

Supplementary Data

TITLE

Sensitization of glioblastoma tumor micro-environment to chemo- and immunotherapy by Galectin-1 intranasal knock-down strategy

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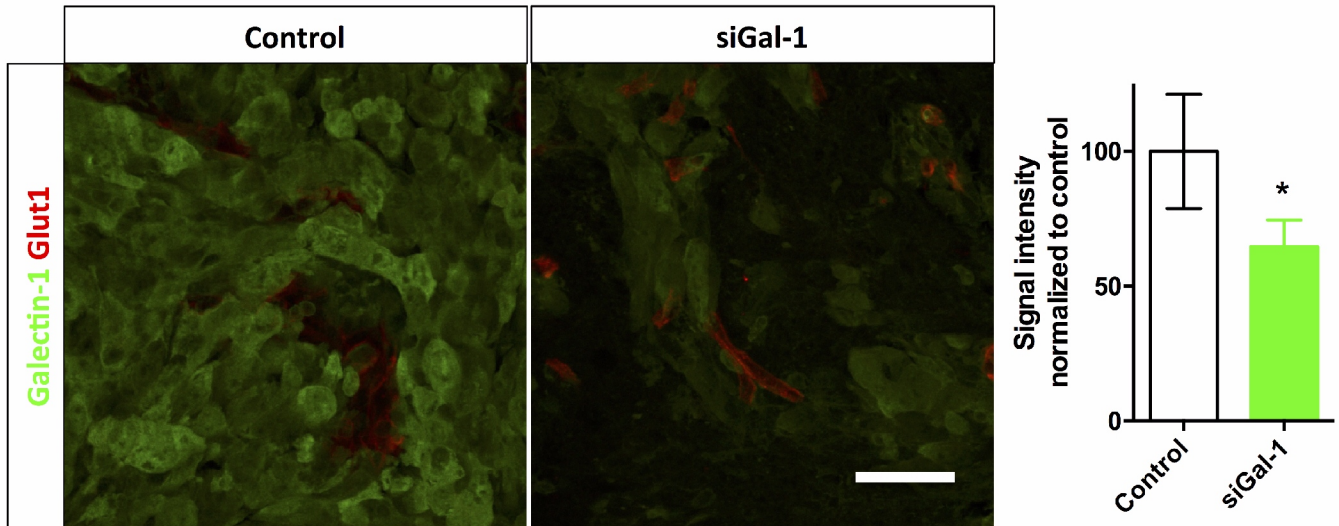
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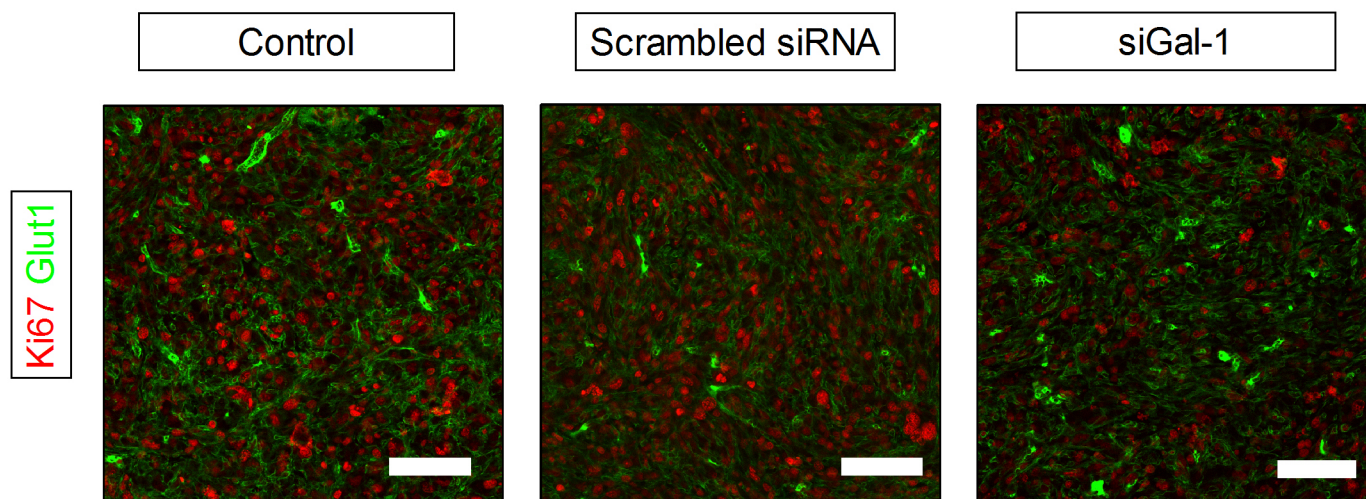
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Supplementary Figure 1



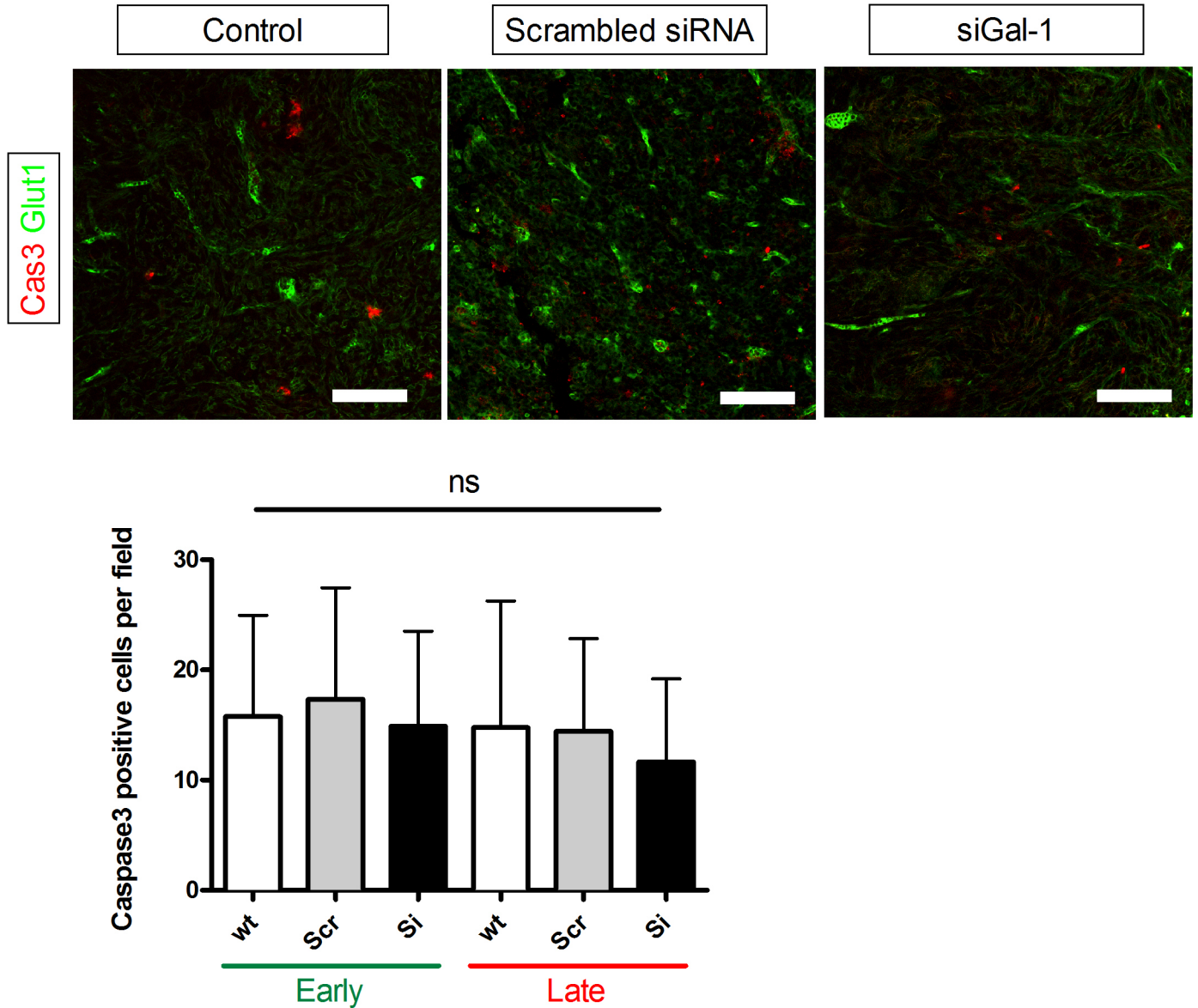
Supplementary Figure 1: Intratumoral Gal-1 reduction. Mice were inoculated with 0.5×10^6 GL261 cells which induces a lethal GBM after 15-20 days. Representative pictures of late stage tumors are shown for control mice or mice treated with siGal-1 as described in Figure 1A. Green: Gal-1 and Red: Glut-1. On the right side, the quantification of Gal-1 intensity in at least 3 slices/mouse, demonstrates a significant reduction of Gal-1 in the tumor. ($n=5/\text{group}$, $* = p<0.05$, unpaired t – test). Scale bar: $100\mu\text{m}$.

Supplementary Figure 2



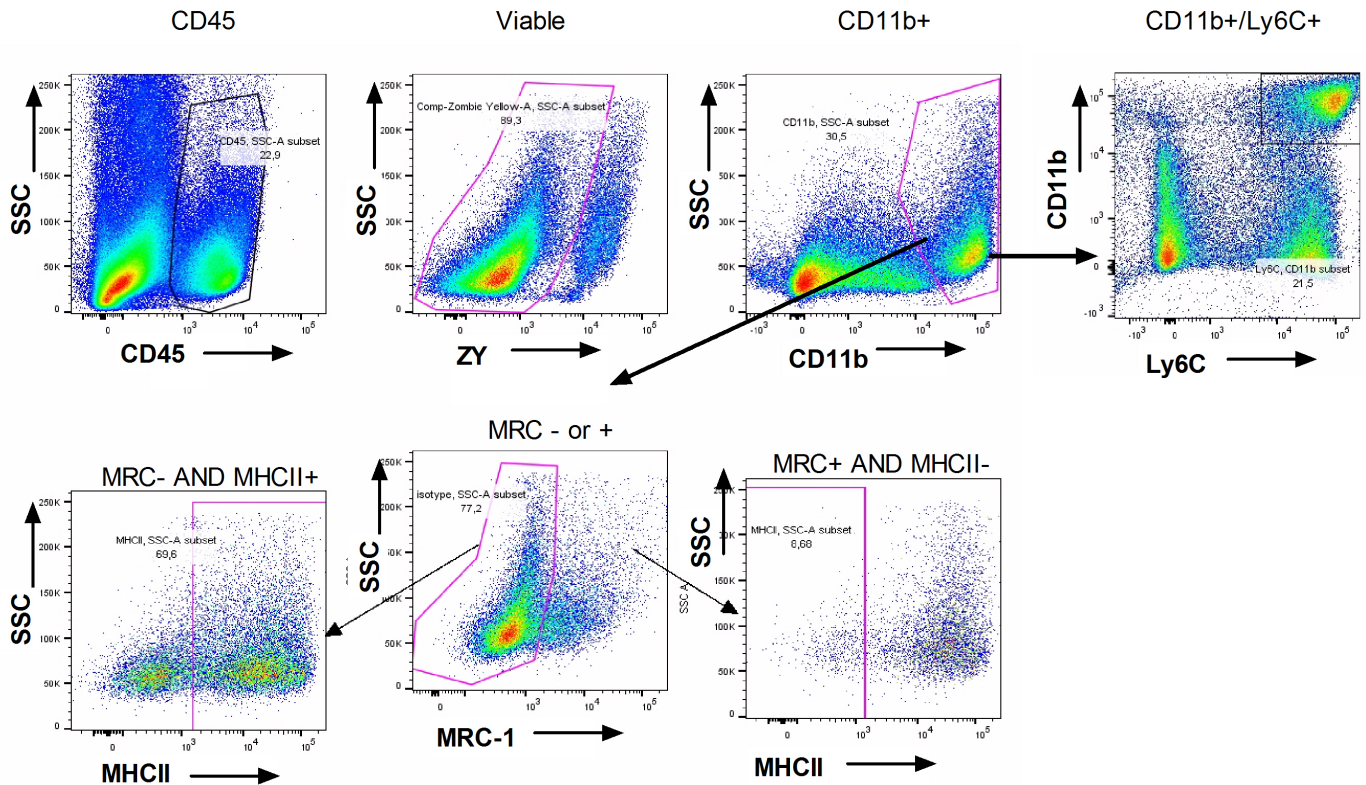
Supplementary Figure 2: Ki67 staining pictures. Mice were inoculated with 0.5×10^6 GL261 cells which induces a lethal GBM after 15-20 days. Representative pictures of late stage tumors are shown for control, untreated mice; mice treated with scrambled siRNA loaded chitosan nanoparticles, or mice treated with siGal-1. Scale bar: 250 μ m.

Supplementary Figure 3



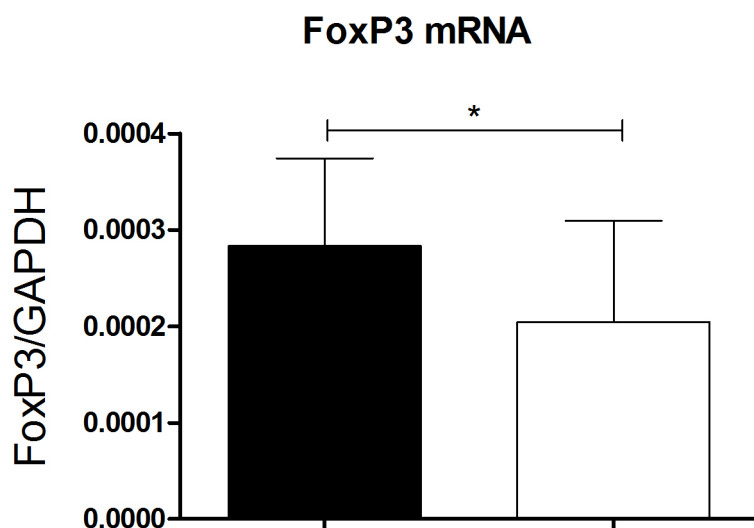
Supplementary Figure 3: Cas3 staining pictures. Mice were inoculated with 0.5×10^6 GL261 cells which induces a lethal GBM after 15-20 days. Representative pictures of late stage tumors are shown for control, untreated mice; mice treated with scrambled siRNA loaded chitosan nanoparticles, or mice treated with siGal-1. Moreover, quantification is provided of multiple sections/mice ($n=4$ /group, at least 3 independent fields/mouse). Scale bar 200 μ m.

Supplementary Figure 4



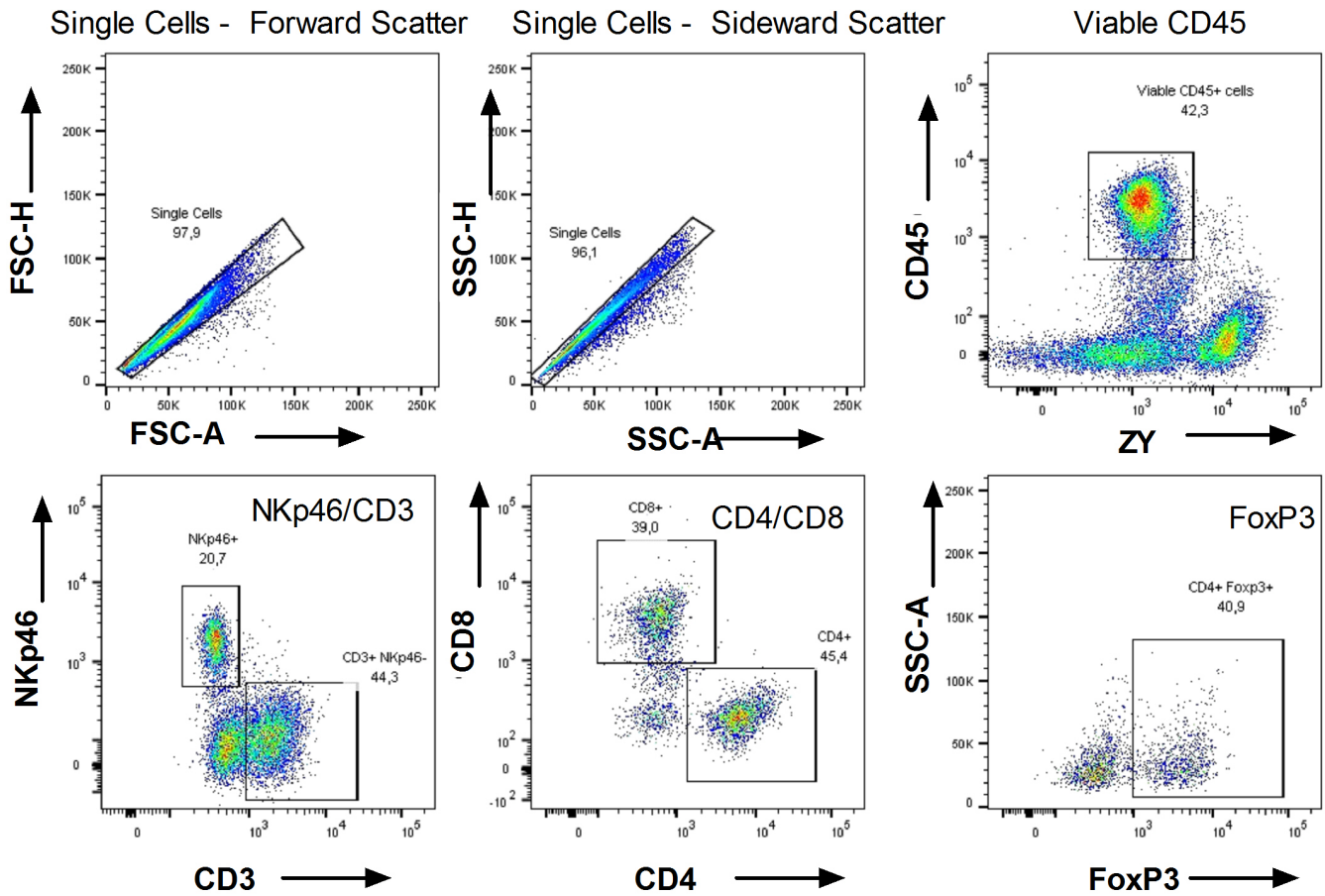
Supplementary Figure 4: Gating strategy for myeloid cell population. Flow cytometry was performed on isolated mononuclear brain infiltrating cells of mice and gated for CD45 positive (leukocytes), viable cells (ZY negative), and CD11b positive (myeloid). Subsequently, we looked either into Ly6C for monocytic MDSCs and Ly6G for granulocytic MDSCs. Furthermore, we also analysed the CD11b positive cells for MRC1 expression (middle, down row), and we monitored for MRC1 negative cells for MHCII positive (M1 macrophage phenotype), on the other side, we monitored the MRC1 positive cells for MHCII negative (M2 macrophage phenotype).

Supplementary Figure 5



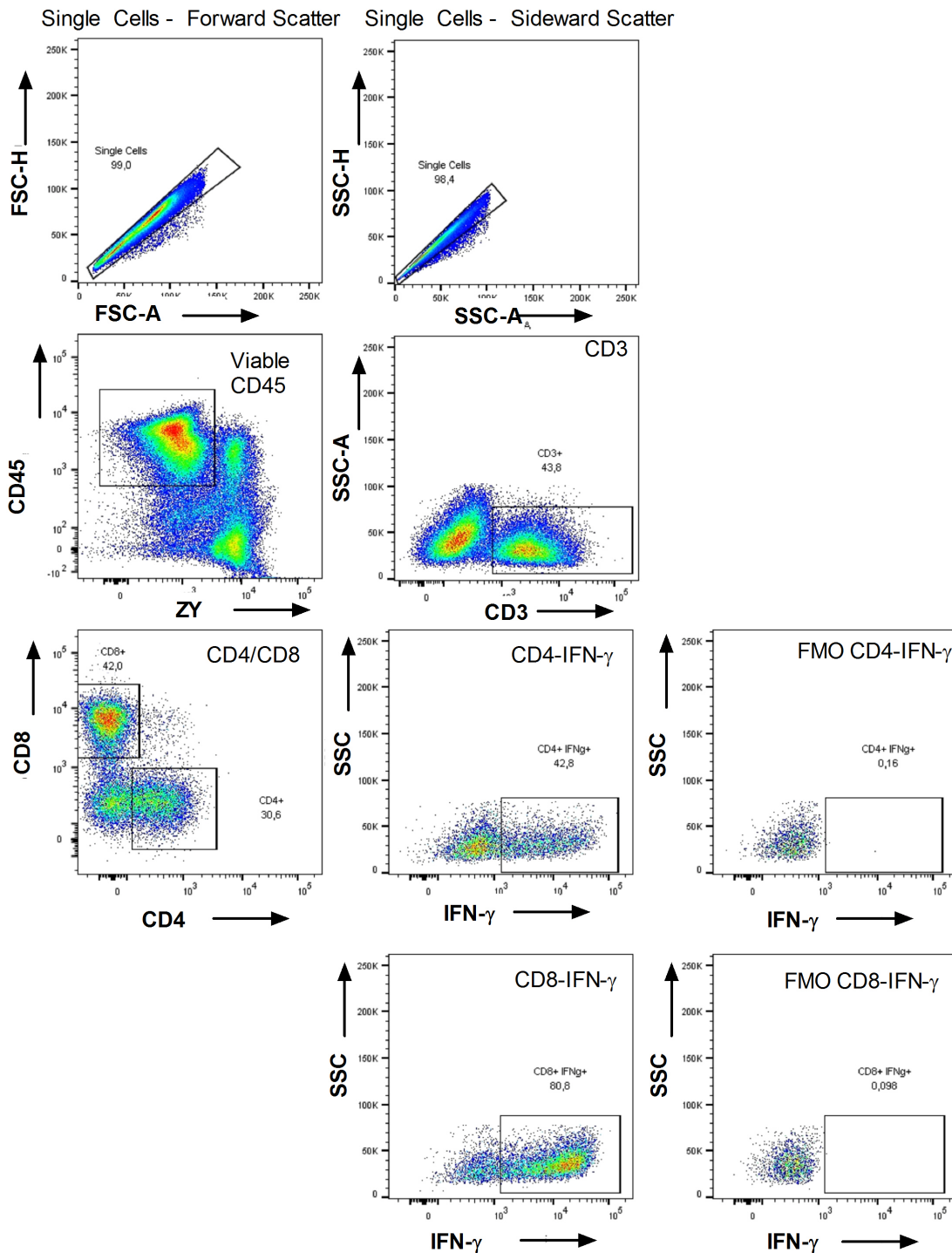
Supplementary Figure 5: mRNA analysis for FoxP3. Mice were inoculated with 0.5×10^6 GL261 cells which induces a lethal GBM after 15-20 days. RT-qPCR was performed on isolated GBM specimen of mice that were left untreated, or treated with siGal-1 on day 4, 8, 12 and 15 after tumor inoculation, and brains were isolated at day 20, which revealed a significant reduction for *foxP3* transcription factor expression (n = 10/group, * = p < 0.05).

Supplementary Figure 6



Supplementary Figure 6: Gating strategy for Treg. Flow cytometry was performed on isolated mononuclear brain infiltrating cells of mice and gated for single cells (via SSC and FSC), CD45 positive (leukocytes), viable cells (ZY negative), and CD3 positive (lymphoid). Subsequently, we looked into CD4 and CD8 for T cells, and within the CD4 gate, we monitored the expression of FoxP3 expression.

Supplementary Figure 7



Supplementary Figure 7: Gating strategy for Lymphoid cell populations. Flow cytometry was performed on isolated mononuclear brain infiltrating cells of mice and gated for single cells (via SSC and FSC), CD45 positive (leukocytes), viable cells (ZY negative), and CD3 positive (lymphoid). Subsequently, we looked into CD4 and CD8 for T cells, and for both cell populations, we monitored the expression of IFN- γ expression, as guided by Fluorescence Minus One, to determine the proper gating strategy.