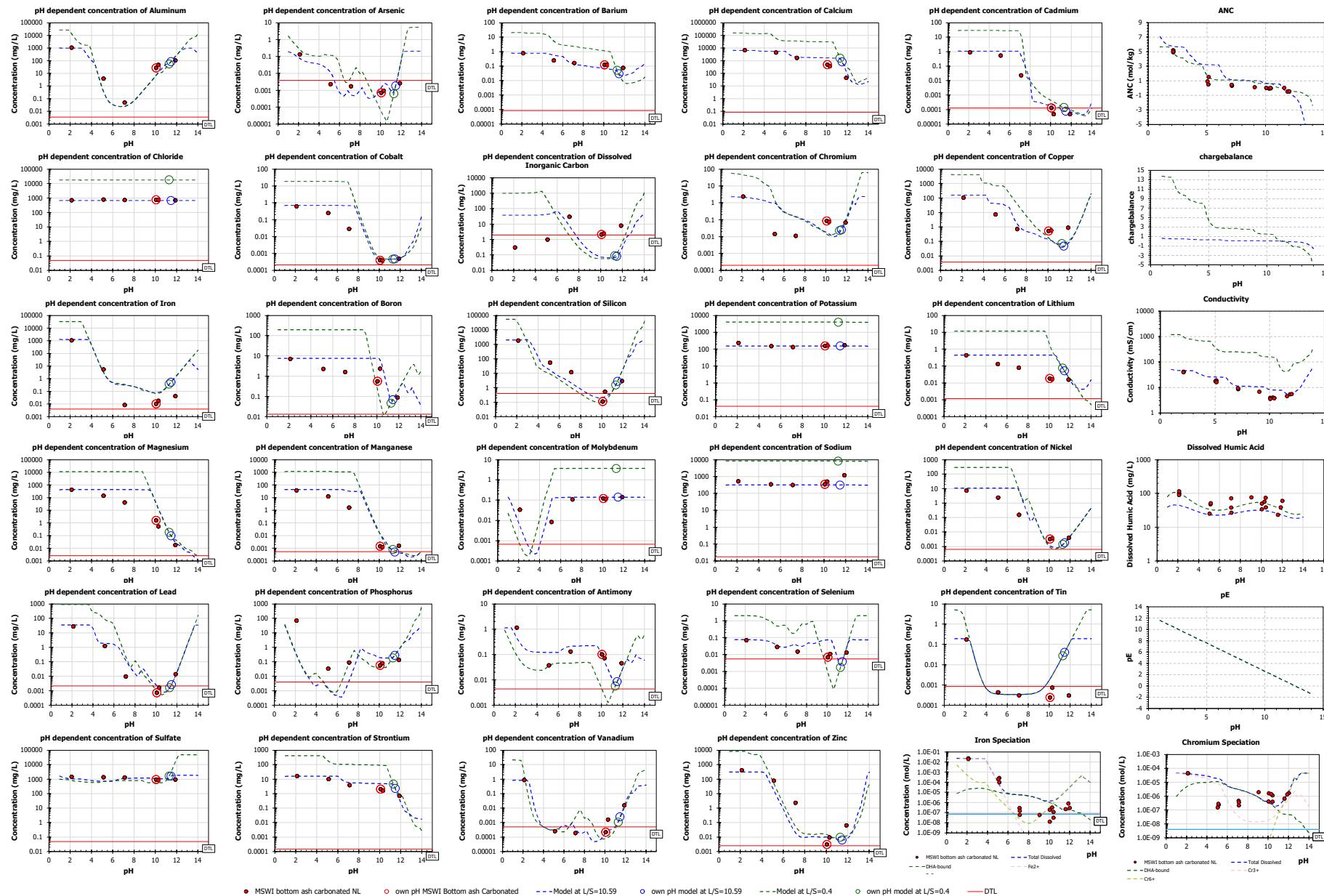


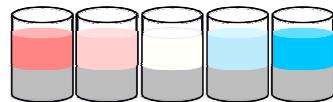
## MSWI BOTTOM ASH CARBONATED NL

## COMPARISON pH DEPENDENCE WITH MODEL



**Object Name** pH Dependent Leaching Test Model  
MSWI Bottom ash Carbonated NL

### pH Dependent Leaching Test Scenario



#### Lab Test                      Extra L/S Simulation

##### Lab Test

##### Model Parameters

Available Content								
Entity	Unit	Default	Entity	mg/kg	Entity	mg/kg	Entity	mg/kg
c0		-5.079	Al	1.083E+04	B	79.54	Po4	1948
c1		0.6359	As	2.203	Si	2.175E+04	Sb	12.02
c2		-0.2942	Ba	8.447	K	1620	Se	0.7975
c3		0.05090	Ca	7.186E+04	Li	4.664	Sn	2.036
c4		-0.003732	Cd	11.76	Mg	4501	So4	1.949E+04
c5		9.770E-05	Cl	7190	Mn	464.3	Sr	167.4
Clay	mg/kg	3000	Co	7.496	Mo	1.474	V	8.971
Hydrous Ferric Oxid	mg/kg	500.0	CO32-	6.294E+04	Na	3343	Zn	3313
L/S	L/kg	10.59	Cr	24.99	Ni	114.7		
pE		2.170	Cu	1688	NO3	6.200E-08		
pH		10.43	F	1.900E-08	Pb	366.1		
Solid Humic Acid	mg/kg	800.0	Fe	1.422E+04				
Simulated Low L/S	L/kg	0.4000						

##### Solid Solutions

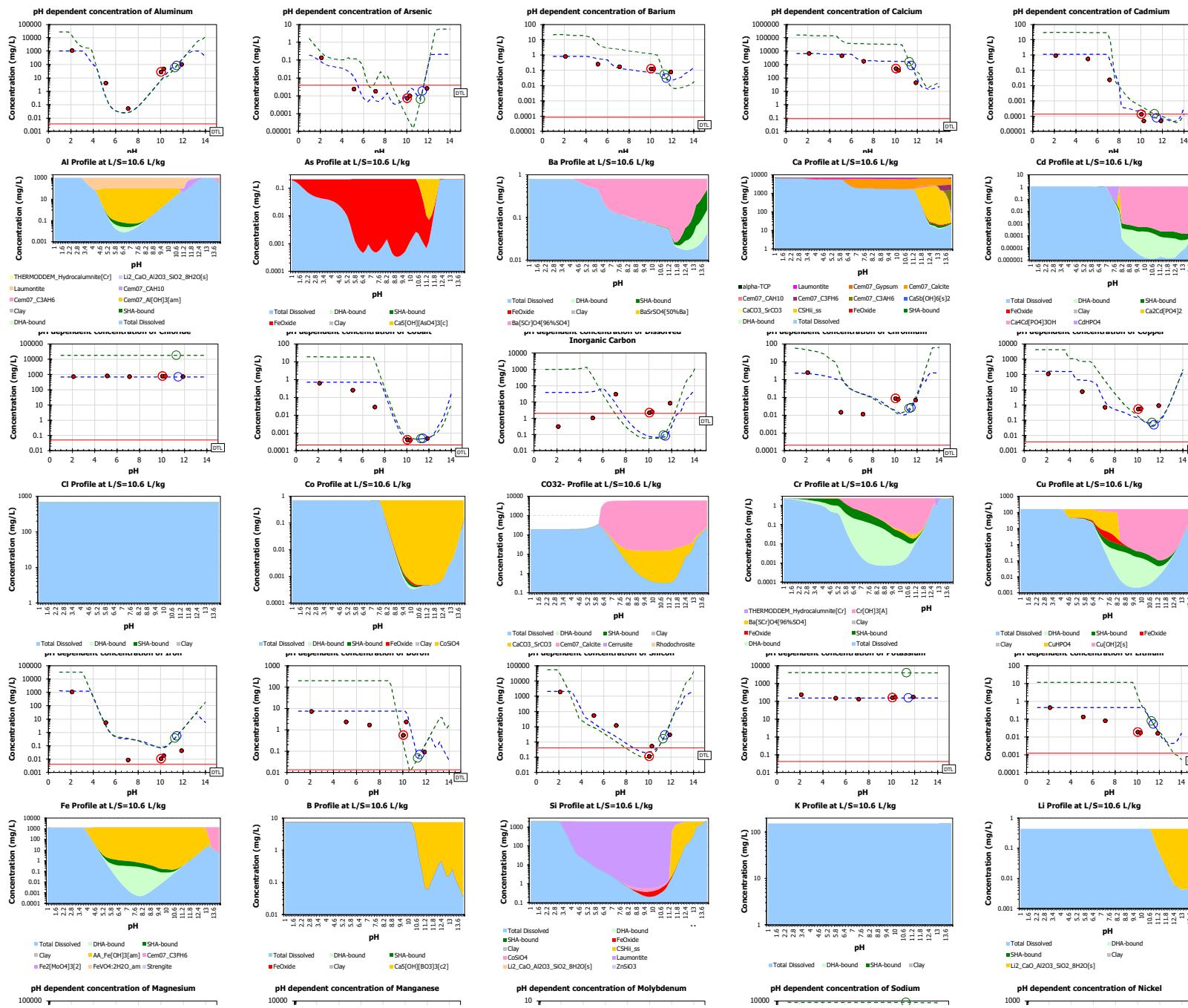
Name	End Member	Log(K)	Reaction
CSHii_ss	Cem07_Jenn_ss	-7.799	Cem07_Jenn_ss + 1.33333 H+ -> 1 CSHii_ss + 1.66667 Ca+2 + 1.76667 H2O + 1 H2SiO4-2
	Cem07_Tob_II_ss	10.36	Cem07_Tob_II_ss -> 1 CSHii_ss + 0.83333 Ca+2 + 0.33333 H+ + 0.16667 H2O + 1 H2SiO4-2

##### 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+	Cu[OH]2[s]	-8.640	Cu[OH]2[s] + 2 H+ -> 1 Cu+2 + 2 H2O
alpha-TCP	25.50	alpha-TCP -> 3 Ca+2 + 2 PO4-3	CuHPO4	26.00	CuHPO4 -> 1 Cu+2 + 1 H+ + 1 PO4-3
Ba[Scr]O4[96%SO4]	9.790	Ba[Scr]O4[96%SO4] -> 1 Ba+2 + 0.04 CrO4-2 + 0.96 SO4-2	Exp2_Ca[OH]Sb[OH]6	1.000	Exp2_Ca[OH]Sb[OH]6 + 1 H+ -> 1 Ca+2 + 1 H2O + 1 Sb[OH]6-
BaSrSO4[50%Ba]	8.221	BaSrSO4[50%Ba] -> 0.5 Ba+2 + 1 SO4-2 + 0.5 Sr+2	Exp2_Ca2[OH]2[SeO4]	-8.000	Exp2_Ca2[OH]2[SeO4] + 2 H+ -> 2 Ca+2 + 2 H2O + 1 SeO4-2
Ca2Cd[PO4]2	32.95	Ca2Cd[PO4]2 -> 2 Ca+2 + 1 Cd+2 + 2 PO4-3	Fe2[MoO4]3[2]	86.35	Fe2[MoO4]3[2] + 8 H2O -> 2 Fe[OH]4- + 8 H+ + 3 MoO4-2
Ca4Cd[PO4]3OH	39.23	Ca4Cd[PO4]3OH + 1 H+ -> 4 Ca+2 + 1 Cd+2 + 1 H2O + 3 PO4-3	FeVO4:2H2O_am	23.48	FeVO4:2H2O_am + 2 H2O -> 1 Fe[OH]4- + 1 VO2+
Ca5[OH][AsO4]3[c]	26.13	Ca5[OH][AsO4]3[c] + 1 H+ -> 3 AsO4-3 + 5 Ca+2 + 1 H2O	Laumontite	118.0	Laumontite + 8 H2O -> 2 Al[OH]4- + 1 Ca+2 + 8 H+ + 4 H2SiO4-2
Ca5[OH][BO3]3[c2]	-53.00	Ca5[OH][BO3]3[c2] + 7 H+ -> 5 Ca+2 + 3 H2BO3- + 1 H2O	Li2_Ca_O_Al2O3_SiO2_8H2O[s]	22.69	Li2_Ca_O_Al2O3_SiO2_8H2O[s] -> 2 Al[OH]4- + 1 Ca+2 + 3 H2O + 1 H2SiO4-2 + 2 Li+
CaCO3_SrCO3	19.85	CaCO3_SrCO3 -> 2 CO3-2 + 1 Ca+2 + 1 Sr+2	Manganite	-25.27	Manganite + 3 H+ + 1 e- -> 2 H2O + 1 Mn+2
CaSb[OH]6[s]2	19.41	CaSb[OH]6[s]2 -> 1 Ca+2 + 2 Sb[OH]6-	Ni[OH]2[s]	-10.80	Ni[OH]2[s] + 2 H+ -> 2 H2O + 1 Ni+2
CdHPO4	26.48	CdHPO4 -> 1 Cd+2 + 1 H+ + 1 PO4-3	NiHPO4	25.00	NiHPO4 -> 1 H+ + 1 Ni+2 + 1 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	Pb2V2O7	0.9500	Pb2V2O7 + 3 H+ -> 1.5 H2O + 1 Pb+2 + 1 VO2+
Cem07_C3AH6	-35.14	Cem07_C3AH6 + 4 H+ -> 2 Al[OH]4- + 3 Ca+2 + 4 H2O	Pb3[VO4]2	-3.070	Pb3[VO4]2 + 4 H+ -> 2 H2O + 1.5 Pb+2 + 1 VO2+
Cem07_C3FH6	-30.82	Cem07_C3FH6 + 4 H+ -> 3 Ca+2 + 2 Fe[OH]4- + 4 H2O	PbHPO4	28.00	PbHPO4 -> 1 H+ + 1 PO4-3 + 1 Pb+2
Cem07_CAH10	7.505	Cem07_CAH10 -> 2 Al[OH]4- + 1 Ca+2 + 6 H2O	Rhodochrosite	10.41	Rhodochrosite -> 1 CO3-2 + 1 Mn+2
Cem07_Calcite	8.485	Cem07_Calcite -> 1 CO3-2 + 1 Ca+2	Sn[OH]2[s]	1.447	Sn[OH]2[s] + 2 H+ -> 2 H2O + 1 Sn+2
Cem07_Gypsum	4.583	Cem07_Gypsum -> 1 Ca+2 + 2 H2O + 1 SO4-2	Strengite	48.00	Strengite + 2 H2O -> 1 Fe[OH]4- + 4 H+ + 1 PO4-3
Cerrusite	13.13	Cerrusite -> 1 CO3-2 + 1 Pb+2	THERMODDEM_Hydrocalumnite[Cr]	-25.34	THERMODDEM_Hydrocalumnite[Cr] + 4 H+ -> 2 Al[OH]4- + 4 Ca+2 + 1 CrO4-2 + 13 H2O
CoSiO4	6.289	CoSiO4 + 2 H+ -> 2 Co+2 + 1 H2SiO4-2	ZnSiO3	18.69	ZnSiO3 + 1 H2O -> 1 H2SiO4-2 + 1 Zn+2
Cr[OH]3[A]	68.13	Cr[OH]3[A] + 1 H2O -> 1 CrO4-2 + 5 H+ + 3 e-			

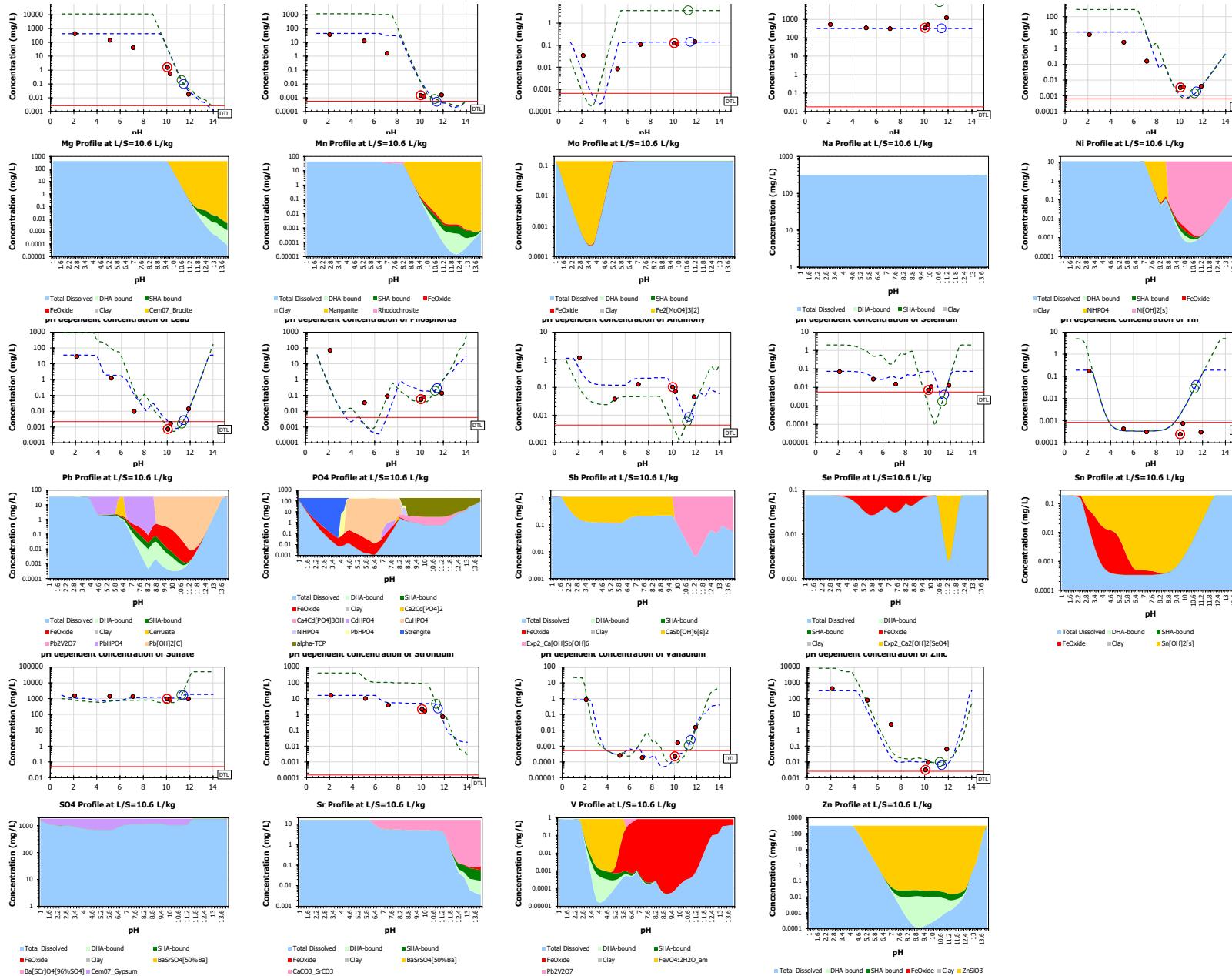
**MSWI BOTTOM ASH CARBONATED NL**

## **COMPARISON AND PARTITIONING**



## MSWI BOTTOM ASH CARBONATED NL

## COMPARISON AND PARTITIONING



## Model Comparison: residuals - Concentration

Name MSWI Bottom ash Carbonated NL

### 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 Devi:** 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	6	5	4	3	2	1	Total Avg
	pH	2.13	5.13	7.13	10.1	10.3	11.9	Deviation
Al	-0.03	-1.03	-0.26	-0.58	-0.58	0.33	0.33	0.23
As	-0.25	0.69	-0.50	0.32	0.37	1.01	1.01	0.24
Ba	0.01	0.37	-0.09	-0.24	-0.26	-0.54	-0.54	0.12
Ca	-0.01	0.11	0.06	0.54	0.66	0.51	0.51	0.17
Cd	0.09	0.32	1.33	0.23	0.62	0.16	0.16	0.25
Cl	-0.02	-0.05	-0.02	-0.05	-0.04	-0.01	-0.01	0.01
Co	0.07	0.46	1.39	0.02	0.01	0.03	0.03	0.24
CO32-	-	-	-	-	-	-	-	-
Cr	-0.04	1.83	1.14	-0.67	-0.71	0.04	0.04	0.40
Cu	0.18	0.77	0.36	-0.57	-0.68	-1.05	-1.05	0.27
Fe	0.07	-0.23	1.58	0.88	0.67	1.51	1.51	0.41
B	0.03	0.52	0.66	1.13	0.43	0.31	0.31	0.25
Si	0.05	-0.52	-0.74	0.29	-0.32	0.68	0.68	0.20
K	-0.19	0.01	0.06	-0.01	-0.04	-0.05	-0.05	0.03
Li	0.01	0.53	0.74	1.38	1.42	0.11	0.11	0.36
Mg	0.00	0.49	1.01	1.38	1.40	0.24	0.24	0.38
Mn	0.08	0.55	1.30	0.99	0.78	-0.55	-0.55	0.33
Mo	-0.82	1.04	0.11	0.06	0.09	0.00	0.00	0.22
Na	-0.22	-0.03	0.01	-0.03	-0.20	-0.57	-0.57	0.11
Ni	0.17	0.65	1.63	-0.43	-0.59	0.01	0.01	0.32
Pb	0.10	0.18	1.07	0.50	-0.04	-0.06	-0.06	0.20
Sb	-0.41	0.50	0.21	-0.07	-0.15	-0.31	-0.31	0.13
Se	0.03	0.11	0.36	1.03	0.83	0.30	0.30	0.23
Sn	0.01	-0.08	0.04	0.91	0.62	2.54	2.54	0.46
SO4	-0.14	-0.28	-0.06	0.06	0.10	0.30	0.30	0.08
Sr	-0.01	0.20	0.17	0.38	0.49	-0.23	-0.23	0.12
V	-0.01	0.06	0.28	-0.07	-0.77	-0.01	-0.01	0.14
Zn	-0.12	-0.69	-2.06	0.52	0.02	-0.91	-0.91	0.40
Avg Deviation	0.04	0.12	0.17	0.12	0.12	0.14	0.14	0.23

Yellow = own pH

All residuals within + 1 or - 1 are considered to represent a good fit.