

AssayMax™ Human NDFIP1 ELISA Kit

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For any questions regarding troubleshooting or performing the assay, please contact our support team at support@assaypro.com.

Thank you for choosing Assaypro.

Assay Summary

Step 1. Add 50 μ l of Standard or Sample per well. Incubate 2 hours.

Step 2. Wash, then add 50 μ l of Biotinylated Antibody per well. Incubate 1 hour.

Step 3. Wash, then add 50 μ l of SP Conjugate per well. Incubate 30 minutes.

Step 4. Wash, then add 50 μ l of Chromogen Substrate per well. Incubate 25 minutes.

Step 5. Add 50 μ l of Stop Solution per well. Read at 450 nm immediately.

Symbol Key



Consult instructions for use.

Assay Template

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AssayMax™ Human NEDD4 Family-Interacting Protein 1 (NDFIP1) ELISA Kit

Catalog No. EN2550-1
Sample insert for reference use only

Introduction

NEDD4 Family-Interacting Protein 1 (NDFIP1, N4WBP5) belongs to a small group of evolutionarily conserved proteins with three transmembrane domains. NEDD4 proteins are ubiquitin ligases important for protein degradation and trafficking, as well as cell growth, development, and gene regulation. Regulation of NEDD4 family catalytic activities is important for controlling cellular protein homeostasis to prevent diseases. NDFIP1 is a 221 amino acid protein with a mass of 26 kDa that contains 2 N-terminal PY motifs and 3 transmembrane regions. This Golgi-associated protein is a potential target for ubiquitination by the NEDD4 family of proteins. NDFIP1 may act as an adaptor for NEDD4-like proteins and their targets to control ubiquitin-dependent protein sorting and trafficking (1). NDFIP1 activates NEDD4-1 by relief of WW domain linker autoinhibition (2). In the brain, both NDFIP1 and NEDD4 are co-expressed in the same neuron, and their interaction is crucial for increasing the survival of cortical neurons during injury (3).

Principle of the Assay

The AssayMax™ Human NDFIP1 ELISA (Enzyme-Linked Immunosorbent Assay) Kit is designed for detection of NDFIP1 in human CSF, cell culture, cell lysate, and tissue samples. This assay employs a quantitative sandwich enzyme immunoassay technique that measures human NDFIP1 in approximately 4 hours. A polyclonal antibody specific for human NDFIP1 has been pre-coated onto a 96-well microplate with removable strips. NDFIP1 in standards and samples is sandwiched by the immobilized antibody and a biotinylated polyclonal antibody specific for human NDFIP1, which is recognized by a streptavidin-peroxidase (SP) conjugate. All unbound material is washed away and a peroxidase enzyme substrate is added. The color development is stopped and the intensity of the color is measured.

Caution and Warning

 This product is for Research Use Only and is not intended for use in diagnostic procedures.

- Prepare all reagents (diluent buffer, wash buffer, standard, biotinylated antibody, and SP conjugate), as instructed, prior to running the assay.
- Prepare all samples prior to running the assay. The dilution factors for the samples are suggested in this insert. However, the user should determine the optimal dilution factor.
- Spin down the SP conjugate vial, the biotinylated antibody vial, and the standard diluent vial before opening and using contents.
- The Stop Solution is an acidic solution.
- The kit should not be used beyond the expiration date.

Reagents

- Human NDFIP1 Microplate: A 96-well polystyrene microplate (12 strips of 8 wells) coated with a polyclonal antibody against human NDFIP1.
- Sealing Tapes: Each kit contains 3 precut, pressure sensitive sealing tapes that can be cut to fit the format of the individual assay.
- Human NDFIP1 Standard: Human NDFIP1 in a buffered protein base (9.6 ng, lyophilized).
- **Biotinylated Human NDFIP1 Antibody (50x):** A 50-fold concentrated biotinylated polyclonal antibody against human NDFIP1 (120 µl).
- MIX Diluent Concentrate (10x): A 10-fold concentrated buffered protein base (30 ml).
- Standard Diluent (1x): A buffered protein base with stabilizer (2 ml).
- Wash Buffer Concentrate (20x): A 20-fold concentrated buffered surfactant (30 ml, 2 bottles).
- SP Conjugate (100x): A 100-fold concentrate (80 μl).
- Chromogen Substrate (1x): A stabilized peroxidase chromogen substrate tetramethylbenzidine (7 ml).
- Stop Solution (1x): A 0.5 N hydrochloric acid solution to stop the chromogen substrate reaction (11 ml).

Storage Condition

- Upon arrival, immediately store components of the kit at recommended temperatures up to the expiration date.
- Store Standard, SP Conjugate, and Biotinylated Antibody at -20°C.
- Store Microplate, Diluent Concentrate (10x), Standard Diluent (1x), Wash Buffer, Stop Solution, and Chromogen Substrate at 2-8°C.
- Unused microplate wells may be returned to the foil pouch with the desiccant packs and resealed. May be stored for up to 30 days in a vacuum desiccator.

Other Supplies Required

- Microplate reader capable of measuring absorbance at 450 nm
- Pipettes (1-20 μl, 20-200 μl, 200-1000 μl, and multiple channel)
- Deionized or distilled reagent grade water

Sample Collection, Preparation, and Storage

- **CSF:** Collect cerebrospinal fluid (CSF) using sample pot. Centrifuge samples at 3000 x g for 10 minutes. A 2-fold sample dilution is suggested into MIX Diluent; however, user should determine optimal dilution factor depending on application needs. The undiluted samples can be stored at -80°C for up to 3 months. Avoid repeated freeze-thaw cycles.
- Cell Culture Supernatant: Centrifuge cell culture media at 1500 rpm for 10 minutes at 4°C to remove debris and collect supernatant. If necessary, dilute samples into MIX Diluent; user should determine optimal dilution factor depending on application needs. The undiluted samples can be stored at -80°C. Avoid repeated freeze-thaw cycles.
- Cell Lysate: Rinse cell with cold PBS and then scrape the cell into a tube with 5 ml of cold PBS and 0.5 M EDTA. Centrifuge suspension at 1500 rpm for 10 minutes at 4°C and aspirate supernatant. Resuspend pellet in ice-cold Lysis Buffer (PBS, 1% Triton X-100, protease inhibitor cocktail). For every 1 x 10⁶ cells, add approximately 100 μl of ice-cold Lysis Buffer. Incubate on ice for 60 minutes. Centrifuge at 13000 rpm for 30 minutes at 4°C and collect supernatant. If necessary, dilute samples into MIX Diluent; user should determine optimal dilution factor depending on application needs. The undiluted samples can be stored at -80°C. Avoid repeated freeze-thaw cycles.
- **Tissue:** Extract tissue samples with 0.1 M phosphate-buffered saline (pH 7.4) containing 1% Triton X-100 and centrifuge at 14000 x g for 20 minutes. Collect the supernatant and measure the protein concentration. If necessary, dilute samples into MIX Diluent; user should determine optimal dilution factor depending on application needs. Store remaining extract at -80°C. Avoid repeated freeze-thaw cycles.

Applicable samples may also include biofluids, cell culture, and tissue homogenates. If necessary, user should determine optimal dilution factor depending on application needs.

Refer to Dilution Guidelines for further instruction.

	Guidelines for Dilutions of 100-fold or Greater (for reference only; please follow the insert for specific dilution suggested)						
100x			10000x				
A)	4 μl sample : 396 μl buffer (100x) = 100-fold dilution	A) B)	4 µl sample ։ 396 µl buffer (100x) 4 µl of A ։ 396 µl buffer (100x)				
	Assuming the needed volume is less than or equal to 400 μ l.		= 10000-fold dilution Assuming the needed volume is less than or equal to 400 μl.				
	1000x		100000x				
A) B)	4 μl sample : 396 μl buffer (100x) 24 μl of A : 216 μl buffer (10x) = 1000-fold dilution	A) B) C)	4 μl sample : 396 μl buffer (100x) 4 μl of A : 396 μl buffer (100x) 24 μl of B : 216 μl buffer (10x) = 100000-fold dilution				
	Assuming the needed volume is less than or equal to 240 μl.		Assuming the needed volume is less than or equal to 240 μl.				

Reagent Preparation

- Freshly dilute all reagents and bring all reagents to room temperature before use.
- MIX Diluent Concentrate (10x): Dilute the MIX Diluent Concentrate 10fold with reagent grade water to produce a 1x solution. When diluting
 the concentrate, make sure to rinse the bottle thoroughly to extract any
 precipitates left in the bottle. Mix the 1x solution gently until the crystals
 have completely dissolved. Store for up to 30 days at 2-8°C.
- Human NDFIP1 Standard: Reconstitute the Human NDFIP1 Standard (9.6 ng) with 0.3 ml of Standard Diluent to generate a 32 ng/ml standard stock solution. Allow the vial to sit for 10 minutes with gentle agitation prior to making dilutions. Prepare duplicate or triplicate standard points by serially diluting from the standard stock solution (32 ng/ml) 2-fold with equal volume of MIX Diluent to produce 16, 8, 4, 2, 1, 0.5, and 0.25 ng/ml solutions. MIX Diluent serves as the zero standard (0 ng/ml). Aliquot remaining stock solution to limit repeated freeze-thaw cycles. This solution should be stored at -20°C and used within 30 days.

Standard Point	Dilution	[NDFIP1] (ng/ml)
P1	1 part Standard (32 ng/ml) + 1 part MIX Diluent	16
P2	1 part P1 + 1 part MIX Diluent	8.0
Р3	1 part P2 + 1 part MIX Diluent	4.0
P4	1 part P3 + 1 part MIX Diluent	2.0
P5	1 part P4 + 1 part MIX Diluent	1.0
P6	1 part P5 + 1 part MIX Diluent	0.5
P7	1 part P6 + 1 part MIX Diluent	0.25
P8	MIX Diluent	0.0

- Biotinylated Human NDFIP1 Antibody (50x): Spin down the antibody briefly and dilute the desired amount of the antibody 50-fold with MIX Diluent to produce a 1x solution. The undiluted antibody should be stored at -20°C.
- Wash Buffer Concentrate (20x): Dilute the Wash Buffer Concentrate 20fold with reagent grade water to produce a 1x solution. When diluting
 the concentrate, make sure to rinse the bottle thoroughly to extract any
 precipitates left in the bottle. Mix the 1x solution gently until the crystals
 have completely dissolved.
- SP Conjugate (100x): Spin down the SP Conjugate briefly and dilute the
 desired amount of the conjugate 100-fold with MIX Diluent to produce a
 1x solution. The undiluted conjugate should be stored at -20°C.

Assay Procedure

- Prepare all reagents, standard solutions, and samples as instructed. Bring all reagents to room temperature before use. The assay is performed at room temperature (20-25°C).
- Remove excess microplate strips from the plate frame and return them
 immediately to the foil pouch with desiccants inside. Reseal the pouch
 securely to minimize exposure to water vapor and store in a vacuum
 desiccator.
- Add 50 µl of Human NDFIP1 Standard or sample to each well. Gently tap
 plate to thoroughly coat the wells. Break any bubbles that may have
 formed. Cover wells with a sealing tape and incubate for 2 hours. Start
 the timer after the last addition.
- Wash the microplate manually or automatically using a microplate washer. Invert the plate and decant the contents; hit 4-5 times on absorbent material to completely remove the liquid. If washing manually, wash five times with 200 µl of Wash Buffer per well. Invert the plate each time and decant the contents; hit 4-5 times on absorbent material to completely remove the liquid. If using a microplate washer,

- wash six times with 300 µl of Wash Buffer per well; invert the plate and hit 4-5 times on absorbent material to completely remove the liquid.
- Add 50 µl of Biotinylated Human NDFIP1 Antibody to each well. Gently tap plate to thoroughly coat the wells. Break any bubbles that may have formed. Cover wells with a sealing tape and incubate for 1 hour.
- Wash the microplate as described above.
- Add 50 µl of SP Conjugate to each well. Gently tap plate to thoroughly coat the wells. Break any bubbles that may have formed. Cover wells with a sealing tape and incubate for 30 minutes. Turn on the microplate reader and set up the program in advance.
- Wash the microplate as described above.
- Add 50 µl of Chromogen Substrate to each well. Gently tap plate to thoroughly coat the wells. Break any bubbles that may have formed. Incubate in ambient light for 25 minutes or until the optimal blue color density develops.
- Add 50 µl of Stop Solution to each well. The color will change from blue to yellow. Gently tap plate to ensure thorough mixing. Break any bubbles that may have formed.
- Read the absorbance on a microplate reader at a wavelength of 450 nm immediately. If wavelength correction is available, subtract readings at 570 nm from those at 450 nm to correct optical imperfections.
 Otherwise, read the plate at 450 nm only. Please note that some unstable black particles may be generated at high concentration points after stopping the reaction for about 10 minutes, which will reduce the readings.

Data Analysis

- Calculate the mean value of the duplicate or triplicate readings for each standard and sample.
- To generate a standard curve, plot the graph using the standard concentrations on the x-axis and the corresponding mean 450 nm absorbance (OD) on the y-axis. The best fit line can be determined by regression analysis using log-log or four-parameter logistic curve fit.
- Determine the unknown sample concentration from the Standard Curve and multiply the value by the dilution factor.

Typical Data

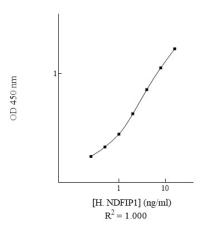
The typical data is provided for reference only. Individual laboratory
means may vary from the values listed. Variations between laboratories
may be caused by technique differences.

Standard Point	ng/ml	OD	Average OD
P1	16	1.736	1.783
	10	1.830	1.705
P2	8.0	1.110	1.146
ΓZ	8.0	1.182	1.140
P3	4.0	0.713	0.690
P3	4.0	0.667	0.090
P4	2.0	0.401	0.394
P4		0.387	0.594
P5	1.0	0.253	0.245
P3	1.0	0.237	0.245
P6	0.5	0.187	0.182
PO	0.5	0.177	0.162
P7	0.25	0.150	0.146
F /	0.25	0.142	0.140
P8	0.0	0.116	0.114
го	0.0	0.112	0.114

Standard Curve

• The curve is provided for illustration only. A standard curve should be generated each time the assay is performed.

Human NDFIP1 Standard Curve



Reference Value

 These cell lines were tested in house (n=10). The cell line averages are provided for reference only.

Cell Lysate	Dilution Factor	Average Value (ng/mg cell lysate)
293T (human embryonic kidney)	4x	3.722
HeLa (human cervical cancer)	1x	0.081
Jurkat E6-1 (human T-cell leukemia)	4x	3.282

Performance Characteristics

- This assay recognizes both natural and recombinant human NDFIP1.
- The minimum detectable dose of human NDFIP1 as calculated by 2SD from the mean of a zero standard was established to be 0.2 ng/ml.
- Intra-assay precision was determined by testing three cell lysate samples twenty times in one assay.
- Inter-assay precision was determined by testing three cell lysate samples in twenty assays.

	Intra-Assay Precision			Inter	-Assay Pred	ision
Sample	1	2	3	1	2	3
n	20	20	20	20	20	20
CV (%)	5.9%	5.2%	4.8%	11.3%	9.6%	10.2%
Average CV (%)	5.3%				10.4%	

Spiking Recovery

 Recovery was determined by spiking two cell lysate samples with different NDFIP1 concentrations.

Sample	Unspiked Sample (ng/ml)	Spiking Value (ng/ml)	Expected	Observed	Recovery (%)
		7.387	8.757	9.193	105%
1 – 293T	1.370	1.596	2.966	2.729	92%
		0.454	1.824	1.631	89%
		7.387	8.700	7.413	85%
2 – Jurkat E6-1	1.313	1.596	2.909	2.581	89%
		0.454	1.767	1.558	88%
Average Recovery (%)					91%

Troubleshooting

Issue	Causes	Course of Action			
	Use of improper	Check the expiration date listed before use.			
	components	 Do not interchange components from different lots. 			
		Check that the correct wash buffer is being used.			
		 Check that all wells are empty after aspiration. 			
	Improper wash step	 Check that the microplate washer is dispensing properly. 			
		 If washing by pipette, check for proper pipetting 			
_		technique.			
Low Precision	Splashing of reagents while loading wells	Pipette properly in a controlled and careful manner.			
J.e	Inconsistent volumes	 Pipette properly in a controlled and careful manner. 			
₹	loaded into wells	Check pipette calibration.			
و ا		Check pipette for proper performance.			
	Insufficient mixing of	 Thoroughly agitate the lyophilized components after 			
	reagent dilutions	reconstitution.			
		Thoroughly mix dilutions.			
		Check the microplate pouch for proper sealing.			
	Improperly sealed	Check that the microplate pouch has no punctures.			
	microplate	Check that three desiccants are inside the microplate			
	Microplate was left	pouch prior to sealing.			
l =	unattended between	 Each step of the procedure should be performed uninterrupted. 			
eu S	steps	uninterrupteu.			
Sig	Omission of step	Consult the provided procedure for complete list of steps.			
년 도	Steps performed in	Consult the provided procedure for the correct order.			
Ξ̈́	incorrect order				
₽Ş	Insufficient amount of	Check pipette calibration.			
≯ isr	reagents added to	 Check pipette for proper performance. 			
ly Low o	wells				
Unexpectedly Low or High Signal Intensity	Wash step was skipped	 Consult the provided procedure for all wash steps. 			
ţe	Improper wash buffer	 Check that the correct wash buffer is being used. 			
) e	Improper reagent	Consult reagent preparation section for the correct			
, d	preparation	dilutions of all reagents.			
<u>u</u>	Insufficient or	Consult the provided procedure for correct incubation			
ر ا	prolonged incubation	time.			
	periods	Sandwich ELISA: If samples generate OD values higher			
		than the highest standard point (P1), dilute samples			
≝		further and repeat the assay.			
ě	Non-optimal sample	Competitive ELISA: If samples generate OD values lower			
5	dilution	than the highest standard point (P1), dilute samples			
2		further and repeat the assay.			
arc		 User should determine the optimal dilution factor for 			
ğ		samples.			
Deficient Standard Curve Fit	Contamination of	 A new tip must be used for each addition of different 			
it S	reagents	samples or reagents during the assay procedure.			
e.	Contents of wells	Verify that the sealing film is firmly in place before placing			
fici	evaporate	the assay in the incubator or at room temperature.			
De.		Pipette properly in a controlled and careful manner.			
_	Improper pipetting	Check pipette calibration.			
		Check pipette for proper performance.			

Insufficient mixing of reagent dilutions	Thoroughly agitate the lyophilized components after reconstitution. Thoroughly mix dilutions.
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References

- (1) Harvey K et al. (2002) J Biol Chem. 277(11):9307-9317.
- (2) Jiang H et al. (2019) J Biol Chem. 294(46):17421-17436.
- (3) Sang Q et al. (2006) J Neurosci. 26(27):7234-7244.

Version 1.0

10 May 2021