

# Human HSP47/SERPINH1 Sandwich ELISA Kit Datasheet

Please read it entirely before use

Catalogue Number: KE00365

Size: 96T

**Sensitivity:** 0.04 ng/mL **Range:** 0.47-30 ng/mL

**Usage:** For the quantitative detection of human HSP47/SERPINH1 concentration in cell lysate.

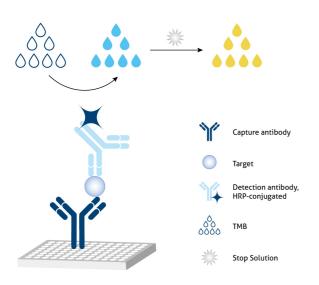
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## 1. Background

HSP47 is also named as Serpin H1, colligen, CBP1, CBP2, HSP47, SERPINH2 and belongs to the serpin family. This gene encodes a member of the serpin superfamily of serine proteinase inhibitors. The encoded protein is localized to the endoplasmic reticulum. It binds specifically to collagen and can be involved as a chaperone in the biosynthetic pathway of collagen. The Serpin Family H Member 1 (SERPINH1) gene encodes a protein called HSP47, which is required for the correct folding and secretion of collagen. SERPINH1 has been implicated in several collagen-related diseases, such as, brittle bone disease, keloid scars, and lung fibrosis. SERPINH1 also correlates with the grade of glioma, and regulates proliferation, migration and invasion of glioma cell lines. SERPINH1 regulates the Wnt/β-catenin signaling pathway and modulates gastric cancer (GC) progression SERPINH1 is aberrantly expressed in cervical cancer, breast cancer, glioblastoma and colorectal cancer. SERPINH1 promotes invasion and metastasis of breast cancer cells by regulating the expression of several extracellular matrix (ECM) proteins .

## 2. Principle



# Sandwich ELISA structure (Detection antibody labeled with HRP)

A capture antibody is pre-coated onto the bottom of wells which binds to analyte of interest. A detection antibody labeled with HRP also binds to the analyte. 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)	1 plate	Unopened Kit:
Protein standard - 60 ng/bottle; lyophilized	2 bottles	•
Detection antibody, HRP-conjugated (100×) - 120 µL/vial*	1 vial	Store at 2-8°C for 6 months or -
Sample Diluent PT 3B1 - 30 mL/bottle	1 bottle	20°C for 12 months.
Detection Diluent - 30 mL/bottle	1 bottle	Opened Kit:
Wash Buffer Concentrate (20×) - 30 mL/bottle	1 bottle	All reagents stored at 2-8°C for
Extraction Reagent - 30 mL/bottle	1 bottle	
Tetramethylbenzidine Substrate (TMB) - 12 mL/bottle	1 bottle	7 days.
Stop Solution - 12 mL/bottle	1 bottle	Please use a new standard
Plate Cover Seals	4 pieces	for each assay.

<sup>\*</sup> 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 Cell Lysate:

- 1) Collect cells and wash by centrifuging at 500 x g for 5 minutes before resuspension in pre-cooled PBS buffer. Perform this step three times.
- 2) Count cells and then discard the supernatant.
- 3) Add protease inhibitor cocktail to the Extraction Reagent to a final concentration immediately prior to performing cell lysis.
- 4) Add 1 mL of Extraction reagent (containing protease inhibitor cocktail) Per 1 x 107 cells, Incubate cell suspension on ice for 30 minutes, use ultrasound to treat the samples.
- 5) Centrifuge cell lysate at 10,000 x g for 5 minutes at 4°C.
- 6) Measure the concentration of total protein in cell lysate using BCA assay. Where possible, keep samples on ice to avoid protein degradation.

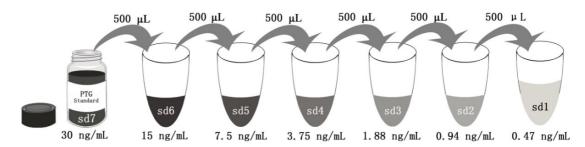
## 7. Regent Preparation

- **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, HRP-conjugated(1X):** Dilute 100X Detection Antibody, HRP-conjugated 1:100 using Detection Diluent prior to assay. Suggested 1:100 dilution:  $10 \,\mu$ L 100X Detection Antibody, HRP-conjugated + 990  $\mu$ L Detection Diluent (Centrifuge the 100X Detection Antibody solution, HRP-conjugated for a few seconds prior to use).
- **7.3 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: 1:16 or 1:32 is recommended for cell lysate.

#### 7.4 Standard Serial Dilution:

Add 2 mL Sample Diluent PT 3B1 in protein standard.



Add # µL of Standard diluted in the previous step	_	500 μL					
# μL of Sample Diluent PT 3B1	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, HRP-conjugated 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.

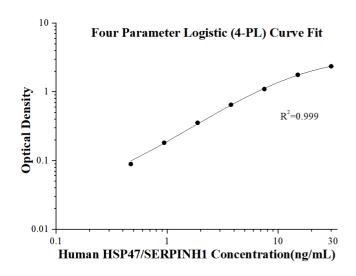
- 8.1 Take out the required number of microplate strips and return excess strips to the foil pouch containing the drying reagent pack and reseal; store at 4°C immediately. Microplate strips should be used in one week.
- 8.2 Preset the layout of the microplate, including control group, standard group 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  $\mu$ 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 Add 100  $\mu$ L of 1X Detection antibody, HRP-conjugated solution (refer to Reagent Preparation7.2) to each well. Seal plate with cover seal and incubate for 40 minutes at 37°C.
- 8.6 Repeat wash step in 8.4.
- 8.7 Signal development: Add 100  $\mu$ 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.
- 8.8 Quenching color development: Add 100  $\mu$ L 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.
- 8.9 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).
- 8.10 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 Detection antibody, HRP-conjugated Solution	100 µL	40 min	4 times	Cover Wells incubate at 37°C
3	TMB Substrate	100 µL	15-20 min	Do not wash	Incubate in the dark at 37°C
4	Stop Solution 100 µL 0 min Do not wash -				
5	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.



(ng/mL)	0.D	Average	Corrected
0	0.0319 0.0345	0.0332	-
0.47	0.1238 0.1216	0.1227	0.0895
0.94	0.232 0.1995	0.21575	0.18255
1.88	0.4098 0.3686	0.3892	0.356
3.75	0.7014 0.6679	0.68465	0.65145
7.5	1.1456 1.1245	1.13505	1.10185
15	1.8233 1.8033	1.8133	1.7801
30	2.3785 2.4251	2.4018	2.3686

#### 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	n	Mean (ng/mL)	SD	CV%
1	20	13.78	1.10	8.00
2	20	3.36	0.29	8.57
3	20	0.79	0.06	7.95

Inter-assay Precision				
Sample	n	Mean (ng/mL)	SD	CV%
1	24	15.77	1.08	6.86
2	24	3.85	0.28	7.39
3	24	0.96	0.09	9.59

## 9.3 Recovery

The recovery of human HSP47/SERPINH1 spiked to three different levels throughout the range of the assay in cell lysate was evaluated.

Sample Type		Average% of Expected	Range (%)
Cell lysate	1:32	109	104-118

## 9.4 Sample values

#### Cell lysates

	Human HSP47/SERPINH1 (ng/mL)	Total protein (mg/mL)
HepG2 cell lysate	17.99	2.50
HeLa cell lysate	56.94	3.20
MCF-7 cell lysate	35.11	3.50

## 9.5 Sensitivity

The minimum detectable dose of human HSP47/SERPINH1 is 0.04 ng/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 cell lysate samples were initially diluted 1:8.)

		Cell lysate
	Average% of Expected	100
1:2	Range (%)	-
1./	Average% of Expected	101
1:4	Range (%)	91-112
1:8	Average% of Expected	118
	Range (%)	108-129
1:16	Average% of Expected	118
	Range (%)	110-126

## 9.7 Specificity

This kit specifically recognizes native and recombinant human HSP47/SERPINH1.

#### 10. References

- 1. Duarte BDP, Bonatto D. (2018) J Cancer Res Clin Oncol. 144(12):2319-2328.
- 2. Tian S, et al. (2020) Aging (Albany NY). 12(4):3574-3593.
- 3. Zhao D, et al. (2014) J Neurooncol. 118(1):39-47.
- 4. Tian S, et al. (2020) Aging (Albany NY). 12(4):3574-3593.
- 5. Zhu J, et al. (2015) Cancer Res. 75(8):1580-91.