

## ARCHITECT c8000 new Type/ c16000

### NEFA-HR(2)

<b>● General</b> Assay: <input type="text" value="NEFA-HR(2)"/> Type: <input type="text" value="Photometric"/>					
<b>● Reaction definition</b> Reaction mode: <input type="text" value="End up"/>					
Wavelength: <input type="text" value="548"/> / <input type="text" value="660"/>		Read times Main: <input type="text" value="31"/> - <input type="text" value="33"/>			
Last required read: <input type="text" value="33"/>		Flex: <input type="text"/> - <input type="text"/>			
Absorbance range: <input type="text" value="0.1000"/> - <input type="text" value="3.0000"/>		Color correction: <input type="text"/> - <input type="text"/>			
Sample blank type: <input type="text" value="self"/>		Blank: <input type="text" value="14"/> - <input type="text" value="16"/>			
<b>● Reagent/Sample</b>					
Reagent: <input type="text" value="NEFA-HR(2)"/>		R1 Reagent volume: <input type="text" value="180"/>	R2 Reagent volume: <input type="text" value="90"/>		
Diluent: <input type="text" value="saline"/>		Water volume: <input type="text"/>	Water volume: <input type="text"/>		
Diluent dispense mode: <input type="text" value="Type 0"/>		Dispense mode: <input type="text" value="Type 0"/>	Dispense mode: <input type="text" value="Type 0"/>		
Dilution name: <input type="text" value="NORMAL"/>		Sample: <input type="text" value="4.0"/>	sample: <input type="text"/>	Diluent: <input type="text"/>	
		Water: <input type="text"/>	Default dilution: <input checked="" type="radio"/>		
<b>● Validity checks</b> Reaction check: <input type="text" value="None"/>					
Maximum absorbance variation:					
<b>● Calibration</b> Assay: <input type="text" value="NEFA-HR(2)"/> Calibration method: <input type="text" value="Linear"/>					
<b>● Calibrators</b>					
Calibrator set: <input type="text" value="NEFA Standard"/>		Blank: <input type="text" value="Water"/>	Concentration: <input type="text" value="0,0"/>		
Replicates: <input type="text" value="3"/>		Cal 1: <input type="text" value="CAL"/>	Concentration: <input type="text" value="*1"/>		
		<input type="text"/>			
		<input type="text"/>			
		<input type="text"/>			
		<input type="text"/>			
<b>● Volumes</b> Calibrator: NEFA Standard					
Calibrator level		Sample	Diluted sample	Diluent	Water
Blank: Water		<input type="text" value="4.0"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>
Cal 1: CAL		<input type="text" value="4.0"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>
Cal 2: None					
Cal 3: None					
Cal 4: None					
Cal 5: None					
Cal 6: None					
<b>Result units</b> Result units: <input type="text" value="mmol/L"/> Correlation factor: <input type="text" value="1,0000"/>					
Decimal places: <input type="text" value="1"/>			Intercept: <input type="text" value="0,0000"/>		

\*1: input calibrator concentration, find on the vial

val UK:0608D1kg