# **Human Kidney Injury Molecule 1 ELISA**

Cat. No.: BA1001

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.<sup>1</sup>

### 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 ( $H_2SO_4$ ).

The absorbance values are measured at 450 nm (optionally 450/630 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 µL of Standards, diluted QCs and Samples to the wells



Incubate OVER NIGHT at 4°C, NO shaking

3-times wash the wells (350  $\mu$ L/well)



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



Incubate for 2 hours at 25°C, shaking at 500 rpm

3-times wash the wells (350 µL/well)



Add 100  $\mu\text{L}$  of Substrate Solution to the wells



Incubate for 25 min in the dark at 25°C,

NO shaking

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



Read the signal at 450 nm (450/630 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

- Glassware and test tubes
- 2. Microtiter plate washer
- 3. Precision pipettes (various volumes) with tips
- 4. Orbital shaker
- Microtiter plate reader capable of measuring absorbance at 450 nm or 450/630 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
- To prevent cross sample contamination, use disposable labware and pipette tips
- To protect laboratory stuff, wear protective gloves and protective clothing
- The substrate solution should remain colourless, keep it protected from light
- The test should be performed at standard laboratory conditions (temperature 25°C ±2°C).

### STORAGE CONDITIONS

- The kit must be stored at 2 8°C.
- 2. The opened components can be stored for one week at  $2-8^{\circ}$ 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°C ±2°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 pg/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.

Buffer   Std1   Standard 1000 pg/mL   See CofA   1000 pg/mL   (lyophilised)   See CofA   1000 pg/mL   Std2   300 μL of Std1   300 μL   500 pg/mL   Std3   300 μL of Std2   300 μL   250 pg/mL   Std4   300 μL of Std3   300 μL   125 pg/mL   Std5   300 μL of Std4   300 μL   62.5 pg/mL   Std6   300 μL of Std5   300 μL   31.25 pg/mL   Std7   300 μL of Std6   300 μL   15.625 pg/mL				
Std1         Standard 1000 pg/mL (lyophilised)         See CofA         1000 pg/mL           Std2         300 μL of Std1         300 μL         500 pg/mL           Std3         300 μL of Std2         300 μL         250 pg/mL           Std4         300 μL of Std3         300 μL         125 pg/mL           Std5         300 μL of Std4         300 μL         62.5 pg/mL           Std6         300 μL of Std5         300 μL         31.25 pg/mL           Std7         300 μL of Std6         300 μL         15.625 pg/mL		Volume of Standard	Dilution	Concentration
(Iyophilised)       Std2     300 μL of Std1     300 μL     500 pg/ml       Std3     300 μL of Std2     300 μL     250 pg/ml       Std4     300 μL of Std3     300 μL     125 pg/ml       Std5     300 μL of Std4     300 μL     62.5 pg/ml       Std6     300 μL of Std5     300 μL     31.25 pg/ml       Std7     300 μL of Std6     300 μL     15.625 pg/ml			Buffer	
Std2       300 μL of Std1       300 μL       500 pg/ml         Std3       300 μL of Std2       300 μL       250 pg/ml         Std4       300 μL of Std3       300 μL       125 pg/ml         Std5       300 μL of Std4       300 μL       62.5 pg/ml         Std6       300 μL of Std5       300 μL       31.25 pg/ml         Std7       300 μL of Std6       300 μL       15.625 pg/ml	Std1	Standard 1000 pg/mL	See CofA	1000 pg/mL
Std3       300 μL of Std2       300 μL       250 pg/ml         Std4       300 μL of Std3       300 μL       125 pg/ml         Std5       300 μL of Std4       300 μL       62.5 pg/ml         Std6       300 μL of Std5       300 μL       31.25 pg/ml         Std7       300 μL of Std6       300 μL       15.625 pg/ml		(lyophilised)		
Std4       300 μL of Std3       300 μL       125 pg/ml         Std5       300 μL of Std4       300 μL       62.5 pg/ml         Std6       300 μL of Std5       300 μL       31.25 pg/ml         Std7       300 μL of Std6       300 μL       15.625 pg/ml	Std2	300 μL of Std1	300 μL	500 pg/mL
Std5       300 μL of Std4       300 μL       62.5 pg/ml         Std6       300 μL of Std5       300 μL       31.25 pg/ml         Std7       300 μL of Std6       300 μL       15.625 pg/ml	Std3	300 μL of Std2	300 μL	250 pg/mL
Std6       300 μL of Std5       300 μL       31.25 pg/ml         Std7       300 μL of Std6       300 μL       15.625 pg/ml	Std4	300 μL of Std3	300 μL	125 pg/mL
Std7 300 μL of Std6 300 μL 15.625 pg/ml	Std5	300 μL of Std4	300 μL	62.5 pg/mL
200   2	Std6	300 μL of Std5	300 μL	31.25 pg/mL
Blank _ 250 μL 0 ng/mL	Std7	300 μL of Std6	300 μL	15.625 pg/mL
	Blank	-	250 μL	0 ng/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.

# BioLab Assays

### Preparation of Wash Buffer 1x

Prepare a working solution of Wash Buffer by adding 50 mL of Wash Buffer 15x conc. to 700 mL of deionized/ distilled water (dH $_2$ O). Mix well. Store at 4°C for two weeks or at -20°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°C. Lipemic or haemolytic samples may cause false results.

Recommended dilution of samples is 1:2, i.e., 70  $\mu L$  of sample + 70  $\mu L$  of Dilution Buffer for singlets and 150  $\mu L$  of sample + 150  $\mu L$  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  $\mu$ L of set of Standards, Quality Controls, diluted Samples and Dilution Buffer = Blank into each well. Incubate OVER NIGHT at 4°C  $\pm$ 2°C, NO shaking.
- Wash the wells 3-times with 1x Wash Buffer (350 μL/well). When finished, tap the plate against the paper towel to remove the liquid completely.
- 4. Pipette 100  $\mu$ L of HRP-labelled Antibody Conjugate into each well. Incubate for **2 hours** at 25°C  $\pm$ 2°C, shaking at **500 rpm**.
- 5. Wash the wells as described in point 3.
- Pipette 100 μL Substrate solution, incubate for 25 min, at 25°C ±2°C. Avoid exposure to the light during this step.
- 7. Pipette 100 μL of STOP solution.
- 8. Read the signal at 450 or 450/630 nm within 15 min.

## Plate layout

	1	2	3	4	5	6	7	8	9	10	11	12
Α	Std 1		Bckg	Sa 4	Sa 8	Sa						
В	Std 2					12	16	20	24	28	32	36
С	Std 3		Sa 1	Sa 5	Sa 9	Sa						
D	Std 4					13	17	21	25	29	33	37
Е	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
Н	QCB				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 pg/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

### 3. Accuracy

# Dilution linearity

Sample	Dilution	Measured concentration (pg/mL)	Expected concentration (pg/mL)	Yield (%)
1		509	-	-
	2x	230	254	90
	4x	106	127	83
	8x	52	64	82
2		448	-	-
	2x	196	224	87
	4x	99	112	89
	8x	54	56	96

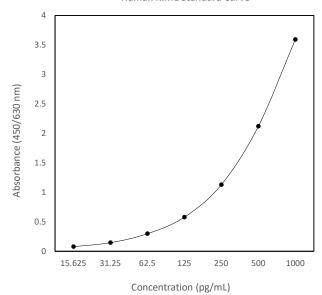
#### Spiking Recovery

Sample	Spike (ng/mL)	Measured concentration (pg/mL)	Expected concentration (pg/mL)	Yield (%)
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.

# Human KIM1 Standard Curve



### RESOURCES

<sup>1</sup> 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, López-Hernández 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.