

## SUPPLEMENTAL INFORMATION

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## Supplemental Methods

### *Patient Descriptions – MS Patients*

Patient **MS-1** has relapsing-remitting MS (RRMS) with 3 attacks in the 12 months prior to lumbar puncture (LP); brain and cervical spinal cord MRIs were consistent with MS and showed an increase in lesion load in the 7 months prior to LP; she received a 3-day course of 1000 mg intravenous methylprednisolone 2 weeks prior to LP; EDSS was 1.0.

Patient **MS-2** experienced an attack of multifocal neurological symptoms 2 years prior to LP associated with multiple T2 hyperintense foci on MRI suggestive of demyelinating disease, consistent with the diagnosis of a clinically isolated syndrome (CIS). She was treated with a course of glucocorticoids, and gradually improved. LP revealed OCB. This patient developed a clinical relapse while this manuscript was under preparation. Therefore, this patient now carries a diagnosis of MS; EDSS at the time of her LP was 2.0.

Patient **MS-3** has RRMS for >3 years with typical MRI and CSF findings; treatment was with glatiramer acetate (GA) continuously for 15 months prior to LP; EDSS was 2.0. This patient participated in our studies exclusively for research purposes; no WBC, WBC differentiation, and OCB analysis was obtained. Her previous LP revealed CSF-restricted OCB.

Patient **MS-4** has RRMS with 2 attacks within the two years prior to LP; MRI and CSF findings showed changes typical of MS; EDSS was 1.5.

Patient **MS-5** had an initial episode of neurological symptoms 2 months prior to LP; brain MRI performed 1 week prior to LP revealed a single contrast-enhancing lesion; a diagnosis of clinically definite MS was reached by McDonald-criteria (1); EDSS was 1.5.

Patient **MS-6** presented with left face numbness and diplopia 1 month prior to LP; her MS diagnosis was confirmed by repeat brain MRI showing numerous new T2 hyperintense lesions after 6 months; her EDSS was 2.0.

*Patient Descriptions OND-Patients (see MRI images below)*

Patient **OND-1** has a single asymptomatic non-enhancing cerebellar lesion; her CSF is “negative” with respect to OCB and other inflammatory markers (IgG-Index, white cell count), and a clinical and imaging follow-up after 14-months revealed an unchanged clinical and imaging situation. This patient may have radiologically isolated syndrome (RIS) (2) but cannot be classified as having either CIS, or MS (1).

Patient **OND-2**'s most likely diagnosis is a low-grade cervical spinal cord glioma. In particular, her lesion extends beyond the spinal cord circumference and thus appears tumor-like, has grown very slowly over the course of 2 years, does not show pathological contrast enhancement, and is a solitary lesion of considerable size. If this lesion were indeed of a demyelinating nature, a significant clinical neurological deficit would be expected; however, the patient displays only very little neurological symptoms. In this clinical constellation a biopsy of the cervical cord lesion to confirm the diagnosis of glioma is contraindicated given the very high risk of complications.

Patient **OND-3** had a single episode with C3-C4 level myelitis 2 years prior to LP; MRI findings improved over time; CSF was normal; NMO-IgG antibodies were negative. He had been treated with IFN- $\beta$  1a for a presumed CIS, but treatment was discontinued shortly after the LP because the patient did not meet diagnostic criteria for MS (1) and normal CSF findings placed him in a lower risk group to develop MS in the future.

Patient **OND-4** had a single episode of a high cervical cord transverse myelitis 5 months prior to LP. CSF findings and brain MRI were normal. The patient did not meet diagnostic criteria for MS (1).

Patient **OND-5** presented with generalized body pain and on brain MRI had innumerable small foci of increased T2 signal without enhancement; these were largely atypical for demyelinating

disease and were thought to more likely represent microvascular events. CSF was normal. The patient did not meet diagnostic criteria for MS (1).

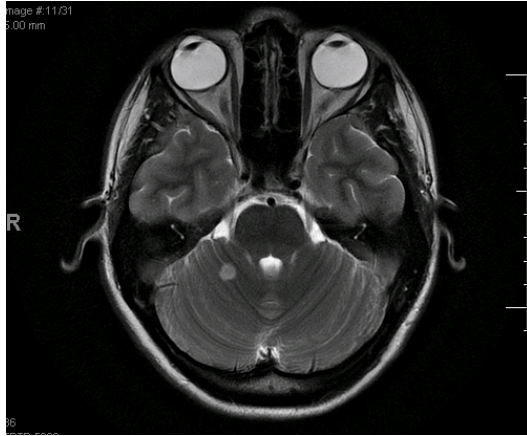
Patient **OND-6** had a slowly progressive clinical myelopathy, and underwent an anterior cervical discectomy and fusion C5/6 in 2009 for spinal canal stenosis. In 2005 cervical spine MRI revealed spinal stenosis from C4 to C6; spinal cord signal abnormality was not noted. In 2008 cervical spine MRI revealed progressive spinal stenosis at C5/6 and a newly described abnormal T2 signal hyperintensity at C4-5, and to a lesser degree at level C5-7; pathological contrast enhancement was not noted. In 2012 cervical spine MRI revealed persistent cervical spinal cord T2 signal abnormality at level C5-6. Her brain MRI revealed no abnormal findings in 2008 and 2012. There is no evidence for a clinical or radiographic dissemination in time or space. Her CSF analysis in 2011 revealed 5 OCB, a normal IgG-Index of 0.6 and normal cell count. The patient did not meet diagnostic criteria for MS (1).

Patient **OND-7** has chronic migraine; in 2009 she developed right shoulder and arm weakness lasting a few weeks. Brain MRI revealed non-specific foci of increased T2 and FLAIR signal; cervical spine MRI was unremarkable. CSF was normal, and an extensive diagnostic workup was unrevealing. The patient did not meet diagnostic criteria for MS (1).

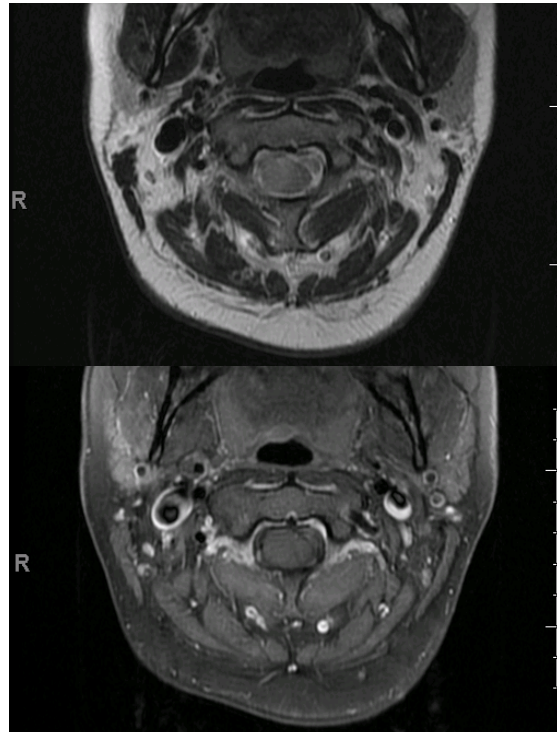
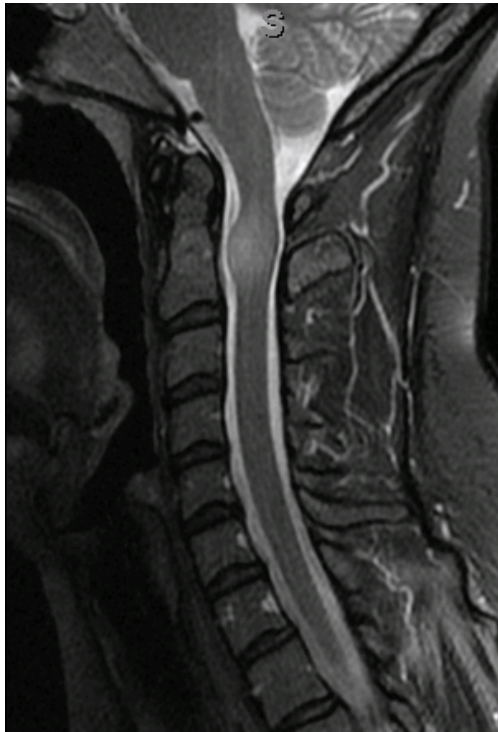
## References

1. Polman, C.H., Reingold, S.C., Banwell, B., Clanet, M., Cohen, J.A., Filippi, M., Fujihara, K., Havrdova, E., Hutchinson, M., Kappos, L., et al. 2011. Diagnostic criteria for multiple sclerosis: 2010 revisions to the McDonald criteria. *Ann Neurol* 69:292-302.
2. Okuda, D.T., Mowry, E.M., Beheshtian, A., Waubant, E., Baranzini, S.E., Goodin, D.S., Hauser, S.L., and Pelletier, D. 2009. Incidental MRI anomalies suggestive of multiple sclerosis: the radiologically isolated syndrome. *Neurology* 72:800-805.

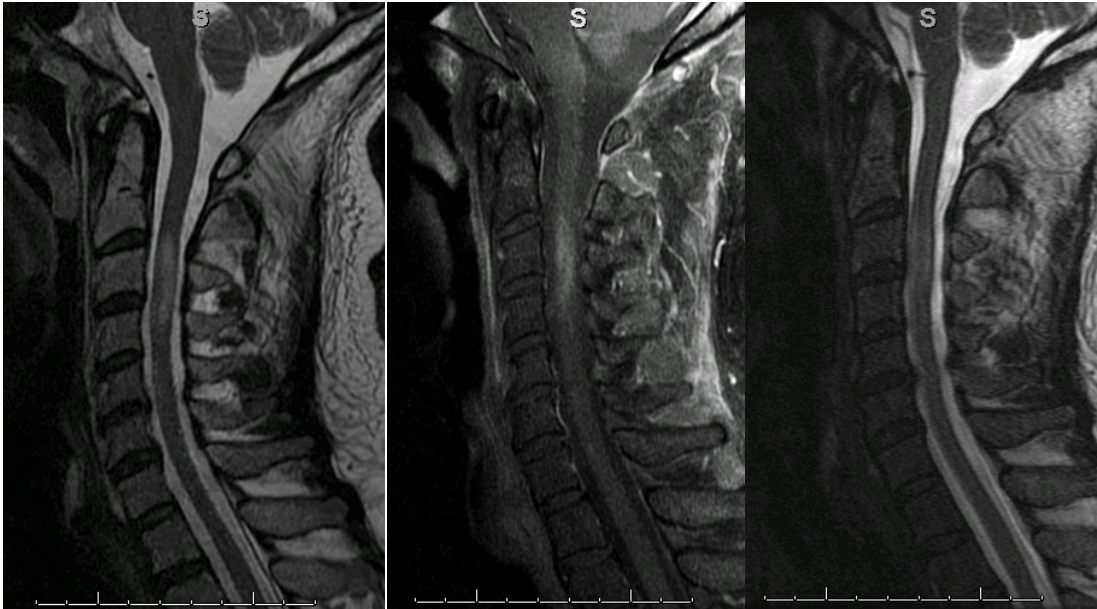
*MRI Images of OND Patients*



Patient **OND-1**: Brain MRI 09/2010. Left panel shows axial T2 with a single focus of increased T2 signal in the left cerebellar hemisphere. This lesion did not show pathological contrast enhancement. 14 months later there was no change to the patient's MRI findings.



Cervical spinal cord MRI (04/2012) of patient **OND-2**. Left panel, sagittal T2; right upper panel axial T2 at level C2; right lower panel, axial T1 post gadolinium at level C2. The C2 level lesion has increased slightly in size since 2009.

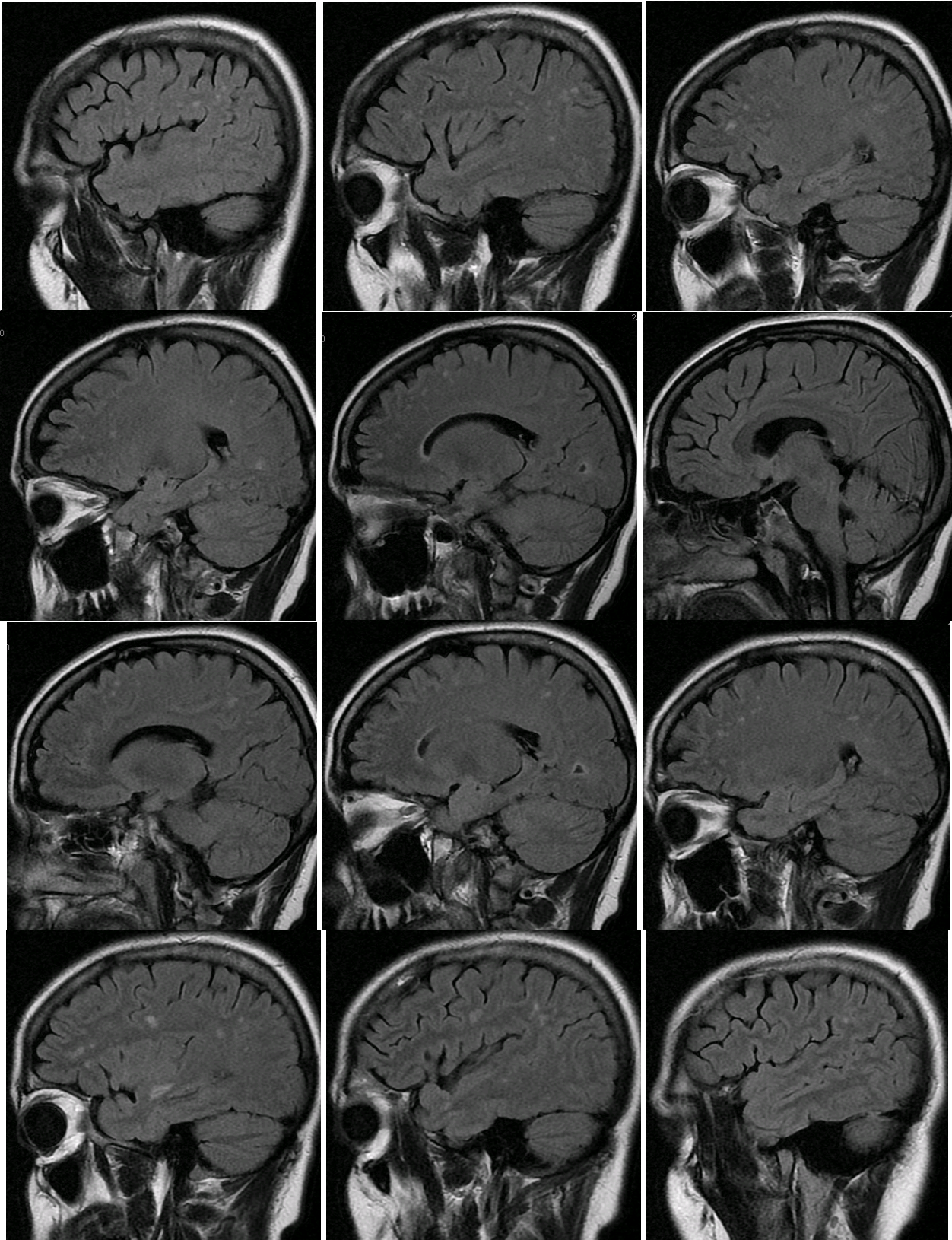


Patient **OND-3**: Left panel: Sagittal T2 C cord 07/2009. Middle panel: Sagittal T1+gadolinium 08/2009. Right panel: Sagittal T2 C cord 09/2009.



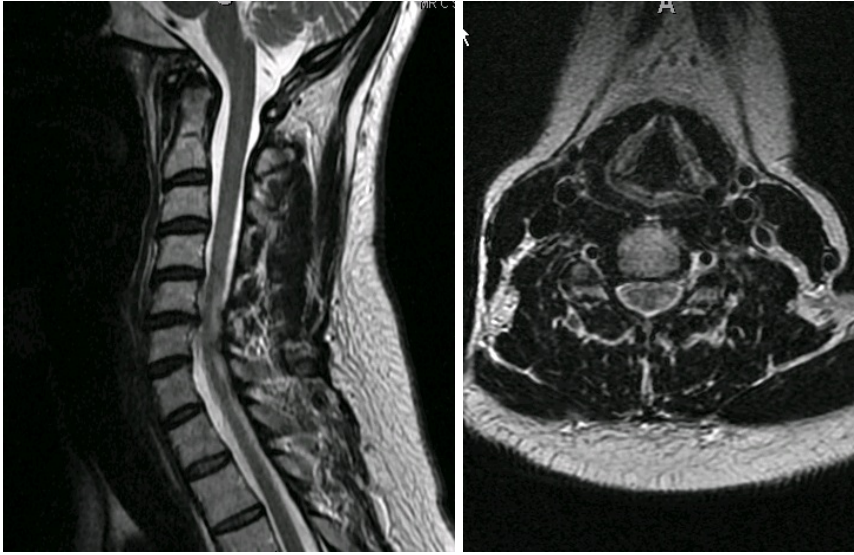
Patient **OND-4**: Left panel: Axial T2 C cord 10/2010 showing a left posterior focus of increased T2 signal at level C1; there was very subtle gadolinium enhancement (not shown). Right panel: Axial T2 C cord 02/2011 showing the same lesion with less T2 hyperintensity 4 months later. Brain MRI was normal (not shown).



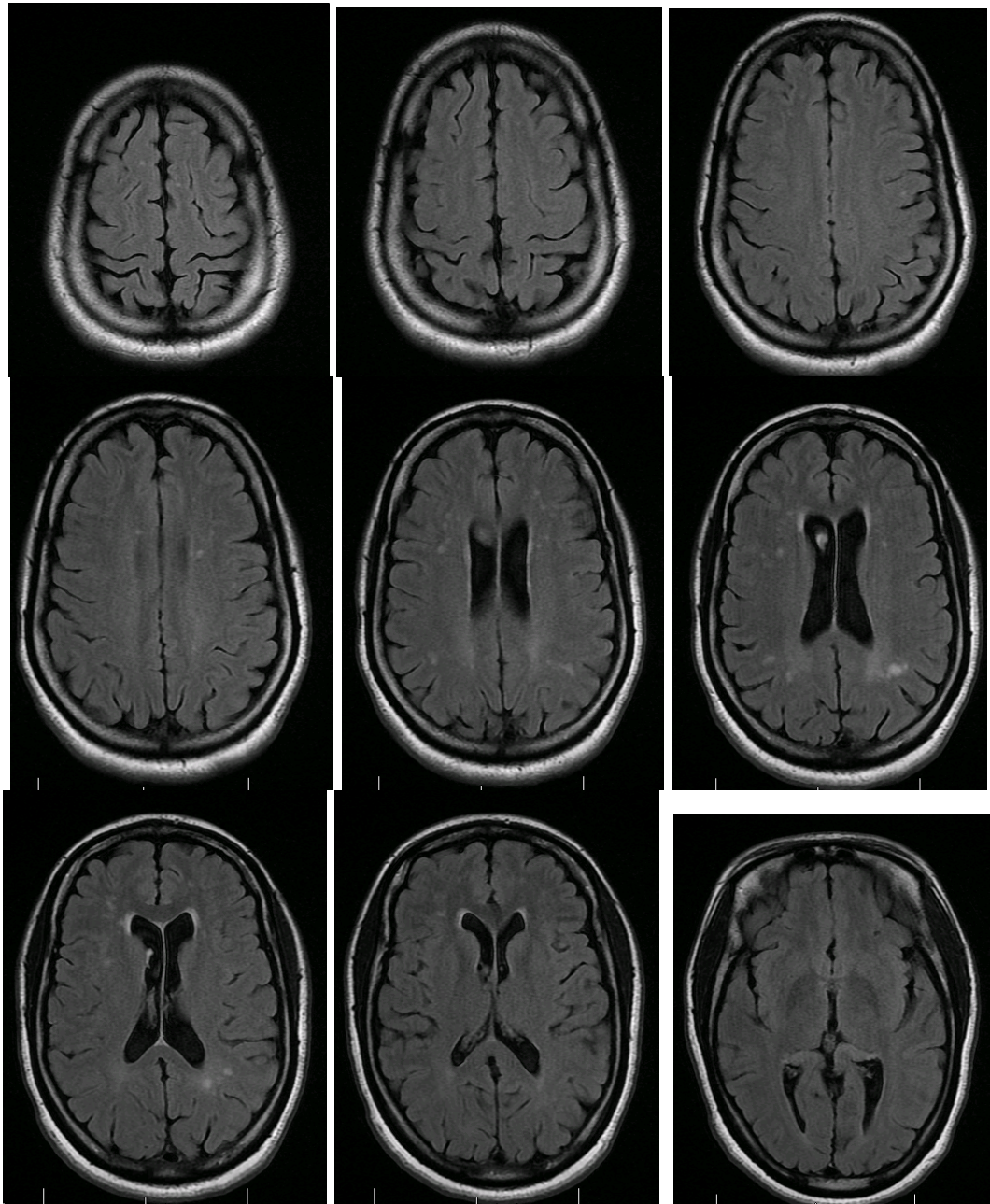


Patient **OND-5**: Sagittal FLAIR 02/2011: Top left panel to bottom right, left to right sections, respectively. There was no pathological contrast enhancement after gadolinium (not shown).





Patient **OND-6**: Left panel:  
Sagittal T2 C cord 07/2009.  
Right panel: Sagittal  
T1+gadolinium 08/2009.



Patient **OND- 7**: Axial FLAIR sequences showing numerous foci of abnormal T2 signal which prompted further MS diagnostic work-up in this patient. As described above, a MS diagnosis could not be established in this patient.

## Supplemental Tables

ID	CSF	PBMC	CSF	PBMC	CSF	PBMC
	(all reads)	(all reads)	(V,J,CDR3)	(V,J,CDR3)	(non-red)	(non-red)
MS-1	6,660	735,584	5,661	507,411	270	17,151
MS-2	27,130	307,641	20,752	235,515	383	9,657
MS-3	34,813	397,723	25,658	243,301	625	10,986
MS-4	11,981	423,533	6,164	347,616	170	10,123
MS-5	30,907	296,397	26,226	256,809	494	9,348
MS-6	34,550	438,860	30,286	344,805	432	13,260
OND-1	21,854	415,570	6,759	341,955	74	11,034
OND-2	11,073	380,129	3,933	308,411	107	17,750
OND-3	18,021	393,717	4,059	324,620	89	9,484
OND-6	26,965	435,258	14,262	339,707	81	5,620
OND-4	-*	317,020	-*	277,157	-*	10,015
OND-5	-*	350,603	-*	273,079	-*	7,549
OND-7	-*	307,517	-*	272,077	-*	7,498

**Supplemental Table 1:** Numbers of sequences obtained by deep repertoire sequencing (“all reads”). “V, J, CDR3” are numbers of sequences with identifiable IGHV, IGHJ and H-CDR3 sequences. Clonally non-redundant datasets (“non-red”) for IGHV usage analyses were generated by considering only one representative from a set of reads bearing the same IGHV-segment, IGHVJ-segment, H-CDR3 length, and H-CDR3 amino acid composition that differed by less than two residues from another member in the clonal family. \*No IgG-VH sequences could be amplified from these patients’ mRNA. Presence of viable mRNA was confirmed by RT-PCR of GAPDH.

Patient ID	Figure	CDR3	IGHV	mut	IGHJ
MS-1		CARVYYKWNDEWSWFDPW	IGHV1-3	28	IGHJ5
		CAREPGTSLHCSSGGNCYSKDasNFW	IGHV1-69	30	IGHJ3
		CARGRYSRSSRYFDLW	IGHV1-69	21	IGHJ4
	2A	CARIEPSSSRGSLFFFDYW	IGHV2-70	18	IGHJ4
		CTTDEGGNSGYFFEYW	IGHV3-15	34	IGHJ4
	5A	CAKAGGYSGSSGRRNFYSFDSW	IGHV3-23	10	IGHJ4
		CARGTGGGWYYRIHFDDYW	IGHV3-23	23	IGHJ4
		CAKVRFVEWLHLFDYW	IGHV3-23	12	IGHJ4
	5C	CARVVNYDILTGYFRDAFDIW	IGHV3-23	16	IGHJ3
		CARDQGGYYGNFFLDYW	IGHV3-48	30	IGHJ4
		CARQKSTRTRTRYMYGLDVW	IGHV3-48	23	IGHJ6
		CAGRRYFESSGPPEFPW	IGHV3-48	16	IGHJ5
	3A	<b>CARHGTSLTSPYSRFFYW</b>	<b>IGHV4-39</b>	<b>18</b>	<b>IGHJ4</b>
		<b>CARSGTKKVMGHFDYW</b>	<b>IGHV4-39</b>	<b>8</b>	<b>IGHJ4</b>
	<b>CARSGPGGEPVYFDPW</b>	<b>IGHV4-59/61</b>	<b>34</b>	<b>IGHJ5</b>	
	<b>CARVRRAGGRSWFDPW</b>	<b>IGHV4-59/61</b>	<b>14</b>	<b>IGHJ5</b>	
	CASHRGQWLVTW	IGHV5-a	27	IGHJ5	
OND-1		CAREDCSAASCYWAYLHHW	IGHV1-18	19	IGHJ1
		CARGEAPTQLQLGTFDFW	IGHV1-46	26	IGHJ4
		CAKDSAGPVLVWKPWFDPW	IGHV3-23	10	IGHJ5
		CARDSMPDKDFADYFPYW	IGHV3-30/33rn	32	IGHJ1
		CTRAGPPELWDYFYGLDVW	IGHV3-49	8	IGHJ6
		CVRDRDWNYGDCW	IGHV3-7	18	IGHJ4
		CVKSPTYSSGQGYFDYW	IGHV3-9	11	IGHJ4
		<b>CVRDRAVAAHDAFDIW</b>	<b>IGHV4-59/61</b>	<b>33</b>	<b>IGHJ3</b>
OND-2		CARTFYDSSEMGGSW	IGHV1-18	10	IGHJ5
		CAHCSGGTCAFDYW	IGHV1-2	3	IGHJ4
		CAREFDSW	IGHV1-2	9	IGHJ4
		CANSTSSETIWFDPW	IGHV1-2	23	IGHJ5
		CGRGERKATPMEVW	IGHV1-8	33	IGHJ6
		CARRRTLGHCSSSSCGRAFDIW	IGHV2-5	17	IGHJ3
		CARDLKGREVMGPHPPDFDYW	IGHV3-30/33rn	15	IGHJ4
		CARDPTQTIPTGFSGYFDYW	IGHV3-30/33rn	20	IGHJ4
		CARARDYYDSSGYGYW	IGHV3-48	13	IGHJ4
		CARDGPTGALDYW	IGHV3-48	12	IGHJ4
		CAKWYNSGYRAFDIW	IGHV3-53/66	17	IGHJ3
		CARGVDSWSGGPGDW	IGHV3-7	23	IGHJ4
		CVLRNSGTLTYFASW	IGHV3-9	21	IGHJ4
		CARWDNYYDSGYFDYW	IGHV4-30-4/31	4	IGHJ4
		CARAYDFWSGYEYAMDVW	IGHV4-34	8	IGHJ6
	CARGRGTMASASSGWDYW	IGHV4-34	36	IGHJ4	
OND-3		CAKAKRHCSNAGCPVMSASDYW	IGHV1-18	16	IGHJ4
		CARGSYDKPFDYW	IGHV1-2	32	IGHJ4
		CARSLKQLVTRTFGYW	IGHV1-2	14	IGHJ4

	CASAQYPDDGLSTYYYYGIDVW	IGHV1-69	13	IGHJ6
	CARGTQIGDLDGYFGLDVW	IGHV1-69	21	IGHJ6
	CARGYGDYGRVDFW	IGHV1-69	23	IGHJ4
	CARDAPKLVAGNKHWDYW	IGHV3-21	14	IGHJ4
	CAKVFRGAVAGSFDYW	IGHV3-23	14	IGHJ4
	CATDLGSSLTGYYYFW	IGHV3-30/33m	22	IGHJ5
	CMVGNHYHSGRHDYW	IGHV3-49	40	IGHJ4
	CARGNPLTNLRRGTGFDPW	IGHV4-34	24	IGHJ5
	<b>CARQGFSYGLRTMAHFDFW</b>	<b>IGHV4-39</b>	<b>30</b>	<b>IGHJ2</b>
	<b>CARQGYDHWSGYSSWFDPW</b>	<b>IGHV4-39</b>	<b>13</b>	<b>IGHJ5</b>
MS-2	CARGPYYYDISGDSYAHYGMVDW	IGHV3-13	12	IGHJ6
	CTRERSYSYLRSYFDYW	IGHV3-15	17	IGHJ4
	CVP IKGAEANYW	IGHV3-15	17	IGHJ4
	CTRGLLWLDGDYRDYW	IGHV3-30/33m	20	IGHJ4
	CAGGQTGYCTGGNSQRCYGMVDW	IGHV3-53/66	34	IGHJ6
	CARVDATSWYYFDSW	IGHV3-53/66	22	IGHJ4
	CAKVKGRQLVHYYYGMVDW	IGHV3-53/66	21	IGHJ6
	CARDEPFDYDTSSYSPFDSW	IGHV3-7	3	IGHJ4
	CAREGVYDNLLETDADFVW	IGHV3-7	19	IGHJ3
	CARVPRRFSAYDV	IGHV3-7	6	IGHJ5
3C	CTTESWYTFYSW	IGHV3-72	32	IGHJ5
	CVRAFTSSWDKGGSDVDFDPW	IGHV4-30-4/31	12	IGHJ5
	CARDERITLIPGAFDVW	IGHV4-30-4/31	7	IGHJ3
	CGRDGRGRELLPFPFDSW	IGHV4-30-4/31	16	IGHJ4
	CARDLDYSGSNLAYW	IGHV4-30-4/31	16	IGHJ4
	CARDLSGSGSYFPVDAW	IGHV4-30-4/31	18	IGHJ5
	CARDPAESGYFSGTFDIW	IGHV4-30-4/31	16	IGHJ3
	CARDRVGWVRGVFPFPAW	IGHV4-30-4/31	8	IGHJ5
	CARETGLFKSFYFDYW	IGHV4-30-4/31	16	IGHJ4
	CARGLTGWYDPDW	IGHV4-30-4/31	29	IGHJ4
	CARGRGYSSNWYPLRFDSW	IGHV4-30-4/31	8	IGHJ4
	CARRASPHHYDGSGEDYW	IGHV4-30-4/31	29	IGHJ4
	CAHTKDAGMLTWLEHW	IGHV4-30-4/31	34	IGHJ4
	CARVDRTAGYYFDNW	IGHV4-30-4/31	14	IGHJ4
	CARVPRGGPNPFI RGTFDYW	IGHV4-30-4/31	14	IGHJ4
	CARVVDYFDHW	IGHV4-30-4/31	17	IGHJ4
	CARGRGFYDGTGHQNYFDPW	IGHV4-34	14	IGHJ5
	<b>CAGDGSYSYTRFDQW</b>	<b>IGHV4-39</b>	<b>7</b>	<b>IGHJ4</b>
	<b>CARGLPRLSPHSW</b>	<b>IGHV4-39</b>	<b>31</b>	<b>IGHJ4</b>
	<b>CARHGLPPYWNVDVSTHQYFDSW</b>	<b>IGHV4-39</b>	<b>19</b>	<b>IGHJ5</b>
	<b>CARHGTGDSW</b>	<b>IGHV4-39</b>	<b>30</b>	<b>IGHJ4</b>
	<b>CARHGTGDSWYRGIRVGDW</b>	<b>IGHV4-39</b>	<b>29</b>	<b>IGHJ4</b>
	<b>CARHPAYSSNWYLP IYFHHW</b>	<b>IGHV4-39</b>	<b>18</b>	<b>IGHJ1</b>
	<b>CARHPDSSDNTGRAFFNPFYDW</b>	<b>IGHV4-39</b>	<b>19</b>	<b>IGHJ4</b>
	<b>CARHPPGSSGFYGHDAFDVW</b>	<b>IGHV4-39</b>	<b>8</b>	<b>IGHJ3</b>
	<b>CARHPTPFIAGPGTFDYW</b>	<b>IGHV4-39</b>	<b>11</b>	<b>IGHJ4</b>



MS-2	CARHRDYDSSGFYRRRAFAMW	IGHV4-39	12	IGHJ3
	CARHRGIAVAVQLNPSRNYFDYW	IGHV4-39	13	IGHJ4
	CVRHSGGHYYDSDSVYKWMSEYFDYW	IGHV4-39	16	IGHJ4
	CTRHSYQLQGVMPINWFDPW	IGHV4-39	20	IGHJ5
	CARHSPHYFYGSGSYQGWFGPW	IGHV4-39	23	IGHJ5
	CAAIRGRMAAAYHW	IGHV4-39	26	IGHJ4
	CVRLGPDYGFNFDFYW	IGHV4-39	13	IGHJ4
	CARLGYCSSISCYADYW	IGHV4-39	11	IGHJ4
	CARLGYCSSISCYAESW	IGHV4-39	15	IGHJ5
	CARLMKGSKDIAVAGTFDYW	IGHV4-39	16	IGHJ4
	CARLNDYDSGDYYWNFDFYW	IGHV4-39	18	IGHJ4
	CARLQYGSYPIYYYGMDVW	IGHV4-39	13	IGHJ6
	CVALRYFDWSVGKLFDFYW	IGHV4-39	15	IGHJ4
	CARPGGSDGYNPFDFW	IGHV4-39	15	IGHJ4
	CARPVAGEDYGGNYWFDPW	IGHV4-39	15	IGHJ5
	CARQGYYYDSSGLFDYW	IGHV4-39	16	IGHJ4
	CARQPNRFLSDRSGGDYW	IGHV4-39	13	IGHJ5
	CARQPNRFYYDSSGGDYW	IGHV4-39	10	IGHJ5
	CTKRSSWYGEHYGMDVW	IGHV4-39	20	IGHJ6
	CAKSPPLGNSGAFDIW	IGHV4-39	15	IGHJ3
	CARVEMSTIRGVGLLDYW	IGHV4-39	13	IGHJ4
	CARVGPPTSDAVPGGLFYWYFDLW	IGHV4-39	25	IGHJ2
	CVAVRYFAWSVGKLFDFYW	IGHV4-39	16	IGHJ4
	CARDTFRQTTMVTRWFDPW	IGHV4-4	18	IGHJ5
	CARTVRYVDWRNKLLYHFDHW	IGHV4-4	29	IGHJ4
	CARVPLLRDGGIYYGLDVW	IGHV4-4	14	IGHJ6
	CARAYDTNSQGPFDWF	IGHV4-59/61	22	IGHJ4
	CARGSDILTGYSLGGWFDPW	IGHV4-59/61	13	IGHJ5
	CARLRRDGPQEYFHHW	IGHV4-59/61	22	IGHJ1
	CAKRAAYCDGKRCRAFDYW	IGHV4-59/61	16	IGHJ4
	CASRMGGQQLVWGYW	IGHV4-59/61	17	IGHJ4
	CARSVGASGTLTGYPFHHW	IGHV4-59/61	19	IGHJ1
CARYFAPGHYSGDGAFDIW	IGHV4-59/61	19	IGHJ3	
CARRIAPLRGTYSMFAFDIW	IGHV5-51	16	IGHJ3	
CARPQRSSSSTFDHW	IGHV5-a	25	IGHJ4	
MS-3	CARGGYCTGGSCYEPEFDYW	IGHV1-18	13	IGHJ4
	CAHWESGYSYSRGGGSRRRYFDSW	IGHV2-5	11	IGHJ4
	CTTVKLGWRSSYYDTAFDYW	IGHV3-15	16	IGHJ4
	CVKDQGDYIWGTYPSTFDYW	IGHV3-23	20	IGHJ4
	CANRYLVDSSGSYRDDPFDIW	IGHV3-23	5	IGHJ3
	CAKTRYADYYVGNFYFNW	IGHV3-23	20	IGHJ4
	CTRARVLAARSRYADYNYGMDVW	IGHV3-30/33rn	24	IGHJ6
	CARKPDRGYNLDDYGEYKPTSPFDYW	IGHV3-30/33rn	21	IGHJ4
	CARDAPDCGGDCYSLPTVRFDFYW	IGHV3-7	10	IGHJ4
	CARDEETFVWGSYRDQTNYYYYGMDVW	IGHV3-7	8	IGHJ6
	CVRDHQWLVLGRRCDSW	IGHV3-7	17	IGHJ5

MS-3	CAKDWTDAGAVTNVFDYW	IGHV3-7	24	IGHJ4
	CARDYGSATYYASYHGMGVW	IGHV3-7	13	IGHJ6
	CARGIAAYSGGRYDSHFDSW	IGHV3-7	21	IGHJ4
	CARGQGSGRYYRLRFDYW	IGHV3-7	11	IGHJ4
	CARRGATTPRGRTLDSW	IGHV3-7	9	IGHJ4
	CARVLTVRGVSSQGFDSW	IGHV3-7	18	IGHJ4
	CARVPPWVGTITPLFDYW	IGHV3-7	6	IGHJ4
	CARGYHSFDMW	IGHV3-72	15	IGHJ3
	CARDAKWFGESNYDMDVW	IGHV4-30-4/31	12	IGHJ6
	CARDKTWLGELGIEHNAFDVW	IGHV4-30-4/31	13	IGHJ3
	CARDLHDYGDYFDYW	IGHV4-30-4/31	10	IGHJ4
	CARDYDYGDKWFDPW	IGHV4-30-4/31	14	IGHJ5
	CAREGGAVAGILVW	IGHV4-30-4/31	14	IGHJ5
2B	CAREGIPDYDYGMVW	IGHV4-30-4/31	22	IGHJ6
	CARESAHDSSYFYFDSW	IGHV4-30-4/31	10	IGHJ4
	CARGPAVLRYPDRLLHFDYW	IGHV4-30-4/31	34	IGHJ4
	CASNVDVLTAYYSGGSYFDHW	IGHV4-30-4/31	12	IGHJ4
	CASQYYDILTAYYVGSWFDPW	IGHV4-30-4/31	13	IGHJ5
	CASRSGYTNNNNWFDPW	IGHV4-30-4/31	11	IGHJ5
	CVRSKYDILTGYDYGKGFHFIW	IGHV4-30-4/31	10	IGHJ3
	CARTGMRNWFDLW	IGHV4-30-4/31	17	IGHJ5
	CAKTSGWFGVDSW	IGHV4-30-4/31	19	IGHJ4
	CARVPAAIDTLKFLFDSW	IGHV4-30-4/31	6	IGHJ4
	CARVPDYESDPVIWGSYFDAW	IGHV4-30-4/31	25	IGHJ5
	CRWGGDMDVW	IGHV4-34	20	IGHJ6
	CARGQYSEWDLTRPHYYYAMVW	IGHV4-34	14	IGHJ6
	<b>CASEEYDIMNGRYGLSNWFGAW</b>	<b>IGHV4-39</b>	<b>12</b>	<b>IGHJ5</b>
	<b>CARFPAQILPGYYVGRAGDYW</b>	<b>IGHV4-39</b>	<b>12</b>	<b>IGHJ4</b>
	<b>CAKGEEGYDTSGIFQOW</b>	<b>IGHV4-39</b>	<b>31</b>	<b>IGHJ1</b>
	<b>CAGGGHWPQVDFW</b>	<b>IGHV4-39</b>	<b>23</b>	<b>IGHJ4</b>
	<b>CARHEIKQTPPSTSKFDPW</b>	<b>IGHV4-39</b>	<b>17</b>	<b>IGHJ5</b>
	<b>CARHELLQTPPSTSKFGPW</b>	<b>IGHV4-39</b>	<b>30</b>	<b>IGHJ5</b>
	<b>CARHIALPARAMYHDTIGTPGPLDFW</b>	<b>IGHV4-39</b>	<b>17</b>	<b>IGHJ5</b>
	<b>CARHNYDILTGYYSRPHYSWSW</b>	<b>IGHV4-39</b>	<b>6</b>	<b>IGHJ4</b>
	<b>CARHNYDIL</b>	<b>IGHV4-39</b>	<b>6</b>	<b>IGHJ4</b>
	<b>CARHQTGDDDYFDYW</b>	<b>IGHV4-39</b>	<b>16</b>	<b>IGHJ4</b>
	<b>CARHVLAAGGTLPWGPKFSPKYFDPW</b>	<b>IGHV4-39</b>	<b>21</b>	<b>IGHJ5</b>
	<b>CARLNNFSTSTSYDPPYYFYAMVW</b>	<b>IGHV4-39</b>	<b>11</b>	<b>IGHJ6</b>
	<b>CARLSSDYINWDFW</b>	<b>IGHV4-39</b>	<b>12</b>	<b>IGHJ5</b>
	<b>CARLYYDVLTAAYGLPSYLDYW</b>	<b>IGHV4-39</b>	<b>10</b>	<b>IGHJ4</b>
	<b>CARPHYDILTAYYVNASWFDPW</b>	<b>IGHV4-39</b>	<b>13</b>	<b>IGHJ5</b>
	<b>CARPHYDILTGRYNVAHWDFPW</b>	<b>IGHV4-39</b>	<b>8</b>	<b>IGHJ5</b>
	<b>CASPPEQGIW</b>	<b>IGHV4-39</b>	<b>20</b>	<b>IGHJ3</b>
	<b>CARPPGTSVSWYFDLW</b>	<b>IGHV4-39</b>	<b>18</b>	<b>IGHJ2</b>
	<b>CASPRAIGATGPFYDW</b>	<b>IGHV4-39</b>	<b>21</b>	<b>IGHJ4</b>
	<b>CVRPSINYDSSGYFHDADFIDW</b>	<b>IGHV4-39</b>	<b>13</b>	<b>IGHJ3</b>

MS-3		<b>CVRQGFYGSPTYYPEDW</b>	IGHV4-39	32	IGHJ4	
		<b>CARQGYDILNAYYGRPHYFDYW</b>	IGHV4-39	18	IGHJ4	
		<b>CARQPGDYDSTEYFDYW</b>	IGHV4-39	14	IGHJ4	
		<b>CARQYYDEYSPNWFDSW</b>	IGHV4-39	16	IGHJ5	
		<b>CASQYYDILTGRYSAGQYFDYW</b>	IGHV4-39	7	IGHJ4	
		<b>CARRGYDMLTAYYGEGNWFDPW</b>	IGHV4-39	20	IGHJ5	
		<b>CARRPTIYPGSESAYRVVSW</b>	IGHV4-39	14	IGHJ5	
		<b>CASSSSWKGHFQHW</b>	IGHV4-39	18	IGHJ1	
		<b>CVGSSSSWKGFDPW</b>	IGHV4-39	15	IGHJ5	
		<b>CASSTGTTLRRSYFDSW</b>	IGHV4-39	35	IGHJ4	
		<b>CASSTGTTLRRSHFADW</b>	IGHV4-39	18	IGHJ4	
		<b>CARTSLLVPDTGPW</b>	IGHV4-39	15	IGHJ5	
		<b>CATVVPGNYVSSGYFPDYW</b>	IGHV4-39	10	IGHJ4	
		CARADCDISGYSSWYFDYW	IGHV4-4	21	IGHJ4	
		CARAYYDISGYSSWYFDYW	IGHV4-4	23	IGHJ4	
		<b>CAKANVRHGFRLVGNYPMDAW</b>	IGHV4-59/61	19	IGHJ6	
		<b>CARDLGNDFDW</b>	IGHV4-59/61	13	IGHJ4	
		<b>CARDLGQFDSW</b>	IGHV4-59/61	18	IGHJ4	
		<b>CAREMEGSLW</b>	IGHV4-59/61	25	IGHJ5	
		<b>CARGKEGDLRDSFYFDYW</b>	IGHV4-59/61	8	IGHJ4	
		<b>CARHFSDLVGTISYWRENYYYGMDVW</b>	IGHV4-59/61	11	IGHJ6	
		<b>CARHRANGGMDVW</b>	IGHV4-59/61	28	IGHJ6	
		<b>CAGNRLGYDDNGHPGMDLW</b>	IGHV4-59/61	23	IGHJ6	
		<b>CARRPGTGPIPYLYYGMDVW</b>	IGHV4-59/61	16	IGHJ6	
		<b>CARTEYNWFDPW</b>	IGHV4-59/61	26	IGHJ5	
		<b>CARTYGSPTYSRDYYGMDVW</b>	IGHV4-59/61	10	IGHJ6	
		<b>CARVKVNFDSGYSYTRSFYW</b>	IGHV4-59/61	7	IGHJ4	
		<b>CARVRSAMDVW</b>	IGHV4-59/61	9	IGHJ6	
	MS-4		CARRERYSDSWYDYW	IGHV1-3	10	IGHJ4
		2C	CAKSDDYDFHNIDSW	IGHV3-23	21	IGHJ4
		CARDVFDAWYDHRFDW	IGHV3-53/66	15	IGHJ4	
		CARDFYYTSGRYALDLW	IGHV3-7	23	IGHJ3	
		CARDQYSDGWPGTLGPRRLYYGVAVW	IGHV3-7	13	IGHJ6	
		<b>CARDLGRGGYDYVWGTYSRVFDYW</b>	IGHV4-39	8	IGHJ4	
		<b>CARDWYNTGWSPFYFDYW</b>	IGHV4-39	15	IGHJ4	
		<b>CARSWGFGWEYFDYW</b>	IGHV4-39	20	IGHJ4	
		CARDSDSGSWYARYFDVW	IGHV4-4	16	IGHJ2	
		<b>CAREIRFCNTASCHKWIDPW</b>	IGHV4-59/61	20	IGHJ5	
		<b>CAVGEMTTIVGSYYYFGMDVW</b>	IGHV4-59/61	14	IGHJ6	
		<b>CAQGEVVGTSPSHFYYPMDVW</b>	IGHV4-59/61	12	IGHJ6	
MS-5		CARDLGGGAIYYYSYMDVW	IGHV1-18	13	IGHJ6	
		CARERSRNAVVSEGAFDVW	IGHV1-18	42	IGHJ3	
		CAREPYTAMAASFYDW	IGHV1-2	1	IGHJ4	
		CARGLDGYNWNVGYW	IGHV1-2	1	IGHJ4	
		CARVSIKSGSYLHDIW	IGHV1-2	7	IGHJ4	
		CARDGVYGARNYPWFYDLW	IGHV1-3	17	IGHJ2	

MS-5	CIRELSGGQFDNW	IGHV1-46	26	IGHJ4
	CGSGSTPDFMDVW	IGHV1-46	24	IGHJ6
	CARGDYGDYGYYYYYMDVW	IGHV1-69	12	IGHJ6
	CARGIYIAGAGGKTYFDYW	IGHV1-69	27	IGHJ4
	CATVTFPGYMDVW	IGHV1-69	25	IGHJ6
	CALACRNAVCPQRDFDFNYW	IGHV1-8	19	IGHJ4
	CTTDRTVGATRDFGYW	IGHV3-15	4	IGHJ4
	CAKDGWELHQDAFDVW	IGHV3-23	17	IGHJ3
	CAKDRAVGVGSGDSFESW	IGHV3-23	18	IGHJ3
	CAMHRYSFAYSVIDVW	IGHV3-23	8	IGHJ4
	CARSPARLLDVW	IGHV3-23	15	IGHJ6
	CVRGSRALGGTGDYW	IGHV3-30/33rn	24	IGHJ4
	CARLFYISKQGYDMDVW	IGHV3-30/33rn	0	IGHJ6
	CARDRQYQLLPAPTWFDPW	IGHV3-48	16	IGHJ5
	CAREPGDFWSGDYFFDCW	IGHV3-48	20	IGHJ4
	CARGGSLLYGYCSDGNCYLDYW	IGHV3-53/66	8	IGHJ4
	CASSSLR	IGHV3-7	14	IGHJ5
2E	CVGFNPPIDYW	IGHV3-7	13	IGHJ4
	CARSAVAGNYW	IGHV3-7	18	IGHJ4
	CARNADYDILTGYRPGNFDFW	IGHV3-7	21	IGHJ4
	CARDSSRWSFDVW	IGHV3-72	31	IGHJ6
4A	CSRMYNWNFDYW	IGHV3-72	19	IGHJ4
	CTRRGCTSVSCSSIW	IGHV3-73	14	IGHJ5
	CARDPHDYGGNRFDYW	IGHV3-74	3	IGHJ4
	CSRQHNFWTGSPPYMDVW	IGHV3-74	12	IGHJ6
	CAEFGRHGDYW	IGHV3-74	19	IGHJ4
	CAKEILRSKYDLWSGYKPFDIW	IGHV3-9	17	IGHJ3
	CAKEPDSSGWSTGGFDPW	IGHV3-9	13	IGHJ5
3B	CASQSGDIVSSWYMDVW	IGHV4-30-2	1	IGHJ6
	CARCHYGSKFFDYW	IGHV4-30-4/31	18	IGHJ4
	CARGCTGGNCYFDYW	IGHV4-30-4/31	15	IGHJ4
	CASPRTNYDPGNPMPFDIW	IGHV4-30-4/31	12	IGHJ3
	CARRTSRREGVNWFDPW	IGHV4-30-4/31	17	IGHJ5
	CARSSGPRFDPW	IGHV4-30-4/31	30	IGHJ5
	CARAFSTDLWGGYYKGNWFDPW	IGHV4-34	11	IGHJ5
	CASALSTDLWSGYKGNWFDPW	IGHV4-34	3	IGHJ5
	<b>CARDGNDFWSGHHAGYFDLW</b>	<b>IGHV4-39</b>	<b>22</b>	<b>IGHJ4</b>
4C	<b>CAREGPMVRGVTRTFDCW</b>	<b>IGHV4-39</b>	<b>21</b>	<b>IGHJ4</b>
5B	<b>CARHHPSWATTGPDSW</b>	<b>IGHV4-39</b>	<b>28</b>	<b>IGHJ4</b>
	<b>CVRPSGGRNWFYFDVW</b>	<b>IGHV4-39</b>	<b>24</b>	<b>IGHJ2</b>
	<b>CARQLIARSQVDIVVVVTGTTFDYW</b>	<b>IGHV4-39</b>	<b>13</b>	<b>IGHJ4</b>
	<b>CARQYPSWSTTGPDIW</b>	<b>IGHV4-39</b>	<b>4</b>	<b>IGHJ4</b>
	<b>CARRRHSGVTDSYIWGSEREDAFDIW</b>	<b>IGHV4-39</b>	<b>10</b>	<b>IGHJ3</b>
	CARDQGSSGWGDAFELW	IGHV4-4	18	IGHJ3
	CATGGPRMDYW	IGHV4-4	22	IGHJ4
	<b>CARARWFGELRPPAPYMDVW</b>	<b>IGHV4-59/61</b>	<b>14</b>	<b>IGHJ6</b>

MS-5	<b>CARFGARYCDSPRCQGYYYYYYMDVW</b>	<b>IGHV4-59/61</b>	<b>25</b>	<b>IGHJ6</b>
	<b>CARFIDSSIWGSFPDYW</b>	<b>IGHV4-59/61</b>	<b>5</b>	<b>IGHJ4</b>
	<b>CAGRGFWSPPYYYYMDVW</b>	<b>IGHV4-59/61</b>	<b>2</b>	<b>IGHJ6</b>
	<b>CARVGDDFWSGFWPLW</b>	<b>IGHV4-59/61</b>	<b>30</b>	<b>IGHJ5</b>
	<b>CARVGTTSYYYYYMDVW</b>	<b>IGHV4-59/61</b>	<b>5</b>	<b>IGHJ6</b>
	3D <b>CARVKEDFWSGYTFDYW</b>	<b>IGHV4-59/61</b>	<b>13</b>	<b>IGHJ4</b>
	<b>CARVNVETRDSTSAPYYGAVDFW</b>	<b>IGHV4-59/61</b>	<b>20</b>	<b>IGHJ4</b>
OND-6	CARDAAGTYDFWSGYYSW	IGHV3-7	9	IGHJ4
	CARGGDGYNFYHYHGMVW	IGHV4-30-4/31	11	IGHJ6
MS-6	CAREHRYGSFATGDYFDYW	IGHV1-2	13	IGHJ4
	CARERVVSSSSNWGYW	IGHV1-2	19	IGHJ4
	CARDDGWWSGFHGDYRWFESW	IGHV1-69	25	IGHJ5
	CASGDLGHCTRTHCYEDQYYYYYMDVW	IGHV1-69	23	IGHJ6
	CARPVTGTRGAFDIW	IGHV1-69	15	IGHJ3
	CASSAYCSGGSCLNWFDPW	IGHV1-69	21	IGHJ5
	2D CASGDLGHCTTSNCYEDQYYYYYMDVW	IGHV1-69	14	IGHJ6
	CARRRSPPYRSGMDVW	IGHV3-13	9	IGHJ6
	CVREYNWNDGNWIDPW	IGHV3-21	26	IGHJ5
	3E CAKRWGAGQFDYW	IGHV3-23	20	IGHJ4
	4B CAREGFGVFDYW	IGHV3-30/33rn	20	IGHJ4
	5D CAKGIGRDYDFWSGYFYW	IGHV3-30/33rn	10	IGHJ4
	CAKKDVNYYGIDVW	IGHV3-30/33rn	15	IGHJ6
	CATEGVDCRAGNCYWGFYYW	IGHV3-48	22	IGHJ4
	CATEGVDCRAGNCYW	IGHV3-48	21	IGHJ6
	CARDRAVVTDYDSSGYQNDVDVFDLW	IGHV3-49	24	IGHJ3
	<b>CARDRELDGYNSPFFDYW</b>	<b>IGHV4-59/61</b>	<b>19</b>	<b>IGHJ4</b>
<b>CARTYNWNYWSGLDYW</b>	<b>IGHV4-59/61</b>	<b>13</b>	<b>IGHJ4</b>	

**Supplemental Table 2:** Characteristics of IgG-VH clusters containing clonally related sequences present either exclusively in the CSF or in both, CSF, and PB. IgG-VH sequences were clustered based on usage of identical IGHV and IGHJ germline segments and H-CDR3 aminoacid sequences using a defined distance metric (see Methods). “CDR3” is the most frequent H-CDR3 aminoacid sequence present per cluster. IGHV and IGHJ are the closest germline segments per network and “Mut” are average numbers of nucleotide mutations present in IGHV ranging from CDR1 to FR3 among all sequences present in the cluster. IgG-VH using IGHV4-39 and IGHV4-59/61 which appear to be overrepresented in MS CSF are in bold letters. Per cluster all IgG-VH sequences that contained high-quality sequence information ranging from the 5’ end of H-CDR1 to the 3’ end of H-CDR3 were selected from the database to generate IgG-VH lineage trees (Figures 2 to 5); column “Figure” indicates respective trees in Figures 2 to 5 and clusters in Figure 1.



IGHV	MS-1	MS-2	MS-3	MS-4	MS-5	MS-6	OND-1	OND-2	OND-3	OND-6	p
IGHV1-18	0	0.26	3.04	0	1.01	0	17.57	0	5.62	0	n.s.
IGHV1-2	0	0.52	0	0	1.01	12.96	0	6.54	2.25	0	n.s.
IGHV1-24	0	0	0	0	0	0	0	1.87	0	0	n.s.
IGHV1-3	8.15	0	0	5.88	0.4	0	0	0.93	0	0	n.s.
IGHV1-46	0	0	0	0	1.62	0	50	0.93	0	0	n.s.
IGHV1-69	14.07	0.26	0	0	1.82	64.81	0	0.93	25.84	1.23	n.s.
IGHV1-8	0	0	0	0	0.61	0	0	0.93	0	0	n.s.
IGHV2-26	0	0.26	0	0	0	0	0	0	0	0	n.s.
IGHV2-5	0	0.26	1.6	0	0	0	0	1.87	0	0	n.s.
IGHV2-70	10.37	0	0	0	0	0	0	0	0	0	n.s.
IGHV3-11	0	0	0	0	0.2	0.46	0	1.87	4.49	0	n.s.
IGHV3-13	0	4.44	0	0	0.2	0.46	0	0.93	0	0	n.s.
IGHV3-15	2.96	1.04	0.16	0	0.4	0	0	0	0	0	n.s.
IGHV3-20	0	0	0	0	0.2	0	0	0	0	0	n.s.
IGHV3-21	0	0	0	0	1.01	2.08	0	0.93	5.62	0	n.s.
IGHV3-23	21.85	0.78	4.8	17.06	2.83	0.23	1.35	0.93	4.49	0	n.s.
IGHV3-	0.37	0.78	0.48	0	1.62	14.12	2.7	22.43	0	4.94	n.s.
IGHV3-48	20	0	0	0	1.42	0	0	17.76	0	0	n.s.
IGHV3-49	0	0	0	0	0.2	0.69	10.81	0.93	5.62	0	n.s.
IGHV3-53/66	0.37	2.61	0	8.82	2.43	0	0	10.28	0	0	n.s.
IGHV3-7	0	2.87	10.24	18.82	9.51	0.23	1.35	0.93	0	0	n.s.
IGHV3-72	0	0	0.16	0	7.49	0	0	0.93	0	0	n.s.
IGHV3-74	0	0	0	0	0.81	0.23	0	1.87	0	0	n.s.
IGHV3-9	0	0	0	0	1.62	0	9.46	14.95	0	0	n.s.
IGHV3-h	2.22	0	0	0	0	0	0	2.8	0	0	n.s.
IGHV4-30-2	0	0.78	0.16	0	0.4	0	0	0	0	0	n.s.
IGHV4-30-	0	19.84	16	0	20.04	0	0	1.87	1.12	93.83	n.s.
IGHV4-34	0	1.04	1.6	0	0.81	0	0	3.74	29.21	0	n.s.
<b>IGHV4-39</b>	<b>8.89</b>	<b>42.56</b>	<b>52.48</b>	<b>18.24</b>	<b>23.68</b>	<b>0</b>	<b>0</b>	<b>0</b>	<b>14.61</b>	<b>0</b>	<b>0.04</b>
IGHV4-4	0	3.13	0.64	10.59	0.81	0	0	0	0	0	n.s.
<b>IGHV4-59/61</b>	<b>10.74</b>	<b>13.32</b>	<b>8.32</b>	<b>20.59</b>	<b>17.21</b>	<b>3.7</b>	<b>6.76</b>	<b>1.87</b>	<b>1.12</b>	<b>0</b>	<b>0.01</b>
IGHV4-b	0	0	0.32	0	0	0	0	0	0	0	n.s.
IGHV5-51	0	4.96	0	0	0.61	0	0	0	0	0	n.s.
IGHV5-a	0	0.26	0	0	0	0	0	0	0	0	n.s.
IGHV7-4-1	0	0	0	0	0	0	0	0.93	0	0	n.s.

**Supplemental Table 3:** Percent usage per IGHV germline segment in the CSF of MS and OND patients. Comparison was made between MS and OND and p-values were calculated using a resampling-based permutation test. Only germline segments are represented which were found in at least one CSF sample. IGHV4-39 and IGHV4-59/61 appear significantly overrepresented in MS CSF; n.s., not significant.

IGHV	MS-1	MS-2	MS-3	MS-4	MS-5	MS-6	OND-1	OND-2	OND-3	OND-6	p
IGHV1-18	0	0	1.18	0	3.28	0	12.50	6.25	7.69	0	n.s.
IGHV1-2	0	0	0	0	4.92	11.11	0	18.75	15.38	0	n.s.
IGHV1-3	5.88	0	0	8.33	1.64	0	0	0	0	0	n.s.
IGHV1-46	0	0	0	0	3.28	0	12.50	0	0	0	n.s.
IGHV1-69	11.76	0	0	0	4.92	27.78	0	0	23.08	0	n.s.
IGHV1-8	0	0	0	0	1.64	0	0	6.25	0	0	n.s.
IGHV2-5	0	0	1.18	0	0	0	0	6.25	0	0	n.s.
IGHV2-70	5.88	0	0	0	0	0	0	0	0	0	n.s.
IGHV3-13	0	1.41	0	0	0	5.56	0	0	0	0	n.s.
IGHV3-15	5.88	2.82	1.18	0	1.64	0	0	0	0	0	n.s.
IGHV3-21	0	0	0	0	0	5.56	0	0	7.69	0	n.s.
IGHV3-23	23.53	0	3.53	8.33	6.56	5.56	12.50	0	7.69	0	n.s.
IGHV3-30/33rn	0	1.41	2.35	0	3.28	16.67	12.50	12.50	7.69	0	n.s.
IGHV3-48	17.65	0	0	0	3.28	11.11	0	12.50	0	0	n.s.
IGHV3-49	0	0	0	0	0	5.56	12.50	0	7.69	0	n.s.
IGHV3-53/66	0	4.23	0	8.33	1.64	0	0	6.25	0	0	n.s.
IGHV3-7	0	4.23	11.76	16.67	6.56	0	12.50	6.25	0	50.00	n.s.
IGHV3-72	0	1.41	1.18	0	3.28	0	0	0	0	0	n.s.
IGHV3-73	0	0	0	0	1.64	0	0	0	0	0	n.s.
IGHV3-74	0	0	0	0	4.92	0	0	0	0	0	n.s.
IGHV3-9	0	0	0	0	3.28	0	12.50	6.25	0	0	n.s.
IGHV4-30-2	0	0	0	0	1.64	0	0	0	0	0	n.s.
IGHV4-30-4/31	0	21.13	18.82	0	8.20	0	0	6.25	0	50.00	n.s.
IGHV4-34	0	1.41	2.35	0	3.28	0	0	12.50	7.69	0	n.s.
IGHV4-39	11.76	45.07	38.82	25.00	14.75	0	0	0	15.38	0	0.06
IGHV4-4	0	4.23	2.35	8.33	3.28	0	0	0	0	0	0.07
<b>IGHV4-59/61</b>	<b>11.76</b>	<b>9.86</b>	<b>15.29</b>	<b>25.00</b>	<b>13.11</b>	<b>11.11</b>	<b>12.50</b>	<b>0</b>	<b>0</b>	<b>0</b>	<b>0.02</b>
IGHV5-51	0	1.41	0	0	0	0	0	0	0	0	n.s.
IGHV5-a	5.88	1.41	0	0	0	0	0	0	0	0	n.s.

Supplemental Table 4: Percent usage per IGHV germline segment amongst **IgG-VH clusters** (Figure 1, Table S2) of closely related or identical IgG-VH with at least one CSF clone in MS and OND patients. Comparison was made between MS and OND and p-values were calculated using a resampling-based permutation test. Only germline segments are represented which were found in at least one B cell cluster. IGHV4-59/61 is significantly overrepresented in MS B cell clusters, IGHV4-39 and IGHV4-4 reach marginal statistical significance; n.s., not significant ( $p>0.05$ ).

IGHV	MS-1	MS-2	MS-3	MS4	MS-5	MS-6	OND-1	OND-2	OND-3	OND-4	OND-5	OND-6	OND-7	p
IGHV1-14	0	0	0	0	0	0.01	0	0.01	0.01	0	0	0	0	n.s.
IGHV1-17	0	0	0.01	0	0	0	0	0	0.01	0	0	0	0	n.s.
IGHV1-18	5.77	4.79	5.52	5.28	5.57	4.30	4.22	5.52	5.62	7.72	4.58	3.54	3.31	n.s.
IGHV1-2	2.76	2.34	2.37	4.56	2.11	3.58	2.83	5.30	6.66	0.63	0.64	5.30	5.01	n.s.
IGHV1-24	1.59	0.86	1.51	1.08	0.80	1.13	1.25	1.93	1.19	1.79	0.15	0.87	1.09	n.s.
IGHV1-3	2.05	3.55	2.93	1.51	2.47	1.24	2.59	1.68	0	2.30	2.92	0	0.02	n.s.
IGHV1-45	0	0	0	0.13	0	0	0	0	0	0	0.11	0	0	n.s.
IGHV1-46	1.96	1.68	3.90	2.03	2.01	1.62	5.87	1.70	1.21	4.62	2.23	2.10	0.56	n.s.
IGHV1-58	0.14	0.04	0.04	0.30	0.03	0.14	0.13	0.11	0.02	0.11	0.07	0	0	n.s.
IGHV1-67	0	0	0	0	0	0	0.01	0	0	0	0	0	0	n.s.
IGHV1-68	0	0	0	0	0	0	0	0	0.01	0	0	0	0	n.s.
IGHV1-69	11.80	4.73	1.55	4.72	4.93	6.50	5.88	3.91	11.20	4.48	4.67	4.27	7.53	n.s.
IGHV1-8	0.07	0	1.56	0.64	1.85	2.05	0.35	0.65	1.37	0	1.44	2.46	1.41	n.s.
IGHV1-c	0	0	0	0	0	0	0	0	0	0	0	0.02	0.16	n.s.
IGHV1-f	0.43	0.80	0	0	0	1.05	0.01	0	0.50	0	0	0.02	2.50	n.s.
IGHV2-10	0	0	0	0	0	0.01	0	0	0	0	0	0.20	0	n.s.
IGHV2-26	0.17	0.12	0.07	0.16	0.16	0.23	0.16	0.22	0.12	1.17	0.29	0.02	0.35	n.s.
IGHV2-5	2.58	1.97	4.11	1.89	2.15	2.35	3.34	3.31	1.60	3.50	1.39	1.42	0.10	n.s.
IGHV2-70	0.75	0.56	0.35	0.63	0.18	0.43	0.39	0.78	0.59	0.70	0.11	0.16	0.45	n.s.
IGHV3-11	1.11	1.43	1.98	1.99	1.39	2.23	2.37	2.06	2.16	0.15	0.11	1.23	2.25	n.s.
IGHV3-13	0.24	0.40	0.44	0.77	0.15	0.44	0.29	0.41	0.24	0.13	0.11	1.09	0	n.s.
IGHV3-15	2.15	3.32	1.86	1.62	1.87	1.79	3.43	1.50	1.44	1.40	1.42	2.86	2.24	n.s.
IGHV3-20	0.45	0.58	0.64	0.28	0.18	0.26	0.47	0.30	0.03	0	0.73	0.04	0.12	n.s.
IGHV3-21	1.54	3.48	3.59	2.61	4.49	2.33	2.50	2.92	2.56	3.28	2.33	2.79	2.12	n.s.
<b>IGHV3-23</b>	<b>6.92</b>	<b>11.05</b>	<b>11.78</b>	<b>10.89</b>	<b>9.14</b>	<b>9.98</b>	<b>7.59</b>	<b>5.37</b>	<b>5.70</b>	<b>4.06</b>	<b>9.18</b>	<b>10.30</b>	<b>5.15</b>	<b>0.04</b>
IGHV3-	12.21	13.51	4.18	10.14	9.45	9.65	8.66	12.97	11.33	15.72	8.10	14.13	9.89	n.s.
IGHV3-35	0.01	0	0	0	0	0	0	0	0	0	0	0	0	n.s.
IGHV3-41	0	0.01	0	0	0	0	0	0	0.05	0	0	0	0	n.s.
IGHV3-43	0.41	0.14	0.66	0.60	0.33	0.27	0.30	0.15	0.36	0.52	0.67	0.20	0.87	n.s.
IGHV3-47	0	0	0.02	0	0	0	0	0	0	0	0	0	0	n.s.
IGHV3-48	2.62	1.82	2.84	1.89	4.50	4.56	2.59	3.49	1.33	2.42	2.79	3.90	3.47	n.s.
IGHV3-49	1.14	1.11	1.10	0.74	0.44	1.93	0.98	1.03	0.45	0.92	1.62	1.16	1.02	n.s.
IGHV3-53/66	4.47	2.59	3.24	3.23	2.35	0.79	1.79	2.92	1.92	2.25	3.98	2.95	2.10	n.s.
IGHV3-64	0.50	1.53	0.54	0.58	0	0.38	2.23	0.84	0	0.40	0.49	0.27	1.06	n.s.
IGHV3-65	0	0	0.02	0	0	0	0	0	0	0	0.04	0	0	n.s.
IGHV3-7	2.68	4.63	4.15	5.71	6.42	6.06	6.18	3.78	7.89	1.72	4.56	7.74	5.00	n.s.
IGHV3-71	0	0.01	0	0	0	0	0	0.01	0	0	0	0	0	n.s.
IGHV3-72	1.10	0.84	0.66	0.34	0.81	0.69	1.76	0.77	0.06	0.03	0.55	1.03	2.24	n.s.
IGHV3-73	1.21	0.58	0.63	0.31	0.17	0.72	0.70	0.34	1.50	0.75	1.02	0.30	0.57	n.s.
<b>IGHV3-74</b>	<b>3.51</b>	<b>3.31</b>	<b>5.02</b>	<b>5.01</b>	<b>3.40</b>	<b>3.24</b>	<b>3.43</b>	<b>2.65</b>	<b>1.23</b>	<b>3.07</b>	<b>2.10</b>	<b>1.12</b>	<b>2.32</b>	<b>0.02</b>
IGHV3-9	0.08	0	1.80	2.29	4.69	2.87	2.88	2.06	6.24	0	0.42	3.67	0.52	n.s.
IGHV3-h	0.20	0.13	0.04	0.14	0.04	0.41	0.29	0.41	0.02	0.21	0.38	0.16	0.09	n.s.
IGHV4-28	0.02	0	0.07	0.05	0	0.01	0	0.01	0.01	0	0.07	0	0.09	n.s.
IGHV4-30-2	0.75	0.29	0.01	0	1.26	0.59	0.04	0.06	0.62	1.98	1.71	0.30	0.12	n.s.
IGHV4-30-	4.89	3.07	1.15	1.69	2.85	3.78	1.31	2.34	4.85	2.72	0.49	5.07	0.40	n.s.
IGHV4-34	4.34	2.71	5.45	5.78	2.94	2.06	2.43	6.33	3.25	3.60	7.81	3.59	3.19	n.s.
IGHV4-39	3.31	8.19	6.38	6.90	6.05	6.58	3.12	3.84	6.74	5.37	8.61	4.48	8.34	n.s.
IGHV4-4	1.43	2.18	1.85	2.09	3.05	1.75	1.73	2.26	0.83	2.00	1.24	1.71	1.35	n.s.
IGHV4-55	0	0	0	0	0	0.02	0	0.01	0.02	0.01	0.02	0	0.02	n.s.
IGHV4-59/61	6.01	5.97	7.04	6.79	8.65	4.96	6.21	8.33	5.80	10.14	10.96	3.22	5.92	n.s.
IGHV4-b	1.24		0.01	0.01		2.29	2.58	1.49	0.01	3.62	3.41	0.25	12.40	n.s.
IGHV5-51	3.67	3.17	2.90	2.55	2.56	2.47	1.93	3.48	2.22	2.79	1.46	1.73	3.31	n.s.
IGHV5-a	0.69	1.50	1.98	0.57	0	0	0.34	0.83	0	2.15	1.00	0	0	n.s.
IGHV6-1	1.02	0.97	1.77	1.53	0.53	0.79	0.51	0.47	0.99	1.56	1.88	4.34	1.35	n.s.
IGHV7-4-1	0.03	0	2.30	0.01	0	1.46	4.31	1.48	0	0	2.13	0	0	n.s.

Supplemental Table 5: Percent usage per IGHV germline segment in the PBMC of MS and OND patients. Comparison was made between MS and OND and p-values were calculated using a resampling-based permutation test. Only germline segments are represented which were found in at least one PBMC sample. IGHV3-23 and IGHV3-74 appear to be significantly overrepresented among PBMC IgG-VH in MS; n.s., not significant ( $p>0.05$ ).

## Supplemental Figures

### *Legends*

Supplemental Figure 1: Comparison of sequence counts shown in Table S1. Shown are numbers of IgG-VH sequences per disease group (MS or OND) after sequencing and initial quality control (“all reads”), numbers of sequences with identifiable IGHV, IHGJ, and CDR3 (“V, J, CDR3”), and numbers of non-redundant IgG-VH datasets (“non-red”; see Methods). Numbers of non-redundant CSF IgG-VH sequences were significantly higher in MS patients compared to OND (unpaired 2-tailed t-test).

Supplemental Figure 2: Representative multiple alignments of IgG-VH sequence clusters. Shown are examples from patients MS-1 (Figure 1, cluster N), and MS-5 (Figure 1, cluster M). Alignments of translated amino acid sequences corresponding to the indicated clusters; sequences are labeled by unique identifier, sequencing ID, and compartment (i.e. CSF or PBMC). Closest IGHV germline segments and average similarity to germline are: MS-1 IGHV3-23, 94%; MS-5, IGHV4-39 90%.

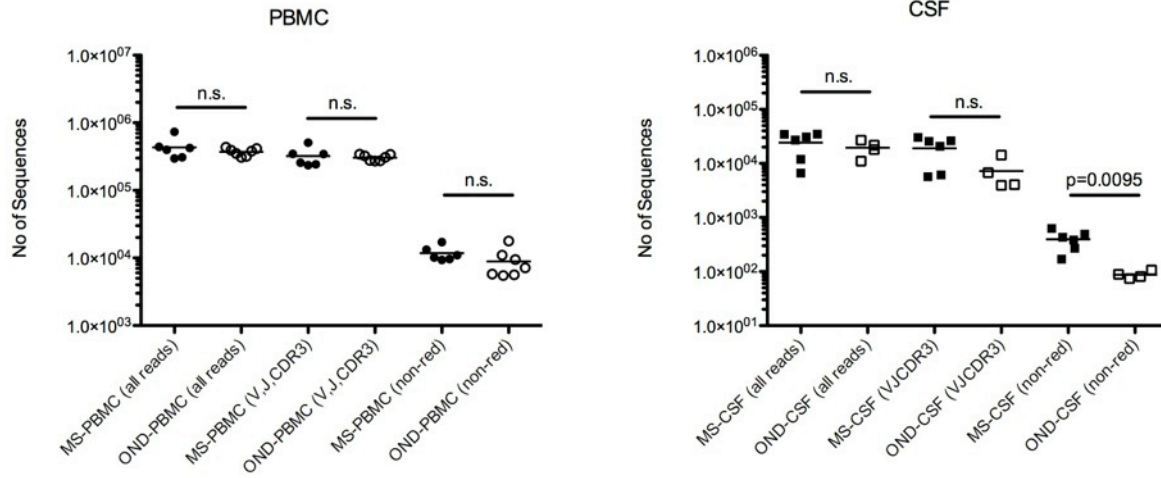
Supplemental Figure 3: Comparisons of IGHV usage profiles in PBMC of MS and OND patients. Only IGHV are shown that were represented at >0.5% in the respective PBMC sample; in PB IGHV3-23 and IGHV3-74 were over-represented ( $p=0.04$  and  $p=0.02$ , respectively) in MS patients compared with OND. Shown values are mean + SEM.

Supplemental Figure 4: Distribution of CDR3 length (A) and IGHV mutations (B) in MS and OND IgG-VH networks. IGHV mutations shown are nucleotide mutations. No significant differences could be observed between MS and OND (unpaired 2-tailed t-test).

Supplemental Figure 5: Strategy of clustering of related B cells based on a Levenshtein distance of 1 in the H-CDR3. Identical H-CDR3 are placed in individual nodes unless found in both compartments which is when they are placed in overlapping, differently colored nodes (blue=CSF, red=PB); H-CDR3 differing by a single aminoacid are directly connected by edges. Shown on the left is a resulting network and on the right the alignment of H-CDR3 sequences represented in this IgG-VH cluster.



Figures



Supplemental Figure 1

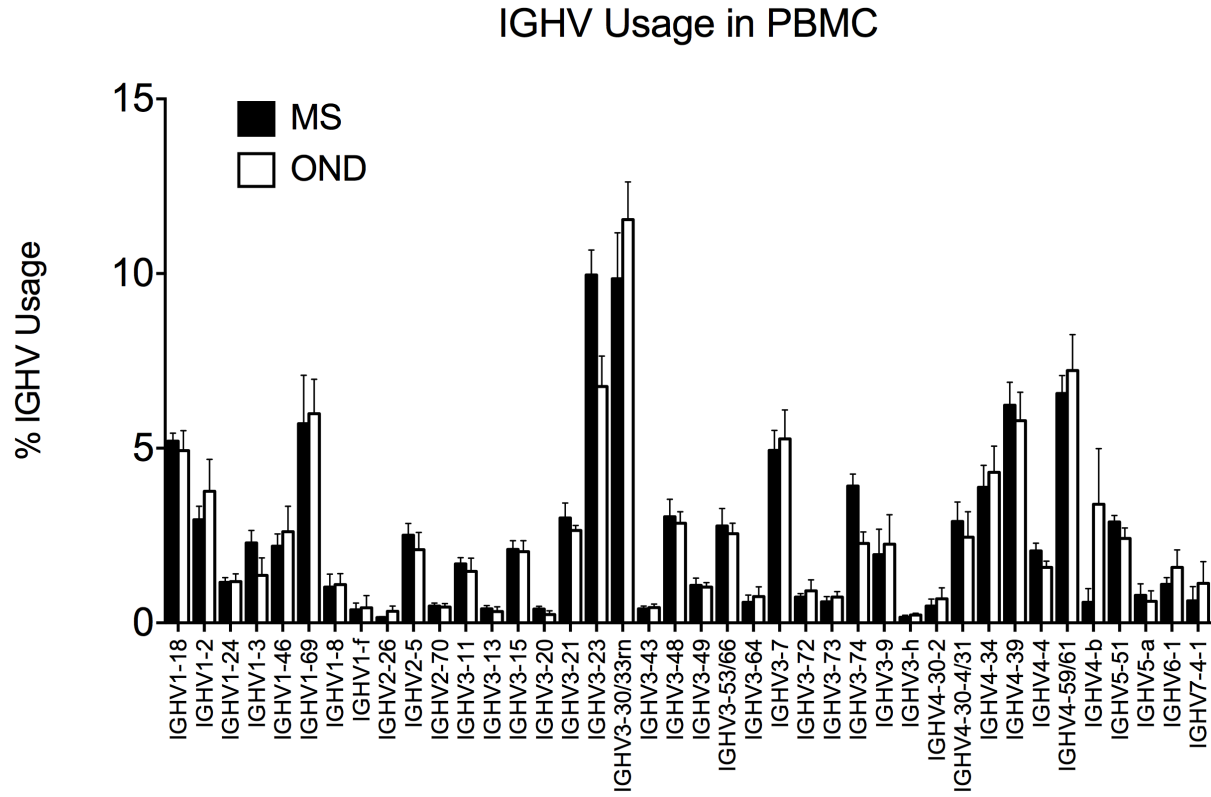
**MS-1**  
**IGHV3-23**

	<u>CDR1</u>	<u>CDR2</u>	<u>CDR3</u>
Consensus -	GSLRLSCAASGFTFSSYAMNWRQAPGKLEWVSTISGSGDSTYYADSVKGRLLTIARDNSKNTLYLQMNSLRADDTAVVYCAKAGGYSGSSGRRNFYSFDSWGQG		
GYUR81E02GJ1N8_1710_CSF			GPVRLSCAASGFTFSSYAMNWRQAPGKLEWVSTISGSGDSTYYADSVKGRLLTIARDNSKNTLYLQMNSLRADDTAVVYCAKAGGYSGSSGRRNFYSFDSWGQG
GYUR81E01D6KYU_1710_CSF			GPVRLSCAASGFTFSSYAMNWRQAPGKLEWVSTISGSGDSTYYADSVKGRLLTIARDNSKNTLYLQMNSLRADDTAVVYCAKAGGYSGSSGRRNFYSFDSWGQG
GYUR81E01ICRFYV_1710_CSF			GSLRLSCAASGFTFSSYAMNWRQAPGKLEWVSTISGSGDSTYYADSVKGRLLTIARDNSKNTLYLQMNSLRADDTAVVYCAKAGGYSGSSGRRNFYSFDSWGQG
GYUR81E02F2XU8_1710_CSF			GSLRLSCAASGFTFSSYAMNWRQAPGKLEWVSTISGSGDSTYYADSVKGRLLTIARDNSKNTLYLQMNSLRADDTAVVYCAKAGGYSGSSGRRNFYSFDSWGQG
GYUR81E02JNEG0_1710_CSF			GSLRLSCAASGFTFSSYAMNWRQAPGKLEWVSTISGSGDSTYYADSVKGRLLTIARDNSKNTLYLQMNSLRADDTAVVYCAKAGGYSGSSGRRNFYSFDSWGQG
GYUR81E01DYQJ6_1710_CSF			GSLRLSCAASGFTFSSYAMNWRQAPGKLEWVSTISGSGDSTYYADSVKGRLLTIARDNSKNTLYLQMNSLRADDTAVVYCAKAGGYSGSSGRRNFYSFDSWGQG
GYUR81E02GV21B_1710_CSF			GSLRLSCAASGFTFSSYAMNWRQAPGKLEWVSTISGSGDSTYYADSVKGRLLTIARDNSKNTLYLQMNSLRADDTAVVYCAKAGGYSGSSGRRNFYSFDSWGQG
GYUR81E01BIH4Y_1710_CSF			GSLRLSCAASGFTFSSYAMNWRQAPGKLEWVSTISGSGDSTYYADSVKGRLLTIARDNSKNTLYLQMNSLRADDTAVVYCAKAGGYSGSSGRRNFYSFDSWGQG
GYUR81E01ETVEL_1710_PBM			GSLRLSCAASGFTFSSYAMNWRQAPGKLEWVSTISGSGDSTYYADSVKGRLLTIARDNSKNTLYLQMNSLRADDTAVVYCAKAGGYSGSSGRRNFYSFDSWGQG
GYUR81E01EIVRZ_1710_CSF			GSLRLSCAASGFTFSSYAMNWRQAPGKLEWVSTISGSGDSTYYADSVKGRLLTIARDNSKNTLYLQMNSLRADDTAVVYCAKAGGYSGSSGRRNFYSFDSWGQG
GYUR81E01EGYFW_1710_CSF			GSLRLSCAASGFTFSSYAMNWRQAPGKLEWVSTISGSGDSTYYADSVKGRLLTIARDNSKNTLYLQMNSLRADDTAVVYCAKAGGYSGSSGRRNFYSFDSWGQG
GYUR81E02H5U04_1710_CSF			GSLRLSCAASGFTFSSYAMNWRQAPGKLEWVSTISGSGDSTYYADSVKGRLLTIARDNSKNTLYLQMNSLRADDTAVVYCAKAGGYSGSSGRRNFYSFDSWGQG
GYUR81E01BKKUT_1710_PBM			GSLRLSCAASGFTFSSYAMNWRQAPGKLEWVSTISGSGDSTYYADSVKGRLLTIARDNSKNTLYLQMNSLRADDTAVVYCAKAGGYSGSSGRRNFYSFDSWGQG
GYUR81E01AVVRS_1710_CSF			GSLRLSCAASGFTFSSYAMNWRQAPGKLEWVSTISGSGDSTYYADSVKGRLLTIARDNSKNTLYLQMNSLRADDTAVVYCAKAGGYSGSSGRRNFYSFDSWGQG
GYUR81E01EQSLO_1710_PBM			GSLRLSCAASGFTFSSYAMNWRQAPGKLEWVSTISGSGDSTYYADSVKGRLLTIARDNSKNTLYLQMNSLRADDTAVVYCAKAGGYSGSSGRRNFYSFDSWGQG
GYUR81E01AVB20_1710_CSF			GSLRLSCAASGFTFSSYAMNWRQAPGKLEWVSTISGSGDSTYYADSVKGRLLTIARDNSKNTLYLQMNSLRADDTAVVYCAKAGGYSGSSGRRNFYSFDSWGQG
GYUR81E02J25JE_1710_CSF			GSLRLSCAASGFTFSSYAMNWRQAPGKLEWVSTISGSGDSTYYADSVKGRLLTIARDNSKNTLYLQMNSLRADDTAVVYCAKAGGYSGSSGRRNFYSFDSWGQG
GYUR81E01BIN04_1710_CSF			GSLRLSCAASGFTFSSYAMNWRQAPGKLEWVSTISGSGDSTYYADSVKGRLLTIARDNSKNTLYLQMNSLRADDTAVVYCAKAGGYSGSSGRRNFYSFDSWGQG
GYUR81E01BFANW_1710_CSF			GSLRPSCAASGFTFSSYAMNWRQAPGKLEWVSTISGSGDSTYYADSVKGRLLTIARDNSKNTLYLQMNSLRADDTAVVYCAKAGGYSGSSGRRNFYSFDSWGQG

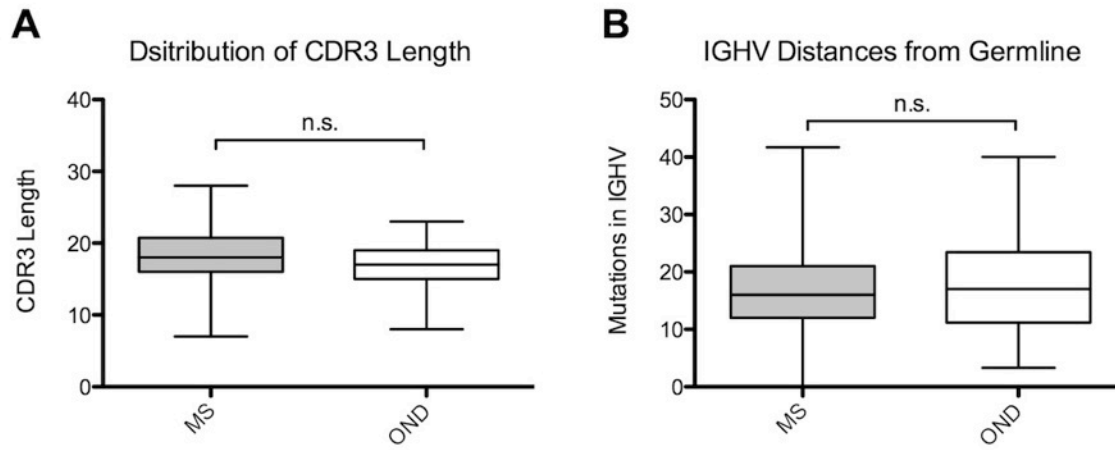
**MS-5**  
**IGHV4-39**

	<u>CDR1</u>	<u>CDR2</u>	<u>CDR3</u>
Consensus -	ETLSLTCNVSGGSIITSHYWGIRPPGKLEWIGTVVYSGSTFFYNPILKSRVTISVDTSKNHFSRLRSTVTAADAAYVYCAREGPMVGVTRTFDCWGQG		
G57NQVN02FDL7N_3811_CSF			ETPPLTCNASGGSIITSHYWGIRPPGKLEWIGTVVYSGSTFFYNPILKSRVTISVDTSKNHFSRLRSTVTAADAAYVYCAREGPMVGVTRTFDCWGQG
G57NQVN02GG6QT_3811_CSF			ETLSLTCNVSGGSIITSHYWGIRPPGKLEWIGTVVYSGSTFFYNPILKSRVTISVDTSKNHFSRLRSTVTAADAAYVYCAREGPMVGVTRTFDCWGQG
G57NQVN02HRU7S_3811_CSF			ETLSLTCNVSGGSIITSHYWGIRPPGKLEWIGTVVYSGSTFFYNPILKSRVTISVDTSKNHFSRLRSTVTAADAAYVYCAREGPMVGVTRTFDCWGQG
G50A0C102HYEWX_3811_PBM			ETLSLTCNVSGGSIITSHYWGIRPPGKLEWIGTVVYSGSTFFYNPILKSRVTISVDTSKNHFSRLRSTVTAADAAYVYCAREGPMVGVTRTFDCWGQG
G57NQVN02HKV95_3811_CSF			ETLSLTCNVSGGSIITSHYWGIRPPGKLEWIGTVVYSGSTFFYNPILKSRVTISVDTSKNHFSRLRSTVTAADAAYVYCAREGPMVGVTRTFDCWGQG
G57NQVN02FJLID_3811_CSF			ETLSLTCNVSGGSIITSHYWGIRPPGKLEWIGTVVYSGSTFFYNPILKSRVTISVDTSKNHFSRLRSTVTAADAAYVYCAREGPMVGVTRTFDCWGQG
G57NQVN02G01TB_3811_CSF			ETLSLTCNVSGGSIITSHYWGIRPPGKLEWIGTVVYSGSTFFYNPILKSRVTISVDTSKNHFSRLRSTVTAADAAYVYCAREGPMVGVTRTFDCWGQG
G57NQVN02HQRNS_3811_CSF			ETLSLTCNVSGGSIITSHYWGIRPPGKLEWIGTVVYSGSTFFYNPILKSRVTISVDTSKNHFSRLRSTVTAADAAYVYCAREGPMVGVTRTFDCWGQG
G57NQVN02HXITC_3811_CSF			ETLSLTCNVSGGSIITSHYWGIRPPGKLEWIGTVVYSGSTFFYNPILKSRVTISVDTSKNHFSRLRSTVTAADAAYVYCAREGPMVGVTRTFDCWGQG
G57NQVN02JQ02P_3811_CSF			ETLSLTCNVSGGSIITSHYWGIRPPGKLEWIGTVVYSGSTFFYNPILKSRVTISVDTSKNHFSRLRSTVTAADAAYVYCAREGPMVGVTRTFDCWGQG
G57NQVN02HA9S6_3811_CSF			ETLSLTCNVSGGSIITSHYWGIRPPGKLEWIGTVVYSGSTFFYNPILKSRVTISVDTSKNHFSRLRSTVTAADAAYVYCAREGPMVGVTRTFDCWGQG
G57NQVN02JRNCG_3811_CSF			ETLSLTCNVSGGSIITSHYWGIRPPGKLEWIGTVVYSGSTFFYNPILKSRVTISVDTSKNHFSRLRSTVTAADAAYVYCAREGPMVGVTRTFDCWGQG
G57NQVN02H2ANC_3811_CSF			ETLSLTCNVSGGSIITSHYWGIRPPGKLEWIGTVVYSGSTFFYNPILKSRVTISVDTSKNHFSRLRSTVTAADAAYVYCAREGPMVGVTRTFDCWGQG
G57NQVN02FNQP3_3811_CSF			ETLSLTCNVSGGSIITSHYWGIRPPGKLEWIGTVVYSGSTFFYNPILKSRVTISVDTSKNHFSRLRSTVTAADAAYVYCAREGPMVGVTRTFDCWGQG
G57NQVN02I8G0S_3811_CSF			ETLSLTCNVSGGSIITSHYWGIRPPGKLEWIGTVVYSGSTFFYNPILKSRVTISVDTSKNHFSRLRSTVTAADAAYVYCAREGPMVGVTRTFDCWGQG
G57NQVN02IDWEC_3811_CSF			ETLSLTCNVSGGSIITSHYWGIRPPGKLEWIGTVVYSGSTFFYNPILKSRVTISVDTSKNHFSRLRSTVTAADAAYVYCAREGPMVGVTRTFDCWGQG
G57NQVN02G39MV_3811_CSF			ETLSLTCNVSGGSIITSHYWGIRPPGKLEWIGTVVYSGSTFFYNPILKSRVTISVDTSKNHFSRLRSTVTAADAAYVYCAREGPMVGVTRTFDCWGQG
G57NQVN02H4KF3_3811_CSF			ETLSLTCNVSGGSIITSHYWGIRPPGKLEWIGTVVYSGSTFFYNPILKSRVTISVDTSKNHFSRLRSTVTAADAAYVYCAREGPMVGVTRTFDCWGQG
G57NQVN02IY9XR_3811_CSF			ETLSLTCNVSGGSIITSHYWGIRPPGKLEWIGTVVYSGSTFFYNPILKSRVTISVDTSKNHFSRLRSTVTAADAAYVYCAREGPMVGVTRTFDCWGQG
G57NQVN02H2FUP_3811_CSF			ETLSLTCNVSGGSIITSHYWGIRPPGKLEWIGTVVYSGSTFFYNPILKSRVAISVDTSKNHFSRLRSTVTAADAAYVYCAREGPMVGVTRTFDCWGQG

Supplemental Figure 2



Supplemental Figure 3



Supplemental Figure 4

