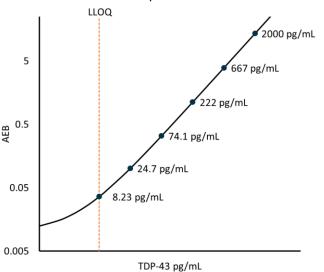
Simoa™ TDP-43 Kit SR-X™ Data Sheet

Item 103293

Description

The TAR DNA binding protein of 43 kDa (TDP-43 or TARDBP) is a highly conserved and ubiquitously expressed nuclear protein with roles in transcription and splicing regulation. It is also the major component of ubiquitinpositive cytoplasmic inclusions found in the brains of patients with frontotemporal lobar degeneration (FTLD) and amyotrophic lateral sclerosis (ALS). In addition, TDP-43-containing aggregates are found in a significant number of patients with Alzheimer's Disease (AD) and other neuromuscular disorders. The majority of TDP-43 protein found in cytoplasmic inclusions is truncated, and it has been shown that the C-terminal domain is intrinsically prone to aggregation. Mutations in the Cterminal region of the TDP-43 gene have been associated with both ALS and FTLD, and are thought to facilitate ubiquitination and phosphorylation of the TDP-43 protein, leading to the formation of pathological inclusions and eventual neurodegeneration. Analysis of TDP-43 levels in plasma may allow the indexing of TDP-43 pathology within the brain to aid in the diagnosis of different forms of dementia and distinguish between TDP-43 proteinopathy and tauopathy. The Simoa TDP-43 assay has been developed with a full-length protein calibrator and antibodies against AA 203 - 209 and the C-terminal region; it is expected to detect both full-length and pathological, truncated forms of the protein.

Calibration Curve: Calibrator concentrations and Lower Limit of Quantification depicted.



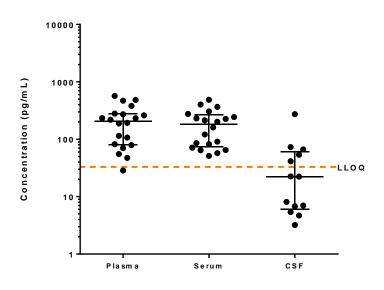
Lower Limit of Quantification (LLOQ): Triplicate measurements of serially diluted calibrator were read back on the calibration curve over 6 runs each for 1 reagent lot across 2 instruments (6 runs total).

Limit of Detection (LOD): Calculated as 2.5 standard deviations from the mean of background signal read back on each calibration curve over 6 runs each for 1 reagent lot across 2 instruments (6 runs total).

Analytical LLOQ	8.23 pg/mL pooled CV 11% mean recovery 112%
LOD	0.780 pg/mL range 0.019-1.59 pg/mL
Dynamic range (serum and plasma)	0 - 8000 pg/mL
Diluted Sample volume*	100 μL per measurement
Tests per kit	96

^{*}See Kit Instruction for details

Endogenous Sample Reading: Healthy donor matched EDTA plasma (n=20), and serum (n=20) were measured. 13 CSF samples were measured. Bars depict median with interquartile range. Orange line represents functional LLOQ.



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Sample Type	Mean TDP-43 pg/mL	Median TDP-43 pg/mL	% Above LOD
Serum	191	182	100%
Plasma	228	206	100%
CSF	102	22.3	100%

Precision: Measurements of 1 plasma-based panel, 3 serum-based panels and 2 calibrator-based controls. Triplicate measurements were made for 6 runs each for 2 reagent lots across 2 instruments (12 runs total, 36 measurements).

Sample	Mean (pg/mL)	Within run CV	Btwn run CV	Btwn inst CV	Btwn lot CV
Control 1	176	5.8%	5.5%	4.0%	1.1%
Control 2	1837	4.7%	4.9%	1.0%	4.4%
Panel 1	69.6	13.7%	9.9%	3.1%	7.5%
Panel 2	82.9	12.2%	14.2%	0.0%	9.4%
Panel 3	363	8.1%	10.9%	7.0%	0.1%
Panel 4	1941	4.9%	14.4%	10.2%	1.4%

Spike and Recovery: 2 serum, 2 EDTA plasma and 2 CSF samples were spiked at high and low concentrations within the range of the assay and analyzed on SR-X.

Observed recovery was consistently low in serum and plasma, but results from dilutional linearity and immune-depletion experiments support specificity of the assay signal.

Dilution Linearity: 2 endogenous EDTA plasma, 2 endogenous serum and 2 spiked CSF samples were diluted 2X serially with Sample Diluent from MRD to 32x in serum/plasma and from MRD to 64x in CSF.

Spike and Recovery (Serum/Plasma)	Mean = 52% Range: 38-60%
Dilution Linearity	Mean = 111%
(Serum/Plasma) (32x)	Range: 91–123%
Spike and Recovery	Mean = 86%
(CSF)	Range: 74-96%
Dilution Linearity	Mean = 111%
(CSF) (64x)	Range: 100-121%

Immuno-depletion: 1 Serum and 2 plasma samples were separately incubated with TDP-43 beads and Antibody isotype control beads prior to analysis on SR-X. Mean depletion was 100%.

The Simoa TDP-43 assay kit is formulated for use on either the SR-X or HD-1 platform. Data in this document was obtained from runs on the SR-X platform unless otherwise noted. Some differences in performance claims between the HD-1 and SR-X may be observed when comparing datasheets for the two platforms. This may be due to experiments run at different time-points with different reagent lots and different samples, or may be due to minor differences in antibody and analyte behavior in the different assay formats.