

# **Neuroprotection against ischemic stroke requires a specific class of early responder T cells in mice**

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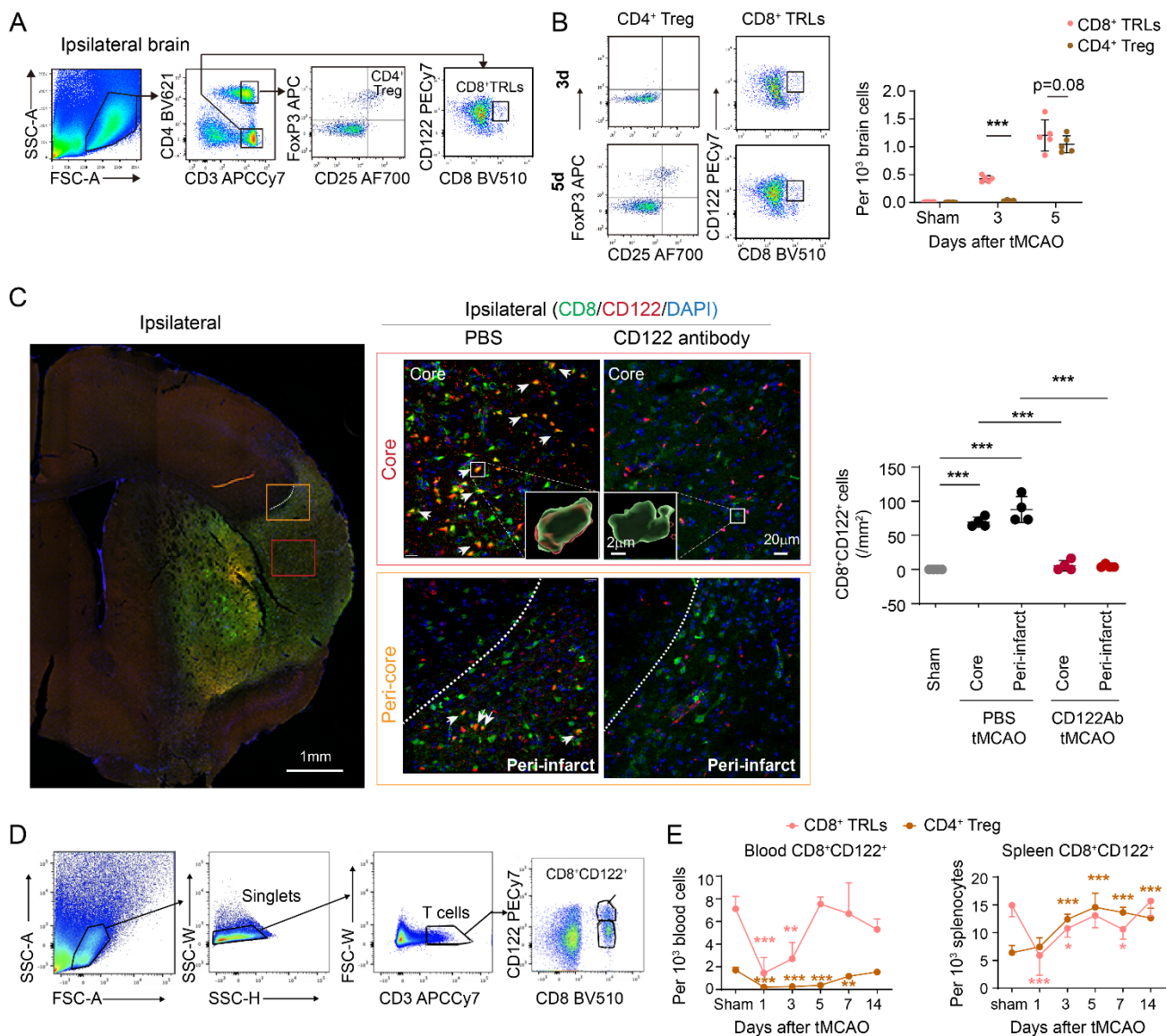
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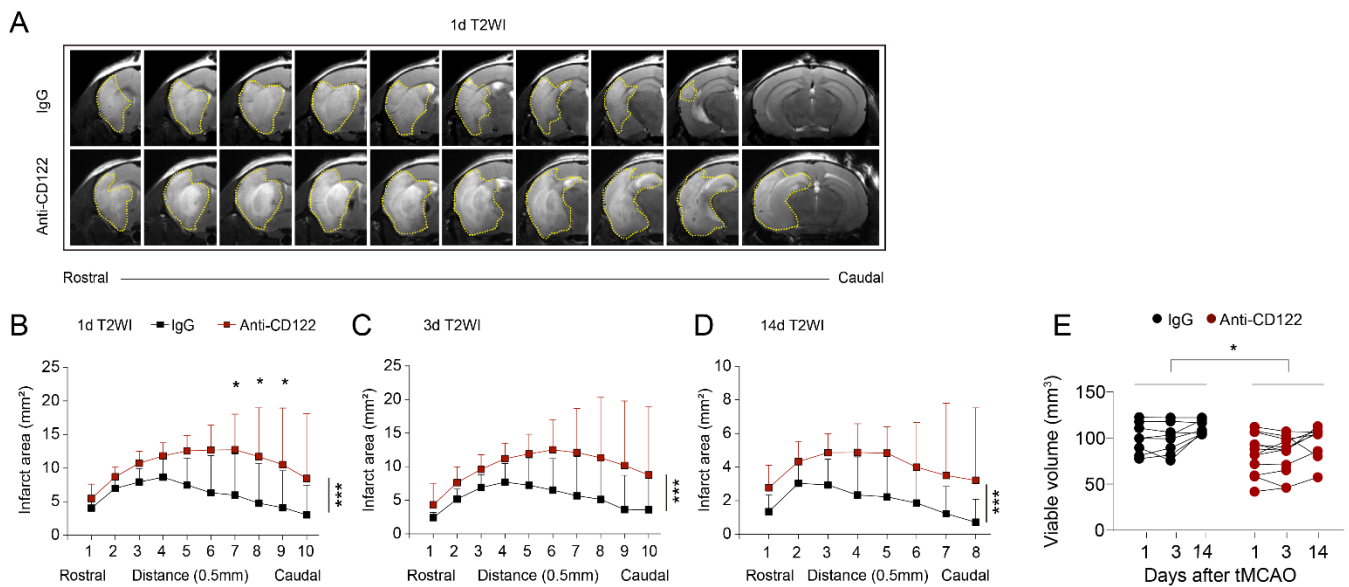
## Supplemental method:

**Antibodies used for flow cytometry:** The following antibodies were used: anti-CD45 PerCP-Cy5.5 (45-0451, Thermo Fisher Scientific, 1:400); anti-CD3 APC (17-0032, Thermo Fisher Scientific, 1:400); anti-CD3 eFluor450 (48-0031, Thermo Fisher Scientific, 1:400); anti-CD3 FITC (11-0031, Thermo Fisher Scientific, 1:400); anti-CD3 PerCP (100325, BioLegend, 1:400); anti-CD3 APC eFluor 780 (47-0032, Thermo Fisher Scientific, 1:400); anti-CD19 FITC (11-0193, Thermo Fisher Scientific, 1:400); anti-CD8 FITC (11-0081, Thermo Fisher Scientific, 1:400); anti-CD8 eFluor450 (48-0081, Thermo Fisher Scientific, 1:400); anti-CD8 BUV737 (612759, BD Biosciences, 1:400); anti-CD8 BV510 (563068, BD Biosciences, 1:400); anti-CD8 BV421 (100737, BioLegend, 1:400); anti-CD8 Pacific blue (MCD0828, Thermo Fisher Scientific, 1:400); anti-NK1.1 PE (12-5941, Thermo Fisher Scientific, 1:400); anti-NK1.1 APC (17-5941, Thermo Fisher Scientific, 1:400); anti-CD45.1 PerCP-Cy5.5 (110728, BioLegend, 1:400); anti-CD4 FITC (11-0041, Thermo Fisher Scientific, 1:400); anti-CD4 APC (17-0042, Thermo Fisher Scientific, 1:400); anti-CD4 Pacific blue (100428, BioLegend, 1:400); anti-CD25 Alexa Fluor700 (102024, BioLegend, 1:400); anti-CD122 PE-Cy7 (25-1222, Thermo Fisher Scientific, 1:400); anti-CD11b BUV737 (612801, BD Biosciences, 1:400); anti-CD11c BV421 (117329, BioLegend, 1:400); anti-Ly6G BUV395 (563978, BD Biosciences, 1:400); anti-F4/80 PE (157303, BioLegend, 1:400); anti-Foxp3 APC (17-5773, Thermo Fisher Scientific, 1:200); anti-IL-10 APC (17-7101, Thermo Fisher Scientific, 1:200); anti-IL-10 PE (12-7101, Thermo Fisher Scientific, 1:200); anti-HELIOS FITC (137214, BioLegend, 1:200); anti-CD49d BV510 (745007, BD Biosciences, 1:400); anti-CD103 APC (121413, BioLegend, 1:400); anti-CD45.1 PerCP-Cy5.5 (110728, BioLegend, 1:400); anti-LIFR PE (FAB5990P, R&D system, 1:100); anti-CXCR3 BV510 (745033, BD Biosciences, 1:200); anti-CCR6 BV421 (129817, BioLegend, 1:200); anti-CCR6 APC (129813, BioLegend, 1:200); anti-CXCR6 APC (151105, BioLegend, 1:200); anti-CCR5 PE Dazzle594 (359126, BioLegend, 1:200); anti-CCR7 BV605 (120125, BioLegend, 1:200); anti-CXCR5 PE Dazzle594 (145521, BioLegend, 1:200); anti-CCR4 PE (131203, BioLegend, 1:200); anti-CX3CR1 FITC (149019, BioLegend, 1:200). For flow cytometric staining of CCR10, anti-rabbit-BV421 (406410, BioLegend, 1:1000) was used as a secondary antibody after incubation with rabbit anti mouse CCR10 (ab1656, abcam, 1:400).

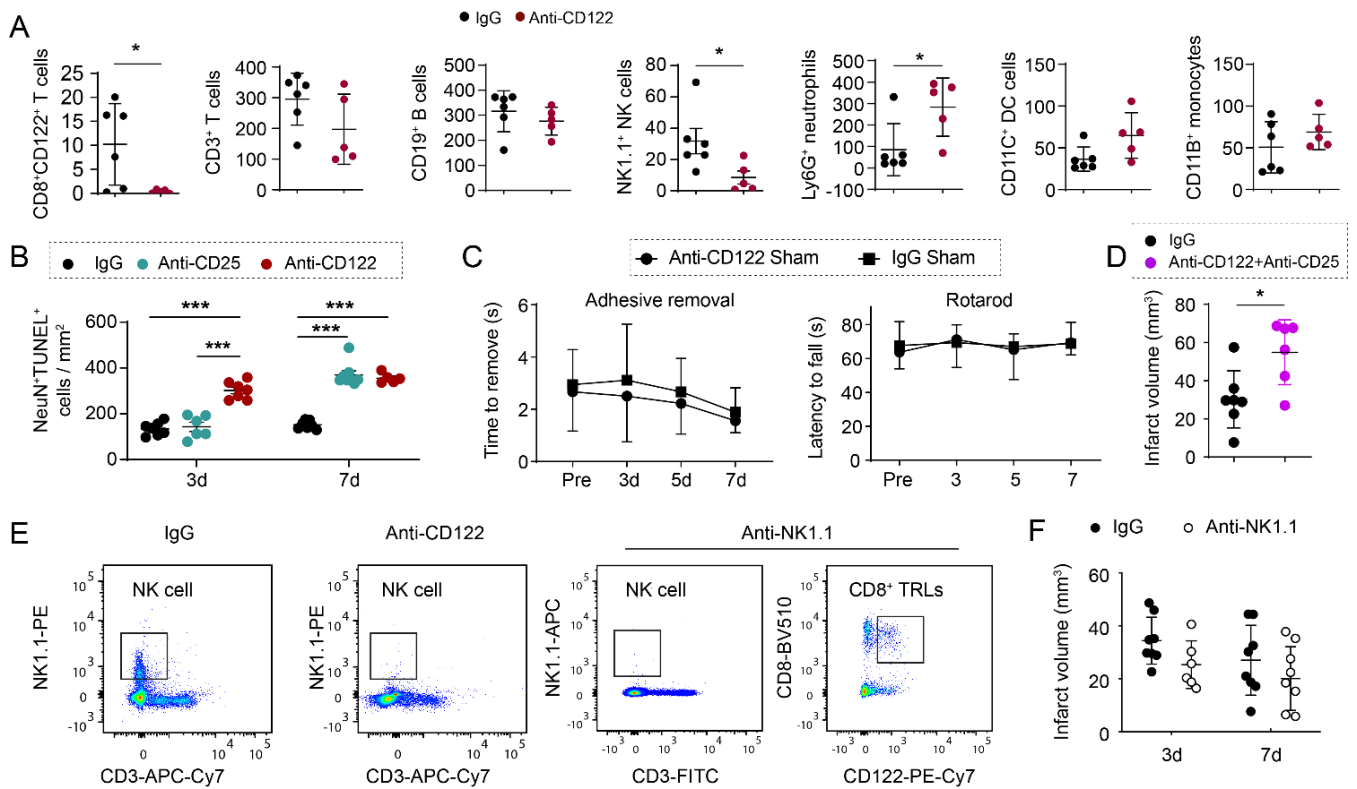


**Supplemental Fig. 1. Changes in the numbers of CD8<sup>+</sup>CD122<sup>+</sup> TRLs after tMCAO. (A)** Gating strategy for CD8<sup>+</sup>CD122<sup>+</sup> TRLs and CD4<sup>+</sup> Tregs in the ischemic brain. **(B)** Brain infiltration of CD4<sup>+</sup>CD25<sup>+</sup>Foxp3<sup>+</sup> Tregs and CD8<sup>+</sup>CD122<sup>+</sup> TRLs in sham and at 3d and 5d after tMCAO. n=4-5/group. Two-way ANOVA and *post hoc* Fisher's LSD **(C)** Triple-label immunofluorescence (CD8/CD122/DAPI) detects CD8<sup>+</sup>CD122<sup>+</sup> TRLs in the infarct core and peri-infarct areas (inner boarder) 5d after tMCAO. Anti-CD122 mAb treatment (100 μg, 2d prior to 60 min tMCAO) diminished the infiltration of CD8<sup>+</sup>CD122<sup>+</sup> TRLs into the ischemic brain. The inserts are 3D rendered of the selected cells using Imaris. The number of CD122<sup>+</sup>CD8<sup>+</sup> TRLs were quantified. n=4/group. One-way ANOVA and *post hoc* Bonferroni. **(D)** Gating strategy for CD8<sup>+</sup>CD122<sup>+</sup> TRLs in the blood and spleen. **(E)** Quantification of blood and spleen

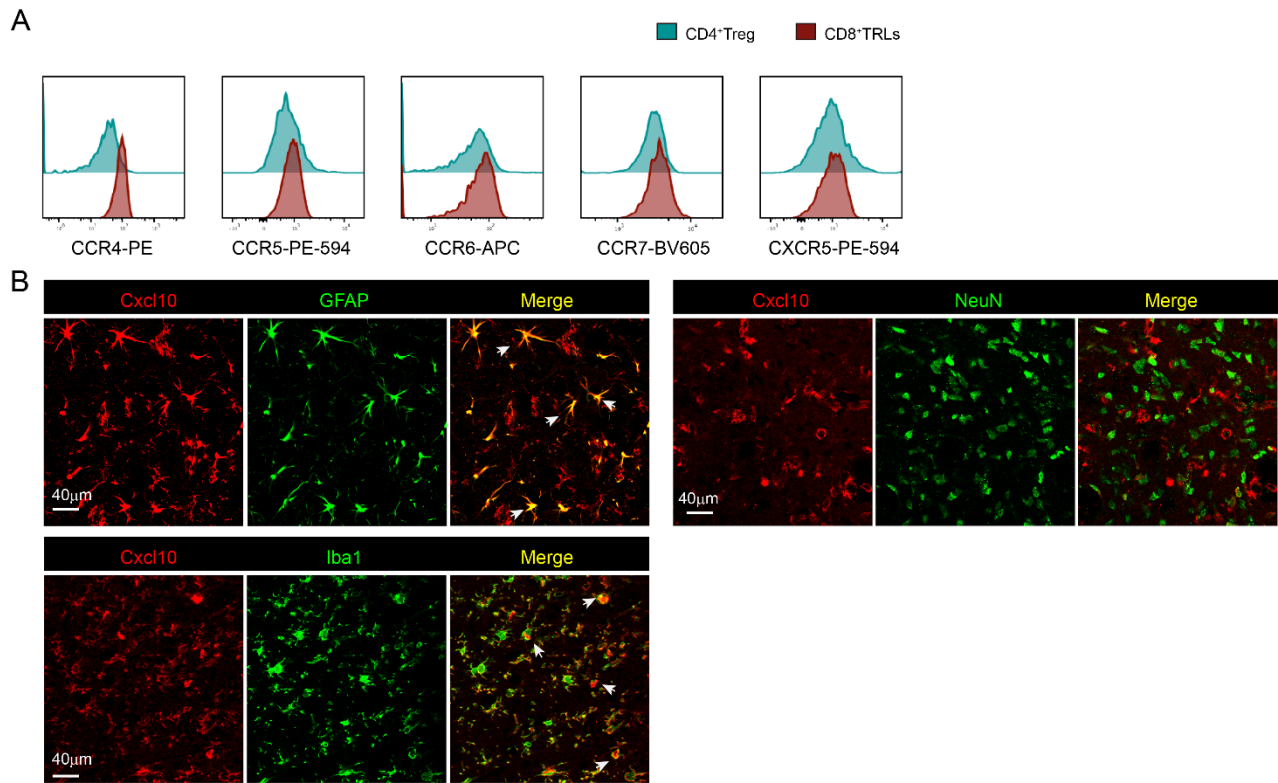
CD8<sup>+</sup>CD122<sup>+</sup> TRLs and CD4<sup>+</sup> Tregs at indicated time points after tMCAO. n=3-8/group. \* p<0.05, \*\*p<0.01, \*\*\*p<0.001 vs. sham controls. One way ANOVA and Dunnett.



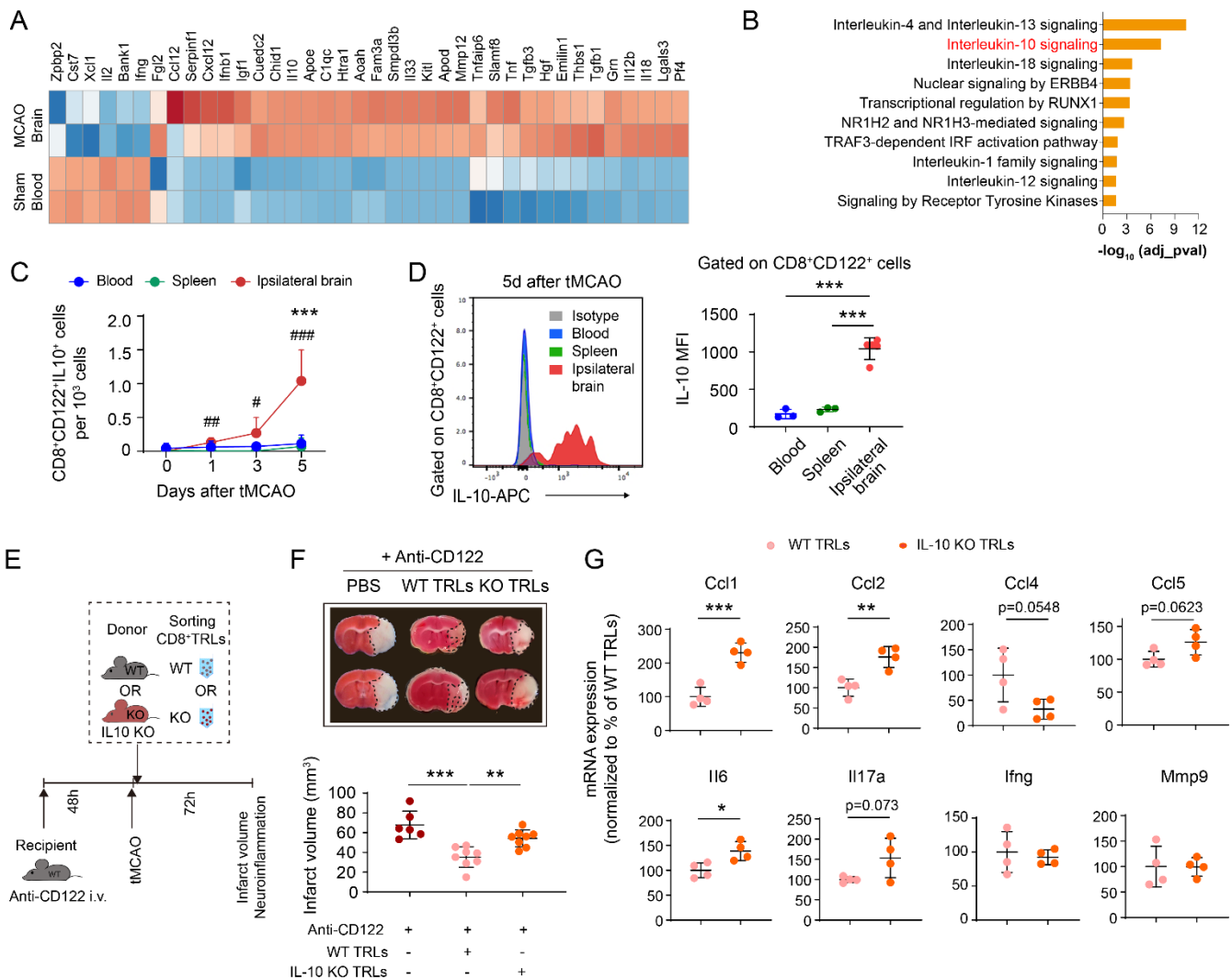
**Supplemental Fig. 2. Depletion of CD8<sup>+</sup> TRLs worsens stroke outcomes.** Mice were treated with CD122 mAb (100  $\mu$ g) or IgG isotype control (100  $\mu$ g) and subjected to tMCAO or sham operation.  $n=8$  mice for IgG group.  $n=12$  mice for anti-CD122 group. **(A)** Shown are rostral to caudal T2 weighted images (T2WI) taken at 1d after tMCAO in 10 coronal brain slices. *Dashed line* illustrates the boundary of brain infarct. **(B-D)** Infarct area was calculated on T2WI from 8-10 coronal sections encompassing the ischemic lesion at 1, 3, and 14d after tMCAO. Two-way ANOVA and *post hoc* Bonferroni. **(E)** Viable tissue volume was calculated on T2WI images at 1, 3, and 14d after tMCAO. Mixed-effects repeated measures ANOVA and *post hoc* Bonferroni. All data are mean $\pm$ SD, \* $p<0.05$ , \*\*\* $p<0.001$ .



**Supplemental Fig. 3. Impact of CD8<sup>+</sup>CD122<sup>+</sup> TRLs on stroke outcomes.** **(A)** Effect of anti-CD122 treatment on the number of immune cells in blood 3d after tMCAO. n=5-6/group. Two-tailed Student's t test. **(B)** Quantification of NeuN<sup>+</sup>TUNEL<sup>+</sup> neurons in the peri-infarct areas 3d and 7d after tMCAO in IgG Ab (100  $\mu$ g), anti-CD122 mAb (100  $\mu$ g), or anti-CD25 mAb (100  $\mu$ g) treated (ip, 2d before tMCAO) mice. n=5-8/group. One-way ANOVA and *post hoc* Dunnett. **(C)** Sensorimotor function was analyzed with the adhesive removal test and rotarod test in sham mice injected with anti-CD122 or isotype IgG. n=6/group. Two-way repeated measures ANOVA. **(D)** Quantification of infarct volumes in MAP2 stained sections 3d after tMCAO in anti-CD122 mAb + anti-CD25 mAb (100  $\mu$ g for each mAb) treated mice or IgG treated mice. n=6-7/group. Two-tailed Student's t test. **(E-F)** Depletion of NK cells with anti-NK1.1 mAb showed no effect on infarct volume 3d after 60 min tMCAO. Mice were treated with isotype IgG (100  $\mu$ g), anti-CD122 (100  $\mu$ g), or anti-NK1.1 mAb (100  $\mu$ g for NK cell depletion) 2d prior to 60 min tMCAO. **(E)** Flow cytometry plot showing NK cell depletion by anti-NK1.1 or anti-CD122 mAb 3d after stroke. **(F)** Infarct volume was assessed by MAP2 staining. n=6-8/group. Two-tailed Student's t test. All data are mean $\pm$ SD, \* p<0.05, \*\*\* p<0.001.



**Supplemental Fig. 4. Signals for CD8<sup>+</sup> TRL recruitment and activation.** Mice were subjected to 60 min tMCAO. **(A)** Flow cytometric analysis of CCR4<sup>+</sup>, CCR5<sup>+</sup>, CCR6<sup>+</sup>, CCR7<sup>+</sup> and CXCR5<sup>+</sup> cells among CD8<sup>+</sup> TRLs and CD4<sup>+</sup> Tregs in the ischemic brain. The plots are representative from 3 animals in each group. **(B)** Expression of CXCL10 (red) in astrocytes (GFAP, green), microglia (Iba1, green) and neurons (NeuN, green) was assessed in the ischemic brain by immunostaining 3d after stroke. The arrows point to selective double-labeled cells (CXCL10/GFAP, CXCL10/Iba1).

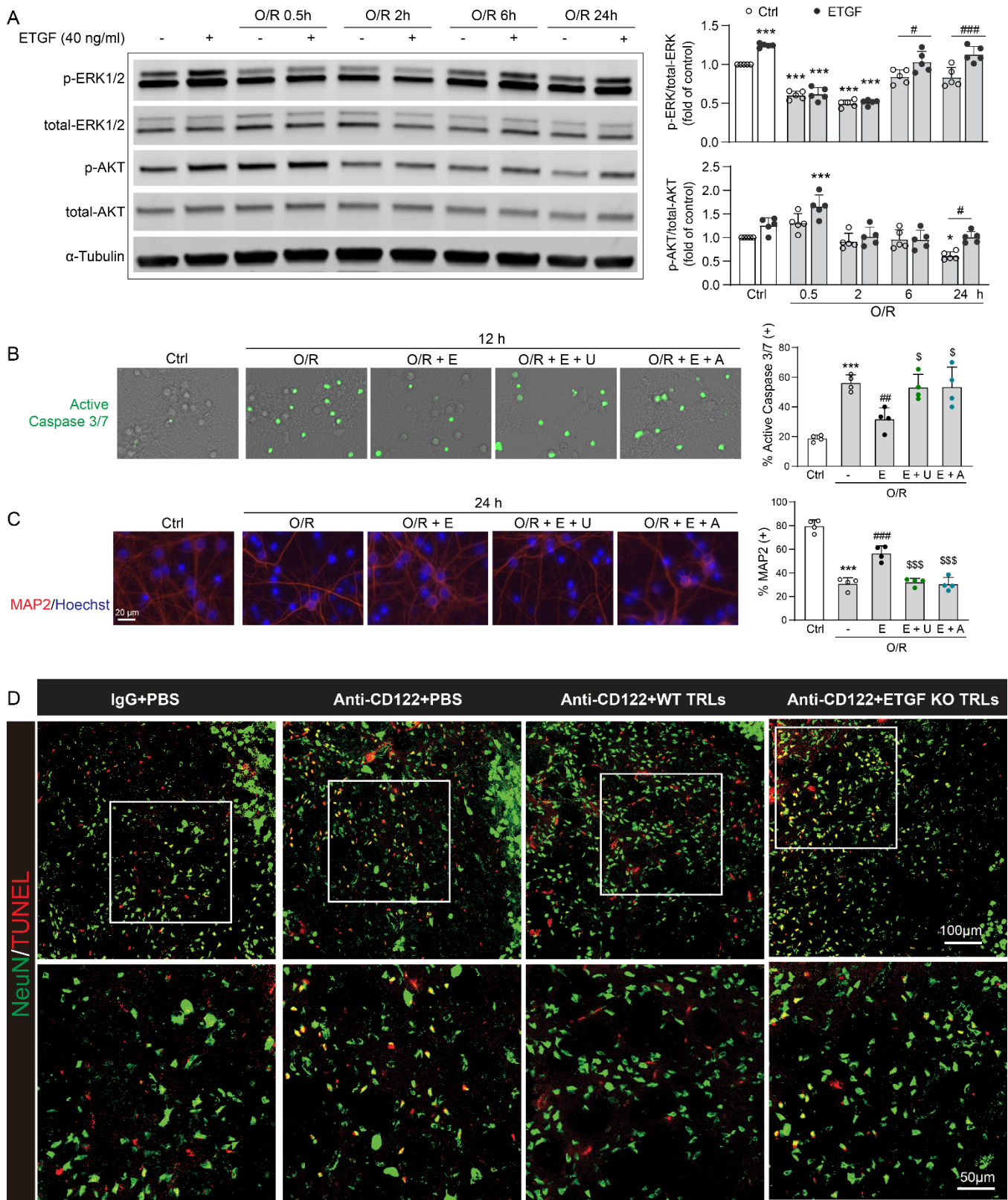


**Supplemental Fig. 5. CD8<sup>+</sup> TRLs ameliorate post-stroke inflammation in an IL-10 dependent manner.** (A) Heatmap showing immunoregulation-related DEGs that are higher (red) or lower (blue) in brain infiltrating TRLs from stroke mice vs. blood TRLs from sham mice in RNA-seq analysis. (B) IPA pathway analysis for signaling cascades that were activated in brain infiltrating CD8<sup>+</sup> TRLs. (C) The number of IL-10<sup>+</sup>CD8<sup>+</sup>CD122<sup>+</sup> TRLs in the blood, spleen and ipsilateral brains at indicated time points after tMCAO. n=3-8/group. \*\*\*p<0.001 vs. blood. #p<0.05, ##p<0.01, ###p<0.001 vs. spleen. One-way ANOVA and *post hoc* Bonferroni. (D) Quantification of the mean fluorescence intensity (MFI) of IL-10 in CD8<sup>+</sup> TRLs 5d after tMCAO. n=3-5/group. One-way ANOVA and *post hoc* Bonferroni. (E-F) Mice were treated with anti-CD122 mAb (100 µg) 2d prior to 60 min tMCAO. CD8<sup>+</sup> TRLs prepared from WT or IL-10 KO mice were transferred (1×10<sup>6</sup> cells, *i.v.*) into CD8<sup>+</sup> TRL-depleted mice 2h after stroke. Infarct volumes



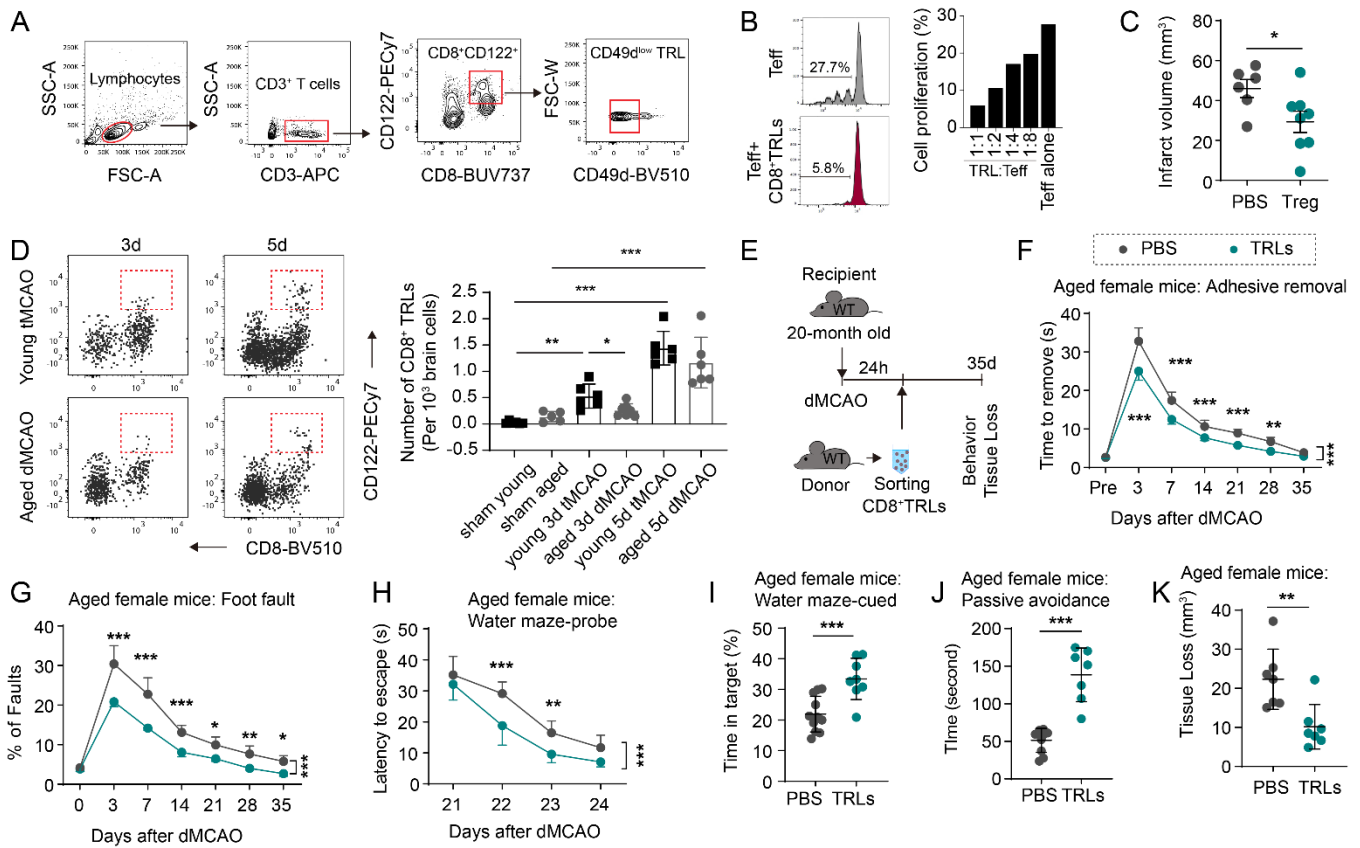
were quantified by TTC staining 3d after stroke. n=6-8/group. One-way ANOVA and *post hoc* Bonferroni.

**(G)** Brain inflammation was quantified by RT-PCR analysis for *Ccl1*, *Ccl2*, *Ccl4*, *Ccl5*, *Il6*, *Il17a*, *Ifng*, and *Mmp9* expression at 3d after tMCAO. n=4/group. Two-tailed Student's t test. All data are mean±SD, \*p<0.05, \*\*p<0.01, \*\*\*p<0.001.



**Supplemental Fig. 6. CD8<sup>+</sup> TRLs ameliorate neuronal injury in an ETGF dependent manner.** Primary cortical neurons were pre-treated with ETGF (E, 40 ng/ml) or PBS and subjected to OGD followed by oxygen-glucose restoration (O/R). Cells were collected at 0.5, 2, 6, 12 and 24h after O/R for Western

blotting, cell viability or cell death assays. **(A)** Western blotting for phosphorylated- or total-ERK1/2 and phosphorylated- or total-AKT were performed at the indicated time points after OGD. Data are from 5 independent experiments. **(B)** Apoptotic cell death was assessed by immunofluorescent staining for the active form of caspase-3/7 (green), and data were expressed as percentages of active caspase 3/7<sup>+</sup> cells among total number of cells at 12h after O/R. While ETGF (E, 40 ng/ml) reduced apoptosis after O/R, administration of AKT inhibitor VIII (A, 5  $\mu$ M) or ERK inhibitor U0126 (U, 5  $\mu$ M) abolished the neuroprotective effect of ETGF. Data are from 4 independent experiments. **(C)** Neuronal survival was measured by MAP2 staining 24h after O/R. While ETGF (E, 40 ng/ml) increased cell survival after O/R, administration of AKT inhibitor VIII (A, 5  $\mu$ M) or ERK inhibitor U0126 (U, 5  $\mu$ M) abolished the neuroprotective effect of ETGF. Data are from 4 independent experiments. All data are mean  $\pm$  SD. One way ANOVA and *post hoc* Bonferroni. \*\*\* $p$ <0.001 vs. non-OGD control. # $p$ <0.05, ##  $p$ <0.01, ### $p$ <0.001 vs. OGD control or as indicated. \$  $p$ <0.05, \$\$\$ $p$ <0.001 vs. OGD+ETGF. O/R: OGD/reperfusion. **(D)** Mice were treated with anti-CD122 mAb (100  $\mu$ g) 2d prior to 60 min tMCAO. CD8<sup>+</sup> TRLs prepared from WT or ETGF KO mice were transferred ( $1 \times 10^6$  cells, *i.v.*) into CD8<sup>+</sup> TRL-depleted mice 2h after tMCAO. Representative images demonstrating TUNEL (red) colabeling with the neuronal marker NeuN (green) in peri-infarct areas 3d after stroke. The images in white boxes in the upper panel were enlarged in the lower panel.



**Supplemental Fig. 7. Effect of CD8<sup>+</sup> TRLs on post-stroke brain injury and functional recovery. (A)**

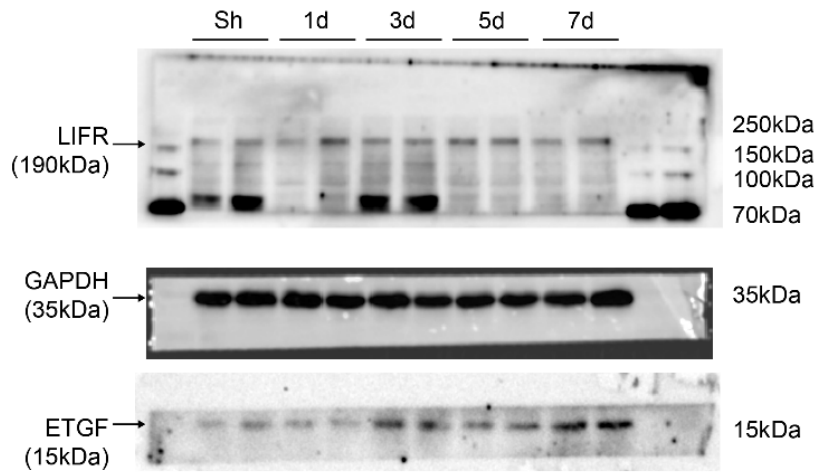
CD8<sup>+</sup> TRLs (CD8<sup>+</sup>CD122<sup>+</sup>CD49d<sup>low</sup>) were sorted from pooled spleens of healthy young donor mice. **(B)**

T effector cells (Teffs) suppression test. Teffs were labeled with CFSE (1 μM, 37°C, 10 min) and then plated at 2 × 10<sup>5</sup> cells per well in a U-bottom 96-well plate in the presence of CD3/CD28 activation beads to stimulate their proliferation. TRLs were added at a ratio of 1:1, 1:2, 1:4, or 1:8 to the number of Teffs, and cells were incubated for 3d. Suppression of Teff proliferation was determined by CFSE dilution on a flow cytometer. Left: Representative plots of suppression assay using CFSE-labeled Teffs incubated without or with CD8<sup>+</sup> TRLs (1:1). Right: The histogram indicates Teff proliferation rate. **(C)** CD8<sup>+</sup> TRLs (CD8<sup>+</sup>CD122<sup>+</sup>CD49d<sup>low</sup>) were sorted from pooled spleens of healthy aged male donor mice (20-month-old). Young male stroke mice were treated intravenously with aged CD8<sup>+</sup> TRLs or PBS at 2h after tMCAO. Quantification of MAP2 staining at 3d after tMCAO. n=6-8. **(D)** Comparison of the amounts of CD8<sup>+</sup> TRL infiltration into the brain in young (12-week-old) and aged (20-month-old) mice after tMCAO and dMCAO, respectively. n=4-6/group. **(E-K)** Aged female (20-month-old) stroke mice were treated intravenously with 1x10<sup>6</sup> FACS-sorted CD8<sup>+</sup> TRLs or PBS at 24h after dMCAO. n=7-11/group. **(E)** Experimental design for

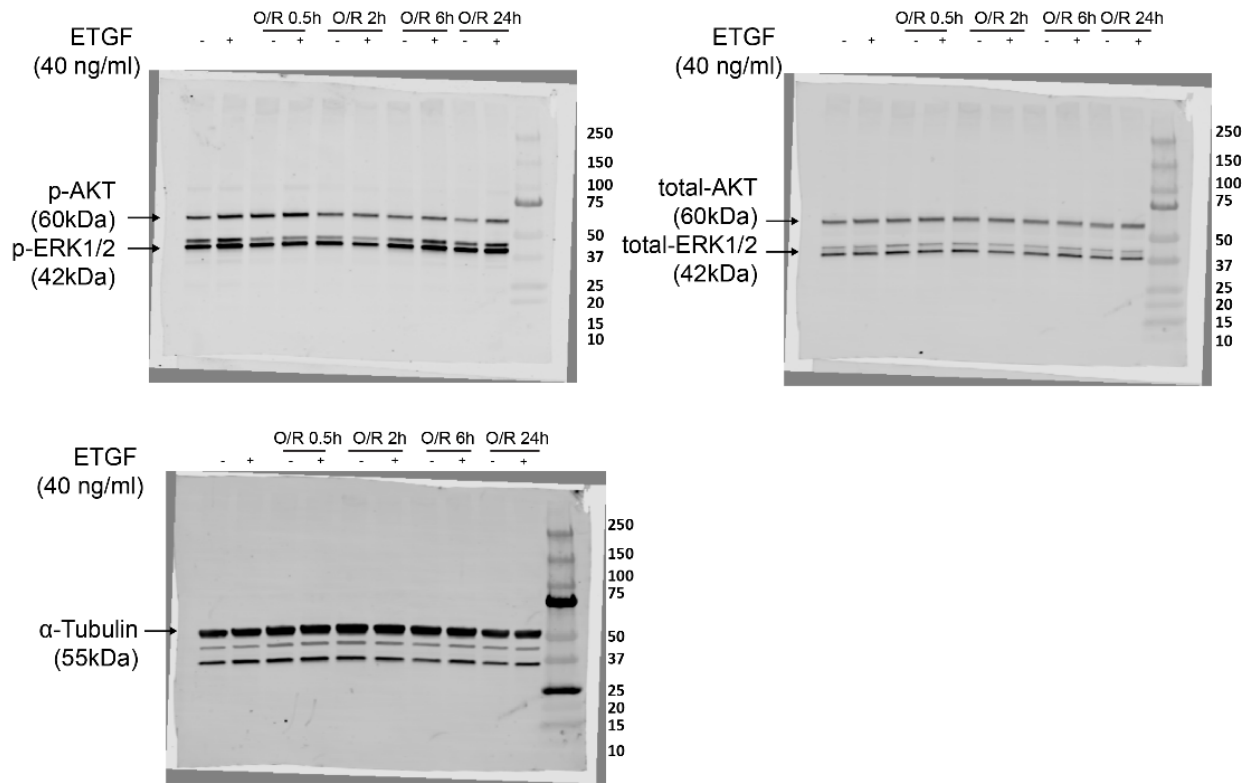
aged female mice. Donor mice (WT) provide CD8<sup>+</sup> TRLs. Recipient mice (20-month old WT) receive TRLs or PBS 24h after dMCAO. Behavior and tissue loss are assessed 35 days later. **(F)** Line graph showing the time to remove adhesive removal in aged female mice. TRLs significantly reduce the time to remove adhesive compared to PBS at various time points (\*\*\*). **(G)** Line graph showing the percentage of foot faults in aged female mice. TRLs significantly reduce the percentage of foot faults compared to PBS at various time points (\*\*\*). **(H)** Line graph showing the latency to escape in aged female mice. TRLs significantly reduce the latency to escape compared to PBS at various time points (\*\*\*). **(I)** Scatter plot showing the time in target (%) in aged female mice. TRLs significantly increase the time in target compared to PBS (\*\*\*). **(J)** Scatter plot showing the time (second) in aged female mice. TRLs significantly increase the time compared to PBS (\*\*\*). **(K)** Scatter plot showing the tissue loss (mm<sup>3</sup>) in aged female mice. TRLs significantly reduce tissue loss compared to PBS (\*\*).

Fig. 8J-P and Fig S7F-K. Sensorimotor dysfunction was assessed by the adhesive removal **(F)** and foot-fault tests **(G)** up to 35d after dMCAO. **(H-I)** Spatial learning and memory were assessed at 21d–25d after dMCAO in the Morris water maze. **(J)** Non-spatial memory was assessed at 35d after dMCAO using the passive avoidance test. **(K)** Quantification of MAP2 staining at 35d after dMCAO. \* $p < 0.05$ , \*\* $p < 0.01$ , \*\*\* $p < 0.001$ . Two-tailed Student's t test or Welch's t test (C, D, I, J, K), or two-way ANOVA and *post hoc* Bonferroni (F, G, H).

**A** Full unedited gel for Figure 5B



**B** Full unedited gel for Figure S6A



**Supplemental Fig. 8. Uncut western blot gel images.**

**Table S4. Mortality and exclusion rates.**

<b>Group</b>	<b>Surgery</b>	<b>Treatment</b>	<b>Mortality</b>	<b>Exclusion rate</b>
WT young	Sham	- or anti-CD122	0	0
WT young	tMCAO	- or IgG	6.04%	4.95%
WT young	tMCAO	Anti-CD122	8.86%	7.59%
WT young	tMCAO	CD122 Ab+WT TRLs	6.06%	6.06%
WT young	tMCAO	Anti-NK1.1	6.25%	6.25%
WT young	tMCAO	Anti-CD25	10%	5%
WT young	tMCAO	Anti-CD25+anti-CD122	14.29%	0
WT young	tMCAO	CD122 Ab+LIF-treated WT TRLs	0	0
WT young	tMCAO	CD122 Ab+LIF inhibitor-treated WT TRLs	0	0
WT young	tMCAO	CD122 Ab+ETGF	11.11%	11.11%
WT young	tMCAO	CD122 Ab+CXCL10 KO TRLs	0	0
WT young	tMCAO	CD122 Ab+CXCR3 KO TRLs	10%	0%
WT young	tMCAO	CD122 Ab+ETGF KO TRLs	14.29%	0%
WT young	tMCAO	CD122 Ab+IL10 KO TRLs	0	11.11%
WT young	tMCAO	Young TRLs	6%	4%
WT young	tMCAO	Aged TRLs	0%	0%
Rag1 KO	tMCAO	-	0	10%
Rag1 KO	tMCAO	Teff+TRLs	11.11%	0%
Rag1 KO	tMCAO	Teff	10%	10%

Rag1 KO	tMCAO	TRLs	0	11.11%
Aged WT male	Sham	-	0%	0%
Aged WT male	dMCAO	- or PBS	7.7%	0%
Aged WT male	dMCAO	TRLs	9%	0%
Aged female	dMCAO	PBS	9%	0%
Aged female	dMCAO	TRLs	0%	0%

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**Table S5. Primers for RT-PCR.**

<b>Gene</b>	<b>Forward primer</b>	<b>Reverse primer</b>
<i>Ccl1</i>	AGTTCTTGGCTCCACCAGAC	CATCCTGTATCCACACGGCA
<i>Ccl2</i>	TGACCCCAAGAAGGAATGGG	ACCTTAGGGCAGATGCAGTT
<i>Ccl4</i>	CTTCTGTGCTCCAGGGTTCTC	CTGCCTCTTTTGGTCAGGAATACCA
<i>Ccl5</i>	AAGTGTGTGCCAACCCAGAG	CCCATTTTCCCAGGACCGAG
<i>Cxcl9</i>	ATG AAGTCCGCTGTTCTTTTCC	GTCTCTTATGTAGTCTTCCTT G
<i>Cxcl10</i>	CTAGCTCAGGCTCGTCAGTT	CCCTTGGGAAGATGGTGGTTA
<i>Cxcl11</i>	ATGAACAGGAAGGTCACAGC	GATGTCACATGTTTTGACGC
<i>Il1a</i>	AAGACAAGCCTGTGTTGCTGAAGG	TCCCAGAAGAAAATGAGGTCGGTC
<i>Il6</i>	TCCTACCCCAACTTCCAATGCTC	TTGGATGGTCTTGGTCCTTAGCC
<i>Il10</i>	CCAAGCCTTATCGGAAATGA	TTTTCACAGGGGAGAAATCG
<i>Il17a</i>	CCTGGACTCTCCACCGCAA	TTCCCTCCGCATTGACACAG
<i>Mmp9</i>	CCAGCCGACTTTTGTGGTCT	TGGCCTTTAGTGTCTGGCTG
<i>Tnf</i>	AGAAGTTCCCAAATGGCCTC	CCACTTGGTGGTTTGCTACG
<i>Ifng</i>	ATGAACGCTACACACTGCATC	CCATCCTTTTGCCAGTTCCTC
<i>Tgfb1</i>	TGCGCTTGCAGAGATTA AAA	CGTCAA AAGACAGCCACTCA
<i>Tgfa</i>	CTTCTTGGTGCAAAGGCTCG	TCGACTGACGAATGGGCTTG
<i>Gapdh</i>	CCCTTAAGAGGGATGCTGCC	TACGGCCAAATCCGTT CACA

**Table S6. Statistical Analysis**

FIGURE 1	n	DATA STRUCTURE	TEST USED	STATISTIC	P VALUE
1A	Day0: n=4, Day1: n=8, Day3: n=5, Day5: n=5, Day7: n=6, Day14: n=7, Identical number for both groups at all time points.	Normal distribution	Unpaired t test	t(day0)=0.000, t(day1)=3.953, t(day3)=15.32, t(day5)=9.161, t(day7)=2.4, t(day14)=9.133	p(day0)>0.9999, p(day1)=0.0014, p(day3)<0.0001, p(day5)=0.0007, p(day7)=0.0394, p(day14)<0.0001
1B	CD103: n=4, IL-10: n=4, HELIOS: n=4, Identical number for both groups.	Normal distribution	Unpaired t test	t(CD103)=3.700, t(IL-10)=6.216, t(HELIOS)=10.21	p(CD103)=0.0101, p(IL-10)=0.0078, p(HELIOS)<0.0001
1E	IgG: n=8, Anti-CD122: n=12	Normal distribution	2-way repeated ANOVA, Bonferroni post hoc	F(1,18)=4.495	p=0.0481, p(baseline)=/, p(ischemia)>0.9999, p(reperfusion)=0.2743
1G	IgG: n=8, Anti-CD122: n(day1)=12; n(day3)=12; n(day14)=9	Normal distribution	Mixed-effects repeated measurement, Bonferroni post hoc	F(1,18)=6.265	p=0.0222, p(day1)=0.0473, p(day3)=0.0642, p(day14)=0.0281
1H	IgG: n=8, Anti-CD122: n=12	Normal distribution	2-way repeated ANOVA, Bonferroni post hoc	F(1,18)=10.62	p=0.0044, p(day1)=0.0090, p(day3)=0.0047
1J	IgG+PBS: n=6, Anti-CD122+PBS: n=6, Anti-CD122+CD8* TRL: n=6	Normal distribution	1-way ANOVA, Dunnett post hoc	F(2,15)=11.21	p=0.0011, p(IgG+PBS vs Anti-CD122+PBS)=0.0013, p(Anti-CD122+PBS vs Anti-CD122+CD8* TRL)=0.0027
1K	Day3: n(IgG)=7, n(anti-CD25)=7, n(anti-CD122)=8; Day7: n(IgG)=8, n(anti-CD25)=7, n(anti-CD122)=8	Normal distribution	1-way ANOVA, Dunnett post hoc	Day 3 F(2,19)=5.317, Day 7, F(2, 20)=6.403	Day3: p=0.0147, p(IgG vs anti-CD25)=0.4843, p(IgG vs anti-CD122)=0.0091; Day7: p=0.0071 p(IgG vs anti-CD25)=0.0130, p(IgG vs anti-CD122)=0.0096
1L	IgG: n=10, Anti-CD25: n=9, Anti-CD122: n=8	Normal distribution	2-way repeated ANOVA, Dunnett post hoc	F(2,24)=16.47	p<0.0001; Pre: p(IgG vs anti-CD25)=0.8867, p(IgG vs anti-CD122)=0.7962; day3: p(IgG vs anti-CD25)=0.4484, p(IgG vs anti-CD122)<0.0001; day5: p(IgG vs anti-CD25)=0.0001, p(IgG vs anti-CD122)<0.0001; day7: p(IgG vs anti-CD25)<0.0001, p(IgG vs anti-CD122)<0.0001
1M	IgG: n=10, Anti-CD25: n=10, Anti-CD122: n=8	Normal distribution	2-way repeated ANOVA, Dunnett post hoc	F(2,25)=9.829	p=0.0007; Pre: p(IgG vs anti-CD25)=0.9713, p(IgG vs anti-CD122)=0.9986; day3: p(IgG vs anti-CD25)=0.5733, p(IgG vs anti-CD122)<0.0001; day5: p(IgG vs anti-CD25)=0.0424, p(IgG vs anti-CD122)<0.0001; day7: p(IgG vs anti-CD25)=0.0053, p(IgG vs anti-CD122)=0.0167
1N	IgG: n=10, Anti-CD25: n=9, Anti-CD122: n=8	Normal distribution	2-way repeated ANOVA, Dunnett post hoc	F(2,24)=7.172	p=0.0036; Pre: p(IgG vs anti-CD25)=0.8974, p(IgG vs anti-CD122)=0.9961; day3: p(IgG vs anti-CD25)=0.9995, p(IgG vs anti-CD122)=0.0328; day5: p(IgG vs anti-CD25)=0.0006, p(IgG vs anti-CD122)=0.0015;

					day7: p(IgG vs anti-CD25)=0.0017, p(IgG vs anti-CD122)=0.3357
FIGURE 2	n	DATA STRUCTURE	TEST USED	STATISTIC	P VALUE
2D	CD4 <sup>+</sup> Treg: n=3, CD8 <sup>+</sup> TRL: n=3	Normal distribution	Unpaired t test	t=15.48	p=0.0001
FIGURE 3	n	DATA STRUCTURE	TEST USED	STATISTIC	P VALUE
3A	1d: sham=3, Cxcl9=3, Cxcl10=3, Cxcl11=3; 3d: sham=3, Cxcl9=3, Cxcl10=3, Cxcl11=3	Normal distribution	Unpaired t test	1d: t(CXCL9)=4.370, t(CXCL10)=2.735, t(CXCL11)=2.067; 3d: t(CXCL9)=4.158, t(CXCL10)=3.545, t(CXCL11)=5.532	1d: p(CXCL9)=0.012, p(CXCL10)=0.0522, p(CXCL11)=0.1076 ; 3d: p(CXCL9)=0.0142, p(CXCL10)=0.0239, p(CXCL11)=0.0052
3B	Blood: n(sham)=6, n(tMCAO)=6; Brain: n(sham)=6, n(tMCAO)=5	Normal distribution	Unpaired t test	Blood: t=1.431; Brain: t=2.691	p(blood)=0.1828; p(brain)=0.0248;
3E	n=6 for both groups	Normal distribution	Unpaired t test	t=5.980	P=0.0001
3F	WT: n=3; Cxcr3 KO: n=3	Normal distribution	Unpaired t test	t=3.228	p=0.0321
3H	IgG+PBS: n=8; Anti-CD122: n(PBS)=8, n(WT TRL)=6, n(Cxcr3 KO TRL)=6	Normal distribution	1-way ANOVA, Bonferroni post hoc	F(3,24)=13.62	p<0.0001; p(IgG+PBS vs Anti- CD122+PBS)<0.0001, p(Anti-CD122+PBS vs WT TRL)=0.0006, p(IgG+PBS vs Anti- CD122+WT TRL)>0.9999, p(IgG+PBS vs Anti- CD122+CXCR3 KO TRL)=0.0045, p(Anti-CD122+WT TRL vs Anti- CD122+CXCR3 KO TRL)=0.024, p(Anti-CD122+PBS vs Anti- CD122+CXCR3 KO TRL)>0.9999
3I	TNFα: n(ctrl)=3, n(Teff)=3, n(Teff+WT)=3, n(Teff+CXCR3 KO)=3; IL-4: n(ctrl)=3, n(Teff)=3, n(Teff+WT)=3, n(Teff+CXCR3 KO)=3	Normal distribution	1-way ANOVA, Bonferroni post hoc	TNFα: F(3,8)=974, IL-4:F(3,8)=313	p(TNFα)<0.0001, p(Teff vs Teff+WT)<0.0001, p(Teff vs Teff+CXCR3 KO)=0.0002, p(Teff+WT vs Teff+CXCR3 KO)=0.0592; p(IL-4)<0.0001, p(Teff vs Teff+WT)=0.0164, p(Teff vs Teff+CXCR3 KO)=0.0044, p(Teff+WT vs Teff+CXCR3 KO)>0.9999;
FIGURE 4	n	DATA STRUCTURE	TEST USED	STATISTIC	P VALUE
4D(II1a)	1d: sham=7, IgG=5, anti-CD122=3; 3d: sham=3, IgG=3, anti-CD122=3	Normal distribution	2-way ANOVA, Bonferroni post hoc	F(2,18)=22.96	p<0.0001; 1d: p(sham vs IgG)=0.0005, p(sham vs anti-CD122)=0.0009, p(IgG vs anti-CD122)>0.9999; 3d: p(sham vs IgG)>0.9999, p(sham vs anti-CD122)=0.0002, p(IgG vs anti-CD122)=0.0007
4D(Tnf)	1d: sham=7, IgG=5, anti-CD122=3; 3d: sham=3, IgG=3, anti-CD122=3	Normal distribution	2-way ANOVA, Bonferroni post hoc	F(2,18)=18.93	p<0.0001; 1d: p(sham vs IgG)=0.0033, p(sham vs anti-CD122)=0.0056, p(IgG vs anti-CD122)>0.9999; 3d: p(sham vs IgG)=0.2952, p(sham vs anti-CD122)=0.0004, p(IgG vs anti-CD122)=0.0183
4D(Ifnγ)	1d: sham=7, IgG=4, anti-CD122=3; 3d: sham=3, IgG=3, anti-CD122=3	Normal distribution	2-way ANOVA, Bonferroni post hoc	F(2,17)=19.30	p<0.0001; 1d: p(sham vs IgG)=0.6568, p(sham vs anti-CD122)=0.2584, p(IgG vs anti-CD122)>0.9999; 3d: p(sham vs IgG)>0.9999, p(sham vs anti-CD122)<0.0001, p(IgG vs anti-CD122)<0.0001
4D(II6)	1d: sham=7, IgG=5, anti-CD122=3;	Normal distribution	2-way ANOVA, Bonferroni post hoc	F(2,18)=8.730	P=0.0022;

	3d: sham= 3, IgG=3, anti-CD122=3				1d: p(sham vs IgG)=0.1745, p(sham vs anti-CD122)=0.0694, p(IgG vs anti-CD122)>0.9999; 3d: p(sham vs IgG)>0.9999, p(sham vs anti-CD122)=0.0102, p(IgG vs anti-CD122)=0.0646
4D(II10)	1d: sham=7, IgG=4, anti-CD122=3; 3d: sham= 3, IgG=3, anti-CD122=3	Normal distribution	2-way ANOVA, Bonferroni post hoc	F(2,17)=1.691	p=0.2139; 1d: p(sham vs IgG)=0.7743, p(sham vs anti-CD122)=0.0521, p(IgG vs anti-CD122)=0.5208; 3d: p(sham vs IgG)>0.9999, p(sham vs anti-CD122)>0.9999, p(IgG vs anti-CD122)>0.9999
4D(Tgfb1)	1d: sham=7, IgG=4, anti-CD122=3; 3d: sham= 3, IgG=3, anti-CD122=3	Normal distribution	2-way ANOVA, Bonferroni post hoc	F(2,17)=9.128	p=0.0020; 1d: p(sham vs IgG)=0.0019, p(sham vs anti-CD122)=0.0222, p(IgG vs anti-CD122)>0.9999; 3d: p(sham vs IgG)=0.4335, p(sham vs anti-CD122)=0.1945, p(IgG vs anti-CD122)>0.9999
4F	n=3 for both groups at each cell type	Normal distribution	Unpaired t test	t(neutrophil)=1.967; t(DC)=1.155; t(macrophage)=1.716; t(T cell)=1.138; t(B cell)=1.709	p(neutrophil)=0.1206; p(DC)=0.3125; p(macrophage)=0.1614; p(T cell)=0.3187; p(B cell)=0.1627
4G	naïve=9; Teff=8; Teff+TRL=8; TRL=8	Normal distribution	1-way ANOVA, Bonferroni post hoc	F(3,29)=9.948	p=0.0001; p( naïve vs Teff)=0.0202, p( naïve vs TRL)=0.0404, p(Teff vs Teff+TRL)=0.0054

**FIGURE 5**      **n**      **DATA STRUCTURE**      **TEST USED**      **STATISTIC**      **P VALUE**

5B	LIFR: n=4 ETGF: n=4	Normal distribution	1-way ANOVA, Dunnett post hoc	F(LIFR)=2.865; F(ETGF)=6.919	p(LIFR)=0.0602, p(1d vs sham)=0.1376, p(3d vs sham)=0.0467, p(5d vs sham)=0.0373, p(7d vs sham)=0.5236; p(ETGF)=0.0023, p(1d vs sham)=0.9987, p(3d vs sham)=0.0065, p(5d vs sham)=0.0215, p(7d vs sham)=0.0469
5C	Blood: sham=3, tMCAO=3; Spleen: sham=3, tMCAO=4; Brain=3	Normal distribution	1-way ANOVA, Dunnett post hoc	F(4,11)=124.9	P<0.0001; P<0.0001 for Brain tMCAO vs all other groups.
5E	n=6 for all groups	Normal distribution	1-way ANOVA, Bonferroni post hoc	F(3,20)=16.13	p<0.0001; p(blood sham vs blood tMCAO)>0.9999; p(brain sham vs brain tMCAO)<0.0001

**FIGURE 6**      **n**      **DATA STRUCTURE**      **TEST USED**      **STATISTIC**      **P VALUE**

6A(Liffr)	n(ctrl)=12; n(Cl)=6; n(IP)=7	Non normal distribution	Kruskal-Wallis test, Dunn's post hoc	Kruskal-Wallis statistic=19.69	p<0.0001; p(ctrl vs Cl)>0.9999, p(ctrl vs IP)<0.0001, p(Cl vs IP)=0.0011
6A(Etgff)	n(ctrl)=16; n(Cl)=8; n(IP)=16	Non normal distribution	Kruskal-Wallis test, Dunn's post hoc	Kruskal-Wallis statistic=29.46	p<0.0001; p(ctrl vs Cl)=0.2702, p(ctrl vs IP)=0.0002, p(Cl vs IP)<0.0001
6A(II10)	n(ctrl)=6; n(Cl)=6; n(IP)=12	Non normal distribution	Kruskal-Wallis test, Dunn's post hoc	Kruskal-Wallis statistic=16.13	p=0.0003; p(ctrl vs Cl)>0.9999, p(ctrl vs IP)=0.0123, p(Cl vs IP)=0.0009
6B	n(CL)=6; n(IP)=6; n(LIFab)=4	Normal distribution	1-way ANOVA, Bonferroni post hoc	ETGF: F(2,13)=8.904; IL10: F(2,13)= 4.550	p=0.0037; p(CL vs IL)=0.0051; p(CL vs LIFab)>0.9999; p(IP vs LIFab)=0.0241 IL10:

					$p=0.0318$ ; $p(\text{CL vs IP})=0.0446$ ; $p(\text{CL vs LIFab})>0.9999$ ; $p(\text{IP vs LIFab})=0.1279$
6E	$n=3$ for both groups	Normal distribution	Unpaired t test;	$t(\text{ETGF MFI})=5.383$ ; $t(\text{IL10 MFI})=3.472$ ;	$p(\text{ETGF MFI})=0.0058$ ; $p(\text{IL10 MFI})=0.0255$ ;
6F	$n(\text{PBS})=5$ ; $n(\text{non-treated CD8}^+ \text{ TRLs})=6$ ; $n(\text{LIF-treated CD8}^+ \text{ TRLs})=6$ ; $n(\text{LIFR inhibitor-treated CD8}^+ \text{ TRLs})=6$	Normal distribution	1-way ANOVA, Bonferroni post hoc	$F(3,19)=15.68$	$p<0.0001$ ; $p(\text{PBS vs non-treated treated CD8}^+ \text{ TRLs})=0.0088$ ; $p(\text{PBS vs LIF-treated treated CD8}^+ \text{ TRLs})<0.0001$ ; $p(\text{PBS vs LIFR inhibitor-treated treated CD8}^+ \text{ TRLs})=0.6027$ ; $p(\text{non-treated CD8}^+ \text{ TRLs vs LIF-treated CD8}^+ \text{ TRLs})=0.0452$ $p(\text{LIF-treated CD8}^+ \text{ TRLs vs LIFR inhibitor-treated treated CD8}^+ \text{ TRLs})=0.0004$
6G	For both 1d and 3d: $n(\text{PBS})=5$ ; $n(\text{non-treated CD8}^+ \text{ TRLs})=6$ ; $n(\text{LIF-treated CD8}^+ \text{ TRLs})=6$ ; $n(\text{LIFR inhibitor-treated CD8}^+ \text{ TRLs})=6$	Normal distribution	1-way ANOVA, Bonferroni post hoc	1d: $F(3,19)=11.16$ ; 3d: $F(3,19)=8.546$	1d: $p=0.0002$ ; $p(\text{PBS vs non-treated treated CD8}^+ \text{ TRLs})=0.0508$ ; $p(\text{PBS vs LIF-treated treated CD8}^+ \text{ TRLs})=0.0002$ ; $p(\text{PBS vs LIFR inhibitor-treated treated CD8}^+ \text{ TRLs})>0.9999$ ; $p(\text{non-treated CD8}^+ \text{ TRLs vs LIF-treated CD8}^+ \text{ TRLs})=0.0924$ ; $p(\text{LIF-treated CD8}^+ \text{ TRLs vs LIFR inhibitor-treated treated CD8}^+ \text{ TRLs})=0.0014$ 3d: $p=0.0008$ ; $p(\text{PBS vs non-treated treated CD8}^+ \text{ TRLs})=0.1495$ ; $p(\text{PBS vs LIF-treated treated CD8}^+ \text{ TRLs})=0.0011$ ; $p(\text{PBS vs LIFR inhibitor-treated treated CD8}^+ \text{ TRLs})=0.9681$ ; $p(\text{non-treated CD8}^+ \text{ TRLs vs LIF-treated CD8}^+ \text{ TRLs})=0.1417$ $p(\text{LIF-treated CD8}^+ \text{ TRLs vs LIFR inhibitor-treated treated CD8}^+ \text{ TRLs})=0.0034$
<b>FIGURE 7</b>	<b>n</b>	<b>DATA STRUCTURE</b>	<b>TEST USED</b>	<b>STATISTIC</b>	<b>P VALUE</b>
7C	$n=4$ for both groups	Normal distribution	unpaired t test	$t(\% \text{ETGF})=4.683$ ; $t(\text{ETGF MFI})=2.982$	$p(\% \text{ETGF})=0.0034$ ; $p(\text{ETGF MFI})=0.0246$
7F	control=5, OGD=7, OGD WT TRL CM=5, OGD KO TRL CM=7	Normal distribution	1-way ANOVA, Bonferroni post hoc	$F(3,20)=38.72$	$p<0.0001$ ; $p(\text{control vs OGD})<0.0001$ , $p(\text{control vs OGD WT TRL CM})=0.0001$ , $p(\text{control vs OGD KO TRL CM})<0.0001$ , $p(\text{OGD vs OGD WT TRL CM})=0.0075$ , $p(\text{OGD vs OGD KO TRL CM})>0.9999$ , $p(\text{OGD WT TRL CM vs OGD KO TRL CM})=0.0223$
7G	$N=6-7$	Normal distribution	1-way ANOVA, Bonferroni post hoc	$F(9,55)=56.63$	$p<0.0001$ ; $p(\text{OGD vs ctrl})<0.0001$ , $p(\text{OGD vs OGD ETGF})<0.001$ , $p(\text{OGD ETGF vs OGD ETGF U2.5})=0.0085$ , $p(\text{OGD ETGF vs OGD ETGF U5})<0.0001$ , $p(\text{OGD ETGF vs OGD ETGF A2.5})=0.0139$ , $p(\text{OGD ETGF vs OGD ETGF A5})<0.0001$ ,

7H	IgG=8, Anti-CD122=6, ETGF KO TRLs=6	Normal distribution	1-way ANOVA, Bonferroni post hoc	F(2,17)=3.816	p=0.0428; p(IgG vs Anti-Cd122)=0.0404, p(IgG vs KO TRL)=0.6034, p(Anti-Cd122 vs KO TRL)=0.5979
7I	WT=5, ETGF=7,	Normal distribution	Unpaired t test	t=2.357	p=0.0402
7K	n(IgG)=4, n(anti- CD122)=4, n(anti- CD122+WT)=6, n(anti-CD122+KO)=4	Normal distribution	1-way ANOVA, Bonferroni post hoc	cell density: F(3,14)=55.13; Tunel*/NeuN*%: F(3,14)=130.5	Tunel*/NeuN* density: p<0.0001, p(IgG vs anti-CD122)<0.0001, p(IgG vs CD122+WT)<0.9999, p(IgG vs anti-CD122+KO)<0.0001, p(anti-CD122 vs anti- CD122+WT)<0.0001, p(anti-CD122 vs anti- CD122+KO)>0.9999, p(anti-CD122+WT vs anti- CD122+KO)<0.0001; Tunel*/NeuN*%: p<0.0001; p(IgG vs anti-CD122)<0.0001, p(IgG vs CD122+WT)<0.9999, p(IgG vs anti-CD122+KO)<0.0001, p(anti-CD122 vs anti- CD122+WT)<0.0001, p(anti-CD122 vs anti- CD122+KO)>0.9999, p(anti-CD122+WT vs anti- CD122+KO)<0.0001
<b>FIGURE 8</b>	<b>n</b>	<b>DATA STRUCTURE</b>	<b>TEST USED</b>	<b>STATISTIC</b>	<b>P VALUE</b>
8C	PBS=6, TRL0.5=8, TRL1=7, TRL2=6	Normal distribution	1-way ANOVA, Dunnett post hoc	F(3,23)=2.577	p=0.0784, p(PBS vs TRLs 1)=0.0482, p(PBS vs TRLs 0.5)=0.1269, p(PBS vs TRLs 2)=0.7370
8D	PBS=6, TRL=7	Non normal distribution	Mann-Whitney test	U(0d)=10, U(1d)=2.5, U(2d)=6, U(3d)=13	p(0d)=0.1026; p(1d)=0.0035, p(2d)=0.0326, p(3d)=0.2861
8E	PBS=8, TRL=12	Normal distribution	2-way repeated ANOVA, Bonferroni post hoc	F(1,18)=21.43	p=0.0002; p(pre)>0.9999, p(3d)=0.0067, p(5d)=0.0156, p(7d)=0.0208, p(10d)=0.0010, p(14d)=0.0185
8F	PBS=8, TRL=12	Normal distribution	2-way repeated ANOVA, Bonferroni post hoc	F(1,18)=8.364	p=0.0097; p(pre)>0.9999, p(3d)=0.1572, p(5d)=0.0716, p(7d)=0.2492, p(10d)=0.0349, p(14d)=0.4164
8H	PBS=8, TRL=12	Normal distribution	2-way repeated ANOVA, Bonferroni post hoc	F(1,18)=5.497	p=0.0307; p(pre)>0.9999, p(10d)=0.7436, p(11d)=0.2756, p(12d)=0.0350, p(13d)=0.0264
8I	PBS=8, TRL=12	Normal distribution	Unpaired t test	t=1.090	p=0.2901
8J	PBS=10, TRLs=10	Normal distribution	2-way repeated ANOVA, Bonferroni post hoc	F(1,18)=28.93	p<0.0001, p(pre)>0.9999, p(3d)<0.0001, p(7d)<0.0001, p(14d)=0.1029, p(21d)=0.0041, p(28d)<0.0002, p(35d)=0.0003

8K	PBS=10, TRLs=10	Normal distribution	2-way repeated ANOVA, Bonferroni post hoc	F(1,18)=102.2	p<0.0001, p(pre)>0.9999, p(3d)<0.0001, p(7d)<0.0001, p(14d)<0.0001, p(21d)<0.0001, p(28d)=0.0244, p(35d)=0.2560
8L	PBS=10, TRLs=10	Normal distribution	2-way repeated ANOVA, Bonferroni post hoc	F(1,18)=115.6	p<0.0001, p(pre)>0.9999, p(3d)<0.0001, p(7d)<0.0001, p(14d)<0.0001, p(21d)<0.0001, p(28d)<0.0001, p(35d)<0.0001
8M	PBS=10, TRLs=10	Normal distribution	2-way repeated ANOVA, Bonferroni post hoc	F(1,18)=29.21	p<0.0001, p(21d)=0.0602, p(22d)<0.0001, p(23d)=0.0350, p(24d)=0.3231
8N	PBS=10, TRLs=10	Normal distribution	Unpaired t test	t=4.295	P=0.0004
8O	PBS=10, TRLs=10	Normal distribution	Unpaired t test	t=4.971	p<0.0001
8P	PBS=10, TRLs=10	Normal distribution	Unpaired t test	t=2.955	P=0.0085
<b>FIGURE S1</b>	<b>n</b>	<b>DATA STRUCTURE</b>	<b>TEST USED</b>	<b>STATISTIC</b>	<b>P VALUE</b>
S1 B	d0: CD8=4, CD4=5; d3: CD8=5, CD4=4; d5: CD8=5, CD4=5	Normal distribution	2-way ANOVA, Fisher's LSD	F(1,22)=12.53	P(group)=0.0018; D0: p(CD4 vs CD8)=0.9342 d3: p(CD4 vs CD8)=0.0004 d5: p(CD4 vs CD8)=0.0826
S1 C	n=4	Normal distribution	1-way ANOVA, Bonferroni post hoc	F(4,15)=72.32	P<0.0001, p(ctrl vs 5d core)<0.0001, p(ctrl vs 5d peri) <0.0001, p(5d core vs CD122Ab core)<0.0001 p(5d peri-infarct vs CD122Ab peri-infarct)<0.0001
S1 E blood	CD8*TRLs: d0=3, d1=8, d3=7, d5=5, d7=6, d14=4; CD4* Treg: d0=3, d1=5, d3=5, d5=4, d7=4, d14=5	Normal distribution	1-way ANOVA, Dunnett post hoc	F(CD8* TRLs)=15.20; F(CD4* Treg)=73.65	p(CD8* TRLs)<0.0001; p(d0 vs d1)<0.0001; p(d0 vs d3)=0.0019; p(d0 vs d5)=0.9926; p(d0 vs d7)=0.9910; p(d0 vs d14)=0.3897. p(CD4* Treg)<0.0001; p(d0 vs d1)<0.0001; p(d0 vs d3)<0.0001; p(d0 vs d5)<0.0001; p(d0 vs d7)=0.0013; p(d0 vs d14)=0.4580.
S1 E spleen	CD8* TRLs: d0=5, d1=6, d3=6, d5=5, d7=6, d14=7; CD4* Treg: d0=3, d1=3, d3=6, d5=4, d7=3, d14=5	Normal distribution	1-way ANOVA, Dunnett post hoc	F(CD8* TRLs)=13.05; F(CD4* Treg)=14.78	p(CD8* TRLs)<0.0001; p(d0 vs d1)<0.0001; p(d0 vs d3)=0.0346; p(d0 vs d5)=0.6306; p(d0 vs d7)=0.0265; p(d0 vs d14)=0.9684; p(CD4* Treg)<0.0001; p(d0 vs d1)=0.8671; p(d0 vs d3)=0.0002; p(d0 vs d5)<0.0001; p(d0 vs d7)=0.0001; p(d0 vs d14)=0.0002
<b>FIGURE S2</b>	<b>n</b>	<b>DATA STRUCTURE</b>	<b>TEST USED</b>	<b>STATISTIC</b>	<b>P VALUE</b>
S2 B	IgG=8; Anti-CD122=12	Normal distribution	2-way ANOVA, Bonferroni post hoc	F(1,180)=41.87	p<0.0001; p(1)>0.9999; p(2)>0.9999; p(3)>0.9999; p(4)>0.9999; p(5)=0.2560; p(6)=0.0518; p(7)0.0308; p(8)=0.0238; p(9)=0.0499; p(10)=0.1712
S2 C	IgG=8; Anti-CD122=12	Normal distribution	2-way ANOVA, Bonferroni post hoc	F(1,180)=33.71	p<0.0001; p(1)>0.9999; p(2)>0.9999; p(3)>0.9999; p(4)>0.9999; p(5)=0.6246; p(6)=0.1683;

					$p(7)=0.1035$ ; $p(8)=0.1371$ ; $p(9)=0.0887$ ; $p(10)=0.3808$
S2 D	IgG=8; Anti-CD122=9	Normal distribution	2-way ANOVA, Bonferroni post hoc	$F(1,120)=28.48$	$p<0.0001$ ; $p(1)>0.9999$ ; $p(2)>0.9999$ ; $p(3)=0.6710$ ; $p(4)=0.1894$ ; $p(5)=0.1588$ ; $p(6)=0.4506$ ; $p(7)=0.3311$ ; $p(8)=0.2091$
S2 E	IgG: d1=8, d3=8, d14=8; Anti-CD122: d1=12, d3=12, d14=9	Normal distribution	Mixed-effects analysis, Bonferroni post hoc	$F(1,18)=5.734$	$p=0.0277$ ; $p(d1)=0.1611$ , $p(d3)=0.1244$ , $p(d14)=0.1132$
<b>FIGURE S3</b>	<b>n</b>	<b>DATA STRUCTURE</b>	<b>TEST USED</b>	<b>STATISTIC</b>	<b>P VALUE</b>
S3A	IgG=6, anti-CD122=5 for each panel	Normal distribution	Unpaired t test	$t(TRL)=2.568$ ; $t(CD3^+ T)=1.637$ ; $t(CD19^+ B)=0.9097$ ; $t(NK)=2.406$ ; $t(neutrophil)=2.566$ ; $t(DC)=2.203$ ; $t(monocyte)=1.124$	$p(TRL)=0.0303$ ; $p(CD3^+ T)=0.1360$ ; $p(CD19^+ B)=0.3867$ ; $p(NK)=0.0395$ ; $p(neutrophil)=0.0304$ ; $p(DC)=0.0550$ ; $p(monocyte)=0.2900$
S3B	3d: IgG=7, anti- CD25=6, anti- CD122=7; 7d: IgG=5, anti- CD25=8, anti- CD122=5	Normal distribution	1-way ANOVA, Dunnnett post hoc	$F(3d)=40.09$ ; $F(7d)=61.23$	$p(3d)<0.0001$ , $p(IgG$ vs anti- CD25) $>0.9999$ , $p(IgG$ vs anti- CD122) $<0.0001$ , $p(anti-CD25$ vs anti- CD122) $<0.0001$ ; $p(7d)<0.0001$ , $p(IgG$ vs anti- CD25) $<0.0001$ , $p(IgG$ vs anti- CD122) $<0.0001$ , $p(anti-CD25$ vs anti- CD122) $>0.9999$
S3C Adhesive	IgG sham=6; Anti-CD122 sham=6	Normal distribution	2-way ANOVA, Bonferroni post hoc	$F(1,10)=0.7864$	$p=0.3960$ ; $p>0.9999$ for all time points between two groups.
S3C Rotarod	IgG sham=6; Anti-CD122 sham=6	Normal distribution	2-way ANOVA, Bonferroni post hoc	$F(1,10)=0.0182$	$p=0.8953$ ; $p>0.9999$ for all time points between two groups.
S3D	IgG=7, anti- CD122+anti-CD25=6	Normal distribution	Unpaired t test	$t=2.779$	$p=0.0179$
S3F	3d: IgG=8, anti-NK 1.1=6; 7d: IgG=8, anti- NK1.1=8	Normal distribution	Student's t test	3d $t=1.882$ 7d $t=1.094$	$p(3d)=0.082$ ; $p(7d)=0.2924$
<b>FIGURE S5</b>	<b>n</b>	<b>DATA STRUCTURE</b>	<b>TEST USED</b>	<b>STATISTIC</b>	<b>P VALUE</b>
S5 C	d0: blood=3, spleen=3, ipsilateral brain=4; d1: blood=7, spleen=6, ipsilateral brain=8; d3: blood=7, spleen=6, ipsilateral brain=5; d5: blood=5, spleen=5, ipsilateral brain=5;	Non normal distribution	1-way ANOVA, Bonferroni post hoc for each timepoint	1d: $F(2,18)=6.223$ ; 3d: $F(2,15)=6.254$ ; 5d: $F(2,12)=18.59$	1d: $p=0.0088$ , $p(blood$ vs spleen) $=0.4105$ , $p(blood$ vs ipsilateral brain) $=0.1911$ , $p(spleen$ vs ipsilateral brain) $=0.0077$ ; 3d: $p=0.0106$ , $p(blood$ vs spleen) $>0.9999$ $p(blood$ vs ipsilateral brain) $=0.0555$ $p(spleen$ vs ipsilateral brain) $=0.0111$ ; 5d: $p=0.0002$ , $p(blood$ vs spleen) $>0.9999$ , $p(blood$ vs ipsilateral brain) $=0.0007$ $p(spleen$ vs ipsilateral brain) $=0.0005$
S5D	Blood=3, spleen=3, ipsilateral brain=5	Normal distribution	1-way ANOVA, Bonferroni post hoc	$F(2,8)=82.96$	$p<0.0001$ ; $p(blood$ vs spleen) $>0.9999$ ; $p(blood$ vs ipsilateral brain) $<0.0001$ ; $p(spleen$ vs ipsilateral brain) $<0.0001$
S5 F	Anti-CD122=6, WT TRL=8, KO TRL=8	Normal distribution	1-way ANOVA, Bonferroni post hoc	$F(2,19)=15.89$	$p<0.0001$ ; $p(anti-CD122$ vs WT TRL) $<0.0001$ , $p(anti-CD122$ vs KO TRL) $=0.0948$ , $p(WT TRL$ vs KO TRL) $=0.0075$
S5 G	$n=4$ for both WT TRL and KO TRL	Normal distribution	Unpaired t test	$t(CCL1)=6.453$ ; $t(CCL2)=4.549$ ;	$p(CCL1)=0.0007$ ; $p(CCL2)=0.0039$ ;



					t(CCL4)=2.380; t(CCL5)=2.286; t(IL6)=3.195; t(IL17A)=2.170; t(IFNg)=0.4778; t(MMP9)=0.0276.	p(CCL4)=0.0548; p(CCL5)=0.0623; p(il6)=0.0187; p(IL17A)=0.0730; p(IFNg)=0.6497 p(MMP9)=0.9788
FIGURE S6	n	DATA STRUCTURE	TEST USED	STATISTIC	P VALUE	
S6A wb erk	N=5	Normal distribution	1-way ANOVA, Bonferroni post hoc	F (9,40)=53.53	p<0.0001; p(ctrl vs ETGF)=0.0012, p(ctrl vs OGD 0.5h)<0.0001, p(ctrl vs OGD 0.5h ETGF)<0.0001, p(ctrl vs OGD 2h)<0.0001, p(ctrl vs OGD 2h ETGF)<0.0001; p(OGD 6h vs OGD 6h ETGF)=0.0295, p(OGD 24h vs OGD 24h ETGF)<0.0001	
S6A wb Akt	N=5	Normal distribution	1-way ANOVA, Bonferroni post hoc	F (9,40)=13.39	p<0.0001; p(ctrl vs OGD 0.5h ETGF)<0.0001, p(OGD 24h vs OGD 24h ETGF)=0.024	
S6B	N=4	Normal distribution	1-way ANOVA, Bonferroni post hoc	F (4,15)=15.67	p<0.0001; p(ctrl vs OGD)<0.0001, p(OGD vs OGD ETGF)=0.0068, p(OGD ETGF vs OGD ETGF U)=0.0183, p(OGD ETGF vs OGD ETGF A)=0.0163	
S6C	N=4	Normal distribution	1-way ANOVA, Bonferroni post hoc	F (4,15)=69.16	p<0.0001; p(ctrl vs OGD)<0.0001, p(OGD vs OGD ETGF)<0.0001, p(OGD ETGF vs OGD ETGF U)<0.0001, p(OGD ETGF vs OGD ETGF A)<0.0001	
FIGURE S7	n	DATA STRUCTURE	TEST USED	STATISTIC	P VALUE	
S7C	PBS=6, Treg=8	Normal distribution	Unpaired t test	t=2.273	P=0.0422	
S7D	Sham young=6, sham aged=5, young 3=6, aged 3=7, young 5=6, aged=6	Normal distribution	Sham: Welch's t test; 3d: unpaired t test; 5d: Welch's t test	Sham: Welch-corrected t=2.504; 3d: t=2.701; 5d: Welch-corrected t=1.159	p(sham)=0.0597, p(3d)=0.0206, p(5d)=0.2771	
S7F	PBS=10; TRLs=7	Normal distribution	2-way ANOVA, Bonferroni post hoc	F(1,15)=57.71	p<0.0001; p(Pre)>0.9999; p(3d)<0.0001; p(7d)<0.0001; p(14d)=0.0009; p(21d)=0.0004; p(28d)=0.0074; p(35d)>0.9999	
S7G	PBS=10; TRLs=7	Normal distribution	2-way ANOVA, Bonferroni post hoc	F(1,15)=71.72	p<0.0001; p(Pre)>0.9999; p(3d)<0.0001; p(7d)<0.0001; p(14d)<0.0001; p(21d)=0.0105; p(28d)=0.0075; p(35d)=0.0321	
S7H	PBS=11; TRLs=8	Normal distribution	2-way ANOVA, Bonferroni post hoc	F(1,17)=19.08	P=0.0004; p(21d)=0.5889; p(22d)<0.0001; p(23d)=0.0052; p(24d)=0.1114	
S7I	PBS=11; TRLs=8	Normal distribution	Unpaired t test	t=3.981	p=0.0010	
S7J	PBS=10; TRLs=7	Normal distribution	Unpaired t test	t=6.882	p<0.0001	
S7K	PBS=7; TRLs=7	Normal distribution	Unpaired t test	t=3.355	p=0.0057	