Interleukin-10 Improves Stroke Outcome by Controlling the Detrimental Interleukin-17A Response

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**Interleukin-10 improves stroke outcome
by controlling the detrimental Interleukin-17A response**

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Keywords

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Abstract

Background: Lymphocytes have dichotomous functions in ischemic stroke. Regulatory T cells are protective, while IL-17A from innate lymphocytes promotes the infarcts growth. With recent advances of T cell-subtype specific transgenic mouse models it now has become possible to study the complex interplay of T cell subpopulations in ischemic stroke.

Methods: In a murine model of experimental stroke we analyzed the effects of IL-10 on the functional outcome for up to 14 days post-ischemia and defined the source of IL-10 in ischemic brains based on immunohistochemistry, flow cytometry, and bone marrow chimeric mice. We used neutralizing IL-17A antibodies, intrathecal IL-10 injections, and transgenic mouse models which harbor a deletion of the IL-10R on distinct T cell subpopulations to further explore the interplay between IL-10 and IL-17A pathways in the ischemic brain.

Results: We demonstrate that IL-10 deficient mice exhibit significantly increased infarct sizes on days three and seven and enlarged brain atrophy and impaired neurological outcome on day fourteen following tMCAO. In ischemic brains IL-10 producing immune cells included regulatory T cells, macrophages, and microglia. Neutralization of IL-17A following stroke reversed the worse outcome in IL-10 deficient mice and intracerebral treatment with recombinant IL-10 revealed that IL-10 controlled IL-17A positive lymphocytes in ischemic brains. Importantly, IL-10 acted differentially on $\alpha\beta$ and $\gamma\delta$ T cells. IL-17A producing CD4⁺ $\alpha\beta$ T cells were directly controlled via their IL-10-receptor (IL-10R), whereas IL-10 by itself had no direct effect on the IL-17A production in $\gamma\delta$ T cells. The control of the IL-17A production in $\gamma\delta$ T cells depended on an intact IL10R signaling in regulatory T cells (Tregs).

Conclusions: Taken together, our data indicate a key function of IL-10 in restricting the detrimental IL-17A-signaling in stroke and further supports that IL-17A is a therapeutic opportunity for stroke treatment.

Background

Worldwide, stroke is one of the leading causes of mortality and sustained disability, with 5.5 million deaths and 11 million disability-adjusted life-years in 2016 [1].

Inflammation is a key component of stroke pathophysiology, with dichotomous effects on the affected tissue. In the early stages, highly conserved pro-inflammatory pathways lead to a worsening of the initial tissue damage, whereas protective mechanisms are involved in tissue repair and ischemic tolerance in the long term [2].

Among the inflammatory cascades, which drive the early sterile immune response, IL-17A holds an important role. Following stroke, IL-17A is mainly produced by innate-like $\gamma\delta$ T cells. By driving the expression of neutrophil attracting chemokines (e.g. C-X-C motif chemokines (CXCL-1)), IL-17A is crucial for the amplification of the local inflammatory response in the first hours, resulting in a worsened neurological outcome [3, 4]. However, IL-17A levels and numbers of infiltrating neutrophils already decrease 4 days after ischemia [5]. Immunological mechanisms that limit the IL-17A-dependent inflammatory response are largely unknown. Among potential anti-inflammatory pathways, IL-10 holds a prominent role, due to its strong immunomodulatory and neuroprotective effects [6, 7]. In stroke, IL-10 has been shown to negatively regulate proinflammatory cytokines (e.g. IFN- γ , TNF- α), and antigen-presenting cells, including microglia [6]. Although IL-10 and IL-17A have opposing effects on stroke outcome, it is largely unknown whether these two cytokines interact. Studies from models of autoimmunity [8, 9] indicate that regulatory CD4⁺ T cells and IL-10 represent a fundamental mechanism for regulating IL-17A. It has been shown that IL-10 can directly inhibit the IL-17A production of CD4⁺ effector T cells [8]. Interestingly, disruption of IL-10 signaling on CD4⁺ Tregs and T regulatory type 1 cells (T_r1) also reduced their capacity to suppress the Th17 immune response which suggests an indirect suppression via an autocrine feed-forward loop [9, 10].

Compared to the regulation of $\alpha\beta$ T cells less is known about IL-10 effects on $\gamma\delta$ T cells. However, Park et al. suggested that Treg cell-derived IL-10 directly suppresses TCR $\gamma\delta^+$ T cells by limiting their proliferation and IL-17A production in a model of inflammatory bowel disease [11].

The goal of this study was to analyze the role of the IL-10 signaling for the control of the detrimental IL-17A response in stroke. Our data show that IL-10 plays a key role in regulating IL-17A production in $\alpha\beta$ and $\gamma\delta$ T cells in the ischemic brain. Here, IL-10 acts differentially as CD4 $^+$ T cells are directly controlled, whereas intact IL-10R signaling on Tregs is required to suppress the IL-17A production in $\gamma\delta$ T cells.

Materials and Methods

Animals

All animal experiments were approved by local animal care committees (Behörde für Justiz und Verbraucherschutz der Freien und Hansestadt Hamburg Lebensmittelsicherheit und Veterinärwesen, Hamburg, Germany). Experiments were conducted in accordance with the Guide for the Care and Use of Laboratory Animals published by the US National Institutes of Health (publication No. 83–123, revised 1996). All procedures were performed in accordance with the ARRIVE guidelines (Animal Research: Reporting of In Vivo Experiments).

10-18 weeks old male C57BL/6 mice provided by the animal facility of the University Medical Center Hamburg-Eppendorf were studied. All mice were backcrossed on a C57BL/6 background for >10 generations. To evaluate effects on a global IL-10 deficiency and to perform bone marrow transplantation we used *Il10* $^{-/-}$ mice and respective WT littermate controls. Subsets of Tregs were studied in male double reporter *Foxp3-IRES-mRFP (FIR) x Il10-IRES-GFP-enhanced reporter (tiger)* mice. To deplete the IL-10R on CD4 $^+$, Foxp3 $^+$, or $\gamma\delta$ T cells we took advantage of the Cre-loxP system and crossed *CD4* Cre , *Foxp3* Cre , and *TCR δ* CreER

with *Il10Ra^{fl/fl}* mice respectively. *CD4^{Cre}* [12], *Foxp3^{Cre}* [13], *Il10Ra^{fl/fl}* mice [14], *FIR (Foxp3-IRES-mRFP)* [15], and tiger (*Il10-IRES-GFP-enhanced*) [16] reporter mice were kindly provided by S. Huber and are described elsewhere. *TCRδ^{CreER}* [17] mice were kindly provided by I. Prinz and are described elsewhere. For the induction of Cre-recombinase *TCRδ^{CreER}* animals were treated three times with 2 mg Tamoxifen within 5 days at the age of 8-10 weeks. Two weeks later the animals were used for experiments. CD45.1⁺ WT and *Il10^{-/-}* mice were obtained from the Jackson Laboratory. For all animal experiments, mice were randomized and coded by an independent researcher, to achieve blinding of the experiments. Age-matched male wildtype (WT) littermates served as controls. To reduce the variability of our outcome parameters (neurological outcome, inflammatory response) caused by sex differences and to thereby decrease group sizes, only male mice were used throughout the study. Recent studies have demonstrated profound effects of sex-differences on infarct sizes and the inflammatory response. To validate our studies in females, further studies are required.

Transient Middle Cerebral Artery Occlusion (tMCAO)

We randomized all mice and conducted transient middle cerebral artery occlusion (tMCAO), using 6-0 nylon monofilament with silicone-coated tips (Doccol Corp.) as described previously [4]. Mice were anesthetized and monitored for heart rate, respiratory rate, oxygen saturation, and rectal body temperature. An appropriate reduction (> 80%) in regional cerebral blood flow was confirmed by transcranial laser Doppler flowmetry (Perimed) in the area of the cerebral cortex supplied by the MCA. After 45 minutes, the monofilament was retracted to allow reperfusion for 3, 7, or 14 days. In the sham group, carotid arteries were visualized without insertion of the filament. Following surgery, animals were kept on heating pads and sterile saline was given via subcutaneous injection for rehydration. Exclusion criteria were defined in a score, which included weight loss, general condition, spontaneous behavior, and impairment of wound healing. We anesthetized all mice (20 to 25 g, 12 weeks; TVH, University Medical

Center Hamburg-Eppendorf) using isoflurane 1% to 2% v/v oxygen. For analgesia we injected buprenorphine 0.03 mg/kg BW intraperitoneally every 12 hours for 24 hours. Mice were excluded from the study if they (i) died during the surgical procedure, (ii) experienced subarachnoid hemorrhage, (iii) had a Bederson score less than one after reawakening, or (iv) due to <80% reduction in regional cerebral blood flow during MCAO.

Antibody treatment

Animals were treated intravenously with 500 µg of mouse monoclonal anti-murine IL-17A antibody (Clone MM17F3; 16.6 mg/kg of body weight) or with 500 µg of isotype-control antibody (IgG1, BioXCell, MOPC-21, BE0083) 3h and 3 days after reperfusion.

Intracerebroventricular injections

Within 3 hours after induction of MCA occlusion we performed an intracerebral injection of 200 ng control vehicle [bovine serum albumin (BSA) in 2 µl artificial cerebrospinal fluid consisting of mM: 126 NaCl; 2.5 KCl; 1.2 NaH₂PO₄; 1.3 MgCl₂ and 2.4 CaCl₂ at pH 7.4] or 200 ng IL-10 [recombinant mouse protein (rmIL-10 (carrier-free), Biolegend #575802) in 2µl artificial cerebrospinal fluid].

Intracerebral injections were performed as previously described [18]: Briefly, mice were anesthetized with isoflurane (4% for induction; 2.5% for maintenance) in 100 % oxygen. The mice were placed in a stereotactic frame (Stoelting, 51500U) and a 1 cm long incision of the scalp was made over the midline. A cranial burr hole (0.9 mm) was drilled 2.2 mm lateral to the midline and 0.2 mm anterior to the bregma. A 10 µl Hamilton syringe (Hamilton, 1701RN) was connected to a 26-gauge needle (Hamilton, 26G, Point Style 4, 12°) and inserted into a motorized stereotaxic injector (Stoelting, integrated stereotaxic injector [ISI]). The 26-gauge needle was then slowly introduced 3.7 mm deep into the left hemisphere and 2 µl of IL-10 or control vehicle was infused at a rate of 0.5 µl/min. The needle was left in place for another 10

min and then slowly withdrawn. Then a suture for the scalp incision was made. Animals recovered for a period of 60 min under a heating lamp with free access to food and water.

Scoring

After stroke induction, we repeatedly scored mice using the Bederson score on a scale from 0–5. This was done immediately after reawakening and every day until sacrifice. A composite neurological score was assigned 14 days after tMCAO. Bederson Score: 0 no deficit, 1 preferential turning, 2 circlings, 3 longitudinal rollings, 4 no movement, 5 death. Composite neurological score: Each mouse was rated on neurologic function scales unique to the mouse. Scores range from 0 (healthy) to 59 (the worst performance in all categories) and represent the sum of the results of all categories. These include general deficits: Hair (0-2), eyes (0-3), ears (0-2), posture (0-3), spontaneous activity (0-3), and focal deficits: circling (0-5), gait (0-4), body symmetry (0-3), climbing on a surface held at 45° (0-4), forelimb symmetry (0-4), compulsory circling (0-3), whisker response to a light touch (0-4), gripping (0-3), beam walking (0-9), balance on a beam (0-3), and balance on a round stick (0-4).

Analysis of Infarct Size by Triphenyltetrazolium chloride Staining and Magnetic Resonance Imaging (MRI)

The infarct size was analyzed 7 days post-reperfusion by harvesting brains and cutting them into 1 mm slices (Braintree Scientific, 1 mm) followed by vital staining using 2% (wt/vol) 2,3,5-triphenyl-2-hydroxy-tetrazolium chloride in phosphate buffer. We determined infarct volumes in a blinded fashion using NIH ImageJ software.

MRI comprised T2-weighted imaging and was performed on a dedicated 7T MR small animal imaging system (ClinScan; Bruker) 3 and 14 days post-reperfusion.

Bone marrow chimeras

For the generation of bone marrow chimeras WT or *Il10*^{-/-} mice were reconstituted with either *Il10*^{-/-} or WT bone marrow respectively. 6-8-week-old male recipients (CD45.2⁺) were exposed to whole-body irradiation (9 Gy; 1 Gy min⁻¹) using a cesium-137 gamma irradiator (BIOBEAM 2000, Leipzig, Germany), as described before [19]. After 24h, mice were reconstituted with bone marrow cells derived from tibiae and femurs from CD45.2⁺ *Il10*^{-/-} or CD45.1⁺ WT mice respectively. Every recipient received 10⁷ bone marrow cells intravenously. We assessed reconstitution and distribution between CD45.1 and CD45.2 cells by FACS analysis of peripheral blood cells of recipient mice six weeks after grafting. tMCAO was induced in recipient mice six weeks after transplantation.

Cell Sorting

Immune cells were sorted using a BD FACS Aria IIIu and collected in RPMI with 25% fetal calf serum.

RNA Isolation and Quantitative Real-Time Polymerase Chain Reaction

We isolated RNA from brain tissue at indicated time points following tMCAO. Hemispheres were separated and homogenized in TRIzol Reagent (1ml per 100 mg tissue), chloroform was added, samples were centrifuged at 12000 g for 15 min at 4 °C and the upper aqueous phase was collected. RNA was precipitated by the addition of isopropyl alcohol, washed and dissolved in TE-Buffer. We isolated total RNA from cells using QIA-Shredder spin columns and the RNeasy Micro Kit (QIAGEN) and transcribed complementary DNA using Maxima First Strand cDNA Synthesis Kit for RT-qPCR (Fermentas). Real-time PCR primers were obtained from Applied Biosystems (Carlsbad, CA): Il6: Mm00446190_m1; Il23: Mm00518984_m1; Il1b: Mm00434228_m1; Tgfb: Mm01178820_m1; Cxcl1 Mm00433859_m1; Mmp3: Mm00442991_m1; BActin: Mm00607939_s1, Sdha: Mm01352366_m1. Probe mixtures were

purchased from Fermentas (Waltham, MA). The relative gene expression was calculated using the $\Delta\Delta C_t$ method and the samples were normalized to the control population and the expression of Sdha or β -Actin. Samples were randomized and coded by an independent researcher, so experiments were carried out blindly.

Antibodies and flow cytometry

Cell types were analyzed by flow cytometry, as previously described [4]. Briefly, mice were euthanized and perfused with phosphate-buffered saline. Only ipsilesional hemispheres were dissected, digested for 30 min at 37°C (1 mg/ml collagenase (Roche), 0.1 mg/ml DNase I (Roche) in DMEM) and pressed through a cell strainer. Cells were incubated with standard erythrocyte lysis buffer on ice and separated from myelin and debris by Percoll gradient (GE Healthcare) centrifugation, followed by surface staining. In case of further intracellular staining T cells were stimulated with phorbol 12-myristate 13-acetate (100 ng/mL; Sigma- Aldrich) and ionomycin (1 μ g/mL; Sigma-Aldrich) in the presence of brefeldin A (3 μ g/mL; eBioscience) at 37°C in 10% CO₂ for 4 hours before surface staining. To achieve fixation, permeabilization and intracellular staining True-Nuclear™ Transcription Factor Buffer Set (#424401) was used according to the manufacturer's instructions. Mouse antibodies were as follows: from Biolegend CD45.2 (104; #109805), CD45.1 (A20, #110707), B220 (RA3-6B2, #103245, #103237), CD3 (145-2C11, #100308; 17A2, #100228), NK1.1 (PK136, #108745, #108713), TCR- $\gamma\delta$ (GL-3, #118118), CD4 (RM4-5, #100553, #100551), CD8 (53.6-7, #100732, #100741, #100722), CD49b (HMA2, #103506), CD11b (M1/70, #101216), MHC II (M5/ 114.15.2; #107605), Ly6g (1 A8; #127627), F4/80 (BM8, # 123133), Lag-3 (C9B7W, #125210), PD-1 (29F.1A12, #135220), IFN- γ (XMG1.2, #505815, #505839), CD210 (1B1.3a, #112705); from eBioscience CD45 (30-F11, #47-0451-82, #56-0451-82), CD19 (eBio1D3, #25-0193-81), CD11c (N 418, #25-0114-82), TIGIT (GIGD7, #46-9501-82) and IL-17A (eBio17B1, #17-7177-81); from BDBioscience CD4 (GK1.5, #563790) and ROR γ T (Q31-378, #562607).

TrueCount tubes (Becton Dickinson), containing fluorescence beads were used to determine the total cell counts per hemisphere according to the manufacturer's protocol. 30% of each sample volume was used for an isotype control. Data were acquired with a Fortessa FACS system (BD Biosciences) and analyzed with FlowJo. Doublets were excluded with FSC-A and FSC-H linearity.

Immunohistochemistry and Immunofluorescence

Immunofluorescence for GFP and CD45 co-expressing cells were performed on 20 µm thick fresh frozen brain sections as previously described [20]. Cryostat cut sections were air-dried, rinsed in TBS for 10 min before blocked with 10% fetal calf serum in TBS + 0.5% Triton for 30 min at RT. Next, sections were incubated with rabbit anti-mouse GFP (ab290, Abcam) overnight at 4°C. The following day the sections were rinsed for 10 min in TBS + 0.5% Triton before incubation with the cross-absorbed species-specific secondary antibody Alexa Fluor 488 goat anti-rabbit (#A11070, Invitrogen) diluted 1:200 for 2 hours. Next, the sections were rinsed in TBS for 10 min before incubation for 2 hours with PE-conjugated rat anti-mouse CD45 (#553081 BD Biosciences). After a 2 x 10 min rinse in TBS the sections were transferred to TBS containing 40,6-diamidino-2-phenylindole (DAPI) (D1306, Invitrogen), rinsed in water, and mounted in gelvatol. Respective Isotype controls were devoid of signal.

Statistical Analysis

Statistical analyses were performed using the appropriate test indicated in the figure legends. Briefly, the Student t test was used to compare infarct volumes, the Mann-Whitney U test for the comparison of clinical scores, and the 1-way ANOVA test for multiple comparisons with the Bonferroni post hoc test. P values <0.05 were considered statistically significant.

Results

IL-10 is protective following transient middle cerebral artery occlusion in mice.

To investigate the role of IL-10 in acute ischemic brain damage and the subsequent recovery, we analyzed the neurological outcomes of Interleukin-10 knockout (*Il10*^{-/-}) mice and wild type (WT) littermate controls up to 14 days after tMCAO. On day 3 and day 7 *Il10*^{-/-} mice exhibited significantly increased infarct volumes (Figure 1A and B). Underlining the impact of IL-10 not only on the initial damage but also on the regeneration we found that *Il10*^{-/-} mice showed significantly increased brain atrophy and worse neurological scores on day 14 when compared to littermate controls (Figure 1C and D, and Additional file: Figure SIA). Mortality did not differ significantly between groups (Additional file: Figure SIB).

IL-10 is upregulated in infiltrating T cells and macrophages.

To visualize the spatial distribution and cellular source of IL-10 in post-stroke brains, we performed immunohistochemistry using a reporter mouse expressing a green fluorescent protein [GFP] under the control of the IL-10 promotor. 14 days following tMCAO, we detected GFP-positive CD45⁺ cells in the penumbra area of the infarct indicating that immune cells contribute to the local IL-10 production in ischemic brains (Figure 1E).

Next, we studied whether the IL-10 production from invading immune cells or brain resident cells including microglia, astrocytes, or neurons is decisive for the protective IL-10 effect following stroke. We generated bone marrow chimeric mice using CD45.1⁺ WT or CD45.2⁺ *Il10*^{-/-} donor mice which were injected into CD45.2⁺ *Il10*^{-/-} or WT recipient mice, respectively. Following reconstitution >98% leukocytes expressed congenic markers of the respective donor (Additional File: Figure SIIA and B). On day 3 after tMCAO we observed that *Il10*^{-/-} mice reconstituted with WT bone marrow displayed significantly reduced infarct volumes when compared to WT littermate controls reconstituted with *Il10*^{-/-} bone marrow cells. Underscoring

the importance of IL-10 from peripheral immune cells, we further found that *Il10*^{-/-} mice reconstituted with WT bone marrow displayed significantly improved neurological scores 14 days post-stroke, whereas brain atrophy did not differ significantly between groups (Figure 2A-C).

To further investigate the temporal dynamics and the cellular source of IL-10, we measured mRNA levels of *Il10* transcripts in microglia and infiltrating macrophages on days 3, 7, and 14 post-stroke. Notably, macrophages showed significantly increased IL-10 level on days 7 and 14, whereas microglia showed a significant upregulation only on day 14, suggesting that the IL-10 production of macrophages and microglia is delayed following ischemia (Figure 2D).

Besides microglia and macrophages, lymphocytes are an important source of IL-10. As the staining of IL-10 by hand is challenging we took advantage of the double-knock-in reporter *Fir-tiger* mouse model, which enables the simultaneous detection of IL-10 (green fluorescent protein [GFP]) and Foxp3 (monomeric red fluorescent protein [mRFP]) by flow cytometry. CD4⁺ T cells exhibited the highest frequency of IL-10 producers when compared to B and NK cells as well as CD8⁺ T cells (Additional File I: Figure SIIC) [21]. Interestingly, in CD4⁺ T cells, we identified two different IL-10⁺ subpopulations that peak at different timepoints following stroke. On day 3 following tMCAO, IL-10⁺ Foxp3⁻ CD4⁺ T cells were significantly increased in frequency compared to sham surgery. Notably, these CD4⁺ T cells did not co-express the Tr1 cell markers CD49b and Lag-3, while Tr1 cells were detectable at the same time point in spleens of respective mice (Additional File I: Figure SIID) [21]. On day 7 the IL-10⁺ Foxp3⁻ CD4⁺ T cells decreased in frequency, while IL-10⁺ Foxp3⁺ Tregs appeared in ischemic hemispheres reaching significance on day 14 when compared to sham surgery (Figure 2E).

Combined, this data suggests that the IL-10 production of infiltrating immune cells, in particular T cells and macrophages, is decisive for IL-10-mediated neuroprotection in stroke.

IL-10 controls the IL-17A response in the inflamed brain.

In models of autoimmunity including inflammatory bowel disease [8] or nephritis [9], IL-10 effects largely depend on the inhibition of IL-17A pathways. IL-17A dependent inflammatory pathologies in turn are induced by the prototypical pro-inflammatory cytokines IL-1 β and IL-23 among others. To investigate whether IL-1 β and IL-23 are also upregulated following ischemia, we investigated their expression in microglia and macrophages sorted from ischemic hemispheres. In microglia, we observed a significant increase in *Il23a* and *Il1b* mRNA levels on day 14 (Figure 3A). Macrophages showed a significant upregulation of *Il23a* and *Il1b* on day 7 and of *Il1b* on day 14, respectively (Figure 3B). Overall, these results demonstrate a sustained upregulation of IL-17A-polarizing cytokines in the ischemic brain.

To investigate whether IL-10 is capable to control IL-17A pathways in the presence of IL-1 β and IL-23, we next analyzed IL-17A level in $\alpha\beta$ and $\gamma\delta$ T cells from ischemic brains of *Il10*^{-/-} and WT mice. On day 7 we observed a significantly increased frequency of IL-17A⁺ CD4⁺ and IL-17A⁺ $\gamma\delta$ T cells in the post-ischemic CNS of *Il10*^{-/-} mice when compared to WT littermate controls (Figure 3C). Notably, in contrast to IL-17A positive $\gamma\delta$ T cells, we detected almost no Th17 cells on day 7 in WT littermates, indicating that physiological IL-10 levels strictly suppress the IL-17A production in CD4⁺ T cells in ischemic brains.

The main IL-17A effect in the ischemic brain is the induction of neutrophil-attracting chemokines (e.g. CXCL-1) [4, 22] as well as Matrix Metalloproteinases [4, 23]. Consistent with increased IL-17A levels, we detected a significant upregulation of *Cxcl1* and *Mmp3* transcripts in the whole brain mRNA of *Il10*^{-/-} mice at day 3 post-stroke (Figure 3D).

Next, we performed intracerebral injections of 200 ng recombinant IL-10 or Bovine Serum Albumin (BSA) in WT mice 3 h following tMCAO to evaluate the capacity of IL-10 to suppress IL-17A production locally in the ischemic brain. Intracerebral IL-10 treatment significantly lowered the presence of IL-17A producing $\gamma\delta$ T cells when compared to a BSA control 3 days post tMCAO (Figure 3E).

To show that the dysregulation of IL-17A is decisive for the worse outcome in *Il10*^{-/-} mice, we treated *Il10*^{-/-} mice and littermate controls with 500 µg IL-17A neutralizing antibody or IgG control 3 h and 3 days post tMCAO. Anti-IL17A treatment rendered the infarct volume and the neurological scores of *Il10*^{-/-} and WT littermate controls almost equal 7 days post-tMCAO (Figure 4A and B). Importantly, treatment with the IgG control antibody did not rescue the worse outcome of *Il10*^{-/-} when compared to littermate controls (Figure 4C and D).

Differential effects of IL-10 on IL-17A⁺ brain infiltrating CD4⁺ and γδ T cells

To identify whether IL-10 directly inhibits the IL-17A response through the IL-10 receptor on T cells, we next employed transgenic mice, in which the IL-10 signaling is abrogated in mature CD4⁺ T cells. To generate a deletion of the α-chain of the IL-10R in CD4⁺ T cells which is essential for IL-10 signaling, we bred *CD4*^{Cre} with *Il10Ra*^{fl/fl} mice. We observed a significant increase in frequency of IL-17A⁺ brain-infiltrating CD4⁺ and γδ T cells of *CD4*^{Cre} *x* *Il10Ra*^{fl/fl} mice at day 7 following tMCAO when compared to *CD4*^{Cre} *x* *Il10Ra*^{wt/wt} control mice (Figure 5 A and B). As γδ T cells from *CD4*^{Cre} *x* *Il10Ra*^{fl/fl} mice have an intact IL-10 receptor, these data suggest that γδ T cells are inhibited through alternative pathways which are indirectly controlled by IL-10 whereas IL-10 exerts its inhibitory functions directly on CD4⁺ T cells.

Notably, IL-17A⁺ γδ T cells display a significantly lower IL-10R expression when compared to IL-17A⁻ γδ T cells. This finding further supports the hypothesis that IL-17A⁺ γδ T cells are not directly inhibited by IL-10 (Figure 5C).

To confirm that IL-10 is not affecting γδ T cells directly, we bred *TCRδ*^{CreER} *x* *Il10Ra*^{fl/fl} mice. In line with our finding in *CD4*^{Cre} *x* *Il10Ra*^{fl/fl} mice, the selective depletion of the IL-10 receptor in γδ T cells had no direct effect on their IL-17A production (Figure 5D).

In sum, these data suggest that CD4 T cells play a central role for the control of the detrimental IL-17A response in ischemic brains, and that the IL-10 signaling in CD4 T cells is essential for their protective function. That the targeted depletion of the IL-10 receptor on γδ T cells does

not alter their IL-17A production, in turn, clearly indicates that the IL-17A production in these innate lymphocytes is not directly controlled by IL-10.

IL-10R expression on Tregs is obligatory to inhibit IL-17A production in $\gamma\delta$ T cells

To further investigate whether Treg or non-Treg CD4 T cells are capable to react to IL-10 and thus potentially control $\gamma\delta$ T cells, we next analyzed expression levels of the IL-10 receptor in double reporter *Fir-tiger* mice. On day 7 following tMCAO, we observed a significantly higher expression of the IL-10 receptor in the Foxp3⁺ compared to Foxp3⁻ compartment (Figure 6A), further supporting that IL-10 can activate Tregs.

To further test whether Tregs are controlling $\gamma\delta$ T cells in ischemic brains and if IL-10 confers Tregs the ability to inhibit the IL-17A production in $\gamma\delta$ T cells, we bred *Foxp3^{Cre}* x *Il10Ra^{fl/fl}* mice to generate a Treg-selective deletion of the IL-10R α . Seven days following tMCAO we analyzed the frequency of IL-17A⁺ CD4 and $\gamma\delta$ T cells derived from ischemic hemispheres (Figure 6B and C). Upon deletion of the IL-10R α in Tregs, CD4⁺ T cells displayed no skewing towards a Th17 phenotype. However, we observed a significantly increased IL-17A production in $\gamma\delta$ T cells in *Foxp3^{Cre}* x *Il10Ra^{fl/fl}* mice (Figure 6C). In summary, our data show that IL-10, which by itself had no direct effect on $\gamma\delta$ T cells, is able to suppress the IL-17A production of $\gamma\delta$ T cells via Tregs.

Discussion

IL-17A is seen as a decisive factor for the early detrimental postischemic inflammatory response following stroke [3, 4]. One of the main effects of the IL-17A axis in ischemic brains is the induction of CXC-chemokines, which lead to a massive infiltration of neutrophils [4, 22]. However, in stroke, the initial IL-17A driven inflammatory response is short-lasting, indicating that the sterile inflammatory response is effectively controlled [4].

Mechanisms that limit the detrimental IL-17A response in stroke are not well understood. In this study, we tested whether the anti-inflammatory cytokine IL-10 controls the early IL-17A response in T cells following experimental stroke. We found that IL-10 expressed by invading leukocytes plays a key role in the regulation of the IL-17A production in both in $\alpha\beta$ and $\gamma\delta$ T cells in ischemic hemispheres. Interestingly, the effects of IL-10 on CD4⁺ and $\gamma\delta$ T cells differ. While IL-10 directly inhibits the IL-17A production in CD4⁺ T cells, signaling via the IL-10R endows Tregs with the capacity to downregulate IL-17A in $\gamma\delta$ T cells. However, abrogation of direct IL-10R signaling in $\gamma\delta$ T cells does not impact their IL-17A production, underlining that IL-10 controls the IL-17A production of $\alpha\beta$ and $\gamma\delta$ T cells differentially.

IL-10 has been implicated in several studies to provide neuroprotection in the brain [6, 24]. However, in stroke, only Benakis et al. found that IL-10 is protective through the control of IL-17A pathways [22]. The authors showed that the modulation of the gut microbiome diminished the frequency of IL-17A⁺ $\gamma\delta$ T cells in the gut in an IL-10 dependent manner which was associated with an attenuation of the IL-17A driven postischemic inflammatory reaction in the brain.

Beyond these observations, our data indicate that IL-10 not only controls the IL-17A driven postischemic inflammatory response in the peripheral immune compartment but also directly in the ischemic brain. We showed that intracerebral injections of recombinant IL-10 reduce the frequency of IL-17A producing $\gamma\delta$ T cells in the ischemic brain following tMCAO.

A strict regulation of IL-17A level in the CNS may also have physiological functions. Recent studies showed that IL-17A⁺ $\gamma\delta$ T cells populate the meninges from perinatal stages in order to build a self-renewing pool which controls neuronal signaling, behavior, and cognition under homeostatic conditions [25]. The fact that IL-17A can have important homeostatic functions in the brain on the one hand and be proinflammatory on the other hand is explained by the physiology of IL-17A signaling. Outcomes of the IL-17A receptor activation depend on the cooperative stimulation by other cytokines, such as IL-1 β and TNF- α [26] which are

upregulated in the ischemic brain tissue and can lead to synergistic activation of downstream signaling pathways [20]. IL-17A alone is only a modest activator of inflammatory pathways. During post stroke inflammation, IL-10 seems to directly inhibit the IL-17A production in Th17 cells in ischemic brains, which is a known mechanism from other models of autoimmunity [8]. Accordingly, the targeted deletion of the IL-10R in CD4⁺ T cells leads to increased frequencies in Th17 cells in ischemic brains. In contrast to Th17 cells, IL-10 does not directly control IL-17A⁺ $\gamma\delta$ T cells. Signaling via the IL-10R rather endows Tregs with the capacity to downregulate IL-17A in $\gamma\delta$ T cells. In this study, we did not identify the molecular factor, which enables IL-10R⁺ Tregs to inhibit IL-17A⁺ $\gamma\delta$ T cells. Possible candidates include the B- and T-lymphocyte attenuator and the programmed cell death protein 1 as it has been reported that they negatively regulate IL-17A in a model of psoriatic-like skin inflammation [27].

Translating findings from animal models to human patients is still challenging. However, we previously showed in autptic brain tissue that the expression of IL-17A in T cells goes along with a massive neutrophil infiltration, indicating that the IL-17A-neutrophil axis contributes to human pathophysiology as well [4]. Regarding the potent and pleiotropic immunosuppressive properties of IL-10 on the systemic immune compartment, IL-10 itself might not represent an ideal therapeutic target in stroke. Stroke patients have a significantly increased susceptibility for pneumonia due to aspirations and an immunosuppressive state [28]. Our data show that the worse outcome in *Il10*^{-/-} mice is largely driven by increased IL-17A levels in the brain, which can be counteracted via neutralization of IL-17A. With this in mind, a targeted antagonization of IL-17A seems promising, irrespective of individual IL-10 levels pre-stroke. Besides, several experimental studies have already shown that the neutralization of IL-17A is protective in stroke [3, 4, 22]. Furthermore, IL-17A antibodies are already approved in the treatment of psoriasis [29] and severe adverse effects on the immune status have, thus far, not been apparent. However, prior to a clinical application of IL-17A antibodies in stroke, long-term data on the effects of IL-17A neutralization in both sexes should be collected in order to adjust for potential

negative effects. This is necessary to exclude that the neutralization has negative long-term consequences on post-stroke regeneration by inhibiting the recently discovered beneficial IL-17A effects on memory and behavior [25].

Conclusion

In conclusion, our study provides additional insights into the complex interplay between different T cell subpopulations during post-stroke neuroinflammation. We have shown that IL-10 is pivotal for the control of pathogenic IL-17A⁺ $\alpha\beta$ and $\gamma\delta$ T cells and that an excessive IL-17A production is harmful. Thus, our study improves the understanding of the IL-17A dependent postischemic inflammatory response and may pave the way towards new immunomodulatory therapies for stroke treatment.

Additional file 1: Supplementary Figures 1 and 2

List of Abbreviations

IL-17A: Interleukin-17A; tMCAO: transient Middle Cerebral Artery Occlusion; IL-10: Interleukin-10; IL-10R: Interleukin-10-receptor; Tregs: regulatory T cells; T_R1: T regulatory type 1 cells; WT: wildtype; BSA: bovine serum albumin; MRI: Magnetic Resonance Imaging; *Il10*^{-/-}: Interleukin-10 knockout; *FIR*: *Foxp3-IRES-mRFP*; *tiger*: *Il10-IRES-GFP-enhanced reporter*; CXCL-1: C-X-C Motif Chemokine ligand 1; MMP-3: Matrix Metalloproteinase-3

Declarations

Ethics Approval

All animal experiments were approved by local animal care committees (Behörde für Justiz und Verbraucherschutz der Freien und Hansestadt Hamburg Lebensmittelsicherheit und Veterinärwesen, Hamburg, Germany, permission number: G59/17, N079/19).

Consent for publication

Not applicable.

Availability of data and materials

The datasets used and analyzed during the current study are available from the corresponding author on reasonable request.

Competing interests

The authors declare that they have no competing interests.

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Author Contributions

MG, MP, TM, and SH designed the study. MP, BHC, PL, JV, EJ, LJ and LB conducted various experiments and acquired and analyzed data. TB, BR, IS, KD, IP, RAF, YK, TR and CG

provided important reagents and valuable feedback on the manuscript. MG and MP wrote and edited the manuscript. All authors read and approved the final manuscript.

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Disclosures

None.

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Figures and Figure legends

Figure 1

Figure 1: *I110* deficiency leads to increased infarct sizes, enhanced brain atrophy, and poorer long-term outcome following tMCAO.

MRI was used to quantify infarct volume at day 3 (A) and cortical atrophy volume at day 14 (C) after tMCAO in WT control and *I110*^{-/-} mice (representative T2 image). (B) Triphenyltetrazolium chloride (TTC) staining was used for the evaluation of infarct volume at

day 7 (representative TTC brain slices). (D) Composite neurological score was performed on day 14 after tMCAO. (E) *Il10/GFP*-positive cells were visualized in the penumbra area of the ischemic hemisphere by GFP counterstaining in *FIR-tiger* mice 14 days after tMCAO (green, *Il10/GFP*-positive cells; red, CD45-positive cells; blue, 40,6-diamidino-2-phenylindole nuclear staining; scale bar 20 μ m). Infarct data are presented as mean $\pm$ SEM of 8-10 WT and 6-10 *Il10*^{-/-} mice (A and B), atrophy data as mean $\pm$ SEM of 11 WT and 13 *Il10*^{-/-} mice (C), and neurological score of 12 WT and 11 *Il10*^{-/-} mice (C). Statistical significances were analyzed by Student *t* test (A-C) and Mann-Whitney *U* test (D). *P<0.05, **P<0.01 and ***P<0.001.

Figure 2

Figure 2: IL-10 from invading immune cells is neuroprotective

MRI was used to quantify infarct volume at day 3 and cortical atrophy volume at day 14 (A, B) after tMCAO in the chimeric mice (representative T2 image). (C) Composite neurological score was performed on day 14 after tMCAO. (D) Relative gene expression of *Il10* in brain resident

CD45^{intermed}/CD11b⁺ microglia and central nervous system-infiltrating CD45^{high}/CD11b⁺/CD11c⁻/MHCII⁻/Ly6g⁻/F4/80⁺ macrophages purified 3, 7, and 14 days after tMCAO by fluorescence activated cell sorting from ischemic hemispheres of C57Bl/6 mice. Expression levels were normalized to corresponding levels of microglia after sham surgery or blood macrophages. (E) Flow cytometric analysis of IL-10 produced by CD4⁺ Tregs (Foxp3⁺) or non-Tregs (Foxp3⁻). Infarct and atrophy data are presented as mean $\pm$ SEM of 16 WT mice which received *Il10*^{-/-} and 10-11 *Il10*^{-/-} mice which received WT bone marrow cells (A, B), neurological score as mean $\pm$ SEM of 15 WT mice which received *Il10*^{-/-} and 10 *Il10*^{-/-} mice which received WT bone marrow cells (C), RT-qPCR gene expression data as mean $\pm$ SEM of 3-7 (D) and Flow cytometric data as mean of $\pm$ SEM of 5-10 mice in each group (E). Significances analyzed by Student *t* test (A, B), Mann-Whitney *U* test (C), and 1-way ANOVA with Bonferroni post hoc test (D, E). *P<0.05, **P<0.01 and ****P<0.0001. MΦ indicates macrophages; and MG, microglia.

Figure 3

Figure 3: IL-10 controls the IL-17A axis in the proinflammatory milieu of the postischemic brain

Relative gene expression of *Il23a* and *Il1b* in brain resident CD45^{intermed}/CD11b⁺ microglia (A) and central nervous system-infiltrating CD45^{high}/CD11b⁺/CD11c⁻/MHCII⁻/Ly6g⁻/F4/80⁺ macrophages (B) purified 3, 7 and 14 days after tMCAO by fluorescence activated cell sorting

from ischemic hemispheres. Expression levels were normalized to corresponding levels of splenic macrophages and microglia after sham operation. (C) Flow cytometric analysis of IL-17A produced by CD4⁺ and $\gamma\delta$ T cells isolated from ischemic hemispheres of WT controls and *Il10*^{-/-} mice 7 days after tMCAO. (D) Relative gene expression of *Cxcl1* and *Mmp3* in ischemic hemispheres of WT and *Il10*^{-/-} mice 3d after tMCAO. (E) Frequency of IL-17A producing $\gamma\delta$ T cells purified from ischemic brains was analyzed 3 days following tMCAO in control mice (BSA) and mice receiving IL-10 intracerebral 3h after tMCAO induction. (A, B) RT-qPCR gene expression data is represented as mean $\pm$ SEM of 4-7 WT, Flow cytometric data as mean $\pm$ SEM of 6 WT and 5 *Il10*^{-/-} (C), RT-qPCR gene expression data as mean $\pm$ SEM of 8 WT and 8 *Il10*^{-/-} mice per group (D) and flow cytometric data as mean $\pm$ SEM of 7 WT mice in each treatment group (E). Statistical significances were analyzed by 1-way ANOVA with Bonferroni post hoc test (A, B) and by Student t test (C-E). *P<0.05, **P<0.01, ***P<0.001, ****P<0.0001.

Figure 4

Figure 4: Neutralization of IL-17A abolishes the worse outcome of *Il10*^{-/-} mice

Triphenyltetrazolium chloride staining was used for evaluation of infarct volume at day 7 of IL-17A antibody (A) or IgG control (C) treated *Il10*^{-/-} and WT control mice. Bederson score was performed 7 days after tMCAO IL-17A antibody (B) or in IgG control (D) treated group. Infarct data are presented as mean $\pm$ SEM of 9-11 WT and 6-9 *Il10*^{-/-} mice per group. Bederson score as mean $\pm$ SEM of 9-11 WT and 7-10 *Il10*^{-/-} mice per group. Statistical significances were analyzed by Student t test (A, C) and Mann-Whitney *U* test (B, D). **P*<0.05, ***P*<0.01.

Figure 5

Figure 5: T cell-specific blockade of IL-10R leads to increased frequencies of IL-17A⁺ CD4⁺ and γδ T cells

Flow cytometric analysis of IL-17A produced by CD4⁺ (A) and γδ (B) T cells isolated from ischemic hemispheres of *CD4^{Cre}Il10Ra^{wt/wt}* controls and *CD4^{Cre}Il10Ra^{fl/fl}* 7 days after tMCAO. (C) Flow cytometric analysis of IL-10R (CD210) on IL-17A⁺ and IL-17A⁻ γδ T cells isolated from ischemic hemispheres of C57BL/6 mice 7 days after tMCAO. (D) Flow cytometric analysis of IL-17A produced by γδ T cells isolated from ischemic hemispheres of *TCRδ^{CreER}Il10Ra^{wt/wt}* controls and *TCRδ^{CreER}Il10Ra^{fl/fl}* 7 days after tMCAO. Flow cytometric data is represented as mean ± SEM of 9 *CD4^{Cre}Il10Ra^{wt/wt}* and 8 *CD4^{Cre}Il10Ra^{fl/fl}* (A, B), 6 WT (C) and 6 *TCRδ^{CreER}Il10Ra^{wt/wt}* and 7 *TCRδ^{CreER}Il10Ra^{fl/fl}* (D) mice per group. Statistical

Figure 6

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mechanism of protection from ischemic brain injury induced by IL-10. Flow cytometric data is presented as mean $\pm$ SEM of 6 *FIR-tiger* (A) and 6 *Foxp3^{Cre}Il10Ra^{wt/wt}* and 7 *Foxp3^{Cre}Il10Ra^{fl/fl}* mice per group (B, C). Statistical significances were analyzed by Student t test (A–C). **P<0.01.

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