

Human TNF-RII



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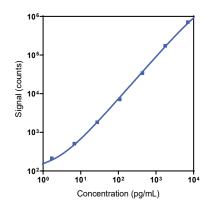
Meso Scale Discovery, a divistion of Meso Scale Diagnostics

, LLC. 1601 Research Boulevard Rockville, MD 20850-3173 USA

| | Product Options | Catalog Number | Description |
|--|-----------------|--|---|
| | Multiplex | K151AEM, K251AEM | U-PLEX Immuno-Oncology Group 1 (human) |
| | Singleplex | K151AHRK-1/-2/-4 | U-PLEX Human TNF-RII Assay with SECTOR™ plates |
| | | K151AHRK-21/-22/-24 | U-PLEX Human TNF-RII Assay with QuickPlex Ultra™ plates |
| | | K251AHRK-2/-4 | U-PLEX Human TNF-RII Assay with 384-well plates |
| | Antibody Set | B21AHR-2/-3 | U-PLEX Human TNF-RII Antibody Set |
| | Protocol | U-PLEX Product Inserts are available at www.mesoscale.com. | |

The U-PLEX® platform was designed to provide ultimate flexibility for the detection of biomarkers in a wide variety of sample types. This datasheet provides the representative performance of the U-PLEX Human TNF-RII Assay tested on U-PLEX 96-well SECTOR plates run as a multiplex. The data do not represent the product specifications. Under your experimental conditions, the assay may perform differently from the representative data. U-PLEX assays are offered in either singleplex or multiplex; both are available on 96- or 384-well plates. See a U-PLEX product insert for instrument compatibility.

Representative Calibration Curve and Sensitivity



| Assay | Median LLOD (pg/mL) | LLOD Range (pg/mL) | |
|---------|------------------------|-----------------------|--|
| TNF-RII | 1.60 | 1.04-2.45 | |

The Calibrator curve was fitted with a 4-parameter logistic model with a $1/Y^2$ weighting. The lower limit of detection (LLOD) is a calculated concentration corresponding to 2.5 standard deviations above the background (zero Calibrator).

Precision

| Control | Average Conc. (pg/mL) | Average Intra-run Conc. (%CV) | Inter-run Conc. (%CV) | |
|---------|-----------------------|-------------------------------|-----------------------|--|
| High | 1,180 | 2.5 | 5.8 | |
| Mid | 265 | 2.2 | 7.4 | |
| Low | 64 | 2.0 | 7.4 | |

Controls were made by spiking Calibrator into assay diluent at 3 levels within the quantitative range of the assay. Average intra-run concentration %CV is the average %CV of the control replicates within an individual run. Inter-run concentration %CV is the variability of controls across multiple runs.

For Research Use Only. Not for use in diagnostic procedures.





MSD® U-PLEX Human TNF-RII

Tested Samples

| Sample Type | Serum (N = 9) | EDTA Plasma (N = 9) | Citrate Plasma (N = 9) | Normal Lysate (N = 5) | Tumor Lysate (N = 5) |
|----------------|------------------|------------------------|---------------------------|--------------------------|-------------------------|
| Median (pg/mL) | 6,100 | 5,690 | 4,950 | 341 | 489 |
| Range (pg/mL) | 3,910–9,590 | 3,740-8,910 | 3,160–7,610 | 212–553 | 74–1,490 |
| % Detected | 100 | 100 | 100 | 100 | 100 |

Normal serum and plasma samples were diluted 100-fold prior to testing in the assay. Lysates were tested at a protein concentration of 0.5 mg/mL.

Parallelism

| Serum | | | EDTA Plasma | | |
|---------------|--------------------|------------------|---------------|--------------------|------------------|
| Fold Dilution | Average % Recovery | % Recovery Range | Fold Dilution | Average % Recovery | % Recovery Range |
| 50 | 93 | 88–97 | 50 | 90 | 48–98 |
| 200 | 102 | 99–106 | 200 | 105 | 100–111 |
| 400 | 105 | 95–111 | 400 | 108 | 101–118 |

Samples were tested at different dilutions. One hundred-fold diluted samples were tested to determine the expected concentration of the analyte.

% Recovery = (measured concentration / expected concentration) x 100

Spike Recovery

| | Ser | um | EDTA Plasma | | |
|-------------|---|---------|-------------------|------------------|--|
| Spike Level | Spike Level Average % Recovery % Recovery Range | | Average% Recovery | % Recovery Range | |
| High | 103 | 95–107 | 106 | 97–124 | |
| Mid | 119 | 107–130 | 106 | 96–126 | |
| Low | 109 | 95–129 | 97 | 91–101 | |

Samples were diluted 100-fold prior to addition of spike. The expected concentration of the analyte in spiked samples was calculated by addition of the Calibrator spike concentration to the unspiked sample concentration.

% Recovery = (measured concentration / expected concentration) x 100

Specificity

To assess specificity, the TNF-RII Antibody Set was tested individually against a larger panel of analytes for nonspecific binding: APRIL/TNFSF13, BAFF-R/TNFRSF13C, BCMA/TNFRSF17, CD20, CD27, CD276/B7-H3, CD28, CD40L (soluble), CTACK, CTLA-4, ENA-78, Eotaxin, Eotaxin-2, Eotaxin-3, EPO, E-Selectin, FGF (basic), FLT3L, Fractalkine, G-CSF, Galectin-9, GITR/TNFRSF18, GITRL/TNFSF18, GM-CSF, gp130 (soluble), Granzyme A, Granzyme B, GR0- α , HAVCR2/TIM-3, HVEM/TNFRSF14, I-309, ICOS, ICOSL/B7-H2, IFN- α 2a, IFN- β , IFN- γ , IL-1 α , IL-1 β , IL-1RA, IL-10, IL-12/IL-23p40, IL-12p70, IL-13, IL-15, IL-16, IL-17A, IL-17A/F, IL-17C, IL-17D, IL-17E/IL 25, IL-17F, IL-18, IL-2, IL-21, IL-22, IL-23, IL-27, IL-29/IFN-A1, IL-2R α , IL-3, IL-31, IL-33, IL-4, IL-5, IL-6, IL-7, IL-8, IL-9, IP-10, I-TAC, LAG-3, LIGHT/TNFSF14, MCP-1, MCP-2, MCP-4, M-CSF, MDC, MIF, MIG, MIP-1 α , MIP-5, MMP-1, MMP-2, MMP-7, MMP-9, Nectin-4, OX40/TNFRSF4, PD1, PD-L1, PD-L2, Pentraxin 3, Perforin, PIGF, P-Selectin, RAGE (soluble), RANKL/TNFSF11, RANTES, S100A12, TARC, Tie-2, TIGIT, TLR-1, TNF-RI, TNF-RI, TNF- α , TNF- β , TP0, TRAIL, TSLP, VEGF-A, VEGF-D, VEGFR-1/FIt-1 and YKL-40. Nonspecific binding was less than 2.0%.

% Nonspecificity = (nonspecific signal / specific signal) x 100

Diluent Compatibility

Diluents 58 and 3 are provided with this assay. MSD offers a range of assay and antibody diluents for separate purchase. Depending on your assay needs, other diluents may be tested.

Assay Components

Calibrator: TNF-RII is included in Calibrator 29. The human TNF-RII Calibrator is a full-length recombinant protein expressed in E. coli.

Antibodies: The U-PLEX Human TNF-RII Assay uses a mouse monoclonal antibody for capture and a mouse monoclonal antibody for detection.

Assay generation: A

Note: This datasheet contains representative assay performance data. In custom multiplex formats, the assay may perform differently from the representative data shown.

Note: MSD recommends that samples be diluted 100-fold prior to analysis in this assay.

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