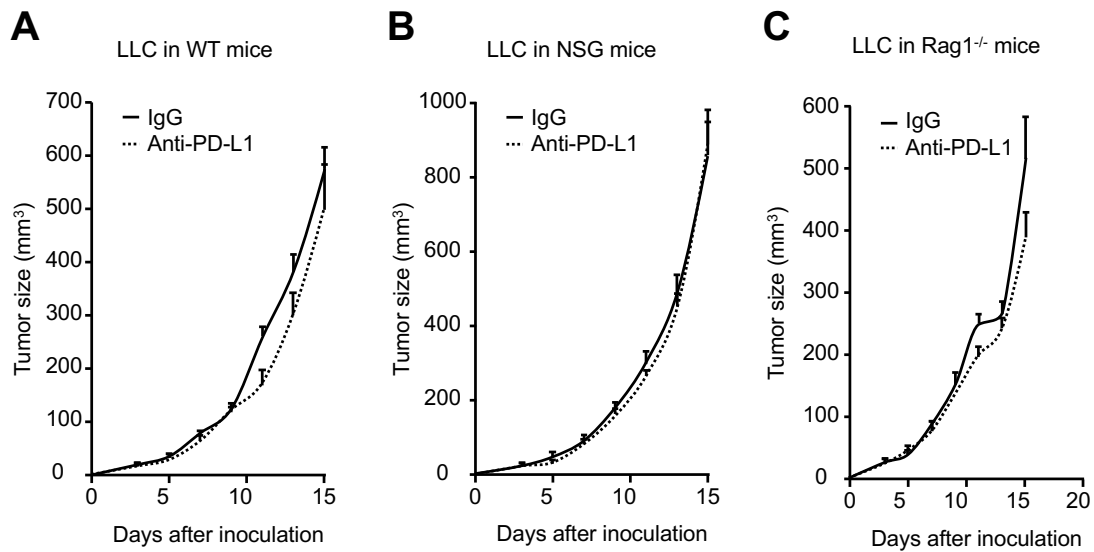
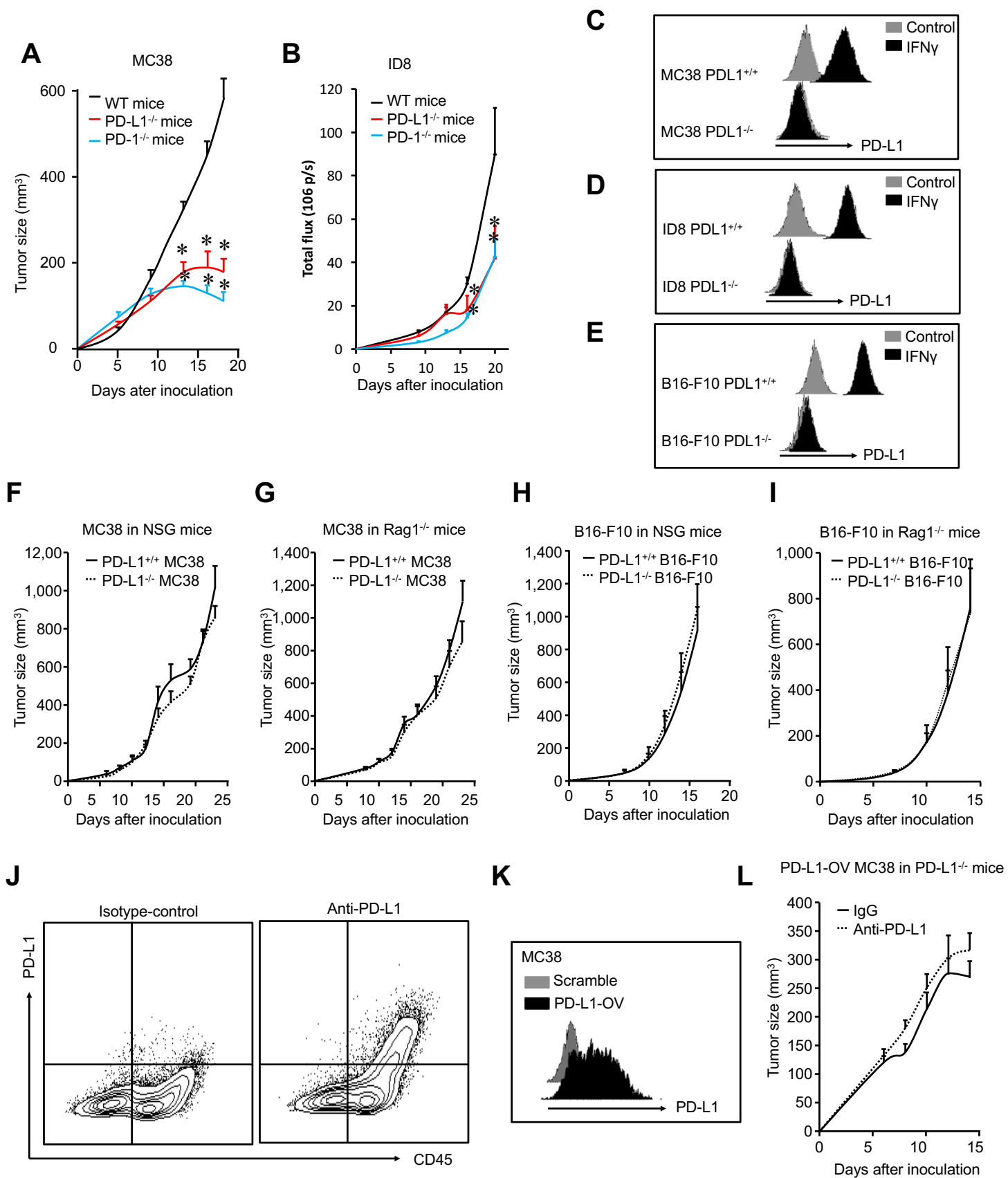
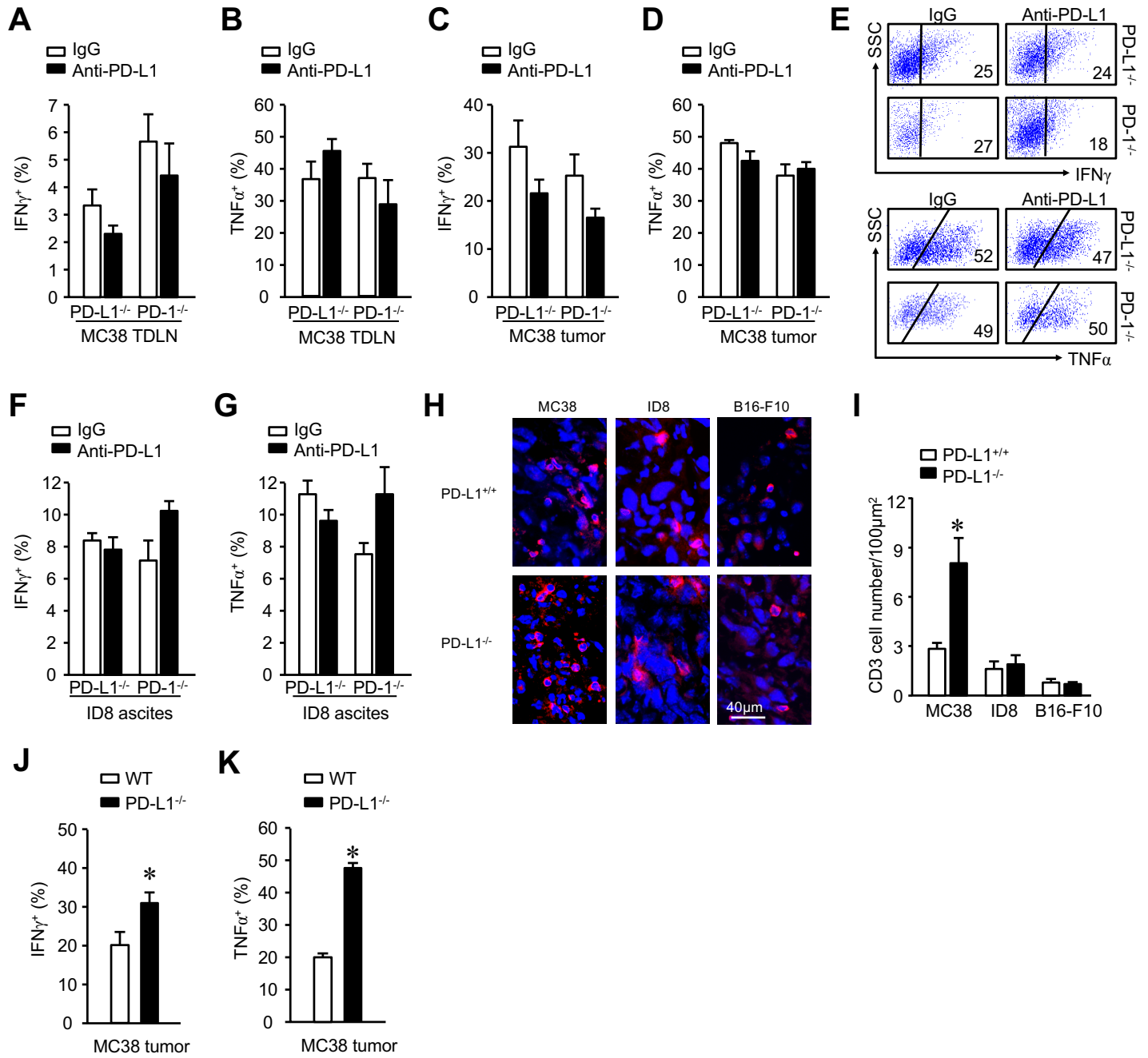


**Figure 1S**

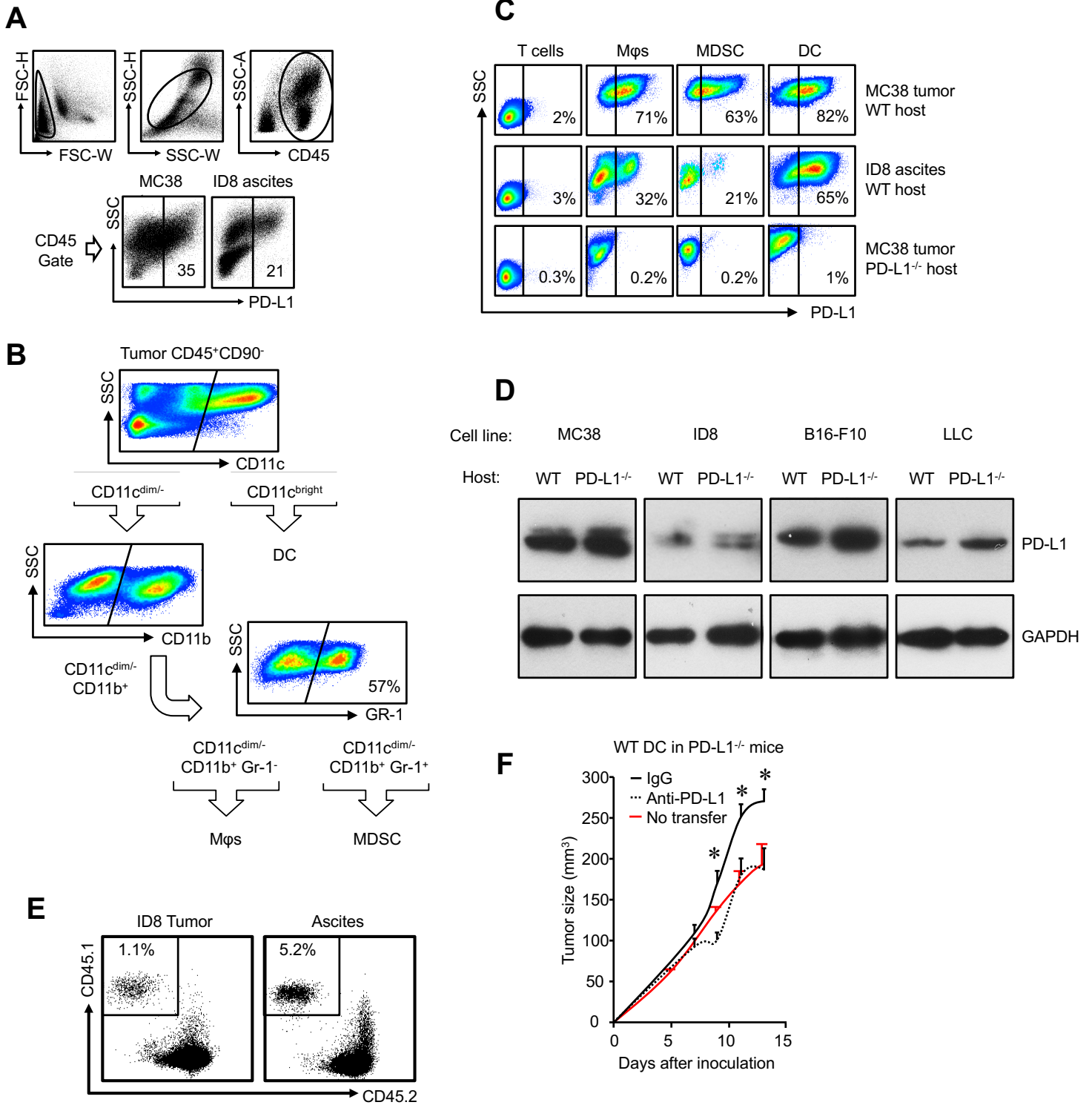


**Figure 2S**

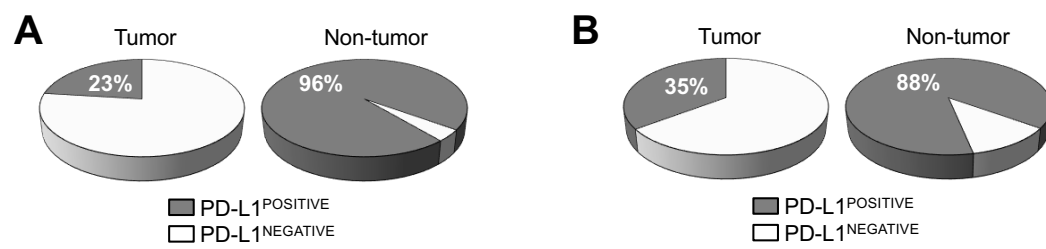
**Figure 3S**



**Figure 4S**



**Figure 5S**



## Supplementary figure legends

### Figure S1. Effect of anti-PD-L1 treatment on LLC volume in mice.

(A-C) WT, NSG, and Rag1<sup>-/-</sup> mice were inoculated with LLC tumor cells. The mice were treated from day 3 every 3 days with anti-PD-L1 or isotype control (rlgG1). Tumor volume was monitored. n = 5-7. Wilcoxon test was used for two-way comparisons.

### Figure S2. Expression and effect of tumor PD-L1 on tumor growth.

(A-B) WT, PD-L1<sup>-/-</sup>, and PD-1<sup>-/-</sup> mice were inoculated with MC38 (A), and ID8 (B) tumor cells. Tumor volume was monitored. n = 5-7. T-test was used for two-way comparisons (\*P < 0.05). (C-E) CRISPR PD-L1 homozygous PD-L1 knockout (PD-L1<sup>-/-</sup>) tumor cell clones were made for MC38 (C), ID8 (D), and B16-F10 (E). PD-L1<sup>-/-</sup> and control tumor cells were stimulated for 48 hours with or without IFN $\gamma$  (10ng/ml). PD-L1 expression was analyzed by FACS. One of 4 experiments is shown. (F-I) PD-L1<sup>-/-</sup> MC38 (F, G), PD-L1<sup>-/-</sup> B16-F10 (H, I) and wild type control tumor cells were inoculated into NSG and Rag1<sup>-/-</sup> mice. Tumor load was monitored. n=5 mice per group. (J) PD-L1<sup>-/-</sup> MC38 cells were inoculated into wild type mice. Single cells were prepared from tumor tissues on day 15. PD-L1 expression was analyzed by FACS on CD45<sup>+</sup> immune cells and CD45<sup>-</sup> tumor cells. One of 5 is shown. (K-L) PD-L1 overexpressed (PD-L1-OV) or scrambled MC38 cells were analyzed for PD-L1 expression by FACS (K). PD-L1<sup>-/-</sup> mice were inoculated with PD-L1-OV MC38 cells (L, n = 8) and treated with anti-PD-L1 or isotype control (rlgG1). Tumor volume was monitored. Wilcoxon test was used for two-way comparisons. (\*P < 0.05).

### Figure S3. Effect of anti-PD-L1 on T cell effector cytokine expression.

(A-G) T cell effector cytokines were analyzed with intracellular staining in MC38 tumor draining lymph nodes (TDLN) (A, B), MC38 tumor tissues (C-E), and ID8 tumor ascites (F, G) in PD-L1<sup>-/-</sup> and PD-1<sup>-/-</sup> mice. Data are expressed as the mean  $\pm$  SEM (n = 3-5 per group). Representative original flow cytometry data are shown (E). (H, I) T cell infiltration was analyzed with immunofluorescence staining for CD3 in tumors from PD-L1<sup>+/+</sup> and PD-L1<sup>-/-</sup> mice. Data are expressed as the mean  $\pm$  SEM (I, n = 10 per group). Representative staining images are shown (H). T-test was used for two-way comparisons (\*P < 0.05). (J, K) T cell effector cytokines were analyzed with intracellular staining in MC38 tumor in WT and PD-L1<sup>-/-</sup> mice, n = 5. T-test was used for two-way comparisons (\*P < 0.05).

### Figure S4. APC subsets in tumor tissues and tumor PD-L1 expression *in vivo*.

(A-C) PD-L1 expression in tumor infiltrating immune cells. (A) PD-L1 expression on CD45<sup>+</sup> immune cells in MC38 tumor tissues. (B) FACS gating on different immune cell subsets in MC38 tumor tissues. Representative flow cytometry data showed gates on CD45<sup>+</sup>CD90<sup>-</sup> cells, DCs, macrophages, and MDSCs. (C) The percentages of PD-L1 expression in each immune cell subset in MC38 and ID8 ascites. One of five replicates is shown. (D) PD-L1 expression on tumor cells *in vivo*. Tumor cells were isolated from wild type and PD-L1<sup>-/-</sup> mice. PD-L1 expression was detected by Western blots in tumor cells. Experiments were performed in triplicates; representative replicate is shown. (E) CD45.1<sup>+</sup> peritoneal APCs were injected into ID8 tumor bearing CD45.2<sup>+</sup> mice. 48 hours after APC injection, CD45.1<sup>+</sup> APCs were examined in ID8 tumor tissues and ascites by flow cytometry. n = 5. (F) MC38 tumor bearing PD-L1<sup>-/-</sup> mice were adoptively transferred with (black) or without (red) WT DCs. Mice were treated with anti-PD-L1 or isotype IgG1. Tumor volume was monitored. Wilcoxon test was used for two-way comparisons (n = 7, \*P < 0.05).

**Figure S5. PD-L1<sup>+</sup> APCs and tumor cells in patients with melanoma and ovarian cancer.**

(A-B) Percentages of patients with PD-L1<sup>+</sup> tumor cells or PD-L1<sup>+</sup> non-tumor cells were shown in melanoma (A) and ovarian cancer (B) tissues.

**Supplementary Table 1. Characteristics of melanoma patients and PD-L1 expression pattern**

Patient	Gender	Response	tissue type	Tumor	Non-tumor		(%)	
					positivity	score	mDC	MQ
1	M	CR	sinus primary	negative	Positive	4	50	30
2	M	CR	lymph node metastasis	single cells	Positive	3	20	0
3	M	CR	skin primary	negative	Positive	3	5	10
4	M	CR	subcutaneous metastasis	negative	Positive	3	20	5
5	M	CR	lymph node metastasis	negative	Positive	3	50	5
6	F	CR	lymph node metastasis	negative	Positive	4	20	50
7	F	CR	skin primary	positive	Positive	3	30	20
8	M	CR	lymph node metastasis	negative	Positive	4	50	50
9	M	CR	lymph node metastasis	single cells	Positive	3	20	40
10	M	CR	skin primary	single cells	Positive	4	50	40
11	M	PR	sinus metastasis	negative	Positive	2	20	0
12	M	PR	brain metastasis	negative	Positive	1	1	5
13	M	PR	subcutaneous metastasis	negative	Positive	3	30	20
14	M	PR	soft tissue metastasis	negative	Positive	1	5	1
15	M	PR	lymph node metastasis	negative	Positive	2	20	10
16	F	PR	skin primary	negative	Positive	1	1	10
17	F	SD	uvea primary	negative	Positive	3	50	10
18	M	SD	skin primary	positive	Positive	1	5	1
19	M	SD	lymph node metastasis	negative	Positive	3	30	30
20	M	SD	soft tissue metastasis	positive	Positive	1	1	1
21	F	SD	lymph node metastasis	negative	Positive	3	20	50
22	M	PD	lymph node metastasis	negative	Positive	2	5	0
23	F	PD	lymph node metastasis	negative	Positive	2	30	20
24	F	PD	skin/vulvar primary	negative	negative	0	0	0
25	M	PD	lung metastasis	negative	Positive	1	1	1
26	F	PD	dermal/subcutaneous metastasis	negative	Positive	1	1	1



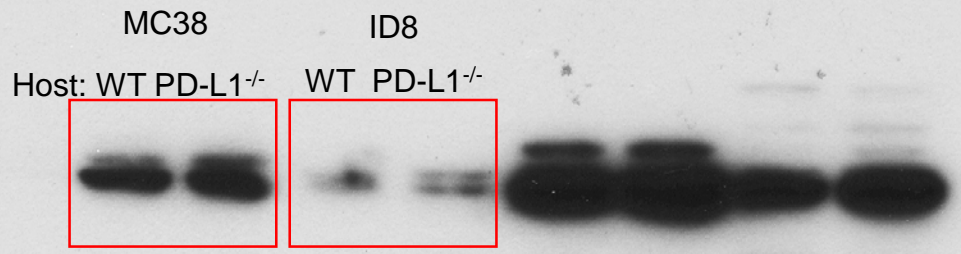
**Supplementary Table 2. Characteristics of ovarian cancer patients and PD-L1 expression pattern**

Patient	Gender	Response	Tumor	Non-tumor		(%)	
				positivity	score	mDC	MQ
1	F	CR	Negative	Positive	4	20	50
2	F	CR	Negative	Positive	3	10	20
3	F	PR	Negative	Positive	4	40	40
4	F	SD	Positive	Positive	4	10	20
5	F	SD	Negative	Positive	2	1	10
6	F	SD	Negative	Positive	4	40	50
7	F	SD	Negative	Positive	2	1	10
8	F	SD	Negative	Positive	2	10	5
9	F	PD	Positive	Positive	3	20	40
10	F	PD	Negative	Positive	1	1	1
11	F	PD	Negative	Positive	2	5	5
12	F	PD	Positive	Negative	0	0	0
13	F	PD	Positive	Negative	0	0	0
14	F	PD	Negative	Positive	2	5	5
15	F	PD	Negative	Positive	2	5	5
16	F	PD	Positive	Positive	2	5	10
17	F	PD	Negative	Positive	2	5	1

B5H1  
Tissue

1

Fig S4D. Anti-PD-L1



3  
1

PD-L1<sup>-/-</sup> cell line  
→

Fig S4D. Anti-PD-L1 B16-F10

Host: WT PD-L1<sup>-/-</sup>

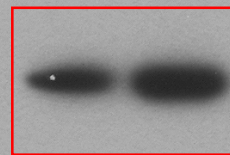
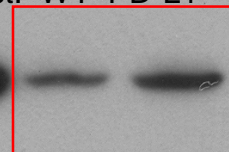


Fig S4D. Anti-PD-L1 LLC

LLC

Host: WT PD-L1<sup>-/-</sup>



1  
B16-F10

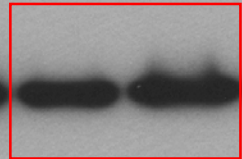
2  
LLC

3  
control

Fig S4D. Anti-GAPDH

B16-F10

Host: WT PD-L1<sup>-/-</sup>



2 GAPDH  
- 11.3  
tissue #

MC38

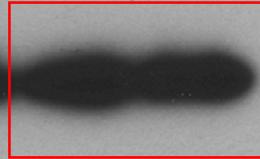
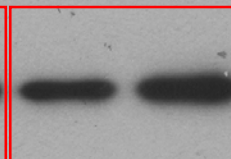
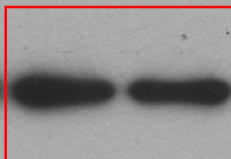
ID8

LLC

Host: WT PD-L1<sup>-/-</sup>

Host: WT PD-L1<sup>-/-</sup>

Host: WT PD-L1<sup>-/-</sup>



4  
- 2#

6 cell line

