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Supplemental Information

Interleukin-34 Limits the Therapeutic Effects of Immune Checkpoint Blockade

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1 **Transparent Methods**

2 **Cell lines**

3 The ovarian cancer cell line OV2944-HM-1 (HM-1) was purchased from the Japanese
4 Collection of Research Bioresources. The colon cancer cell line CT26 used in this study
5 was kindly provided by Dr. Hidemitsu Kitamura, Hokkaido University. The breast cancer
6 cell line 4T1 was purchased from American Type Culture Collection. HM-1 cell line was
7 maintained in α MEM (Fujifilm Wako Pure Chemical Industries). CT26 and 4T1 cell lines
8 were maintained in RPMI-1640 (Fujifilm Wako Pure Chemical Industries). All culture
9 media were supplemented with 10% fetal bovine serum (Sigma Aldrich), 1%
10 Penicillin/Streptomycin (Nacalai Tesque), and 1% Non-Essential Amino Acid (Nacalai
11 Tesque). Cells were maintained in a 5% CO₂/air environment at 37°C.

12

13 **Mice and *in vivo* assay**

14 Six to eight-week-old female B6C3F1 and BALB/c mice were purchased from Japan SLC,
15 Inc. The mice were maintained under specific pathogen-free conditions in the animal
16 facility at Hokkaido University. For *in vivo* assay, 2×10^5 tumor cells were inoculated s.c.
17 into the right flank of syngeneic female mice. Antibody treatment (anti-PD-1 (RMP1-14),
18 250 μ g/mouse; CTLA-4 (UC10-4F10), 250 μ g/mouse; or IL-34 (C054-35), 200 μ g/mouse

19 was started when tumor size reached 5 mm in diameter. Anti-PD-1 and anti-CTLA-4
20 antibodies were kindly provided by Dr. Hideo Yagita (Juntendo University). Anti-IL-34
21 antibody was purchased from BioLegend. Detailed information about antibodies is
22 described in Supplementary Table 1. All animal procedures were approved by the
23 Hokkaido University Animal Care Committee (Approval number: 14-0171).

24

25 **Generation of *Il34* knockout and *Il34* overexpression cell lines**

26 *Il34*^{KO} cell line was generated by using IL-34 CRISPR/Cas9 KO Plasmid (m) (Santa Cruz
27 Biotechnology, Inc.). The plasmids were transfected by using *TransIT-X2* (Mirus) or Neon[®]
28 Transfection system (Thermo Fisher Scientific). Cells were selected by GFP expression
29 48 hours after transfection. For the generation of *Il34* overexpression CT26 cell line,
30 mouse *Il34* coding sequence was cloned into pLenti-EF1a-C-Myc-DDK-IRES-Puro vector
31 (Origene). Lenti-X293T cells were transfected with lentiviral vector and two packaging
32 plasmids pCMV-VSV-G-RSV-Rev and pCAG-HIVgp using *TransIT-X2*. The complex was
33 added in HEK293T cells and incubated 3 days. After collection of HEK293T medium,
34 CT26 was cultured with 1:1 mixture of HEK293T medium and fresh medium, following
35 selection by puromycin.

36

37 **Quantitative PCR analysis**

38 Total RNA was extracted using TRIsure reagent (Bioline). cDNA was synthesized using
39 ReverTra Ace[®] qPCR RT Master Mix (Toyobo). Quantitative PCR was performed on cDNA
40 using KAPA SYBR[®] FAST qPCR Master Mix (2X) ABI Prism[®] (Kapa Biosystems) on a
41 StepOnePlus Real-Time PCR System (Thermo Fisher Scientific). The primers are listed
42 in Supplementary Table 2.

43

44 **Cell viability assay**

45 To assess cell viability, MTT assay was performed using MTT Cell count kit (Nacalai
46 Tesque). Absorbance at a test wavelength of 570 nm and a reference wavelength of 650
47 nm was measured by using a Multiskan FC (Thermo Fisher Scientific). Cell proliferation
48 was observed up to 4 days.

49

50 **Enzyme-linked immunosorbent assay (ELISA)**

51 The production of IL-34 in cell lines was measured with ELISA. Culture supernatants were
52 collected at 48 h after seeding the cells at a density of 1×10^6 in 6-well plate. The IL-34
53 contents was measured with LEGEND MAX Mouse IL-34 ELISA kit with Pre-Coated
54 Plates (Biolegend).

55

56 **Isolation of tumor-infiltrating immune cells from solid tumor**

57 Isolation of tumor-infiltrating immune cells from solid tumors was performed by using BD
58 Horizon™ Dri Tumor & Tissue (Becton, Dickinson, and Company). The recovered tumor-
59 infiltrating cells were used as samples for flow cytometry or RNA extraction.

60

61 **Flow cytometry**

62 Cells were washed and blocked with FcR Blocking Reagent (TONBO biosciences) and
63 stained with 4',6-diamidino-2-phenylindole (DAPI, Cayman Chemical Company) and the
64 antibodies against following molecules; CD3ε, CD4, CD8α, F4/80, CD11b, CD11c, CD45,
65 CD115, IA-IE (MHC classII), iNOS, Arginase 1 and PD-L1 (BioLegend). Data were
66 acquired using BD FACSCanto II, BD FACSAria, or BD FACSCelesta flow cytometer, and
67 analyzed using FlowJo software. Detailed information about antibodies is described in
68 Supplementary Table 1.

69

70 **Next-generation sequencing and data analysis**

71 Total RNA was extracted using PureLink™ RNA Mini Kit (Life Technologies). Next-
72 generation sequencing was performed at Kazusa DNA Research Institute (illumina HiSeq

73 2500). The quality and concentration of the RNA was verified with Agilent 2100
74 Bioanalyzer and Quantus Fluorometer (Promega), respectively. All the samples showed
75 RIN values > 8. Sequencing libraries were prepared using Agilent SureSelect Strand-
76 Specific RNA Library Prep for Illumina according to the manufacturer's instructions. Briefly,
77 poly-A RNA was purified from 300 ng total RNA per sample using oligo dT magnetic beads.
78 The libraries were PCR amplified for 13 cycles and purified with AMPure XP beads.
79 Sequencing of the libraries was conducted on the Illumina HiSeq2500 system performing
80 paired-end 100 bp reads. The reads were mapped to mouse reference genome mm10
81 with Tophat (v2.1.0), and calculated FPKM (fragments per kilobase of exon per million
82 reads mapped) value with cufflinks (v2.2.1). The FPKM values were normalized by CD45,
83 and shown as global z-score.

84

85 **Immunohistochemistry staining**

86 For DAB staining, immunohistochemistry staining was performed on paraffin-embedded
87 tumor tissue sections. PD-L1 was stained using DAB (Dojindo) followed by hematoxylin
88 counterstaining (Fujifilm Wako Pure Chemical Industries). PD-L1 staining was kindly
89 performed by Dr. Yutaka Hatanaka, Research Division of Genome Companion
90 Diagnostics, Hokkaido University Hospital. For multiple immunofluorescent staining, Opal

91 4-color fluorescent IHC kit (Perkin-Elmer) was used. Tumor sections were objectively
92 judged by two independent researchers at 600× magnification for each section. More than
93 6 tumor areas in each section were randomly selected for evaluation. FV1000 OLYMPUS
94 software was used for quantification of immunofluorescent staining. Detailed information
95 about antibodies is described in Supplementary Table 1.

96

97 **PDX model**

98 PDX model was performed at DNA Link, Inc. Firstly, HuNSG mice were generated as
99 previously reported by The Jackson Laboratory (Shultz et al., 2005). In brief, human fetal
100 liver CD34⁺-purified HSC were purchased from Stem Express and intravenously injected
101 into three-week-old female NSG mice (10⁵ cells/mouse), 4h post-140 cGy total body
102 irradiation using the RS-2000 irradiator (Rad Source). The engraftment levels of human
103 CD45⁺ cells were determined 12 weeks post-HSC transplantation by flow cytometric
104 quantification of peripheral blood. HuNSG mice that had over 25% of human CD45⁺ cells
105 in the peripheral blood were considered as engrafted and humanized. PDX models were
106 generated using tumor tissues from patients who underwent surgery as the primary
107 treatment strategy for lung cancer at Samsung Medical Center. Twelve weeks post-human
108 HSC transplantation, 30-40 μl finely minced tumors were injected s.c. into the left flank of

109 HuNSG mice. Treatment was started when the tumor volumes reached 70-120 mm³.
110 Treatment with anti-human IL-34 (BioLegend; 250 µg per injection, 3 times a week for 4
111 weeks), anti-human PD-1 (Selleckchem; 10 mg/kg for the first dose, followed by 5 mg/kg
112 dose every 5 days), antibodies combination, or saline was administered intraperitoneally.
113 Vehicle control saline (Sigma Aldrich) was administered 3 times per week until the
114 endpoint. Tumor size was measured by caliper twice a week, and volumes (mm³) were
115 calculated by $(\text{length} \times \text{width}^2)/2$.

116 For histological analysis, tumor tissues were fixed with 4% formaldehyde, embedded
117 with paraffin and sections were stained with hematoxylin and eosin.

118 All animal experiments were performed under the guidelines approved by the
119 Institutional Animal Care and Use Committee of Seoul National University Biomedical
120 Research Institute.

121

122 **Statistics**

123 Statistical analysis was performed with JMP[®] 14 (SAS Institute Inc.). Significance was
124 determined by Student's *t*-test, Tukey's multiple comparison test, or Steel-Dwass
125 nonparametric multiple comparison test. p-Value was considered statistically significant
126 when < 0.05 .

127 **Supplemental Reference**

128 Shultz, L.D., Lyons, B.L., Burzenski, L.M., Gott, B., Chen, X., Chaleff, S., Kotb, M.,

129 Gillies, S.D., King, M., Mangada, J., et al. (2005). Human Lymphoid and Myeloid Cell

130 Development in NOD/LtSz- *scid* *IL2R* γ ^{null} Mice Engrafted with Mobilized Human

131 Hemopoietic Stem Cells. *J. Immunol.* *174*, 6477–6489.

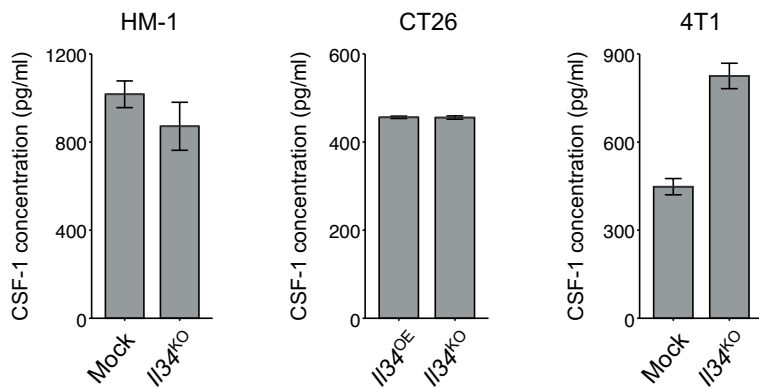


Figure S1 : The expression of CSF-1 from various murine cancer cell lines. Related to Figure 1, 2.

CSF-1 concentration in supernatants of HM-1, CT26 and 4T1 cell lines (n=3/cell line).

Data represent mean \pm SEM.

Figure S1

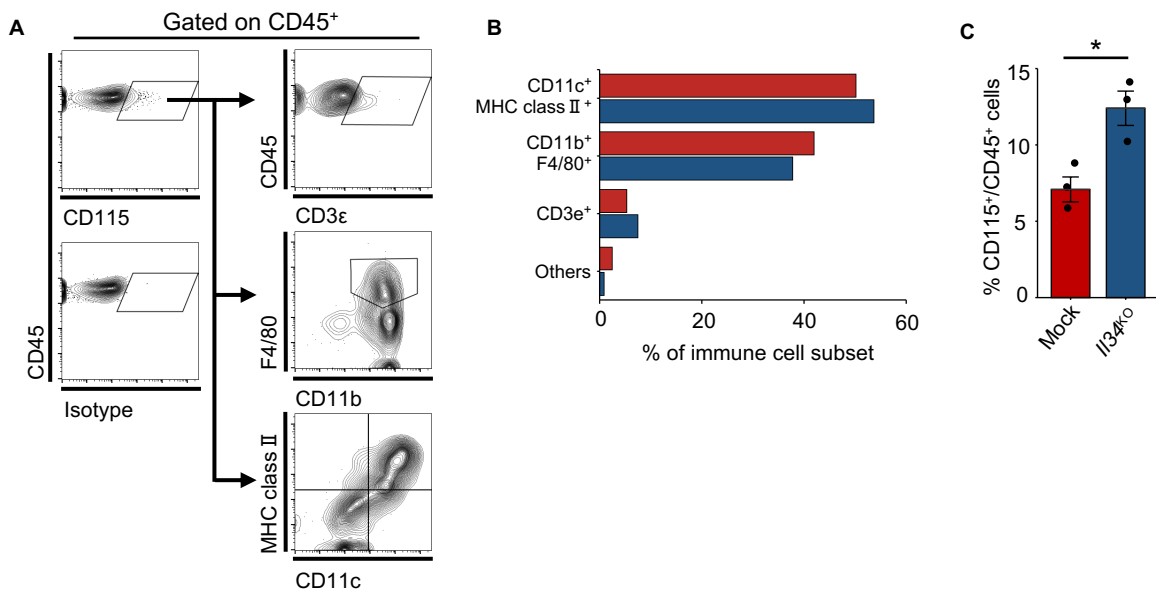


Figure S2 : Identification of the immune cell subset expressing CD115 in mock-HM-1 and I/34^{KO}-HM-1 tumors. Related to Figure 1.

(A) Representative flow cytometry profiles showing CD115⁺ cells within tumor-infiltrating CD45⁺ cells in HM-1 tumor.

(B) Bar graph shows the cell type expressing CD115 within tumor-infiltrating CD45⁺ cells (n=3/group).

(C) Bar graph represent the frequency of CD115⁺ cells within tumor-infiltrating CD45⁺ cells in mock- or I/34^{KO}-HM-1 tumors (n=3/group). Data represent mean ± SEM. *p<0.05; two-tailed Student's t-test.

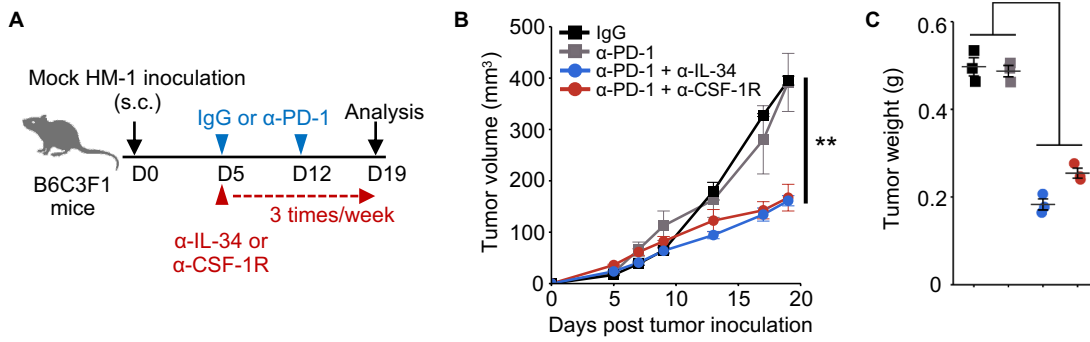


Figure S3 : IL-34 blockade equally enhanced anti-tumor efficacy of α-PD-1 treatment comparing with CSF-1R blockade in HM-1 model. Related to Figure 1.

(A) Schematic of the α-PD-1 mAb treatment in combination with α-IL-34 or α-CSF-1R mAb. The timeline shows the procedure of tumor inoculation and antibody treatment.

(B) Mock HM-1 tumor growth in B6C3F1 mice treated with the indicated antibodies (n=3-4/group).

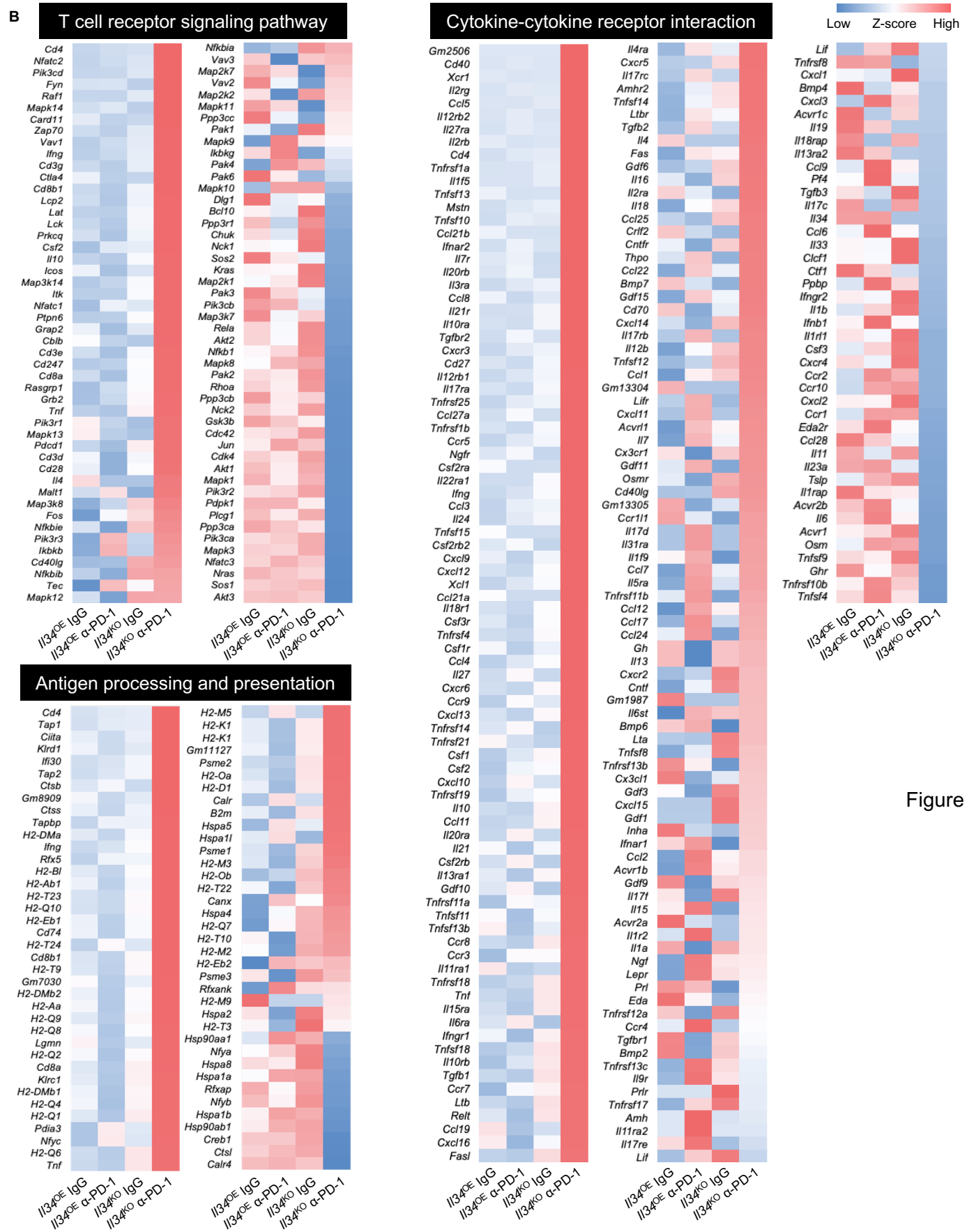
(C) Tumor weight on day 19 after tumor inoculation (n=3-4/group).

Data represent mean ± SEM. **p<0.01; Tukey's multiple comparison test.

Canonical Pathway	p-value
T cell receptor signaling pathway	3.68E-10
Graft-versus-host disease	4.47E-10
Antigen processing and presentation	9.52E-10
Systemic lupus erythematosus	3.84E-09
Allograft rejection	5.42E-09
Asthma	3.76E-08
Cell adhesion molecules (CAMs)	3.86E-08
Hematopoietic cell lineage	1.05E-07
Type I diabetes mellitus	2.25E-07
Cytokine-cytokine receptor interaction	2.45E-07

(A) The list of gene-set clusters enhanced in *Il34*^{KO} CT26 group compared to *Il34*^{OE} CT26 group.

(B) Heatmap shows the differentially of gene expression on selected several gene-set clusters.



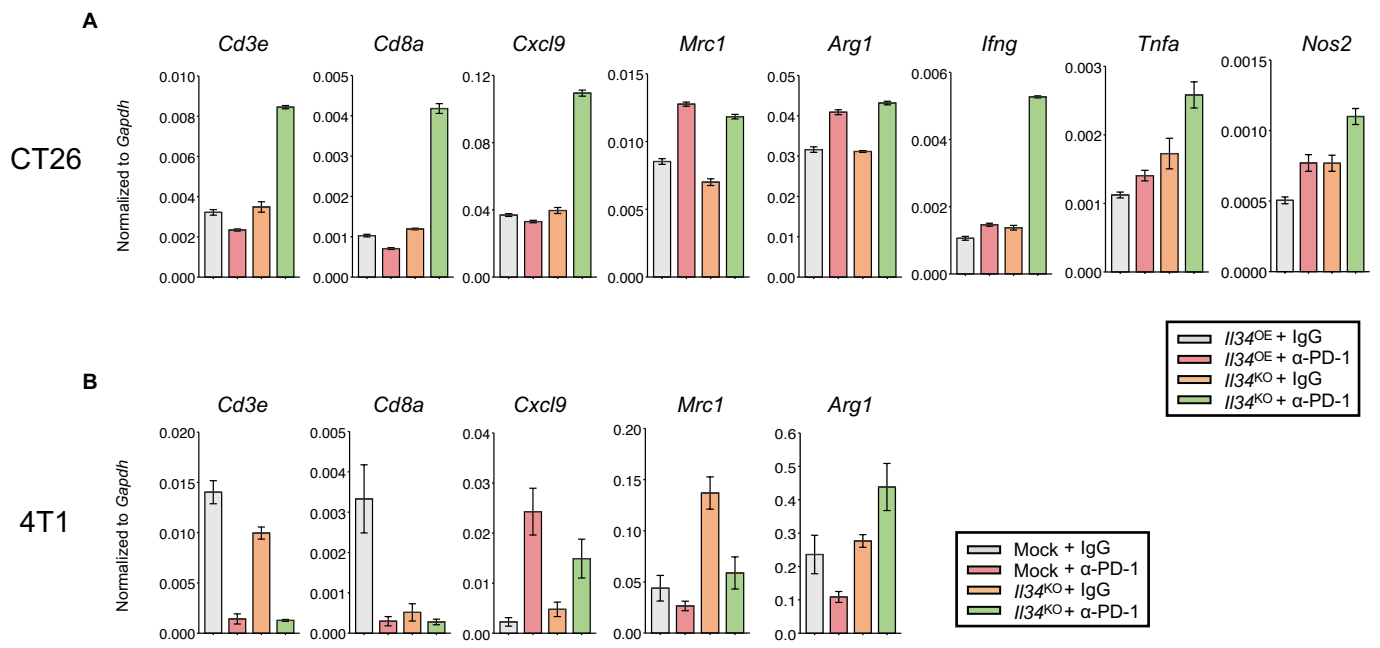


Figure S5 : qPCR analysis in CT26 and 4T1 tumors. Related to Figure 2.

(A) Expression of selected genes were evaluated by qPCR analysis in CT26 tumor samples used for NGS analysis (Fig. 2D).

(B) Gene expression displayed in (A) were analyzed by qPCR in 4T1 tumors (n=3/group).

Data represent mean \pm SEM of technical triplicate.

Key resources table

REAGENT & RESOURCE	SOURCE	IDENTIFIER
Antibodies		
Anti-mouse CD3 ϵ (145-2C11) APC	BioLegend	Cat#100236; RRID: AB_2561456
Anti-mouse CD4 (RM4-5) APC-Cy7	BioLegend	Cat#00525; RRID: AB_312726
Anti-mouse CD8 α (53-6.7) FITC	BioLegend	Cat#00706; RRID: AB_312745
Anti-mouse CD11b (M1/70) FITC	BioLegend	Cat#101206; RRID: AB_312789
Anti-mouse CD11c (N418) APC	BioLegend	Cat#117309; RRID: AB_313778
Anti-mouse CD45 (30-F11) Pacific blue	BioLegend	Cat#103126; RRID: AB_493535
Anti-mouse CD45 (30-F11) FITC	BioLegend	Cat#103108; RRID: AB_312973
Anti-mouse CD45 (30-F11) PE	BioLegend	Cat#103106; RRID: AB_312971
Anti-mouse CD45 (30-F11) APC	BioLegend	Cat#103112; RRID: AB_312977
Anti-mouse CD45 (30-F11) PE-Cy7	BioLegend	Cat#103114; RRID: AB_312979
Anti-mouse CD45 (30-F11) APC-Cy7	BioLegend	Cat# 103116; RRID: AB_312981
Anti-mouse F4/80 (BM8) APC	BioLegend	Cat#123116; RRID: AB_893481
Anti-mouse CD274 (MIH5) APC	BioLegend	Cat#124311; RRID: AB_10612935
Anti-mouse CD273 (TY25) PE	BioLegend	Cat#107205; RRID: AB_2299418
Anti-mouse CD80 (16-10A1) FITC	BioLegend	Cat#104705; RRID: AB_313126
Anti-mouse CD86 (GL-1) FITC	BioLegend	Cat#105005; RRID: AB_313148
Anti-mouse CD115 (AFS98) APC	BioLegend	Cat#125509; RRID: AB_2085222
Anti-mouse I-A/I-E (M5/114.15.2) FITC	BioLegend	Cat#107606; RRID: AB_313321
Anti-mouse iNOS (CXNFT) APC, eBioscience™	Invitrogen™	Cat#17-5920-80; RRID: AB_2573244
Anti-human/mouse Arginase 1 (A1exF5) PE, eBioscience™	Invitrogen™	Cat#12-3697-80; RRID: AB_2734839
Purified anti-mouse CD16/CD32 (2.4G2) (Fc Block)	TONBO bioscience	Cat#70-0161; RRID: AB_2621487
Purified anti-mouse CD45 (30-F11)	BioLegend	Cat#10302; RRID: AB_312967
Purified anti-mouse F4/80 (BM8)	BioLegend	Cat#123101; RRID: AB_893504
Purified anti-mouse Areginase-1 (D4E3M)	CST	Cat#93668; RRID: AB_2800207
Purified anti-mouse Nos2 (Rabbit polyclonal)	Abcam	Cat#ab15323; RRID: AB_301857
Purified anti-mouse PD-1 (RMP1-14)	Dr. Hideo Yagita (Juntendo University, Tokyo)	N/A

Purified anti-mouse CTLA-4 (UC10-4F10)	Dr. Hideo Yagita (Juntendo University, Tokyo)	N/A
Purified anti-mouse IL-34 (C054-35)	BioLegend	Cat#147202; RRID: AB_2563031
Purified anti-mouse CSF-1R (AFS98)	Bioxell	Cat#BE0213; RRID: AB_2687699
ChromPure Rat igG, whole molecule	Jackson Immuno Research LABORATPRIES, INC.	Cat#012-000-003; RRID: AB_2337136
Purified anti-human IL-34 (1D12)	Millipore	Cat#MABT493
Purified anti-human CD274 (E1L3N)	CST	Cat#13684; RRID: AB_2687655
Purified anti-human PD-1 (monoclonal)	Selleckchem	Cat#A2002; RRID: AB_2810223
Purified anti-human IL-34 (E0320E7)	BioLegend	Cat#361302; RRID: AB_2563033

Cell Culture Regents

RPMI-1640with L-Glutamine and Phenol Red	Fujifilm Wako Pure Chemical Industries	Cat#189-02025
D-MEM (high Glucose) with L-Glutamine and Phenol Red	Fujifilm Wako Pure Chemical Industries	Cat#044-29765
MEM α with L-Glutamine and Phenol Red	Fujifilm Wako Pure Chemical Industries	Cat#135-15175
Defined fetal bovine serum	Sigma Aldrich	Cat#F7524
Penicillin-Streptomycin Mixed Solution (100x)	Nacali Tesque	Cat#26253-84
MEM Non-Essential Amino Acid Solution (100x)	Nacali Tesque	Cat#06344-56
2.5g/l-Trypsin/1mmol/l-EDTA Solution, with Phenol Red	Nacali Tesque	Cat#32777-15

Critical Commercial Regents

TransIT-X2 [®] Dynamic Delivery System	Takara	Cat#V6104
LEGEND MAX [™] Mouse IL-34 ELISA Kit	BioLegend	Cat#439107
eBioscience [™] Fixation/Permeabilization Concentrate	Invitrogen [™]	Cat#00-5123-43
eBioscience [™] Fixation/Permeabilization Diluent	Invitrogen [™]	Cat#00-5223-56
eBioscience [™] Permeabilization Buffer (10X)	Invitrogen [™]	Cat#00-8333-56

BD Horizon™ Dri Tumor & Tissue Dissociation Reagent (TTDR)	BD bioscience	Cat#661563
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Experimental Models: Cell Lines

OV2944-HM-1	Japanese Collection of Research Bioresources	Cat#JCRB1208
CT26	Dr. Hidemitsu Kitamura (Hokkaido University, Hokkaido)	N/A
4T1	ATCC	Cat#CRL-2539

Experimental Models: Organisms/Strains

B6C3F1	Japan SLC, Inc.	N/A
Balb/c	Japan SLC, Inc.	N/A

Plasmids

pCAG-VSVG	Addgene	Cat#8454
pCMV-VSV-G-RSV-Rev	Addgene	Cat#35616
IL-34 CRISPR/Cas9 KO Plasmid (m)	Santa Cruz Biotechnology, Inc.	Cat#sc-429354
pLenti-EF1a-C-Myc-DDK-IRES-Puro	Origene	Cat#PS100085

Primer list for quantitative PCR analysis

Species	Gene	Forward (5'-3')	Reverse (5'-3')
	Gapdh	TCAAATGGGGTGAGGCCGGT	TTGCTGACAATCTTGAGTGA
	Cd3e	AAGTAATGAGCTGGCTGCGT	TCGTCACTGTCTAGAGGGCA
	Cd8a	GGATTGGACTTCGCCTGTGA	TGGGACATTTGCAAACACGC
	Mrc1	CTCTGTTCAGCTATTGGACGC	TGGCACTCCCAAACATAATTTGA
	lfng	AAGACAATCAGGCCATCAGCA	AGCGACTCCTTTTCCGCTTC
Mouse	Tnfa	TTCTATGGCCCAGACCCTCA	CTTGGTGGTTTGCTACGACG
	Nos2	ACATCGACCCGTCCACAGTAT	CAGAGGGGTAGGCTTGTCTC
	Arg1	GTATGACGTGAGAGACCACG	CTCGCAAGCCAATGTACACG
	Cxcl9	AATGCACGATGCTCCTGCA	AGGTCTTTGAGGGATTTGTAGTG
	Cxcl10	AGTGCTG CCGTCATTTTCTG	TCCCTATGGCCCTCATTCTCA
	Cxcl11	GTAATTTACCCGAGTAACGGC	CACCTTTGTCGTTTATGAGCCTT