

ELISA PRODUCT INFORMATION & MANUAL

Human NANS ELISA Kit (Colorimetric) NBP3-18741 Sample Insert for reference use only

Enzyme-linked Immunosorbent Assay for quantitative detection. For research use only.

Not for diagnostic or therapeutic procedures.

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 12 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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Human NANS ELISA Kit (Colorimetric)

Catalog No. NBP3-18741
Sample insert for reference use only

Introduction

Sialic acid synthase (SAS), also known as N-acetylneuraminate synthase or NANS, consists of 359 amino acids having a molecular mass of 40 kDa. The human enzyme uses N-acetylmannosamine 6-phosphate and mannose 6-phosphate as substrates to generate phosphorylated forms of N-acetylneuraminic acid (Neu5Ac) and 2-keto-3-deoxy-D-glycero-D-galactonononic acid (KDN), respectively, but exhibits much higher activity toward the Neu5Ac phosphate product. Sialic acids are frequently the terminal sugars on secreted and cell surface glycoproteins and glycolipids, and their presence can have considerable influence on the biological properties of a cell (1). Changes in amount, type, or linkage of sialic acid in tumour cell glycoconjugates can affect tumour growth and metastasis (2). SAS/NANS-mediated synthesis of sialic acid is required for early brain development and skeletal growth (3).

Principle of the Assay

The Human NANS ELISA Kit (Colorimetric) is designed for detection of NANS in human **cell lysate and tissue samples**. This assay employs a quantitative **sandwich enzyme immunoassay** technique that measures human NANS in approximately 4 hours. A polyclonal antibody specific for human NANS has been pre-coated onto a 96-well microplate with removable strips. NANS in standards and samples is sandwiched by the immobilized antibody and a biotinylated polyclonal antibody specific for human NANS, 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 NANS Microplate: A 96-well polystyrene microplate (12 strips of 8 wells) coated with a polyclonal antibody against human NANS.
- **Sealing Tapes:** Each kit contains 3 precut, pressure sensitive sealing tapes that can be cut to fit the format of the individual assay.
- Human NANS Standard: Human NANS in a buffered protein base (64 ng, lyophilized).
- **Biotinylated Human NANS Antibody (50x):** A 50-fold concentrated biotinylated polyclonal antibody against human NANS (120 µl).
- **EIA Diluent Concentrate (10x):** A 10-fold concentrated buffered protein base (20 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

- 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 (10 mM Tris pH 8.0, 130 mM NaCl, 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. 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. 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 Assuming the needed volume is less than or equal to 400 μl. | A) B) | 4 μl sample : 396 μl buffer (100x) 4 μl of A : 396 μl buffer (100x) = 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 Assuming the needed volume is less than | 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. | | or equal to 240 µl. | | |

Reagent Preparation

• Freshly dilute all reagents and bring all reagents to room temperature before use.

- **EIA Diluent Concentrate (10x):** Dilute the EIA Diluent Concentrate 10-fold 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 NANS Standard:** Reconstitute the Human NANS Standard (64 ng) with 0.5 ml of **Standard Diluent** to generate a 128 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 (128 ng/ml) 2-fold with equal volume of **EIA Diluent** to produce 64, 32, 16, 8, 4, 2, and 1 ng/ml solutions. EIA 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 | [NANS] (ng/ml) |
|-------------------|--|-------------------|
| P1 | 1 part Standard (128 ng/ml) + 1 part EIA Diluent | 64 |
| P2 | 1 part P1 + 1 part EIA Diluent | 32 |
| Р3 | 1 part P2 + 1 part EIA Diluent | 16 |
| P4 | 1 part P3 + 1 part EIA Diluent | 8.0 |
| P5 | 1 part P4 + 1 part EIA Diluent | 4.0 |
| P6 | 1 part P5 + 1 part EIA Diluent | 2.0 |
| P7 | 1 part P6 + 1 part EIA Diluent | 1.0 |
| P8 | EIA Diluent | 0.0 |

- Biotinylated Human NANS Antibody (50x): Spin down the antibody briefly and dilute the desired amount of the antibody 50-fold with EIA 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 EIA 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 μ l of Human NANS 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 μ l of Biotinylated Human NANS 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 12 minutes or until the optimal blue color density develops.
- Add 50 μ l 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

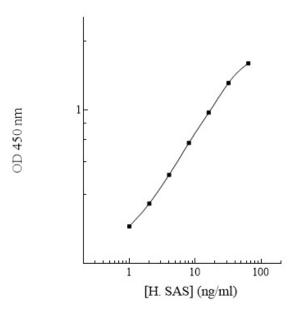
• The typical data is provided for reference only. Individual laboratory means may vary from the values listed. Variations between laboratories may be caused by technique differences.

| Standard Point | ng/ml | OD | Average OD |
|----------------|-------|-------|------------|
| P1 | 64 | 2.036 | 2.008 |
| PI | 04 | 1.980 | 2.006 |
| P2 | 32 | 1.520 | 1.495 |
| PZ | 32 | 1.470 | 1.493 |
| P3 | 16 | 0.976 | 0.957 |
| ro | 10 | 0.938 | 0.937 |
| P4 | 8.0 | 0.620 | 0.608 |
| F4 | | 0.596 | 0.008 |
| P5 | 4.0 | 0.385 | 0.377 |
| LO | 4.0 | 0.369 | 0.377 |
| P6 | 2.0 | 0.251 | 0.245 |
| r u | 2.0 | 0.239 | 0.243 |
| P7 | 1.0 | 0.177 | 0.174 |
| Γ/ | 1.0 | 0.171 | 0.174 |
| P8 | 0.0 | 0.086 | 0.084 |
| го | 0.0 | 0.082 | 0.064 |

Standard Curve

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

Human SAS Standard Curve



Reference Value

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

| Cell Culture Lysate | Dilution Factor | Average Value (ng/mg cell lysate) |
|-------------------------------------|--------------------|-----------------------------------|
| 293T (human embryonic kidney) | 10x | 87.273 |
| A549 (human adenocarcinoma) | 10x | 50.897 |
| Jurkat E6-1 (human T-cell leukemia) | 10x | 23.519 |

Performance Characteristics

- This assay recognizes both natural and recombinant human NANS.
- The minimum detectable dose of human NANS as calculated by 2SD from the mean of a zero standard was established to be 0.49 ng/ml.
- Intra-assay precision was determined by testing three lysate samples twenty times in one assay.
- Inter-assay precision was determined by testing three lysate samples in twenty assays.

| | Intra-Assay Precision | | | Inter-Assay Precision | | |
|-------------------|-----------------------|------|------|-----------------------|-------|------|
| Sample | 1 | 2 | 3 | 1 | 2 | 3 |
| n | 20 | 20 | 20 | 20 | 20 | 20 |
| CV (%) | 3.8% | 5.2% | 4.0% | 9.2% | 10.1% | 9.9% |
| Average CV (%) | 4.3% | | | | 9.7% | |

Recovery

| Standard Added Value | 4 – 32 ng/ml |
|----------------------|--------------|
| Recovery % | 87 – 116% |
| Average Recovery % | 105% |

Linearity

• Lysate samples were serially diluted to test for linearity.

| Average Percentage of Expected Value (%) | | |
|--|-------------------------------|--|
| Canada Dilatian | 293T (human embryonic kidney) | |
| Sample Dilution | Cell Culture Lysate | |
| 5x | 102% | |
| 10x | 97% | |
| 20x | 101% | |

Troubleshooting

| Issue | Causes | Course of Action |
|---------------|---|---|
| | Use of improper components | Check the expiration date listed before use. Do not interchange components from different lots. |
| _ | Improper wash step | Check that the correct wash buffer is being used. Check that all wells are empty after aspiration. Check that the microplate washer is dispensing properly. If washing by pipette, check for proper pipetting technique. |
| cisio | Splashing of reagents while loading wells | Pipette properly in a controlled and careful manner. |
| Low Precision | Inconsistent volumes loaded into wells | Pipette properly in a controlled and careful manner. Check pipette calibration. Check pipette for proper performance. |
| 1 | Insufficient mixing of reagent dilutions | Thoroughly agitate the lyophilized components after reconstitution. Thoroughly mix dilutions. |
| | Improperly sealed microplate | Check the microplate pouch for proper sealing. Check that the microplate pouch has no punctures. Check that three desiccants are inside the microplate pouch prior to sealing. |

| l _ | Microplate was left | Each step of the procedure should be performed |
|--|-----------------------------|---|
| пa | unattended between steps | uninterrupted. |
| Sig | Omission of step | Consult the provided procedure for complete list of steps. |
| ج | Steps performed in | Consult the provided procedure for complete list of steps. Consult the provided procedure for the correct order. |
| Ξ̈́ | incorrect order | • consult the provided procedure for the correct order. |
| ₽ | Insufficient amount of | Check pipette calibration. |
| Unexpectedly Low or High Signal Intensity | reagents added to wells | Check pipette for proper performance. |
| <u>≥</u> ⊆ | Wash step was skipped | Consult the provided procedure for all wash steps. |
| ied Eed | Improper wash buffer | Check that the correct wash buffer is being used. |
| ect | Improper reagent | Consult reagent preparation section for the correct |
| S × | preparation | dilutions of all reagents. |
| ne | Insufficient or | Consult the provided procedure for correct incubation |
| > | prolonged incubation | time. |
| | periods | |
| | | Sandwich ELISA: If samples generate OD values higher the phich act the plant as int (P1) dilute as years. |
| | | than the highest standard point (P1), dilute samples further and repeat the assay. |
| يب | Non-optimal sample | Competitive ELISA: If samples generate OD values lower |
| 正 | dilution | than the highest standard point (P1), dilute samples |
| Ž | | further and repeat the assay. |
| Deficient Standard Curve Fit | | User should determine the optimal dilution factor for |
| <u>5</u> | | samples. |
| qa | Contamination of | A new tip must be used for each addition of different |
| an | reagents | samples or reagents during the assay procedure. |
| St | Contents of wells | Verify that the sealing film is firmly in place before placing |
| l t | evaporate | the assay in the incubator or at room temperature. |
| <u>i</u> | | Pipette properly in a controlled and careful manner. Charles is a trackly at the set of the s |
|)ef | Improper pipetting | Check pipette calibration.Check pipette for proper performance. |
| | | |
| | Insufficient mixing of | Thoroughly agitate the lyophilized components after reconstitution. |
| | reagent dilutions | Thoroughly mix dilutions. |
| | | 3 , |

References

- (1) Lawrence SM et al. (2000) J Biol Chem. 275(23):17869-17877.
- (2) Takano R et al. (1994) Glycobiology. 4(5):665-674.
- (3) van Karnebeek CD et al. (2016) Nat Genet. 48(7):777-784.

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