

Predicted Maximum Permitted Concentration of Certain Metals in Foods - Chicken - The IWMF at the TTAL site

ASR	Assessment Height (mAG)	Antimony	Arsenic	Cadmium	Chromium	Lead	Mercuric chloride	Methyl mercury	Mercury
Maximum Permitted Concentration (ppm)		1	1.4	0.2	1	6			0.5
TT1	0-19.2	0.00E+00	0.00E+00	1.24E-10	0.00E+00	0.00E+00	4.01E-06	1.07E-08	4.02E-06
TT2	0-19.2	0.00E+00	0.00E+00	8.87E-11	0.00E+00	0.00E+00	1.45E-05	3.89E-08	1.46E-05
TT3	0-19.2	0.00E+00	0.00E+00	1.40E-10	0.00E+00	0.00E+00	2.24E-05	6.00E-08	2.25E-05
TT4	0-19.2	0.00E+00	0.00E+00	5.58E-10	0.00E+00	0.00E+00	1.26E-04	3.38E-07	1.27E-04
TT5	0-19.2	0.00E+00	0.00E+00	2.42E-10	0.00E+00	0.00E+00	6.82E-06	1.83E-08	6.84E-06
TT6	0-19.2	0.00E+00	0.00E+00	2.11E-10	0.00E+00	0.00E+00	2.18E-05	5.85E-08	2.19E-05
TM1	0-19.2	0.00E+00	0.00E+00	1.11E-11	0.00E+00	0.00E+00	6.80E-07	1.82E-09	6.81E-07
	19.2-41	0.00E+00	0.00E+00	1.11E-11	0.00E+00	0.00E+00	6.80E-07	1.82E-09	6.81E-07
	41-78.6	0.00E+00	0.00E+00	1.11E-11	0.00E+00	0.00E+00	6.80E-07	1.82E-09	6.81E-07
	78.6-120	0.00E+00	0.00E+00	1.11E-11	0.00E+00	0.00E+00	6.80E-07	1.82E-09	6.81E-07
TM2	0-19.2	0.00E+00	0.00E+00	2.67E-11	0.00E+00	0.00E+00	5.27E-07	1.41E-09	5.28E-07
	19.2-41	0.00E+00	0.00E+00	2.67E-11	0.00E+00	0.00E+00	5.27E-07	1.41E-09	5.28E-07
	41-78.6	0.00E+00	0.00E+00	2.67E-11	0.00E+00	0.00E+00	5.27E-07	1.41E-09	5.28E-07
	78.6-120	0.00E+00	0.00E+00	2.67E-11	0.00E+00	0.00E+00	5.27E-07	1.41E-09	5.28E-07
TM3	0-19.2	0.00E+00	0.00E+00	1.97E-11	0.00E+00	0.00E+00	7.99E-07	2.14E-09	8.01E-07
	19.2-41	0.00E+00	0.00E+00	1.97E-11	0.00E+00	0.00E+00	7.99E-07	2.14E-09	8.01E-07
	41-78.6	0.00E+00	0.00E+00	1.97E-11	0.00E+00	0.00E+00	7.99E-07	2.14E-09	8.01E-07
	78.6-120	0.00E+00	0.00E+00	1.97E-11	0.00E+00	0.00E+00	7.99E-07	2.14E-09	8.01E-07
TM4	0-19.2	0.00E+00	0.00E+00	2.82E-11	0.00E+00	0.00E+00	8.50E-07	2.28E-09	8.52E-07
	19.2-41	0.00E+00	0.00E+00	2.82E-11	0.00E+00	0.00E+00	8.50E-07	2.28E-09	8.52E-07
	41-78.6	0.00E+00	0.00E+00	2.82E-11	0.00E+00	0.00E+00	8.50E-07	2.28E-09	8.52E-07
	78.6-120	0.00E+00	0.00E+00	2.82E-11	0.00E+00	0.00E+00	8.50E-07	2.28E-09	8.52E-07
TM5	0-19.2	0.00E+00	0.00E+00	2.04E-11	0.00E+00	0.00E+00	9.51E-07	2.55E-09	9.54E-07
	19.2-41	0.00E+00	0.00E+00	2.04E-11	0.00E+00	0.00E+00	9.51E-07	2.55E-09	9.54E-07
	41-78.6	0.00E+00	0.00E+00	2.04E-11	0.00E+00	0.00E+00	9.51E-07	2.55E-09	9.54E-07
	78.6-120	0.00E+00	0.00E+00	2.04E-11	0.00E+00	0.00E+00	9.51E-07	2.55E-09	9.54E-07
TM6	0-19.2	0.00E+00	0.00E+00	2.16E-11	0.00E+00	0.00E+00	1.19E-06	3.19E-09	1.19E-06
	19.2-41	0.00E+00	0.00E+00	2.16E-11	0.00E+00	0.00E+00	1.19E-06	3.19E-09	1.19E-06
	41-78.6	0.00E+00	0.00E+00	2.16E-11	0.00E+00	0.00E+00	1.19E-06	3.19E-09	1.19E-06
	78.6-120	0.00E+00	0.00E+00	2.16E-11	0.00E+00	0.00E+00	1.19E-06	3.19E-09	1.19E-06
SLW1	0-19.2	0.00E+00	0.00E+00	3.76E-10	0.00E+00	0.00E+00	1.54E-06	4.12E-09	1.54E-06
SLW2	0-19.2	0.00E+00	0.00E+00	4.01E-10	0.00E+00	0.00E+00	1.71E-06	4.58E-09	1.71E-06
SLW3	0-19.2	0.00E+00	0.00E+00	1.42E-10	0.00E+00	0.00E+00	3.60E-07	9.64E-10	3.61E-07
TC1	0-19.2	0.00E+00	0.00E+00	9.98E-11	0.00E+00	0.00E+00	2.04E-07	5.46E-10	2.04E-07
	19.2-41	0.00E+00	0.00E+00	9.98E-11	0.00E+00	0.00E+00	2.04E-07	5.46E-10	2.04E-07
	41-78.6	0.00E+00	0.00E+00	9.98E-11	0.00E+00	0.00E+00	2.04E-07	5.46E-10	2.04E-07
	78.6-120	0.00E+00	0.00E+00	9.98E-11	0.00E+00	0.00E+00	2.04E-07	5.46E-10	2.04E-07
TC2	0-19.2	0.00E+00	0.00E+00	9.13E-11	0.00E+00	0.00E+00	1.36E-07	3.64E-10	1.36E-07
	19.2-41	0.00E+00	0.00E+00	9.13E-11	0.00E+00	0.00E+00	1.36E-07	3.64E-10	1.36E-07
	41-78.6	0.00E+00	0.00E+00	9.13E-11	0.00E+00	0.00E+00	1.36E-07	3.64E-10	1.36E-07
	78.6-120	0.00E+00	0.00E+00	9.13E-11	0.00E+00	0.00E+00	1.36E-07	3.64E-10	1.36E-07
TC3	0-19.2	0.00E+00	0.00E+00	8.29E-11	0.00E+00	0.00E+00	1.19E-07	3.19E-10	1.19E-07
	19.2-41	0.00E+00	0.00E+00	8.29E-11	0.00E+00	0.00E+00	1.19E-07	3.19E-10	1.19E-07
	41-78.6	0.00E+00	0.00E+00	8.29E-11	0.00E+00	0.00E+00	1.19E-07	3.19E-10	1.19E-07
	78.6-120	0.00E+00	0.00E+00	8.29E-11	0.00E+00	0.00E+00	1.19E-07	3.19E-10	1.19E-07
TC4	0-19.2	0.00E+00	0.00E+00	9.13E-11	0.00E+00	0.00E+00	1.19E-07	3.19E-10	1.19E-07
	19.2-41	0.00E+00	0.00E+00	9.13E-11	0.00E+00	0.00E+00	1.19E-07	3.19E-10	1.19E-07
	41-78.6	0.00E+00	0.00E+00	9.13E-11	0.00E+00	0.00E+00	1.19E-07	3.19E-10	1.19E-07
	78.6-120	0.00E+00	0.00E+00	9.13E-11	0.00E+00	0.00E+00	1.19E-07	3.19E-10	1.19E-07
TC5	0-19.2	0.00E+00	0.00E+00	9.97E-11	0.00E+00	0.00E+00	1.36E-07	3.64E-10	1.36E-07
	19.2-41	0.00E+00	0.00E+00	9.97E-11	0.00E+00	0.00E+00	1.36E-07	3.64E-10	1.36E-07
	41-78.6	0.00E+00	0.00E+00	9.97E-11	0.00E+00	0.00E+00	1.36E-07	3.64E-10	1.36E-07
	78.6-120	0.00E+00	0.00E+00	9.97E-11	0.00E+00	0.00E+00	1.36E-07	3.64E-10	1.36E-07
TC6	0-19.2	0.00E+00	0.00E+00	9.97E-11	0.00E+00	0.00E+00	1.36E-07	3.64E-10	1.36E-07
	19.2-41	0.00E+00	0.00E+00	9.97E-11	0.00E+00	0.00E+00	1.36E-07	3.64E-10	1.36E-07
	41-78.6	0.00E+00	0.00E+00	9.97E-11	0.00E+00	0.00E+00	1.36E-07	3.64E-10	1.36E-07
	78.6-120	0.00E+00	0.00E+00	9.97E-11	0.00E+00	0.00E+00	1.36E-07	3.64E-10	1.36E-07
AP1	0-19.2	0.00E+00	0.00E+00	1.08E-10	0.00E+00	0.00E+00	2.04E-07	5.46E-10	2.04E-07
AP2	0-19.2	0.00E+00	0.00E+00	1.08E-10	0.00E+00	0.00E+00	2.55E-07	6.83E-10	2.56E-07
AP3	0-19.2	0.00E+00	0.00E+00	1.59E-10	0.00E+00	0.00E+00	5.98E-07	1.60E-09	5.99E-07
AP4	0-19.2	0.00E+00	0.00E+00	7.52E-11	0.00E+00	0.00E+00	2.21E-07	5.92E-10	2.21E-07
	19.2-41	0.00E+00	0.00E+00	7.52E-11	0.00E+00	0.00E+00	2.21E-07	5.92E-10	2.21E-07
	41-78.6	0.00E+00	0.00E+00	7.52E-11	0.00E+00	0.00E+00	2.21E-07	5.92E-10	2.21E-07
	78.6-120	0.00E+00	0.00E+00	7.52E-11	0.00E+00	0.00E+00	2.21E-07	5.92E-10	2.21E-07
AP5	0-19.2	0.00E+00	0.00E+00	7.50E-11	0.00E+00	0.00E+00	1.87E-07	5.01E-10	1.87E-07
AP6	0-19.2	0.00E+00	0.00E+00	7.49E-11	0.00E+00	0.00E+00	1.87E-07	5.01E-10	1.87E-07
AP7	0-19.2	0.00E+00	0.00E+00	8.34E-11	0.00E+00	0.00E+00	2.55E-07	6.83E-10	2.56E-07
	19.2-41	0.00E+00	0.00E+00	8.34E-11	0.00E+00	0.00E+00	2.55E-07	6.83E-10	2.56E-07
	41-78.6	0.00E+00	0.00E+00	8.34E-11	0.00E+00	0.00E+00	2.55E-07	6.83E-10	2.56E-07
AP8	0-19.2	0.00E+00	0.00E+00	7.50E-11	0.00E+00	0.00E+00	1.87E-07	5.01E-10	1.87E-07
KT1	0-19.2	0.00E+00	0.00E+00	4.14E-11	0.00E+00	0.00E+00	3.40E-08	9.10E-11	3.41E-08
	19.2-41	0.00E+00	0.00E+00	4.14E-11	0.00E+00	0.00E+00	3.40E-08	9.10E-11	3.41E-08
	41-78.6	0.00E+00	0.00E+00	4.14E-11	0.00E+00	0.00E+00	3.40E-08	9.10E-11	3.41E-08
	78.6-120	0.00E+00	0.00E+00	4.14E-11	0.00E+00	0.00E+00	3.40E-08	9.10E-11	3.41E-08
KT2	0-19.2	0.00E+00	0.00E+00	3.32E-11	0.00E+00	0.00E+00	3.40E-08	9.10E-11	3.41E-08
	19.2-41	0.00E+00	0.00E+00	3.32E-11	0.00E+00	0.00E+00	3.40E-08	9.10E-11	3.41E-08
	41-78.6	0.00E+00	0.00E+00	3.32E-11	0.00E+00	0.00E+00	3.40E-08	9.10E-11	3.41E-08
	78.6-120	0.00E+00	0.00E+00	3.32E-11	0.00E+00	0.00E+00	3.40E-08	9.10E-11	3.41E-08
KT3	0-19.2	0.00E+00	0.00E+00	3.32E-11	0.00E+00	0.00E+00	3.40E-08	9.10E-11	3.41E-08
KT4	19.2-41	0.00E+00	0.00E+00	3.32E-11	0.00E+00	0.00E+00	3.40E-08	9.10E-11	3.41E-08
KT5	0-19.2	0.00E+00	0.00E+00	3.32E-11	0.00E+00	0.00E+00	3.40E-08	9.10E-11	3.41E-08
	19.2-41	0.00E+00	0.00E+00	1.66E-11	0.00E+00	0.00E+00	3.40E-08	9.10E-11	3.41E-08
	41-78.6	0.00E+00	0.00E+00	1.66E-11	0.00E+00	0.00E+00	3.40E-08	9.10E-11	3.41E-08
	78.6-120	0.00E+00	0.00E+00	1.66E-11	0.00E+00	0.00E+00	3.40E-08	9.10E-11	3.41E-08
	120-175	0.00E+00	0.00E+00	1.66E-11	0.00E+00	0.00E+00	3.40E-08	9.10E-11	3.41E-08
KT6	0-19.2	0.00E+00	0.00E+00	2.49E-11	0.00E+00	0.00E+00	3.40E-08	9.10E-11	3.41E-08
KT7	19.2-41	0.00E+00	0.00E+00	2.49E-11	0.00E+00	0.00E+00	3.40E-08	9.10E-11	3.41E-08
KT8	0-19.2	0.00E+00	0.00E+00	1.65E-11	0.00E+00	0.00E+00	1.70E-08	4.55E-11	1.70E-08
	19.2-41	0.00E+00	0.00E+00	1.65E-11	0.00E+00	0.00E+00	1.70E-08	4.55E-11	1.70E-08
	41-78.6	0.00E+00	0.00E+00	1.65E-11	0.00E+00	0.00E+00	1.70E-08	4.55E-11	1.70E-08
	78.6-120	0.00E+00	0.00E+00	1.65E-11	0.00E+00	0.00E+00	1.70E-08	4.55E-11	1.70E-08

Predicted Maximum Permitted Concentration of Certain Metals in Foods - Pork - The IWMF at the TTAL site

ASR	Assessment Height (mAG)	Antimony	Arsenic	Cadmium	Chromium	Lead	Mercuric chloride	Methyl mercury	Mercury
Maximum Permitted Concentration (ppm)		1	1.4	0.2	1	6			0.5
TT1	0-19.2	0.00E+00	0.00E+00	1.20E-08	0.00E+00	0.00E+00	9.56E-08	2.57E-10	9.59E-08
TT2	0-19.2	0.00E+00	0.00E+00	2.77E-09	0.00E+00	0.00E+00	3.46E-07	9.26E-10	3.47E-07
TT3	0-19.2	0.00E+00	0.00E+00	4.61E-09	0.00E+00	0.00E+00	5.34E-07	1.43E-09	5.35E-07
TT4	0-19.2	0.00E+00	0.00E+00	3.17E-11	0.00E+00	0.00E+00	3.01E-06	8.04E-09	3.01E-06
TT5	0-19.2	0.00E+00	0.00E+00	2.40E-08	0.00E+00	0.00E+00	1.63E-07	4.39E-10	1.63E-07
TT6	0-19.2	0.00E+00	0.00E+00	1.29E-08	0.00E+00	0.00E+00	5.20E-07	1.39E-09	5.21E-07
TM1	0-19.2	0.00E+00	0.00E+00	9.22E-10	0.00E+00	0.00E+00	1.63E-08	4.71E-11	1.63E-08
	19.2-41	0.00E+00	0.00E+00	9.22E-10	0.00E+00	0.00E+00	1.63E-08	4.72E-11	1.63E-08
	41-78.6	0.00E+00	0.00E+00	9.22E-10	0.00E+00	0.00E+00	1.63E-08	4.75E-11	1.63E-08
	78.6-120	0.00E+00	0.00E+00	9.22E-10	0.00E+00	0.00E+00	1.63E-08	4.80E-11	1.63E-08
TM2	0-19.2	0.00E+00	0.00E+00	2.77E-09	0.00E+00	0.00E+00	1.26E-08	3.73E-11	1.27E-08
	19.2-41	0.00E+00	0.00E+00	2.77E-09	0.00E+00	0.00E+00	1.26E-08	3.75E-11	1.27E-08
	41-78.6	0.00E+00	0.00E+00	2.77E-09	0.00E+00	0.00E+00	1.26E-08	3.77E-11	1.27E-08
	78.6-120	0.00E+00	0.00E+00	2.77E-09	0.00E+00	0.00E+00	1.27E-08	3.83E-11	1.27E-08
TM3	0-19.2	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	1.91E-08	5.47E-11	1.92E-08
	19.2-41	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	1.91E-08	5.48E-11	1.92E-08
	41-78.6	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	1.91E-08	5.51E-11	1.92E-08
	78.6-120	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	1.91E-08	5.56E-11	1.92E-08
TM4	0-19.2	0.00E+00	0.00E+00	2.77E-09	0.00E+00	0.00E+00	2.03E-08	5.79E-11	2.04E-08
	19.2-41	0.00E+00	0.00E+00	2.77E-09	0.00E+00	0.00E+00	2.03E-08	5.80E-11	2.04E-08
	41-78.6	0.00E+00	0.00E+00	2.77E-09	0.00E+00	0.00E+00	2.03E-08	5.83E-11	2.04E-08
	78.6-120	0.00E+00	0.00E+00	2.77E-09	0.00E+00	0.00E+00	2.03E-08	5.88E-11	2.04E-08
TM5	0-19.2	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	2.27E-08	6.39E-11	2.28E-08
	19.2-41	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	2.27E-08	6.40E-11	2.28E-08
	41-78.6	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	2.28E-08	6.44E-11	2.28E-08
	78.6-120	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	2.28E-08	6.55E-11	2.28E-08
TM6	0-19.2	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	2.84E-08	7.98E-11	2.85E-08
	19.2-41	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	2.84E-08	8.00E-11	2.85E-08
	41-78.6	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	2.84E-08	8.04E-11	2.85E-08
	78.6-120	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	2.85E-08	8.14E-11	2.85E-08
SLW1	0-19.2	0.00E+00	0.00E+00	4.15E-08	0.00E+00	0.00E+00	3.69E-08	1.09E-10	3.70E-08
SLW2	0-19.2	0.00E+00	0.00E+00	4.42E-08	0.00E+00	0.00E+00	4.09E-08	1.20E-10	4.11E-08
SLW3	0-19.2	0.00E+00	0.00E+00	1.57E-08	0.00E+00	0.00E+00	8.80E-09	3.25E-11	8.83E-09
TC1	0-19.2	0.00E+00	0.00E+00	1.11E-08	0.00E+00	0.00E+00	5.01E-09	1.96E-11	5.03E-09
	19.2-41	0.00E+00	0.00E+00	1.11E-08	0.00E+00	0.00E+00	5.01E-09	1.96E-11	5.03E-09
	41-78.6	0.00E+00	0.00E+00	1.11E-08	0.00E+00	0.00E+00	5.02E-09	1.97E-11	5.04E-09
	78.6-120	0.00E+00	0.00E+00	1.11E-08	0.00E+00	0.00E+00	5.02E-09	1.97E-11	5.04E-09
TC2	0-19.2	0.00E+00	0.00E+00	1.01E-08	0.00E+00	0.00E+00	3.39E-09	1.53E-11	3.41E-09
	19.2-41	0.00E+00	0.00E+00	1.01E-08	0.00E+00	0.00E+00	3.39E-09	1.53E-11	3.41E-09
	41-78.6	0.00E+00	0.00E+00	1.01E-08	0.00E+00	0.00E+00	3.40E-09	1.54E-11	3.41E-09
	78.6-120	0.00E+00	0.00E+00	1.01E-08	0.00E+00	0.00E+00	3.40E-09	1.54E-11	3.41E-09
TC3	0-19.2	0.00E+00	0.00E+00	9.22E-09	0.00E+00	0.00E+00	2.98E-09	1.38E-11	2.99E-09
	19.2-41	0.00E+00	0.00E+00	9.22E-09	0.00E+00	0.00E+00	2.98E-09	1.38E-11	2.99E-09
	0-19.2	0.00E+00	0.00E+00	1.01E-08	0.00E+00	0.00E+00	2.98E-09	1.38E-11	2.99E-09
	19.2-41	0.00E+00	0.00E+00	1.01E-08	0.00E+00	0.00E+00	2.98E-09	1.38E-11	2.99E-09
TC4	0-19.2	0.00E+00	0.00E+00	1.01E-08	0.00E+00	0.00E+00	3.00E-09	1.44E-11	3.01E-09
	19.2-41	0.00E+00	0.00E+00	1.01E-08	0.00E+00	0.00E+00	3.00E-09	1.44E-11	3.01E-09
	41-78.6	0.00E+00	0.00E+00	1.01E-08	0.00E+00	0.00E+00	3.01E-09	1.50E-11	3.02E-09
	78.6-120	0.00E+00	0.00E+00	1.01E-08	0.00E+00	0.00E+00	3.01E-09	1.50E-11	3.02E-09
TC5	0-19.2	0.00E+00	0.00E+00	1.11E-08	0.00E+00	0.00E+00	3.41E-09	1.57E-11	3.42E-09
	19.2-41	0.00E+00	0.00E+00	1.11E-08	0.00E+00	0.00E+00	3.41E-09	1.58E-11	3.42E-09
	41-78.6	0.00E+00	0.00E+00	1.11E-08	0.00E+00	0.00E+00	3.42E-09	1.62E-11	3.43E-09
	78.6-120	0.00E+00	0.00E+00	1.11E-08	0.00E+00	0.00E+00	3.42E-09	1.62E-11	3.43E-09
TC6	0-19.2	0.00E+00	0.00E+00	1.11E-08	0.00E+00	0.00E+00	3.45E-09	1.76E-11	3.47E-09
	19.2-41	0.00E+00	0.00E+00	1.11E-08	0.00E+00	0.00E+00	3.45E-09	1.76E-11	3.47E-09
	41-78.6	0.00E+00	0.00E+00	1.11E-08	0.00E+00	0.00E+00	3.45E-09	1.76E-11	3.47E-09
	78.6-120	0.00E+00	0.00E+00	1.11E-08	0.00E+00	0.00E+00	3.45E-09	1.76E-11	3.47E-09
AP1	0-19.2	0.00E+00	0.00E+00	1.20E-08	0.00E+00	0.00E+00	5.03E-09	2.02E-11	5.05E-09
AP2	0-19.2	0.00E+00	0.00E+00	1.20E-08	0.00E+00	0.00E+00	6.24E-09	2.35E-11	6.27E-09
AP3	0-19.2	0.00E+00	0.00E+00	1.75E-08	0.00E+00	0.00E+00	1.45E-08	4.75E-11	1.45E-08
AP4	0-19.2	0.00E+00	0.00E+00	8.30E-09	0.00E+00	0.00E+00	5.43E-09	2.11E-11	5.45E-09
	19.2-41	0.00E+00	0.00E+00	8.30E-09	0.00E+00	0.00E+00	5.43E-09	2.11E-11	5.45E-09
	41-78.6	0.00E+00	0.00E+00	8.30E-09	0.00E+00	0.00E+00	5.43E-09	2.12E-11	5.45E-09
	78.6-120	0.00E+00	0.00E+00	8.30E-09	0.00E+00	0.00E+00	5.43E-09	2.13E-11	5.45E-09
AP5	0-19.2	0.00E+00	0.00E+00	8.30E-09	0.00E+00	0.00E+00	4.62E-09	1.89E-11	4.64E-09
AP6	0-19.2	0.00E+00	0.00E+00	8.30E-09	0.00E+00	0.00E+00	4.62E-09	1.92E-11	4.64E-09
AP7	0-19.2	0.00E+00	0.00E+00	9.22E-09	0.00E+00	0.00E+00	6.23E-09	2.31E-11	6.25E-09
	19.2-41	0.00E+00	0.00E+00	9.22E-09	0.00E+00	0.00E+00	6.23E-09	2.29E-11	6.25E-09
	41-78.6	0.00E+00	0.00E+00	9.22E-09	0.00E+00	0.00E+00	6.23E-09	2.30E-11	6.25E-09
AP8	0-19.2	0.00E+00	0.00E+00	8.30E-09	0.00E+00	0.00E+00	4.62E-09	1.92E-11	4.64E-09
KT1	0-19.2	0.00E+00	0.00E+00	4.61E-09	0.00E+00	0.00E+00	8.68E-10	4.64E-12	8.73E-10
	19.2-41	0.00E+00	0.00E+00	4.61E-09	0.00E+00	0.00E+00	8.69E-10	4.69E-12	8.74E-10
	41-78.6	0.00E+00	0.00E+00	4.61E-09	0.00E+00	0.00E+00	8.69E-10	4.69E-12	8.74E-10
	78.6-120	0.00E+00	0.00E+00	4.61E-09	0.00E+00	0.00E+00	8.71E-10	4.75E-12	8.75E-10
KT2	0-19.2	0.00E+00	0.00E+00	3.69E-09	0.00E+00	0.00E+00	8.68E-10	4.64E-12	8.73E-10
	19.2-41	0.00E+00	0.00E+00	3.69E-09	0.00E+00	0.00E+00	8.69E-10	4.69E-12	8.74E-10
	41-78.6	0.00E+00	0.00E+00	3.69E-09	0.00E+00	0.00E+00	8.69E-10	4.69E-12	8.74E-10
	78.6-120	0.00E+00	0.00E+00	3.69E-09	0.00E+00	0.00E+00	8.71E-10	4.75E-12	8.75E-10
KT3	0-19.2	0.00E+00	0.00E+00	3.69E-09	0.00E+00	0.00E+00	8.68E-10	4.64E-12	8.73E-10
KT4	0-19.2	0.00E+00	0.00E+00	3.69E-09	0.00E+00	0.00E+00	8.69E-10	4.69E-12	8.74E-10
KT5	0-19.2	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	8.68E-10	4.64E-12	8.73E-10
	19.2-41	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	8.69E-10	4.69E-12	8.74E-10
	41-78.6	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	8.69E-10	4.69E-12	8.74E-10
	78.6-120	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	8.71E-10	4.75E-12	8.75E-10
KT6	0-19.2	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	8.74E-10	4.91E-12	8.79E-10
	19.2-41	0.00E+00	0.00E+00	2.77E-09	0.00E+00	0.00E+00	8.68E-10	4.64E-12	8.73E-10
	41-78.6	0.00E+00	0.00E+00	2.77E-09	0.00E+00	0.00E+00	8.69E-10	4.69E-12	8.74E-10
KT7	0-19.2	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	8.64E-10	4.47E-12	8.69E-10
KT8	0-19.2	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	4.59E-10	3.39E-12	4.63E-10
	19.2-41	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	4.59E-10	3.39E-12	4.63E-10
	41-78.6	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	4.59E-10	3.39E-12	4.63E-10
78.6-120	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	4.62E-10	3.50E-12	4.66E-10	

Predicted Maximum Permitted Concentration of Certain Metals in Foods - Produce - The IWMF at the TTAL site

ASR	Assessment Height (mAG)	Antimony	Arsenic	Cadmium	Chromium	Lead	Mercuric chloride	Methyl mercury	Mercury
Maximum Permitted Concentration (ppm)		1	1.4	0.1	1	6			0.5
TT1	0-19.2	2.30E-07	1.36E-04	1.35E-05	1.36E-04	1.36E-04	1.81E-04	8.50E-06	1.89E-04
TT2	0-19.2	8.94E-07	3.13E-05	3.13E-06	3.16E-05	3.13E-05	6.49E-04	2.94E-05	6.79E-04
TT3	0-19.2	1.38E-06	5.22E-05	5.21E-06	5.27E-05	5.22E-05	1.00E-03	4.54E-05	1.05E-03
TT4	0-19.2	7.81E-06	2.27E-10	2.88E-08	2.66E-06	3.99E-07	5.64E-03	2.55E-04	5.89E-03
TT5	0-19.2	3.77E-07	2.71E-04	2.71E-05	2.72E-04	2.71E-04	3.08E-04	1.47E-05	3.23E-04
TT6	0-19.2	1.33E-06	1.46E-04	1.46E-05	1.47E-04	1.46E-04	9.77E-04	4.45E-05	1.02E-03
TM1	0-19.2	3.99E-08	1.04E-05	1.04E-06	1.05E-05	1.04E-05	3.41E-05	2.44E-06	3.66E-05
	19.2-41	3.99E-08	1.04E-05	1.04E-06	1.05E-05	1.04E-05	3.43E-05	2.47E-06	3.67E-05
	41-78.6	3.99E-08	1.04E-05	1.04E-06	1.05E-05	1.04E-05	3.45E-05	2.55E-06	3.71E-05
	78.6-120	3.99E-08	1.04E-05	1.04E-06	1.05E-05	1.04E-05	3.51E-05	2.70E-06	3.78E-05
TM2	0-19.2	2.63E-08	3.13E-05	3.12E-06	3.13E-05	3.13E-05	2.73E-05	2.13E-06	2.94E-05
	19.2-41	2.63E-08	3.13E-05	3.12E-06	3.13E-05	3.13E-05	2.74E-05	2.16E-06	2.96E-05
	41-78.6	2.63E-08	3.13E-05	3.12E-06	3.13E-05	3.13E-05	2.77E-05	2.24E-06	2.99E-05
	78.6-120	2.63E-08	3.13E-05	3.12E-06	3.13E-05	3.13E-05	2.82E-05	2.39E-06	3.06E-05
TM3	0-19.2	4.41E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	3.95E-05	2.68E-06	4.21E-05
	19.2-41	4.41E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	3.96E-05	2.71E-06	4.23E-05
	41-78.6	4.41E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	3.98E-05	2.79E-06	4.26E-05
	78.6-120	4.41E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	4.04E-05	2.94E-06	4.33E-05
TM4	0-19.2	4.73E-08	3.13E-05	3.12E-06	3.13E-05	3.13E-05	4.17E-05	2.78E-06	4.45E-05
	19.2-41	4.73E-08	3.13E-05	3.12E-06	3.13E-05	3.13E-05	4.18E-05	2.81E-06	4.47E-05
	41-78.6	4.73E-08	3.13E-05	3.12E-06	3.13E-05	3.13E-05	4.21E-05	2.89E-06	4.50E-05
	78.6-120	4.73E-08	3.13E-05	3.12E-06	3.13E-05	3.13E-05	4.27E-05	3.05E-06	4.57E-05
TM5	0-19.2	5.36E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	4.58E-05	2.85E-06	4.86E-05
	19.2-41	5.36E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	4.59E-05	2.88E-06	4.88E-05
	41-78.6	5.36E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	4.63E-05	2.99E-06	4.93E-05
	78.6-120	5.36E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	4.74E-05	3.30E-06	5.07E-05
TM6	0-19.2	7.04E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	5.72E-05	3.55E-06	6.07E-05
	19.2-41	7.04E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	5.74E-05	3.59E-06	6.09E-05
	41-78.6	7.04E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	5.78E-05	3.72E-06	6.15E-05
	78.6-120	7.04E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	5.88E-05	3.99E-06	6.28E-05
SLW1	0-19.2	5.78E-08	4.70E-04	4.68E-05	4.70E-04	4.69E-04	7.87E-05	5.92E-06	8.46E-05
SLW2	0-19.2	6.31E-08	5.01E-04	5.00E-05	5.01E-04	5.01E-04	8.64E-05	6.30E-06	9.27E-05
SLW3	0-19.2	1.37E-08	1.77E-04	1.77E-05	1.78E-04	1.77E-04	2.53E-05	3.33E-06	2.86E-05
TC1	0-19.2	7.36E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	1.57E-05	2.28E-06	1.80E-05
	19.2-41	7.36E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	1.57E-05	2.28E-06	1.80E-05
	41-78.6	7.36E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	1.59E-05	2.32E-06	1.82E-05
	78.6-120	7.36E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	1.59E-05	2.32E-06	1.82E-05
	120-175	7.36E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	1.59E-05	2.32E-06	1.82E-05
	19.2-41	4.20E-09	1.15E-04	1.14E-05	1.15E-04	1.15E-04	1.27E-05	2.15E-06	1.49E-05
TC2	19.2-41	4.20E-09	1.15E-04	1.14E-05	1.15E-04	1.15E-04	1.27E-05	2.15E-06	1.49E-05
	41-78.6	4.20E-09	1.15E-04	1.14E-05	1.15E-04	1.15E-04	1.28E-05	2.18E-06	1.50E-05
	78.6-120	4.20E-09	1.15E-04	1.14E-05	1.15E-04	1.15E-04	1.28E-05	2.18E-06	1.50E-05
	120-175	4.20E-09	1.15E-04	1.14E-05	1.15E-04	1.15E-04	1.28E-05	2.18E-06	1.50E-05
TC3	0-19.2	3.15E-09	1.04E-04	1.04E-05	1.04E-04	1.04E-04	1.15E-05	1.99E-06	1.35E-05
	19.2-41	3.15E-09	1.04E-04	1.04E-05	1.04E-04	1.04E-04	1.15E-05	1.99E-06	1.35E-05
	41-78.6	3.15E-09	1.04E-04	1.04E-05	1.04E-04	1.04E-04	1.15E-05	1.99E-06	1.35E-05
TC4	0-19.2	4.20E-09	1.15E-04	1.14E-05	1.15E-04	1.15E-04	1.15E-05	1.99E-06	1.35E-05
	19.2-41	4.20E-09	1.15E-04	1.14E-05	1.15E-04	1.15E-04	1.15E-05	1.99E-06	1.35E-05
	41-78.6	4.20E-09	1.15E-04	1.14E-05	1.15E-04	1.15E-04	1.18E-05	2.07E-06	1.39E-05
	78.6-120	4.20E-09	1.15E-04	1.14E-05	1.15E-04	1.15E-04	1.22E-05	2.18E-06	1.43E-05
TC5	120-175	4.20E-09	1.15E-04	1.14E-05	1.15E-04	1.15E-04	1.27E-05	2.33E-06	1.51E-05
	0-19.2	6.31E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	1.32E-05	2.27E-06	1.54E-05
	19.2-41	6.31E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	1.32E-05	2.29E-06	1.55E-05
	41-78.6	6.31E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	1.36E-05	2.40E-06	1.60E-05
	78.6-120	6.31E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	1.41E-05	2.55E-06	1.67E-05
	120-175	6.31E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	1.50E-05	2.78E-06	1.77E-05
TC6	0-19.2	6.31E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	1.32E-05	2.27E-06	1.54E-05
	19.2-41	6.31E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	1.32E-05	2.29E-06	1.55E-05
	41-78.6	6.31E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	1.36E-05	2.40E-06	1.60E-05
	78.6-120	6.31E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	1.41E-05	2.55E-06	1.67E-05
	120-175	6.31E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	1.50E-05	2.78E-06	1.77E-05
	19.2-41	9.46E-09	1.36E-04	1.35E-05	1.36E-04	1.36E-04	1.64E-05	2.46E-06	1.88E-05
AP1	0-19.2	1.26E-08	1.36E-04	1.35E-05	1.36E-04	1.36E-04	1.86E-05	2.56E-06	2.12E-05
AP2	0-19.2	2.21E-08	1.98E-04	1.98E-05	1.98E-04	1.98E-04	3.58E-05	3.78E-06	3.96E-05
AP3	0-19.2	1.05E-08	9.39E-05	9.37E-06	9.40E-05	9.39E-05	1.69E-05	2.43E-06	1.93E-05
AP4	19.2-41	1.05E-08	9.39E-05	9.37E-06	9.40E-05	9.39E-05	1.69E-05	2.43E-06	1.93E-05
	41-78.6	1.05E-08	9.39E-05	9.37E-06	9.40E-05	9.39E-05	1.70E-05	2.46E-06	1.95E-05
	78.6-120	1.05E-08	9.39E-05	9.37E-06	9.40E-05	9.39E-05	1.71E-05	2.49E-06	1.96E-05
	19.2-41	7.36E-09	9.39E-05	9.37E-06	9.40E-05	9.39E-05	1.54E-05	2.36E-06	1.77E-05
AP5	0-19.2	6.31E-09	9.39E-05	9.37E-06	9.40E-05	9.39E-05	1.56E-05	2.44E-06	1.81E-05
AP6	0-19.2	1.05E-08	1.04E-04	1.04E-05	1.04E-04	1.04E-04	1.82E-05	2.43E-06	2.06E-05
AP7	19.2-41	1.05E-08	1.04E-04	1.04E-05	1.04E-04	1.04E-04	1.81E-05	2.40E-06	2.05E-05
	41-78.6	1.05E-08	1.04E-04	1.04E-05	1.04E-04	1.04E-04	1.81E-05	2.42E-06	2.06E-05
	78.6-120	1.05E-08	1.04E-04	1.04E-05	1.04E-04	1.04E-04	1.81E-05	2.42E-06	2.06E-05
AP8	0-19.2	7.36E-09	9.39E-05	9.37E-06	9.40E-05	9.39E-05	1.56E-05	2.44E-06	1.81E-05
KT1	0-19.2	1.05E-09	5.22E-05	5.20E-06	5.22E-05	5.22E-05	3.99E-06	7.65E-07	4.75E-06
	19.2-41	1.05E-09	5.22E-05	5.20E-06	5.22E-05	5.22E-05	4.04E-06	7.81E-07	4.82E-06
	41-78.6	1.05E-09	5.22E-05	5.20E-06	5.22E-05	5.22E-05	4.04E-06	7.81E-07	4.82E-06
	78.6-120	1.05E-09	5.22E-05	5.20E-06	5.22E-05	5.22E-05	4.04E-06	7.81E-07	4.82E-06
	120-175	1.05E-09	5.22E-05	5.20E-06	5.22E-05	5.22E-05	4.10E-06	7.96E-07	4.89E-06
	0-19.2	1.05E-09	4.17E-05	4.16E-06	4.18E-05	4.17E-05	3.99E-06	7.65E-07	4.75E-06
KT2	19.2-41	1.05E-09	4.17E-05	4.16E-06	4.18E-05	4.17E-05	4.04E-06	7.81E-07	4.82E-06
	41-78.6	1.05E-09	4.17E-05	4.16E-06	4.18E-05	4.17E-05	4.04E-06	7.81E-07	4.82E-06
	78.6-120	1.05E-09	4.17E-05	4.16E-06	4.18E-05	4.17E-05	4.10E-06	7.96E-07	4.89E-06
	120-175	1.05E-09	4.17E-05	4.16E-06	4.18E-05	4.17E-05	3.99E-06	7.65E-07	4.75E-06
KT3	0-19.2	1.05E-09	4.17E-05	4.16E-06	4.18E-05	4.17E-05	4.04E-06	7.81E-07	4.82E-06
KT4	0-19.2	1.05E-09	4.17E-05	4.16E-06	4.18E-05	4.17E-05	3.99E-06	7.65E-07	4.75E-06
KT5	0-19.2	1.05E-09	2.09E-05	2.08E-06	2.09E-05	2.09E-05	3.99E-06	7.65E-07	4.75E-06
	19.2-41	1.05E-09	2.09E-05	2.08E-06	2.09E-05	2.09E-05	4.04E-06	7.81E-07	4.82E-06
	41-78.6	1.05E-09	2.09E-05	2.08E-06	2.09E-05	2.09E-05	4.04E-06	7.81E-07	4.82E-06
	78.6-120	1.05E-09	2.09E-05	2.08E-06	2.09E-05	2.09E-05	4.10E-06	7.96E-07	4.89E-06
	120-175	1.05E-09	2.09E-05	2.08E-06	2.09E-05	2.09E-05	4.26E-06	8.43E-07	5.11E-06
	0-19.2	1.05E-09	3.13E-05	3.12E-06	3.13E-05	3.13E-05	3.99E-06	7.65E-07	4.75E-06
KT6	19.2-41	1.05E-09	3.13E-05	3.12E-06	3.13E-05	3.13E-05	4.04E-06	7.81E-07	4.82E-06
	78.6-120	1.05E-09	3.13E-05	3.12E-06	3.13E-05	3.13E-05	3.82E-06	7.19E-07	4.54E-06
KT7	0-19.2	0.00E+00	2.09E-05	2.08E-06	2.09E-05	2.09E-05	3.06E-06	6.85E-07	3.75E-06
KT8	19.2-41	0.00E+00	2.09E-05	2.08E-06	2.09E-05	2.09E-05	3.06E-06	6.85E-07	3.75E-06
	41-78.6	0.00E+00	2.09E-05	2.08E-06	2.09E-05				

Predicted Maximum Permitted Concentration of Certain Metals in Foods - Chicken - The IWMF at the artificial island near SKC

ASR	Assessment Height (mAG)	Antimony	Arsenic	Cadmium	Chromium	Lead	Mercuric chloride	Methyl mercury	Mercury
Maximum Permitted Concentration (ppm)		1	1.4	0.2	1	6			0.5
SKC1	0-19.2	0.00E+00	0.00E+00	1.46E-10	0.00E+00	0.00E+00	3.11E-05	8.34E-08	3.12E-05
SL1	0-19.2	0.00E+00	0.00E+00	1.70E-10	0.00E+00	0.00E+00	1.48E-06	3.96E-09	1.48E-06
SL2	0-19.2	0.00E+00	0.00E+00	2.54E-10	0.00E+00	0.00E+00	1.77E-06	4.74E-09	1.77E-06
SLW1	0-19.2	0.00E+00	0.00E+00	1.91E-10	0.00E+00	0.00E+00	5.80E-07	1.55E-09	5.81E-07
SLW2	0-19.2	0.00E+00	0.00E+00	2.33E-10	0.00E+00	0.00E+00	5.97E-07	1.60E-09	5.98E-07
SLW3	0-19.2	0.00E+00	0.00E+00	1.58E-10	0.00E+00	0.00E+00	5.80E-07	1.55E-09	5.81E-07
TC1	0-19.2	0.00E+00	0.00E+00	1.25E-10	0.00E+00	0.00E+00	3.93E-07	1.05E-09	3.94E-07
	19.2-41	0.00E+00	0.00E+00	1.25E-10	0.00E+00	0.00E+00	3.93E-07	1.05E-09	3.94E-07
	41-78.6	0.00E+00	0.00E+00	1.25E-10	0.00E+00	0.00E+00	3.93E-07	1.05E-09	3.94E-07
	78.6-120	0.00E+00	0.00E+00	1.25E-10	0.00E+00	0.00E+00	3.93E-07	1.05E-09	3.94E-07
	120-175	0.00E+00	0.00E+00	1.25E-10	0.00E+00	0.00E+00	3.93E-07	1.05E-09	3.94E-07
TC2	0-19.2	0.00E+00	0.00E+00	1.50E-10	0.00E+00	0.00E+00	3.93E-07	1.05E-09	3.94E-07
	19.2-41	0.00E+00	0.00E+00	1.50E-10	0.00E+00	0.00E+00	3.93E-07	1.05E-09	3.94E-07
	41-78.6	0.00E+00	0.00E+00	1.50E-10	0.00E+00	0.00E+00	3.93E-07	1.05E-09	3.94E-07
	78.6-120	0.00E+00	0.00E+00	1.50E-10	0.00E+00	0.00E+00	3.93E-07	1.05E-09	3.94E-07
	120-175	0.00E+00	0.00E+00	1.50E-10	0.00E+00	0.00E+00	3.93E-07	1.05E-09	3.94E-07
TC3	0-19.2	0.00E+00	0.00E+00	1.58E-10	0.00E+00	0.00E+00	3.93E-07	1.05E-09	3.94E-07
	19.2-41	0.00E+00	0.00E+00	1.58E-10	0.00E+00	0.00E+00	3.93E-07	1.05E-09	3.94E-07
	0-19.2	0.00E+00	0.00E+00	1.66E-10	0.00E+00	0.00E+00	4.10E-07	1.10E-09	4.11E-07
TC4	19.2-41	0.00E+00	0.00E+00	1.66E-10	0.00E+00	0.00E+00	4.10E-07	1.10E-09	4.11E-07
	41-78.6	0.00E+00	0.00E+00	1.66E-10	0.00E+00	0.00E+00	4.10E-07	1.10E-09	4.11E-07
	78.6-120	0.00E+00	0.00E+00	1.66E-10	0.00E+00	0.00E+00	4.10E-07	1.10E-09	4.11E-07
	120-175	0.00E+00	0.00E+00	1.66E-10	0.00E+00	0.00E+00	4.10E-07	1.10E-09	4.11E-07
	0-19.2	0.00E+00	0.00E+00	1.75E-10	0.00E+00	0.00E+00	4.78E-07	1.28E-09	4.79E-07
TC5	19.2-41	0.00E+00	0.00E+00	1.75E-10	0.00E+00	0.00E+00	4.78E-07	1.28E-09	4.79E-07
	41-78.6	0.00E+00	0.00E+00	1.75E-10	0.00E+00	0.00E+00	4.78E-07	1.28E-09	4.79E-07
	78.6-120	0.00E+00	0.00E+00	1.75E-10	0.00E+00	0.00E+00	4.78E-07	1.28E-09	4.79E-07
	120-175	0.00E+00	0.00E+00	1.75E-10	0.00E+00	0.00E+00	4.78E-07	1.28E-09	4.79E-07
	0-19.2	0.00E+00	0.00E+00	1.34E-10	0.00E+00	0.00E+00	6.14E-07	1.64E-09	6.15E-07
TC6	19.2-41	0.00E+00	0.00E+00	1.34E-10	0.00E+00	0.00E+00	6.14E-07	1.64E-09	6.15E-07
	41-78.6	0.00E+00	0.00E+00	1.34E-10	0.00E+00	0.00E+00	6.14E-07	1.64E-09	6.15E-07
	78.6-120	0.00E+00	0.00E+00	1.34E-10	0.00E+00	0.00E+00	5.97E-07	1.60E-09	5.98E-07
	120-175	0.00E+00	0.00E+00	1.34E-10	0.00E+00	0.00E+00	5.97E-07	1.60E-09	5.98E-07
	0-19.2	0.00E+00	0.00E+00	1.83E-10	0.00E+00	0.00E+00	4.10E-07	1.10E-09	4.11E-07
AP1	0-19.2	0.00E+00	0.00E+00	1.83E-10	0.00E+00	0.00E+00	4.10E-07	1.10E-09	4.11E-07
AP2	0-19.2	0.00E+00	0.00E+00	1.83E-10	0.00E+00	0.00E+00	3.76E-07	1.01E-09	3.77E-07
AP3	0-19.2	0.00E+00	0.00E+00	1.58E-10	0.00E+00	0.00E+00	5.46E-07	1.46E-09	5.47E-07
AP4	0-19.2	0.00E+00	0.00E+00	1.91E-10	0.00E+00	0.00E+00	3.08E-07	8.26E-10	3.09E-07
	19.2-41	0.00E+00	0.00E+00	1.91E-10	0.00E+00	0.00E+00	3.08E-07	8.26E-10	3.09E-07
	41-78.6	0.00E+00	0.00E+00	1.91E-10	0.00E+00	0.00E+00	3.08E-07	8.26E-10	3.09E-07
	78.6-120	0.00E+00	0.00E+00	1.91E-10	0.00E+00	0.00E+00	3.08E-07	8.26E-10	3.09E-07
	0-19.2	0.00E+00	0.00E+00	1.66E-10	0.00E+00	0.00E+00	3.25E-07	8.71E-10	3.26E-07
AP5	0-19.2	0.00E+00	0.00E+00	1.66E-10	0.00E+00	0.00E+00	3.25E-07	8.71E-10	3.26E-07
AP6	0-19.2	0.00E+00	0.00E+00	1.66E-10	0.00E+00	0.00E+00	3.08E-07	8.26E-10	3.09E-07
AP7	19.2-41	0.00E+00	0.00E+00	1.66E-10	0.00E+00	0.00E+00	3.08E-07	8.26E-10	3.09E-07
	41-78.6	0.00E+00	0.00E+00	1.66E-10	0.00E+00	0.00E+00	3.08E-07	8.26E-10	3.09E-07
	0-19.2	0.00E+00	0.00E+00	1.83E-10	0.00E+00	0.00E+00	3.08E-07	8.26E-10	3.09E-07
AP8	0-19.2	0.00E+00	0.00E+00	1.83E-10	0.00E+00	0.00E+00	3.08E-07	8.26E-10	3.09E-07
KT1	0-19.2	0.00E+00	0.00E+00	1.16E-10	0.00E+00	0.00E+00	2.57E-07	6.89E-10	2.58E-07
	19.2-41	0.00E+00	0.00E+00	1.16E-10	0.00E+00	0.00E+00	2.57E-07	6.89E-10	2.58E-07
	41-78.6	0.00E+00	0.00E+00	1.16E-10	0.00E+00	0.00E+00	2.57E-07	6.89E-10	2.58E-07
	78.6-120	0.00E+00	0.00E+00	1.16E-10	0.00E+00	0.00E+00	2.57E-07	6.89E-10	2.58E-07
	0-19.2	0.00E+00	0.00E+00	1.16E-10	0.00E+00	0.00E+00	2.57E-07	6.89E-10	2.58E-07
KT2	19.2-41	0.00E+00	0.00E+00	1.16E-10	0.00E+00	0.00E+00	2.57E-07	6.89E-10	2.58E-07
	41-78.6	0.00E+00	0.00E+00	1.16E-10	0.00E+00	0.00E+00	2.57E-07	6.89E-10	2.58E-07
	78.6-120	0.00E+00	0.00E+00	1.16E-10	0.00E+00	0.00E+00	2.57E-07	6.89E-10	2.58E-07
	0-19.2	0.00E+00	0.00E+00	1.16E-10	0.00E+00	0.00E+00	2.74E-07	7.35E-10	2.75E-07
	19.2-41	0.00E+00	0.00E+00	1.16E-10	0.00E+00	0.00E+00	2.74E-07	7.35E-10	2.75E-07
KT3	0-19.2	0.00E+00	0.00E+00	1.16E-10	0.00E+00	0.00E+00	2.74E-07	7.35E-10	2.75E-07
KT4	19.2-41	0.00E+00	0.00E+00	1.16E-10	0.00E+00	0.00E+00	2.57E-07	6.89E-10	2.58E-07
	0-19.2	0.00E+00	0.00E+00	8.30E-11	0.00E+00	0.00E+00	1.87E-07	5.00E-10	1.87E-07
	19.2-41	0.00E+00	0.00E+00	8.30E-11	0.00E+00	0.00E+00	1.87E-07	5.00E-10	1.87E-07
	41-78.6	0.00E+00	0.00E+00	8.30E-11	0.00E+00	0.00E+00	1.87E-07	5.00E-10	1.87E-07
	78.6-120	0.00E+00	0.00E+00	8.30E-11	0.00E+00	0.00E+00	1.87E-07	5.00E-10	1.87E-07
KT5	120-175	0.00E+00	0.00E+00	8.30E-11	0.00E+00	0.00E+00	1.87E-07	5.00E-10	1.87E-07
	0-19.2	0.00E+00	0.00E+00	9.96E-11	0.00E+00	0.00E+00	2.40E-07	6.44E-10	2.41E-07
	19.2-41	0.00E+00	0.00E+00	9.96E-11	0.00E+00	0.00E+00	2.40E-07	6.44E-10	2.41E-07
	0-19.2	0.00E+00	0.00E+00	5.81E-11	0.00E+00	0.00E+00	1.36E-07	3.63E-10	1.36E-07
	19.2-41	0.00E+00	0.00E+00	4.98E-11	0.00E+00	0.00E+00	1.19E-07	3.18E-10	1.19E-07
KT7	41-78.6	0.00E+00	0.00E+00	4.98E-11	0.00E+00	0.00E+00	1.19E-07	3.18E-10	1.19E-07
	78.6-120	0.00E+00	0.00E+00	4.98E-11	0.00E+00	0.00E+00	1.19E-07	3.18E-10	1.19E-07
	0-19.2	0.00E+00	0.00E+00	2.99E-11	0.00E+00	0.00E+00	1.31E-06	3.50E-09	1.31E-06
	SL3	0-19.2	0.00E+00	0.00E+00	2.99E-11	0.00E+00	1.31E-06	3.50E-09	1.31E-06
	CC1	0-19.2	0.00E+00	0.00E+00	6.36E-11	0.00E+00	0.00E+00	1.63E-06	4.37E-09
CC2	0-19.2	0.00E+00	0.00E+00	6.04E-11	0.00E+00	0.00E+00	9.16E-07	2.45E-09	9.18E-07
CC3	0-19.2	0.00E+00	0.00E+00	7.26E-11	0.00E+00	0.00E+00	1.83E-06	4.91E-09	1.84E-06
CC4	0-19.2	0.00E+00	0.00E+00	3.41E-11	0.00E+00	0.00E+00	4.24E-07	1.14E-09	4.25E-07

Predicted Maximum Permitted Concentration of Certain Metals in Foods - Pork - The IWMF at the artificial island near SKC

ASR	Assessment Height (mAG)	Antimony	Arsenic	Cadmium	Chromium	Lead	Mercuric chloride	Methyl mercury	Mercury
Maximum Permitted Concentration (ppm)		1	1.4	0.2	1	6			0.5
SKC1	0-19.2	0.00E+00	0.00E+00	9.29E-10	0.00E+00	0.00E+00	7.42E-07	1.98E-09	7.44E-07
SL1	0-19.2	0.00E+00	0.00E+00	1.84E-08	0.00E+00	0.00E+00	3.54E-08	1.04E-10	3.56E-08
SL2	0-19.2	0.00E+00	0.00E+00	2.77E-08	0.00E+00	0.00E+00	4.24E-08	1.24E-10	4.25E-08
SLW1	0-19.2	0.00E+00	0.00E+00	2.12E-08	0.00E+00	0.00E+00	1.43E-08	5.59E-11	1.43E-08
SLW2	0-19.2	0.00E+00	0.00E+00	2.58E-08	0.00E+00	0.00E+00	1.47E-08	5.87E-11	1.48E-08
SLW3	0-19.2	0.00E+00	0.00E+00	1.75E-08	0.00E+00	0.00E+00	1.43E-08	5.74E-11	1.43E-08
TC1	0-19.2	0.00E+00	0.00E+00	1.38E-08	0.00E+00	0.00E+00	9.77E-09	4.22E-11	9.81E-09
	19.2-41	0.00E+00	0.00E+00	1.38E-08	0.00E+00	0.00E+00	9.78E-09	4.27E-11	9.82E-09
	41-78.6	0.00E+00	0.00E+00	1.38E-08	0.00E+00	0.00E+00	9.80E-09	4.35E-11	9.84E-09
	78.6-120	0.00E+00	0.00E+00	1.38E-08	0.00E+00	0.00E+00	9.84E-09	4.53E-11	9.89E-09
TC2	120-175	0.00E+00	0.00E+00	1.38E-08	0.00E+00	0.00E+00	9.92E-09	4.86E-11	9.97E-09
	0-19.2	0.00E+00	0.00E+00	1.66E-08	0.00E+00	0.00E+00	9.77E-09	4.22E-11	9.81E-09
	19.2-41	0.00E+00	0.00E+00	1.66E-08	0.00E+00	0.00E+00	9.78E-09	4.27E-11	9.82E-09
	41-78.6	0.00E+00	0.00E+00	1.66E-08	0.00E+00	0.00E+00	9.80E-09	4.35E-11	9.84E-09
TC3	78.6-120	0.00E+00	0.00E+00	1.66E-08	0.00E+00	0.00E+00	9.84E-09	4.53E-11	9.89E-09
	120-175	0.00E+00	0.00E+00	1.66E-08	0.00E+00	0.00E+00	9.92E-09	4.86E-11	9.97E-09
	0-19.2	0.00E+00	0.00E+00	1.75E-08	0.00E+00	0.00E+00	9.77E-09	4.24E-11	9.82E-09
	19.2-41	0.00E+00	0.00E+00	1.75E-08	0.00E+00	0.00E+00	9.78E-09	4.29E-11	9.83E-09
TC4	0-19.2	0.00E+00	0.00E+00	1.84E-08	0.00E+00	0.00E+00	1.02E-08	4.35E-11	1.02E-08
	19.2-41	0.00E+00	0.00E+00	1.84E-08	0.00E+00	0.00E+00	1.02E-08	4.40E-11	1.02E-08
	41-78.6	0.00E+00	0.00E+00	1.84E-08	0.00E+00	0.00E+00	1.02E-08	4.54E-11	1.03E-08
	78.6-120	0.00E+00	0.00E+00	1.84E-08	0.00E+00	0.00E+00	1.03E-08	4.97E-11	1.04E-08
TC5	120-175	0.00E+00	0.00E+00	1.84E-08	0.00E+00	0.00E+00	1.05E-08	5.52E-11	1.05E-08
	0-19.2	0.00E+00	0.00E+00	1.94E-08	0.00E+00	0.00E+00	1.18E-08	5.02E-11	1.19E-08
	19.2-41	0.00E+00	0.00E+00	1.94E-08	0.00E+00	0.00E+00	1.19E-08	5.04E-11	1.19E-08
	41-78.6	0.00E+00	0.00E+00	1.94E-08	0.00E+00	0.00E+00	1.19E-08	5.12E-11	1.19E-08
TC6	78.6-120	0.00E+00	0.00E+00	1.94E-08	0.00E+00	0.00E+00	1.19E-08	5.37E-11	1.20E-08
	120-175	0.00E+00	0.00E+00	1.94E-08	0.00E+00	0.00E+00	1.20E-08	5.68E-11	1.21E-08
	0-19.2	0.00E+00	0.00E+00	1.47E-08	0.00E+00	0.00E+00	1.51E-08	5.88E-11	1.51E-08
	19.2-41	0.00E+00	0.00E+00	1.47E-08	0.00E+00	0.00E+00	1.51E-08	5.90E-11	1.51E-08
AP1	41-78.6	0.00E+00	0.00E+00	1.47E-08	0.00E+00	0.00E+00	1.51E-08	5.98E-11	1.52E-08
	78.6-120	0.00E+00	0.00E+00	1.47E-08	0.00E+00	0.00E+00	1.48E-08	6.12E-11	1.48E-08
	120-175	0.00E+00	0.00E+00	1.47E-08	0.00E+00	0.00E+00	1.48E-08	6.43E-11	1.49E-08
	0-19.2	0.00E+00	0.00E+00	2.03E-08	0.00E+00	0.00E+00	1.02E-08	4.54E-11	1.03E-08
AP2	0-19.2	0.00E+00	0.00E+00	2.03E-08	0.00E+00	0.00E+00	9.42E-09	4.33E-11	9.46E-09
AP3	0-19.2	0.00E+00	0.00E+00	1.75E-08	0.00E+00	0.00E+00	1.35E-08	5.41E-11	1.35E-08
AP4	0-19.2	0.00E+00	0.00E+00	2.12E-08	0.00E+00	0.00E+00	7.71E-09	3.51E-11	7.74E-09
	19.2-41	0.00E+00	0.00E+00	2.12E-08	0.00E+00	0.00E+00	7.71E-09	3.51E-11	7.74E-09
	41-78.6	0.00E+00	0.00E+00	2.12E-08	0.00E+00	0.00E+00	7.71E-09	3.53E-11	7.75E-09
	78.6-120	0.00E+00	0.00E+00	2.12E-08	0.00E+00	0.00E+00	7.72E-09	3.56E-11	7.75E-09
AP5	0-19.2	0.00E+00	0.00E+00	1.84E-08	0.00E+00	0.00E+00	8.11E-09	3.62E-11	8.15E-09
AP6	0-19.2	0.00E+00	0.00E+00	1.84E-08	0.00E+00	0.00E+00	8.16E-09	3.84E-11	8.20E-09
AP7	0-19.2	0.00E+00	0.00E+00	1.84E-08	0.00E+00	0.00E+00	7.67E-09	3.34E-11	7.70E-09
	19.2-41	0.00E+00	0.00E+00	1.84E-08	0.00E+00	0.00E+00	7.67E-09	3.35E-11	7.70E-09
	41-78.6	0.00E+00	0.00E+00	1.84E-08	0.00E+00	0.00E+00	7.67E-09	3.37E-11	7.71E-09
AP8	0-19.2	0.00E+00	0.00E+00	2.03E-08	0.00E+00	0.00E+00	7.76E-09	3.73E-11	7.80E-09
KT1	0-19.2	0.00E+00	0.00E+00	1.29E-08	0.00E+00	0.00E+00	6.25E-09	2.16E-11	6.27E-09
	19.2-41	0.00E+00	0.00E+00	1.29E-08	0.00E+00	0.00E+00	6.25E-09	2.16E-11	6.27E-09
	41-78.6	0.00E+00	0.00E+00	1.29E-08	0.00E+00	0.00E+00	6.25E-09	2.16E-11	6.27E-09
	78.6-120	0.00E+00	0.00E+00	1.29E-08	0.00E+00	0.00E+00	6.25E-09	2.16E-11	6.28E-09
KT2	0-19.2	0.00E+00	0.00E+00	1.29E-08	0.00E+00	0.00E+00	6.25E-09	2.16E-11	6.27E-09
	19.2-41	0.00E+00	0.00E+00	1.29E-08	0.00E+00	0.00E+00	6.25E-09	2.16E-11	6.27E-09
	41-78.6	0.00E+00	0.00E+00	1.29E-08	0.00E+00	0.00E+00	6.25E-09	2.16E-11	6.27E-09
	78.6-120	0.00E+00	0.00E+00	1.29E-08	0.00E+00	0.00E+00	6.25E-09	2.16E-11	6.28E-09
KT3	0-19.2	0.00E+00	0.00E+00	1.29E-08	0.00E+00	0.00E+00	6.66E-09	2.26E-11	6.68E-09
KT4	19.2-41	0.00E+00	0.00E+00	1.29E-08	0.00E+00	0.00E+00	6.66E-09	2.26E-11	6.68E-09
	0-19.2	0.00E+00	0.00E+00	1.29E-08	0.00E+00	0.00E+00	6.25E-09	2.16E-11	6.27E-09
KT5	0-19.2	0.00E+00	0.00E+00	9.22E-09	0.00E+00	0.00E+00	4.56E-09	1.67E-11	4.57E-09
	19.2-41	0.00E+00	0.00E+00	9.22E-09	0.00E+00	0.00E+00	4.56E-09	1.67E-11	4.57E-09
	41-78.6	0.00E+00	0.00E+00	9.22E-09	0.00E+00	0.00E+00	4.56E-09	1.67E-11	4.57E-09
	78.6-120	0.00E+00	0.00E+00	9.22E-09	0.00E+00	0.00E+00	4.56E-09	1.67E-11	4.57E-09
KT6	120-175	0.00E+00	0.00E+00	9.22E-09	0.00E+00	0.00E+00	4.56E-09	1.68E-11	4.58E-09
	0-19.2	0.00E+00	0.00E+00	1.11E-08	0.00E+00	0.00E+00	5.85E-09	2.05E-11	5.87E-09
KT7	19.2-41	0.00E+00	0.00E+00	1.11E-08	0.00E+00	0.00E+00	5.85E-09	2.05E-11	5.87E-09
KT8	0-19.2	0.00E+00	0.00E+00	6.45E-09	0.00E+00	0.00E+00	3.35E-09	1.36E-11	3.36E-09
	19.2-41	0.00E+00	0.00E+00	5.53E-09	0.00E+00	0.00E+00	2.94E-09	1.25E-11	2.96E-09
	41-78.6	0.00E+00	0.00E+00	5.53E-09	0.00E+00	0.00E+00	2.94E-09	1.25E-11	2.96E-09
	78.6-120	0.00E+00	0.00E+00	5.53E-09	0.00E+00	0.00E+00	2.94E-09	1.25E-11	2.96E-09
SL3	0-19.2	0.00E+00	0.00E+00	2.77E-09	0.00E+00	0.00E+00	3.13E-08	8.96E-11	3.13E-08
CC1	0-19.2	0.00E+00	0.00E+00	6.45E-09	0.00E+00	0.00E+00	3.90E-08	1.10E-10	3.91E-08
CC2	0-19.2	0.00E+00	0.00E+00	6.45E-09	0.00E+00	0.00E+00	2.19E-08	6.26E-11	2.20E-08
CC3	0-19.2	0.00E+00	0.00E+00	7.37E-09	0.00E+00	0.00E+00	4.38E-08	1.21E-10	4.39E-08
CC4	0-19.2	0.00E+00	0.00E+00	3.69E-09	0.00E+00	0.00E+00	1.02E-08	3.17E-11	1.02E-08

Predicted Maximum Permitted Concentration of Certain Metals in Foods - Produce - The IWMF at the artificial island near SKC

ASR	Assessment Height	Antimony	Arsenic	Cadmium	Chromium	Lead	Mercuric chloride	Methyl mercury	Mercury
Maximum Permitted Concentration (ppm)		1	1.4	0.1	1	6			0.5
SKC1	0-19.2	1.93E-06	1.04E-05	1.05E-06	1.11E-05	1.05E-05	1.39E-03	6.29E-05	1.45E-03
SL1	0-19.2	6.10E-08	2.09E-04	2.08E-05	2.09E-04	2.09E-04	7.56E-05	5.67E-06	8.13E-05
SL2	0-19.2	8.09E-08	3.13E-04	3.12E-05	3.13E-04	3.13E-04	8.99E-05	6.62E-06	9.65E-05
SLW1	0-19.2	1.79E-08	2.40E-04	2.39E-05	2.40E-04	2.40E-04	4.45E-05	6.41E-06	5.09E-05
SLW2	0-19.2	1.79E-08	2.92E-04	2.91E-05	2.92E-04	2.92E-04	4.70E-05	6.95E-06	5.40E-05
SLW3	0-19.2	1.58E-08	1.98E-04	1.98E-05	1.98E-04	1.98E-04	4.60E-05	6.84E-06	5.28E-05
TC1	0-19.2	1.26E-08	1.57E-04	1.56E-05	1.57E-04	1.56E-04	3.44E-05	5.54E-06	3.99E-05
	19.2-41	1.26E-08	1.57E-04	1.56E-05	1.57E-04	1.56E-04	3.49E-05	5.68E-06	4.06E-05
	41-78.6	1.26E-08	1.57E-04	1.56E-05	1.57E-04	1.56E-04	3.56E-05	5.89E-06	4.15E-05
	78.6-120	1.26E-08	1.57E-04	1.56E-05	1.57E-04	1.56E-04	3.75E-05	6.40E-06	4.39E-05
	120-175	1.26E-08	1.57E-04	1.56E-05	1.57E-04	1.56E-04	4.07E-05	7.33E-06	4.81E-05
TC2	0-19.2	1.26E-08	1.88E-04	1.87E-05	1.88E-04	1.88E-04	3.44E-05	5.54E-06	3.99E-05
	19.2-41	1.26E-08	1.88E-04	1.87E-05	1.88E-04	1.88E-04	3.49E-05	5.68E-06	4.06E-05
	41-78.6	1.26E-08	1.88E-04	1.87E-05	1.88E-04	1.88E-04	3.56E-05	5.89E-06	4.15E-05
	78.6-120	1.26E-08	1.88E-04	1.87E-05	1.88E-04	1.88E-04	3.75E-05	6.40E-06	4.39E-05
	120-175	1.26E-08	1.88E-04	1.87E-05	1.88E-04	1.88E-04	4.07E-05	7.33E-06	4.81E-05
TC3	0-19.2	1.16E-08	1.98E-04	1.98E-05	1.98E-04	1.98E-04	3.46E-05	5.60E-06	4.02E-05
	19.2-41	1.16E-08	1.98E-04	1.98E-05	1.98E-04	1.98E-04	3.50E-05	5.72E-06	4.08E-05
	0-19.2	1.16E-08	2.09E-04	2.08E-05	2.09E-04	2.09E-04	3.54E-05	5.64E-06	4.10E-05
TC4	0-19.2	1.16E-08	2.09E-04	2.08E-05	2.09E-04	2.09E-04	3.58E-05	5.76E-06	4.16E-05
	19.2-41	1.16E-08	2.09E-04	2.08E-05	2.09E-04	2.09E-04	3.73E-05	6.18E-06	4.35E-05
	41-78.6	1.16E-08	2.09E-04	2.08E-05	2.09E-04	2.09E-04	4.15E-05	7.37E-06	4.89E-05
	78.6-120	1.16E-08	2.09E-04	2.08E-05	2.09E-04	2.09E-04	4.70E-05	8.93E-06	5.60E-05
	120-175	1.16E-08	2.09E-04	2.08E-05	2.09E-04	2.09E-04	4.07E-05	6.44E-06	4.72E-05
TC5	0-19.2	1.47E-08	2.19E-04	2.19E-05	2.19E-04	2.19E-04	4.07E-05	6.44E-06	4.72E-05
	19.2-41	1.47E-08	2.19E-04	2.19E-05	2.19E-04	2.19E-04	4.09E-05	6.48E-06	4.74E-05
	41-78.6	1.47E-08	2.19E-04	2.19E-05	2.19E-04	2.19E-04	4.17E-05	6.71E-06	4.84E-05
	78.6-120	1.47E-08	2.19E-04	2.19E-05	2.19E-04	2.19E-04	4.42E-05	7.41E-06	5.16E-05
	120-175	1.47E-08	2.19E-04	2.19E-05	2.19E-04	2.19E-04	4.73E-05	8.29E-06	5.56E-05
TC6	0-19.2	2.10E-08	1.67E-04	1.67E-05	1.67E-04	1.67E-04	4.68E-05	6.71E-06	5.35E-05
	19.2-41	2.10E-08	1.67E-04	1.67E-05	1.67E-04	1.67E-04	4.70E-05	6.76E-06	5.37E-05
	41-78.6	2.10E-08	1.67E-04	1.67E-05	1.67E-04	1.67E-04	4.78E-05	6.99E-06	5.48E-05
	78.6-120	2.10E-08	1.67E-04	1.67E-05	1.67E-04	1.67E-04	4.95E-05	7.65E-06	5.72E-05
	120-175	2.00E-08	1.67E-04	1.67E-05	1.67E-04	1.67E-04	5.26E-05	8.53E-06	6.12E-05
AP1	0-19.2	1.16E-08	2.30E-04	2.29E-05	2.30E-04	2.30E-04	3.73E-05	6.18E-06	4.35E-05
AP2	0-19.2	1.05E-08	2.30E-04	2.29E-05	2.30E-04	2.30E-04	3.58E-05	6.11E-06	4.19E-05
AP3	0-19.2	1.47E-08	1.98E-04	1.98E-05	1.98E-04	1.98E-04	4.33E-05	6.45E-06	4.98E-05
	0-19.2	8.41E-09	2.40E-04	2.39E-05	2.40E-04	2.40E-04	2.89E-05	4.89E-06	3.38E-05
	19.2-41	8.41E-09	2.40E-04	2.39E-05	2.40E-04	2.40E-04	2.89E-05	4.89E-06	3.38E-05
AP4	41-78.6	8.41E-09	2.40E-04	2.39E-05	2.40E-04	2.40E-04	2.91E-05	4.94E-06	3.40E-05
	78.6-120	8.41E-09	2.40E-04	2.39E-05	2.40E-04	2.40E-04	2.93E-05	5.01E-06	3.43E-05
	0-19.2	8.41E-09	2.09E-04	2.08E-05	2.09E-04	2.09E-04	2.97E-05	4.92E-06	3.46E-05
AP6	0-19.2	8.41E-09	2.09E-04	2.08E-05	2.09E-04	2.09E-04	3.18E-05	5.54E-06	3.74E-05
	0-19.2	8.41E-09	2.09E-04	2.08E-05	2.09E-04	2.09E-04	2.71E-05	4.39E-06	3.15E-05
	19.2-41	8.41E-09	2.09E-04	2.08E-05	2.09E-04	2.09E-04	2.73E-05	4.44E-06	3.17E-05
AP7	41-78.6	8.41E-09	2.09E-04	2.08E-05	2.09E-04	2.09E-04	2.75E-05	4.49E-06	3.20E-05
	0-19.2	8.41E-09	2.30E-04	2.29E-05	2.30E-04	2.30E-04	3.11E-05	5.51E-06	3.66E-05
	0-19.2	6.31E-09	1.46E-04	1.46E-05	1.46E-04	1.46E-04	1.63E-05	1.88E-06	1.82E-05
KT1	19.2-41	6.31E-09	1.46E-04	1.46E-05	1.46E-04	1.46E-04	1.63E-05	1.88E-06	1.82E-05
	41-78.6	6.31E-09	1.46E-04	1.46E-05	1.46E-04	1.46E-04	1.63E-05	1.88E-06	1.82E-05
	78.6-120	6.31E-09	1.46E-04	1.46E-05	1.46E-04	1.46E-04	1.64E-05	1.90E-06	1.83E-05
	0-19.2	6.31E-09	1.46E-04	1.46E-05	1.46E-04	1.46E-04	1.63E-05	1.88E-06	1.82E-05
KT2	19.2-41	6.31E-09	1.46E-04	1.46E-05	1.46E-04	1.46E-04	1.63E-05	1.88E-06	1.82E-05
	41-78.6	6.31E-09	1.46E-04	1.46E-05	1.46E-04	1.46E-04	1.63E-05	1.88E-06	1.82E-05
	78.6-120	6.31E-09	1.46E-04	1.46E-05	1.46E-04	1.46E-04	1.64E-05	1.90E-06	1.83E-05
	0-19.2	6.31E-09	1.46E-04	1.46E-05	1.46E-04	1.46E-04	1.71E-05	1.91E-06	1.90E-05
KT3	19.2-41	6.31E-09	1.46E-04	1.46E-05	1.46E-04	1.46E-04	1.71E-05	1.91E-06	1.90E-05
	0-19.2	6.31E-09	1.46E-04	1.46E-05	1.46E-04	1.46E-04	1.63E-05	1.88E-06	1.82E-05
KT4	0-19.2	4.20E-09	1.04E-04	1.04E-05	1.04E-04	1.04E-04	1.31E-05	1.72E-06	1.48E-05
	19.2-41	4.20E-09	1.04E-04	1.04E-05	1.04E-04	1.04E-04	1.31E-05	1.72E-06	1.48E-05
	41-78.6	4.20E-09	1.04E-04	1.04E-05	1.04E-04	1.04E-04	1.31E-05	1.72E-06	1.48E-05
	78.6-120	4.20E-09	1.04E-04	1.04E-05	1.04E-04	1.04E-04	1.32E-05	1.74E-06	1.49E-05
	120-175	4.20E-09	1.04E-04	1.04E-05	1.04E-04	1.04E-04	1.32E-05	1.75E-06	1.50E-05
KT6	0-19.2	5.26E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	1.56E-05	1.85E-06	1.74E-05
	19.2-41	5.26E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	1.56E-05	1.85E-06	1.74E-05
	0-19.2	3.15E-09	7.31E-05	7.29E-06	7.31E-05	7.30E-05	1.10E-05	1.66E-06	1.27E-05
KT7	0-19.2	3.15E-09	6.26E-05	6.25E-06	6.27E-05	6.26E-05	1.02E-05	1.63E-06	1.19E-05
	19.2-41	3.15E-09	6.26E-05	6.25E-06	6.27E-05	6.26E-05	1.02E-05	1.63E-06	1.19E-05
	41-78.6	3.15E-09	6.26E-05	6.25E-06	6.27E-05	6.26E-05	1.02E-05	1.63E-06	1.19E-05
	78.6-120	3.15E-09	6.26E-05	6.25E-06	6.27E-05	6.26E-05	1.02E-05	1.63E-06	1.19E-05
	0-19.2	7.15E-08	3.13E-05	3.12E-06	3.14E-05	3.13E-05	6.47E-05	4.44E-06	6.92E-05
CC1	0-19.2	7.99E-08	7.31E-05	7.29E-06	7.31E-05	7.30E-05	7.82E-05	4.79E-06	8.30E-05
CC2	0-19.2	3.57E-08	7.31E-05	7.29E-06	7.31E-05	7.30E-05	4.51E-05	3.04E-06	4.82E-05
CC3	0-19.2	9.04E-08	8.35E-05	8.33E-06	8.36E-05	8.35E-05	8.62E-05	4.91E-06	9.11E-05
CC4	0-19.2	1.37E-08	4.17E-05	4.16E-06	4.18E-05	4.17E-05	2.37E-05	2.18E-06	2.58E-05

Predicted Maximum Permitted Concentration of Certain Metals in Foods - Chicken - Co-exist Scenario

ASR	Assessment Height (mAG)	Antimony	Arsenic	Cadmium	Chromium	Lead	Mercuric chloride	Methyl mercury	Mercury
Maximum Permitted Concentration (ppm)		1	1.4	0.2	1	6			0.5
TT1	0-19.2	0.00E+00	0.00E+00	1.24E-10	0.00E+00	0.00E+00	4.01E-06	1.07E-08	4.02E-06
TT2	0-19.2	0.00E+00	0.00E+00	8.87E-11	0.00E+00	0.00E+00	1.45E-05	3.89E-08	1.46E-05
TT3	0-19.2	0.00E+00	0.00E+00	1.40E-10	0.00E+00	0.00E+00	2.24E-05	6.00E-08	2.25E-05
TT4	0-19.2	0.00E+00	0.00E+00	5.58E-10	0.00E+00	0.00E+00	1.26E-04	3.38E-07	1.27E-04
TT5	0-19.2	0.00E+00	0.00E+00	2.42E-10	0.00E+00	0.00E+00	6.82E-06	1.83E-08	6.84E-06
TT6	0-19.2	0.00E+00	0.00E+00	2.11E-10	0.00E+00	0.00E+00	2.18E-05	5.85E-08	2.19E-05
	0-19.2	0.00E+00	0.00E+00	1.11E-11	0.00E+00	0.00E+00	6.80E-07	1.82E-09	6.81E-07
	19.2-41	0.00E+00	0.00E+00	1.11E-11	0.00E+00	0.00E+00	6.80E-07	1.82E-09	6.81E-07
	41-78.6	0.00E+00	0.00E+00	1.11E-11	0.00E+00	0.00E+00	6.80E-07	1.82E-09	6.81E-07
	78.6-120	0.00E+00	0.00E+00	1.11E-11	0.00E+00	0.00E+00	6.80E-07	1.82E-09	6.81E-07
	0-19.2	0.00E+00	0.00E+00	2.67E-11	0.00E+00	0.00E+00	5.27E-07	1.41E-09	5.28E-07
	19.2-41	0.00E+00	0.00E+00	2.67E-11	0.00E+00	0.00E+00	5.27E-07	1.41E-09	5.28E-07
	41-78.6	0.00E+00	0.00E+00	2.67E-11	0.00E+00	0.00E+00	5.27E-07	1.41E-09	5.28E-07
	78.6-120	0.00E+00	0.00E+00	2.67E-11	0.00E+00	0.00E+00	5.27E-07	1.41E-09	5.28E-07
	0-19.2	0.00E+00	0.00E+00	1.97E-11	0.00E+00	0.00E+00	7.99E-07	2.14E-09	8.01E-07
	19.2-41	0.00E+00	0.00E+00	1.97E-11	0.00E+00	0.00E+00	7.99E-07	2.14E-09	8.01E-07
	41-78.6	0.00E+00	0.00E+00	1.97E-11	0.00E+00	0.00E+00	7.99E-07	2.14E-09	8.01E-07
	78.6-120	0.00E+00	0.00E+00	1.97E-11	0.00E+00	0.00E+00	7.99E-07	2.14E-09	8.01E-07
	0-19.2	0.00E+00	0.00E+00	2.82E-11	0.00E+00	0.00E+00	8.50E-07	2.28E-09	8.52E-07
	19.2-41	0.00E+00	0.00E+00	2.82E-11	0.00E+00	0.00E+00	8.50E-07	2.28E-09	8.52E-07
	41-78.6	0.00E+00	0.00E+00	2.82E-11	0.00E+00	0.00E+00	8.50E-07	2.28E-09	8.52E-07
	78.6-120	0.00E+00	0.00E+00	2.82E-11	0.00E+00	0.00E+00	8.50E-07	2.28E-09	8.52E-07
	0-19.2	0.00E+00	0.00E+00	2.04E-11	0.00E+00	0.00E+00	9.51E-07	2.55E-09	9.54E-07
	19.2-41	0.00E+00	0.00E+00	2.04E-11	0.00E+00	0.00E+00	9.51E-07	2.55E-09	9.54E-07
	41-78.6	0.00E+00	0.00E+00	2.04E-11	0.00E+00	0.00E+00	9.51E-07	2.55E-09	9.54E-07
	78.6-120	0.00E+00	0.00E+00	2.04E-11	0.00E+00	0.00E+00	9.51E-07	2.55E-09	9.54E-07
	0-19.2	0.00E+00	0.00E+00	2.16E-11	0.00E+00	0.00E+00	1.19E-06	3.19E-09	1.19E-06
	19.2-41	0.00E+00	0.00E+00	2.16E-11	0.00E+00	0.00E+00	1.19E-06	3.19E-09	1.19E-06
	41-78.6	0.00E+00	0.00E+00	2.16E-11	0.00E+00	0.00E+00	1.19E-06	3.19E-09	1.19E-06
	78.6-120	0.00E+00	0.00E+00	2.16E-11	0.00E+00	0.00E+00	1.19E-06	3.19E-09	1.19E-06
SLW1	0-19.2	0.00E+00	0.00E+00	5.68E-10	0.00E+00	0.00E+00	2.12E-06	5.68E-09	2.12E-06
SLW2	0-19.2	0.00E+00	0.00E+00	6.34E-10	0.00E+00	0.00E+00	2.31E-06	6.18E-09	2.31E-06
SLW3	0-19.2	0.00E+00	0.00E+00	3.00E-10	0.00E+00	0.00E+00	9.40E-07	2.52E-09	9.43E-07
	0-19.2	0.00E+00	0.00E+00	2.25E-10	0.00E+00	0.00E+00	5.98E-07	1.60E-09	5.99E-07
	19.2-41	0.00E+00	0.00E+00	2.25E-10	0.00E+00	0.00E+00	5.98E-07	1.60E-09	5.99E-07
	41-78.6	0.00E+00	0.00E+00	2.25E-10	0.00E+00	0.00E+00	5.98E-07	1.60E-09	5.99E-07
	78.6-120	0.00E+00	0.00E+00	2.25E-10	0.00E+00	0.00E+00	5.98E-07	1.60E-09	5.99E-07
	120-175	0.00E+00	0.00E+00	2.25E-10	0.00E+00	0.00E+00	5.98E-07	1.60E-09	5.99E-07
	0-19.2	0.00E+00	0.00E+00	2.41E-10	0.00E+00	0.00E+00	5.30E-07	1.42E-09	5.31E-07
	19.2-41	0.00E+00	0.00E+00	2.41E-10	0.00E+00	0.00E+00	5.30E-07	1.42E-09	5.31E-07
	41-78.6	0.00E+00	0.00E+00	2.41E-10	0.00E+00	0.00E+00	5.30E-07	1.42E-09	5.31E-07
	78.6-120	0.00E+00	0.00E+00	2.41E-10	0.00E+00	0.00E+00	5.30E-07	1.42E-09	5.31E-07
	120-175	0.00E+00	0.00E+00	2.41E-10	0.00E+00	0.00E+00	5.30E-07	1.42E-09	5.31E-07
	0-19.2	0.00E+00	0.00E+00	2.41E-10	0.00E+00	0.00E+00	5.13E-07	1.37E-09	5.14E-07
	19.2-41	0.00E+00	0.00E+00	2.41E-10	0.00E+00	0.00E+00	5.13E-07	1.37E-09	5.14E-07
	0-19.2	0.00E+00	0.00E+00	2.57E-10	0.00E+00	0.00E+00	5.30E-07	1.42E-09	5.31E-07
	19.2-41	0.00E+00	0.00E+00	2.57E-10	0.00E+00	0.00E+00	5.30E-07	1.42E-09	5.31E-07
	41-78.6	0.00E+00	0.00E+00	2.57E-10	0.00E+00	0.00E+00	5.30E-07	1.42E-09	5.31E-07
	78.6-120	0.00E+00	0.00E+00	2.57E-10	0.00E+00	0.00E+00	5.30E-07	1.42E-09	5.31E-07
	120-175	0.00E+00	0.00E+00	2.57E-10	0.00E+00	0.00E+00	5.30E-07	1.42E-09	5.31E-07
	0-19.2	0.00E+00	0.00E+00	2.74E-10	0.00E+00	0.00E+00	6.15E-07	1.65E-09	6.16E-07
	19.2-41	0.00E+00	0.00E+00	2.74E-10	0.00E+00	0.00E+00	6.15E-07	1.65E-09	6.16E-07
	41-78.6	0.00E+00	0.00E+00	2.74E-10	0.00E+00	0.00E+00	6.15E-07	1.65E-09	6.16E-07
	78.6-120	0.00E+00	0.00E+00	2.74E-10	0.00E+00	0.00E+00	6.15E-07	1.65E-09	6.16E-07
	120-175	0.00E+00	0.00E+00	2.74E-10	0.00E+00	0.00E+00	6.15E-07	1.65E-09	6.16E-07
	0-19.2	0.00E+00	0.00E+00	2.34E-10	0.00E+00	0.00E+00	7.51E-07	2.01E-09	7.53E-07
	19.2-41	0.00E+00	0.00E+00	2.34E-10	0.00E+00	0.00E+00	7.51E-07	2.01E-09	7.53E-07
	41-78.6	0.00E+00	0.00E+00	2.34E-10	0.00E+00	0.00E+00	7.51E-07	2.01E-09	7.53E-07
	78.6-120	0.00E+00	0.00E+00	2.34E-10	0.00E+00	0.00E+00	7.51E-07	2.01E-09	7.53E-07
	120-175	0.00E+00	0.00E+00	2.34E-10	0.00E+00	0.00E+00	7.51E-07	2.01E-09	7.53E-07
AP1	0-19.2	0.00E+00	0.00E+00	2.91E-10	0.00E+00	0.00E+00	6.15E-07	1.65E-09	6.16E-07
AP2	0-19.2	0.00E+00	0.00E+00	2.91E-10	0.00E+00	0.00E+00	6.32E-07	1.69E-09	6.33E-07
AP3	0-19.2	0.00E+00	0.00E+00	3.17E-10	0.00E+00	0.00E+00	1.14E-06	3.07E-09	1.15E-06
	0-19.2	0.00E+00	0.00E+00	2.66E-10	0.00E+00	0.00E+00	5.30E-07	1.42E-09	5.31E-07
	19.2-41	0.00E+00	0.00E+00	2.66E-10	0.00E+00	0.00E+00	5.30E-07	1.42E-09	5.31E-07
	41-78.6	0.00E+00	0.00E+00	2.66E-10	0.00E+00	0.00E+00	5.30E-07	1.42E-09	5.31E-07
	78.6-120	0.00E+00	0.00E+00	2.66E-10	0.00E+00	0.00E+00	5.30E-07	1.42E-09	5.31E-07
AP5	0-19.2	0.00E+00	0.00E+00	2.41E-10	0.00E+00	0.00E+00	5.13E-07	1.37E-09	5.14E-07
AP6	0-19.2	0.00E+00	0.00E+00	2.41E-10	0.00E+00	0.00E+00	5.13E-07	1.37E-09	5.14E-07
AP7	0-19.2	0.00E+00	0.00E+00	2.49E-10	0.00E+00	0.00E+00	5.64E-07	1.51E-09	5.65E-07
	19.2-41	0.00E+00	0.00E+00	2.49E-10	0.00E+00	0.00E+00	5.64E-07	1.51E-09	5.65E-07
	41-78.6	0.00E+00	0.00E+00	2.49E-10	0.00E+00	0.00E+00	5.64E-07	1.51E-09	5.65E-07
AP8	0-19.2	0.00E+00	0.00E+00	2.57E-10	0.00E+00	0.00E+00	4.96E-07	1.33E-09	4.97E-07
	0-19.2	0.00E+00	0.00E+00	1.58E-10	0.00E+00	0.00E+00	2.92E-07	7.82E-10	2.93E-07
	19.2-41	0.00E+00	0.00E+00	1.58E-10	0.00E+00	0.00E+00	2.92E-07	7.82E-10	2.93E-07
	41-78.6	0.00E+00	0.00E+00	1.58E-10	0.00E+00	0.00E+00	2.92E-07	7.82E-10	2.93E-07
	78.6-120	0.00E+00	0.00E+00	1.58E-10	0.00E+00	0.00E+00	2.92E-07	7.82E-10	2.93E-07
	0-19.2	0.00E+00	0.00E+00	1.49E-10	0.00E+00	0.00E+00	2.92E-07	7.82E-10	2.93E-07
	19.2-41	0.00E+00	0.00E+00	1.49E-10	0.00E+00	0.00E+00	2.92E-07	7.82E-10	2.93E-07
	41-78.6	0.00E+00	0.00E+00	1.49E-10	0.00E+00	0.00E+00	2.92E-07	7.82E-10	2.93E-07
	78.6-120	0.00E+00	0.00E+00	1.49E-10	0.00E+00	0.00E+00	2.92E-07	7.82E-10	2.93E-07
	0-19.2	0.00E+00	0.00E+00	1.49E-10	0.00E+00	0.00E+00	3.09E-07	8.27E-10	3.10E-07
	19.2-41	0.00E+00	0.00E+00	1.49E-10	0.00E+00	0.00E+00	3.09E-07	8.27E-10	3.10E-07
KT4	0-19.2	0.00E+00	0.00E+00	1.49E-10	0.00E+00	0.00E+00	2.92E-07	7.82E-10	2.93E-07
	0-19.2	0.00E+00	0.00E+00	9.96E-11	0.00E+00	0.00E+00	2.21E-07	5.92E-10	2.21E-07
	19.2-41	0.00E+00	0.00E+00	9.96E-11	0.00E+00	0.00E+00	2.21E-07	5.92E-10	2.21E-07
	41-78.6	0.00E+00	0.00E+00	9.96E-11	0.00E+00	0.00E+00	2.21E-07	5.92E-10	2.21E-07
	78.6-120	0.00E+00	0.00E+00	9.96E-11	0.00E+00	0.00E+00	2.21E-07	5.92E-10	2.21E-07
	120-175	0.00E+00	0.00E+00	9.96E-11	0.00E+00	0.00E+00	2.21E-07	5.92E-10	2.21E-07
KT6	0-19.2	0.00E+00	0.00E+00	1.25E-10	0.00E+00	0.00E+00	2.75E-07	7.36E-10	2.76E-07
	19.2-41	0.00E+00	0.00E+00	1.25E-10	0.00E+00	0.00E+00	2.75E-07	7.36E-10	2.76E-07
KT7	0-19.2	0.00E+00	0.00E+00	7.47E-11	0.00E+00	0.00E+00	1.70E-07	4.55E-10	1.70E-07
	0-19.2	0.00E+00	0.00E+00	6.64E-11	0.00E+00	0.00E+00	1.36E-07	3.64E-10	1.36E-07
	19.2-41	0.00E+00	0.00E+00	6.64E-11	0.00E+00	0.00E+00	1.36E-07	3.64E-10	1.36E-07
	41-78.6	0.00E+00	0.00E+00	6.64E-11	0.00E+00	0.00E+00	1.36E-07	3.64E-10	1.36E-07
	78.6-120	0.00E+00	0.00E+00	6.64E-11	0.00E+00	0.00E+00	1.36E-07	3.64E-10	1.36E-07
SKC1	0-19.2	0.00E+00	0.00E+00	1.46E-10	0.00E+0				

Predicted Maximum Permitted Concentration of Certain Metals in Foods - Pork - Co-exist Scenario

ASR	Assessment Height (mAG)	Antimony	Arsenic	Cadmium	Chromium	Lead	Mercuric chloride	Methyl mercury	Mercury
Maximum Permitted Concentration (ppm)		1	1.4	0.2	1	6			0.5
TT1	0-19.2	0.00E+00	0.00E+00	1.20E-08	0.00E+00	0.00E+00	9.56E-08	2.57E-10	9.59E-08
TT2	0-19.2	0.00E+00	0.00E+00	2.77E-09	0.00E+00	0.00E+00	3.46E-07	9.26E-10	3.47E-07
TT3	0-19.2	0.00E+00	0.00E+00	4.61E-09	0.00E+00	0.00E+00	5.34E-07	1.43E-09	5.35E-07
TT4	0-19.2	0.00E+00	0.00E+00	3.17E-11	0.00E+00	0.00E+00	3.01E-06	8.04E-09	3.01E-06
TT5	0-19.2	0.00E+00	0.00E+00	2.40E-08	0.00E+00	0.00E+00	1.63E-07	4.39E-10	1.63E-07
TT6	0-19.2	0.00E+00	0.00E+00	1.29E-08	0.00E+00	0.00E+00	5.20E-07	1.39E-09	5.21E-07
	0-19.2	0.00E+00	0.00E+00	9.22E-10	0.00E+00	0.00E+00	1.63E-08	4.71E-11	1.63E-08
	19.2-41	0.00E+00	0.00E+00	9.22E-10	0.00E+00	0.00E+00	1.63E-08	4.72E-11	1.63E-08
	41-78.6	0.00E+00	0.00E+00	9.22E-10	0.00E+00	0.00E+00	1.63E-08	4.75E-11	1.63E-08
	78.6-120	0.00E+00	0.00E+00	9.22E-10	0.00E+00	0.00E+00	1.63E-08	4.80E-11	1.63E-08
TM1	0-19.2	0.00E+00	0.00E+00	2.77E-09	0.00E+00	0.00E+00	1.26E-08	3.73E-11	1.27E-08
	19.2-41	0.00E+00	0.00E+00	2.77E-09	0.00E+00	0.00E+00	1.26E-08	3.75E-11	1.27E-08
	41-78.6	0.00E+00	0.00E+00	2.77E-09	0.00E+00	0.00E+00	1.26E-08	3.77E-11	1.27E-08
	78.6-120	0.00E+00	0.00E+00	2.77E-09	0.00E+00	0.00E+00	1.27E-08	3.83E-11	1.27E-08
TM2	0-19.2	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	1.91E-08	5.47E-11	1.92E-08
	19.2-41	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	1.91E-08	5.48E-11	1.92E-08
	41-78.6	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	1.91E-08	5.51E-11	1.92E-08
	78.6-120	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	1.91E-08	5.56E-11	1.92E-08
TM3	0-19.2	0.00E+00	0.00E+00	2.77E-09	0.00E+00	0.00E+00	2.03E-08	5.79E-11	2.04E-08
	19.2-41	0.00E+00	0.00E+00	2.77E-09	0.00E+00	0.00E+00	2.03E-08	5.80E-11	2.04E-08
	41-78.6	0.00E+00	0.00E+00	2.77E-09	0.00E+00	0.00E+00	2.03E-08	5.83E-11	2.04E-08
	78.6-120	0.00E+00	0.00E+00	2.77E-09	0.00E+00	0.00E+00	2.03E-08	5.88E-11	2.04E-08
TM4	0-19.2	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	2.27E-08	6.39E-11	2.28E-08
	19.2-41	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	2.27E-08	6.40E-11	2.28E-08
	41-78.6	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	2.28E-08	6.44E-11	2.28E-08
	78.6-120	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	2.28E-08	6.55E-11	2.28E-08
TM5	0-19.2	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	2.84E-08	7.98E-11	2.85E-08
	19.2-41	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	2.84E-08	8.00E-11	2.85E-08
	41-78.6	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	2.84E-08	8.04E-11	2.85E-08
	78.6-120	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	2.85E-08	8.14E-11	2.85E-08
TM6	0-19.2	0.00E+00	0.00E+00	2.77E-09	0.00E+00	0.00E+00	5.12E-08	1.65E-10	5.13E-08
SLW1	0-19.2	0.00E+00	0.00E+00	7.00E-08	0.00E+00	0.00E+00	5.57E-08	1.79E-10	5.58E-08
SLW2	0-19.2	0.00E+00	0.00E+00	3.32E-08	0.00E+00	0.00E+00	2.31E-09	9.00E-11	2.32E-08
SLW3	0-19.2	0.00E+00	0.00E+00	2.49E-08	0.00E+00	0.00E+00	1.48E-08	6.19E-11	1.49E-08
	19.2-41	0.00E+00	0.00E+00	2.49E-08	0.00E+00	0.00E+00	1.48E-08	6.24E-11	1.49E-08
	41-78.6	0.00E+00	0.00E+00	2.49E-08	0.00E+00	0.00E+00	1.48E-08	6.33E-11	1.49E-08
	78.6-120	0.00E+00	0.00E+00	2.49E-08	0.00E+00	0.00E+00	1.49E-08	6.51E-11	1.49E-08
TC1	120-175	0.00E+00	0.00E+00	2.49E-08	0.00E+00	0.00E+00	1.50E-08	6.84E-11	1.50E-08
	0-19.2	0.00E+00	0.00E+00	2.67E-08	0.00E+00	0.00E+00	1.32E-08	5.76E-11	1.32E-08
	19.2-41	0.00E+00	0.00E+00	2.67E-08	0.00E+00	0.00E+00	1.32E-08	5.81E-11	1.33E-08
	41-78.6	0.00E+00	0.00E+00	2.67E-08	0.00E+00	0.00E+00	1.32E-08	5.90E-11	1.33E-08
	78.6-120	0.00E+00	0.00E+00	2.67E-08	0.00E+00	0.00E+00	1.33E-08	6.08E-11	1.33E-08
TC2	120-175	0.00E+00	0.00E+00	2.67E-08	0.00E+00	0.00E+00	1.33E-08	6.41E-11	1.34E-08
	0-19.2	0.00E+00	0.00E+00	2.67E-08	0.00E+00	0.00E+00	1.28E-08	5.63E-11	1.28E-08
	19.2-41	0.00E+00	0.00E+00	2.67E-08	0.00E+00	0.00E+00	1.28E-08	5.67E-11	1.28E-08
	41-78.6	0.00E+00	0.00E+00	2.67E-08	0.00E+00	0.00E+00	1.28E-08	5.74E-11	1.28E-08
	78.6-120	0.00E+00	0.00E+00	2.67E-08	0.00E+00	0.00E+00	1.28E-08	5.78E-11	1.28E-08
TC3	120-175	0.00E+00	0.00E+00	2.67E-08	0.00E+00	0.00E+00	1.32E-08	5.96E-11	1.33E-08
	0-19.2	0.00E+00	0.00E+00	2.86E-08	0.00E+00	0.00E+00	1.33E-08	6.42E-11	1.34E-08
	19.2-41	0.00E+00	0.00E+00	2.86E-08	0.00E+00	0.00E+00	1.33E-08	6.42E-11	1.34E-08
	41-78.6	0.00E+00	0.00E+00	2.86E-08	0.00E+00	0.00E+00	1.33E-08	6.42E-11	1.34E-08
	78.6-120	0.00E+00	0.00E+00	2.86E-08	0.00E+00	0.00E+00	1.33E-08	6.42E-11	1.34E-08
TC4	120-175	0.00E+00	0.00E+00	2.86E-08	0.00E+00	0.00E+00	1.35E-08	7.03E-11	1.36E-08
	0-19.2	0.00E+00	0.00E+00	3.04E-08	0.00E+00	0.00E+00	1.53E-08	6.60E-11	1.53E-08
	19.2-41	0.00E+00	0.00E+00	3.04E-08	0.00E+00	0.00E+00	1.53E-08	6.63E-11	1.53E-08
	41-78.6	0.00E+00	0.00E+00	3.04E-08	0.00E+00	0.00E+00	1.53E-08	6.75E-11	1.54E-08
	78.6-120	0.00E+00	0.00E+00	3.04E-08	0.00E+00	0.00E+00	1.54E-08	7.05E-11	1.55E-08
TC5	120-175	0.00E+00	0.00E+00	3.04E-08	0.00E+00	0.00E+00	1.55E-08	7.44E-11	1.56E-08
	0-19.2	0.00E+00	0.00E+00	2.58E-08	0.00E+00	0.00E+00	1.85E-08	7.47E-11	1.86E-08
	19.2-41	0.00E+00	0.00E+00	2.58E-08	0.00E+00	0.00E+00	1.85E-08	7.49E-11	1.86E-08
	41-78.6	0.00E+00	0.00E+00	2.58E-08	0.00E+00	0.00E+00	1.85E-08	7.61E-11	1.86E-08
	78.6-120	0.00E+00	0.00E+00	2.58E-08	0.00E+00	0.00E+00	1.85E-08	7.81E-11	1.83E-08
TC6	120-175	0.00E+00	0.00E+00	2.58E-08	0.00E+00	0.00E+00	1.83E-08	8.20E-11	1.84E-08
AP1	0-19.2	0.00E+00	0.00E+00	3.23E-08	0.00E+00	0.00E+00	1.53E-08	6.58E-11	1.53E-08
AP2	0-19.2	0.00E+00	0.00E+00	3.23E-08	0.00E+00	0.00E+00	1.57E-08	6.68E-11	1.57E-08
AP3	0-19.2	0.00E+00	0.00E+00	3.50E-08	0.00E+00	0.00E+00	2.79E-08	1.02E-10	2.80E-08
	0-19.2	0.00E+00	0.00E+00	2.95E-08	0.00E+00	0.00E+00	1.32E-08	5.63E-11	1.32E-08
	19.2-41	0.00E+00	0.00E+00	2.95E-08	0.00E+00	0.00E+00	1.32E-08	5.63E-11	1.32E-08
	41-78.6	0.00E+00	0.00E+00	2.95E-08	0.00E+00	0.00E+00	1.32E-08	5.66E-11	1.32E-08
	78.6-120	0.00E+00	0.00E+00	2.95E-08	0.00E+00	0.00E+00	1.32E-08	5.69E-11	1.32E-08
AP4	120-175	0.00E+00	0.00E+00	2.67E-08	0.00E+00	0.00E+00	1.27E-08	5.52E-11	1.28E-08
AP5	0-19.2	0.00E+00	0.00E+00	2.67E-08	0.00E+00	0.00E+00	1.28E-08	5.77E-11	1.29E-08
AP6	0-19.2	0.00E+00	0.00E+00	2.77E-08	0.00E+00	0.00E+00	1.39E-08	5.65E-11	1.40E-08
	19.2-41	0.00E+00	0.00E+00	2.77E-08	0.00E+00	0.00E+00	1.39E-08	5.65E-11	1.40E-08
	41-78.6	0.00E+00	0.00E+00	2.77E-08	0.00E+00	0.00E+00	1.39E-08	5.67E-11	1.40E-08
	78.6-120	0.00E+00	0.00E+00	2.86E-08	0.00E+00	0.00E+00	1.24E-08	5.66E-11	1.25E-08
AP7	0-19.2	0.00E+00	0.00E+00	1.75E-08	0.00E+00	0.00E+00	7.13E-09	2.62E-11	7.16E-09
AP8	0-19.2	0.00E+00	0.00E+00	1.75E-08	0.00E+00	0.00E+00	7.13E-09	2.63E-11	7.16E-09
	19.2-41	0.00E+00	0.00E+00	1.75E-08	0.00E+00	0.00E+00	7.13E-09	2.63E-11	7.16E-09
	41-78.6	0.00E+00	0.00E+00	1.75E-08	0.00E+00	0.00E+00	7.13E-09	2.63E-11	7.16E-09
	78.6-120	0.00E+00	0.00E+00	1.75E-08	0.00E+00	0.00E+00	7.14E-09	2.64E-11	7.16E-09
KT1	0-19.2	0.00E+00	0.00E+00	1.66E-08	0.00E+00	0.00E+00	7.13E-09	2.62E-11	7.16E-09
	19.2-41	0.00E+00	0.00E+00	1.66E-08	0.00E+00	0.00E+00	7.13E-09	2.63E-11	7.16E-09
	41-78.6	0.00E+00	0.00E+00	1.66E-08	0.00E+00	0.00E+00	7.13E-09	2.63E-11	7.16E-09
	78.6-120	0.00E+00	0.00E+00	1.66E-08	0.00E+00	0.00E+00	7.13E-09	2.63E-11	7.16E-09
KT2	0-19.2	0.00E+00	0.00E+00	1.66E-08	0.00E+00	0.00E+00	7.14E-09	2.64E-11	7.16E-09
	19.2-41	0.00E+00	0.00E+00	1.66E-08	0.00E+00	0.00E+00	7.14E-09	2.64E-11	7.16E-09
	41-78.6	0.00E+00	0.00E+00	1.66E-08	0.00E+00	0.00E+00	7.14E-09	2.64E-11	7.16E-09
	78.6-120	0.00E+00	0.00E+00	1.66E-08	0.00E+00	0.00E+00	7.14E-09	2.64E-11	7.16E-09
KT3	0-19.2	0.00E+00	0.00E+00	1.66E-08	0.00E+00	0.00E+00	7.54E-09	2.73E-11	7.56E-09
	19.2-41	0.00E+00	0.00E+00	1.66E-08	0.00E+00	0.00E+00	7.54E-09	2.74E-11	7.56E-09
	41-78.6	0.00E+00	0.00E+00	1.66E-08	0.00E+00	0.00E+00	7.54E-09	2.74E-11	7.56E-09
	78.6-120	0.00E+00	0.00E+00	1.66E-08	0.00E+00	0.00E+00	7.54E-09	2.74E-11	7.56E-09
KT4	0-19.2	0.00E+00	0.00E+00	1.66E-08	0.00E+00	0.00E+00	7.13E-09	2.62E-11	7.16E-09
	19.2-41	0.00E+00	0.00E+00	1.11E-08	0.00E+00	0.00E+00	5.43E-09	2.13E-11	5.45E-09
	41-78.6	0.00E+00	0.00E+00	1.11E-08	0.00E+00	0.00E+00	5.43E-09	2.14E-11	5.46E-09
	78.6-120	0.							

Predicted Maximum Permitted Concentration of Certain Metals in Foods - Produce - Co-exist Scenario

ASR	Assessment Height (mAG)	Antimony	Arsenic	Cadmium	Chromium	Lead	Mercuric chloride	Methyl mercury	Mercury
Maximum Permitted Concentration (ppm)		1	1.4	0.1	1	6			0.5
TT1	0-19.2	2.30E-07	1.36E-04	1.35E-05	1.36E-04	1.36E-04	1.81E-04	8.50E-06	1.89E-04
TT2	0-19.2	8.94E-07	3.13E-05	3.13E-06	3.16E-05	3.13E-05	6.49E-04	2.94E-05	6.79E-04
TT3	0-19.2	1.38E-06	5.22E-05	5.21E-06	5.27E-05	5.22E-05	1.00E-03	4.54E-05	1.05E-03
TT4	0-19.2	7.81E-06	2.27E-10	2.88E-08	2.66E-06	3.99E-07	5.64E-03	2.55E-04	5.89E-03
TT5	0-19.2	3.77E-07	2.71E-04	2.71E-05	2.72E-04	2.71E-04	3.08E-04	1.47E-05	3.23E-04
TT6	0-19.2	1.33E-06	1.46E-04	1.46E-05	1.47E-04	1.46E-04	9.77E-04	4.45E-05	1.02E-03
	0-19.2	3.99E-08	1.04E-05	1.04E-06	1.05E-05	1.04E-05	3.41E-05	2.44E-06	3.66E-05
	19.2-41	3.99E-08	1.04E-05	1.04E-06	1.05E-05	1.04E-05	3.43E-05	2.47E-06	3.67E-05
	41-78.6	3.99E-08	1.04E-05	1.04E-06	1.05E-05	1.04E-05	3.45E-05	2.55E-06	3.71E-05
	78.6-120	3.99E-08	1.04E-05	1.04E-06	1.05E-05	1.04E-05	3.51E-05	2.70E-06	3.78E-05
	0-19.2	2.63E-08	3.13E-05	3.12E-06	3.13E-05	3.13E-05	2.73E-05	2.13E-06	2.94E-05
	19.2-41	2.63E-08	3.13E-05	3.12E-06	3.13E-05	3.13E-05	2.74E-05	2.16E-06	2.96E-05
	41-78.6	2.63E-08	3.13E-05	3.12E-06	3.13E-05	3.13E-05	2.77E-05	2.24E-06	2.99E-05
	78.6-120	2.63E-08	3.13E-05	3.12E-06	3.13E-05	3.13E-05	2.82E-05	2.39E-06	3.06E-05
	0-19.2	4.41E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	3.95E-05	2.68E-06	4.21E-05
	19.2-41	4.41E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	3.96E-05	2.71E-06	4.23E-05
	41-78.6	4.41E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	3.98E-05	2.79E-06	4.26E-05
	78.6-120	4.41E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	4.04E-05	2.94E-06	4.33E-05
	0-19.2	4.73E-08	3.13E-05	3.12E-06	3.13E-05	3.13E-05	4.17E-05	2.78E-06	4.45E-05
	19.2-41	4.73E-08	3.13E-05	3.12E-06	3.13E-05	3.13E-05	4.18E-05	2.81E-06	4.47E-05
	41-78.6	4.73E-08	3.13E-05	3.12E-06	3.13E-05	3.13E-05	4.21E-05	2.89E-06	4.50E-05
	78.6-120	4.73E-08	3.13E-05	3.12E-06	3.13E-05	3.13E-05	4.27E-05	3.05E-06	4.57E-05
	0-19.2	5.36E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	4.58E-05	2.85E-06	4.86E-05
	19.2-41	5.36E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	4.59E-05	2.88E-06	4.88E-05
	41-78.6	5.36E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	4.63E-05	2.99E-06	4.93E-05
	78.6-120	5.36E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	4.74E-05	3.30E-06	5.07E-05
	0-19.2	7.04E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	5.72E-05	3.55E-06	6.07E-05
	19.2-41	7.04E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	5.74E-05	3.59E-06	6.09E-05
	41-78.6	7.04E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	5.78E-05	3.72E-06	6.15E-05
	78.6-120	7.04E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	5.88E-05	3.99E-06	6.28E-05
SLW1	0-19.2	7.57E-08	7.10E-04	7.08E-05	7.10E-04	7.09E-04	1.23E-04	1.23E-05	1.36E-04
SLW2	0-19.2	8.09E-08	7.93E-04	7.91E-05	7.94E-04	7.93E-04	1.34E-04	1.33E-05	1.47E-04
SLW3	0-19.2	2.94E-08	3.76E-04	3.75E-05	3.75E-04	3.76E-04	7.14E-05	1.02E-05	8.16E-05
	0-19.2	2.00E-08	2.82E-04	2.81E-05	2.82E-04	2.82E-04	5.02E-05	7.83E-06	5.80E-05
	19.2-41	2.00E-08	2.82E-04	2.81E-05	2.82E-04	2.82E-04	5.07E-05	7.97E-06	5.87E-05
	41-78.6	2.00E-08	2.82E-04	2.81E-05	2.82E-04	2.82E-04	5.16E-05	8.22E-06	5.98E-05
	78.6-120	2.00E-08	2.82E-04	2.81E-05	2.82E-04	2.82E-04	5.34E-05	8.73E-06	6.21E-05
	120-175	2.00E-08	2.82E-04	2.81E-05	2.82E-04	2.82E-04	5.67E-05	9.66E-06	6.63E-05
	0-19.2	1.68E-08	3.03E-04	3.02E-05	3.03E-04	3.03E-04	4.72E-05	7.70E-06	5.49E-05
	19.2-41	1.68E-08	3.03E-04	3.02E-05	3.03E-04	3.03E-04	4.77E-05	7.84E-06	5.55E-05
	41-78.6	1.68E-08	3.03E-04	3.02E-05	3.03E-04	3.03E-04	4.85E-05	8.08E-06	5.66E-05
	78.6-120	1.68E-08	3.03E-04	3.02E-05	3.03E-04	3.03E-04	5.03E-05	8.59E-06	5.89E-05
	120-175	1.68E-08	3.03E-04	3.02E-05	3.03E-04	3.03E-04	5.36E-05	9.52E-06	6.32E-05
	0-19.2	1.47E-08	3.03E-04	3.02E-05	3.03E-04	3.03E-04	4.62E-05	7.60E-06	5.38E-05
	19.2-41	1.47E-08	3.03E-04	3.02E-05	3.03E-04	3.03E-04	4.66E-05	7.72E-06	5.43E-05
	0-19.2	1.58E-08	3.24E-04	3.23E-05	3.24E-04	3.23E-04	4.69E-05	7.63E-06	5.46E-05
	19.2-41	1.58E-08	3.24E-04	3.23E-05	3.24E-04	3.23E-04	4.74E-05	7.76E-06	5.51E-05
	41-78.6	1.58E-08	3.24E-04	3.23E-05	3.24E-04	3.23E-04	4.91E-05	8.25E-06	5.74E-05
	78.6-120	1.58E-08	3.24E-04	3.23E-05	3.24E-04	3.23E-04	5.37E-05	9.55E-06	6.33E-05
	120-175	1.58E-08	3.24E-04	3.23E-05	3.24E-04	3.23E-04	5.98E-05	1.13E-05	7.11E-05
	0-19.2	2.10E-08	3.44E-04	3.43E-05	3.45E-04	3.44E-04	5.40E-05	8.72E-06	6.27E-05
	19.2-41	2.10E-08	3.44E-04	3.43E-05	3.45E-04	3.44E-04	5.42E-05	8.78E-06	6.30E-05
	41-78.6	2.10E-08	3.44E-04	3.43E-05	3.45E-04	3.44E-04	5.54E-05	9.12E-06	6.45E-05
	78.6-120	2.10E-08	3.44E-04	3.43E-05	3.45E-04	3.44E-04	5.84E-05	9.97E-06	6.84E-05
	120-175	2.10E-08	3.44E-04	3.43E-05	3.45E-04	3.44E-04	6.24E-05	1.11E-05	7.35E-05
	0-19.2	2.73E-08	2.92E-04	2.91E-05	2.92E-04	2.92E-04	6.00E-05	8.99E-06	6.90E-05
	19.2-41	2.73E-08	2.92E-04	2.91E-05	2.92E-04	2.92E-04	6.03E-05	9.06E-06	6.93E-05
	41-78.6	2.73E-08	2.92E-04	2.91E-05	2.92E-04	2.92E-04	6.15E-05	9.40E-06	7.09E-05
	78.6-120	2.73E-08	2.92E-04	2.91E-05	2.92E-04	2.92E-04	6.37E-05	1.02E-05	7.39E-05
	120-175	2.63E-08	2.92E-04	2.91E-05	2.92E-04	2.92E-04	6.77E-05	1.13E-05	7.90E-05
AP1	0-19.2	2.10E-08	3.65E-04	3.64E-05	3.66E-04	3.65E-04	5.37E-05	8.64E-06	6.23E-05
AP2	0-19.2	2.31E-08	3.65E-04	3.64E-05	3.66E-04	3.65E-04	5.45E-05	8.68E-06	6.31E-05
AP3	0-19.2	3.68E-08	3.97E-04	3.96E-05	3.97E-04	3.96E-04	7.92E-05	1.02E-05	8.95E-05
	0-19.2	1.89E-08	3.34E-04	3.33E-05	3.34E-04	3.34E-04	4.58E-05	7.33E-06	5.32E-05
	19.2-41	1.89E-08	3.34E-04	3.33E-05	3.34E-04	3.34E-04	4.58E-05	7.33E-06	5.32E-05
	41-78.6	1.89E-08	3.34E-04	3.33E-05	3.34E-04	3.34E-04	4.61E-05	7.40E-06	5.35E-05
	78.6-120	1.89E-08	3.34E-04	3.33E-05	3.34E-04	3.34E-04	4.65E-05	7.51E-06	5.40E-05
AP5	0-19.2	1.58E-08	3.03E-04	3.02E-05	3.03E-04	3.03E-04	4.51E-05	7.29E-06	5.24E-05
AP6	0-19.2	1.47E-08	3.03E-04	3.02E-05	3.03E-04	3.03E-04	4.76E-05	7.99E-06	5.55E-05
	0-19.2	1.89E-08	3.13E-04	3.12E-05	3.13E-04	3.13E-04	4.54E-05	6.84E-06	5.22E-05
	19.2-41	1.89E-08	3.13E-04	3.12E-05	3.13E-04	3.13E-04	4.54E-05	6.85E-06	5.23E-05
	41-78.6	1.89E-08	3.13E-04	3.12E-05	3.13E-04	3.13E-04	4.57E-05	6.91E-06	5.26E-05
AP8	0-19.2	1.58E-08	3.24E-04	3.23E-05	3.24E-04	3.23E-04	4.68E-05	7.95E-06	5.47E-05
	0-19.2	7.36E-09	1.98E-04	1.98E-05	1.98E-04	1.98E-04	2.03E-05	2.65E-06	2.30E-05
	19.2-41	7.36E-09	1.98E-04	1.98E-05	1.98E-04	1.98E-04	2.04E-05	2.66E-06	2.31E-05
	41-78.6	7.36E-09	1.98E-04	1.98E-05	1.98E-04	1.98E-04	2.04E-05	2.66E-06	2.31E-05
	78.6-120	7.36E-09	1.98E-04	1.98E-05	1.98E-04	1.98E-04	2.05E-05	2.70E-06	2.32E-05
	0-19.2	7.36E-09	1.88E-04	1.87E-05	1.88E-04	1.88E-04	2.03E-05	2.65E-06	2.30E-05
	19.2-41	7.36E-09	1.88E-04	1.87E-05	1.88E-04	1.88E-04	2.04E-05	2.66E-06	2.31E-05
	41-78.6	7.36E-09	1.88E-04	1.87E-05	1.88E-04	1.88E-04	2.04E-05	2.66E-06	2.31E-05
	78.6-120	7.36E-09	1.88E-04	1.87E-05	1.88E-04	1.88E-04	2.05E-05	2.70E-06	2.32E-05
KT3	0-19.2	7.36E-09	1.88E-04	1.87E-05	1.88E-04	1.88E-04	2.11E-05	2.68E-06	2.38E-05
KT4	0-19.2	7.36E-09	1.88E-04	1.87E-05	1.88E-04	1.88E-04	2.12E-05	2.70E-06	2.39E-05
	0-19.2	7.36E-09	1.88E-04	1.87E-05	1.88E-04	1.88E-04	2.03E-05	2.65E-06	2.30E-05
	0-19.2	5.26E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	1.71E-05	2.49E-06	1.96E-05
	19.2-41	5.26E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	1.72E-05	2.50E-06	1.97E-05
	41-78.6	5.26E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	1.72E-05	2.50E-06	1.97E-05
	78.6-120	5.26E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	1.73E-05	2.54E-06	1.98E-05
	120-175	5.26E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	1.75E-05	2.60E-06	2.01E-05
KT6	0-19.2	6.31E-09	1.57E-04	1.56E-05	1.57E-04	1.56E-04	1.96E-05	2.61E-06	2.22E-05
	19.2-41	6.31E-09	1.57E-04	1.56E-05	1.57E-04	1.56E-04	1.96E-05	2.63E-06	2.23E-05
KT7	0-19.2	4.20E-09	9.39E-05	9.37E-06	9.40E-05	9.39E-05	1.48E-05	2.39E-06	1.72E-05
	0-19.2	3.15E-09	8.35E-05	8.33E-06	8.36E-05	8.35E-05	1.33E-05	2.32E-06	1.56E-05
	19.2-41	3.15E-09	8.35E-05	8.33E-06	8.36E-05	8.35E-05	1.33E-05	2.32E-06	1.56E-05
	41-78.6	3.15E-09	8.35E-05	8.33E-06	8.36E-05	8.35E-05	1.33E-05	2.32E-06	1.56E-05
	78.6-120	3.15E-09	8.35E-05	8.33E-06	8.36E-05	8.35E-05	1.34E-05	2.35E-06	1.58E-05
SKC1	0-19.2	1.93E-06	1.04E-05	1.05E-06	1.11E-05				