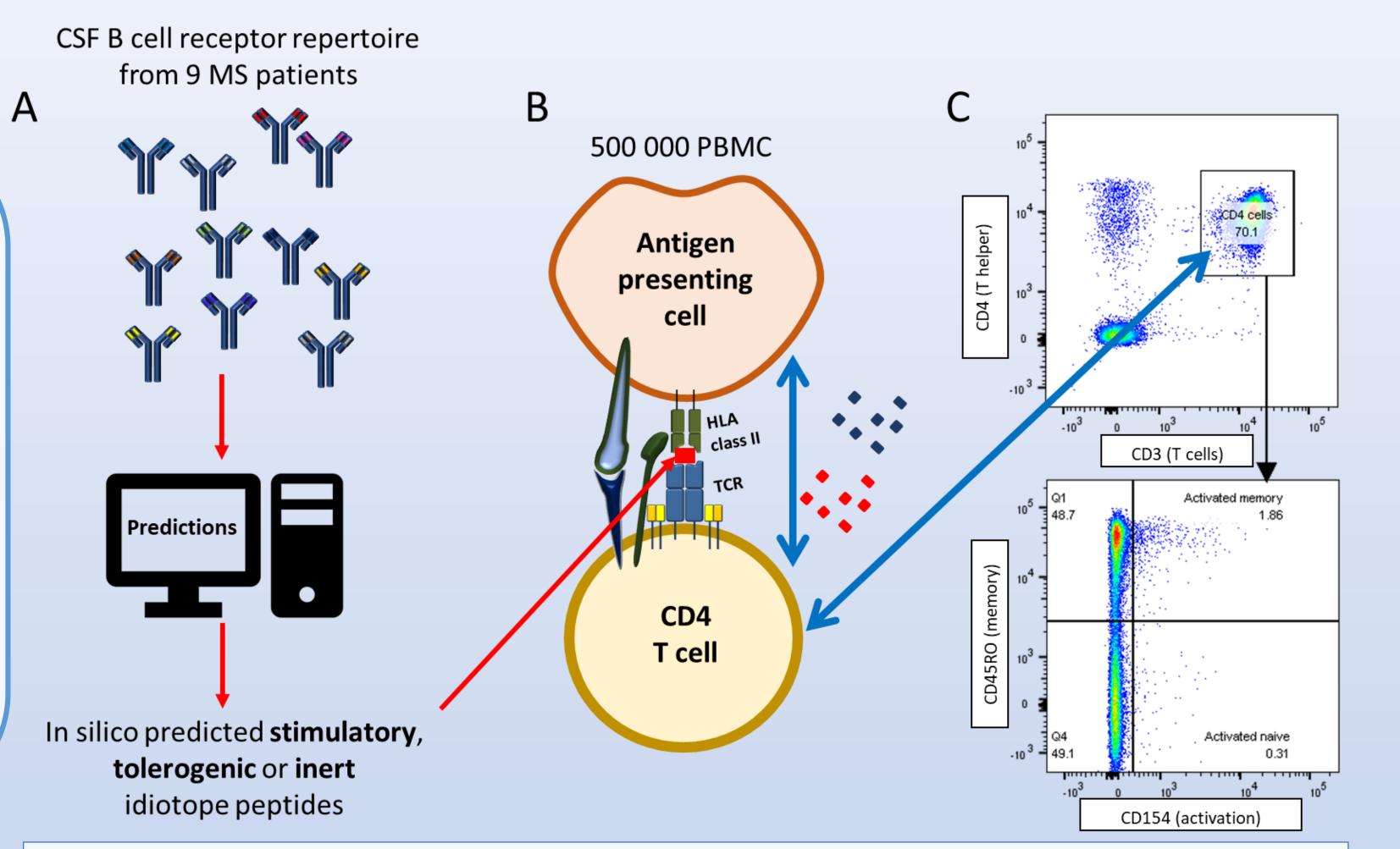
CD4⁺ T cells in the blood of MS patients respond to predicted epitopes from B cell receptors found in spinal fluid Rune A. Høglund^{1,2}, A. Lossius^{1,2,3}, E. J. Homan⁴, R.Bremel⁴ and T. Holmøy^{1,2}

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Introduction

The pathogenic role of B cells in multiple sclerosis (MS) is not known. We have previously demonstrated that B cells from cerebrospinal fluid (CSF) of MS patients can activate T cells through specific recognition of antigenic determinants (idiotopes) from their own B cells receptors (BCRs). The aim of this study was to evaluate these



findings in MS patients using *in silico* prediction models to accurately and quickly identify immunogenic idiotopes of immunoglobulin heavy-chain variable (IGHV) gene repertoires.

Materials and methods

Translated CSF IGHV repertoires were run through bioinformatic predictions *in silico* to predict HLA-DR affinity and endosomal processing, and calculate T cell exposed motif frequency. Predictions and transcript frequencies guided selection of idiotope-peptides from nine MS patients. PBMC from these patients were stimulated *in vitro* with the idiotope peptides in presence of anti-CD40 for 12 hours. Flow cytometry was utilized to identify activated cells using relevant surface makers (Table 1 and Figure 1). Unstimulated cells or insulin peptides were negative controls, and CD3/CD28 beads or EBNA-1 peptides were positive controls **Figure 1.** A) IGHV amino acid sequences (1079 [SD=1213]) from MS patients were run through predictive models to identify likely antigenic or inert peptides. B) PBMC were stimulated with idiotope peptides. C) Activated CD4⁺CD45RO⁺ memory T cells were be detected by surface expression of CD154.



In all nine MS patients, we found blood memory CD4⁺T cells were activated by predicted idiotope-peptides (Figure 2). Responses were mainly towards peptides affiliated with the CDR3 region, and no robust responses were seen towards peptides with low predicted HLA-DR affinity. Activated memory CD4⁺T cells also expressed the chemokine receptor CCR6, affiliated with a Th17 phenotype and allowing passage into the central nervous system (Figure 3).

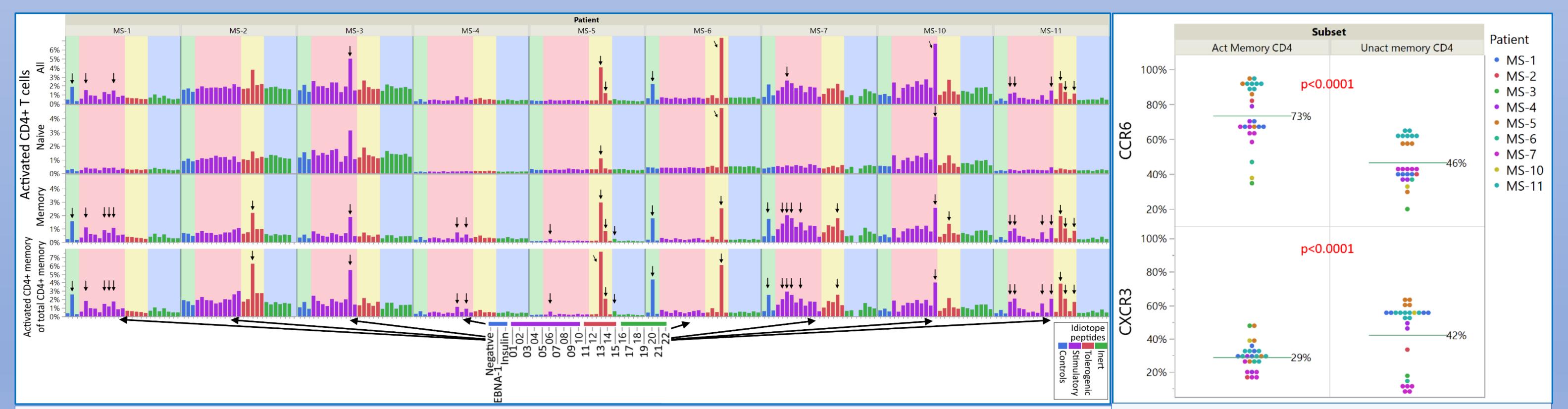


Figure 2. CD3⁺CD4⁺CD8⁻ T cells were assayed for the activation marker CD154. Activated cells among all CD4⁺ cells, CD45RO⁺ memory- or CD45RO⁻ naïve cells are presented as proportions of CD4⁺ cells (upper three panels) or proportion of memory cells (lower panel). Responses were deemed positive (arrows) if the proportion of CD154⁺ cells were 3x higher than in unstimulated (negative) wells.

Figure 3. CCR6 and CXCR3 expression among activated (CD154⁺) or un-activated (CD154⁻) memory (CD45RO⁺) CD4⁺ T cells responding to predicted idiotope peptides. P-values are results of full factorial mixed model.

Table 1: Surface markers andfluorochromes used in experiment

| Laser | Fluorochromes | Marker |
|--------------------|-----------------|-----------------------|
| 407 nm (Violet) | BV421 | CD154 (activation) |
| 488 nm (Blue) | PE | CXCR3 |
| | PE-Cy7 | CD45RO (memory) |
| | PerCP-Cy5.5 | CD4 |
| | FITC | CD3 |
| 633 nm (Red) | APC | CCR6 |
| | APC-H7 | CD14 |
| | | CD8 |
| | LIVE/DEAD™ | Viability |
| | Fixable Near-IR | (dump channel) |

Conclusions

This *in vitro* study suggests that MS patients have a memory T cell repertoire capable of recognizing frequent BCRs found in endogenous CSF, and that these T cells express chemokine receptors allowing them to reach the B cells presenting these peptides. It further indicates that antigenic properties of BCR idiotopes can be predicted *in silico* using HLA affinity and endosomal processing predictions.







Disclosures: RDB and EJH holds equity in IoGenetics, the company responsible for designing the bioinformatics models used in this project. All other authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest. The study was supported by grants from the Norwegian Research Council (grant 250864/F20), the Fritz and Ingrid Nilsen's endowment for MS research and an unrestricted research grant from Biogen Norway.