(A) sLeA expression on murine and human tumor cells



(B) Binding of different sLeA-targeting Ab clones to sLeA+ tumor cells





(C) Binding of different sLeA-targeting Ab clones to synthetic sLeA - indirect ELISA



Figure S1

(A) Murine tumor cells were transduced to stably express the human enzyme Fucosyltransferase III (FUT3), which synthesizes sLeA. *FUT3*-transduced murine tumor cells and human tumor cells (HT-29 and OVCAR3) were labeled with an Anti-sLeA primary Ab (clone 5B1-hlgG1) followed by Alexa-488-conjugated-goat anti-human IgG. The panel shows a representative experiment (n>3), showing similar results. (B) To validate binding of 5B1 and 7E3 Ab clones to surface sLeA - cells were labeled with an Anti-sLeA primary Abs followed by Alexa-488-conjugated-goat anti-human IgG. The panel shows a representative experiment (n>3), showing similar results. (B) To validate binding of 5B1 and 7E3 Ab clones to surface sLeA - cells were labeled with an Anti-sLeA primary Abs followed by Alexa-488-conjugated-goat anti-human IgG. The panel shows a representative experiment (n>3), showing similar results. (C) Assay plates were coated and incubated with synthetic sLeA, blocked and incubated with serial dilutions of clones 5B1-hlgG1, 7E3-hlgG1 or non-relevant hlgG1 isotype control Abs. HRP-conjugated goat anti-human IgG Abs, followed by addition of TMB substrate were used for detection.

(A) Binding of Anti-sLeA Ab variants (murine Fc)

Clone 5B1 100 10¹ 10² 10^{-1} 103 sLeA





(B) Binding of Anti-sLeA Ab variants to sLA+ cells (human Fc)



FUT3+ cells - hlgG1 (WT) FUT3+ cells - hlgG1-N297A FUT3+ cells - hlgG1-GA FUT3+ cells - hlgG1-ALIE FUT3+ cells - hlgG1-GAALIE

Figure S2

Murine tumor cells were transduced to stably express the human enzyme Fucosyltransferase III (FUT3), which synthesizes sLeA. To validate binding of various sLeA-targeting Abs with a murine Fc (A) or human Fc (B) - cells were labeled with an Anti-sLeA primary Abs followed by Alexa-488-conjugated-goat anti-human IgG or anti-mouse IgG and analyzed by flow cytometry. Both panels show a representative experiment (n>3), showing similar results.

(A) sLeA-targeting Abs protect mice from sLeA+ tumor challenge comparably to gp75-targeting Abs



(B) 5B1 Ab-mediated anti-tumor effect relies on activating FcyRs





Figure S3

WT C57BL/6 mice (**A**, **B**) or Fc γ R-humanized mice (**C**) were inoculated IV with 5*10⁵ B16-FUT3 tumor cells. 100 µg of anti-sLeA Abs (5B1-mlgG2a or 5B1-mlgG1 or 5B1-hlgG1 or 5B1-hlgG1-N297A), anti-gp75 (TA99-mlgG2a) or isotype-matched control Abs were administered IP on days 1,4,7 and 11. 14 days post-inoculation, mice were euthanized, lungs were excised and fixed, and metastatic foci were counted. For all panels, data pooled from n=2-3 experiments, n≥11/group. Panels A, B show representative images of three excised lungs from each group. * p<0.05, ** p<0.01, **** p<0.0001. For all panels, the box extends from 25th to 75th percentile, the line within the box represents the median value and the whiskers correlate to 5-95 percentiles.



Figure S4

Binding affinity of hIgG1 Fc variants to human FcqRs was determined by SPR analysis. Representative SPR sensorgrams of hIgG1-WT and hIgG1-GAALIE variants binding to the various classes of hFcqRs (see table 1 for the affinity measurements and fold change in affinity over hIgG1-WT).

(A) Antibody half-life in vivo



(B) ADCC in vivo - platelet depletion



Figure S5

(A) Half-life of hlgG1 variants in FcyR-humanized mice – FcyR-humanized mice were injected IV with 100 μ g of hlgG1 Fc variants, bled at indicated time points, and serum IgG levels were determined by ELISA. n=4/group. (B) ADCC in vivo – FcyR-humanized mice were injected intravenously with 10 μ g of hlgG1 Fc variants (clone 6A6, targeting a platelet-associated antigen). hlgG1-N297A (was included as control. Mice were bled at the indicated time points and platelet numbers were analyzed using an Advia 120 hematology system. Values represent the mean (± SEM) percentage change in platelet number relative to 0h. n=3/group. For hlgG1-WT vs. hlgG1-GAALIE at 4h and 24h - * p<0.05.

(A) Validation of NK cell depletion in murine lungs



(B) NK cell depletion does not hamper anti-tumor effect of sLeA-targeting Abs in FcyR-humanized mice



Figure S6

FcyR-humanized mice were inoculated IV with $5*10^5$ B16-FUT3 tumor cells. 100 µg of anti-sLeA Abs (5B1-hlgG1-GAALIE with G236A/A330L/I332E mutations) or isotype-matched control Abs were administered IP on days 1,4,7 and 11. NK cells were depleted using Intra-nasal administrations of 300 µg of anti-NK1.1 Abs on days -3, -1, 1, 4, 7, and 11. For all panels, 14 days post-inoculation, mice were euthanized, and lungs were excised. **(A) Validation of NK cell depletion** – Lungs were perfused, dissociated and stained for flow cytometry. NK cells were gated as follows: Live/CD45+/CD3-/CD19-/B220-/NKp46+/CD49b+ cells, and are presented as Mean ± SEM. n≥4/group. **** p<0.0001 (one-way ANOVA with Bonferroni's post-test). The panel summarizes the data obtained for all mice (left), and shows representative dot plots for a single mouse from each group (right). **(B) NK cell depletion does not hamper anti-tumor effect of sLeA-targeting Abs** – Lungs were excised and fixed, and metastatic foci were counted. Data pooled from n=2 experiments, n≥11/group, and shows representative images of three excised lungs from each group. ** p<0.01, *** p<0.001 (one-way ANOVA with Bonferroni's post-test). The box extends from 25th to 75th percentile, the line within the box represents the median value and the whiskers correlate to 5-95 percentiles.

anti-sLeA and anti-gp75 Abs demonstrate differential hFcyR-engagement requirements to confer anti-tumor activity



Figure S7

FcγR-humanized mice were inoculated IV with 5*10⁵ B16-FUT3 tumor cells. 100 μg of anti-sLeA Abs (5B1-hlgG1 or 5B1-hlgG1-GA or 5B1-hlgG1-ALIE) or antigp75 (TA99-hlgG1 or TA99-hlgG1-GA or TA99-hlgG1-ALIE) or isotype-matched control Abs were administered IP on days 1,4,7 and 11. 14 days postinoculation, mice were euthanized, lungs were excised and fixed, and metastatic foci were counted. For 5B1 - data pooled from n=2 experiments, n=10-12/group. For TA99 - n=5/group. * p<0.05, *** p<0.001, **** p<0.0001 (one-way ANOVA with Bonferroni's post-test). The box extends from 25th to 75th percentile, the line within the box represents the median value and the whiskers correlate to 5-95 percentiles.