Supplemental Information

HLA-DR15 Molecules Jointly Shape an Autoreactive

T Cell Repertoire in Multiple Sclerosis

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Table S1. Information of PBMC samples from healthy donors and RRMS patients for immunopeptidome analysis and proliferation testing, Related to Figures 1, 3, 4, 5, S1, S4, and S5

| ID | Gender | Age# | Treatment | HLA-DRB1* | HLA-DRB3* | HLA-DRB4* | HLA-DRB5* |
|-------------------------------|--------|------|-------------|----------------------|-----------|-----------|---------------|
| НD-1 ^{†, Ψ} | F | 49 | - | 15:01 / 15:01 | -/- | -/- | 01:01 / 01:01 |
| HD-2 ^{†, Ψ} | M | 41 | - | 13:02 / 15:01 | 03:01 / - | -/- | 01:01 / - |
| НД-3 ^{†, Ψ} | M | 22 | - | 03:01 / 15:01 | 01:01 / - | -/- | 01:01 / - |
| HD-4 ^Ψ | N.A. | N.A. | - | +£ | | | + £ |
| HD-5 ^Ψ | N.A. | N.A. | - | +£ | | | +£ |
| HD-6 ^Ψ | N.A. | N.A. | - | +£ | | | +£ |
| HD-7 ^Ψ | N.A. | N.A. | - | +£ | | | +£ |
| HD-8 [*] | N.A. | N.A. | - | +£ | | | +£ |
| RRMS-1 ^{†, Ψ} | F | 25 | nihil | 04:01 / 15:01 | -/- | 01:01 / - | 01:01 / - |
| RRMS-2 ^{†, Ψ} | F | 49 | nihil | 04:03 / 15:01 | -/- | 01:01 / - | 01:01 / - |
| RRMS-3 ^{†, Ψ} | M | 27 | nihil | 03:01 / 15:01 | 01:01 / - | -/- | 01:01 / - |
| RRMS-4 ^Ψ | F | 26 | nihil | +/- | | | +/- |
| RRMS_NAT-1 ^{†, Ψ, §} | M | 54 | Natalizumab | 15:01 / 15:01 | -/- | -/- | 01:01 / 01:01 |
| RRMS_NAT-2 ^{†, Ψ, §} | M | 43 | Natalizumab | 11:01 / 15:01 | 02:01 / - | -/- | 01:01 / - |
| RRMS_NAT-3 ^{†, Ψ, §} | M | 25 | Natalizumab | 01:01 / 15:01 | -/- | -/- | 01:01 / - |
| RRMS_NAT-4* | M | 29 | Natalizumab | +/- | | | +/- |
| RRMS_NAT-5* | M | 38 | Natalizumab | +/- | | | +/- |
| RRMS_NAT-6* | F | 35 | Natalizumab | +/- | | | +/- |
| RRMS_NAT-7* | M | 47 | Natalizumab | +/- | | | +/- |
| RRMS_NAT-8 ^w | F | 23 | Natalizumab | +/- | | | +/- |
| RRMS_NAT-9 ^w | M | 54 | Natalizumab | +/+ | | | +/+ |
| RRMS_NAT-10 [*] | M | 50 | Natalizumab | +/- | | | +/- |

^{*}The age (in years) when the sample was collected

[†]Samples used for immunopeptidome analyses, B cells and monocytes isolation, and RNAseq

[♥]Samples used for proliferation testing

[§]Samples used for T cell cloning

[£]Determined by flow cytometer using fluorochrome-conjugated anti-DR2a and DR2b-specific antibodies N.A., not available

Table S4. Information of thymic and MS brain tissues for immunopeptidome analysis, Related to Figure 2

| ID | Gender | Age# | Feature | HLA-DRB1*15:01 | HLA-DRB5*01:01 |
|--------------------------|--------|----------|----------------------------------------------------------------|----------------|----------------|
| Thymus_1 [†] | M | 4 days | Immunologically healthy children undergoing cardiac surgery | +/- | +/- |
| Thymus_2 ^{†, §} | M | 6 months | Immunologically healthy children undergoing cardiac surgery | +/- | +/- |
| Thymus_3 ^{†, §} | M | 2 months | Immunologically healthy children undergoing cardiac surgery | +/- | +/- |
| Thymus_4 ^{†, §} | M | 6 months | Immunologically healthy children undergoing cardiac surgery | +/- | +/- |
| MS brain_1 [†] | M | 48 years | White matter lesion with high inflammation | +/- | +/- |
| MS brain_2 [†] | F | 53 years | White matter lesion with high inflammation | +/- | +/- |
| MS brain_3 [†] | M | 46 years | White matter lesion with high inflammation | +/- | +/- |
| MS brain_4 [†] | F | 45 years | White matter lesion with high inflammation | +/- | +/- |

[#]The age when the sample was collected

[†]Samples used for immunopeptidome analyses

[§]Samples used for TECs isolation and RNAseq

Table S5. Information of selected HLA-DR-SPs for analysis of CD4⁺ T cell reactivity, Related to Figure 3

The five most common HLA-DR-SPs presented by DR2a and DR2b on B cells were selected and synthesized for further functional testing. The name of the peptides is composed by the source protein and the location of the peptide sequence within the source protein. The HLA-DR-SP pool included all five individual HLA-DR-SPs.

| Name | Sequence | Source protein |
|----------------------------------|----------------------------------------------------------------|---------------------------------------------------------------------------------------------|
| HLA-DRB1 ₍₅₇₋₇₀₎ | DRYFYNQEESVRFD | HLA-DRB1*15:01 |
| HLA-DRB1 ₍₇₂₋₈₆₎ | DVGEFRAVTELGRPD | HLA-DRB1*15:01 |
| HLA-DRB5 ₍₇₂₋₈₆₎ | DVGEYRAVTELGRPD | HLA-DRB5*01:01 |
| HLA-DRB1/5 ₍₁₈₄₋₁₉₉) | FQTLVMLETVPRSGEV | HLA-DRB1*15:01 / HLA-DRB5*01:01 |
| HLA-DRA ₍₇₀₋₈₅) | LEEFGRFASFEAQGAL | HLA-DRA*01:01 |
| HLA-DR-SPs pool | HLA-DRB1 ₍₅₇₋₇₀₎ + HLA-DRB1 ₍₇₂₋₈₆₎ + HI | LA-DRB5 ₍₇₂₋₈₆) + HLA-DRB1/5 ₍₁₈₄₋₁₉₉) + HLA-DRA ₍₇₀₋₈₅₎ |

Table S6. Information of source samples of CSF-infiltrating CD4+ T cells, Related to Figure 4

| ID | Gender | Age# | Disease | Treatment | HLA-DRB1*15:01 | HLA-DRB5*01:01 |
|---------|--------|------|---------|-----------|----------------|----------------|
| Donor 1 | M | 45 | RRMS | nihil | +/- | +/- |
| Donor 2 | M | 22 | RRMS | nihil | +/- | +/- |
| Donor 3 | F | 27 | RRMS | nihil | +/- | +/- |
| Donor 4 | F | 34 | RRMS | nihil | +/- | +/- |
| Donor 5 | M | 22 | RRMS | nihil | +/- | +/- |
| Donor 6 | F | 37 | RRMS | nihil | +/- | +/- |
| Donor 7 | M | 27 | RRMS | nihil | +/- | +/- |
| Donor 8 | F | 37 | RRMS | nihil | +/- | +/- |
| Donor 9 | F | 34 | RRMS | nihil | +/- | +/- |

[#]The age (in years) when the sample was collected