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# Human Mucosal addressin cell adhesion molecule 1 (MAdCAM1) ELISA Kit

Catalog No.: abx251091

Size: 96T

Range: 0.312 ng/ml - 20 ng/ml

Sensitivity: < 0.188 ng/ml

Storage: Store at 4°C for up to 6 months.

Application: For quantitative detection of MAdCAM1 in Human Serum, Plasma, Tissue Homogenates and other biological fluids.

Introduction: Mucosal vascular addressin cell adhesion molecule 1 (MAdCAM1) is a protein that in humans is encoded by the MADCAM1 gene. MAdCAM1 is an extracellular protein of the endothelium of venules and ligand to the homing receptors of lymphocytes. The task of these ligands and their receptors is to determine which tissue the lymphocyte will enter next. They carry carbohydrates in order to be recognized by L-selectin. MADCAM1 interacts preferentially with the leukocyte beta7 integrin LPAM-1 (alpha4 / beta7), L-selectin, and VLA-4 (alpha4 / beta1) on myeloid cells to direct leukocytes into mucosal and inflamed tissues. It is a member of the immunoglobulin superfamily and is similar to ICAM-1 and VCAM-1.

## Principle of the Assay

This kit is based on sandwich enzyme-linked immuno-sorbent assay technology. MAdCAM1 Antibody is pre-coated onto 96-well plates. The standards, test samples and biotin detection antibody are added to the wells and washed with wash buffer. Biotin conjugated MAdCAM1 antibody is used as a detection antibody. TMB substrate is used to visualize HRP enzymatic reaction. TMB is catalyzed by HRP to produce a blue colour product that changes into yellow after adding stop solution. The intensity of the color yellow is proportional to the MAdCAM1 amount bound on the plate. The O.D. absorbance is measured spectrophotometrically at 450nm in a microplate reader, and then the concentration of MAdCAM1 can be calculated.

## Kit components

- 1. One pre-coated 96 well plate
- 2. Standard: 2 tubes
- 3. Sample/Standard Diluent Buffer: 20 ml
- 4. Biotin conjugated antibody(Concentrated): 120 μl, Dilution 1:100
- 5. Antibody diluent buffer: 10 ml
- 6. HRP streptavidin conjugate (SABC) (Concentrated): 120 µl, Dilution 1:100
- 7. SABC diluent buffer: 10 ml
- 8. TMB substrate: 10 ml
- 9. Stop solution: 10 ml
- 10. Wash buffer (25X): 30 ml

## **Material Required But Not Provided**

- 1. 37°C incubator
- 2. Microplate reader (wavelength: 450nm)
- 3. Precision pipette and disposable pipette tips
- 4. Automated plate washer
- 5. ELISA shaker
- 6. 1.5ml tubes to prepare standard/sample dilutions
- 7. Absorbent filter papers
- 8. 100 ml and 1 L volume graduated cylinders

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### **Protocol**

### A. Preparation of sample and reagents

### 1. Sample

Store samples to be assayed within 24 hours at 2-8°C. Alternatively, aliquot and store at -20°C or -80°C for long term. Avoid repeated freeze-thaw cycles.

- **Serum:** Samples should be collected into a serum separator tube. Coagulate the blood at room temperature (~2 hr) or overnight at 4°C. Centrifuge at approximately 1000 × g for 20 min. Analyze the serum immediately or aliquot and store at -20°C or -80°C.
- Plasma: Collect plasma using EDTA-Na<sub>2</sub> as an anticoagulant. Centrifuge for 15 minutes at 1000 x g within 30 minutes of collection. Assay immediately or aliquot and store at -20°C. Avoid hemolysis and high cholesterol samples.
- Tissue homogenates: The preparation of tissue homogenates will vary depending upon tissue type this is just an example. Rinse tissues with ice-cold PBS to remove the excess of blood. Weigh before homogenization. Finely mince tissues and homogenize with a tissue homogenizer on ice in PBS and sonicate the cell suspension. Centrifuge the homogenates at 5000 × g for 5 min and collect the supernatant. Assay immediately or aliquot and store at -20°C.
- Other biological fluids: Centrifuge at approximately 1000 × g for 20 min to remove precipitant. Analyze immediately or aliquot and store at -20°C or -80°C.

#### Note:

- » Fresh samples or recently obtained samples are recommended to prevent degradation and denaturalization that may lead to erroneous results. It is recommended to store samples to be used within 5 days at 4°C, within 1 month at -20°C and within 2 months at -80°C.
- » Samples should be clear and transparent. Samples must be diluted so that the expected concentration falls within the kit's range.
- » Please bring sample slowly to room temperature. Sample hemolysis will influence the result. Hemolyzed specimen should not be used. Samples that contain NaN<sub>3</sub> cannot be detected as it interferes with HRP.

## General Sample guideline:

Estimate the concentration of the target in the sample and select the correct dilution factor to make the diluted target concentration fall near the middle of the kit's range. Generally, for high concentration, dilute 1:100, for medium concentration, dilute 1:10 and for low concentration, dilute 1:2. Very low concentrations do not need dilution. Dilute the sample with the provided Sample Diluent Buffer and mix thoroughly. Several trials may be necessary to determine the optimal dilution factor.

### 2. Wash buffer

Dilute the concentrated Wash buffer 25-fold (1/25) with distilled water (i.e. add 30 ml of concentrated wash buffer into 720 ml of distilled water).

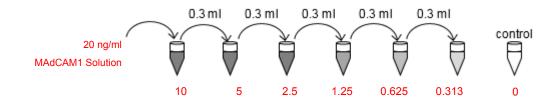
### 3. Standard

Preparation of the MAdCAM1 standard: standard solution should be prepared no more than 2 hours prior to the experiment. Two tubes of standard are included in each kit. Use one tube for each experiment. (Note: Do not dilute the standard directly in the plate).

- a.) 20 ng/ml standard solution. Add 1 ml of Sample/Standard diluent buffer into one Standard tube, keep the tube at room temperature for 10 min, mix thoroughly and avoid foaming or bubbles.
- b.) 10 ng/ml  $\rightarrow$  0.3125 ng/ml standard solutions: Label 6 tubes with 10 ng/ml, 5 ng/ml, 2.5 ng/ml, 1.25 ng/ml, 0.625 ng/ml and 0.3125 ng/ml. Aliquot 0.3 ml of the Sample / Standard diluent buffer into each tube. Add 0.3 ml of the above 20 ng/ml standard solution into 1st tube and mix thoroughly. Transfer 0.3 ml from 1st tube to 2nd tube and mix thoroughly, and so on.

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**Note:** The standard solutions are best used within 2 hours. The standard solution can be stored at 4°C for up to 12 hours, or at -20 °C for up to 48 hours. Avoid repeated freeze-thaw cycles.

- 4. Preparation of Biotin conjugated antibody working solution: prepare no more than 1 hour before the experiment.
- a.) Calculate the total volume of the working solution: 0.1 ml / well × quantity of wells. (Allow 0.1-0.2 ml more than the total volume).
- b.) Dilute the Biotin conjugated antibody with antibody diluent buffer at 1/100 and mix thoroughly. i.e. Add 1 µl of Biotin conjugated antibody into 99 µl of antibody diluent buffer.
- 5. Preparation of HRP Streptavidin Conjugate (SABC) working solution: prepare no more than 30 min. before the experiment
- a.) Calculate the total volume of the working solution: 0.1 ml / well × quantity of wells. (Allow 0.1-0.2 ml more than the total volume).
- b.) Dilute the SABC with SABC diluent buffer at 1/100 and mix thoroughly. i.e. Add 1 µl of SABC into 99 µl of SABC diluent buffer.

## **B.** Assay Procedure

Equilibrate the SABC working solution to room temperature and TMB substrate at 37°C, for at least 30 minutes prior to use. It is recommended to plot a standard curve for each test.

- 1. Wash the plate two times before adding standard, samples and buffers. Any strips that are not being used should be kept dry and stored at 4°C. Set standard, test sample and control (zero) wells on the pre-coated plate and record their positions. It is recommended to measure each standard and sample in duplicate.
- 2. Add 100 µl of the prepared standards solutions into the standard wells.
- 3. Add 100 µl of Sample / Standard diluent buffer into the control (zero) well.
- 4. Add 100 µl of appropriately diluted sample into test sample wells.
- Cover the plate and incubate at 37°C for 90 minutes.
- 6. Remove the cover and discard the contents by clapping the plate on absorbent filter papers or any other absorbent material. **Do** not wash the plate and do NOT let the wells dry out completely at any time.
- 7. Add 100 µl of prepared Biotin conjugated antibody working solution into each well (standard, test sample and zero well). Add the solution at the bottom of each well without touching the side walls. Seal the plate with a cover and incubate at 37°C for 60 minutes.
- 8. Remove the cover, and wash the plate 3 times. Discard the solution without touching the side walls. Blot the plate on an absorbent material. Fill each well completely with wash buffer and soak for at least 1-2 min. Discard the contents and clap the plate on absorbent filter papers or other absorbent material. Repeat this procedure for a total of three times.

**Please note:** For automated washing, discard the solution in all wells and wash three times overfilling the wells with Wash buffer. After the final wash invert the plate and tap on absorbent filter papers or other absorbent material. It is recommended that the washer be set for a soaking time of 1 min.

- 9. Add 100 µl of SABC working solution into each well, cover the plate and incubate at 37°C for 30 minutes.
- 10. Remove the cover and wash the plate 5 times with Wash buffer. Allow the wash buffer to remain in the wells 1-2 min for each wash. Discard the washing buffer and blot the plate onto absorbent filter papers or other absorbent material.

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11. Add 90 µl of TMB substrate into each well. Cover the plate and incubate at 37°C in dark conditions for 15-20 minutes (incubation time is for reference only, do not exceed 30 minutes). When an apparent gradient appears in the standard wells the reaction can

be terminated.

12. Add 50 µl of Stop solution into each well. There should be a color change to yellow. Gently tap the plate to ensure thorough

mixing.

13. Ensure that there are no fingerprints or water on the bottom of the plate, and that the fluid in the wells is free of bubbles. Measure

the absorbance at 450nm immediately.

For calculation, (the relative O.D.450) = (the O.D.450 of each well) - (the O.D.450 of Zero well). The standard curve can be plotted as the relative O.D.450 of each standard solution (Y) vs. the respective concentration of the standard solution (X). Log-log curve fitting is

recommended for data analysis. The Human MAdCAM1 concentration of the samples can be interpolated from the standard curve.

Note: If the samples measured were diluted, multiply the dilution factor by the concentrations from interpolation to obtain the

concentration before dilution.

C. Precautions

1. Ensure that the plate remains dry until starting the assay.

Before using the kit, centrifuge the tubes briefly to bring down the contents trapped in the lid.

3. Wash buffer may crystallize and separate. If this happens, please warm the tube and mix gently to dissolve.

Avoid foaming or bubbles when mixing or reconstituting components.

It is recommended to assay all standards, controls and samples in duplicate or triplicate.

Do NOT let the plate dry out completely during the assay as this will inactivate the biological material on the plate.

Ensure plates are properly sealed or covered during incubation steps.

Complete removal of all solutions and buffers during wash steps is necessary for accurate measurement readings.

To avoid cross contamination do not reuse pipette tips and tubes.

10. Do not use components from a different kit or expired ones.

11. The TMB substrate is light sensitive and should be protected from direct sunlight and UV sources. Unreacted substrate should be colorless or very light yellow in appearance. The product should be allowed to equilibrate to room temperature (25°C) prior to use.

Aspirate the dosage needed with sterilized tips and do not dump the residual solution back into the vial.

D. Precision

Intra-assay Precision (Precision within an assay): 3 samples with low, medium and high levels of MAdCAM1 were tested 20 times on

one plate, respectively.

Inter-assay Precision (Precision between assays): 3 samples with low, medium and high levels of MAdCAM1 were tested on 3

different plates, 8 replicates in each plate.

CV (%) = SD/meanX100

Intra-Assay: CV<8%

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Inter-Assay: CV<10%

## E. Typical Data & Standard Curve

Typical Standard Curve Data provided for demonstration purposes only. A new standard curve must be generated for each assay performed.

Concentration ng/ml	0	0.3125	0.625	1.25	2.5	5	10	20
OD450	0.036	0.124	0.218	0.345	0.595	1.098	1.409	2.038

