

The NGAL Test™ Application Note for Roche Modular P

NGAL

 Code (ACN)

ANALYZE

 Ser/PI Sample type

 Assay/Time/Point

 Wave (2nd/Primary)

Sample Volume			
Normal	3,0	0,0	0
Decrease	15,0	3,0	105
Increase	6,0	0,0	0

Reagent Vol.	Diluent	BCN	OBS	Timing
R1	150	0	xxx	<input style="width: 50px;" type="text"/>
R2	<input style="width: 50px;" type="text"/>	<input style="width: 50px;" type="text"/>	<input style="width: 50px;" type="text"/>	<input style="width: 50px;" type="text" value="T2"/>
R3	50	0	xxx	<input style="width: 50px;" type="text" value="T3"/>
R4	<input style="width: 50px;" type="text"/>	<input style="width: 50px;" type="text"/>	<input style="width: 50px;" type="text"/>	<input style="width: 50px;" type="text"/>

Diluent	
<input type="radio"/> Water	Code <input style="width: 50px;" type="text"/>
<input checked="" type="radio"/> Diluent	OBS <input style="width: 50px;" type="text"/>
	(Saline) <input style="width: 50px;" type="text"/>

Abs. Limit	<input style="width: 50px;" type="text" value="32000"/>	Increase/Decrease	<input style="width: 50px;" type="text" value="Increase"/>	Upper/Lower	
Prozone Limit	<input style="width: 50px;" type="text" value="0"/>	<input style="width: 50px;" type="text" value="0"/>	<input style="width: 50px;" type="text" value="0"/>	<input style="width: 50px;" type="text" value="0"/>	<input style="width: 50px;" type="text" value="0"/>
Cell Detergent	<input style="width: 100%;" type="text" value="Detergent 1"/>				
Twin Test	<input style="width: 100%;" type="text"/>				

Reagent Bottle Size	
A	<input style="width: 50px;" type="text"/>
B	<input style="width: 50px;" type="text"/>
C	<input style="width: 50px;" type="text"/>
D	<input style="width: 50px;" type="text"/>
E	<input style="width: 50px;" type="text"/>

CALIBRATION

Calibration Type	<input style="width: 100%;" type="text" value="SPLINE"/>
Point	<input style="width: 50px;" type="text" value="6"/>
Span	<input style="width: 50px;" type="text" value="6"/>
Weight	<input style="width: 50px;" type="text" value="0"/>
Update Type	<input style="width: 50px;" type="text" value="none"/>
Isozyme Q Channel	<input style="width: 50px;" type="text"/>

SD Limit	<input style="width: 50px;" type="text" value="999"/>				
Duplicate Limit	<input style="width: 50px;" type="text" value="99"/>	%	<input style="width: 50px;" type="text" value="32000"/>	Abs	
Sensitivity Limit	<input style="width: 50px;" type="text" value="-99999"/>		<input style="width: 50px;" type="text" value="99999"/>		
S1-Abs Limit	<input style="width: 50px;" type="text" value="-32000"/>		<input style="width: 50px;" type="text" value="32000"/>		

Auto Calibration	
Timeout	
<input type="checkbox"/> Blank	<input style="width: 50px;" type="text"/>
<input type="checkbox"/> Span	<input style="width: 50px;" type="text"/>
<input type="checkbox"/> 2Point	<input style="width: 50px;" type="text"/>
<input type="checkbox"/> Full	<input style="width: 50px;" type="text"/>
Change Over	
<input type="checkbox"/> Module	<input style="width: 50px;" type="text"/>
<input checked="" type="checkbox"/> Lot	<input style="width: 50px;" type="text" value="full"/>
<input type="checkbox"/> Bottle	<input style="width: 50px;" type="text"/>

RANGE

App Code	<input type="text" value="xxx"/>	Factor	<input type="text"/>
Unit	<input type="text" value="ng/mL"/> Unit 2 <input type="text"/>	<input type="text"/>	<input type="text"/>
	Unit 3 <input type="text"/>	<input type="text"/>	<input type="text"/>
Report Name	<input type="text" value="NGAL"/>		
Data Mode	<input type="text" value="Active"/>		
Technical Limit	<input type="text" value="25"/>	<input type="text" value="5000"/>	
Repeat Limit	<input type="text" value="-99999"/>	<input type="text" value="999999"/>	
<input type="checkbox"/> Control Interval Time		<input type="text" value="0,0"/>	

Qualitative		
(1)	<input type="text" value="0"/>	<input type="text" value="--"/>
(2)	<input type="text" value="0"/>	<input type="text" value="-"/>
(3)	<input type="text" value="0"/>	<input type="text" value="+/-"/>
(4)	<input type="text" value="0"/>	<input type="text" value="++"/>
(5)	<input type="text" value="0"/>	<input type="text" value="+++"/>
(6)	<input type="text" value="0"/>	<input type="text" value="++++"/>

Expected Ranges

Male			
<input type="text" value="#"/>	<input type="text" value="Years"/>	<input type="text" value="#"/>	<input type="text" value="#"/>
<input type="text" value="#"/>	<input type="text" value="Years"/>	<input type="text" value="#"/>	<input type="text" value="#"/>
		<input type="text" value="#"/>	<input type="text" value="#"/>

Female			
<input type="text" value="#"/>	<input type="text" value="Years"/>	<input type="text" value="#"/>	<input type="text" value="#"/>
<input type="text" value="#"/>	<input type="text" value="Years"/>	<input type="text" value="#"/>	<input type="text" value="#"/>
		<input type="text" value="#"/>	<input type="text" value="#"/>

Default	
<input type="text" value="Sex"/>	<input type="text" value="Male"/> <input type="text" value="Female"/>
<input type="radio"/>	
Range	
<input type="text" value="Range 1"/>	<input type="text" value="Range 3"/>

OTHERS

Std	(1)	(2)	(3)	(4)	(5)	(6)
Calib. Code	<input type="text" value="501"/>	<input type="text" value="xxx"/>	<input type="text" value="xxx"/>	<input type="text" value="xxx"/>	<input type="text" value="xxx"/>	<input type="text" value="xxx"/>
Concentration	<input type="text" value="0,00"/>	<input type="text" value="150"/>	<input type="text" value="600"/>	<input type="text" value="1500"/>	<input type="text" value="3000"/>	<input type="text" value="5000"/>
Position						
Conversion Factor						
Sample Volume	<input type="text" value="3,0"/>	<input type="text" value="3,0"/>	<input type="text" value="3,0"/>	<input type="text" value="3,0"/>	<input type="text" value="3,0"/>	<input type="text" value="3,0"/>
Diluent S. Volume	<input type="text" value="0,0"/>	<input type="text" value="0,0"/>	<input type="text" value="0,0"/>	<input type="text" value="0,0"/>	<input type="text" value="0,0"/>	<input type="text" value="0,0"/>
Diluent Volume	<input type="text" value="0"/>	<input type="text" value="0"/>	<input type="text" value="0"/>	<input type="text" value="0"/>	<input type="text" value="0"/>	<input type="text" value="0"/>

xxx: To be defined by operator.

BioPorto Diagnostics A/S

The Application Note applies to the quantitative determination of NGAL in human urine and EDTA plasma. The performance data shown were obtained by the manufacturer. Each individual laboratory should validate its test system.

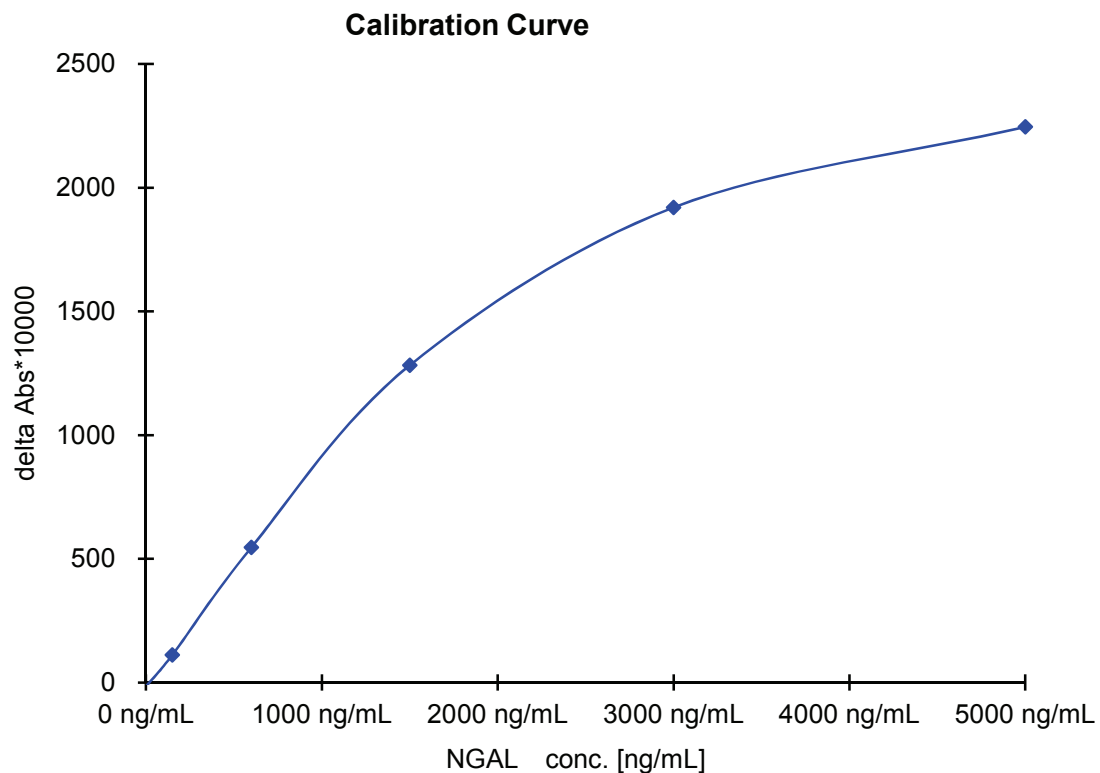
The NGAL Test™ Performance on Roche Modular-P

Sensitivity

Reagent	The NGAL Test™ Lot.RD23
calibrator	Master Calibrator
instrument	Roche Modular-P
parameter	3/150/50 P18-34 570/800nm Spline

	0 ng/mL	150 ng/mL	600 ng/mL	1500 ng/mL	3000 ng/mL	5000 ng/mL
Lot. RD23	-15,5	111,0	546,5	1282,0	1920,0	2245,5

Unit:delta Abs*10000



0 ng/mL	0,0
150 ng/mL	146,0
600 ng/mL	601,5
1500 ng/mL	1539,5
3000 ng/mL	3023,5
5000 ng/mL	4870,5

Unit:ng/mL

<i>Recovery</i>	
0 ng/mL	-
150 ng/mL	97%
600 ng/mL	100%
1500 ng/mL	103%
3000 ng/mL	101%
5000 ng/mL	97%

Re-run	
Normal	4923,0
Re-run	5031,0
<i>Recovery</i>	<i>102%</i>

Within Run Imprecision / Specificity

Sample	QC fluid -25ng/mL / Control Low,High / QC Panel Low,
Reagent	The NGAL Test™ Lot.RD23
Calibrator	Master Calibrator
Instrument	Roche Modular-P
Parameter	3/150/50 P18-34 570/800nm Spline

	QC Fluid 25ng/mL	Control Low Lot.100831	Control High Lot.100831
1	24	195	498
2	20	201	500
3	26	194	492
4	24	194	490
5	19	195	505
6	24	195	492
7	23	192	489
8	24	198	487
9	15	195	495
10	23	214	491
11	24	-	-
12	26	-	-
13	18	-	-
14	26	-	-
15	19	-	-
16	25	-	-
17	18	-	-
18	18	-	-
19	24	-	-
20	20	-	-
N	20	10	10
Mean	22,0	197,3	493,9
Max	26,0	214,0	505,0
Min	15,0	192,0	487,0
SD	3,2927	6,3605	5,5867
CV%	14,97%	3,22%	1,13%

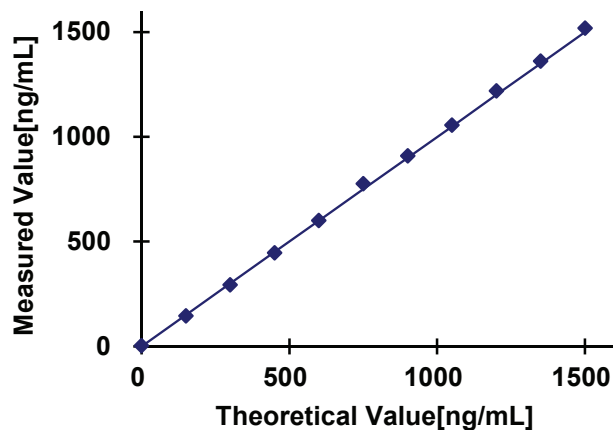
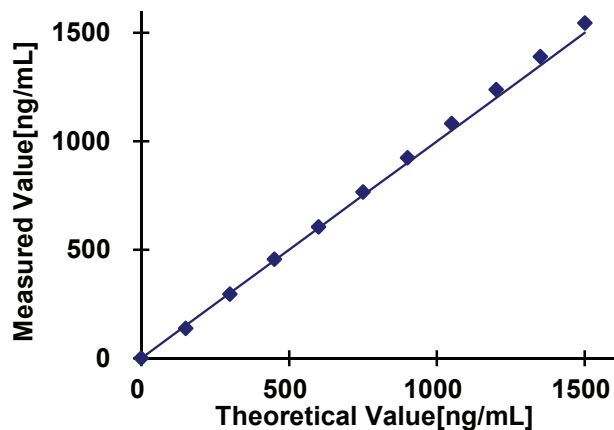
Unit:ng/mL

Specificity

	H917	Roche Modular-P	Recovery
Control Low	194,9	197,3	101%
Control High	485,2	493,9	102%
Panel Low	110,2	113,3	103%
Panel Medium	313,4	321,4	103%
Panel High	873,2	909,4	104%

Linearity ~ 1500ng/mL

Sample	Calibrator
Dilution	Saline, Calibrator diluent
Calibrator	Master Calibrator
Instrument	Roche Modular-P
Parameter	3/150/50 P18-34 570/800nm Spline



Dil.levels	Theoretical	Measured	Recovery
1500ng/mL - Saline			
10,0	0,0	0,0	
9,0	1,0	137,5	92%
8,0	2,0	295,0	98%
7,0	3,0	456,0	101%
6,0	4,0	605,0	101%
5,0	5,0	766,0	102%
4,0	6,0	924,5	103%
3,0	7,0	1081,5	103%
2,0	8,0	1238,5	103%
1,0	9,0	1390,5	103%
0,0	10,0	1545,5	103%

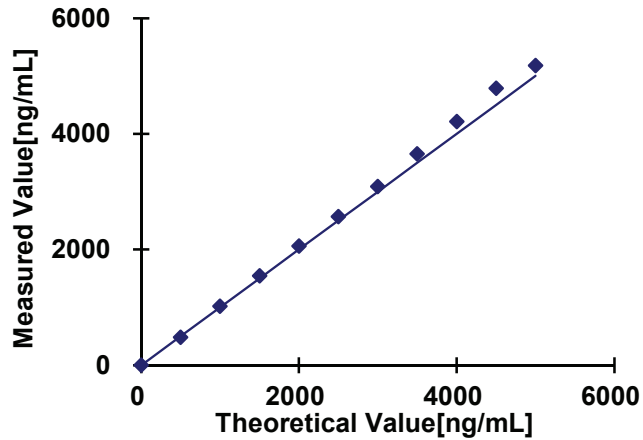
Unit:ng/mL

Dil.levels	Theoretical	Measured	Recovery
1500ng/mL - Calibrator diluent			
10,0	0,0	0,0	4,0
9,0	1,0	146,0	97%
8,0	2,0	293,0	98%
7,0	3,0	447,0	99%
6,0	4,0	601,5	100%
5,0	5,0	777,0	104%
4,0	6,0	909,0	101%
3,0	7,0	1055,5	101%
2,0	8,0	1220,0	102%
1,0	9,0	1361,5	101%
0,0	10,0	1518,5	101%

Unit:ng/mL

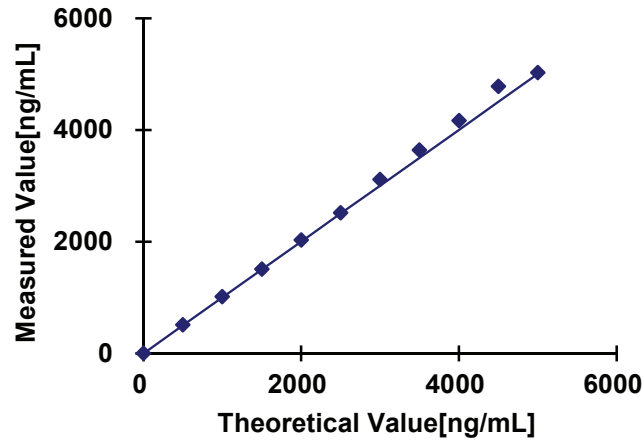
Linearity ~ 5000ng/mL & Prozone

Sample	Calibrator, QC fluid
Dilution	Saline, Calibrator diluent
Calibrator	Master Calibrator
Instrument	Roche Modular-P
Parameter	3/150/50 P18-34 570/800nm Spline



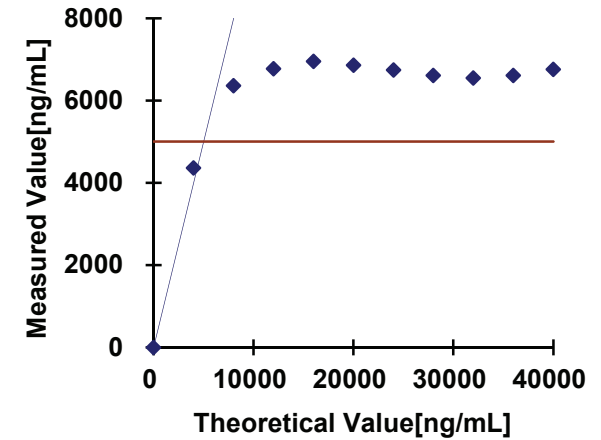
Dil.levels	Theoretical	Measured	Recovery
5000ng/mL - Saline			
10,0	0,0	0,0	
9,0	1,0	484,5	97%
8,0	2,0	1018,0	102%
7,0	3,0	1544,5	103%
6,0	4,0	2061,5	103%
5,0	5,0	2569,0	103%
4,0	6,0	3090,5	103%
3,0	7,0	3655,5	104%
2,0	8,0	4210,5	105%
1,0	9,0	4790,5	106%
0,0	10,0	5182,5	104%

Unit:ng/mL



Dil.levels	Theoretical	Measured	Recovery
5000ng/mL - Calibrator diluent			
10,0	0,0	0,0	
9,0	1,0	510,0	102%
8,0	2,0	1018,0	102%
7,0	3,0	1510,5	101%
6,0	4,0	2028,0	101%
5,0	5,0	2518,5	101%
4,0	6,0	3114,0	104%
3,0	7,0	3641,5	104%
2,0	8,0	4164,0	104%
1,0	9,0	4777,0	106%
0,0	10,0	5028,5	101%

Unit:ng/mL



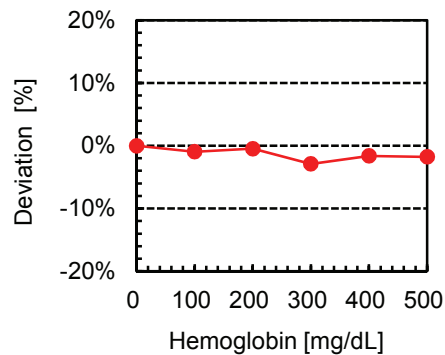
Dil.levels	Theoretical	Measured	Mark
Prozone - Calibrator diluent			
10,0	0,0	0,0	0,5
9,0	1,0	4361,5	
8,0	2,0	6360,0	>5000
7,0	3,0	6768,5	>5000
6,0	4,0	6953,0	>5000
5,0	5,0	6855,5	Error
4,0	6,0	6742,0	Error
3,0	7,0	6612,0	Error
2,0	8,0	6544,5	Error
1,0	9,0	6605,5	Error
0,0	10,0	6755,0	Error

Unit:ng/mL

Interferences

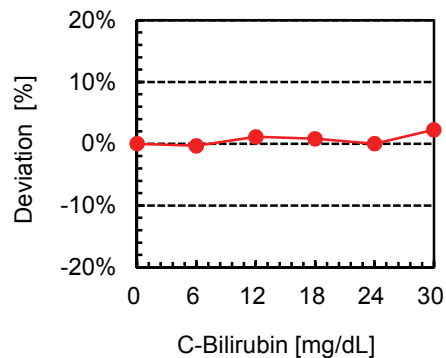
Sample	Hemoglobin, Bilirubin, Intrafat
Dilution	Calibrator diluent
Calibrator	Master Calibrator
Instrument	Roche Modular-P
Parameter	3/150/50 P18-34 570/800nm Spline

Hemoglobin :



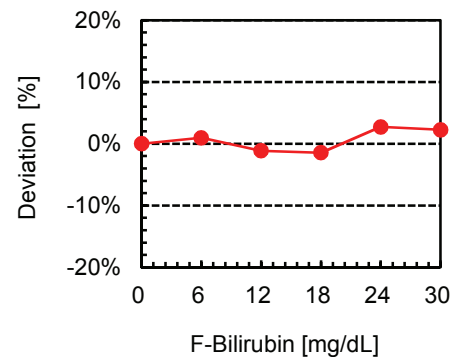
mg/dL	<i>Deviation</i>
0	0,0%
100	-1,0%
200	-0,5%
300	-2,9%
400	-1,6%
500	-1,8%

C-Bilirubin :



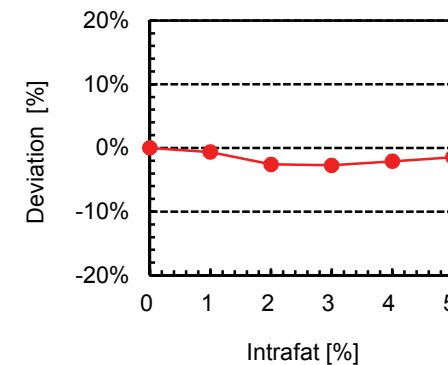
mg/dL	<i>Deviation</i>
0	0,0%
6	-0,3%
12	1,1%
18	0,8%
24	0,0%
30	2,3%

F-Bilirubin :



mg/dL	<i>Deviation</i>
0	0,0%
6	1,0%
12	-1,1%
18	-1,4%
24	2,7%
30	2,3%

Intrafat :



%	<i>Deviation</i>
0	0,0%
1	-0,6%
2	-2,6%
3	-2,7%
4	-2,1%
5	-1,4%