

P249

A novel signature of lipoxin A₄ and prostaglandin E₂ in plasma associated with disease severity in patients with relapsing-remitting and secondary progressive multiple sclerosis

J.Y. Broos^{1,2}, H. Kropshofer³, G. Karlsson³, J. Maca⁴, H.d. Vries¹, M. Giera², G. Kooij¹

¹Amsterdam UMC, location VUmc, MS center Amsterdam, Amsterdam Neuroscience, Amsterdam, Netherlands, ²Leiden University Medical Center, Center for Proteomics & Metabolomics, Leiden, Netherlands, ³Novartis Pharma AG, Basel, Switzerland, ⁴Novartis Pharmaceuticals Corporation, East Hanover, NJ, United States

Introduction: In multiple sclerosis (MS), chronic neuroinflammation may be due to deregulation of resolution of inflammation. This protective process is normally orchestrated by specialised pro-resolving lipid mediators derived from omega-3/-6 fatty acids. Failed resolution might be caused by a lack of pro-resolving lipid mediators, such as lipoxin A₄ (LXA₄), while pro-inflammatory lipid mediators, such as prostaglandin E₂ (PGE₂), may persist.

Objective: To explore whether LXA₄ and PGE₂ can be used as candidate biomarkers of disease activity in patients with relapsing-remitting MS (RRMS) and secondary progressive MS (SPMS).

Methods: In this post hoc analysis, involving baseline (BL) samples from the Phase 4 study LONGTERMS (N=28 RRMS-SPMS converters) and the Phase 3 study EXPAND (N=55 randomly selected patients with SPMS), LXA₄ and PGE₂ levels were quantified by quantitative liquid chromatography with tandem mass spectrometry. Lipids were categorised as lo (not detectable) or hi (detectable) or as signatures LXA₄(lo)+PGE₂(hi) vs LXA₄(hi)+PGE₂(lo). Mean BL characteristics – such as disease duration (DD), Expanded Disability Status Scale (EDSS) score, normalised brain volume (NBV), T2 lesion volume (LV) & 9-Hole Peg Test (9-HPT) were analysed by lipid category.

Results: In RRMS, 14/28 patients were PGE₂-positive at BL, while LXA₄ was only detectable in 4/28 patients. In SPMS, samples were PGE₂(hi) and LXA₄(hi) in 13/55 and 6/55 of the patients, respectively. SPMS patients with PGE₂(hi) vs PGE₂(lo) showed a trend towards a longer DD (20.6 vs 17.6 years), higher EDSS (5.7 vs 5.5), lower NBV (1384 vs 1425 cm³), higher LV (20.5 vs 17.1 cm³), and weaker performance in the 9-HPT (42.5 vs 32.4 seconds). In contrast, SPMS patients with LXA₄(hi) vs LXA₄(lo) trended towards higher NBV (1439 vs 1414 cm³), lower LV (16.4 vs 18.1 cm³), and a better performance in the 9-HPT (31.6 vs 35.2 seconds). SPMS patients with the signature LXA₄(lo)+PGE₂(hi) shared a trend of advanced disease severity compared to patients with LXA₄(hi)+PGE₂(lo), based on EDSS (5.8 vs 5.6), NBV (1383 vs 1438 cm³), LV (21.0 vs 16.7 cm³), and 9-HPT (43.3 vs 31.3 seconds). In RRMS patients, the signature LXA₄(lo)+PGE₂(hi) showed a similar trend towards association with higher EDSS, advanced brain atrophy, and low performance in the 9-HPT.

Conclusion: Circulating levels of LXA₄ and PGE₂ appear to be a candidate lipid biosignature of disease severity in RRMS and SPMS. Validation in larger studies is underway.

Disclosure: This study was funded by Novartis Pharma AG, Basel, Switzerland.

Gijs Kooij is partly supported by a grant from Novartis.

Helga de Vries, Jelle Y. Broos and **Martin Giera** have nothing to disclose.

Harald Kropshofer, Jeff Maca and **Goeril Karlsson** are employees of Novartis.