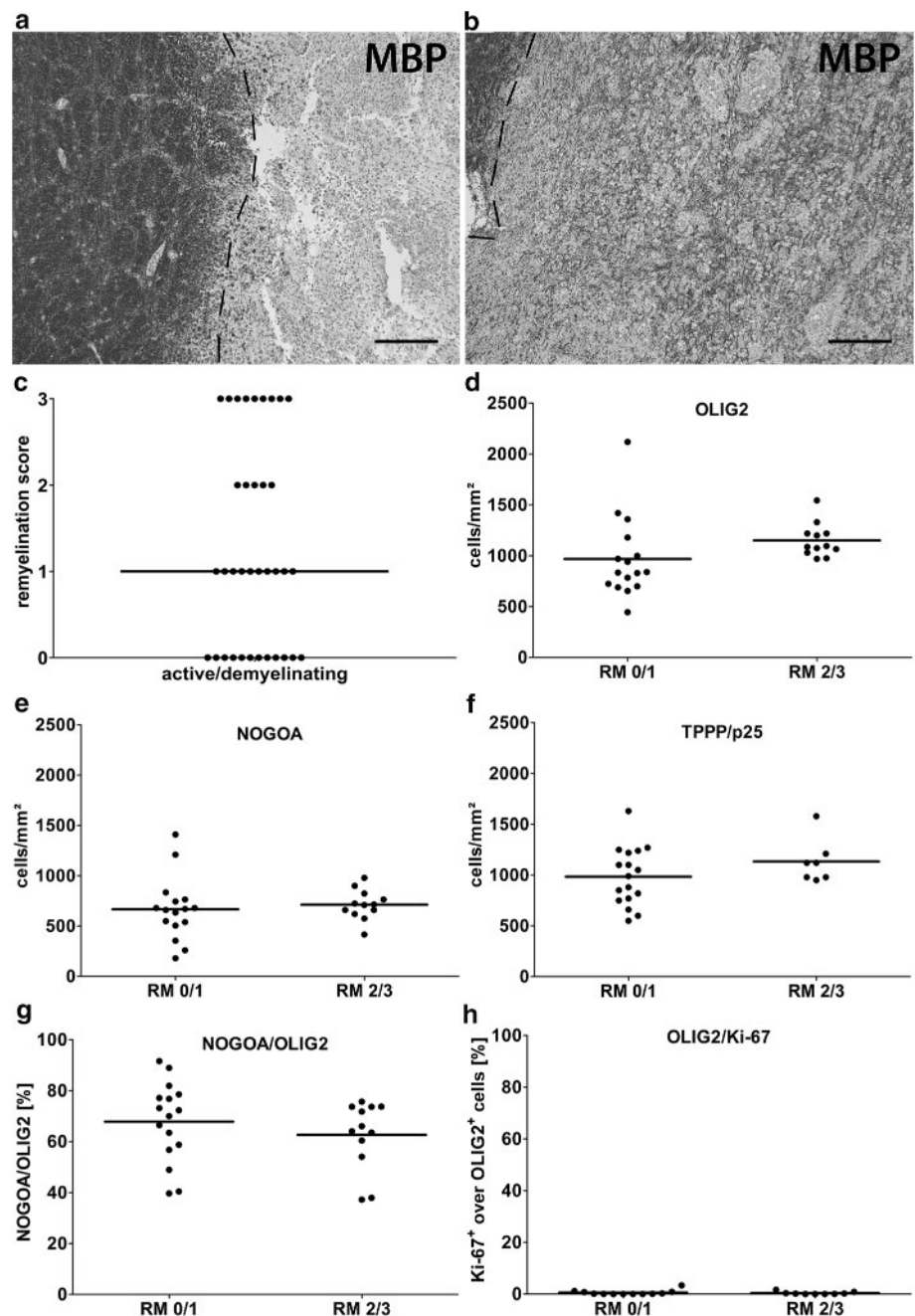


Fig. 3 No differences in oligodendroglial cell numbers in active/demyelinating lesions with and without marked remyelination. **a, b** A lesion with no remyelination (score 0) is shown in **a**, whereas in **b**, a lesion with marked remyelination (score 3) is displayed (IHC for MBP). **c** In 14 out of 36 (= 39%) of the active/demyelinating lesions, we observed marked remyelination (score 2/3). **d–f** Quantification of OLIG2⁺, NOGOA⁺ and TPPP/p25⁺ cells in active/demyelinating lesions with (score 2/3) and without (score 0/1) marked remyelination. **g** No significant difference in the ratio of NOGOA⁺ cells over OLIG2⁺ in lesions with and without marked remyelination. **h** Double IHC for Ki-67 and OLIG2 demonstrated that only few OLIG2⁺ cells express Ki-67. Scale bars in **a** and **b**: 200 μ m. MBP myelin basic protein, OLIG2 oligodendrocyte transcription factor 2, TPPP/p25 tubulin polymerization promoting protein, active/demyelinating active/demyelinating lesions, RM 0/1 remyelination score 0 or 1, RM 2/3 remyelination score 2 or 3



of less than 50% of the lesion area (score 1). The difference in the extent of remyelination between mixed and active as well as inactive lesions was highly significant ($p < 0.001$) (Fig. 4c).

Higher percentages of TMEM119⁺ and iNOS⁺ myeloid cells are associated with less remyelination in mixed lesions

To investigate whether the inflammatory milieu may influence the outcome of remyelination, we examined the composition of inflammatory infiltrates in active, mixed, and

inactive lesions (Fig. 5a–f). Myeloid cells were the dominating inflammatory cell population in all lesion types; however, the numbers of myeloid cells in inactive lesions as well as in the center of mixed lesions were significantly lower compared to active and the rim of mixed lesions (Fig. 5d). Highest numbers of T and B cells were found in active lesions (Fig. 5e, f).

Due to the dominance of the myeloid cell population and based on a plethora of literature describing either detrimental or beneficial effects of myeloid cells on remyelination, we hypothesized that those myeloid cells may determine the outcome of remyelination. We focused our subsequent analyses on

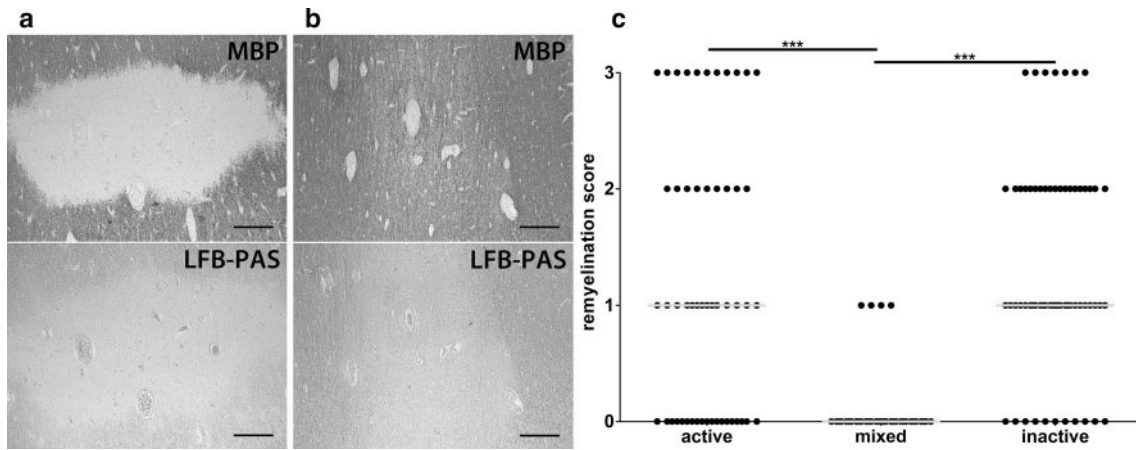


Fig. 4 Almost complete lack of remyelination in mixed lesions. **a**, **b** Pictures display an inactive lesion with no remyelination (score 0) (**a**), whereas **b** shows a completely remyelinated shadow plaque (score 3). Upper panels in **a** and **b** show IHC for MBP, lower panels in **a** and **b** display LFB-PAS staining. **c** Semiquantitative analysis

of remyelination reveals an almost complete lack of remyelination in mixed lesions (**c**). Scale bars in **a** and **b**: 500 μ m. *MBP* myelin basic protein, *LFB-PAS* luxol fast blue–periodic acid Schiff; *active* active lesions, *mixed* mixed active/inactive lesions, *inactive* inactive lesions

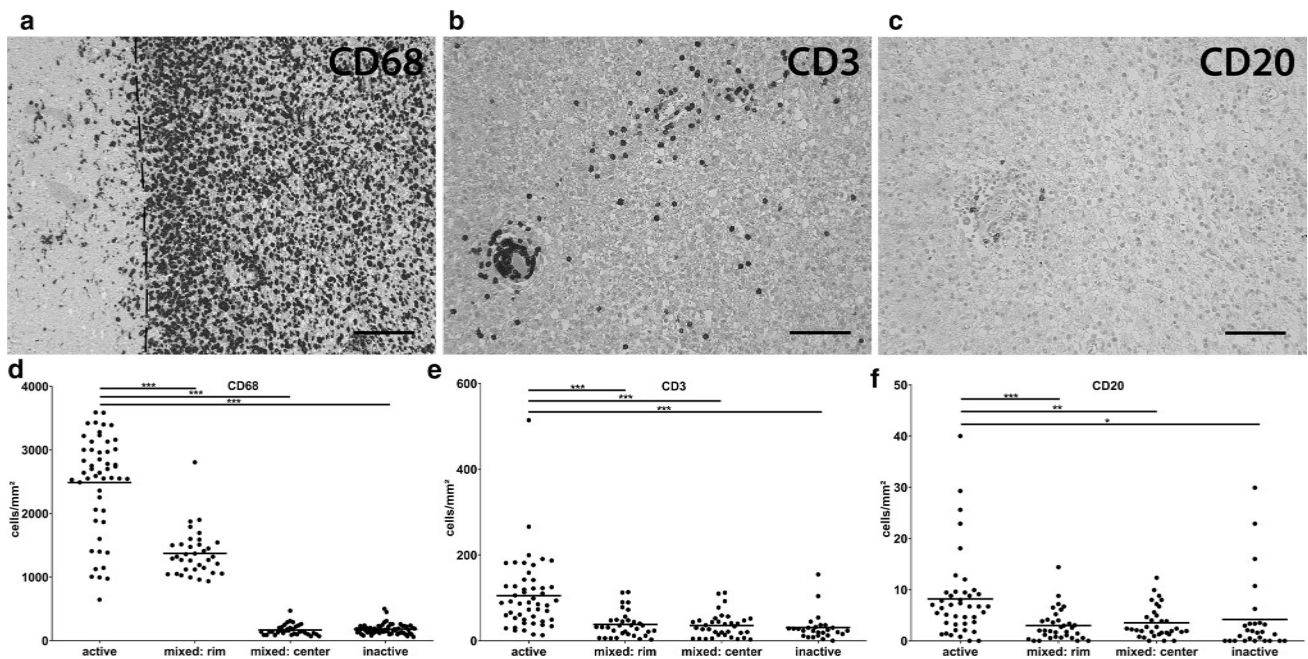


Fig. 5 Inflammatory environment in different lesion types. **a–c** IHC demonstrates numerous CD68⁺ myeloid cells, fewer CD3⁺ T cells and few CD20⁺ B cells in an active lesion. All pictures are taken from the same active lesion. **d–f** Quantification of CD68⁺, CD3⁺ and CD20⁺ cells in different lesion types. Please note the differ-

ent y-axes. Scale bar in **a**: 200 μ m, scale bars in **b** and **c**: 100 μ m. *CD3/20/68* cluster of differentiation 3/20/68, *active* active lesions, *mixed: rim* rim of mixed active/inactive lesions, *mixed: center* center of mixed active/inactive lesions, *inactive* inactive lesions

active lesions and the rim of mixed lesions, since the center of mixed lesions as well as inactive lesions was almost completely depleted of myeloid cells. To characterize the myeloid cell population in more detail, we stained for TMEM119, a marker of homeostatic microglia, iNOS, a pro-inflammatory marker as well as CD163 and CD206, two markers associated with

an anti-inflammatory phenotype of myeloid cells (Fig. 6a–d). The highest absolute number of myeloid cells was identified by CD163 (mean \pm SEM: 906 \pm 70) followed by TMEM119 (mean \pm SEM: 640 \pm 36) and iNOS (mean \pm SEM: 468 \pm 38), whereas fewer myeloid cells expressed CD206 (mean \pm SEM: 128 \pm 12) (Supplementary Fig. 2a–e, online resource). When

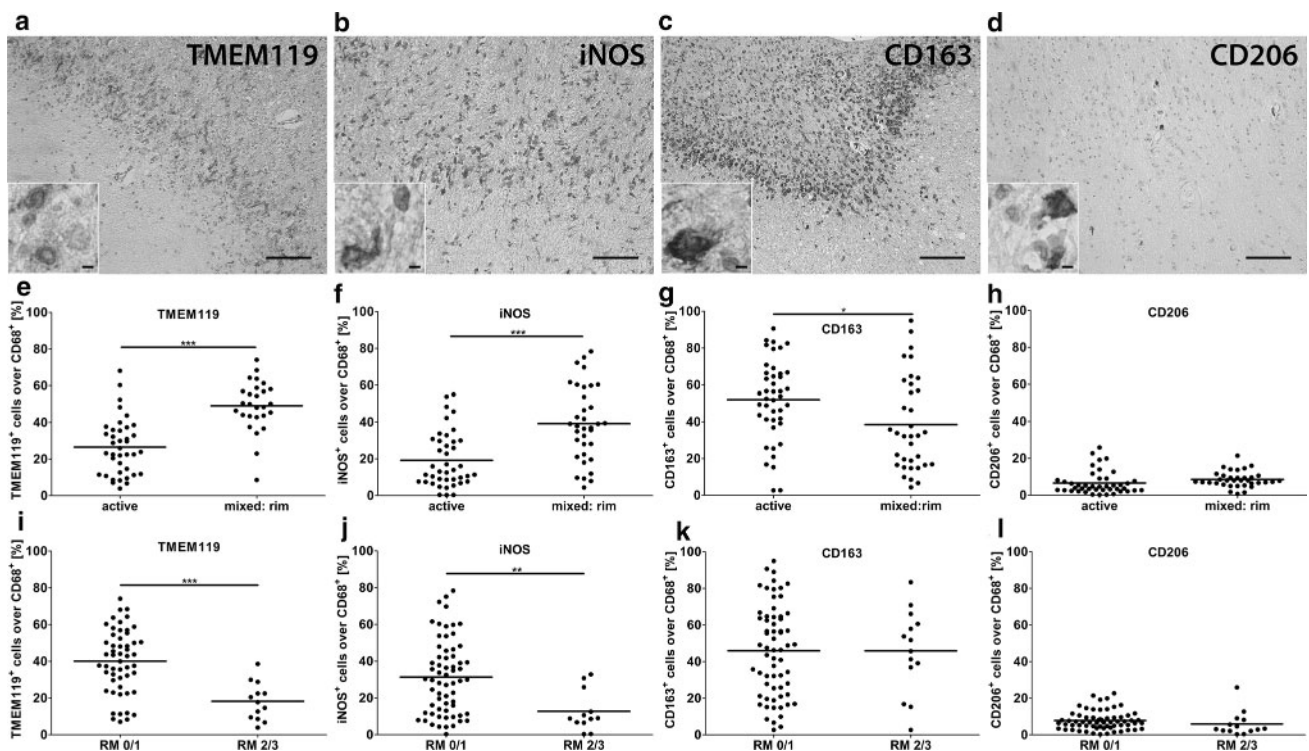


Fig. 6 Relative increase in the numbers of TMEM119⁺ and iNOS⁺ myeloid cells in the rim of mixed lesions. **a–d** IHC for TMEM119, iNOS, CD163 and CD206 was performed. Inserts show labelled cells in higher magnification. **e–g** Quantification demonstrated a significant relative increase in the ratio of TMEM119⁺/CD68⁺ and iNOS⁺/CD68⁺ myeloid cells and a relative decrease of CD163⁺/CD68⁺ myeloid cells in the rim of mixed lesions compared to active lesions. **h** No differences in the number of CD206⁺ over CD68⁺ cells were observed between active lesions and the rim of mixed lesions. **i, j** Comparably, we observed a relative increase in the ratio

of TMEM119⁺/CD68⁺ and iNOS⁺/CD68⁺ in lesions with limited (score 0/1) compared to lesions with marked remyelination (score 2/3). **k, l** No differences were found between lesions with and without marked remyelination with respect to the ratio of CD163⁺/CD68⁺ and CD206⁺/CD68⁺ cells. Scale bars in **a, b** and **d**: 100 μ m, scale bar in **c**: 200 μ m, scale bar in insert in **a–d**: 6.25 μ m. TMEM119 transmembrane protein 119, iNOS inducible nitric oxide synthase, CD68/163/206 cluster of differentiation 68/163/206, active active lesions, mixed: rim rim of mixed active/inactive lesions, RM 0/1 remyelination score 0 or 1, RM 2/3 remyelination score 2 or 3

analyzing the ratio of TMEM119⁺/CD68⁺, iNOS⁺/CD68⁺ and CD163⁺/CD68⁺ cells, we saw a significant increase in the percentage of TMEM119⁺ and iNOS⁺ over CD68⁺ myeloid cells in the rim of mixed lesions compared to active lesions (Fig. 6e, f). The percentage of CD163⁺ cells was significantly reduced in the rim of mixed lesions, whereas no difference in the number of CD206⁺ myeloid cells was observed (Fig. 6g, h). Furthermore, when comparing the ratio of TMEM119⁺, iNOS⁺, CD163⁺ and CD206⁺ over CD68⁺ cells in active and mixed lesions with limited (score 0/1) or marked remyelination (score 2/3), we observed a relative increase in TMEM119⁺ and iNOS⁺ myeloid cells in lesions with limited remyelination.

In summary, these data demonstrate a relative increase of TMEM119⁺ homeostatic microglia and iNOS⁺ pro-inflammatory myeloid cells in the rim of mixed lesions. Furthermore, a relative increase in TMEM119⁺ and iNOS⁺ myeloid cells is associated with reduced remyelination in active and mixed lesions.

Supernatants of M1, but not M0 or M2 polarized microglia inhibit terminal differentiation of hiOL

To further elucidate the impact of pro-inflammatory myeloid cells on oligodendroglial differentiation, we investigated the role of supernatants of activated microglia on hiOL [22]. Fetal and adult human primary microglia were stimulated for 48 h with either LPS and IFN γ to promote an M1 phenotype or with IL-4 and IL-13 to induce an M2 phenotype. In total, we obtained microglia supernatants from three fetal (HFM1-3) and two adult (HAM1, 2) donors. For validation of successful polarization, we determined the expression of typical M1 (*IL-6*, *CXCL10*, *TNF α*) and M2 cytokines (*CD206* and *CD209*) by qRT-PCR (Fig. 7a–e). Supernatants as well as the appropriate medium without the supernatants (controls) were added to differentiating hiOL from three different donors in a dilution of 1:10 from day 4 to 21 of differentiation to examine the effect on early differentiation. No differences in the percentage of O4⁺ cells were observed when comparing appropriate medium controls and

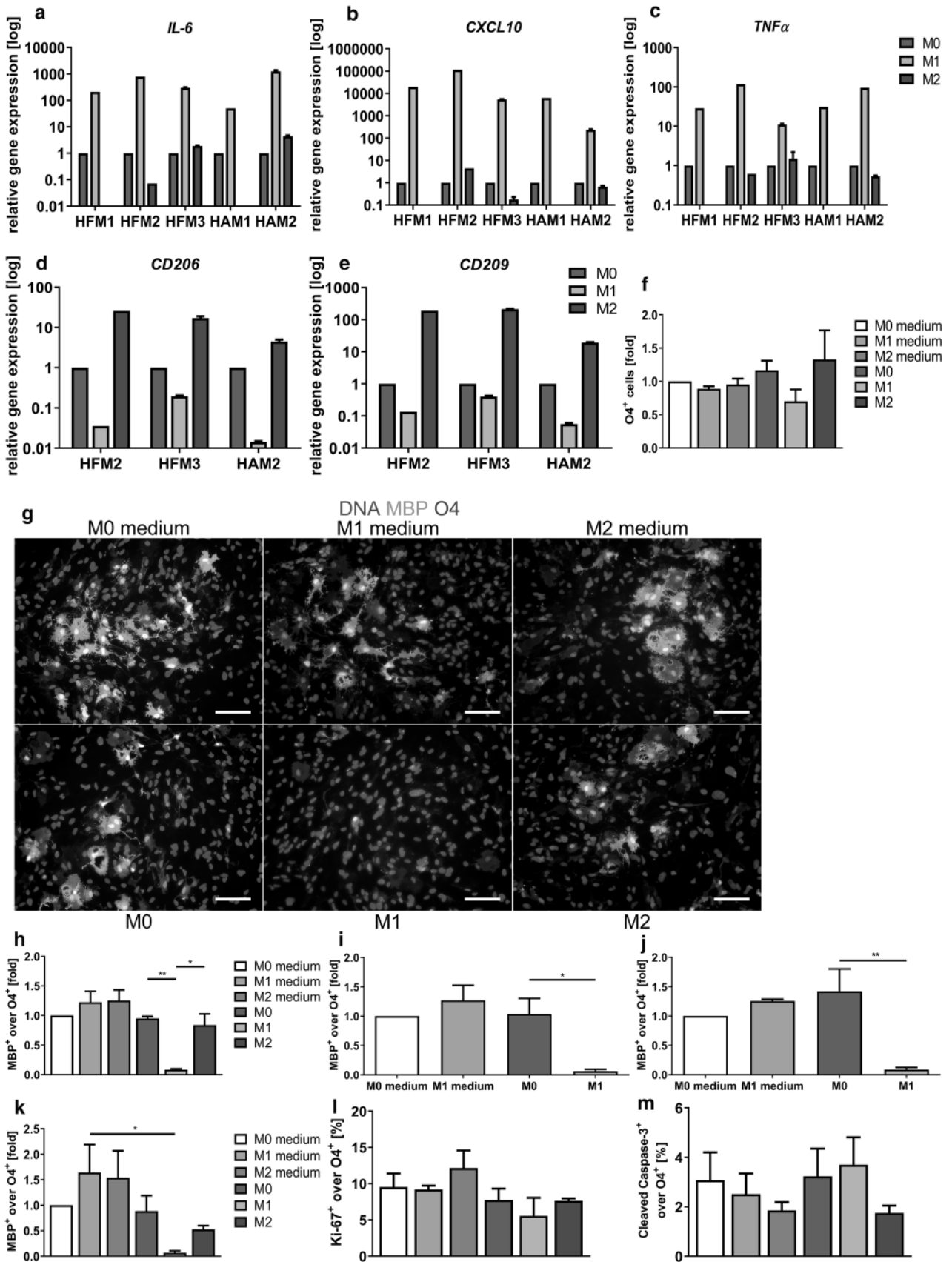


Fig. 7 Supernatants of primary human M1, but not M2 or M0 polarized microglia inhibit the terminal differentiation of hiOL. Every experiment analyzing the effect of microglia supernatants on hiOL was performed with hiOL derived from three different donors. **a–e** To confirm successful polarization into M1 and M2 microglia, qRT-PCR was performed for *IL-6*, *CXCL10* and *TNF α* (M1) and *CD206* and *CD209* (M2). **f** Differentiation of hiOL in the presence of supernatants from M0, M1 or M2 polarized microglia from one fetal donor (HFM2) from day 4 to 21 resulted in no significant differences in the percentage of O4⁺ hiOL when comparing the effect of different supernatants with appropriate controls. **g, h** Culturing O4 sorted hiOL in the presence of supernatants from M1, but not M0 or M2 polarized microglia from one fetal donor (HFM3) from day 21 to 35 impaired significantly the differentiation of O4⁺ hiOL into MBP⁺ mature oligodendrocytes. **i–k** This was confirmed using supernatants from M1 and M0 polarized microglia from another fetal (HFM1) and one adult donor (HAM1) as well as supernatants from M0, M1 and M2 polarized microglia from a second adult donor (HAM2). **l** Analysis of percentages of actively dividing hiOL (Ki-67⁺ over O4⁺ cells) in the presence of supernatants from M0, M1 or M2 polarized microglia from one fetal donor (HFM2) from day 21 to 35 of differentiation did not reveal any significant differences. **m** Percentages of apoptotic hiOL (cleaved caspase 3⁺ over O4⁺ cells) in the presence of supernatants from M0, M1 or M2 polarized microglia from one fetal donor (HFM3) from day 21 to 35 of differentiation were not significantly altered. Scale bar in **g**: 100 μ m. *hiOL* human iPSC-derived oligodendrocytes, *HFM* human fetal microglia, *HAM* human adult microglia, *IL-6* interleukin 6, *CXCL10* C-X-C motif chemokine 10, *TNF α* tumor necrosis factor alpha, *CD206/CD209* cluster of differentiation 206/209, *MBP* myelin basic protein

supernatants (Fig. 7f) as assessed by flow cytometry for the oligodendroglial marker O4 which is expressed by immature and mature oligodendrocytes. To determine the effect of the supernatants on terminal differentiation, untreated O4⁺ hiOL sorted at day 21 of differentiation were exposed to M0, M1, and M2 microglia supernatants or medium controls for 14 days (dilution 1:10). ICC for MBP revealed significantly reduced numbers of mature MBP⁺ hiOL after application of M1 microglia supernatants but not M0 or M2 microglia supernatants from a fetal donor, while O4⁺ hiOL could still be detected (Fig. 7g, h). These results were validated using another fetal as well as two adult donors (Fig. 7i–k). To elucidate whether the reduced presence of MBP⁺ cells is caused by a decreased differentiation into mature hiOL or due to changes in cell death or proliferation, we analyzed the numbers of apoptotic and actively dividing hiOL by double ICC for cleaved caspase 3, respectively, Ki-67 and O4. Neither the percentage of apoptotic nor the percentage of actively dividing hiOL was significantly altered after application of microglia supernatants (Fig. 7l, m). Combined, these data demonstrate that supernatants from M1 polarized human microglia significantly inhibit the terminal differentiation of hiOL into mature MBP⁺ oligodendrocytes but do not affect apoptosis or proliferation of hiOL.

Discussion

Promotion of remyelination represents a new and attractive treatment target in MS. In demyelinating animal models, the proliferation and differentiation of OPC are prerequisites for successful remyelination. In progressive disease stages, MS lesions display a relative preservation of OPC, but lack mature oligodendrocytes suggesting that a differentiation block contributes to remyelination failure in MS. This notion has been recently challenged by suggesting that not OPC, but mature oligodendrocytes are the major remyelinating cells in MS [34, 54, 71]. To further elucidate the mechanisms contributing to impaired remyelination in MS, we performed a detailed analysis of oligodendrocytes and remyelination in different MS lesion types. Our data suggest that different mechanisms contribute to impaired remyelination depending on lesion stage. In active/demyelinating lesions, remyelination occurs only in a subset of lesions despite the presence of high numbers of mature oligodendrocytes in all lesions indicating that an impaired formation of new myelin sheaths rather than a lack of mature oligodendrocytes contributes to insufficient remyelination in this lesion type. Furthermore, in mixed lesions, we observed an almost complete lack of remyelination and a relative increase of TMEM119⁺ microglia cells and iNOS⁺ pro-inflammatory myeloid cells. Supernatants of M1, but not M2 or M0 polarized microglia prevented terminal differentiation of hiOL indicating that pro-inflammatory microglia may contribute to impaired remyelination in mixed lesions.

The generally accepted view that OPC are prerequisite for successful remyelination is, among others, based on studies demonstrating that rodents in which proliferating oligodendrocytes were ablated by X-irradiation did not remyelinate [36]. Additionally, mature human oligodendrocytes transplanted into demyelinated and irradiated spinal cord lesions of rats did not contribute to remyelination [64]. One could speculate that X-irradiation may result in significant damage to the exposed tissue and other factors than the supposed incapability of mature oligodendrocytes may contribute to remyelination failure in these studies. However, in an elegant and more recent study, Crawford and colleagues found no evidence that pre-existing mature oligodendrocytes contribute to remyelination after toxin-induced demyelination using inducible reporter mouse lines and fate mapping [14]. Furthermore, a broad range of studies demonstrated that increased differentiation of OPC into mature myelinating oligodendrocytes results in accelerated remyelination in experimental animal models (for review see [25, 62]). Earlier histological studies using MS tissue samples demonstrated the presence of OPC and an almost complete absence of mature oligodendrocytes in mixed and inactive lesions [11, 39, 68, 70]. These findings together with the results from

animal studies were interpreted as evidence for an impaired oligodendroglial differentiation as contributing factor for remyelination failure in MS. This view has been challenged by experimental studies reporting mature oligodendrocytes being connected to mature and remyelinated myelin sheaths in a cat as well as in a non-human primate model of demyelination and remyelination [19]. Additionally, the results of a recent study measuring the integration of ^{14}C derived from nuclear testing into DNA of oligodendroglial lineage cells suggest that pre-existing oligodendrocytes and not proliferating OPC contribute to remyelination in MS [71]. Our data demonstrate the presence of remyelination and preservation of oligodendrocytes in highly inflammatory active/demyelinating lesions suggesting that oligodendrocytes are not the primary target of the immune attack in MS lesions. Furthermore, despite the presence of high numbers of mature oligodendrocytes, only a subset of lesions showed signs of remyelination. This indicates that not a differentiation block, but rather an impaired formation of new myelin sheaths contributes to remyelination failure in these lesions. Additionally, it raises the question which cells contributed to successful remyelination in active/demyelinating lesions with marked remyelination. We observed only a very low percentage of proliferating OPC in these lesions suggesting that these cells may not be major contributors to remyelination. Considering further the fact that OPC account for approximately 5% of the total oligodendroglial population and the relatively long time period required for differentiation of human OPC into mature oligodendrocytes [7, 12, 39], it appears unlikely that only recently differentiated OPC contributed to successful remyelination in active/demyelinating lesions, but instead favors a scenario in which also mature oligodendrocytes surviving the initial inflammatory attack contribute to remyelination. However, one has to keep in mind that Ki-67 only identifies cells which are actively dividing. The discrepancy between this scenario and experimental rodent studies might be explained by the fact that EAE, lysolecithine or cuprizone induced demyelination are characterized by a marked loss of mature oligodendrocytes [40, 49, 50] which is in contrast to our observations in active/demyelinating MS lesions. However, histology provides only a snapshot, and as long as no reliable markers for recently differentiated oligodendrocytes and remyelination are available, we will not be able to definitively clarify which cell population contributes to remyelination in MS. Importantly, our data also do not exclude the possibility that in other lesion types than active/demyelinating lesions, an oligodendroglial differentiation block contributes to impaired remyelination. Furthermore, remyelination is a complex process. It not only depends on the presence and absence of oligodendrocytes or its progenitors, but also requires axons susceptible for remyelination. Several studies for example have demonstrated that neuronal activity stimulates remyelination [3, 28, 55,

66]. Additionally, also astrocytes may influence the outcome of remyelination. They can be beneficial for remyelination by promotion of oligodendroglial differentiation and providing of cholesterol for new myelin sheaths, but have been also shown to impair remyelination by either activating remyelination inhibiting pathways such as the JAG1-NOTCH1 pathway or by modulating the extracellular matrix [10, 32, 35, 61, 72]. One also has to keep in mind that the biopsies were taken from patients with an untypical clinical presentation and, therefore, might not be representative for MS patients with a more typical disease onset. However, most of the patients diagnosed with an inflammatory demyelinating lesion consistent with MS, developed clinical definitive MS with a relapsing–remitting disease course [59].

The histological appearance of MS lesions changes over time. In acute and RRMS active lesions predominate; however, when MS progresses, the percentage of active lesions decreases, whereas the percentages of mixed and inactive lesions increase [26, 46]. Patients with a more severe disease course have a higher percentage of mixed lesions suggesting that mixed lesions contribute significantly to disease progression [46]. Interestingly, we observed an almost complete absence of remyelination in mixed lesions in contrast to active or inactive lesions. This is in line with an earlier publication by Luchetti and colleagues who reported an inverse correlation between mixed versus remyelinated lesions [46]. Remyelination is frequently occurring at the lesion border; however, mixed lesions are characterized by a rim of myeloid cells at the lesion border suggesting that tissue environment in the rim of mixed lesions prevents successful remyelination. Interestingly, we observed a relative increase in TMEM119⁺ and iNOS⁺ and a decrease of CD163⁺ myeloid cells in the rim of mixed lesions compared to active lesions; however, one has also to keep in mind that TMEM119 is a homeostatic microglia marker whose expression decreases with activation [4]. In EAE, myeloid cells change from an iNOS⁺ pro-inflammatory and tissue-destructive phenotype to an iNOS⁻ tissue repair phenotype and this may be mediated by other immune cells, such as T and B cells [23, 43, 58]. This is in contrast to mixed MS lesions where at least a subset of myeloid cells preserves an iNOS⁺ phenotype. To further understand consequences of pro-inflammatory myeloid cells for oligodendrocytes, we exposed hiOL to the supernatants of pro- and anti-inflammatory microglia. We observed a marked decrease in the terminal differentiation into MBP⁺ oligodendrocytes after exposure to supernatants from pro-inflammatory human microglia, whereas supernatants from microglia polarized into an M0 or an M2 phenotype did not affect oligodendroglial differentiation. This is contrary to results from rodent studies in which supernatants from M0 and M1 polarized microglia had no effect on oligodendroglial differentiation, whereas the supernatants from M2 polarized microglia promoted oligodendroglial

differentiation [52]. Microvesicles isolated from either pro- or anti-inflammatory rat microglia both promoted the differentiation of rat oligodendrocytes [44]. These different results suggest that there might be significant differences between rodent and human oligodendrocytes with respect to the reaction to an inflammatory milieu. However, also the protocols for the polarization into pro- and anti-inflammatory microglial cells were slightly different and this might have also affected the outcome. Furthermore, other immune cells, such as T and B cells, have been reported to influence remyelination, either via direct effects on oligodendrocytes or by modulating myeloid cells [17].

We observed a comparable percentage of lesions with marked remyelination in active and inactive lesions. Therefore, it might be tempting to speculate that active lesions with marked remyelination become remyelinated inactive lesions, whereas active lesions lacking marked remyelination turn into mixed lesions and potentially afterwards into inactive lesions lacking remyelination. However, as mentioned above, histological studies provide only a snapshot and imaging studies demonstrated repeated waves of de- and remyelination in MS lesions [6]. Imaging markers for the different lesions types and longitudinal imaging studies are required to understand how lesions develop over time and how they contribute to disease progression.

In summary, our data suggest that the underlying causes for remyelination failure in MS are lesion stage dependent. Whereas in active/demyelinating lesions, lack of myelin sheath formation contributes to remyelination failure, in inactive and mixed lesions oligodendroglial loss and a hostile tissue environment may prevent successful remyelination. Since pharmacological approaches that focus exclusively on promotion of oligodendroglial differentiation to enhance remyelination might fail, treatment strategies targeting multiple of the steps required for successful remyelination, namely oligodendroglial proliferation, migration, differentiation, myelin sheath formation, and survival should be considered. Our data also suggest that promotion of remyelination in progressive MS in which mixed and inactive lesions dominate may require different treatment approaches compared to RRMS characterized by more active lesions. To understand the molecular mechanisms driving and preventing remyelination and to develop new treatment approaches which successfully promote remyelination in MS, better animal models are required which mimic the pathological hallmarks of the different lesion types.

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Author contributions TK initiated the project. KH, LS, and TK conceived and designed the study. KH and TK analyzed MS tissue samples. LS developed and performed the cell culture experiments and analyzed the data. NWK, MCV, LH, and H generated and characterized the supernatants from inflammatory cells. CT analysed data. JA, GM, and IH discussed results and provided cell lines. KH, LS, and TK wrote the manuscript with critical insight and commentary from all co-authors.

Compliance with ethical standards

Conflict of interest T.K. has a pending patent application for the generation of human oligodendrocytes.

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