



ANALYTICAL PARAMETERS

Analytical Conditions		Reanalysis conditions				Multi-Standards Setting				
R1 volume	80	Serum reac. Smp. Vol (µL)		0		Formula Points	Quadratic	5	Axis conv.	No converte
R2 volume	0	Serum dilut. method. Vol (µL)		None						
R3 volume	20	Serum reac. Smp. Vol (d)		0			FV	Dil method	dil smp vol	Dil Vol
R4 volume	0	Serum dilut. method. Vol (d)		None		BLK	0.00	None	0	0
R1 diluent volume	0	Standards setting				1	*x0.125	None	0	0
R2 diluent volume	0	BLK H		9.9999		2	*x0.125	None	0	0
R3 diluent volume	0	BLK L		9.9999		3	*x0.5	None	0	0
R4 diluent volume	0	STD H		9.9999		4	*x0.75	None	0	0
Serum reac. s.volume	3,5	STD L		9.9999		5	*	None	0	0
Serum diluent method	Standard	FV		9.9999						
		Abnml. (serum) H		9.9999						
Reaction Time	10 min.	Abnml. (serum) L		9.9999						
Reagent 1 stir.	Weak	Calculation method settings				Reation rate method				
Reagent 2 stir.	Weak	M-DET. P.I	0	Prozone		Cycle				3
Reagent 3 stir.	Weak	M-DET. P.m	95	Prozone form	None	Factor				3.0
Reagent 4 stir.	Weak	M-DET. P.n	98	Prozone limit	9999	Reac. Type				INC
Sub-analy. conditions		M-DET. P.p	0	Prozone judge	Upper Limit	E2. corre.				Not dot
Name	FR	M-DET. P.r	0	Judge limit	9999	Blank (µL)				9.9999
Digits	2			M-DET.P.m	0	Blank (d)				-9.999
M-wave. L	658	Check D.P.I	0	M-DET.P.n	0	Sample (µL)				9.9999
S-wave. L.	-	Limit value	0.00	S-DET. P.p	0	Sample (d)				-9.999
Analy. mthd	EPA	Variance	3	S-DET.P.r	0	Endpoint method				
Calc. mtdh	MSTD		10			Re. Absorb (µL)				9.9999
Qualit. judg	Not do					Re. Absorb (d)				-9.999