

Beaverlodge Scenario Approach Using RESRAD-BIOTA

Dr. Sunita Kamboj and Charley Yu

Environmental Science Division Argonne National Laboratory

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Beaver Lodge Exercise Phase II

- Provided water or/and sediment concentrations for five radionuclides (Pb-210, Po-210, Ra-226, Th-230, and U-238).
- Measured activity concentration for fish species provided in some cases
- The samples were collected from 1995 2008 at 6 areas at 15 sites
- Six areas are:
 - Beaver lodge Lake
 - Cluff Lake
 - Gunnar Mine Site
 - Key Lake
 - McClean Lake
 - Rabbit Lake
- Calculate <u>whole body activity concentration</u> and <u>weighted absorbed dose</u> <u>rates</u> for fish and invertebrates.

Methodology and Assumptions

- Used Probabilistic RESRAD-BIOTA
- Used distributions for water concentration, sediment concentration, and tissue concentrations (if available)
- If water or sediment concentration not available used conservative Kd values to estimate activity concentrations
- Used probabilistic Biv values

Input Parameters for Model Run

DOSIMETRY - Weighting factors for absorbed dose rates

		to be consistent with other
Internal alpha	1.00E+01	models
Internal beta gamma	1.00E+00	
Internal low beta	1.00E+00	

DOSIMETRY - Daughters explicitly included

		Po-210			
Pb210	Bi-210 (100%)	(100%)			
Po210	Х	Х	Х	Х	Х
				Pb-214 & Po-	
Ra226	Rn-222	Po-218	Bi-214	214	At-218 & TI-210
Th230					
U-238	Th-234	Pa-234	Pa-234m		
U-235	Th-231				
U-234					

It was assumed U-238, U-234 and U-235 were present in their natural activity concentration ratio of 1:1:0.046.

Pb-210: Bi-210 and Po-210 was assumed to be in secular equilibrium with Pb-210.

Ra-226: Rn-222, Po-218, Bi-214 at the same conc. as Ra-226, Bi-214 and Po-214 at 99.98% of Ra-226 conc. And At-218 and TI-210 at 0.02% of Ra-226 conc.

U-238: Th-234 at the same concentration as U-238, Pa-234m at 99.8% and Pa-234 at 0.33% of U-238 concentration.

U-235: At 4.6% of U-238 concentration. Th-231 at the same concentration as U-235

U-234: At the same concentration as U-238

Input Parameters for Model Run, cont'd

GEOMETRY, MASS, & OCCUPANCY FACTORS

		Mass (g	Occupancy-	
	Geometry (cm)	ww)	Biota run	
	Length x		% water /	Geometry -
	height x width		%sediment	used
Pelagic (e.g. Northern pike &				
Lake Trout)	50 x 15 x 10	1200	87.5/12.5	4
Benthic Fish - Large (White				
sucker & Lake whitefish)	45 x 15 x 10	1191	65/35	4
Benthic Fish - Small (Lake				
chub)	6.8 x 1.5 x 1	4.5	90/10	Ext -2 and Int -3
Benthic Invertebrates	0.34 x 0.17 x			
(Chironomus riparius)	0.15	0.12	62.5/37.5	1
Benthic Invertebrates (Pisidium				
sp.)	2.5 x 1.5 x 1	1.6	75/25	2
Benthic Invertebrates (Caddisfly,				
Nemotaulius sp.)	3.5x1.46x1.46	1.75	75/25	2

Assumptions for Model Run

Species	Biv - used	Pb-210	Po-210	Ra-226	Th-230	Uranium
Pelagic (e.g. Northern pike				Bounded	Bounded	Bounded
& Lake Trout)	Pelagic fish	Exponential	Exponential	Lognormal	Lognormal	Lognormal
Benthic Fish - Large (White				Bonded	Bounded	Bounded
sucker & Lake whitefish)	Benthic fish	Exponential	Exponential	Lognormal	Lognormal	Lognormal
Benthic Fish - Small (Lake	Denthie fieh	E ver an ential	Evenential	Bonded	Bounded	Bounded
cnub)	Benthic fish	Exponential	Exponential	Lognormal	Lognormal	Lognormal
Benthic Invertebrates		Eveneratio	Europontial	Eveneratio	Evenential	Evenential
(Chironomus riparius)	Insect Larvae	Exponential	Exponential	Exponential	Exponential	Exponential
Benthic Invertebrates	Bivalve		Bounded	Bonded		
(Pisidium sp.)	Mollusc	Exponential	Lognormal	Lognormal	Exponential	Exponential
Benthic Invertebrates						
(Caddisfly, Nemotaulius sp.)	Insect Larvae	Exponential	Exponential	Exponential	Exponential	Exponential

*For exponential distribution parameter required is lambda (=1/mean) and for bounded lognormal distribution parameter required are mean, error factor, minimum, and maximum



RESRAD-BIOTA Inputs for Contamination at Beaverlodge Lake



Uncertainty Analysis Setup in the RESRAD-BIOTA Code

Jncertainty Analysis Input Summary						
Sample specifications Parameter distribution	ns Input Rank Correlations	Output specifications				
Sampling parameters	Information about current selection	Uncertainty Analysis Input S	Summary			
Random Seed: 1000	The random seed determines the serie	Sample specifications	Parameter o	listributions	Input Rank Correlations	Output specifications
Number of Observations: 1500	numbers that are generated. Specifica			Charles and the		
Number of <u>R</u> epetitions: 1	generated if the simulation needs to be	Variable De: Water BIV of Pb-210 in	scription Benthic fish-large	- Statistics of Unc	ertain variable	
		Water BIV of Pb-210 in	Benthic fish-small	Water BIV of Ha	-226 in Benthic fish-small	
		Water BIV of Pb-210 in	Bivalve-mollusc	Distribution	BOUNDED LOGNORMAL	✓ Default
		Water BIV of Pb-210 in	Insect-larvae			Mean 80
		Water BIV of Pb-210 in	pelagic fish		I	Error Factor (Erf) 6.5
-Sampling Technique		Water BIV of Po-210 in Water BIV of Po-210 in	Benthic fish-large Renthic fish-small			Minimum .3
 Latin Hypercube Monte Cada 		Water BIV of Po-210 in	Bivalve-mollusc			Maximum 810
O Monte cano		Water BIV of Po-210 in	Caddisfly			
		Water BIV of Po-210 in	pelagic fish			
		Water BIV of Ra-226 in	Benthic fish-large			
		Water BIV of Ha-226 in Water BIV of Ba-226 in	Benthic fish-small Rivalve-mollusc			
Grouping of observations		Water BIV of Ra-226 in	Caddisfly			
 Correlated or Uncorrelated 		Water BIV of Ra-226 in	Insect-larvae			
O Handom		Water BIV of Th-230 in	Benthic fish-large	Previous parame	ter 🔺	
		Water BIV of Th-230 in	Benthic fish-small	Next parameter	-	
 Perform uncertainty analysis C Suppress uncertainty 	ncertainty analysis this session Help	Water BIV of Th-230 in Water BIV of Th-230 in	Bivalve-mollusc Caddisflv	Remove	parameter Distribution H	lelp Restore Default
		Water BIV of Th-230 in	Insect-larvae			
		Perform uncertainty ana	ilysis O Su	ppress uncertainty	analysis this session	lelp <u>O</u> K