

# IMMUNOGLOBULIN M (IgM)

Turbidimetry

The application parameters comprised here constitute a guide to facilitate the validation of our reagents by the instrument. It is advisable to validate the use when there is any change in software or reagent versions.

## Instrument: **ADVIA 1650**

### Reagent preparation

Reagent is ready to use.

### Instrument settings

ANALYTICAL PARAMETERS	Reanalysis conditions	Multi-Standards setting			
<u>Analytical Conditions</u>	Serum reac. smp. vol. (μ) <b>0.00</b>	Formula	Logit Log 3	Axis conv.	No convert.
R1 volume <b>90.00</b>	Serum dilut. method (μ) <b>None</b>	Points	6		
R2 volume <b>0.00</b>	Serum reac. smp. vol. (d) <b>0.00</b>	FV	Dil.Method	Dil.smp.vol	Diluent vol
R3 volume <b>0.00</b>	Serum dilut. method (d) <b>None</b>	BLK			STD-H
R4 volume <b>0.00</b>		1    * <b>Special</b>			STD-L
R1 diluent vol. <b>0.00</b>	<u>Standards setting</u>	2    * <b>Special</b>			
R2 diluent vol. <b>0.00</b>	BLK H <b>9.9999</b>	3    * <b>Special</b>			
R3 diluent vol. <b>0.00</b>	BLK L <b>-9.9999</b>	4    * <b>Special</b>			
R4 diluent vol. <b>0.00</b>	STD H <b>9.9999</b>	5    * <b>Special</b>			
Serum reac. s. vol. <b>7.00</b>	STD L <b>-9.9999</b>				
Serum dil. method <b>Standard</b>	FV <b>0.0000</b>				
	Abnml. (serum) H <b>99999</b>	* assigned value			
Reaction time <b>10 min.</b>	Abnml. (serum) L <b>-9999</b>				
Reagent 1 stir. <b>Weak</b>					
Reagent 2 stir. <b>Weak</b>					
Reagent 3 stir. <b>Weak</b>					
Reagent 4 stir. <b>Weak</b>					
<u>Sub-analy. conditions</u>	<u>Calculation method setting</u>				<u>Reaction rate method</u>
Name <b>IG M</b>	M-DET. P. l <b>0</b>	<b>Prozone</b>			Cycle <b>3</b>
Digits <b>2</b>	M-DET. P. m <b>96</b>	Prozone form.	<b>None</b>		Factor <b>3.0</b>
M-wave. L. <b>340 nm</b>	M-DET. P. n <b>98</b>	Prozone limit	<b>9.999</b>		Reac. Type <b>Inc.</b>
S-wave. L. <b>EPA</b>	S-DET. P. p <b>0</b>	Prozone judge	<b>Upper limit</b>		E2 corre. <b>Not do</b>
Analy. mthd. <b>MSTD</b>	S-DET. P. r <b>0</b>	Judge limit	<b>9.999</b>		Blank (μ) <b>9.9999</b>
Calc. mthd <b>Not do</b>	Check D.P.I. <b>0</b>	M-DET. P. m.	<b>0</b>		Blank (d) <b>-9.999</b>
Qualit. judg.	Limit value <b>0.003</b>	M-DET. P. n.	<b>0</b>		Sample (μ) <b>9.9999</b>
	Variance <b>10.0</b>	S-DET. P. p.	<b>0</b>		Sample (d) <b>-9.999</b>
		S-DET. P. r.	<b>0</b>		
					<u>Endpoint method</u>
					Re. Absorb (μ) <b>9.9999</b>
					Re. Absorb (d) <b>-9.999</b>