



## HOMOGENEOUS ENZYMIC HOMOCYSTEINE REAGENT

### Synchron® DX User Defined Parameters:

#### INSTRUMENT PARAMETERS:

Test Name:	User Defined	Initial	
Reaction Type:	RATE 1	Start Read:	281 sec
		End Read:	296 sec
		Reaction 1	
Units:	µmol/L	Start Read:	357 sec.
Decimal Precision	X.X	End Read:	632 sec.
Reaction Direction:	NEGATIVE		
Calculation Factor:	1.0	Reaction 2	
Math Model:	Linear	Start Read:	n/a
Cal. Time Limit	120 hrs.	End Read:	n/a
Number of Calibrators	2		
#1	0.0	USABLE RESULT RANGE:	
#2	*	Lower Limit	0.7
Primary Wavelength	340 nm	Upper Limit:	50
Secondary Wavelength	380 nm		
Sample Volume:	25 µl	ERROR DETECTION LIMITS:	
		Reagent Blank/Blank	
REAGENTS:		ABS Low Limit:	-1.500
Primary Inject (first)/First Inject		ABS High Limit:	2.200
Compartment/Component:	A	Rate Low Limit:	2.200
Volume/Dispense Volume:	179 µl	Rate High Limit:	-1.500
Add Time/Inject Time:	n/a	Mean Deviation:	2.200
Primary Inject (second)/Second Inject		Reaction /Reaction 1	
Compartment/Component	B	ABS Low Limit:	-1.500
Volume/Dispense Volume:	22 µl	ABS High Limit:	2.200
Add Time/Inject Time:	-180	Rate Low Limit:	2.200
Secondary Inject/Third Inject		Rate High Limit:	-1.500
Compartment/Component:	C	Mean Deviation	2.200
Volume/Dispense Volume:	14 µl		
Add Time/Inject Time	272 sec	SUBSTRATE DEPLETION:	
Blank		Initial Rate:	- 0.060
Start Read:	- 50 sec	Delta ABS:	2.200
End Read:	- 10 sec	MULTI POINT SPAN:	1-2: - 0.001**

\* Value on Calibrator D vial, \*\* Note the minus sign for this parameter.

RESULTS: Results are printed out by the Synchron DX in µmol/L.

Note: Samples with values greater than 50 µmol/L should be diluted 1:3 with saline and rerun. Multiply results by 3.