Rejuvenation of Regeneration in the Aging Central Nervous System

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SUMMARY

Remyelination is a regenerative process in the central nervous system (CNS) that produces new myelin sheaths from adult stem cells. The decline in remyelination that occurs with advancing age poses a significant barrier to therapy in the CNS, particularly for long-term demyelinating diseases such as multiple sclerosis (MS). Here we show that remyelination of experimentally induced demyelination is enhanced in old mice exposed to a youthful systemic milieu through heterochronic parabiosis. Restored remyelination in old animals involves recruitment to the repairing lesions of blood-derived monocytes from the young parabiotic partner, and preventing this recruitment partially inhibits rejuvenation of remyelination. These data suggest that enhanced remyelinating activity requires both youthful monocytes and other factors, and that remyelination-enhancing therapies targeting endogenous cells can be effective throughout life.

INTRODUCTION

Remyelination is a spontaneous regenerative process in the adult CNS that restores saltatory conduction, prevents axonal degeneration, and promotes functional recovery (Duncan et al., 2009; Edgar and Nave, 2009; Smith et al., 1979). Stimulation of remyelination in demyelinating diseases such as MS could alleviate the major underlying causes of disability—impaired conduction by demyelinated neurons and axonal degeneration.

However, like most mammalian tissues, the CNS experiences declining efficiency of regeneration with increasing age (Sim et al., 2002). Reduced remyelination in aged animals occurs in part due to changes in the environmental signals regulating remyelination (Hinks and Franklin, 2000), but also reflects epigenetic changes within aging oligodendrocyte precursor cells (OPCs), which decrease their ability to differentiate into remyelinating oligodendrocytes (Shen et al., 2008; Tang et al., 2000). These age-dependent changes mirror, and may in part determine, a well-recognized feature of many chronically demyelinated MS lesions, which contain oligodendrocyte lineage cells that fail to differentiate into myelinating oligodendrocytes (Kuhlmann et al., 2008).

Recent data suggest that age-associated defects in neural stem cells can be reversed by reactivation of telomerase (Jaskeloff et al., 2011), suggesting that aged OPCs might, in principle, remain competent for efficient remyelination. We therefore investigated whether aged OPCs can indeed be rejuvenated by exogenous factors, reversing the typical age-associated decline in remyelination efficiency.

Using toxin-induced focal demyelination in the mouse spinal cord, together with heterochronic parabiosis (Conboy et al., 2005; Villeda et al., 2011), we tested whether exposure to a youthful systemic environment might improve remyelination by OPCs in the aged CNS. We chose this experimental system for several reasons. First, acute toxin-induced demyelination is better suited to studying the regenerative biology of demyelination than immune-mediated models that are complicated by autoimmunity. Second, because effective remyelination is generally associated with acute demyelination, it is likely that chronic demyelination arises as a consequence of deficiencies in the regenerative response; thus, age-associated delays in remyelination in an acute experimental demyelination model reflect a possible basis for the evolution of chronic demyelination in MS (Confavreux and Vukusic, 2006; Goldschmidt et al., 2009). Third, use of the parabiotic system uniquely tests the relevance of systemic factors to regeneration in the CNS by permitting exposure of aged tissues to blood-borne factors at physiologically relevant levels.

RESULTS

Exposure to a Youthful Systemic Environment Enhances Remyelination in Aged Animals

To examine the impact of the systemic environment on remyelination efficiency after spinal cord demyelination, we surgically joined aged mice to isogenic or congenic young animals through heterochronic parabiosis. Three weeks after animals were parabiotically joined, demyelination was induced in the spinal cord of the old partner by focal injection of the demyelinating toxin lysolecithin. To control for possible effects of the parabiotic condition itself, heterochronous mice were always compared to isochronous pairs (young mice joined to young partners, or old mice joined to old partners).
Chimerism in parabiotic pairs and accessibility of experimentally induced lesions to circulating blood cells was evaluated using pairs in which one partner was transgenically marked by expression of green fluorescent protein (GFP). Cross-circulation was confirmed by flow cytometry at the time of sacrifice and revealed the expected mixing of GFP− and GFP+ cells in the spleens of each partner (Figures S1A and S1B available online) (Wright et al., 2001). GFP+ cells were also detected in the lesions of old WT partners at multiple postlesion time points (Figure S1C). Together with the extravasation of fibrinogen and IgG at the lesion site (Figure S1D), these data point to disruption of the blood-spinal cord barrier in this model, as has been observed in MS. Thus, the spinal cord lesions of old animals in heterochronic pairings were exposed to both circulating cells and soluble factors derived from the young partner.

To ascertain the impact of exposure to youthful blood-borne factors on remyelination activity in aged partners, we first compared lesions at 7 days postlesion (dpl), when the lesions contain an expanding population of OPCs but no oligodendrocytes, and again at 14 dpl when new OPC-derived oligodendrocytes appear. The numbers of proliferating OPCs were identified by coexpression of the OPC transcription factor Nkx2.2 and the proliferation marker Ki67. The density of proliferating OPCs was significantly increased in heterochronic-old animals compared with isochronic-old controls at 7 and 14 dpl (Figure 1A). The enhanced OPC proliferation was associated with a significant increase in the density of CD34+ endothelial cells in heterochronic-old lesions at 14 dpl (Figure 1B); vascular endothelial cells have previously been shown to induce OPC proliferation (Arai and Lo, 2009). Reactive astrocytosis was not different between the three parabiotic conditions (Figure S1E). OPC apoptosis, measured by colabeling of Nkx2.2+ cells with terminal deoxynucleotidyl transferase-mediated deoxyuridine triphosphate nick-end labeling (TUNEL), was not significantly different in heterochronic-old animals compared with isochronic-old controls at either 7 or 14 dpl (Figure 1C).

We next determined the density of Olig2+/CC1+ mature oligodendrocytes in heterochronic-old lesions compared with isochronic-old controls, and found it significantly increased at 14 and also 21 dpl, when in young animals remyelination is complete (Figures 2A and 2B). At 21 dpl, the prevalence of mature oligodendrocytes in heterochronic-old lesions was equivalent to Olig2+/CC1+ cell densities in the isochronic-young group. These results were confirmed using antibodies to transferrin, an independent marker of oligodendrocytes (Connor and Fine, 1986) (Figure S2A). To test whether the enhanced production of Olig2+/CC1+ mature oligodendrocytes in heterochronic-old lesions leads to increased remyelination, we used histological analysis of semithin resin sections to assess differences in remyelination in all three groups at 21 dpl. This technique reliably detects the characteristically thin myelin sheaths that identify remyelination, and revealed that remyelination improved significantly in heterochronic-old animals as compared with isochronic-old controls (Figures 2C and 2D). Lesion volume in heterochronic animals did not differ from lesion volume in isochronic-old animals (Figure S2B), indicating that the parabiotic enhancement of remyelination was not attributable to differences in lesion size. Together, these data demonstrate that exposure of aged animals to a youthful systemic environment increases angiogenesis at the site of demyelinated lesions, promotes OPC proliferation, and reverses the age-associated differentiation block of OPCs, restoring the ability of these cells to form mature, remyelinating oligodendrocytes to levels indistinguishable from those of young animals.

Figure 1. Heterochronic Parabiosis Stimulates OPC Proliferation and Angiogenesis

(A) Proliferating OPCs (Nkx2.2+/Ki67+) within lesions at 7 and 14 dpl. Arrows indicate colocalization. n ≥ 4 pairs were analyzed for each condition and time point. Quantified data are presented as means ± SEM. Data were analyzed by one-way ANOVA followed by Tukey's post test. *p < 0.05, **p < 0.001. Scale bars: (A) and (C), 20 μm; (B), 125 μm. See also Figure S1.
The Young Systemic Environment Enhances the Remyelinating Function of Endogenous, Aged OPCs

In heterochronic GFPyoung/WTold pairs, GFP+ cells were concentrated within the lesioned area of the spinal cord (Figures 3A and 3B). We found no colocalization of the oligodendrocyte lineage marker Olig2 with GFP+ cells in the lesions of old partners (Figure 3C), indicating that enhanced remyelination did not result from engraftment of young OPCs, but was, rather, mediated by endogenous old OPCs, whose differentiation capacity was restored by exposure to a young systemic environment. In addition, to clarify whether enhancement of OPC differentiation in old animals exposed to a young systemic environment reflects a positive influence of the young environment, or dilution of negative inputs from the aged environment, we next asked whether OPC differentiation might be impaired in a young mouse exposed to an old circulation. Demyelinating spinal cord lesions in young mice parabiotically joined to old partners revealed no impairment of OPC differentiation (Figure 2E) or remyelination (Figure 2F) compared to isochronic-young control mice. Thus, in contrast to a recent report indicating the existence of age-regulated systemic factors that actively suppress neurogenesis in the CNS (Villeda et al., 2011), exposure of young OPCs to an aged systemic environment does not dominantly impair regenerative function in young mice.

Monocytes from the Young Partner Are Critical for Rejuvenation of Remyelination

Analysis of the GFP+ cells within the lesions of old partners in heterochronic pairings revealed them to be almost exclusively MAC1+ macrophages (Figures 3D and 3E). Very small numbers of GFP+/CD4+ and GFP+/CD8+ T lymphocytes, GFP+/CD94+ natural killer cells, GFP+/B220+ B cells and GFP+/NIMP-R14+ neutrophils were also present (Figure 3E). The inflammatory infiltrate derived from the parabiotic partner did not differ in extent or composition (Figure 3E) among isochronic-young, isochronic-old, and heterochronic-old animals at 5 dpl. In addition, the mean densities of partner-derived GFP+ macrophages within the lesions (~200 cells/mm²) were comparable to those reported in a parabiosis model of autoimmune spinal cord demyelination (Ajami et al., 2011).

Because the innate immune system (principally macrophages) has been suggested to stimulate remyelination (Kotter et al., 2006; Shechter et al., 2009), we hypothesized that the young blood-derived macrophages recruited to the lesion might play...
Figure 4. Young Macrophages Play a Central Role in the Rejuvenating Effect of Heterochronic Parabiosis

(A) Experimental rationale: in CCR2<sup>−/−</sup> WT<sup>old</sup> parabionts, only old monocytes are recruited to the old partner lesion in response to MCP-1 signaling. Soluble factors retain access to the lesion by virtue of local disruption of the blood-spinal-cord barrier (BBB).

(B) Immunostaining for GFP and MAC1 in heterochronic WT<sup>young</sup>/WT<sup>old</sup> and CCR2<sup>−/−</sup> WT<sup>young</sup>/WT<sup>old</sup> pairs demonstrates the absence of donor-derived macrophages in CCR2<sup>−/−</sup> WT<sup>old</sup> lesions.

(C) Comparison of the inflammatory infiltrate at 5 dpl shows that recruitment of donor-derived MAC1+ IB4+ macrophages predominates in old WT mice paired with young WT mice (n = 3) and is abolished in old WT mice paired with young CCR2<sup>−/−</sup> mice (n = 4, white bars). There are small numbers of CD4<sup>+</sup> T cells, CD8<sup>+</sup> T cells, CD94<sup>+</sup> natural killer cells, B220<sup>+</sup> B cells, and NIMP-R14<sup>+</sup> neutrophils in the lesions of both pairings.
a role in the rejuvenation of remyelination via heterochronic parabiosis. Monocytes/macrophages are recruited from the circulation to sites of inflammation by chemotaxis in response to gradients of MCP-1, which binds the chemokine receptor CCR2 (Charo and Ransohoff, 2006). Given this critical role of CCR2 in macrophage recruitment (Boring et al., 1997; Kurihara et al., 1997), we examined remyelination in single young CCR2-deficient mice and found a significant reduction in CC1+/Olig2+ cells at 21 dpl (Figure S3A), thereby confirming a role for recruited monocytes in remyelination. We next used CCR2-deficient, GFP-expressing mice as young partners in heterochronic parabiotic pairings. This system specifically blocked the recruitment of young macrophages to lesions in old partner animals (Figures 4A and 4B), but did not affect cross-circulation, as measured by splenic chimerism, in either isochronic or heterochronic pairs (Figure S3B). Recruitment of GFP+ nonmacrophage cells, which was very low to begin with, was largely unaffected by CCR2 deficiency of the young partner (Figure 4C), and aged CCR2+ monocytes were still recruited to the lesion from the chimeric circulation.

We found that the density of mature oligodendrocytes was decreased significantly in lesions of old WT animals paired to young CCR2-deficient animals, as compared with old animals paired with young WT partners (Figure 4D). However, compared with isochronic-old controls, heterochronic-old animals, in which the recruitment of young macrophages was abolished by CCR2 deficiency (Figure 4C), exhibited only partial inhibition of the rejuvenating effect of heterochronic parabiosis on OPC differentiation (Figure 4D). Thus, this strategy represents a useful “cell deletion system” that provides clarification of the role of young monocytes, the major population of recruited young-partner cells, in the rejuvenation of remyelination.

**Young Macrophages Enhance Remyelination via Myelin Debris Clearance**

Given the importance of youthful macrophages in stimulating remyelinating activity in aged animals, we next asked how these cells might accomplish this effect in old lesions. We first examined whether young macrophages might produce factors that directly affect OPC function by analyzing the expression of growth factors previously implicated in OPC proliferation and maturation (Mason et al., 2003; Vana et al., 2007). However, analysis of bone-marrow-derived macrophages harvested from isochronic-young, isochronic-old, or heterochronic animals revealed no apparent differences in the production of a number of key OPC-regulatory growth factors (including IGF-1 and PDGF-1A) in any of the experimental conditions (Figures S4A and S4B). Similarly, because immune-modulatory cytokines previously have been implicated as systemic regulators of tissue repair in the skeletal muscle and CNS (Brack et al., 2007; Villeda et al., 2011), we undertook an array-based comparison of 97 cytokines in the serum of isochronic-old, heterochronic-old (WT/WT), and heterochronic-old paired with young CCR2-deficient animals (CCR2−/−/WT). This comparison yielded surprisingly few differences (Figures S4C–S4G), suggesting that modulation of factors within the local microenvironment of the lesion exerts a greater influence on remyelination than alterations in levels of specific circulating factors in the bloodstream.

We next considered whether differences in the phagocytic activity of young versus old macrophages might contribute to their differences in stimulating OPC function. Myelin contains proteins that inhibit OPC differentiation, and previous studies have highlighted the importance of macrophage-mediated removal of myelin debris in remyelination (Baer et al., 2009; Kotter et al., 2006). The efficiency of lipid-rich myelin debris clearance is reflected by the amount of detectable lipid within the lesion area. Staining of lesions with Oil-Red O revealed that lipid levels in the lesions of old partners involved in heterochronic parabiosis were similar to those in young isochronic pairs, and were significantly reduced when compared with those in old isochronic pairs (Figures 4E and 4F). The enhancement of myelin debris clearance after heterochronic pairing was substantially attenuated in pairings involving CCR2-deficient young partners (Figures 4E and 4F). Similar results were obtained using antibodies to a degenerate form of the myelin protein myelin basic protein (MBP) (Figures 4G and 4H). Together, these data are consistent with a model in which young macrophages facilitate OPC differentiation and remyelination by augmenting the clearance of inhibitory myelin debris.

**DISCUSSION**

These experiments provide proof of principle that, despite their intrinsic alterations, aged OPCs remain responsive to exogenous prodifferentiation signals and retain their competence for efficient repair. These data further indicate that signals from the systemic environment can override age-related, intrinsic changes in OPCs, suggesting that these changes are predominantly epigenetic. This finding has fundamental implications for strategic approaches to remyelination-enhancing therapies, validating endogenous OPCs—even when aged, as they are likely to be in many MS patients—as a pharmacological target to stimulate remyelination.

Although the full cast of characters involved in the rejuvenation of aged OPCs has yet to be identified, our data implicates a cell-based mechanism in which young macrophages recruited

(D) Oligodendrocyte (CC1+/Olig2+) density at 21 dpl in CCR2−/−/young/WTold pairs was significantly reduced compared with that in heterochronic old WT pairs, but elevated compared with that of isochronic-old controls (n = 4 pairs per condition). Data were analyzed by one-way ANOVA followed by Dunnett’s post test. (E and F) Oil-Red O staining revealed significantly more lipid-rich myelin debris in the lesions of isochronic-old pairings than in isochronic-young pairings, consistent with the notion that myelin debris have an inhibitory effect on remyelination. Oil-Red O staining in heterochronic-old WT lesions was significantly lower than in isochronic-old controls. This enhanced myelin debris clearance observed in heterochronic pairings was attenuated when the young partner was CCR2 deficient. Note that the staining appears less diffuse in the isochronic-young and the heterochronic groups, suggesting that more myelin debris has been taken up by phagocytic cells. (G and H) Similar observations were made for intralesional degenerated myelin basic protein accumulation. *p < 0.05; **p < 0.001. Data are means ± SEM. Scale bars: (B), 50 μm; (E) and (G), 100 μm. See also Figure S3 and Figure S4.
during the early phase of remyelination alter OPC survival, proliferation, and differentiation to promote a more youthful remyelination response. Central to this function is the greater capacity of young macrophages than old macrophages for efficient clearance of myelin debris. This finding is consistent with prior observations that transplanted young macrophages are more effective than old macrophages at improving wound repair in old animals (Danon et al., 1989). Moreover, this work demonstrates that the CNS maintains its responsiveness to age-regulated circulatory factors, such that age-dependent deficiencies in repair of these tissues can, in part, be reversed by circulating factors.

**EXPERIMENTAL PROCEDURES**

**Animal Procedures**

C57BL/6 (WT) mice, C57BL/6-UBC-GFP transgenic mice, and CCR2-/- mice on the C57BL/6 background were obtained from Jackson Laboratories (USA). Parabiotic pairs with a GFP+ partner were generated with either C57BL/6-UBC-GFP transgenic mice or C57BL/6-CCR2-/- mice. Parabiotic pairs were joined as previously described (Wagers et al., 2002). A focal demyelinating spinal cord lesion was induced as previously described (Woodruff et al., 2004). See Supplemental Experimental Procedures for details.

**Flow Cytometry Analysis of Cross-Circulation**

Spleens from both partners of GFP+/WT parabiotic pairs were removed and homogenized, and red blood cells were lysed. Cells were mixed with propidium iodide (PI) solution to label dead cells, which were excluded from analysis. Samples were subjected to flow cytometry and analyzed using FlowJo software (TreeStar, USA).

**Immunohistochemistry and Quantification of Immunolabeling**

For staining with the anti-Olig2, anti-Ki67, and anti-Nkx2.2 antibodies, antigen retrieval was performed. For staining with the mouse primary antibody anti-Nkx2.2, the Mouse-on-Mouse immunodetection kit (Vector laboratories, BPK-2202) was used with additional labeling with other antibodies or TUNEL. For CD34 immunostaining, slides were pretreated with ice-cold methanol for 10 min before being blocked. Nuclei were counterstained with DAPI. For CD34 immunostaining, slides were pretreated with ice-cold methanol for 10 min before being blocked. Nuclei were counterstained with DAPI. For staining with the mouse primary antibody anti-Nkx2.2, the Mouse-on-Mouse immunodetection kit (Vector laboratories, BPK-2202) was used with additional labeling with other antibodies or TUNEL. For CD34 immunostaining, slides were pretreated with ice-cold methanol for 10 min before being blocked. Nuclei were counterstained with DAPI. For staining with the mouse primary antibody anti-Nkx2.2, the Mouse-on-Mouse immunodetection kit (Vector laboratories, BPK-2202) was used with additional labeling with other antibodies or TUNEL. For CD34 immunostaining, slides were pretreated with ice-cold methanol for 10 min before being blocked. Nuclei were counterstained with DAPI. For staining with the mouse primary antibody anti-Nkx2.2, the Mouse-on-Mouse immunodetection kit (Vector laboratories, BPK-2202) was used with additional labeling with other antibodies or TUNEL.

**Cytokine Arrays**

Blood samples were collected from each partner of the parabiotic pair directly before sacrifice. The RayBio Mouse Cytokine Antibody Array 6 (RayBiotech) was used to detect the presence or absence of 97 cytokines in each serum sample.

**Statistics**

All statistical analysis was performed in GraphPad Prism (GraphPad Software) and differences were considered significant at p < 0.05 (see Supplemental Experimental Procedures for details).

**SUPPLEMENTAL INFORMATION**

Supplemental Information for this article includes four figures and Supplemental Experimental Procedures and can be found with this article online at doi:10.1016/j.stem.2011.11.019.

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