

Human ACE2 Sandwich ELISA Kit Datasheet

Please read it entirely before use

Catalogue Number: KE00254

Size: 5*96T

Sensitivity: 0.15 pg/mL Range: 6.25 - 400 pg/mL

Usage: For the quantitative detection of human ACE2 concentrations in serum, plasma and urine.

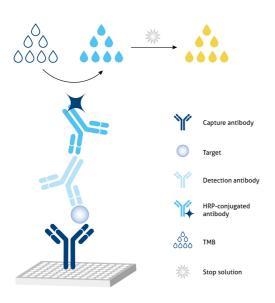
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1. Background

Angiotensin-converting enzyme 2 (ACE2), a critical regulator of the renin-angiotensin system (RAS), belongs to the peptidase M2 family. It plays an important role in cardiovascular homeostasis by regulating vascular tone, fluid, and electrolyte balance. ACE2 functions as a carboxymonopeptidase hydrolyzing the cleavage of a single C-terminal residue from Angiotensin-II (Ang-II), the key peptide hormone of RAS, to form Angiotensin-(1-7) (Ang-(1-7)), which binds to the G-protein-coupled Mas receptor and activates signaling pathways that counteract the pathways activated by Ang-II. ACE2 is expressed in a variety of tissues, including the kidneys, testes, heart, and intestines, and is particularly enriched in lung epithelium. Recent studies have shown that increased ACE2 expression is likely to help prevent secondary fibrosis changes following COVID-19 pneumonia.

2. Principle



Sandwich ELISA structure (HRP conjugated secondary antibody)

A capture antibody is pre-coated onto the bottom of wells which binds to analyte of interest. A detection antibody also binds to the analyte. Horseradish peroxidase (HRP)-conjugated secondary antibody binds to the detection antibody. TMB acts as the HRP substrate and the solution color will change from colorless to blue. A stop solution containing sulfuric acid turns solution yellow. The color intensity is proportional to the quantity of bound protein which is measurable at 450 nm with the correction wavelength set at 630 nm.

3. Required Materials

- 3.1 A microplate reader capable of measuring absorbance at 450 nm with the correction wavelength set at 630 nm.
- 3.2 Calibrated, adjustable precision pipettes and disposable plastic tips. A manifold multi-channel pipette is recommended for large assays.
- 3.3 Plate washer: automated or manual.
- 3.4 Absorbent paper towels.
- 3.5 Glass or plastic tubes to prepare standard and sample dilutions.
- 3.6 Beakers and graduated cylinders.
- 3.7 Log-log or semi-log graph paper or computer and software for ELISA data analysis. A four-parameter logistic (4-PL) curve-fit is recommended.

4. Kit Components and Storage

Microplate - antibody coated 96-well microplate (8 well × 12 strips)	5 plates	Unopened Kit:	
Protein standard - 800 pg/bottle; lyophilized	10 bottles	•	
Detection Antibody (100×) - 600 µL/vial*	1 vial	Store at 2-8°C for 6 months or -	
HRP-conjugated antibody (100×) - 600 µL/vial*	1 vial	20°C for 12 months.	
Sample Diluent PT 3 - 150 mL/bottle	1 bottle	Opened Kit:	
Detection Diluent - 150 mL/bottle	1 bottle	All reagents stored at 2-8°C for	
Wash Buffer Concentrate (20×) - 150 mL/bottle	1 bottle		
Tetramethylbenzidine Substrate (TMB) - 60 mL/bottle	1 bottle	7 days.	
Stop Solution - 60 mL/bottle	1 bottle	Please use a new standard	
Plate Cover Seals	15 pieces	for each assay.	

^{*} Centrifugation immediately before use

5. Safety Notes

- 5.1 Avoid any skin and eye contact with Stop Solution and TMB. In case of contact, wash thoroughly with water.
- 5.2 Do not use the kit after the expiration date.
- 5.3 Do not mix or substitute reagents or materials from other kit lots or other sources.
- 5.4 Be sure to wear protective equipment such as gloves, masks and goggles during the experiment.
- 5.5 When using an automated plate washer, adding a 30 second soak period following the addition of Wash Buffer to improve assay precision

6. Sample Collection and Storage

- 6.1 Serum: Allow blood samples to clot for 30 minutes, followed by centrifugation for 15 minutes at 1000xg. Clear serum can be assayed immediately or aliquoted and stored at -20°C. Avoid repeated freeze-thaw cycles.
- 6.2 Plasma: Use EDTA, heparin, or citrate as an anticoagulant for plasma collection. Centrifuge for 15 minutes at 1000xg within 30 minutes of collection. The plasma can be assayed immediately or aliquoted and stored at -20°C. Avoid repeated freeze-thaw cycles.
- 6.3 Urine: Collect urine samples and centrifuge for 20 minutes at 1000 \times g. Collect the aqueous layer, assay immediately or aliquot and store samples at \leq -20°C. Avoid repeated freeze-thaw cycles.

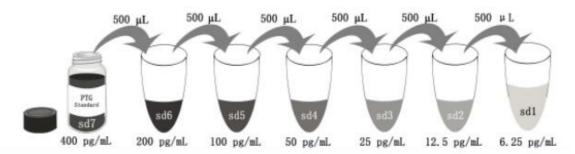
7. Regent Preparation

- 7.1 Wash Buffer (1X): If crystals have formed in the concentrate, warm to room temperature and mix gently until the crystals have completely dissolved. Add 30 mL of Wash Buffer Concentrate(20X) to 570 mL deionized or distilled water to prepare 1X Wash Buffer.
- **7.2 Detection Antibody (1X):** Dilute 100X Detection Antibody 1:100 using Detection Diluent prior to assay. Suggested 1:100 dilution: $10 \,\mu$ L 100X Detection Antibody + 990 μ L Detection Diluent (Centrifuge the 100 X Detection Antibody solution for a few seconds prior to use).
- **7.3 HRP-conjugated antibody (1X):** Dilute 100X HRP-conjugated antibody 1:100 using Detection Diluent prior to assay. Suggested 1:100 dilution: $10 \,\mu$ L 100X HRP-conjugated antibody + 990 μ L Detection Diluent (Centrifuge the 100X HRP-conjugated antibodyy solution for a few seconds prior to use).
- **7.4 Sample Dilution:** Different samples should be diluted with corresponding Sample Diluent, samples may require further dilution if the readout values are higher than the highest standard OD reading. Variations in sample collection, processing and storage may affect the results of the measurement.

Recommended Dilution for different sample types: 1:2 or 1:4 is recommended for human serum and plasma; 1:2 is recommended for urine.

7.5 Standard Serial Dilution:

Add 2 mL Sample Diluent PT 3 in protein standard.



Add # µL of Standard diluted in the previous step	T	500 μL					
# µL of Sample Diluent PT 3	2000 μL	500 μL					
	"sd7"	"sd6"	"sd5"	"sd4"	"sd3"	"sd2"	"sd1"

8. Assay Procedure Summary

Bring all reagents to room temperature before use (Detection antibody and HRP-conjugated antibody can be used immediately). To avoid cross-contamination, change pipette tips between additions of each standard level, between sample additions, and between reagent additions. Also, use separate reservoirs for each reagent.

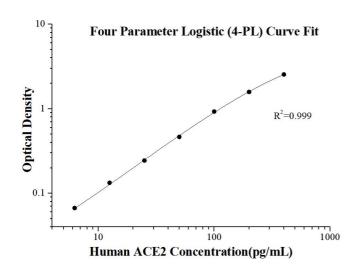
- 8.1 Take out the required number of microplate strips and return excess strips to the foil pouch containing the drying reagent pack and reseal; store at 4°C immediately. Microplate strips should be used in one week.
- 8.2 Preset the layout of the microplate, including control group, standard group and sample group, add 100 µL of each standard and sample to the appropriate wells. (Make sure sample addition is uninterrupted and completed within 5 to 10 minutes, It is recommended to assay all standards, controls, and samples in duplicate).
- 8.3 Seal plate with cover seal, pressing it firmly onto top of microwells. Incubate the plate for 2 hours at 37°C.
- 8.4 Wash
- 1) Gently remove the cover seal. Discard the liquid from wells by aspirating or decanting. Remove any residual solution by tapping the plate a few times on fresh paper towels.
- 2) Wash 4 times with 1X Wash Buffer, using at least 350-400 μ L per well. Following the last wash, firmly tap plates on fresh towels 10 times to remove residual Wash Buffer. Avoid getting any towel fibers in the wells or wells drying out completely.
- $8.5 \text{ Add } 100 \,\mu\text{L}$ of 1X Detection Antibody solution (refer to Reagent Preparation7.2) to each well. Seal plate with cover seal and incubate for 1 hour at 37°C.
- 8.6 Repeat wash step in 8.4.
- $8.7~\text{Add}\ 100~\mu\text{L}$ of 1X HRP-conjugated antibody solution (refer to Reagent Preparation 7.3) to each well. Seal plate with cover seal and incubate the plate for 40 minutes at 37°C .
- 8.8 Repeat wash step in 8.4.
- 8.9 Signal development: Add 100 μ L of TMB substrate solution to each well, protected from light. Incubate for 15 to 20 minutes. Substrate Solution should remain colorless until added to the plate.
- 8.10 Quenching color development: Add 100 µL of Stop Solution to each well in the same order as addition of the TMB substrate. Mix by tapping the side of the plate gently. NB: Avoid skin and eye contact with the Stop solution.
- 8.11 Read results: Immediately after adding Stop solution read the absorbance on a microplate reader at a wavelength of 450 nm. If possible, perform a double wavelength readout (450 nm and 630 nm).
- 8.12 Data analysis: Calculate the average of the duplicate readings (OD value) for each standard and sample, and subtract the average of the zero standard absorbance. Construct a standard curve by plotting the mean absorbance for each standard on the y-axis against the concentration on the x-axis, use four-parameter logistic curve- fit (4-PL) analysis to do this. If the samples have been diluted, the OD readout from the standard curve must be multiplied by the dilution factor used.

Step	Reagent	Volume	Incubation	Wash	Notes
1	Standard and Samples	100 µL	120 min	4 times	Cover Wells incubate at 37°C
2	Diluent Antibody Solution	100 µL	60 min	4 times	Cover Wells incubate at 37°C
3	Diluent HRP Solution	100 µL	40 min	4 times	Cover Wells incubate at 37°C
4	TMB Substrate	100 µL	15-20 min	Do not wash	Incubate in the dark at 37°C
5	Stop Solution	100 µL	0 min	Do not wash	-
6	Read plate at 450 nm and 630 nm immediately after adding Stop solution. DO NOT exceed 5 minutes.				

9. Validation Data

9.1 Standard curve

These standard curves are provided for demonstration only. A standard curve should be generated for each set of samples assayed.



(pg/mL)	0.D	Average	Corrected
0	0.07 0.069	0.0695	-
6.25	0.138 0.135	0.1365	0.067
12.5	0.21 0.196	0.203	0.1335
25	0.325 0.303	0.314	0.2445
50	0.544 0.524	0.534	0.4645
100	0.993 1.003	0.998	0.9285
200	1.673 1.633	1.653	1.5835
400	2.616 2.621	2.6185	2.549

9.2 Precision

Intra-assay Precision (Precision within an assay) Three samples of known concentration were tested 20 times on one plate to assess intra-assay precision.

Inter-assay Precision (Precision between assays) Three samples of known concentration were tested in 24 separate assays to assess inter-assay precision.

Intra-assay Precision					
Sample	Sample n Mean (pg/mL) SD CV%				
1	20	181.9	6.0	3.3	
2	20	46.9	1.5	3.1	
3	20	11.4	0.7	6.0	

	Inter-assay Precision				
Sample	n	Mean (pg/mL)	SD	CV%	
1	24	191.7	12.0	6.3	
2	24	47.6	2.5	5.2	
3	24	12.3	1.0	8.2	

9.3 Recovery

The recovery of human ACE2 spiked to three different levels throughout the range of the assay in various matrices was evaluated.

Sample Type		Average% of Expected	Range (%)
Human placma	1:2	106	90-124
Human plasma	1:4	111	92-121
Urine	1:16	84	77-95
Offile	1:32	86	75-100

9.4 Sample values

Human serum, plasma and urine samples were evaluated for the presence of human ACE2 in this assay.

Sample Type	Mean (pg/mL)	Range (pg/mL)
Human serum (n=16)	33.2	12.5-134.4
Human plasma (n=16)	60.6	10.0-355.8
Urine (n=8)	93.4	7.6-331.0

9.5 Sensitivity

The minimum detectable dose of Human ACE2 is 0.15 pg/mL. This was determined by adding two standard deviations to the concentration corresponding to the mean O.D. of 20 zero standard replicates.

9.6 Linearity

To assess the linearity of the assay, samples were diluted with the appropriate **Sample Diluent** to produce samples with values within the dynamic range of the assay.

(The urine samples were initially diluted 1:8.)

		Human plasma	Urine
1.2	Average% of Expected	100	100
1:2	Range (%)	-	-
1.7	Average% of Expected	92	87
1:4	Range (%)	90-93	85-89
1.0	Average% of Expected	89	85
1:8	Range (%)	80-99	84-85
1:16	Average% of Expected	94	86
1.10	Range (%)	89-99	80-92

9.7 Specificity

This assay recognizes natural and recombinant human ACE2.

The following factors prepared at 50 ng/mL were assayed and exhibited no cross-reactivity or interference.

Recombinant human

Angiotensinogen

HAT

IL-4

IFN-γ

Neprilysin

Renin

TACE

A sample containing 50 ng/mL of the recombinant human ACE reads as 2.71 pg/mL (0.005% cross-reactivity).

10. References

- 1. Li Q. et al. (2020) Clin Sci (Lond). 134:2581-2595.
- 2. Li Y. et al. (2020) Biochem Biophys Res Commun. 526:947-952.
- 3. Hikmet F. et al. (2020) Mol Syst Biol. 16:e9610.
- 4. Chaudhry F. et al. (2020) Open Heart. 7:e001424.