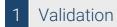


Datasheet for ABIN1690717

Ngal/Mmp-9 Complex ELISA Kit







Overview	
Quantity:	96 tests
Target:	Ngal/Mmp-9 Complex
Reactivity:	Human
Method Type:	Competition ELISA
Detection Range:	0.312-20 ng/mL
Minimum Detection Limit:	0.312 ng/mL
Application:	ELISA
Product Details	
Purpose:	For the quantitative determination of human NGAL/MMP-9 complex concentrations in serum, plasma, cell culture supernates, tissue homogenates and saliva.

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Sample Type:	Serum, Plasma, Saliva, Cell Culture Supernatant, Tissue Homogenate					
Analytical Method:	Quantitative					
Detection Method:	Colorimetric					
Cross-Reactivity (Details):	Limited by current skills and knowledge, it is impossible for us to complete the cross-reactivity detection between the target antigen and all analogues for other species. Therefore, cross reaction may still exist.					
Sensitivity:	0.029 ng/mL					
Components:	 Assay plate (12 × 8 coated Microwells) Standard (freeze dried) Biotin-antibody (100 × concentrate) 					

- HRP-avidin (100 × concentrate)
- · Biotin-antibody Diluent
- · HRP-avidin Diluent
- · Sample Diluent
- Wash Buffer (25 × concentrate)
- · TMB Substrate
- Stop Solution
- · Adhesive Strip (for 96 wells)
- · Instruction manual

Material not included:

- Microplate reader capable of measuring absorbance at 450nm, with the correction wavelength set at 540nm or 570nm.
- An incubator which can provide stable incubation conditions up to 37°C ± 0.5°C.
- · Squirt bottle, manifold dispenser or automated microplate washer.
- · Absorbent paper for blotting the microtiter plate.
- · 100mL and 500mL graduated cylinders.
- Deionized or distilled water.
- · Pipettes and pipette tips.
- · Test tubes for dilution.

Target Details

Target:

Ngal/Mmp-9 Complex

Application Details

Application Notes:

- The supplier is only responsible for the kit itself, but not for the samples consumed during the assay. The user should calculate the possible amount of the samples used in the whole test. Please reserve sufficient samples in advance.
- Samples to be used within 5 days may be stored at 2-8°C, otherwise samples must be stored at -20°C (≤ 1 month) or -80°C (≤ 2 months) to avoid loss of bioactivity and contamination.
- Grossly hemolyzed samples are not suitable for use in this assay.
- If the samples are not indicated in the manual, a preliminary experiment to determine the validity of the kit is necessary.
- Please predict the concentration before assaying. If values for these are not within the range
 of the standard curve, users must determine the optimal sample dilutions for their particular
 experiments.
- Tissue or cell extraction samples prepared by chemical lysis buffer may cause unexpected ELISA results due to the impacts of certain chemicals.
- Owing to the possibility of mismatching between antigens from another resource and antibodies used in this supplier's kits (e.g., antibody targets conformational epitope rather than linear epitope), some native or recombinant proteins from other manufacturers may not be recognized by this supplier's products.

- Influenced by factors including cell viability, cell number and cell sampling time, samples
 from cell culture supernatant may not be recognized by the kit.
- Fresh samples without long time storage are recommended for the test. Otherwise, protein degradation and denaturalization may occur in those samples and finally lead to wrong results.

Comment:

Detection wavelength: 450 nm

Information on standard material:

Depending on the antigen to be detected, standards can be either native or recombinant protein. The recombinant proteins are being expressed in CHO cells in most cases. Please inquire for more information. The formulation of auxiliary material in the standard is considered proprietary information, however it does not contain any poisonous substance. Proclin 300 (1:3000) is used as preservative.

Information on reagents:

In most cases the stop solution provided is 1 N H2SO4. The formulation of wash solution is proprietary information. None of the components contain (sodium) azide, thimerosal, 2-mercaptoethanol (2-ME) or any other poisonous materials. For the sandwich method kits, the sample diluent, antibody diluent, enzyme diluent and standard all contain BSA.

Information on antibodies:

The antibodies provided in different kits vary in regards to clonality and host. Some antibodies are affinity purified, some are Protein A

Sample Volume:

50 μL

Assay Time:

1 - 4.5 h

Plate:

Pre-coated

Protocol:

This assay employs the competitive inhibition enzyme immunoassay technique. The microtiter plate provided in this kit has been pre-coated with the target antigen. Standards or samples are added to the appropriate microtiter plate wells with Horseradish Peroxidase (HRP) conjugated antibody preparation specific for the target antigen. The competitive inhibition reaction is launched between the pre-coated target antigen and the target antigen in samples. A substrate solution is added to the wells and the color develops opposite to the amount of the target antigen in the samples. The color development is stopped and the intensity of the color is measured.

Reagent Preparation:

- HRP-conjugate (1x) Centrifuge the vial before opening.
 HRP-conjugate requires a 100-fold dilution. The suggested dilution is 10μL of HRP-conjugate + 990μL of HRP-conjugate Diluent.
- Wash Buffer (1x) If crystals have formed in the concentrate, warm up to room temperature and mix gently until the crystals have completely dissolved. Dilute 20mL of Wash Buffer Concentrate (25x) into deionized or distilled water to prepare 500mL of Wash Buffer (1x).
- Standard Centrifuge the standard vial at 6000-10000rpm for 30s.

 Reconstitute the Standard with 1ml of Sample Diluent. Do not substitute other diluents. This reconstitution produces a stock solution of 400ng/mL. Mix the standard to ensure complete reconstitution and allow the standard to sit for a minimum of 15 minutes with gentle agitation prior to making dilutions.

Pipette $150\mu L$ of Sample Diluent into each tube. Use the stock solution to produce a 2-fold dilution series. Mix each tube thoroughly before the next transfer. The undiluted Standard serves as the high standard (400ng/mL). Sample Diluent serves as the zero standard (0ng/mL).)

Note:

- Kindly use graduated containers to prepare the reagent. Please don't prepare the reagent directly in the Diluent vials provided in the kit.
- Bring all reagents to room temperature (18-25°C) before use for 30 min.
- Prepare fresh standard for each assay. Use within 4 hours and discard after use.
- · Making serial dilution in the wells directly is not permitted.
- Please carefully reconstitute Standards according to the instruction. Avoid foaming and mix gently until the crystals have completely dissolved. To minimize imprecision caused by pipetting, use small volumes and ensure that pipettors are calibrated. It is recommended to suck more than 10µL when pipetting.
- It is recommended to use distilled water to prepare reagents and samples. Using contaminated water or container for reagent preparation will influence detection result.

Sample Collection:

- **Serum**: Use a serum separator tube (SST) and allow samples to clot for two hours at room temperature or overnight at 4 °C before centrifugation for 15 minutes at 1000 × g. Remove serum and assay immediately or aliquot and store samples at -20 °C or -80 °C. Avoid repeated freeze-thaw cycles.
- Plasma: Collect plasma using EDTA or heparin as an anticoagulant. Centrifuge for 15 minutes at 1000 x g at 2-8 °C within 30 minutes of collection. Assay immediately or aliquot and store samples at -20 °C or -80 °C. Avoid repeated freeze-thaw cycles.
- Tissue Homogenates: Rinse 100 mg tissue with 1× PBS, homogenize in 1mL of 1× PBS and store overnight at -20 °C. After two freeze-thaw cycles to break the cell membranes, centrifuge the homogenates for 5 minutes at 5000 × g, 2-8 °C. Remove and assay the supernate immediately. Alternatively, aliquot and store samples at -20 °C or -80 °C. Centrifuge the sample again after thawing before the assay. Avoid repeated freeze-thaw cycles.

Assay Procedure:

• 1. Prepare all reagents, working standards and samples as directed in the respective

sections.

- 2. Refer to the Assay Layout Sheet to determine the number of wells to be used and put any remaining wells and the desiccant back into the pouch and seal the ziploc, store unused wells at 4°C.
- 3. Set a blank well without any solution.
- 4. Add 50µL of standard or sample per well (except blank well). Then add 50µL of HRP-conjugate (1x) to each well (except blank well). Cover the microtiter plate with adhesive strip. Incubate for 30 minutes at 37°C. A plate layout is provided to record standards and samples assayed.
- 5. Aspirate each well and wash, repeating the process four times for a total of five washes. Wash by filling each well with Wash Buffer (200µL) using a squirt bottle, multi-channel pipette, manifold dispenser or autowasher and let it stand for 2 minutes, complete removal of liquid at each stage is essential for good performance. After the last wash, remove any remaining Wash Buffer by aspirating or decanting. Invert the plate and blot it against clean paper towels.
- 6. Add 90µL of TMB Substrate to each well. Incubate for 20 minutes at 37°C. Protect from light.
- 7. Add 50µL of Stop Solution to each well, gently tap the plate to ensure thorough mixing.
- 8. Determine the optical density of each well within 5 minutes using a microplate reader set to 450nm. If wavelength correction is available, set to 540nm or 570nm. Subtract readings at 540nm or 570nm from the readings at 450nm. This subtraction will correct for optical imperfections in the plate. Readings made directly at 450nm without correction may be higher and less accurate.

Note:

- The experiment's final results will be closely related to validity of the products, operation skills of the end users and the environmental conditions.
- Samples or reagents addition: Please use the freshly prepared Standard. Please carefully add samples to wells and mix gently to avoid foaming. Do not touch the well wall as possible. For each step in the procedure, total dispensing time for addition of reagents or samples to the assay plate should not exced 10 minutes. This will ensure equal elapsed time for each pipetting step, without interruption. Duplication of all standards and specimens, although not required, is recommended. To avoid cross-contamination, change pipette tips between reagent additions. Also, use separate reservoirs for each reagent.
- Incubation: To ensure accurate results, proper adhesion of plate sealers during incubation steps is necessary. Do not allow wells to sit uncovered for extended periods between incubation steps. Once reagents have been added to the well strips, DO NOT let the strips DRY at any time during the assay. Incubation time and temperature must be observed.
- Washing: The wash procedure is critical. Complete removal of liquid at each step is essential
 for good performance. After the last wash, remove any remaining Wash Solution by
 aspirating or decanting and remove any drop of water and fingerprint on the bottom of the
 plate. Insufficient washing will result in poor precision and falsely elevated absorbance
 reading. When using an automated plate washer, adding a 30 second soak period following
 the addition of wash buffer and/or rotating the plate 180 degrees between wash steps may

improve assay precision.

- Controlling of reaction time: Observe the change of color after adding TMB Substrate (e.g. observation once every 10 minutes), TMB Substrate should change from colorless or light blue to gradations of blue. If the color is too deep, add Stop Solution in advance to avoid excessively strong reaction which will result in inaccurate absorbance reading.
- TMB Substrate is easily contaminated. TMB Substrate should remain colorless or light blue until added to the plate. Please protect it from light.
- Stop Solution should be added to the plate in the same order as the TMB Substrate. The
 color developed in the wells will turn from blue to yellow upon addition of the Stop Solution.
 Wells that are green in color indicate that the Stop Solution has not mixed thoroughly with the
 TMB Substrate.

Calculation of Results:

Average the duplicate readings for each standard and sample and subtract the average zero standard optical density.

Create a standard curve by reducing the data using computer software capable of generating a four parameter logistic (4-PL) curve fit. As an alternative, construct a standard curve by plotting the mean absorbance for each standard on the x-axis against the concentration on the y-axis and draw a best fit curve through the points on the graph. The data may be linearized by plotting the log of the target antigen concentration versus the log of the O.D. and the best fit line can be determined by regression analysis. This procedure will produce an adequate but less precise fit of the data.

If samples have been diluted, the concentration read from the standard curve must be multiplied by the dilution factor.

Assay Precision:

Intra-assay precision (precision within an assay): Three samples of known concentration were tested twenty times on one plate to assess precision.

Inter-assay precision (precision between assays): Three samples of known concentration were tested in twenty assays to assess precision.

- Intra-assay: CV% less than 8%
- · Inter-assay: CV% less than 10%

Restrictions:

For Research Use only

Handling

Precaution of Use:

The Stop Solution provided with this kit is an acid solution. Wear eye, hand, face and clothing protection when using this material.

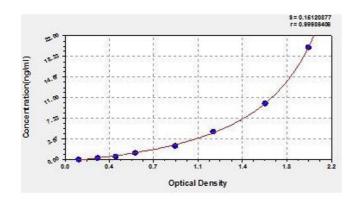
Handling Advice:

- The kit should not be used beyond the expiration date on the kit label.
- Do not mix or substitute reagents with those from other lots or sources.

- If samples generate values higher than the highest standard, dilute the samples with Sample Diluent and repeat the assay.
- Any variation in Sample Diluent, operator, pipetting technique, washing technique, incubation time/temperature and kit age can cause variation in binding.
- This assay is designed to eliminate interference by soluble receptors, binding proteins and other factors present in biological samples. Until all factors have been tested in the Immunoassay, the possibility of interference cannot be excluded.

Storage:	4 °C/-20 °C
Storage Comment:	For unopened kit: All the reagents should be kept according to the labels on vials.
Expiry Date:	6 months

Images



ELISA

Image 1. Typical standard curve





Successfully validated (ELISA (ELISA))

by Affina Biotechnologies

Report Number: 029583

Date: Jan 30 2014

Lot Number:	T30095914
Method validated:	ELISA (ELISA)
Positive Control:	Human individual postmenopausal female serum
Negative Control:	Chicken plasma
Notes:	Standard curve was fit to a logistic equation, which returned a small positive concentration value for the negative control.
Primary Antibody:	- Antigen: NGAL/MMP-9 complex - Catalog number: ABIN1690717 - Supplier: Cusabio - Supplier catalog number: CSB-E08006h - Lot number: T23095910
Controls:	 Positive control: Human individual post-menopausal female serum (Biochemed, 750-NS-FI-POM) Negative control: Chicken plasma (Sigma-Aldrich, p3266) – reconstituted at 1 mg/mL Standard curve: 0, 312, 0.625, 1.25, 2.5, 5, 10, 20 ng/mL NGAL/MP9 complex provided in the ELISA kit Spike control: 10 ng/mL standard premixed with chicken plasma in a 1:1 ratio
Protocol:	 100 µL of standard and samples were added 96-well strip plates provided in the kit. All samples and standards were assayed in duplicate. The microplate was covered and incubated at 37°C for 1 hr. Plate contents were discarded. 100 µL of biotin antibody conjugate was added and incubated at 37°C for 1 hr Content of the wells was discarded and wells were washed 3 times with 200 µl of 1x wash solution with a 2 min soak for each wash. 100 µL of HRP-avidin conjugate was added and incubated at 37°C for 1 hr. Contents of the wells were discarded and wells were washed 5 times with 200 µl of 1x wash solution with a 2 min soak for each wash. 90 µl of TMB substrate was added to each well. The plate was covered and incubated at 37°C for 15 min. 50 µl of the Stop Solution was added per well. The optical density (OD value) of each well was read immediately using a microplate reader set to 450 nm. The duplicate readings for each sample were averaged and the average zero standard optical



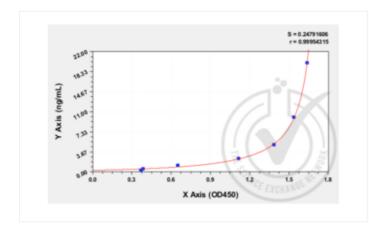
density subtracted. The corrected average-value was tabulated as Average Absorbance. A standard curve was generated by plotting the mean OD value for each standard on the x-axis against the concentration on the Y-axis using CurveExpert 1.4. A logistic equation was used for the best fit through the points on the graph.

· The CurveExpert Analyze feature was used to calculate NGAL/MMP9 concentrations of the samples based on their Average Absorbance values.

Experimental Notes:

none

Images for Validation report #029583



Validation image no. 1 for Ngal/Mmp-9 Complex ELISA Kit (ABIN1690717)

Figure 1: Graph of corrected-average absorbance (OD 450 nm) readings plotted for standard curve samples.

Type	Sample ng/ml	Reading-	Reading- 2	Avg Reading	Avg Absorbance	SD	Calculated
Standard Curve	20	1.873	1.862	1.867	1.636	0.007311	20.006
	10	1.773	1.766	1.769	1.538	0.004773	9.9746
	5	1.619	1.610	1.615	1.384	0.006717	5.12739
	2.5	1.364	1.324	1.344	1.113	0.028314	2.46678
	1.25	0.878	0.883	0.881	0.650	0.003488	0.918345
	0.625	0.594	0.638	0.616	0.385	0.030798	0.57974
	0.312	0.570	0.629	0.600	0.369	0.041974	0.564453
	blank	0.221	0.241	0.231			1)
Spike Control	10	1.737	1.716	1.726	1.495	0.015	8.02297
Positive Control	Human plasma	1.082	1.119	1.101	0.870	0.0266	2.78
Negative control	Chicken Plasma	0.241	0.221	0.231	0.000	0.014	0.31158

Validation image no. 2 for Ngal/Mmp-9 Complex ELISA Kit (ABIN1690717)

Table 1: Table of absorbance readings (OD 450 nm) for standard curve, spike controls and unknown control samples. NGAL/MMP9 is clearly detected in the positive sample. Spike controls indicate little interference in absorbance readings from the two-fold diluted plasma sample. Value for Average Reading is derived from the average of two readings (OD 450nm). The Average Reading for blank sample (no conjugate added) was subtracted from all Average Readings to yield Average Absorbance values for Standards, spike controls and control samples. Standard deviation is included for all samples. The concentration of samples was calculated using the Analyze feature of the CurveExpert 1.4 software for a logistic equation fit.