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Datasheet for ABIN456477

TRAIL ELISA Kit

Overview

Quantity:	96 tests
Target:	TRAIL (TNFSF10)
Binding Specificity:	Soluble
Reactivity:	Human
Method Type:	Sandwich ELISA
Detection Range:	46.9-3000 pg/mL
Minimum Detection Limit:	46.9 pg/mL
Application:	ELISA

Product Details

Purpose:	This immunoassay kit allows for the specific measurement of human soluble tumor necrosis factor-related apoptosis inducing ligand(sTRAIL) concentrations in cell culture supernates, serum, and plasma.
Sample Type:	Cell Culture Supernatant, Serum, Plasma
Analytical Method:	Quantitative
Detection Method:	Colorimetric
Specificity:	This assay recognizes recombinant and natural human sTRAIL.
Cross-Reactivity (Details):	No significant cross-reactivity or interference was observed.
Characteristics:	Homo sapiens, Human, Tumor necrosis factor ligand superfamily member 10, Apo-2 ligand, Apo-2L, TNF-related apoptosis-inducing ligand, Protein TRAIL, TNFSF10, APO2L, TRAIL, CD253

Product Details

Components:	Reagent (Quantity): Assay plate (1), Standard (2), Sample Diluent (1x20ml), Assay Diluent A (1x10ml), Assay Diluent B (1x10ml), Detection Reagent A (1x120µl), Detection Reagent B (1x120µl), Wash Buffer(25 x concentrate) (1x30ml), Substrate (1x10ml), Stop Solution (1x10ml)
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Target Details

Target:	TRAIL (TNFSF10)
Alternative Name:	TNFSF10 (TNFSF10 Products)
Background:	<p>TRAIL (tumor necrosis factor-related apoptosis-inducing ligand) is a recently identified member of the TNF gene superfamily. Five different receptors have been identified for TRAIL. Two receptors, DR4 and DR5, are transmembrane proteins containing death domain similar to FAS and other TNF family receptors. Two other receptors, DcR1 and DcR2, act like decoy proteins for TRAIL binding because they lack the death domain. TRAIL can also bind, though weakly, to osteoprotegerin (OPG), a soluble receptor, which plays a role in osteoclastogenesis. TRAIL induces apoptosis in various tumor cell lines, whereas most primary cells seem to be resistant. TRAIL-mediated apoptosis occurs following its binding to DR4 or DR5 receptors. The mechanism of apoptosis involves activation of caspase-8 and subsequent activation of effector caspases. Also, NF-kappaB and JNK activation play a role in the TRAIL signaling pathway. TRAIL expression is detectable in many normal organs and tissues. Several studies suggest that TRAIL may play a physiological role by contributing to immune privilege, normal cellular development, and inhibition of autoimmune responses. The expression of TRAIL is upregulated in activated T cells, B cells, NK cells, monocytes and macrophages. LPS activation stimulates the release of soluble TRAIL from monocytes and macrophages, suggesting a role of TRAIL in their cytotoxic/phagocytic function. Also, TRAIL expression is increased in transformed cell lines and various diseases including cancer and autoimmune disorders. TRAIL may also be involved in activation-induced T cell death during HIV infection. Certain anti-cancer agents also upregulate TRAIL and TRAIL receptor expression on tumor cells, thus sensitizing cells to apoptosis.</p>
Pathways:	Apoptosis , Positive Regulation of Endopeptidase Activity

Application Details

Sample Volume:	100 µL
Plate:	Pre-coated
Protocol:	This assay employs the quantitative sandwich enzyme immunoassay technique. A monoclonal

antibody specific for sTRAIL has been pre-coated onto a microplate. Standards and samples are pipetted into the wells and any sTRAIL present is bound by the immobilized antibody. An enzyme-linked monoclonal antibody specific for sTRAIL is added to the wells. Following a wash to remove any unbound antibody-enzyme reagent, a substrate solution is added to the wells and color develops in proportion to the amount of sTRAIL bound in the initial step. The color development is stopped and the intensity of the color is measured.

Reagent Preparation:	Bring all reagents to room temperature before use. Wash Buffer - If crystals have formed in the concentrate, warm to room temperature and mix gently until the crystals have completely dissolved. Dilute 20 mL of Wash Buffer Concentrate into deionized or distilled water to prepare 500 mL of Wash Buffer. 3 Standard - Reconstitute the Standard with 1.0 mL of Sample Diluent. This reconstitution produces a stock solution of 3000 pg/ml. Allow the standard to sit for a minimum of 15 minutes with gentle agitation prior to making serial dilutions. The undiluted standard serves as the high standard (3000 pg/ml). The Sample Diluent serves as the zero standard (0 pg/ml). Detection Reagent A and B - Dilute to the working concentration specified on the vial label using Assay Diluent A and B (1:100), respectively.
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Sample Collection:	Cell culture supernates - Remove particulates by centrifugation and assay immediately or aliquot and store samples at ≤ -20 °C. Avoid repeated freeze-thaw cycles. Serum - Use a serum separator tube (SST) and allow samples to clot for 30 minutes before centrifugation for 15 minutes at approximately 1000 x g. Remove serum and assay immediately or aliquot and store samples at -20 °C. Plasma - Collect plasma using EDTA or heparin as an anticoagulant. Centrifuge samples for 15 minutes at 1000 x g at 2 - 8 °C within 30 minutes of collection. Store samples at ≤ -20 °C. Avoid repeated freeze-thaw cycles. Note: Citrate plasma has not been validated for use in this assay.
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Assay Procedure:	<p>Allow all reagents to reach room temperature. Arrange and label required number of strips.</p> <ol style="list-style-type: none">1. Prepare all reagents, working standards and samples as directed in the previous sections.2. Add 100 μL of Standard, Control, or sample per well. Cover with the adhesive strip. Incubate for 2 hours at 37 °C.3. Remove the liquid of each well, don't wash.4. Add 100 μL of Detection Reagent A to each well. Incubate for 1 hour at 37°C. Detection Reagent A may appear cloudy. Warm to room temperature and mix gently until solution appears uniform.5. Aspirate each well and wash, repeating the process three times for a total of three washes. <p>Wash by filling each well with Wash Buffer (350 μL) using a squirt bottle, multi-channel pipette, manifold dispenser or autowasher. Complete removal of liquid at each step is essential to good performance. After the last wash, remove any remaining Wash Buffer by aspirating or</p>
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decanting. Invert the plate and blot it against clean paper towels.

6. Add 100 μ L of Detection Reagent B to each well. Cover with a new adhesive strip. Incubate for 1 hours at 37 °C.

7. Repeat the aspiration/wash as in step

5. 8. Add 90 μ L of Substrate Solution to each well. Incubate for 30 minutes at room temperature. Protect from light.

9. Add 50 μ L of Stop Solution to each well. If color change does not appear uniform, gently tap the plate to ensure thorough mixing.

10. Determine the optical density of each well within 30 minutes, using a microplate reader set to 450 nm.

Important Note:

1. The wash procedure is critical. Insufficient washing will result in poor precision and falsely elevated absorbance readings.

2. It is recommended that no more than 32 wells be used for each assay run if manual pipetting is used since pipetting of all standards, specimens and controls should be completed within 5 minutes. A full plate of 96 wells may be used if automated pipetting is available.

3. Duplication of all standards and specimens, although not required, is recommended.

4. When mixing or reconstituting protein solutions, always avoid foaming. 4

5. 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.

6. To ensure accurate results, proper adhesion of plate sealers during incubation steps is necessary.

Calculation of Results:

Average the duplicate readings for each standard, control, and sample and subtract the average zero standard optical density. Create a standard curve by reducing the data using computer software capable of generating a four parameter logistic (4-PL) curve-fit. As an alternative, construct a standard curve by plotting the mean absorbance for each standard on the y-axis against the concentration on the x-axis and draw a best fit curve through the points on the graph. The data may be linearized by plotting the log of the sTRAIL concentrations versus the log of the O.D. and the best fit line can be determined by regression analysis. This procedure will produce an adequate but less precise fit of the data. If samples have been diluted, the concentration read from the standard curve must be multiplied by the dilution factor.

Restrictions:

For Research Use only

Handling

Handling Advice:	<ol style="list-style-type: none">1. The kit should not be used beyond the expiration date on the kit label.2. Do not mix or substitute reagents with those from other lots or sources.3. If samples generate values higher than the highest standard, further dilute the samples with the Assay Diluent and repeat the assay. Any variation in standard diluent, operator, pipetting technique, washing technique, incubation time or temperature, and kit age can cause variation in binding.4. This assay is designed to eliminate interference by soluble receptors, ligands, binding proteins, and other factors present in biological samples. Until all factors have been tested in the Immunoassay, the possibility of interference cannot be excluded.
Storage:	4 °C/-20 °C
Storage Comment:	The Standard, Detection Reagent A, Detection Reagent B and the 96-well strip plate should be stored at -20 °C upon being received. The other reagents can be stored at 4 °C.