

# Axis-Shield Liquid Stable 2-Part Homocysteine Reagent

## ASSAY PROTOCOL – SIEMENS ADVIA 1650

Type #	*	M-DET.P.1	0	Dil. Sample Vol 1
R1 volume (µL)	150	M-DET.P.m	60	Dil. Sample Vol 2
R2 volume	15	M-DET.p.n	85	Dil. Sample Vol 3
R1 diluent volume	0	S-DET.P.p	0	Dil. Sample Vol 4
R2 diluent volume	0	S-DET.P.r	0	Dil. Sample Vol 5
Serum reac s vol (µL)	36	Check D.P.l	0	Diluent Vol 1
Serum dil. s. vol (µL)	20	Cycle	3	Diluent Vol 2
Serum dil. volume (µL)	60	Factor	3.0	Diluent Vol 3
Serum dil. posit	0	Limit Value	0.000	Diluent Vol 4
Serum dil. method	Special	Variance	10.0	Diluent Vol 5
Urine reac. s vol		Reaction Type	Dec.	Diluent pos. 1
Urine dil. volume		E2 correction	Not do	Diluent pos. 2
Urine dil. posit		Blank u	9,9999	Diluent pos. 3
Urine dil. method		Blank d	-9,999	Diluent pos. 4
Reaction Time	10 min	Sample u	9,9999	Diluent pos. 5
Reagent 1 stir	Weak	Sample d	-9,999	STD BLK (H)
Reagent 2 stir	Strong	Re. Absorb (u)	9,9999	STD BLK (L)
Digits	0	Re. Absorb (d)	-9,999	STD 1 (H)
M-wave. L.	340	Prozone form.	None	STD 1 (L)
S-wave, L.	410	Prozone limit	9,999	STD 2 (H)
Analy mthd	RRA	Prozone judge	Upper Limit	STD 2 (L)
Calc. Mthd	STD	M-DET, P.m (prozone)	0	STD 3 (H)
Quality Judge	Not do	M-DET, P.n (prozone)	0	STD 3 (L)
Blank H	9,999999	S-DET,P.p (prozone)	0	STD 4 (H)
Blank L	-9,9999	S-DET,P.r (prozone)	0	STD 4 (L)
STD H	9,99999	Fornula		STD 5 (H)
STD L	-9,9999	Axis convert		STD 5 (L)
Abnml(serum)H(µmol/L)	*	FV1		
Abnml(serum)L(µmol/L)	*	FV2		
FV(µmol/L)	28.0	FV3		
		FV4		
		FV5		
		Dilute method 1		
		Dilute method 2		
		Dilute method 3		
		Dilute method 4		
		Dilute method 5		

\* User Defined

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<b>Rerun Conditions</b>	
Serum reac. s. vol (u)	3.00
Serum dilute. Method (u)	None
Serum dil.SMP. Vol (u)	0.00
Serum diluent vol (u)	0.00
Serum diluent posi (u)	0
Serum reac. s. vol (d)	3.00
Serum dilute. Method (d)	None
Serum dil. smp. Vol (d)	0.00
Serum diluent vol (d)	0.00
Serum diluent posi (d)	0
Variance (*)	A mark exit, No rerun.
Absorbance (U)	A mark exit, No rerun.
Absorbance (D)	A mark exit, No rerun.
Absorbance limit(u)	A mark exit, No rerun.
Absorbance limit (d)	A mark exit, No rerun.
Cell blank (N)	A mark exit, No rerun.
Absorbance v. limit (H)	A mark exit, No rerun.
Absorbance v. limit (L)	A mark exit, No rerun.
Normal v.limit (h)	A mark exit, No rerun.
Normal v. limit (l)	A mark exit, No rerun.
Reagent shortage	A mark exit, No rerun.
Overflow (/)	A mark exit, No rerun.
Safety (S)	No mark, No rerun.
Prozone (P)	A mark exit, No rerun.
Effect nbr.o.pnts (n)	A mark exit, No rerun.
Calibration (C)	A mark exit, No rerun.
Reanalysis (R)	A mark exit, No rerun.

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