

ARCHITECT INSTRUMENT SETTING

Apolipoprotein A1 - ApoA (BSBE)

Method : Immunoturbidimetric

<input checked="" type="radio"/> Genral	<input type="radio"/> Calibration	<input type="radio"/> Smartwash	<input type="radio"/> Result	<input type="radio"/> Interpretation	
Assay :	<input type="text" value="APOA"/>	Type :	<input type="text" value="Photometric"/>	Version :	<input type="text" value="1"/>
Number :	<input type="text"/>	Assay :	<input type="text" value="Enabled"/>	Date :	<input type="text"/>
				Time :	<input type="text"/>
				Operator :	<input type="text" value="ADMIN"/>

<input checked="" type="radio"/> Reaction Defination	<input type="radio"/> Reagent / Sample	<input type="radio"/> Vallidity Check					
Reaction mode :	<input type="text" value="End up"/>	Read Time					
Wavelength	<table style="width: 100%; border: none;"> <tr> <td style="text-align: center;">Primary</td> <td style="text-align: center;">Secondary</td> </tr> <tr> <td style="text-align: center;"><input type="text" value="340"/></td> <td style="text-align: center;"><input type="text" value="700"/></td> </tr> </table>	Primary	Secondary	<input type="text" value="340"/>	<input type="text" value="700"/>	Main :	<input type="text" value="33 - 33"/>
Primary	Secondary						
<input type="text" value="340"/>	<input type="text" value="700"/>						
Last Required read	<input type="text" value="33"/>	Flex :	<input type="text" value="0-0"/>				
Absorbance range	<input type="text" value="0.0000 - 3.0000"/>	Color correction :	<input type="text" value="0-0"/>				
Sample blank type	<input type="text" value="Self"/>	Blank :	<input type="text" value="10 - 10"/>				

<input type="radio"/> Reaction Defination	<input checked="" type="radio"/> Reagent / Sample	<input type="radio"/> Vallidity Check					
Reagent :	<input type="text" value="APOA"/>	Reagent Volume :	<table style="width: 100%; border: none;"> <tr> <td style="text-align: center;">R1</td> <td style="text-align: center;">R2</td> </tr> <tr> <td style="text-align: center;"><input type="text" value="175"/></td> <td style="text-align: center;"><input type="text" value="58"/></td> </tr> </table>	R1	R2	<input type="text" value="175"/>	<input type="text" value="58"/>
R1	R2						
<input type="text" value="175"/>	<input type="text" value="58"/>						
Diluent	<input type="text" value="Saline"/>	Water volume :	<input type="text" value="0"/>				
Diluent dispense mode :	<input type="text" value="Type0"/>	Dispense mode :	<table style="width: 100%; border: none;"> <tr> <td style="text-align: center;">Type0</td> <td style="text-align: center;">Type0</td> </tr> </table>	Type0	Type0		
Type0	Type0						
Dilution name	Sample	Diluted Sample	Diluent	Water	Dilution Factor	Default dilution	
STANDARD	<input type="text" value="1.6"/>	<input type="text" value="0"/>	<input type="text" value="0"/>	<input type="text" value="0"/>	<input type="text" value="1:0"/>	<input checked="" type="radio"/>	
1:4	<input type="text" value="25"/>	<input type="text" value="2"/>	<input type="text" value="75"/>	<input type="text" value="0"/>	<input type="text" value="1:4"/>	<input type="radio"/>	

<input type="radio"/> Reaction Defination	<input type="radio"/> Reagent / Sample	<input checked="" type="radio"/> Vallidity Check	
Reaction check type :	<input type="text" value="None"/>	Read time B range :	<input type="text"/>
Read time A range :	<input type="text"/>	Minimum absorbance :	<input type="text"/>
Calculation limit :	<input type="text"/>	Rate linearity % :	<input type="text" value="0"/>

