Figure S1.

ChiP analysis of WM164 wild type melanoma cell line stimulated with IL-6 (30ng/mL) and collected at baseline (time 0) or at 0.25, 0.5, 1, 2, 4, 6, 8, 12 and 24 hours after stimulation, assessed with PS5-CTD Pol II (A) and acetyl H3 (B) antibodies, followed by quantitative real-time PCR analysis of the 7 different primers described above (detailed in Table 1). (C) WM164 cells were treated with IFNγ, and then subjected to immunoblot to evaluate HDAC6, PD-L1, ac-Tubulin and GAPDH. (D) WM164 cells were treated with IFNγ, then total RNA was isolated and the expression of PD-L1 analyzed by qRT-PCR. NT, HDAC6KD and STAT3KD WM164 melanoma cells were transfected with either STAT3C-flag plasmid (E) or empty vector (F) and then stimulated with IFNγ (100 ng/mL) or left untreated. Then, PD-L1 was evaluated by flow cytometry.

Figure S2.

WM164 cells were treated with LBH589 10nM or 20nM (A and C) or MGCD0103 1μM or 2μM (B and D) for 24 hours and then stimulated with IL-6 or left untreated. Then, the presence of HDAC6, STAT3, P-STAT3 Y705, P-STAT3 S727, acetylated STAT3, PD-L1 and GAPDH was assessed by flow cytometry (C and D) and immunoblot (E and F).

Figure S3.

(A) NT and HDAC6KD B16-F10 murine melanoma cells were stimulated with IL-6 (30ng/ml) or left untreated. Next, the presence of HDAC6, acetylated tubulin, STAT3, P-STAT3Y705, P-STAT3-S727, acetylated STAT3 and GAPDH was assessed by immunoblot. (B) B16-F10-luc melanoma cell line were treated with the HDAC6 inhibitor Nexturastat for 24 hour and then stimulated with IL-6 (30 ng/mL) or left untreated. The presence of HDAC6, acetylated tubulin,

STAT3, P-STAT3Y705, P-STAT3-S727, acetylated STAT3 and GAPDH was assessed by immunoblot.

Figure S4.

NT and HDAC6KD WM164 human melanoma cells were stimulated with IL-6 (30ng/ml) or left untreated. Next, the presence of PD-L2, TRAIL-R1, TRAIL-R2, B7-H3 and B7-H4 were measured by western blot (A) and flow cytometry (B).







