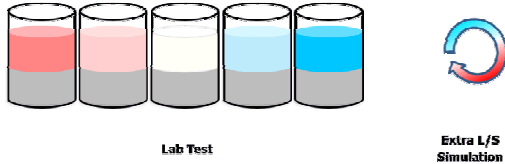


Object Name pH Dependent Leaching Test Model
Marine harbour sediment IT

pH Dependent Leaching Test Scenario



Lab Test

Model Parameters

Entity	Unit	Default	Entity	mg/kg	Entity	mg/kg	Entity	mg/kg
c0		-4.620	Al	2895	B	17.64	PO4	2749
c1		-0.3602	As	3.300	Si	1506	Sb	0.1462
c2		0.02238	Ba	18.54	Hg	2.006E-07	Se	0.1038
c3		-0.001288	Ca	8.060E+04	K	565.0	Sn	0.08355
c4		0.0003533	Cd	0.1088	Li	4.886	SO4	1.244E+04
c5		-1.815E-05	Cl	1500	Mg	2.695E+04	Sr	117.3
Clay	mg/kg	3000	Co	5.899	Mn	167.5	V	7.376
Hydrous Ferric Oxid	mg/kg	1700	CO32-	1.844E+05	Mo	0.9057	Zn	435.3
L/S	L/kg	10.75	Cr	3.972	Na	7764		
pE		1.710	Cu	0.5200	Ni	4.092		
pH		8.641	F	10.00	NO3	6.200E-08		
Solid Humic Acid	mg/kg	1224	Fe	7618	Pb	7.389		
Simulated Low L/S	L/kg	0.4000						

Solid Solutions

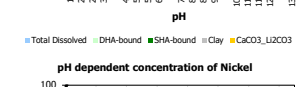
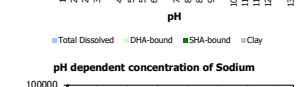
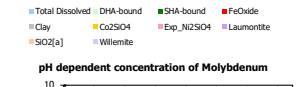
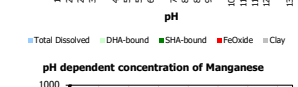
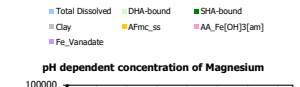
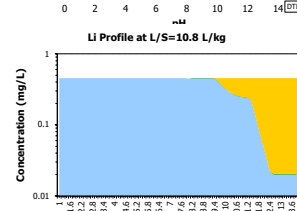
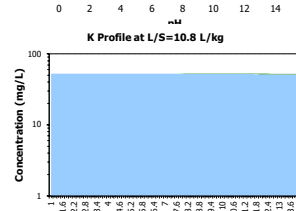
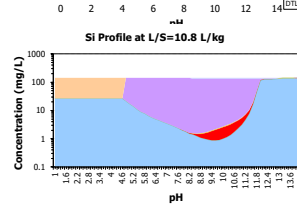
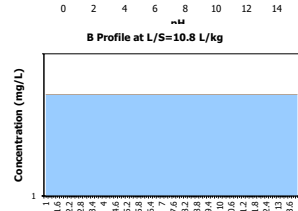
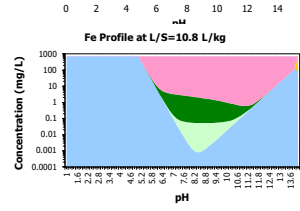
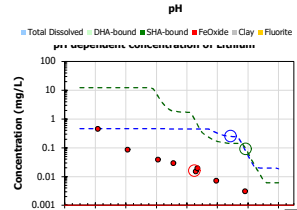
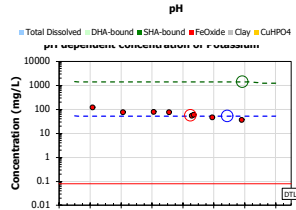
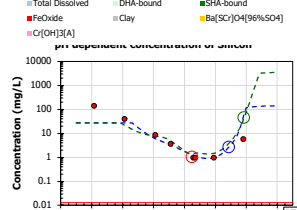
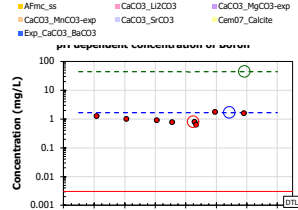
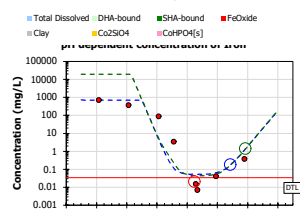
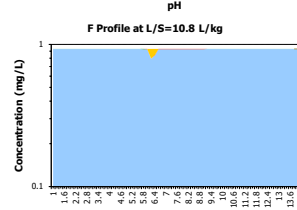
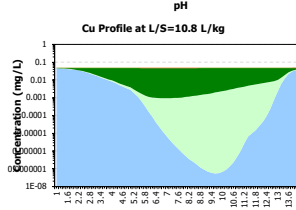
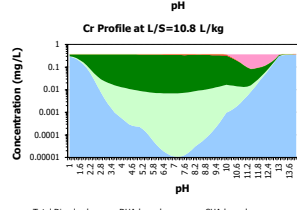
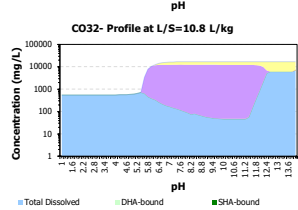
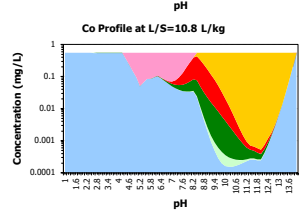
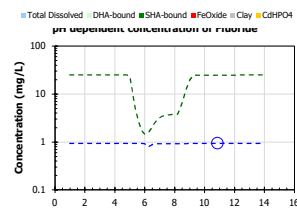
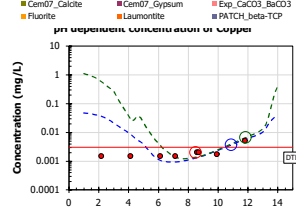
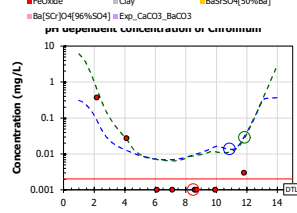
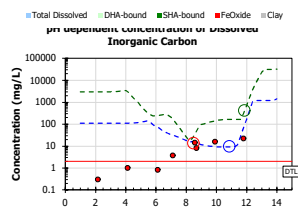
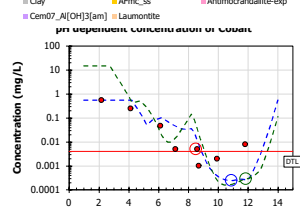
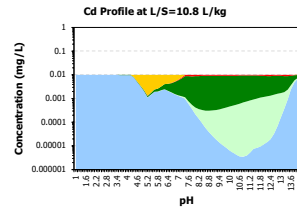
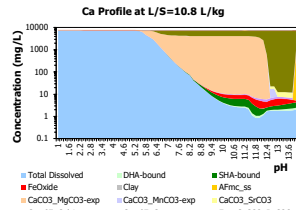
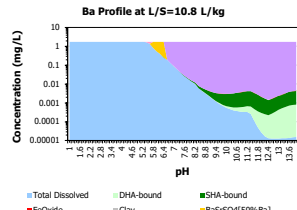
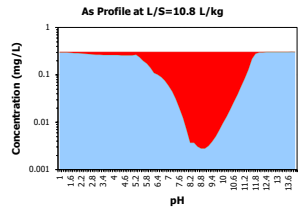
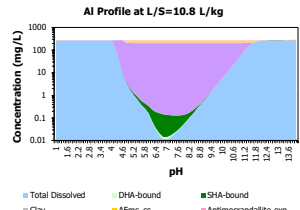
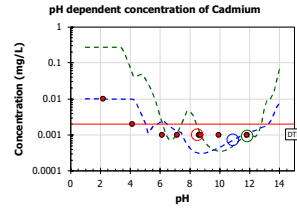
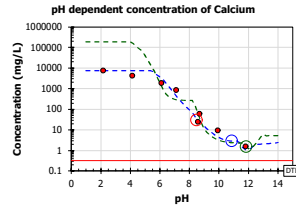
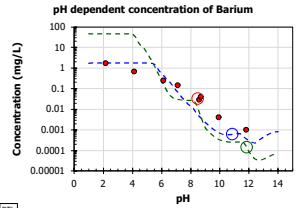
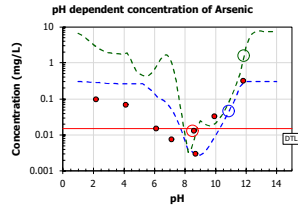
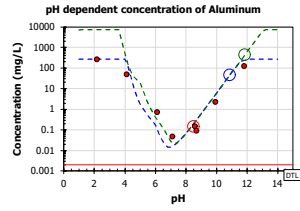
Name	End Member	Log(K)	Reaction
AFmc_ss	Cem07_C4Ach11_ss	-24.50	Cem07_C4Ach11_ss + 4 H+ -> 1 AFmc_ss + 2 Al[OH]4- + 1 CO3-2 + 4 Ca+2 + 9 H2O
	Cem07_C4Fch12_ss	-20.47	Cem07_C4Fch12_ss + 4 H+ -> 1 AFmc_ss + 1 CO3-2 + 4 Ca+2 + 2 Fe[OH]4- + 10 H2O

Minerals

Name	Log(K)	Reaction	Name	Log(K)	Reaction
AA_Fe[OH]3[am]	16.60	AA_Fe[OH]3[am] + 1 H2O -> 1 Fe[OH]4- + 1 H+	CuHPO4	26.00	CuHPO4 -> 1 Cu+2 + 1 H+ + 1 PO4-3
Antimocrandallite-ε	63.00	Antimocrandallite-exp + 8 H2O -> 3 Al[OH]4- + 1 Ca+2 + 3 H+ + 2 Sb[OH]6-	Exp_CaCO3_B	21.30	Exp_CaCO3_BaCO3 -> 1 Ba+2 + 2 CO3-2 + 1 Ca+2
Ba[ScR]O4[96%SO4]	9.790	Ba[ScR]O4[96%SO4] -> 1 Ba+2 + 0.04 CrO4-2 + 0.96 SO4-2	Exp_Ni2SiO4	5.498	Exp_Ni2SiO4 + 2 H+ -> 1 H2SiO4-2 + 2 Ni+2
BaSrSO4[50%Ba]	8.221	BaSrSO4[50%Ba] -> 0.5 Ba+2 + 1 SO4-2 + 0.5 Sr+2	Fe_Vanadate	19.18	Fe_Vanadate + 1 H2O -> 0.5 Fe[OH]4- + 1 VO2+ + 0.5 e-
Ca[OH]Sb[OH]6_exp	4.000	Ca[OH]Sb[OH]6_exp + 1 H+ -> 1 Ca+2 + 1 H2O + 1 Sb[OH]6-	Fe2[MoO4]3[Fluorite]	86.35	Fe2[MoO4]3[Fluorite] + 8 H2O -> 2 Fe[OH]4- + 8 H+ + 3 MoO4-2
CaCO3_Li2CO3	21.30	CaCO3_Li2CO3 -> 2 CO3-2 + 1 Ca+2 + 2 Li+	Fluorite	10.96	Fluorite -> 1 Ca+2 + 2 F-
CaCO3_MgCO3-exp	18.02	CaCO3_MgCO3-exp -> 2 CO3-2 + 1 Ca+2 + 1 Mg+2	Laumontite	118.0	Laumontite + 8 H2O -> 2 Al[OH]4- + 1 Ca+2 + 8 H+ + 4 H2SiO4-2
CaCO3_MnCO3-exp	20.78	CaCO3_MnCO3-exp -> 2 CO3-2 + 1 Ca+2 + 1 Mn+2	Manganite	-25.27	Manganite + 3 H+ + 1 e- -> 2 H2O + 1 Mn+2
CaCO3_SrCO3	19.85	CaCO3_SrCO3 -> 2 CO3-2 + 1 Ca+2 + 1 Sr+2	Ni[OH]2[s]	-10.80	Ni[OH]2[s] + 2 H+ -> 2 H2O + 1 Ni+2
CaSb[OH]6[s]2_exp	19.41	CaSb[OH]6[s]2_exp -> 1 Ca+2 + 2 Sb[OH]6-	NiHPO4	25.00	NiHPO4 -> 1 H+ + 1 Ni+2 + 1 PO4-3
CdHPO4	26.48	CdHPO4 -> 1 Cd+2 + 1 H+ + 1 PO4-3	PATCH_beta-1	28.93	PATCH_beta-TCP -> 3 Ca+2 + 2 PO4-3
Cem07_Al[OH]3[am]	13.76	Cem07_Al[OH]3[am] + 1 H2O -> 1 Al[OH]4- + 1 H+	Pb[OH]2[C]	-8.150	Pb[OH]2[C] + 2 H+ -> 2 H2O + 1 Pb+2
Cem07_Brucite	-16.83	Cem07_Brucite + 2 H+ -> 2 H2O + 1 Mg+2	PbMoO4[c]	15.80	PbMoO4[c] -> 1 MoO4-2 + 1 Pb+2
Cem07_Calcite	8.485	Cem07_Calcite -> 1 CO3-2 + 1 Ca+2	PbOH[Sb[OH]6]_exp1	12.00	PbOH[Sb[OH]6]_exp1 + 1 H+ -> 1 H2O + 1 Pb+2 + 1 Sb[OH]6-
Cem07_Gypsum	4.583	Cem07_Gypsum -> 1 Ca+2 + 2 H2O + 1 SO4-2	Sb[OH]3[s]	32.89	Sb[OH]3[s] + 3 H2O -> 3 H+ + 1 Sb[OH]6- + 2 e-
Co2SiO4	6.289	Co2SiO4 + 2 H+ -> 2 Co+2 + 1 H2SiO4-2	SiO2[a]	24.64	SiO2[a] + 2 H2O -> 2 H+ + 1 H2SiO4-2
CoHPO4[s]	24.48	CoHPO4[s] -> 1 Co+2 + 1 H+ + 1 PO4-3	Willemite	6.289	Willemite + 2 H+ -> 1 H2SiO4-2 + 2 Zn+2
Cr[OH]3[A]	68.13	Cr[OH]3[A] + 1 H2O -> 1 CrO4-2 + 5 H+ + 3 e-	ZnHPO4	24.48	ZnHPO4 -> 1 H+ + 1 PO4-3 + 1 Zn+2

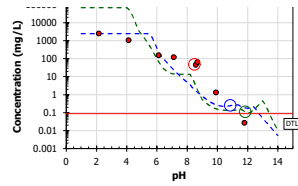
MARINE HARBOUR SEDIMENT IT

COMPARISON AND PARTITIONING

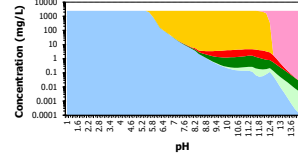


MARINE HARBOUR SEDIMENT IT

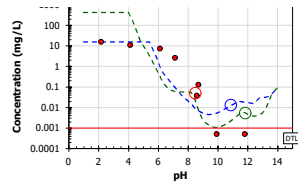
COMPARISON AND PARTITIONING



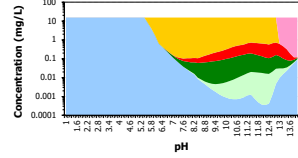
Mg Profile at L/S=10.8 L/kg



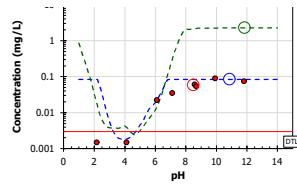
Total Dissolved DHA-bound SHA-bound
 FeOxide Clay CaCO₃/MgCO₃-exp
 Cem07_Bructite



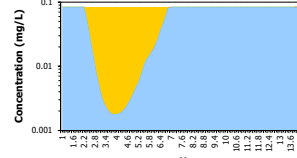
Mn Profile at L/S=10.8 L/kg



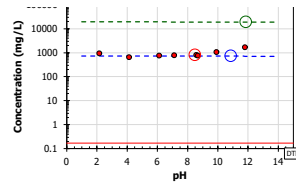
Total Dissolved DHA-bound SHA-bound
 FeOxide Clay CaCO₃/MgCO₃-exp
 Manganite



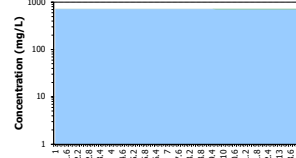
Mo Profile at L/S=10.8 L/kg



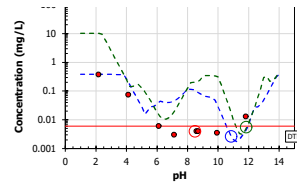
Total Dissolved DHA-bound SHA-bound
 FeOxide Clay PbMoO₄(c)



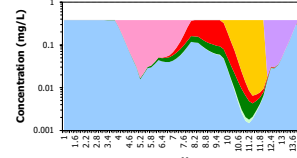
Na Profile at L/S=10.8 L/kg



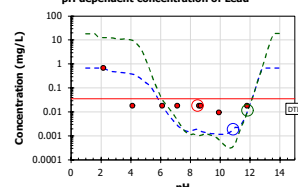
Total Dissolved DHA-bound SHA-bound Clay



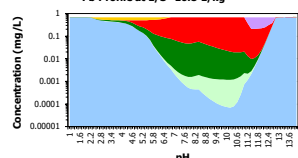
Ni Profile at L/S=10.8 L/kg



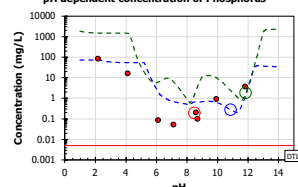
Total Dissolved DHA-bound SHA-bound FeOxide
 Clay Exp_Ni2SiO4 NiHPO4 Ni(OH)2(s)



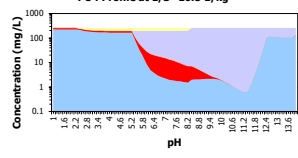
Pb Profile at L/S=10.8 L/kg



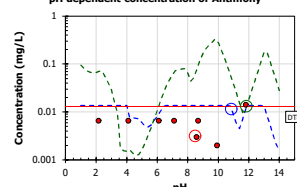
Total Dissolved DHA-bound SHA-bound
 FeOxide Clay PbMoO₄(c)
 PbOH(Sn(OH)₆)-exp-1 = Pb(OH)₂(c)



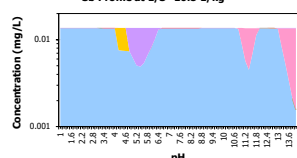
PO4 Profile at L/S=10.8 L/kg



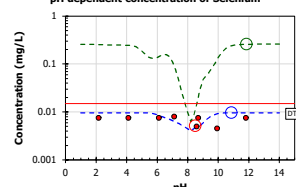
Total Dissolved DHA-bound SHA-bound
 FeOxide Clay CaHPO4
 Ca(OH)Sn(OH)₆-exp-1 = Pb(OH)₂(c)
 PATCH_beta-TCP ZnHPO4



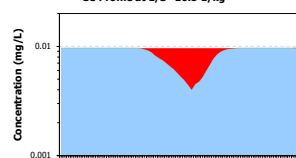
Sb Profile at L/S=10.8 L/kg



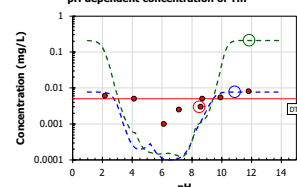
Total Dissolved DHA-bound SHA-bound
 FeOxide Clay Antimondalite-exp
 Ca(OH)Sn(OH)₆-exp-1 = Pb(OH)₂(c)



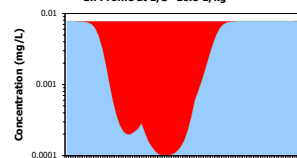
Se Profile at L/S=10.8 L/kg



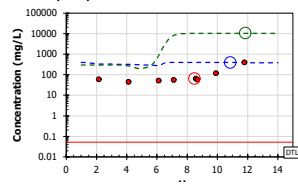
Total Dissolved DHA-bound SHA-bound FeOxide Clay



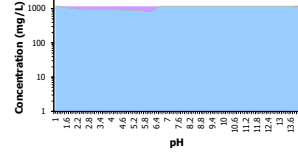
Sn Profile at L/S=10.8 L/kg



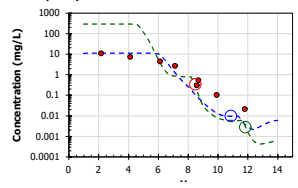
Total Dissolved SHA-bound FeOxide Clay



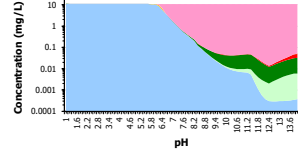
SO4 Profile at L/S=10.8 L/kg



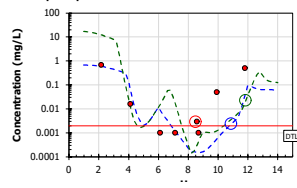
Total Dissolved DHA-bound SHA-bound
 FeOxide Clay BaS₂SO₄(50%Ba)
 Ba(SO₄)₂(96%SO₄) Cem07_Gypsum



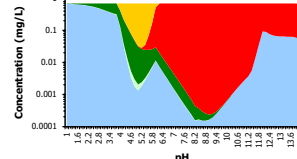
Sr Profile at L/S=10.8 L/kg



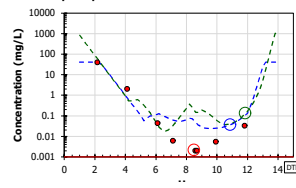
Total Dissolved DHA-bound SHA-bound
 FeOxide Clay BaS₂SO₄(50%Ba)
 CaCO₃_SrCO₃



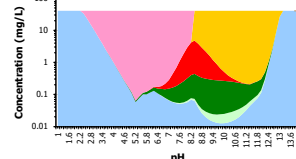
V Profile at L/S=10.8 L/kg



Total Dissolved DHA-bound SHA-bound
 FeOxide Clay Fe₂Vanadate



Zn Profile at L/S=10.8 L/kg



Total Dissolved DHA-bound SHA-bound FeOxide
 Clay Willemite ZnHPO4

Model Comparison: residuals - Concentration

Name Marine harbour sediment IT

Legend

Total Average Deviation Square root of the sum of the squared values of residuals divided by the number of values, over the entire X range.

User Average Deviation Square root of the sum of the squared values of residuals divided by the number of values, over the user defined X range.

Fractional Average Deviation Square root of the sum of the squared values of residuals divided by the number of values, over the fraction.

Note that the Total and User Average Deviation columns are averages as well.

Residual details, concentrations

Residuals as log(model/sample)										
Fraction	8	7	6	5	3	4	2	1	Total Avg	
pH	2.16	4.13	6.12	7.12	8.58	8.70	9.92	11.8	Deviation	
Al	0.01	0.54	-0.85	-0.46	0.20	0.55	0.35	0.27	0.17	
As	0.47	0.59	0.91	0.83	-0.62	-0.01	-0.59	-0.05	0.21	
Ba	0.01	0.40	0.25	-0.34	-0.81	-1.03	-0.74	-0.44	0.21	
Ca	0.00	0.24	0.24	-0.29	0.11	-0.38	-0.31	-0.16	0.09	
Cd	0.00	0.69	0.38	0.11	-0.51	-0.52	-0.36	0.04	0.14	
Cl	-	-	-	-	-	-	-	-	-	
Co	0.01	0.34	0.33	0.96	0.38	0.88	-0.71	-1.48	0.27	
CO32-	-	-	-	-	-	-	-	-	-	
Cr	-0.62	-0.34	0.85	0.84	0.98	1.00	1.20	0.98	0.31	
Cu	1.38	0.75	-0.15	-0.20	-0.18	-0.16	0.11	0.06	0.20	
F	-	-	-	-	-	-	-	-	-	
Fe	0.00	0.29	-1.27	-1.46	0.55	0.88	0.19	0.52	0.28	
B	0.12	0.22	0.26	0.33	0.32	0.42	-0.03	0.02	0.09	
Si	-0.71	-0.16	-0.14	-0.04	0.12	0.09	0.01	0.91	0.15	
Hg	-	-	-	-	-	-	-	-	-	
K	-0.36	-0.15	-0.17	-0.15	-0.01	-0.05	0.06	0.16	0.06	
Li	0.01	0.73	1.08	1.19	1.48	1.38	1.68	1.45	0.43	
Mg	0.01	0.39	0.19	-0.70	-1.43	-1.68	-0.68	0.83	0.32	
Mn	0.01	0.16	-0.90	-1.45	-0.68	-1.27	1.03	1.57	0.37	
Mo	1.75	0.10	0.01	0.38	0.15	0.20	-0.03	0.06	0.23	
Na	-0.10	0.05	-0.01	-0.03	-0.04	-0.02	-0.17	-0.37	0.05	
Ni	0.00	0.42	0.83	1.18	1.38	1.34	1.01	-0.49	0.34	
Pb	-0.01	1.32	0.00	-0.79	-1.03	-1.05	-0.90	-0.26	0.29	
PO4	-	-	-	-	-	-	-	-	-	
Sb	0.32	0.16	0.15	0.32	0.66	0.32	0.83	-0.09	0.15	
Se	0.11	0.11	0.05	-0.10	-0.03	-0.18	0.31	0.11	0.05	
Sn	0.10	-1.34	-0.97	-1.35	-0.42	-0.55	0.12	-0.01	0.28	
SO4	-	-	-	-	-	-	-	-	-	
Sr	0.01	0.17	0.32	-0.33	-0.52	-0.86	-0.87	-0.73	0.20	
Th	-	-	-	-	-	-	-	-	-	
U	-	-	-	-	-	-	-	-	-	
V	-0.06	0.56	0.94	0.08	-1.29	-0.82	-1.95	-1.10	0.37	
Zn	0.01	-0.44	0.45	1.02	1.39	1.31	0.65	0.47	0.30	
Avg Deviat	0.10	0.11	0.12	0.15	0.16	0.17	0.16	0.14	0.22	

Yellow = own pH All residuals within + 1 or - 1 are considered to represent a good fit.