#### **Supplemental figure legends**

**Supplemental Figure 1. MHC class-I expression in B16 tumor cells with or without IFN-γ signaling.** MHC class-I expression on B16 cells after IFN signaling sensitization in vitro (**A**) or IFN signaling deficiency in vivo (**B**). MHC class-I expression on B16 cells after IFN signaling sensitization in vitro (**A**) or IFN signaling deficiency in vivo (**B**).

#### Supplemental Figure 2. Proliferation and apoptosis of B16-OVA cells after

overexpression or knockdown of USP18. USP18 expression in B16-OVA cells was overexpressed or knocked down using lentivirus vector and confirmed by Western blot assay (**A** & **B**). Exponentially growing B16-OVA tumor cells stably expressing shUSP18 (B16-OVA-shUSP18) were pulsed with 10  $\mu$ M BrdU and cell proliferation was analyzed by flow cytometry (**C**). The apoptosis status of B16-OVA-shUSP8 cells was analyzed by flow cytometry staining for Annexin/PI (**D**).

Supplemental Figure 3. Overexpression or knockdown USP18 in B16-OVA cells affects tumor growth.  $3 \times 10^5$  B16-OVA-GFP, B16-OVA-USP18 (2) or B16-OVA-shUSP18(2) tumor cells were intravenously inoculated into C57BL/6 mice. Tumor growth was monitored 3weeks after inoculation. <u>B16-OVA-USP18 (2) and</u> <u>B16-OVA-shUSP18(2) cells lines are from different clone or shRNA.</u>

Supplemental Figure 4. Adhesion, invasion and migration activity of B16 cells after overexpression or knockdown of USP18 expression. The ability of B16-OVA, B16-OVA-USP18 and B16-OVA-shUSP18 tumor cells adhering to fibronectin-coated plates were analyzed (**A**). The invasion and migration ability of B16-OVA, B16-OVA-USP18 and B16-OVA-shUSP18 cells was analyzed in transwell plates coated with fibronectin (**B**) or matrigel (**C**).

Supplemental Figure 5. Overexpression of USP18 in B16 cells does not affect cytokine secretion in vitro but does affect chemokine production in the tumor microenvironment. Cytokine and chemokine secretion from culture supernatant of B16-OVA, B16-OVA-USP18 and B16-shUSP18 tumor cells was analyzed by cytokine array (**A**). Chemokine mRNA level in B16-OVA-GFP and B16-OVA-USP18 tumor microenvironment were analyzed by qRT-PCR (**B**).

# Supplemental Figure 6. ISG15 levels in B16-OVA-GFP, B16-OVA-USP18 or B16-OVA-shUSP18 tumor lysate. ISG15 levels in B16-OVA-GFP,

B16-OVA-USP18 or B16-OVA-shUSP18 tumor lysate was analyzed by Western blot assay.

Supplemental Figure 7. OVA expression in B16-OVA-GFP, B16-OVA-USP18 and B16-OVA-shUSP18 tumor cells. Western blot assay of OVA protein expression in B16-OVA tumor cells with overexpression or knockdown of USP18.

Supplemental Figure 8. Overexpression of USP18 in B16-OVA cells affects
subcutaneous tumorigenesis. C57BL/6 mice received subcutaneous inoculation of 1
× 10<sup>6</sup> B16-OVA-GFP and B16-OVA-USP18 cells. Effector (CD44<sup>+</sup>) T cells in tumor
(A) and spleen (B) was analyzed. CD4<sup>+</sup>Foxp3<sup>+</sup> regulatory T cells located in spleen

were analyzed by flow cytometry (C).



# sFigure 1.





sFigure 2.

GFP





USP18(2)

shUSP18(2)











### sFigure 4.





В



# sFigure 5.



### sFigure 6.

# GFP USP18 shUSP18



# OVA





# sFigure 7.



USP18

54.4

► CD4

82.7

CD8

S.

≁



## sFigure 8.

С