

Comparison of the Simoa and MSD R-PLEX assay to assess serum neurofilament light chain levels in hereditary transthyretin amyloidosis

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INTRODUCTION

Neurofilament light chain (NfL), a biomarker of neuronal damage¹, can be detected in serum and has shown to be a sensitive biomarker for polyneuropathy in patients with hereditary transthyretin (ATTRv) amyloidosis.² A rise in serum NfL (sNfL) precedes the onset of polyneuropathy symptoms.³ Blood biomarkers, such as NfL, have the potential to improve diagnosis. The Single molecule array (Simoa) assay is currently regarded as the gold standard for NfL measurement.^{4,5} It is a highly sensitive and specific immunoassay that allows measurement of sNfL in very low concentrations.⁵ However, the Simoa assay is not widely available, which hampers implementation of NfL measurement into a clinical setting.

OBJECTIVE

We aimed to identify a high performance and well accessible assay for sNfL measurement in ATTRv amyloidosis patients. Here, we present a direct comparison of the Simoa assay with the Meso Scale Discovery (MSD) R-PLEX assay for measuring sNfL.

METHODS

sNfL levels were measured in samples collected between January 2000 and December 2021 from pathogenic transthyretin gene variant (TTRv) carriers and ATTRv amyloidosis patients. In each sample, sNfL levels were measured using both the Simoa assay and the MSD R-PLEX assay allowing direct comparison of the assays (Figure 1A). An additional data set with samples of healthy controls was added. Their sNfL levels were measured with the MSD R-PLEX only (Figure 1B). The internal standard for both assays was measured with the MSD R-PLEX assay allowing direct comparison of the quantitative difference between the two assays (Figure 1C).

RESULTS

A total of 332 samples were evaluated in 72 subjects. sNfL levels measured with the MSD R-PLEX assay (median 128.7pg/mL, interquartile range [IQR] 58.8–279.3) were consistently 4.9 times higher than those measured by the Simoa assay (26.4pg/mL [11.3–50.2]), $p < .0001$. All median sNfL levels per study group are displayed in Figure 1D-E. A strong correlation was found between levels measured with both assays (Pearson correlation coefficient 0.94, $p < .0001$) (Figure 1H). sNfL levels in healthy controls were 33.2 pg/mL [20.7-51.8] (Figure 1F) and correlated with age ($p < .0001$) (Figure 1I). Bland-Altman analysis showed, within the 95% limit of agreement, a mean constant bias of 130.4% for sNfL concentrations determined by the MSD R-PLEX assay compared to the Simoa assay. The quantitative difference as determined with the two assays related to a difference in the concentration of the internal standards provided with the tests (Figure 1G and 1J).

REFERENCES

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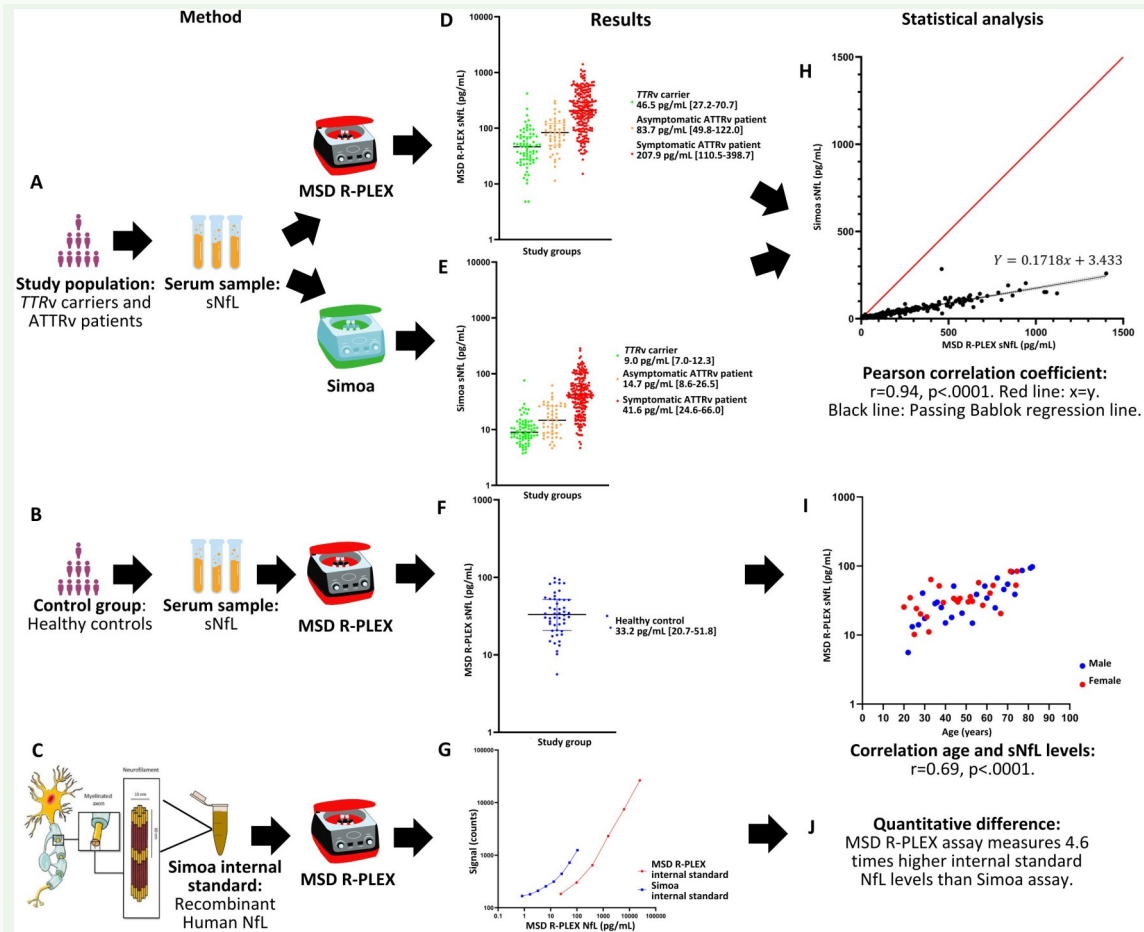


Figure 1. Study design and results.

Abbreviations: ATTRv: hereditary transthyretin amyloidosis; MSD: MesoScale Discovery; NfL: neurofilament light chain; Simoa: single molecule array; sNfL: serum neurofilament light chain; TTRv: transthyretin gene variant.

CONCLUSIONS

- The MSD R-PLEX assay is a robust and sensitive alternative to the Simoa assay for measuring sNfL levels in ATTRv amyloidosis.
- The quantitative difference between the assays is related to the internal standard provided with the tests.

