%	
Water column temperature (lower)	1.0	0.6	-12%	PSS	-7%	-12%	
Sediment PCB concentration (lower)	0.9	0.6	-43%	BI	-32%	-50%	
Sediment PCB concentration (upper)	0.9	0.6	43%	BI	32%	50%	
Lipid content (upper)	0.9	0.9	33%	DC	15%	18%	39%
Lipid content (lower)	0.8	0.9	-31%	DC	-14%	-16%	-39%
K <sub>ow</sub> (lower)	0.7	0.5	-6%	PSS	-4%	-9%	
Dietary absorption efficiency of Lipids (alpha) (lower)	0.7	0.3	-54%	DC	-13%	-52%	-80%
K <sub>ow</sub> (upper)	0.6	0.4	6%	PSS	4%	10%	
Dietary absorption efficiency of NLOM (beta) (lower)	0.5	0.6	-43%	DC	-23%	-36%	-80%
αDOC (proportionality constant for DOC) (lower)	-0.4	-0.2	27%	P/Z	10%	-63%	
β (MAF - proportionality constant for sorption capacity of NLOM) (lower)	0.4	0.2	-11%	SC	-5%	-29%	
β (MAF - proportionality constant for sorption capacity of NLOM) (upper)	0.4	0.2	11%	SC	5%	29%	
DOC (lower)	-0.4	-0.1	7%	P/Z	2%	-18%	
OCsed (lower)	-0.4	-0.3	3%	ES	2%	-8%	
OCsed (upper)	-0.4	-0.2	-2%	ES	-1%	6%	
Porewater, fraction ventilated (lower)	0.3	0.2	-17%	ES	-13%	-55%	-50%
Weight (lower)	0.3	0.5	-25%	DC	-15%	-29%	-77%
Weight (upper)	0.3	0.2	17%	DC	11%	57%	55%
DOC (upper)	-0.3	-0.1	-4%	P/Z	-2%	14%	
A (phytoplankton/algae uptake constant) (lower)	0.3	0.03	9%	P/Z	1%	33%	
A (phytoplankton/algae uptake constant) (upper)	0.24	0.03	-8%	P/Z	-1%	-33%	



PARAMETER	RELATIVE RESPONSE		RESPONSE TO CHANGES IN INPUT VALUES			% CHANGE IN PARAMETER INPUT VALUES	
	MAXIMUM <sup>a</sup>	MEAN <sup>b</sup>	MAXIMUM SPD	SPECIES WITH MAXIMUM SPD	MEAN SPD	MEAN	MAXIMUM <sup>c</sup>
$\alpha_{\text{DOC}}$ (proportionality constant for DOC) (upper)	-0.23	-0.08	-34%	P/Z	-12%	150%	
POC (lower)	-0.22	-0.07	10%	P/Z	3%	-45%	
$\alpha_{\text{POC}}$ (proportionality constant for POC) (lower)	-0.22	-0.08	13%	P/Z	5%	-60%	
POC (upper)	-0.19	-0.07	-8%	P/Z	-3%	41%	
$\alpha_{\text{POC}}$ (proportionality constant for POC) (upper)	-0.15	-0.05	-23%	P/Z	-8%	149%	
Porewater, fraction ventilated (upper)	0.08	0.07	8%	ES	5%	75%	100%
B (phytoplankton/algae uptake constant) (upper)	-0.01	-0.001	-0.36%	P	-0.06%	67%	
B (phytoplankton/algae uptake constant) (lower)	-0.01	-0.001	0.36%	P	0.06%	-67%	
Water density (upper) (seawater)	0.00002	-0.00001	0.00003%	Z	-0.00001 %	2%	

<sup>a</sup> Calculated as the maximum species percent difference divided by the mean or maximum percent change in parameter value. Maximum percent change is used for species-specific parameters only.

<sup>b</sup> Calculated as the mean species percent difference divided by the mean percent change in parameter value.

<sup>c</sup> Percent change for species-specific parameters only.

BI – benthic invertebrate

DC – Dungeness crab

ES – English sole

JF – juvenile fish

P – phytoplankton/algae

PSS – Pacific staghorn sculpin

SC – slender crab

SS – Shiner surfperch

Z – zooplankton

Based on the 10% sensitivity analysis, all parameters that had a maximum SPD equal to or greater than 8% were selected for inclusion in the uncertainty analysis (Tables B-3-1, B-3-2).  $K_{\text{OW}}$  was selected for inclusion in the uncertainty analysis because it was close to the 8% threshold (7% SPD for Pacific staghorn sculpin) and because it is a key chemical-specific parameter with uncertainty. The food ingestion rate ( $G_{\text{D}}$ ) with a maximum SPD of 14% will not be included in the uncertainty analysis because it is calculated by an equation within the model, and Crystal Ball®, the software used to

run the Monte Carlo uncertainty analysis (Section 6.0), cannot test parameters defined by equations.

A 10% change in predicted tissue concentrations was considered sufficient to warrant inclusion in the uncertainty analysis. When maximum SPDs of the plausible range analysis were ranked for target species, 11 parameters were selected for inclusion in the uncertainty analysis. When maximum SPDs of the plausible range analysis were ranked for all species (Table B-3-4), three additional parameters had maximum SPDs of 10% or greater (water column PCB concentration,  $\alpha_{\text{DOC}}$ , and  $\alpha_{\text{POC}}$ ).  $\alpha_{\text{DOC}}$ , and  $\alpha_{\text{POC}}$  were not included in the uncertainty analysis because they are constants within equations in the model rather than true input parameters, and Crystal Ball®, the software used to run the Monte Carlo uncertainty analysis (Section 6.0) cannot test parameters within equations. The phytoplankton uptake constant (A) was included in the uncertainty analysis based on advice from Jon Arnot (2005). A complete discussion of sensitivity analysis results is presented in the main document.

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APPENDIX C. ASSIGNMENT OF PARAMETER  
DISTRIBUTIONS AND CORRELATIONS FOR  
UNCERTAINTY ANALYSIS

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## **Appendix C. Assignment of Parameter Distributions and Correlations for Uncertainty Analysis**

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As discussed in the Food Web Model (FWM) Memorandum 2 (Windward 2005), a probabilistic approach was employed to investigate model uncertainty. Specifically, Monte Carlo simulation was used to investigate the effects of parameter variability and uncertainty on model predictions. As described in Section 5.0 and Appendix B, sensitivity analyses were conducted to identify sensitive parameters. Because these parameters most affect model output, they were further investigated in the uncertainty analysis, as described in this appendix.

Distributions, rather than point estimates, were assigned for input parameters as appropriate (i.e., if data are available and the FWM is sensitive to a given parameter). In the Monte Carlo simulations, the FWM was run 10,000 times using Decisioneering® Crystal Ball 7® software. During each model iteration, different combinations of values for each input parameter were selected based on the probability distribution for each parameter. Output from this uncertainty analysis are distributions of the relative probability of predicted tissue concentrations for each species based on the distributions of FWM input parameter values. In addition to assigning distributions for parameters, correlations may also be defined to prevent improbable combinations of parameter values, such as the combination of an extremely high organism lipid content and an extremely high organism water content.

This appendix describes the process by which distributions were selected for parameters to which the model is sensitive, the assignment of distributions for these parameters, and the assignment of correlations between parameters.

### **C.1 APPROACH FOR DEVELOPMENT OF PARAMETER DISTRIBUTIONS**

As described in Section 6.2.1, the first step of developing the Monte Carlo version of the model is development of parameter distributions. Only parameters identified as sensitive in the sensitivity analysis were included in the uncertainty analysis. The following approach was used to develop parameter distributions:

- 1) When site-specific data were available, the distribution was determined through statistical analysis of the data.<sup>1</sup> Normal distributions were defined in Crystal Ball® by their mean and standard deviation and truncated at zero (with no upper-bound truncation). For non-parametric datasets, a custom

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<sup>1</sup> Distributions were evaluated using ProUCL software (EPA 2004). The distribution that the software determined to best fit the data was selected for use in Monte Carlo simulations.

distribution was assigned based directly on site-specific empirical data.<sup>2</sup> Crystal Ball® randomly selects parameter values from the empirical data for each model iteration.

- 2) A normal or lognormal distribution was assigned if the parameter was biological in nature and/or empirical data for similar parameters exhibited a normal or lognormal distribution (e.g., all other LDW fish species analyzed had normal lipid distributions, so juvenile fish were assigned a normal distribution in the absence of empirical LDW juvenile fish lipid data). The mean and standard deviation of the distributions of parameters from the literature were based on published data when possible.
- 3) For parameters with insufficient data to define a distribution, and/or available data did not conform to a normal or lognormal distribution, a triangle distribution was assigned (MacIntosh et al. 1994). The mode of the triangle was defined as the mean of the data if the data were considered sufficiently relevant and comprehensive. For more uncertain data, the mode was based on consideration of published selections for parameter values used in other food web models (Arnot and Gobas 2004; Gobas and Arnot 2005). The tails of the triangle were defined by the literature values if they were considered sufficient to bound a plausible range.

## **C.2 DEVELOPMENT OF PARAMETER DISTRIBUTIONS**

Distributions define the frequency for which certain parameter values will be used in the multiple iterations of the model. Distributions were developed for the majority of the parameters to which the model was found to be sensitive (Section 5.0 and Appendix B). Distributions were developed by species for species-specific parameters (e.g., Dungeness crab weight, slender crab weight). This section details the decision process for development of parameter distributions, including application of the above approach based on available data and professional judgment.

### **C.2.1 Sensitive parameters without distributions**

Distributions were not developed for eight of the parameters to which the model is sensitive. Four parameters investigated in the sensitivity analysis are calculated by the model. Thus, they could not be varied in an automated manner in the Monte Carlo version of the model and were not included in the uncertainty analysis. These parameters were growth rate, freely dissolved chemical concentration in porewater, the proportionality constant describing similarity in phase partitioning of dissolved

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<sup>2</sup> In the custom distributions used in this assessment, Crystal Ball draws randomly from empirical data. For example, the organism weights for shiner surfperch were assigned a custom distribution because the data were non-parametric. During model iterations, Crystal Ball chooses a value randomly from the 458 empirical values that define the custom distribution (Table C-2-1).

organic carbon (DOC) relative to that of octanol ( $\alpha_{\text{DOC}}$ ), and the proportionality constant describing similarity in phase partitioning of particulate organic carbon (POC) relative to that of octanol ( $\alpha_{\text{POC}}$ ).

Four of the parameters to which the model was found to be sensitive were species dietary absorption efficiencies of lipids for fish (for three fish species and juvenile fish). No distributions were assigned for fish dietary absorption efficiency of lipids because the range of empirical values was too small to define a distribution.

### **C.2.2 Sensitive parameters for which distributions were developed**

Distributions were developed for the remaining 45 parameters identified in the sensitivity analysis and are presented in Table C-2-1. The following sections describe the development of distributions for these parameters, which was dependent on available data for each parameter and consistent with the approaches described in Section C.1.

**Table C-2-1. Uncertainty analysis input values**

PARAMETER DESCRIPTION	UNITS	DISTRIBUTION TYPE	DISTRIBUTION VALUES	CORRELATION COEFFICIENT	SOURCE/NOTES
<b>Environmental Parameters</b>					
Total concentration of PCBs in the water column	ng/L	normal	Mean = 2, SD = 0.4	id	King County water data (2005), standard deviation was selected to include range of empirical data..
Concentration of particulate organic carbon (POC) in the water column	kg/L	normal	mean = $2.9 \times 10^{-7}$ , SD = $6.3 \times 10^{-8}$	with water temperature = 0.0087; with dissolved oxygen = 0.23	Unpublished King County 2005 water data (Mickelson 2006); used standard error from raw data as standard deviation for distribution of estimates of the mean. POC is calculated based on TOC and DOC as described in Appendix A, Table A-2-1. Correlation based on individual samples from 2 locations taken on 10 occasions.
Mean water column temperature	°C	normal	mean = 11.6, SD = 0.0678	with POC = 0.0087; with dissolved oxygen = -0.34	Unpublished King County 2005 water (Mickelson 2006); used standard error from raw data as standard deviation for distribution of estimates of the mean. . Correlation based on individual samples from 2 locations taken on 10 occasions.
Dissolved oxygen concentration in the water column	mg/L	normal	mean = 8.0, SD = 0.36	with POC = 0.0087; with water temperature = -0.34	Unpublished King County 2005 water data (Mickelson 2006); used standard error from raw data as standard deviation for distribution of estimates of the mean. . Correlation based on individual samples from 2 locations taken on 10 occasions.
Concentration of PCBs in sediment	µg/kg dw	normal	mean = 250, SD = 64		SWAC developed from Phase 1 and 2 data as described in Appendix A, section A.2. Standard deviation calculated as described in Appendix C, section C.2.1.2.



**Table C-2-1, cont.**

PARAMETER DESCRIPTION	UNITS	DISTRIBUTION TYPE	DISTRIBUTION VALUES	CORRELATION COEFFICIENT	SOURCE/NOTES
<b>Chemical Parameters</b>					
Octanol-water partition coefficient for PCBs (log $K_{ow}$ )	unitless	normal	Mean = 6.62, SD = 0.186	na	$K_{ow}$ s for each congener from Hawker and Connell (1988); used standard error from raw data as standard deviation for distribution of estimates of the mean.
<b>Biological Parameters</b>					
Proportionality constant expressing the sorption capacity of NLOM relative to that of octanol ( $\beta$ or MAF)	unitless	normal	mean = 0.035, SD = 0.005	na	Mean from Arnot and Gobas (2004); SD from Arnot (2005).
Resistance to chemical uptake through aqueous phase for phytoplankton/algae (A)	day <sup>-1</sup>	normal	mean = $6 \times 10^{-5}$ , SD = $1 \times 10^{-5}$	na	Gobas and Arnot (2005)
Density of lipids	kg/L	triangle	mode = 0.9, min = 0.8, max = 1		Arnot (2006)
<b>Phytoplankton</b>					
Lipid content of organism	%	normal	mean = 0.12, SD = 0.05	id	Mackintosh et al. (2004)
Water content of organism	%	normal	mean = 95.6, SD = 0.55	id	Mackintosh et al. (2004); SD selected to include range in Mackintosh et al. (2004).
<b>Zooplankton</b>					
Organism weight	kg	normal	mean = $1.6 \times 10^{-7}$ , SD = $3.6 \times 10^{-8}$	id	Giles and Cordell (1998); SD selected to include range in Giles and Cordell (1998).
Lipid content	%	normal	mean = 1.2, SD = 0.3	id	Kuroshima et al. (1987); SD selected to include range in Kuroshima et al. (1987).
Water content of organism	%	normal	mean = 90, SD = 1.5	id	Kuroshima et al. (1987); SD selected to include range in Kuroshima et al. (1987).
Dietary absorption efficiency of lipids (alpha)	%	triangle	mode = 72, min = 55, max = 85	id	Arnot and Gobas (2004)

**Table C-2-1, cont.**

PARAMETER DESCRIPTION	UNITS	DISTRIBUTION TYPE	DISTRIBUTION VALUES	CORRELATION COEFFICIENT	SOURCE/NOTES
Dietary absorption efficiency of NLOM (beta)	%	triangle	mode = 72, min = 55, max = 85	id	Arnot and Gobas (2004)
<b>Benthic Invertebrates</b>					
Organism weight	kg	custom <sup>a</sup>	mean = $5.1 \times 10^{-5}$ , raw data	id	LDWG Phase 2 data (n = 10); average from each sample location.
Lipid content	%	normal	mean = 0.89, SD = 0.26	with water content = -0.57	LDWG Phase 2 data.
Water content of organism	%	normal	mean = 88.9, SD = 3.6	with lipid content = -0.57	LDWG Phase 2 data.
Relative fraction of porewater ventilated	unitless	triangle	mode = 0.2, min = 0.05, max = 0.25	id	Winsor et al. (1990)
Dietary absorption efficiency of lipids (alpha)	%	triangle	mode = 75, min = 15, max = 96	id	Arnot and Gobas (2004)
Dietary absorption efficiency of NLOM (beta)	%	triangle	mode = 75, min = 15, max = 96	id	Arnot and Gobas (2004)
<b>Juvenile Fish</b>					
Organism weight	kg	normal	mean = $6 \times 10^{-3}$ , SD = $7 \times 10^{-4}$	id	Derived from all Phase 2 (2004 and 2005) individual $\leq 80$ mm shiner surfperch specimens from the LDW and background locations.
Lipid content	%	normal	mean = 2.5, SD = 0.6	id	LDWG Phase 2 data
Water content of organism	%	normal	mean = 73.9, SD = 2.0	id	LDWG Phase 2 data
Relative fraction of porewater ventilated	unitless	triangle	mode = 0.01, min = 0.005, max = 0.02	id	Gobas and Wilcockson (2003); Gobas and Arnot (2005)
Dietary absorption efficiency of NLOM (beta)	%	normal	mean = 60, SD = 3	id	Arnot and Gobas (2004)

**Table C-2-1, cont.**

PARAMETER DESCRIPTION	UNITS	DISTRIBUTION TYPE	DISTRIBUTION VALUES	CORRELATION COEFFICIENT	SOURCE/NOTES
<b>Slender Crab</b>					
Organism weight	kg	normal	mean = 0.164, SD = 0.0318	id	LDWG Phase 2 data.
Lipid content	%	custom <sup>a</sup>	mean = 1.1, raw data	with water content = 0.6	LDWG Phase 2 data (n = 5).
Water content of organism	%	normal	mean = 83.6, SD = 1.19	with lipid content = 0.6	LDWG Phase 2 data.
Relative fraction of porewater ventilated	unitless	triangle	mode = 0.02, min = 0.01, max = 0.03	id	Gobas and Wilcockson (2003); Gobas and Arnot (2005); Winsor et al. (1990)
Dietary absorption efficiency of lipids (alpha)	%	triangle	mode = 75, min = 15, max = 96	id	Arnot and Gobas (2004)
Dietary absorption efficiency of NLOM (beta)	%	triangle	mode = 75, min = 15, max = 96	id	Arnot and Gobas (2004)
<b>Dungeness Crab</b>					
Organism weight	kg	custom <sup>a</sup>	mean = 0.423, raw data	id	LDWG Phase 2 data (n = 51).
Lipid content	%	normal	mean = 2.6, SD = 1.4	with water content = -0.93	LDWG Phase 1 and 2 data.
Water content of organism	%	normal	mean = 82, SD = 2.9	with lipid content = -0.93	LDWG Phase 1 and 2 data.
Relative fraction of porewater ventilated	unitless	triangle	mode = 0.02, min = 0.01, max = 0.03	id	Gobas and Wilcockson (2003); Gobas and Arnot (2005); Winsor et al. (1990)
Dietary absorption efficiency of lipids (alpha)	%	triangle	mode = 75, min = 15, max = 96	id	Arnot and Gobas (2004)
Dietary absorption efficiency of NLOM (beta)	%	triangle	mode = 75, min = 15, max = 96	id	Arnot and Gobas (2004)

**Table C-2-1, cont.**

PARAMETER DESCRIPTION	UNITS	DISTRIBUTION TYPE	DISTRIBUTION VALUES	CORRELATION COEFFICIENT	SOURCE/NOTES
<b>Pacific Staghorn Sculpin</b>					
Organism weight	kg	custom <sup>a</sup>	mean = 0.060, raw data	id	LDWG Phase 2 data (n = 272).
Lipid content	%	normal	mean = 2.1, SD = 0.37	with water content = -0.80	LDWG Phase 2 data.
Water content of organism	%	normal	mean = 79.0, SD = 0.602	with lipid content = -0.80	LDWG Phase 2 data.
Relative fraction of porewater ventilated	unitless	triangle	mode = 0.05, min = 0.02, max = 0.1	id	
All fish dietary absorption efficiency of NLOM (beta)	%	normal	mean = 60, SD = 3	id	Arnot and Gobas (2004)
<b>Shiner Surfperch</b>					
Organism weight	kg	custom <sup>a</sup>	mean = 0.017, raw data	id	LDWG Phase 2 data (n = 458).
Lipid content	%	normal	mean = 4.6, SD = 1.4	with water content = -0.83	LDWG Phase 1 and 2 data.
Water content of organism	%	normal	mean = 73.9, SD = 2.03	with lipid content = -0.83	LDWG Phase 2 data.
Relative fraction of porewater ventilated	unitless	triangle	mode = 0.01, min = 0.005, max = 0.02	id	Gobas and Wilcockson (2003); Gobas and Arnot (2005)
All fish dietary absorption efficiency of NLOM (beta)	%	normal	mean = 60, SD = 3	id	Arnot and Gobas (2004)
<b>English Sole</b>					
Organism weight	kg	custom <sup>a</sup>	mean = 0.198, raw data		LDWG Phase 2 data (n = 245).
Lipid content	%	normal	mean = 5.5, SD = 1.3	with water content = -0.76	LDWG Phase 2 data.
Water content of organism	%	normal	mean = 75.0, SD = 1.77	with lipid content = -0.76	LDWG Phase 2 data.

**Table C-2-1, cont.**

PARAMETER DESCRIPTION	UNITS	DISTRIBUTION TYPE	DISTRIBUTION VALUES	CORRELATION COEFFICIENT	SOURCE/NOTES
Relative fraction of porewater ventilated	unitless	triangle	mode = 0.1, min = 0.005, max = 0.2	id	Gobas and Wilcockson (2003); Gobas and Arnot (2005)
All fish dietary absorption efficiency of NLOM (beta)	%	normal	mean = 60, SD = 3	id	Arnot and Gobas (2004)

SD – standard deviation

<sup>a</sup> In the custom distributions used in this assessment, Crystal Ball<sup>®</sup> draws randomly from empirical data. During model iterations, Crystal Ball<sup>®</sup> chooses a value randomly from the empirical values that define the custom distribution.

id – inadequate data to evaluate correlations

na – not applicable; no expected correlations with other parameters with distributions

### **C.2.2.1 Biological parameters**

There were two non-species-specific biological parameters to which the model was found to be sensitive. Both the proportionality constant expressing the sorption capacity of non-lipid organic matter relative to that of octanol ( $\beta$ ) and the resistance to chemical uptake through the aqueous phase for phytoplankton/algae (A) were assigned normal distributions. These distributions and their means and standard deviations were based on published models, modeling reports and personal communication with Jon Arnot (Arnot 2005; Arnot and Gobas 2004; Gobas and Arnot 2005).

The majority of site-specific, species-specific lipid content and water content data were normally distributed. Thus, normal distributions based on empirical data were assigned for most of these parameters. One exception was slender crab lipid content, which had a non-parametric distribution. A custom distribution was assigned for this parameter. For species without site-specific water content and lipid data, parameter distributions were assumed to be normal.

For some species, empirical weight data were normally distributed; thus, these weight parameters were assigned normal distributions. For other species, distributions of weight data were not normal or lognormal. This may have occurred because only organisms of a minimum size were targeted during sampling. For species with weight distributions that were not normal or lognormal, custom distributions were assigned.

Few data were available for dietary absorption efficiencies of lipids and NLOM, density of lipids, and for ventilation of porewater. Lacking empirical data to describe distributions, the ranges for these parameters developed for the sensitivity analysis (Section 5.0 and Appendix B) were used for the uncertainty analysis. These ranges were primarily from Arnot and Gobas (2004). For crabs, benthic invertebrates, phytoplankton, and zooplankton, triangle distributions were assigned for dietary absorption efficiencies of lipids and NLOM because the available values from the literature were not consistent with normal or lognormal distributions (e.g., the minimum value was four times lower than the mean, while the maximum was only 25% higher than the mean). Similarly, the fractions of porewater ventilated for all species were assigned triangle distributions. Density of lipids was also assigned a triangle distribution with the same range as used in the sensitivity analysis and as recommended by Jon Arnot (2006). The dietary absorption of NLOM for fish was assigned a normal distribution consistent with the range used in the sensitivity analysis.

### **C.2.1.2 Environmental and Chemical Parameters**

The concentration of POC in the water column, the mean water column temperature, and the dissolved oxygen concentration in the water column were all assigned normal distributions. Because estimates of the mean are normally distributed (central

limit theorem), a normal distribution was assigned for these parameters. The standard error of the original distribution can be used to define a confidence interval on the mean, similar to the way a standard deviation may be used to define a confidence interval for normally distributed data. Thus, the standard error of the empirical data was used as the standard deviation of the distribution of the estimates of the mean.

Fewer empirical data were available for concentration of PCBs in water than for conventional water parameters such as temperature, and dissolved oxygen. Because the mean PCB water concentration was of interest, a normal distribution was assigned. A standard deviation was selected to include the range of empirical data.

The mean of log  $K_{ow}$  was of interest because the model uses only one  $K_{ow}$  for all model calculations for all species. Thus, the average of available data across species was most appropriate. A normal distribution was assigned for the range of estimates of the mean, with a standard deviation equal to the standard error of the estimated of site specific  $K_{ow}$  data (described in Section A.2.6)

The total PCB concentration in sediment was also assigned a normal distribution because it is the distribution of estimates of the mean that are of interest. This is again based on the central limit theorem, which states that estimates of the mean are normally distributed. The standard deviation was selected to describe a 95% confidence interval of the range selected for the sensitivity analysis (125  $\mu\text{g}/\text{kg dw}$  as 5th percentile and 375  $\mu\text{g}/\text{kg dw}$  as 95th percentile, Section B.2.2).

### **C.3. APPROACH AND ASSIGNMENT OF PARAMETER CORRELATIONS**

Correlations were assigned between some of the parameters for which distributions were developed to prevent improbable combinations of values for related parameters. To evaluate correlations, data must be available that can be reasonably matched (in time and location, or by sample specimens) and must be similarly robust in terms of number of samples and data quality. For parameter pairs expected to be correlated for biological or environmental reasons, a correlation test was performed if possible. The calculated correlation coefficients were then included and applied to both parameters in the Monte Carlo model.

Table C-2-1 identifies the parameters found to be correlated and the magnitude of correlation. For many of the uncertain parameters for which correlations were biologically plausible, insufficient data were available to assess correlation (e.g., weight data were collected for each organism, but lipid data were based on composite samples). All parameters for which correlation might be plausible, but for which data were insufficient to evaluate correlations, are noted in Table C-2-1 in the correlation column. Correlation was assessed for several water quality parameters (i.e., POC, mean water temperature, and dissolved oxygen concentration in the water column), and correlation coefficients were included in the model.

For biological parameters, correlations could be assessed only for water and lipid contents for fish, crabs, and benthic invertebrates. Correlations between water and lipid contents were primarily negative, as expected. One exception was slender crab, which exhibited a positive correlation between water and lipid contents. Slender crab lipid and water content data for the model are presented as “whole-body” crabs, but these data are actually calculated based on edible meat and hepatopancreas composite samples (for more detailed description, see Appendix A, Section A.2.3). Correlations for slender crab edible meat lipid and water contents and for hepatopancreas lipid and water contents were also found to be positive, although the sample sizes were small (edible meat  $n = 13$ , hepatopancreas  $n = 5$ ).

#### **C.4 REFERENCES**

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