

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

ASR	Assessment Height (mAG)	Antimony	Arsenic	Cadmium	Chromium	Lead	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	4.02E-06
TT2	0-19.2	0.00E+00	0.00E+00	8.87E-11	0.00E+00	0.00E+00	1.46E-05
TT3	0-19.2	0.00E+00	0.00E+00	1.40E-10	0.00E+00	0.00E+00	2.25E-05
TT4	0-19.2	0.00E+00	0.00E+00	5.58E-10	0.00E+00	0.00E+00	1.27E-04
TT5	0-19.2	0.00E+00	0.00E+00	2.42E-10	0.00E+00	0.00E+00	6.84E-06
TT6	0-19.2	0.00E+00	0.00E+00	2.11E-10	0.00E+00	0.00E+00	2.19E-05
TM1	0-19.2	0.00E+00	0.00E+00	1.11E-11	0.00E+00	0.00E+00	6.81E-07
	19.2-41	0.00E+00	0.00E+00	1.11E-11	0.00E+00	0.00E+00	6.81E-07
	41-78.6	0.00E+00	0.00E+00	1.11E-11	0.00E+00	0.00E+00	6.81E-07
	78.6-120	0.00E+00	0.00E+00	1.11E-11	0.00E+00	0.00E+00	6.81E-07
TM2	0-19.2	0.00E+00	0.00E+00	2.67E-11	0.00E+00	0.00E+00	5.28E-07
	19.2-41	0.00E+00	0.00E+00	2.67E-11	0.00E+00	0.00E+00	5.28E-07
	41-78.6	0.00E+00	0.00E+00	2.67E-11	0.00E+00	0.00E+00	5.28E-07
	78.6-120	0.00E+00	0.00E+00	2.67E-11	0.00E+00	0.00E+00	5.28E-07
TM3	0-19.2	0.00E+00	0.00E+00	1.97E-11	0.00E+00	0.00E+00	8.01E-07
	19.2-41	0.00E+00	0.00E+00	1.97E-11	0.00E+00	0.00E+00	8.01E-07
	41-78.6	0.00E+00	0.00E+00	1.97E-11	0.00E+00	0.00E+00	8.01E-07
	78.6-120	0.00E+00	0.00E+00	1.97E-11	0.00E+00	0.00E+00	8.01E-07
TM4	0-19.2	0.00E+00	0.00E+00	2.82E-11	0.00E+00	0.00E+00	8.52E-07
	19.2-41	0.00E+00	0.00E+00	2.82E-11	0.00E+00	0.00E+00	8.52E-07
	41-78.6	0.00E+00	0.00E+00	2.82E-11	0.00E+00	0.00E+00	8.52E-07
	78.6-120	0.00E+00	0.00E+00	2.82E-11	0.00E+00	0.00E+00	8.52E-07
TM5	0-19.2	0.00E+00	0.00E+00	2.04E-11	0.00E+00	0.00E+00	9.54E-07
	19.2-41	0.00E+00	0.00E+00	2.04E-11	0.00E+00	0.00E+00	9.54E-07
	41-78.6	0.00E+00	0.00E+00	2.04E-11	0.00E+00	0.00E+00	9.54E-07
	78.6-120	0.00E+00	0.00E+00	2.04E-11	0.00E+00	0.00E+00	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
	19.2-41	0.00E+00	0.00E+00	2.16E-11	0.00E+00	0.00E+00	1.19E-06
	41-78.6	0.00E+00	0.00E+00	2.16E-11	0.00E+00	0.00E+00	1.19E-06
	78.6-120	0.00E+00	0.00E+00	2.16E-11	0.00E+00	0.00E+00	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
SLW2	0-19.2	0.00E+00	0.00E+00	4.01E-10	0.00E+00	0.00E+00	1.71E-06
SLW3	0-19.2	0.00E+00	0.00E+00	1.42E-10	0.00E+00	0.00E+00	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
	19.2-41	0.00E+00	0.00E+00	9.98E-11	0.00E+00	0.00E+00	2.04E-07
	41-78.6	0.00E+00	0.00E+00	9.98E-11	0.00E+00	0.00E+00	2.04E-07
	78.6-120	0.00E+00	0.00E+00	9.98E-11	0.00E+00	0.00E+00	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
	19.2-41	0.00E+00	0.00E+00	9.13E-11	0.00E+00	0.00E+00	1.36E-07
	41-78.6	0.00E+00	0.00E+00	9.13E-11	0.00E+00	0.00E+00	1.36E-07
	78.6-120	0.00E+00	0.00E+00	9.13E-11	0.00E+00	0.00E+00	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
	19.2-41	0.00E+00	0.00E+00	8.29E-11	0.00E+00	0.00E+00	1.19E-07
	41-78.6	0.00E+00	0.00E+00	8.29E-11	0.00E+00	0.00E+00	1.19E-07
	78.6-120	0.00E+00	0.00E+00	8.29E-11	0.00E+00	0.00E+00	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
	19.2-41	0.00E+00	0.00E+00	9.13E-11	0.00E+00	0.00E+00	1.19E-07
	41-78.6	0.00E+00	0.00E+00	9.13E-11	0.00E+00	0.00E+00	1.19E-07
	78.6-120	0.00E+00	0.00E+00	9.13E-11	0.00E+00	0.00E+00	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
	19.2-41	0.00E+00	0.00E+00	9.97E-11	0.00E+00	0.00E+00	1.36E-07
	41-78.6	0.00E+00	0.00E+00	9.97E-11	0.00E+00	0.00E+00	1.36E-07
	78.6-120	0.00E+00	0.00E+00	9.97E-11	0.00E+00	0.00E+00	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
	19.2-41	0.00E+00	0.00E+00	9.97E-11	0.00E+00	0.00E+00	1.36E-07
	41-78.6	0.00E+00	0.00E+00	9.97E-11	0.00E+00	0.00E+00	1.36E-07
	78.6-120	0.00E+00	0.00E+00	9.97E-11	0.00E+00	0.00E+00	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
AP2	0-19.2	0.00E+00	0.00E+00	1.08E-10	0.00E+00	0.00E+00	2.56E-07
AP3	0-19.2	0.00E+00	0.00E+00	1.59E-10	0.00E+00	0.00E+00	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
	19.2-41	0.00E+00	0.00E+00	7.52E-11	0.00E+00	0.00E+00	2.21E-07
	41-78.6	0.00E+00	0.00E+00	7.52E-11	0.00E+00	0.00E+00	2.21E-07
	78.6-120	0.00E+00	0.00E+00	7.52E-11	0.00E+00	0.00E+00	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
AP6	0-19.2	0.00E+00	0.00E+00	7.49E-11	0.00E+00	0.00E+00	1.87E-07
AP7	0-19.2	0.00E+00	0.00E+00	8.34E-11	0.00E+00	0.00E+00	2.56E-07
	19.2-41	0.00E+00	0.00E+00	8.34E-11	0.00E+00	0.00E+00	2.56E-07
AP8	41-78.6	0.00E+00	0.00E+00	8.34E-11	0.00E+00	0.00E+00	2.56E-07
KT1	0-19.2	0.00E+00	0.00E+00	7.50E-11	0.00E+00	0.00E+00	1.87E-07
	19.2-41	0.00E+00	0.00E+00	4.14E-11	0.00E+00	0.00E+00	3.41E-08
	41-78.6	0.00E+00	0.00E+00	4.14E-11	0.00E+00	0.00E+00	3.41E-08
	78.6-120	0.00E+00	0.00E+00	4.14E-11	0.00E+00	0.00E+00	3.41E-08
KT2	0-19.2	0.00E+00	0.00E+00	3.32E-11	0.00E+00	0.00E+00	3.41E-08
	19.2-41	0.00E+00	0.00E+00	3.32E-11	0.00E+00	0.00E+00	3.41E-08
	41-78.6	0.00E+00	0.00E+00	3.32E-11	0.00E+00	0.00E+00	3.41E-08
	78.6-120	0.00E+00	0.00E+00	3.32E-11	0.00E+00	0.00E+00	3.41E-08
KT3	0-19.2	0.00E+00	0.00E+00	3.32E-11	0.00E+00	0.00E+00	3.41E-08
KT4	19.2-41	0.00E+00	0.00E+00	3.32E-11	0.00E+00	0.00E+00	3.41E-08
KT5	0-19.2	0.00E+00	0.00E+00	1.66E-11	0.00E+00	0.00E+00	3.41E-08
	19.2-41	0.00E+00	0.00E+00	1.66E-11	0.00E+00	0.00E+00	3.41E-08
	41-78.6	0.00E+00	0.00E+00	1.66E-11	0.00E+00	0.00E+00	3.41E-08
	78.6-120	0.00E+00	0.00E+00	1.66E-11	0.00E+00	0.00E+00	3.41E-08
KT6	0-19.2	0.00E+00	0.00E+00	2.49E-11	0.00E+00	0.00E+00	3.41E-08
	19.2-41	0.00E+00	0.00E+00	2.49E-11	0.00E+00	0.00E+00	3.41E-08
KT7	0-19.2	0.00E+00	0.00E+00	1.66E-11	0.00E+00	0.00E+00	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
	19.2-41	0.00E+00	0.00E+00	1.65E-11	0.00E+00	0.00E+00	1.70E-08
	41-78.6	0.00E+00	0.00E+00	1.65E-11	0.00E+00	0.00E+00	1.70E-08
	78.6-120	0.00E+00	0.00E+00	1.65E-11	0.00E+00	0.00E+00	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	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	9.59E-08
TT2	0-19.2	0.00E+00	0.00E+00	2.77E-09	0.00E+00	0.00E+00	3.47E-07
TT3	0-19.2	0.00E+00	0.00E+00	4.61E-09	0.00E+00	0.00E+00	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
TT5	0-19.2	0.00E+00	0.00E+00	2.40E-08	0.00E+00	0.00E+00	1.63E-07
TT6	0-19.2	0.00E+00	0.00E+00	1.29E-08	0.00E+00	0.00E+00	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
	19.2-41	0.00E+00	0.00E+00	9.22E-10	0.00E+00	0.00E+00	1.63E-08
	41-78.6	0.00E+00	0.00E+00	9.22E-10	0.00E+00	0.00E+00	1.63E-08
	78.6-120	0.00E+00	0.00E+00	9.22E-10	0.00E+00	0.00E+00	1.63E-08
TM2	0-19.2	0.00E+00	0.00E+00	2.77E-09	0.00E+00	0.00E+00	1.27E-08
	19.2-41	0.00E+00	0.00E+00	2.77E-09	0.00E+00	0.00E+00	1.27E-08
	41-78.6	0.00E+00	0.00E+00	2.77E-09	0.00E+00	0.00E+00	1.27E-08
	78.6-120	0.00E+00	0.00E+00	2.77E-09	0.00E+00	0.00E+00	1.27E-08
TM3	0-19.2	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	1.92E-08
	19.2-41	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	1.92E-08
	41-78.6	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	1.92E-08
	78.6-120	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	1.92E-08
TM4	0-19.2	0.00E+00	0.00E+00	2.77E-09	0.00E+00	0.00E+00	2.04E-08
	19.2-41	0.00E+00	0.00E+00	2.77E-09	0.00E+00	0.00E+00	2.04E-08
	41-78.6	0.00E+00	0.00E+00	2.77E-09	0.00E+00	0.00E+00	2.04E-08
	78.6-120	0.00E+00	0.00E+00	2.77E-09	0.00E+00	0.00E+00	2.04E-08
TM5	0-19.2	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	2.28E-08
	19.2-41	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	2.28E-08
	41-78.6	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	2.28E-08
	78.6-120	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	2.28E-08
TM6	0-19.2	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	2.85E-08
	19.2-41	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	2.85E-08
	41-78.6	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	2.85E-08
	78.6-120	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	2.85E-08
SLW1	0-19.2	0.00E+00	0.00E+00	4.15E-08	0.00E+00	0.00E+00	3.70E-08
SLW2	0-19.2	0.00E+00	0.00E+00	4.42E-08	0.00E+00	0.00E+00	4.11E-08
SLW3	0-19.2	0.00E+00	0.00E+00	1.57E-08	0.00E+00	0.00E+00	8.83E-09
TC1	0-19.2	0.00E+00	0.00E+00	1.11E-08	0.00E+00	0.00E+00	5.03E-09
	19.2-41	0.00E+00	0.00E+00	1.11E-08	0.00E+00	0.00E+00	5.03E-09
	41-78.6	0.00E+00	0.00E+00	1.11E-08	0.00E+00	0.00E+00	5.03E-09
	78.6-120	0.00E+00	0.00E+00	1.11E-08	0.00E+00	0.00E+00	5.03E-09
TC2	0-19.2	0.00E+00	0.00E+00	1.01E-08	0.00E+00	0.00E+00	3.41E-09
	19.2-41	0.00E+00	0.00E+00	1.01E-08	0.00E+00	0.00E+00	3.41E-09
	41-78.6	0.00E+00	0.00E+00	1.01E-08	0.00E+00	0.00E+00	3.41E-09
	78.6-120	0.00E+00	0.00E+00	1.01E-08	0.00E+00	0.00E+00	3.41E-09
TC3	0-19.2	0.00E+00	0.00E+00	1.01E-08	0.00E+00	0.00E+00	3.41E-09
	19.2-41	0.00E+00	0.00E+00	1.01E-08	0.00E+00	0.00E+00	3.41E-09
	41-78.6	0.00E+00	0.00E+00	1.01E-08	0.00E+00	0.00E+00	3.41E-09
	78.6-120	0.00E+00	0.00E+00	1.01E-08	0.00E+00	0.00E+00	3.41E-09
TC4	0-19.2	0.00E+00	0.00E+00	9.22E-09	0.00E+00	0.00E+00	2.99E-09
	19.2-41	0.00E+00	0.00E+00	9.22E-09	0.00E+00	0.00E+00	2.99E-09
	41-78.6	0.00E+00	0.00E+00	9.22E-09	0.00E+00	0.00E+00	2.99E-09
	78.6-120	0.00E+00	0.00E+00	9.22E-09	0.00E+00	0.00E+00	2.99E-09
TC5	0-19.2	0.00E+00	0.00E+00	1.01E-08	0.00E+00	0.00E+00	3.02E-09
	19.2-41	0.00E+00	0.00E+00	1.01E-08	0.00E+00	0.00E+00	3.02E-09
	41-78.6	0.00E+00	0.00E+00	1.01E-08	0.00E+00	0.00E+00	3.02E-09
	78.6-120	0.00E+00	0.00E+00	1.01E-08	0.00E+00	0.00E+00	3.02E-09
TC6	0-19.2	0.00E+00	0.00E+00	1.11E-08	0.00E+00	0.00E+00	3.42E-09
	19.2-41	0.00E+00	0.00E+00	1.11E-08	0.00E+00	0.00E+00	3.42E-09
	41-78.6	0.00E+00	0.00E+00	1.11E-08	0.00E+00	0.00E+00	3.42E-09
	78.6-120	0.00E+00	0.00E+00	1.11E-08	0.00E+00	0.00E+00	3.42E-09
AP1	0-19.2	0.00E+00	0.00E+00	1.20E-08	0.00E+00	0.00E+00	5.05E-09
AP2	0-19.2	0.00E+00	0.00E+00	1.20E-08	0.00E+00	0.00E+00	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
AP4	0-19.2	0.00E+00	0.00E+00	8.30E-09	0.00E+00	0.00E+00	5.45E-09
	19.2-41	0.00E+00	0.00E+00	8.30E-09	0.00E+00	0.00E+00	5.45E-09
	41-78.6	0.00E+00	0.00E+00	8.30E-09	0.00E+00	0.00E+00	5.45E-09
	78.6-120	0.00E+00	0.00E+00	8.30E-09	0.00E+00	0.00E+00	5.45E-09
AP5	0-19.2	0.00E+00	0.00E+00	8.30E-09	0.00E+00	0.00E+00	4.64E-09
AP6	0-19.2	0.00E+00	0.00E+00	8.30E-09	0.00E+00	0.00E+00	4.64E-09
AP7	0-19.2	0.00E+00	0.00E+00	9.22E-09	0.00E+00	0.00E+00	6.25E-09
	19.2-41	0.00E+00	0.00E+00	9.22E-09	0.00E+00	0.00E+00	6.25E-09
AP8	0-19.2	0.00E+00	0.00E+00	9.22E-09	0.00E+00	0.00E+00	6.25E-09
KT1	0-19.2	0.00E+00	0.00E+00	8.30E-09	0.00E+00	0.00E+00	4.64E-09
	19.2-41	0.00E+00	0.00E+00	4.61E-09	0.00E+00	0.00E+00	8.73E-10
	41-78.6	0.00E+00	0.00E+00	4.61E-09	0.00E+00	0.00E+00	8.74E-10
	78.6-120	0.00E+00	0.00E+00	4.61E-09	0.00E+00	0.00E+00	8.75E-10
KT2	0-19.2	0.00E+00	0.00E+00	3.69E-09	0.00E+00	0.00E+00	8.73E-10
	19.2-41	0.00E+00	0.00E+00	3.69E-09	0.00E+00	0.00E+00	8.74E-10
	41-78.6	0.00E+00	0.00E+00	3.69E-09	0.00E+00	0.00E+00	8.74E-10
	78.6-120	0.00E+00	0.00E+00	3.69E-09	0.00E+00	0.00E+00	8.75E-10
KT3	0-19.2	0.00E+00	0.00E+00	3.69E-09	0.00E+00	0.00E+00	8.73E-10
	19.2-41	0.00E+00	0.00E+00	3.69E-09	0.00E+00	0.00E+00	8.74E-10
KT4	0-19.2	0.00E+00	0.00E+00	3.69E-09	0.00E+00	0.00E+00	8.73E-10
KT5	0-19.2	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	8.73E-10
	19.2-41	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	8.74E-10
	41-78.6	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	8.74E-10
	78.6-120	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	8.75E-10
KT6	0-19.2	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	8.79E-10
	19.2-41	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	8.79E-10
KT7	0-19.2	0.00E+00	0.00E+00	2.77E-09	0.00E+00	0.00E+00	8.73E-10
KT8	0-19.2	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	8.74E-10
	19.2-41	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	8.69E-10
	41-78.6	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	4.63E-10
	78.6-120	0.00E+00	0.00E+00	1.84E-09	0.00E+00	0.00E+00	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	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	1.89E-04
TT2	0-19.2	8.94E-07	3.13E-05	3.13E-06	3.16E-05	3.13E-05	6.79E-04
TT3	0-19.2	1.38E-06	5.22E-05	5.21E-06	5.27E-05	5.22E-05	1.05E-03
TT4	0-19.2	7.81E-06	2.27E-10	2.88E-08	2.66E-06	3.99E-07	5.89E-03
TT5	0-19.2	3.77E-07	2.71E-04	2.71E-05	2.72E-04	2.71E-04	3.23E-04
TT6	0-19.2	1.33E-06	1.46E-04	1.46E-05	1.47E-04	1.46E-04	1.02E-03
TM1	0-19.2	3.99E-08	1.04E-05	1.04E-06	1.05E-05	1.04E-05	3.66E-05
	19.2-41	3.99E-08	1.04E-05	1.04E-06	1.05E-05	1.04E-05	3.67E-05
	41-78.6	3.99E-08	1.04E-05	1.04E-06	1.05E-05	1.04E-05	3.71E-05
	78.6-120	3.99E-08	1.04E-05	1.04E-06	1.05E-05	1.04E-05	3.78E-05
	0-19.2	2.63E-08	3.13E-05	3.12E-06	3.13E-05	3.13E-05	2.94E-05
TM2	19.2-41	2.63E-08	3.13E-05	3.12E-06	3.13E-05	3.13E-05	2.96E-05
	41-78.6	2.63E-08	3.13E-05	3.12E-06	3.13E-05	3.13E-05	2.99E-05
	78.6-120	2.63E-08	3.13E-05	3.12E-06	3.13E-05	3.13E-05	3.06E-05
	0-19.2	4.41E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	4.21E-05
TM3	19.2-41	4.41E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	4.23E-05
	41-78.6	4.41E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	4.26E-05
	78.6-120	4.41E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	4.33E-05
	0-19.2	4.73E-08	3.13E-05	3.12E-06	3.13E-05	3.13E-05	4.45E-05
TM4	19.2-41	4.73E-08	3.13E-05	3.12E-06	3.13E-05	3.13E-05	4.47E-05
	41-78.6	4.73E-08	3.13E-05	3.12E-06	3.13E-05	3.13E-05	4.50E-05
	78.6-120	4.73E-08	3.13E-05	3.12E-06	3.13E-05	3.13E-05	4.57E-05
	0-19.2	5.36E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	4.86E-05
TM5	19.2-41	5.36E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	4.88E-05
	41-78.6	5.36E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	4.93E-05
	78.6-120	5.36E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	5.07E-05
	0-19.2	7.04E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	6.07E-05
TM6	19.2-41	7.04E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	6.09E-05
	41-78.6	7.04E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	6.15E-05
	78.6-120	7.04E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	6.28E-05
	0-19.2	5.78E-08	4.70E-04	4.68E-05	4.70E-04	4.69E-04	8.46E-05
SLW1	0-19.2	6.31E-08	5.01E-04	5.00E-05	5.01E-04	5.01E-04	9.27E-05
SLW2	0-19.2	1.37E-08	1.77E-04	1.77E-05	1.78E-04	1.77E-04	2.86E-05
SLW3	0-19.2	7.36E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	1.80E-05
TC1	19.2-41	7.36E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	1.80E-05
	41-78.6	7.36E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	1.82E-05
	78.6-120	7.36E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	1.82E-05
	120-175	7.36E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	1.82E-05
	0-19.2	4.20E-09	1.15E-04	1.14E-05	1.15E-04	1.15E-04	1.49E-05
TC2	19.2-41	4.20E-09	1.15E-04	1.14E-05	1.15E-04	1.15E-04	1.49E-05
	41-78.6	4.20E-09	1.15E-04	1.14E-05	1.15E-04	1.15E-04	1.50E-05
	78.6-120	4.20E-09	1.15E-04	1.14E-05	1.15E-04	1.15E-04	1.50E-05
	120-175	4.20E-09	1.15E-04	1.14E-05	1.15E-04	1.15E-04	1.50E-05
	0-19.2	3.15E-09	1.04E-04	1.04E-05	1.04E-04	1.04E-04	1.35E-05
TC3	19.2-41	3.15E-09	1.04E-04	1.04E-05	1.04E-04	1.04E-04	1.35E-05
	0-19.2	4.20E-09	1.15E-04	1.14E-05	1.15E-04	1.15E-04	1.35E-05
	19.2-41	4.20E-09	1.15E-04	1.14E-05	1.15E-04	1.15E-04	1.35E-05
	41-78.6	4.20E-09	1.15E-04	1.14E-05	1.15E-04	1.15E-04	1.39E-05
	78.6-120	4.20E-09	1.15E-04	1.14E-05	1.15E-04	1.15E-04	1.44E-05
TC4	120-175	4.20E-09	1.15E-04	1.14E-05	1.15E-04	1.15E-04	1.51E-05
	0-19.2	6.31E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	1.54E-05
	19.2-41	6.31E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	1.55E-05
	41-78.6	6.31E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	1.60E-05
	78.6-120	6.31E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	1.67E-05
TC5	120-175	6.31E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	1.77E-05
	0-19.2	6.31E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	1.54E-05
	19.2-41	6.31E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	1.54E-05
	41-78.6	6.31E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	1.55E-05
	78.6-120	6.31E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	1.60E-05
TC6	120-175	6.31E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	1.77E-05
	0-19.2	9.46E-09	1.36E-04	1.35E-05	1.36E-04	1.36E-04	1.88E-05
	AP1	0-19.2	1.26E-08	1.36E-04	1.35E-05	1.36E-04	2.12E-05
	AP2	0-19.2	2.21E-08	1.98E-04	1.98E-05	1.98E-04	3.96E-05
	AP3	0-19.2	1.05E-08	9.39E-05	9.37E-06	9.40E-05	9.39E-05
AP4	19.2-41	1.05E-08	9.39E-05	9.37E-06	9.40E-05	9.39E-05	1.93E-05
	41-78.6	1.05E-08	9.39E-05	9.37E-06	9.40E-05	9.39E-05	1.95E-05
	78.6-120	1.05E-08	9.39E-05	9.37E-06	9.40E-05	9.39E-05	1.96E-05
	0-19.2	7.36E-09	9.39E-05	9.37E-06	9.40E-05	9.39E-05	1.77E-05
AP5	0-19.2	6.31E-09	9.39E-05	9.37E-06	9.40E-05	9.39E-05	1.81E-05
AP6	0-19.2	1.05E-08	1.04E-04	1.04E-05	1.04E-04	1.04E-04	2.06E-05
AP7	19.2-41	1.05E-08	1.04E-04	1.04E-05	1.04E-04	1.04E-04	2.05E-05
	41-78.6	1.05E-08	1.04E-04	1.04E-05	1.04E-04	1.04E-04	2.06E-05
	0-19.2	7.36E-09	9.39E-05	9.37E-06	9.40E-05	9.39E-05	1.81E-05
AP8	0-19.2	1.05E-09	5.22E-05	5.20E-06	5.22E-05	5.22E-05	4.75E-06
KT1	19.2-41	1.05E-09	5.22E-05	5.20E-06	5.22E-05	5.22E-05	4.82E-06
	41-78.6	1.05E-09	5.22E-05	5.20E-06	5.22E-05	5.22E-05	4.82E-06
	78.6-120	1.05E-09	5.22E-05	5.20E-06	5.22E-05	5.22E-05	4.89E-06
	0-19.2	1.05E-09	4.17E-05	4.16E-06	4.18E-05	4.17E-05	4.75E-06
KT2	19.2-41	1.05E-09	4.17E-05	4.16E-06	4.18E-05	4.17E-05	4.82E-06
	41-78.6	1.05E-09	4.17E-05	4.16E-06	4.18E-05	4.17E-05	4.82E-06
	78.6-120	1.05E-09	4.17E-05	4.16E-06	4.18E-05	4.17E-05	4.89E-06
	0-19.2	1.05E-09	4.17E-05	4.16E-06	4.18E-05	4.17E-05	4.75E-06
KT3	19.2-41	1.05E-09	4.17E-05	4.16E-06	4.18E-05	4.17E-05	4.82E-06
KT4	0-19.2	1.05E-09	4.17E-05	4.16E-06	4.18E-05	4.17E-05	4.75E-06
KT5	19.2-41	1.05E-09	2.09E-05	2.08E-06	2.09E-05	2.09E-05	4.75E-06
	41-78.6	1.05E-09	2.09E-05	2.08E-06	2.09E-05	2.09E-05	4.82E-06
	78.6-120	1.05E-09	2.09E-05	2.08E-06	2.09E-05	2.09E-05	4.82E-06
	120-175	1.05E-09	2.09E-05	2.08E-06	2.09E-05	2.09E-05	5.11E-06
	0-19.2	1.05E-09	3.13E-05	3.12E-06	3.13E-05	3.13E-05	4.75E-06
KT6	19.2-41	1.05E-09	3.13E-05	3.12E-06	3.13E-05	3.13E-05	4.82E-06
KT7	0-19.2	1.05E-09	2.09E-05	2.08E-06	2.09E-05	2.09E-05	4.54E-06
KT8	19.2-41	1.05E-09	2.09E-05	2.08E-06	2.09E-05	2.09E-05	3.75E-06
	41-78.6	1.05E-09	2.09E-05	2.08E-06	2.09E-05	2.09E-05	3.75E-06
	78.6-120	1.05E-09	2.09E-05	2.08E-06	2.09E-05	2.09E-05	3.89E-06

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	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	3.12E-05
SL1	0-19.2	0.00E+00	0.00E+00	1.70E-10	0.00E+00	0.00E+00	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
SLW1	0-19.2	0.00E+00	0.00E+00	1.91E-10	0.00E+00	0.00E+00	5.81E-07
SLW2	0-19.2	0.00E+00	0.00E+00	2.33E-10	0.00E+00	0.00E+00	5.98E-07
SLW3	0-19.2	0.00E+00	0.00E+00	1.58E-10	0.00E+00	0.00E+00	5.81E-07
TC1	0-19.2	0.00E+00	0.00E+00	1.25E-10	0.00E+00	0.00E+00	3.94E-07
	19.2-41	0.00E+00	0.00E+00	1.25E-10	0.00E+00	0.00E+00	3.94E-07
	41-78.6	0.00E+00	0.00E+00	1.25E-10	0.00E+00	0.00E+00	3.94E-07
	78.6-120	0.00E+00	0.00E+00	1.25E-10	0.00E+00	0.00E+00	3.94E-07
	120-175	0.00E+00	0.00E+00	1.25E-10	0.00E+00	0.00E+00	3.94E-07
TC2	0-19.2	0.00E+00	0.00E+00	1.50E-10	0.00E+00	0.00E+00	3.94E-07
	19.2-41	0.00E+00	0.00E+00	1.50E-10	0.00E+00	0.00E+00	3.94E-07
	41-78.6	0.00E+00	0.00E+00	1.50E-10	0.00E+00	0.00E+00	3.94E-07
	78.6-120	0.00E+00	0.00E+00	1.50E-10	0.00E+00	0.00E+00	3.94E-07
	120-175	0.00E+00	0.00E+00	1.50E-10	0.00E+00	0.00E+00	3.94E-07
TC3	0-19.2	0.00E+00	0.00E+00	1.58E-10	0.00E+00	0.00E+00	3.94E-07
	19.2-41	0.00E+00	0.00E+00	1.58E-10	0.00E+00	0.00E+00	3.94E-07
TC4	0-19.2	0.00E+00	0.00E+00	1.66E-10	0.00E+00	0.00E+00	4.11E-07
	19.2-41	0.00E+00	0.00E+00	1.66E-10	0.00E+00	0.00E+00	4.11E-07
	41-78.6	0.00E+00	0.00E+00	1.66E-10	0.00E+00	0.00E+00	4.11E-07
	78.6-120	0.00E+00	0.00E+00	1.66E-10	0.00E+00	0.00E+00	4.11E-07
	120-175	0.00E+00	0.00E+00	1.66E-10	0.00E+00	0.00E+00	4.11E-07
TC5	0-19.2	0.00E+00	0.00E+00	1.75E-10	0.00E+00	0.00E+00	4.79E-07
	19.2-41	0.00E+00	0.00E+00	1.75E-10	0.00E+00	0.00E+00	4.79E-07
	41-78.6	0.00E+00	0.00E+00	1.75E-10	0.00E+00	0.00E+00	4.79E-07
	78.6-120	0.00E+00	0.00E+00	1.75E-10	0.00E+00	0.00E+00	4.79E-07
	120-175	0.00E+00	0.00E+00	1.75E-10	0.00E+00	0.00E+00	4.79E-07
TC6	0-19.2	0.00E+00	0.00E+00	1.34E-10	0.00E+00	0.00E+00	6.15E-07
	19.2-41	0.00E+00	0.00E+00	1.34E-10	0.00E+00	0.00E+00	6.15E-07
	41-78.6	0.00E+00	0.00E+00	1.34E-10	0.00E+00	0.00E+00	6.15E-07
	78.6-120	0.00E+00	0.00E+00	1.34E-10	0.00E+00	0.00E+00	5.98E-07
	120-175	0.00E+00	0.00E+00	1.34E-10	0.00E+00	0.00E+00	5.98E-07
AP1	0-19.2	0.00E+00	0.00E+00	1.83E-10	0.00E+00	0.00E+00	4.11E-07
AP2	0-19.2	0.00E+00	0.00E+00	1.83E-10	0.00E+00	0.00E+00	3.77E-07
AP3	0-19.2	0.00E+00	0.00E+00	1.58E-10	0.00E+00	0.00E+00	5.47E-07
AP4	0-19.2	0.00E+00	0.00E+00	1.91E-10	0.00E+00	0.00E+00	3.09E-07
	19.2-41	0.00E+00	0.00E+00	1.91E-10	0.00E+00	0.00E+00	3.09E-07
	41-78.6	0.00E+00	0.00E+00	1.91E-10	0.00E+00	0.00E+00	3.09E-07
	78.6-120	0.00E+00	0.00E+00	1.91E-10	0.00E+00	0.00E+00	3.09E-07
AP5	0-19.2	0.00E+00	0.00E+00	1.66E-10	0.00E+00	0.00E+00	3.26E-07
AP6	0-19.2	0.00E+00	0.00E+00	1.66E-10	0.00E+00	0.00E+00	3.26E-07
AP7	0-19.2	0.00E+00	0.00E+00	1.66E-10	0.00E+00	0.00E+00	3.09E-07
	19.2-41	0.00E+00	0.00E+00	1.66E-10	0.00E+00	0.00E+00	3.09E-07
AP8	41-78.6	0.00E+00	0.00E+00	1.66E-10	0.00E+00	0.00E+00	3.09E-07
	0-19.2	0.00E+00	0.00E+00	1.83E-10	0.00E+00	0.00E+00	3.09E-07
KT1	0-19.2	0.00E+00	0.00E+00	1.16E-10	0.00E+00	0.00E+00	2.58E-07
	19.2-41	0.00E+00	0.00E+00	1.16E-10	0.00E+00	0.00E+00	2.58E-07
	41-78.6	0.00E+00	0.00E+00	1.16E-10	0.00E+00	0.00E+00	2.58E-07
	78.6-120	0.00E+00	0.00E+00	1.16E-10	0.00E+00	0.00E+00	2.58E-07
	120-175	0.00E+00	0.00E+00	1.16E-10	0.00E+00	0.00E+00	2.58E-07
KT2	0-19.2	0.00E+00	0.00E+00	1.16E-10	0.00E+00	0.00E+00	2.58E-07
	19.2-41	0.00E+00	0.00E+00	1.16E-10	0.00E+00	0.00E+00	2.58E-07
	41-78.6	0.00E+00	0.00E+00	1.16E-10	0.00E+00	0.00E+00	2.58E-07
	78.6-120	0.00E+00	0.00E+00	1.16E-10	0.00E+00	0.00E+00	2.58E-07
KT3	0-19.2	0.00E+00	0.00E+00	1.16E-10	0.00E+00	0.00E+00	2.75E-07
	19.2-41	0.00E+00	0.00E+00	1.16E-10	0.00E+00	0.00E+00	2.75E-07
KT4	0-19.2	0.00E+00	0.00E+00	1.16E-10	0.00E+00	0.00E+00	2.58E-07
KT5	0-19.2	0.00E+00	0.00E+00	8.30E-11	0.00E+00	0.00E+00	1.87E-07
	19.2-41	0.00E+00	0.00E+00	8.30E-11	0.00E+00	0.00E+00	1.87E-07
	41-78.6	0.00E+00	0.00E+00	8.30E-11	0.00E+00	0.00E+00	1.87E-07
	78.6-120	0.00E+00	0.00E+00	8.30E-11	0.00E+00	0.00E+00	1.87E-07
	120-175	0.00E+00	0.00E+00	8.30E-11	0.00E+00	0.00E+00	1.87E-07
KT6	0-19.2	0.00E+00	0.00E+00	9.96E-11	0.00E+00	0.00E+00	2.41E-07
	19.2-41	0.00E+00	0.00E+00	9.96E-11	0.00E+00	0.00E+00	2.41E-07
KT7	0-19.2	0.00E+00	0.00E+00	5.81E-11	0.00E+00	0.00E+00	1.36E-07
KT8	0-19.2	0.00E+00	0.00E+00	4.98E-11	0.00E+00	0.00E+00	1.19E-07
	19.2-41	0.00E+00	0.00E+00	4.98E-11	0.00E+00	0.00E+00	1.19E-07
	41-78.6	0.00E+00	0.00E+00	4.98E-11	0.00E+00	0.00E+00	1.19E-07
	78.6-120	0.00E+00	0.00E+00	4.98E-11	0.00E+00	0.00E+00	1.19E-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	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	7.44E-07
SL1	0-19.2	0.00E+00	0.00E+00	1.84E-08	0.00E+00	0.00E+00	3.56E-08
SL2	0-19.2	0.00E+00	0.00E+00	2.77E-08	0.00E+00	0.00E+00	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
SLW2	0-19.2	0.00E+00	0.00E+00	2.58E-08	0.00E+00	0.00E+00	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
TC1	0-19.2	0.00E+00	0.00E+00	1.38E-08	0.00E+00	0.00E+00	9.81E-09
	19.2-41	0.00E+00	0.00E+00	1.38E-08	0.00E+00	0.00E+00	9.82E-09
	41-78.6	0.00E+00	0.00E+00	1.38E-08	0.00E+00	0.00E+00	9.84E-09
	78.6-120	0.00E+00	0.00E+00	1.38E-08	0.00E+00	0.00E+00	9.89E-09
	120-175	0.00E+00	0.00E+00	1.38E-08	0.00E+00	0.00E+00	9.97E-09
TC2	0-19.2	0.00E+00	0.00E+00	1.66E-08	0.00E+00	0.00E+00	9.81E-09
	19.2-41	0.00E+00	0.00E+00	1.66E-08	0.00E+00	0.00E+00	9.82E-09
	41-78.6	0.00E+00	0.00E+00	1.66E-08	0.00E+00	0.00E+00	9.84E-09
	78.6-120	0.00E+00	0.00E+00	1.66E-08	0.00E+00	0.00E+00	9.89E-09
	120-175	0.00E+00	0.00E+00	1.66E-08	0.00E+00	0.00E+00	9.97E-09
TC3	0-19.2	0.00E+00	0.00E+00	1.75E-08	0.00E+00	0.00E+00	9.82E-09
	19.2-41	0.00E+00	0.00E+00	1.75E-08	0.00E+00	0.00E+00	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
	19.2-41	0.00E+00	0.00E+00	1.84E-08	0.00E+00	0.00E+00	1.02E-08
	41-78.6	0.00E+00	0.00E+00	1.84E-08	0.00E+00	0.00E+00	1.03E-08
	78.6-120	0.00E+00	0.00E+00	1.84E-08	0.00E+00	0.00E+00	1.04E-08
	120-175	0.00E+00	0.00E+00	1.84E-08	0.00E+00	0.00E+00	1.05E-08
TC5	0-19.2	0.00E+00	0.00E+00	1.94E-08	0.00E+00	0.00E+00	1.19E-08
	19.2-41	0.00E+00	0.00E+00	1.94E-08	0.00E+00	0.00E+00	1.19E-08
	41-78.6	0.00E+00	0.00E+00	1.94E-08	0.00E+00	0.00E+00	1.19E-08
	78.6-120	0.00E+00	0.00E+00	1.94E-08	0.00E+00	0.00E+00	1.20E-08
	120-175	0.00E+00	0.00E+00	1.94E-08	0.00E+00	0.00E+00	1.21E-08
TC6	0-19.2	0.00E+00	0.00E+00	1.47E-08	0.00E+00	0.00E+00	1.51E-08
	19.2-41	0.00E+00	0.00E+00	1.47E-08	0.00E+00	0.00E+00	1.51E-08
	41-78.6	0.00E+00	0.00E+00	1.47E-08	0.00E+00	0.00E+00	1.52E-08
	78.6-120	0.00E+00	0.00E+00	1.47E-08	0.00E+00	0.00E+00	1.48E-08
	120-175	0.00E+00	0.00E+00	1.47E-08	0.00E+00	0.00E+00	1.49E-08
AP1	0-19.2	0.00E+00	0.00E+00	2.03E-08	0.00E+00	0.00E+00	1.03E-08
AP2	0-19.2	0.00E+00	0.00E+00	2.03E-08	0.00E+00	0.00E+00	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
AP4	0-19.2	0.00E+00	0.00E+00	2.12E-08	0.00E+00	0.00E+00	7.74E-09
	19.2-41	0.00E+00	0.00E+00	2.12E-08	0.00E+00	0.00E+00	7.74E-09
	41-78.6	0.00E+00	0.00E+00	2.12E-08	0.00E+00	0.00E+00	7.75E-09
	78.6-120	0.00E+00	0.00E+00	2.12E-08	0.00E+00	0.00E+00	7.75E-09
AP5	0-19.2	0.00E+00	0.00E+00	1.84E-08	0.00E+00	0.00E+00	8.15E-09
AP6	0-19.2	0.00E+00	0.00E+00	1.84E-08	0.00E+00	0.00E+00	8.20E-09
AP7	0-19.2	0.00E+00	0.00E+00	1.84E-08	0.00E+00	0.00E+00	7.70E-09
	19.2-41	0.00E+00	0.00E+00	1.84E-08	0.00E+00	0.00E+00	7.70E-09
AP8	41-78.6	0.00E+00	0.00E+00	1.84E-08	0.00E+00	0.00E+00	7.71E-09
	0-19.2	0.00E+00	0.00E+00	2.03E-08	0.00E+00	0.00E+00	7.80E-09
KT1	0-19.2	0.00E+00	0.00E+00	1.29E-08	0.00E+00	0.00E+00	6.27E-09
	19.2-41	0.00E+00	0.00E+00	1.29E-08	0.00E+00	0.00E+00	6.27E-09
	41-78.6	0.00E+00	0.00E+00	1.29E-08	0.00E+00	0.00E+00	6.27E-09
	78.6-120	0.00E+00	0.00E+00	1.29E-08	0.00E+00	0.00E+00	6.28E-09
	0-19.2	0.00E+00	0.00E+00	1.29E-08	0.00E+00	0.00E+00	6.27E-09
KT2	19.2-41	0.00E+00	0.00E+00	1.29E-08	0.00E+00	0.00E+00	6.27E-09
	41-78.6	0.00E+00	0.00E+00	1.29E-08	0.00E+00	0.00E+00	6.27E-09
	78.6-120	0.00E+00	0.00E+00	1.29E-08	0.00E+00	0.00E+00	6.28E-09
KT3	0-19.2	0.00E+00	0.00E+00	1.29E-08	0.00E+00	0.00E+00	6.68E-09
	19.2-41	0.00E+00	0.00E+00	1.29E-08	0.00E+00	0.00E+00	6.68E-09
KT4	0-19.2	0.00E+00	0.00E+00	1.29E-08	0.00E+00	0.00E+00	6.27E-09
KT5	0-19.2	0.00E+00	0.00E+00	9.22E-09	0.00E+00	0.00E+00	4.57E-09
	19.2-41	0.00E+00	0.00E+00	9.22E-09	0.00E+00	0.00E+00	4.57E-09
	41-78.6	0.00E+00	0.00E+00	9.22E-09	0.00E+00	0.00E+00	4.57E-09
	78.6-120	0.00E+00	0.00E+00	9.22E-09	0.00E+00	0.00E+00	4.57E-09
	120-175	0.00E+00	0.00E+00	9.22E-09	0.00E+00	0.00E+00	4.58E-09
KT6	0-19.2	0.00E+00	0.00E+00	1.11E-08	0.00E+00	0.00E+00	5.87E-09
	19.2-41	0.00E+00	0.00E+00	1.11E-08	0.00E+00	0.00E+00	5.87E-09
KT7	0-19.2	0.00E+00	0.00E+00	6.45E-09	0.00E+00	0.00E+00	3.36E-09
KT8	0-19.2	0.00E+00	0.00E+00	5.53E-09	0.00E+00	0.00E+00	2.96E-09
	19.2-41	0.00E+00	0.00E+00	5.53E-09	0.00E+00	0.00E+00	2.96E-09
	41-78.6	0.00E+00	0.00E+00	5.53E-09	0.00E+00	0.00E+00	2.96E-09
	78.6-120	0.00E+00	0.00E+00	5.53E-09	0.00E+00	0.00E+00	2.96E-09

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

ASR	Assessment Height (mAG)	Antimony	Arsenic	Cadmium	Chromium	Lead	Mercury
Maximum Permitted Concentration (ppm)		1	1.4	2	1	6	0.5
SKC1	0-19.2	5.10E-06	1.33E-07	1.40E-07	2.95E-08	0.00E+00	3.03E-03
SL1	0-19.2	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
SL2	0-19.2	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
SLW1	0-19.2	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
SLW2	0-19.2	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
SLW3	0-19.2	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
TC1	0-19.2	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
	19.2-41	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
	41-78.6	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
	78.6-120	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
	120-175	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
TC2	0-19.2	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
	19.2-41	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
	41-78.6	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
	78.6-120	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
	120-175	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
TC3	0-19.2	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
	19.2-41	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
TC4	0-19.2	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
	19.2-41	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
	41-78.6	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
	78.6-120	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
	120-175	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
TC5	0-19.2	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
	19.2-41	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
	41-78.6	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
	78.6-120	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
	120-175	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
TC6	0-19.2	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
	19.2-41	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
	41-78.6	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
	78.6-120	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
	120-175	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
AP1	0-19.2	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
AP2	0-19.2	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
AP3	0-19.2	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
AP4	0-19.2	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
	19.2-41	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
	41-78.6	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
	78.6-120	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
AP5	0-19.2	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
AP6	0-19.2	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
AP7	0-19.2	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
	19.2-41	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
AP8	41-78.6	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
	0-19.2	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
KT1	0-19.2	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
	19.2-41	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
	41-78.6	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
	78.6-120	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
	0-19.2	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
KT2	19.2-41	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
	41-78.6	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
	78.6-120	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
	0-19.2	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
KT3	19.2-41	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
	0-19.2	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
KT4	0-19.2	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
KT5	0-19.2	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
	19.2-41	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
	41-78.6	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
	78.6-120	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
	120-175	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
KT6	0-19.2	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
	19.2-41	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
KT7	0-19.2	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
KT8	0-19.2	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
	19.2-41	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
	41-78.6	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04
	78.6-120	5.21E-07	7.16E-07	5.72E-07	1.43E-07	0.00E+00	5.66E-04

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

ASR	Assessment Height (mAG)	Antimony	Arsenic	Cadmium	Chromium	Lead	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	1.45E-03
SL1	0-19.2	6.10E-08	2.09E-04	2.08E-05	2.09E-04	2.09E-04	8.13E-05
SL2	0-19.2	8.09E-08	3.13E-04	3.12E-05	3.13E-04	3.13E-04	9.65E-05
SLW1	0-19.2	1.79E-08	2.40E-04	2.39E-05	2.40E-04	2.40E-04	5.09E-05
SLW2	0-19.2	1.79E-08	2.92E-04	2.91E-05	2.92E-04	2.92E-04	5.40E-05
SLW3	0-19.2	1.58E-08	1.98E-04	1.98E-05	1.98E-04	1.98E-04	5.28E-05
TC1	0-19.2	1.26E-08	1.57E-04	1.56E-05	1.57E-04	1.56E-04	3.99E-05
	19.2-41	1.26E-08	1.57E-04	1.56E-05	1.57E-04	1.56E-04	4.06E-05
	41-78.6	1.26E-08	1.57E-04	1.56E-05	1.57E-04	1.56E-04	4.15E-05
	78.6-120	1.26E-08	1.57E-04	1.56E-05	1.57E-04	1.56E-04	4.39E-05
	120-175	1.26E-08	1.57E-04	1.56E-05	1.57E-04	1.56E-04	4.81E-05
TC2	0-19.2	1.26E-08	1.88E-04	1.87E-05	1.88E-04	1.88E-04	3.99E-05
	19.2-41	1.26E-08	1.88E-04	1.87E-05	1.88E-04	1.88E-04	4.06E-05
	41-78.6	1.26E-08	1.88E-04	1.87E-05	1.88E-04	1.88E-04	4.15E-05
	78.6-120	1.26E-08	1.88E-04	1.87E-05	1.88E-04	1.88E-04	4.39E-05
	120-175	1.26E-08	1.88E-04	1.87E-05	1.88E-04	1.88E-04	4.81E-05
TC3	0-19.2	1.16E-08	1.98E-04	1.98E-05	1.98E-04	1.98E-04	4.02E-05
	19.2-41	1.16E-08	1.98E-04	1.98E-05	1.98E-04	1.98E-04	4.08E-05
	0-19.2	1.16E-08	2.09E-04	2.08E-05	2.09E-04	2.09E-04	4.10E-05
TC4	19.2-41	1.16E-08	2.09E-04	2.08E-05	2.09E-04	2.09E-04	4.16E-05
	41-78.6	1.16E-08	2.09E-04	2.08E-05	2.09E-04	2.09E-04	4.35E-05
	78.6-120	1.16E-08	2.09E-04	2.08E-05	2.09E-04	2.09E-04	4.89E-05
	120-175	1.16E-08	2.09E-04	2.08E-05	2.09E-04	2.09E-04	5.60E-05
	0-19.2	1.47E-08	2.19E-04	2.19E-05	2.19E-04	2.19E-04	4.72E-05
TC5	19.2-41	1.47E-08	2.19E-04	2.19E-05	2.19E-04	2.19E-04	4.74E-05
	41-78.6	1.47E-08	2.19E-04	2.19E-05	2.19E-04	2.19E-04	4.84E-05
	78.6-120	1.47E-08	2.19E-04	2.19E-05	2.19E-04	2.19E-04	5.16E-05
	120-175	1.47E-08	2.19E-04	2.19E-05	2.19E-04	2.19E-04	5.56E-05
	0-19.2	2.10E-08	1.67E-04	1.67E-05	1.67E-04	1.67E-04	5.35E-05
TC6	19.2-41	2.10E-08	1.67E-04	1.67E-05	1.67E-04	1.67E-04	5.37E-05
	41-78.6	2.10E-08	1.67E-04	1.67E-05	1.67E-04	1.67E-04	5.48E-05
	78.6-120	2.10E-08	1.67E-04	1.67E-05	1.67E-04	1.67E-04	5.72E-05
	120-175	2.00E-08	1.67E-04	1.67E-05	1.67E-04	1.67E-04	6.12E-05
	AP1	0-19.2	1.16E-08	2.30E-04	2.29E-05	2.30E-04	2.30E-04
AP2	0-19.2	1.05E-08	2.30E-04	2.29E-05	2.30E-04	2.30E-04	4.19E-05
AP3	0-19.2	1.47E-08	1.98E-04	1.98E-05	1.98E-04	1.98E-04	4.98E-05
AP4	0-19.2	8.41E-09	2.40E-04	2.39E-05	2.40E-04	2.40E-04	3.38E-05
	19.2-41	8.41E-09	2.40E-04	2.39E-05	2.40E-04	2.40E-04	3.38E-05
	41-78.6	8.41E-09	2.40E-04	2.39E-05	2.40E-04	2.40E-04	3.40E-05
	78.6-120	8.41E-09	2.40E-04	2.39E-05	2.40E-04	2.40E-04	3.43E-05
	AP5	0-19.2	8.41E-09	2.09E-04	2.08E-05	2.09E-04	2.09E-04
AP6	0-19.2	8.41E-09	2.09E-04	2.08E-05	2.09E-04	2.09E-04	3.74E-05
AP7	0-19.2	8.41E-09	2.09E-04	2.08E-05	2.09E-04	2.09E-04	3.15E-05
	19.2-41	8.41E-09	2.09E-04	2.08E-05	2.09E-04	2.09E-04	3.17E-05
	41-78.6	8.41E-09	2.09E-04	2.08E-05	2.09E-04	2.09E-04	3.20E-05
AP8	0-19.2	8.41E-09	2.30E-04	2.29E-05	2.30E-04	2.30E-04	3.66E-05
KT1	0-19.2	6.31E-09	1.46E-04	1.46E-05	1.46E-04	1.46E-04	1.82E-05
	19.2-41	6.31E-09	1.46E-04	1.46E-05	1.46E-04	1.46E-04	1.82E-05
	41-78.6	6.31E-09	1.46E-04	1.46E-05	1.46E-04	1.46E-04	1.82E-05
	78.6-120	6.31E-09	1.46E-04	1.46E-05	1.46E-04	1.46E-04	1.83E-05
	0-19.2	6.31E-09	1.46E-04	1.46E-05	1.46E-04	1.46E-04	1.82E-05
KT2	19.2-41	6.31E-09	1.46E-04	1.46E-05	1.46E-04	1.46E-04	1.82E-05
	41-78.6	6.31E-09	1.46E-04	1.46E-05	1.46E-04	1.46E-04	1.82E-05
	78.6-120	6.31E-09	1.46E-04	1.46E-05	1.46E-04	1.46E-04	1.83E-05
	0-19.2	6.31E-09	1.46E-04	1.46E-05	1.46E-04	1.46E-04	1.90E-05
KT3	19.2-41	6.31E-09	1.46E-04	1.46E-05	1.46E-04	1.46E-04	1.90E-05
	0-19.2	6.31E-09	1.46E-04	1.46E-05	1.46E-04	1.46E-04	1.82E-05
KT5	0-19.2	4.20E-09	1.04E-04	1.04E-05	1.04E-04	1.04E-04	1.48E-05
	19.2-41	4.20E-09	1.04E-04	1.04E-05	1.04E-04	1.04E-04	1.48E-05
	41-78.6	4.20E-09	1.04E-04	1.04E-05	1.04E-04	1.04E-04	1.48E-05
	78.6-120	4.20E-09	1.04E-04	1.04E-05	1.04E-04	1.04E-04	1.49E-05
	120-175	4.20E-09	1.04E-04	1.04E-05	1.04E-04	1.04E-04	1.50E-05
KT6	0-19.2	5.26E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	1.74E-05
	19.2-41	5.26E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	1.74E-05
KT7	0-19.2	3.15E-09	7.31E-05	7.29E-06	7.31E-05	7.30E-05	1.27E-05
	0-19.2	3.15E-09	6.26E-05	6.25E-06	6.27E-05	6.26E-05	1.19E-05
KT8	19.2-41	3.15E-09	6.26E-05	6.25E-06	6.27E-05	6.26E-05	1.19E-05
	41-78.6	3.15E-09	6.26E-05	6.25E-06	6.27E-05	6.26E-05	1.19E-05
	78.6-120	3.15E-09	6.26E-05	6.25E-06	6.27E-05	6.26E-05	1.19E-05
	0-19.2	3.15E-09	6.26E-05	6.25E-06	6.27E-05	6.26E-05	1.19E-05

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

ASR	Assessment Height (mAG)	Antimony	Arsenic	Cadmium	Chromium	Lead	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	1.89E-04
TT2	0-19.2	8.94E-07	3.13E-05	3.13E-06	3.16E-05	3.13E-05	6.79E-04
TT3	0-19.2	1.38E-06	5.22E-05	5.21E-06	5.27E-05	5.22E-05	1.05E-03
TT4	0-19.2	7.81E-06	2.27E-10	2.88E-08	2.66E-06	3.99E-07	5.89E-03
TT5	0-19.2	3.77E-07	2.71E-04	2.71E-05	2.72E-04	2.71E-04	3.23E-04
TT6	0-19.2	1.33E-06	1.46E-04	1.46E-05	1.47E-04	1.46E-04	1.02E-03
TM1	0-19.2	3.99E-08	1.04E-05	1.04E-06	1.05E-05	1.04E-05	3.66E-05
	19.2-41	3.99E-08	1.04E-05	1.04E-06	1.05E-05	1.04E-05	3.67E-05
	41-78.6	3.99E-08	1.04E-05	1.04E-06	1.05E-05	1.04E-05	3.71E-05
	78.6-120	3.99E-08	1.04E-05	1.04E-06	1.05E-05	1.04E-05	3.78E-05
	120-175	3.99E-08	1.04E-05	1.04E-06	1.05E-05	1.04E-05	3.85E-05
TM2	0-19.2	2.63E-08	3.13E-05	3.12E-06	3.13E-05	3.13E-05	2.94E-05
	19.2-41	2.63E-08	3.13E-05	3.12E-06	3.13E-05	3.13E-05	2.96E-05
	41-78.6	2.63E-08	3.13E-05	3.12E-06	3.13E-05	3.13E-05	2.99E-05
	78.6-120	2.63E-08	3.13E-05	3.12E-06	3.13E-05	3.13E-05	3.06E-05
	120-175	2.63E-08	3.13E-05	3.12E-06	3.13E-05	3.13E-05	3.13E-05
TM3	0-19.2	4.41E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	4.21E-05
	19.2-41	4.41E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	4.23E-05
	41-78.6	4.41E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	4.26E-05
	78.6-120	4.41E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	4.33E-05
	120-175	4.41E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	4.40E-05
TM4	0-19.2	4.73E-08	3.13E-05	3.12E-06	3.13E-05	3.13E-05	4.45E-05
	19.2-41	4.73E-08	3.13E-05	3.12E-06	3.13E-05	3.13E-05	4.47E-05
	41-78.6	4.73E-08	3.13E-05	3.12E-06	3.13E-05	3.13E-05	4.50E-05
	78.6-120	4.73E-08	3.13E-05	3.12E-06	3.13E-05	3.13E-05	4.57E-05
	120-175	4.73E-08	3.13E-05	3.12E-06	3.13E-05	3.13E-05	4.64E-05
TM5	0-19.2	5.36E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	4.86E-05
	19.2-41	5.36E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	4.88E-05
	41-78.6	5.36E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	4.93E-05
	78.6-120	5.36E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	5.07E-05
	120-175	5.36E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	5.21E-05
TM6	0-19.2	7.04E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	6.07E-05
	19.2-41	7.04E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	6.09E-05
	41-78.6	7.04E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	6.15E-05
	78.6-120	7.04E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	6.28E-05
	120-175	7.04E-08	2.09E-05	2.08E-06	2.09E-05	2.09E-05	6.41E-05
SLW1	0-19.2	7.57E-08	7.10E-04	7.08E-05	7.10E-04	7.09E-04	1.36E-04
SLW2	0-19.2	8.09E-08	7.93E-04	7.91E-05	7.94E-04	7.93E-04	1.47E-04
SLW3	0-19.2	2.94E-08	3.76E-04	3.75E-05	3.75E-04	3.76E-04	8.16E-05
TC1	0-19.2	2.00E-08	2.82E-04	2.81E-05	2.82E-04	2.82E-04	5.80E-05
	19.2-41	2.00E-08	2.82E-04	2.81E-05	2.82E-04	2.82E-04	5.87E-05
	41-78.6	2.00E-08	2.82E-04	2.81E-05	2.82E-04	2.82E-04	5.98E-05
	78.6-120	2.00E-08	2.82E-04	2.81E-05	2.82E-04	2.82E-04	6.21E-05
	120-175	2.00E-08	2.82E-04	2.81E-05	2.82E-04	2.82E-04	6.43E-05
TC2	0-19.2	1.68E-08	3.03E-04	3.02E-05	3.03E-04	3.03E-04	5.49E-05
	19.2-41	1.68E-08	3.03E-04	3.02E-05	3.03E-04	3.03E-04	5.55E-05
	41-78.6	1.68E-08	3.03E-04	3.02E-05	3.03E-04	3.03E-04	5.66E-05
	78.6-120	1.68E-08	3.03E-04	3.02E-05	3.03E-04	3.03E-04	5.89E-05
	120-175	1.68E-08	3.03E-04	3.02E-05	3.03E-04	3.03E-04	6.32E-05
TC3	0-19.2	1.47E-08	3.03E-04	3.02E-05	3.03E-04	3.03E-04	5.38E-05
	19.2-41	1.47E-08	3.03E-04	3.02E-05	3.03E-04	3.03E-04	5.43E-05
	41-78.6	1.47E-08	3.03E-04	3.02E-05	3.03E-04	3.03E-04	5.46E-05
	78.6-120	1.47E-08	3.03E-04	3.02E-05	3.03E-04	3.03E-04	5.51E-05
	120-175	1.47E-08	3.03E-04	3.02E-05	3.03E-04	3.03E-04	5.57E-05
TC4	0-19.2	1.58E-08	3.24E-04	3.23E-05	3.24E-04	3.23E-04	5.74E-05
	19.2-41	1.58E-08	3.24E-04	3.23E-05	3.24E-04	3.23E-04	5.74E-05
	41-78.6	1.58E-08	3.24E-04	3.23E-05	3.24E-04	3.23E-04	5.74E-05
	78.6-120	1.58E-08	3.24E-04	3.23E-05	3.24E-04	3.23E-04	5.74E-05
	120-175	1.58E-08	3.24E-04	3.23E-05	3.24E-04	3.23E-04	5.74E-05
TC5	0-19.2	2.10E-08	3.44E-04	3.43E-05	3.44E-04	3.44E-04	6.27E-05
	19.2-41	2.10E-08	3.44E-04	3.43E-05	3.44E-04	3.44E-04	6.30E-05
	41-78.6	2.10E-08	3.44E-04	3.43E-05	3.44E-04	3.44E-04	6.45E-05
	78.6-120	2.10E-08	3.44E-04	3.43E-05	3.44E-04	3.44E-04	6.84E-05
	120-175	2.10E-08	3.44E-04	3.43E-05	3.44E-04	3.44E-04	7.35E-05
TC6	0-19.2	2.73E-08	2.92E-04	2.91E-05	2.92E-04	2.92E-04	6.90E-05
	19.2-41	2.73E-08	2.92E-04	2.91E-05	2.92E-04	2.92E-04	6.93E-05
	41-78.6	2.73E-08	2.92E-04	2.91E-05	2.92E-04	2.92E-04	7.09E-05
	78.6-120	2.73E-08	2.92E-04	2.91E-05	2.92E-04	2.92E-04	7.39E-05
	120-175	2.73E-08	2.92E-04	2.91E-05	2.92E-04	2.92E-04	7.90E-05
AP1	0-19.2	2.10E-08	3.65E-04	3.64E-05	3.66E-04	3.65E-04	6.23E-05
AP2	0-19.2	2.31E-08	3.65E-04	3.64E-05	3.66E-04	3.65E-04	6.31E-05
AP3	0-19.2	3.68E-08	3.97E-04	3.96E-05	3.97E-04	3.96E-04	8.95E-05
AP4	0-19.2	1.89E-08	3.34E-04	3.33E-05	3.34E-04	3.34E-04	5.32E-05
	19.2-41	1.89E-08	3.34E-04	3.33E-05	3.34E-04	3.34E-04	5.32E-05
	41-78.6	1.89E-08	3.34E-04	3.33E-05	3.34E-04	3.34E-04	5.35E-05
	78.6-120	1.89E-08	3.34E-04	3.33E-05	3.34E-04	3.34E-04	5.40E-05
	120-175	1.89E-08	3.34E-04	3.33E-05	3.34E-04	3.34E-04	5.45E-05
AP5	0-19.2	1.58E-08	3.03E-04	3.02E-05	3.03E-04	3.03E-04	5.24E-05
AP6	0-19.2	1.47E-08	3.03E-04	3.02E-05	3.03E-04	3.03E-04	5.55E-05
AP7	0-19.2	1.89E-08	3.13E-04	3.12E-05	3.13E-04	3.13E-04	5.22E-05
	19.2-41	1.89E-08	3.13E-04	3.12E-05	3.13E-04	3.13E-04	5.23E-05
	41-78.6	1.89E-08	3.13E-04	3.12E-05	3.13E-04	3.13E-04	5.26E-05
AP8	0-19.2	1.58E-08	3.24E-04	3.23E-05	3.24E-04	3.23E-04	5.47E-05
	19.2-41	1.58E-08	3.24E-04	3.23E-05	3.24E-04	3.23E-04	5.47E-05
	41-78.6	1.58E-08	3.24E-04	3.23E-05	3.24E-04	3.23E-04	5.47E-05
	78.6-120	1.58E-08	3.24E-04	3.23E-05	3.24E-04	3.23E-04	5.47E-05
	120-175	1.58E-08	3.24E-04	3.23E-05	3.24E-04	3.23E-04	5.47E-05
KT1	0-19.2	7.36E-09	1.98E-04	1.98E-05	1.98E-04	1.98E-04	2.30E-05
	19.2-41	7.36E-09	1.98E-04	1.98E-05	1.98E-04	1.98E-04	2.31E-05
	41-78.6	7.36E-09	1.98E-04	1.98E-05	1.98E-04	1.98E-04	2.31E-05
	78.6-120	7.36E-09	1.98E-04	1.98E-05	1.98E-04	1.98E-04	2.32E-05
	120-175	7.36E-09	1.98E-04	1.98E-05	1.98E-04	1.98E-04	2.32E-05
KT2	0-19.2	7.36E-09	1.88E-04	1.87E-05	1.88E-04	1.88E-04	2.30E-05
	19.2-41	7.36E-09	1.88E-04	1.87E-05	1.88E-04	1.88E-04	2.31E-05
	41-78.6	7.36E-09	1.88E-04	1.87E-05	1.88E-04	1.88E-04	2.31E-05
	78.6-120	7.36E-09	1.88E-04	1.87E-05	1.88E-04	1.88E-04	2.32E-05
	120-175	7.36E-09	1.88E-04	1.87E-05	1.88E-04	1.88E-04	2.32E-05
KT3	0-19.2	7.36E-09	1.88E-04	1.87E-05	1.88E-04	1.88E-04	2.38E-05
	19.2-41	7.36E-09	1.88E-04	1.87E-05	1.88E-04	1.88E-04	2.39E-05
	41-78.6	7.36E-09	1.88E-04	1.87E-05	1.88E-04	1.88E-04	2.39E-05
KT4	0-19.2	7.36E-09	1.88E-04	1.87E-05	1.88E-04	1.88E-04	2.30E-05
	19.2-41	7.36E-09	1.88E-04	1.87E-05	1.88E-04	1.88E-04	2.30E-05
	41-78.6	7.36E-09	1.88E-04	1.87E-05	1.88E-04	1.88E-04	2.30E-05
	78.6-120	7.36E-09	1.88E-04	1.87E-05	1.88E-04	1.88E-04	2.30E-05
	120-175	7.36E-09	1.88E-04	1.87E-05	1.88E-04	1.88E-04	2.30E-05
KT5	0-19.2	5.26E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	1.96E-05
	19.2-41	5.26E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	1.97E-05
	41-78.6	5.26E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	1.97E-05
	78.6-120	5.26E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	1.98E-05
	120-175	5.26E-09	1.25E-04	1.25E-05	1.25E-04	1.25E-04	2.01E-05
KT6	0-19.2	6.31E-09	1.57E-04	1.56E-05	1.57E-04	1.56E-04	2.22E-05
	19.2-41	6.31E-09	1.57E-04	1.56E-05	1.57E-04	1.56E-04	2.23E-05
	41-78.6	6.31E-09	1.57E-04	1.56E-05	1.57E-04	1.56E-04	2.23E-05
KT7	0-19.2	4.20E-09	9.39E-05	9.37E-06	9.40E-05	9.39E-05	1.72E-05
	19.2-41	3.15E-09	8.35E-05	8.33E-06	8.36E-05	8.35E-05	1.56E-05
	41-78.6	3.15E-09	8.35E-05	8.33E-06	8.36E-05	8.35E-05	1.56E-05
KT8	0-19.2	3.15E-09	8.35E-05	8.33E-06	8.36E-05	8.35E-05	1.56E-05
	19.2-41	3.15E-09	8.35E-05	8.33E-06	8.36E-05	8.35E-05	1.56E-05
	41-78.6	3.15E-09	8.35E-05	8.33E-06	8.36E-05	8.35E-05	1.56E-05
SKC1	0-19.2	1.93E-06	1.04E-05	1.05E-06	1.11E-05	1.05E-05	1.45E-03
SL1	0-19.2	6.10E-08	2.09E-04	2.08E-05	2.09E-04	2.09E-04	8.13E-05
SL2	0-19.2	8.09E-08	3.13E-04	3.12E-05	3.13E-04	3.13E-04	9.65E-05