## PromedeusLab

# Human Kidney Injury Molecule 1 ELISA 

Cat. No.: PL1001
Enzyme Immunoassay for the quantitative determination Kidney Injury Molecule 1 (KIM-1) in human urine and serum.

The urinary kidney injury molecule 1 (KIM-1) participates in renal tissue damage and repair and is proposed as a biomarker of early and subclinical AKI. KIM-1 is also elevated in the urine of a significant fraction of patients apparently recovered from an AKI. Besides its potential utility in the early and subclinical diagnosis of renal damage, this study suggests a new application of urinary KIM-1 in the non-invasive follow-up of renal repair after AKI. ${ }^{1}$

## PRINCIPLE OF KIDNEY INJURY MOLECULE 1 ELISA

The microtiter plate is coated with the antibody specifically binding the Kidney Injury Molecule 1. The human serum or plasma is incubated in the plate with the capture antibody.

The specimen is washed out and the specifically bound protein is incubated with HRP-labelled detection antibody. Unbound reagent is then washed out. Horseradish peroxidase (HRP) bound in the complex reacts with the chromogenic substrate (TMB) creating the blue colour. The reaction is stopped by addition of STOP solution $\left(\mathrm{H}_{2} \mathrm{SO}_{4}\right)$.

The absorbance values are measured at 450 nm (optionally $450 / 630 \mathrm{~nm}$ ) and are proportional to the concentration of KIM-1 in the specimen. The concentration of KIM-1 in unknown samples is determined from the calibration curve which is created by plotting the absorbance values against the standard concentration values.

Add $100 \mu \mathrm{~L}$ of Standards, diluted QCs and Samples to the wells


Add $100 \mu \mathrm{~L}$ of HRP-conjugated Antibody to the wells


3-times wash the wells ( $350 \mu \mathrm{~L} /$ well)



Add $100 \mu \mathrm{~L}$ of Stop Solution to the wells

Read the signal at $450 \mathrm{~nm}(450 / 630 \mathrm{~nm})$ within 15 min

Kit Contents

| Item | Qty. |
| :--- | :--- |
| Antibody Coated Microtiter Plate | 96 wells |
| Antibody-HRP Conjugate | 13 mL |
| Master Standard (lyophilized) | 1 vial |
| Quality Control A (human serum, lyophilized) | 1 vial |
| Quality Control B (human serum, lyophilized) | 1 vial |
| Dilution Buffer | 13 mL |
| Wash Buffer 15x conc. | 50 mL |
| Substrate Solution | 13 mL |
| STOP Solution | 13 mL |

MATERIAL REQUIRED BUT NOT SUPPLIED

1. Glassware and test tubes
2. Microtiter plate washer
3. Precision pipettes (various volumes) with tips
4. Orbital shaker
5. Microtiter plate reader capable of measuring absorbance at 450 nm or $450 / 630 \mathrm{~nm}$ with software for data generation

## WARNINGS AND PRECAUTIONS

1. For research use only
2. For professional laboratory use
3. The reagents with different lot numbers should not be mixed
4. To prevent cross sample contamination, use disposable labware and pipette tips
5. To protect laboratory stuff, wear protective gloves and protective clothing
6. The substrate solution should remain colourless, keep it protected from light
7. The test should be performed at standard laboratory conditions (temperature $25^{\circ} \mathrm{C} \pm 2^{\circ} \mathrm{C}$ ).

## STORAGE CONDITIONS

1. The kit must be stored at $2-8^{\circ} \mathrm{C}$.
2. The opened components can be stored for one week at $2-8^{\circ} \mathrm{C}$.

## PREPARATION OF REAGENTS

- Use new pipette tip for pipetting different reagents and samples to prevent cross-contamination.
- All reagents and samples should be allowed to reach the temperature $25^{\circ} \mathrm{C} \pm 2^{\circ} \mathrm{C}$


## Preparation of Standards

Reconstitute lyophilized Human KIM-1 Standard in Dilution Buffer, for the volume information see the Certificate of Analysis. Let it rehydrate for 15 min . The concentration of human KIM-1 in Master Standard is $1000 \mathrm{pg} / \mathrm{mL}$,

Prepare set of Standard solution as follows:
Use the Master Standard to produce a dilution series (as below). Mix each tube thoroughly before the next transfer. The Dilution Buffer serves as Blank.

|  | Volume of Standard | Dilution <br> Buffer | Concentration <br> Std1 <br> Std2Standard $1000 \mathrm{pg} / \mathrm{mL}$ <br> (lyophilised) |
| :--- | :---: | :---: | ---: |
| Std3 | $300 \mu \mathrm{~L}$ of Std1 | $300 \mu \mathrm{~L}$ | $500 \mathrm{pg} / \mathrm{mL}$ |
| Std4 | $300 \mu \mathrm{~L}$ of Std2 | $300 \mu \mathrm{~L}$ | $250 \mathrm{pg} / \mathrm{mL}$ |
| Std5 | $300 \mu \mathrm{~L}$ of Std3 | $300 \mu \mathrm{~L}$ | $125 \mathrm{pg} / \mathrm{mL}$ |
| Std6 | $300 \mu \mathrm{~L}$ of Std4 | $300 \mu \mathrm{~L}$ | $62.5 \mathrm{pg} / \mathrm{mL}$ |
| Std7 | $300 \mu \mathrm{~L}$ of Std5 | $300 \mu \mathrm{~L}$ | $31.25 \mathrm{pg} / \mathrm{mL}$ |
| Blank | - | $300 \mu \mathrm{~L}$ of Std6 | $15.625 \mathrm{pg} / \mathrm{mL}$ |

## Preparation of Quality Control $A$ and $B$

Reconstitute the lyophilized human serum Quality Controls in deionized/distilled water, for the volume information see the Certificate of Analysis. Let the QCs rehydrate for 15 min and dilute them 1:2 in Dilution Buffer, prior to use, see Preparation of samples.

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Preparation of Wash Buffer 1x

Prepare a working solution of Wash Buffer by adding 50 mL of Wash Buffer 15 x conc. to 700 mL of deionized/ distilled water ( $\mathrm{dH}_{2} \mathrm{O}$ ). Mix well. Store at $4^{\circ} \mathrm{C}$ for two weeks or at $-20^{\circ} \mathrm{C}$ for long term storage.

## Preparation of samples

Human serum or plasma may be used with this assay. For long-term storage the samples should be frozen at minimum $-70^{\circ} \mathrm{C}$. Lipemic or haemolytic samples may cause false results.

Recommended dilution of samples is $1: 2$, i.e., $70 \mu \mathrm{~L}$ of sample $+70 \mu \mathrm{~L}$ of Dilution Buffer for singlets and $150 \mu \mathrm{~L}$ of sample $+150 \mu \mathrm{~L}$ of Dilution Buffer for duplicates.

Do not store the diluted samples.

## ASSAY PROCEDURE

1. Prepare the reagents as described in the previous chapter.
2. Pipette $100 \mu \mathrm{~L}$ of set of Standards, Quality Controls, diluted Samples and Dilution Buffer = Blank into each well. Incubate OVER NIGHT at $4^{\circ} \mathrm{C} \pm 2^{\circ} \mathrm{C}$, NO shaking.
3. Wash the wells 3 -times with 1 x Wash Buffer ( $350 \mu \mathrm{~L} /$ well). When finished, tap the plate against the paper towel to remove the liquid completely.
4. Pipette $100 \mu \mathrm{~L}$ of HRP-labelled Antibody Conjugate into each well. Incubate for $\mathbf{2}$ hours at $25^{\circ} \mathrm{C} \pm 2^{\circ} \mathrm{C}$, shaking at $\mathbf{5 0 0} \mathbf{~ r p m}$.
5. Wash the wells as described in point 3 .
6. Pipette $100 \mu \mathrm{~L}$ Substrate solution, incubate for $\mathbf{2 5} \mathbf{~ m i n}$, at $25^{\circ} \mathrm{C} \pm 2^{\circ} \mathrm{C}$. Avoid exposure to the light during this step.
7. Pipette $100 \mu \mathrm{~L}$ of STOP solution.
8. Read the signal at 450 or $450 / 630 \mathrm{~nm}$ within 15 min .

Plate layout

|  | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 | 11 | 12 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| A | Std 1 |  | Bckg | Sa 4 | Sa 8 | Sa | Sa | Sa | Sa | Sa | Sa | Sa |
| B | Std 2 |  |  |  |  | 12 | 16 | 20 | 24 | 28 | 32 | 36 |
| C | Std 3 |  | Sa 1 | Sa 5 | Sa 9 | Sa | Sa | Sa | Sa | Sa | Sa | Sa |
| D | Std 4 |  |  |  |  | 13 | 17 | 21 | 25 | 29 | 33 | 37 |
| E | Std 5 |  | Sa 2 | Sa 6 | Sa | Sa | Sa | Sa | Sa | Sa | Sa | Sa |
| F | Std 6 |  |  |  | 10 | 14 | 18 | 22 | 26 | 30 | 34 | 38 |
| G | QCA |  | Sa3 | Sa 7 | Sa | Sa | Sa | Sa | Sa | Sa | Sa | Sa |
| H |  |  |  |  | 11 | 15 | 19 | 23 | 27 | 31 | 35 | 39 |

## PERFORMANCE CHARACTERISTICS

Samples used in the tests were diluted 1:2 as recommended and assayed. The results are multiplied by the dilution factor.

1. Sensitivity

The limit of detection, defined as a concentration of human KIM-1 giving absorbance higher than absorbance of blank +3 standard deviations, is better than $45 \mathrm{pg} / \mathrm{mL}$ of sample.

## 2. Precision

Intra-assay

| Sample | Mean (pg/mL) | SD | CV (\%) |
| :---: | :---: | :---: | :---: |
| 1 | 519 | 8.8 | 1.7 |
| 2 | 287 | 15.6 | 5.4 |

Inter-assay (Run - to - run)

| Sample | Mean (pg/mL) | SD | CV (\%) |
| :---: | :---: | :---: | :---: |
| 1 | 284 | 16 | 5.7 |
| 2 | 114 | 15 | 13.3 |

Version: ENG. 01
3. Accuracy

Dilution linearity

| Sample | Dilution | Measured <br> concentration <br> $(\mathrm{pg} / \mathrm{mL})$ | Expected <br> concentration <br> $(\mathrm{pg} / \mathrm{mL})$ | Yield <br> $(\%)$ |
| :---: | :---: | :---: | :---: | :---: |
| 1 |  | 509 | - | - |
|  | 2 x | 230 | 254 | 90 |
|  | 4 x | 106 | 127 | 83 |
|  | 8 x | 52 | 64 | 82 |
|  |  | 448 | - | - |
|  | 2 x | 196 | 224 | 87 |
|  | 4 x | 99 | 112 | 89 |
|  | 8 x | 54 | 56 | 96 |

## Spiking Recovery

| Sample | Spike <br> $(\mathrm{ng} / \mathrm{mL})$ | Measured <br> concentration <br> $(\mathrm{pg} / \mathrm{mL})$ | Expected <br> concentration <br> $(\mathrm{pg} / \mathrm{mL})$ | Yield <br> $(\%)$ |
| :---: | :---: | :---: | :---: | :---: |
| 1 | - | 236 | - | - |
|  | 360 | 663 | 596 | 111 |
|  | 180 | 446 | 416 | 107 |
|  | 90 | 341 | 326 | 105 |

## Typical standard curve

The standard curve needs to be measured in every test. Most of the microplate reader can automatically calculate the analyte concentration using 4-parameter algorithm or alternative functions to fit the standard points properly. The concentrations need to be multiplied by the dilution factor, either automatically by reader or manually.


[^0]
[^0]:    RESOURCES
    ${ }^{1}$ Cuesta C, Fuentes-Calvo I, Sancho-Martinez SM, Valentijn FA, Düwel A, Hidalgo-Thomas OA, Agüeros-Blanco C, Benito-Hernández A, Ramos-Barron MA, Gómez-Alamillo C, Arias M, Nguyen TQ, Goldschmeding R, Martínez-Salgado C, Lopez-Hernandez FJ. Urinary KIM-1 Correlates with the Subclinical Sequelae of Tubular Damage Persisting after the Apparent Functional Recovery from Intrinsic Acute Kidney Injury. Biomedicines. 2022 May 10;10(5):1106. doi: 10.3390/biomedicines10051106. PMID: 35625842; PMCID: PMC9139078

