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p38 MAPK-inhibited dendritic cells induce superior antitumour immune responses and overcome regulatory T-cell-mediated immunosuppression

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Dendritic cell (DC)-based cancer immunotherapy is a promising method, but so far has demonstrated limited clinical benefits. Regulatory T cells (Tregs) represent a major obstacle to cancer immunotherapy approaches. Here we show that inhibiting p38 MAPK during DC differentiation enables DCs to activate tumour-specific effector T cells (Teff), inhibiting the conversion of Treg and compromising Treg inhibitory effects on Teff. Inhibition of p38 MAPK in DCs lowers expression of PPAR γ , activating p50 and upregulating OX40L expression in DCs. OX40L/OX40 interactions between DCs and Teff and/or Treg are critical for priming effective and therapeutic antitumour responses. Similarly, p38 MAPK inhibition also augments the T-cell stimulatory capacity of human monocyte-derived DCs in the presence of Treg. These findings contribute to ongoing efforts to improve DC-based immunotherapy in human cancers.

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endritic cell (DC)-based immunotherapy holds great promise for treating malignancies¹. However, the results of most DC-based clinical trials have been disappointing². In cancer patients, $CD4^+CD25^+Foxp3^+$ regulatory T cells (Tregs) accumulate in tumours and secondary lymphoid organs by the mechanisms of expansion of Treg or the conversion of effector T cells (Teff)^{3,4}. Tumour-associated Tregs likely contribute to the suppressive milieu, leading to the suppression of T-cell responses partly via the inhibition of DC functions⁵. Thus, modulating the suppressive function of Treg is essential for induction of effective antitumour immunity^{6,7}.

Among the possible strategies to target Treg *in vivo*, depletion of CD25⁺ T cells by anti-CD25 antibodies may not be a viable approach because both Treg and activated Teff are depleted, inducing conversion of peripheral precursors into Treg^{4,8-10}. A unique surface marker for Treg has not been identified, and an alternative approach to target Treg is the use of cytotoxic T lymphocyte antigen 4 (CTLA4)-blocking antibodies or glucocorticoid-induced tumour necrosis factor receptor (GITR) receptor agonist antibodies to revert Treg-mediated suppression^{8,11}. However, these strategies can result in over-activating of nonspecific T-cell immunity and have been linked to severe autoimmunity in multiple organs, thus limiting their further clinical applications^{12,13}. Therefore, a more desirable approach is needed to modulate the suppressive function of Treg and activate Teff in an antigen-specific manner.

In an earlier study, we demonstrated that lipopolysaccharide (LPS)-induced activation of p38 mitogen-activated protein kinase (MAPK) has a detrimental effect on the generation of immature DCs (iDCs) *in vitro*¹⁴. We and others have also shown that tumour-derived suppressive factors inhibit the differentiation and function of DCs by upregulating p38 MAPK activity in DC precursors in both murine tumour models and cancer patients^{15–18}. These abnormalities in the phenotype and T-cell stimulatory capacity of DCs could be restored by inhibiting p38 MAPK activity in progenitor cells^{15–18}. Based on these observations, we hypothesized that p38 MAPK may attenuate antigen presentation and have an essential role in maintenance of self-tolerance in DC.

In this study, we inhibited p38 MAPK activity in DC precursors using the inhibitor SB202190, and inhibitor SB203580 and/or p38 α -specific small intefering RNA (siRNA) transfection for confirmation. The inhibition of p38 MAPK results in sharply decreased PPAR γ expression in DCs, which reduces functional inhibition of p50 transcriptional activities by PPAR γ . In turn, p50 activation upregulates surface expression of OX40L on DCs, increasing their immunostimulatory potency, activating antigen-specific Teff, inhibiting Treg conversion and function, and facilitating tumour rejection.

Results

p38 MAPK-inhibited DCs activate Teff in the presence of Treg. Among the four different p38 isoenzymes, p38α was the only p38 MAPK isoenzyme detected in the isolated bone marrow (BM) cells and generated iDCs and mature DCs (mDCs) from wild-type C57BL/6 mice (WT-B6 mice) (Supplementary Fig. 1a). We chose to use SB202190 and SB203580, two specific inhibitors of the α- and β-isoforms of p38 MAPK¹⁹. Treatment with 1.5 µM SB202190 or SB203580, but not the inactive analogue SB202474 (ref. 20), was sufficient to inhibit the p38 MAPK activity in BM cells, as determined by our previous studies¹⁶, and by the blockage of the phosphorylation of its downstream kinase MAPKAPK-2 (Supplementary Fig. 1b). To examine the role of p38 MAPK in regulating DC generation and maturation (all generated from BM cells from WT-B6 mice unless indicated specifically), flow cytometry was used to analyse the surface expression of various molecules related to antigen presentation. SB202190- and SB203580-treated, but not SB202474-treated, iDCs expressed higher levels of DC-related molecules compared with control iDCs (Supplementary Fig. 1c). After maturation, SB202190-treated mDCs (mSBDCs) and SB203580-treated mDCs (mSB80DCs) showed significantly higher levels of major histocompatibility complex (MHC) class II, CD80, CD86, CD40 and OX40L than dimethylsulphoxide (DMSO)-treated control mDCs and SB202474-treated mDCs (mSB74DCs) (Fig. 1a,b and Supplementary Fig. 1d). In addition, p38 MAPK inhibition enhanced interleukin (IL)-12 production by mSBDCs and mSB80DCs (Supplementary Fig. 1e). All these cells produced undetectable or very low levels of cytokines such as IL-6, $TNF-\alpha$, IL-10 and interferon (IFN)- γ (ranging from 0 to 50 pg ml⁻¹). In line with these results, knockdown of p38a in DC progenitor cells by p38\alpha-specific siRNA during the differentiation also induced the upregulated expression of DC-related surface molecules as compared with nonspecific siRNA-treated (Si-non-) mDCs (Supplementary Fig. 2). Collectively, these results suggested that p38 MAPK negatively regulated the generation and immunogenicity of BM-derived DCs (BMDCs).

The enhanced immunogenicity of mSBDCs and mSB80DCs may enable them to promote proliferation of activate Teff in the presence of Treg. We used a carboxyfluorescein diacetate succinimidyl ester (CFSE) dilution assay to test the proliferation of Teff. As shown in Fig. 1c and Supplementary Fig. 3, mSBDCs, mSB80DCs or p38a-specific siRNA-treated (Si-p38a-) mDCs induced a profound proliferation of CD4⁺ Teff in the presence of Treg, whereas the proliferation of CD4+ Teff stimulated by mDCs, mSB74DCs or Si-non-mDCs was largely inhibited. Notably, the proliferation of CD4⁺ Teff stimulated by mSBDCs or mSB80DCs in the presence of Treg was even stronger than those stimulated by mDCs in the absence of Treg. We found that CD4⁺ Teff proliferation induced by mSBDCs, mSB80DCs or Si-p38a-mDCs was inhibited significantly by Treg when OX40Lblocking monoclonal antibody (mAb) was added (Fig. 1d and Supplementary Fig. 3). We also screened other molecules that might have contributed to this effect by using functional grade blocking mAbs targeting CD80, CD86, CD40 or IL-12 (Supplementary Fig. 1f-h), and excluded their involvement (Fig. 1d). These data suggest that the enhanced stimulatory function of mSBDCs and mSB80DCs is mediated exclusively by upregulated OX40L expression.

Flow cytometry analysis showed that mSBDCs expressed significantly higher levels of surface OX40L, and Teff stimulated by mSBDCs expressed significantly upregulated OX40L than T cells stimulated by mDCs (Supplementary Fig. 4a,b). As Tregs do not express OX40L²², we next examined the contribution of cell expression of OX40L. We used OX40L-specific siRNA-treated mSBDCs (Si-OX40L-mSBDCs; Supplementary Fig. 4b), and found that CD4⁺ Teff proliferation induced by Si-OX40LmSBDCs was subject to Treg-mediated suppression (Fig. 1e). This Si-OX40L-mSBDC-induced Teff proliferation decreased further to the level of mDC-induced CD4⁺ Teff proliferation in the presence of Treg when OX40L-blocking antibody was added (Supplementary Fig. 4c), indicating that mSBDC-expressed OX40L had a major role in counteracting Treg-mediated suppression of Teff proliferation, and that Teff-expressed OX40L may also be important. These results suggested that mSBDCs may be able to induce tumour-specific T-cell responses even in the presence of Treg.

OX40 costimulation inhibits Treg and activates Teff. In the absence of Treg, the OX40/OX40L signalling appears to only have a minimal effect on proliferation of $CD4^+$ Teff (Fig. 2a and



Figure 1 | OX40L expression by mSBDC is critical for mSBDC-induced Teff proliferation in the presence of Treg. (a) Expression of OX40L on treated murine BMDCs. Gates were preset on CD11c⁺ cells as determined by FACS analysis. Numbers are percentages of OX40L-expressing DCs. (b) Pooled data (n = 3) depicting the percentages of OX40L-positive cells among CD11c⁺ mDCs, mSB74DCs, mSBDCs and mSB80DCs. Error bars represent s.d. (**c-e**) Proliferation of CFSE-labelled wild-type (WT) CD4⁺ CD25⁻ Teff (1 × 10⁵) in the absence or presence of unlabelled CD4⁺ CD25⁺ Treg (1 × 10⁵) induced by various mDCs (1 × 10⁴) plus anti-CD3 antibody (1µg ml⁻¹). Both Teff and Treg were isolated from spleens of naive C57BL/6 mice. (**c**) CD4⁺ CD25⁻ Teff was measured by FACS. Numbers indicate percent CFSE^{lo}/proliferated Teff. (**d**) Control lgG or the specific mAbs were added to the coculture of Teff, Treg and mSBDCs or mSB80DCs. Sixty hours later, Teff proliferation was measured by FACS. (**e**) CD4⁺ CD25⁻ Teff were cocultured with untreated, nonspecific siRNA- (Si-non-) or OX40L-specific siRNA-treated mDCs or mSBBCS. CD4⁺ CD25⁺ Tregs were added as indicated. Sixty hours later, Teff proliferation was measured by FACS. CP4⁺ CD25⁺ Tregs were added as indicated. Sixty hours later, Teff proliferated mDCs or mSBDCs. CD4⁺ CD25⁺ Tregs were added as indicated. Sixty hours later, Teff proliferation was measured by FACS. (**e**) CD4⁺ CD25⁻ Teff were cocultured with untreated, nonspecific siRNA- (Si-non-) or OX40L-specific siRNA-treated mDCs or mSBDCs. CD4⁺ CD25⁺ Tregs were added as indicated. Sixty hours later, Teff proliferation was measured by FACS. Prese were added as indicated. Sixty hours later, Teff proliferation was measured by FACS. Representative results from one of three performed experiments are shown. *P* values were calculated with Student's *t*-test.



Figure 2 | Role of OX40 signalling in counteracting Treg-mediated suppression by mSBDCs. (a-c) Proliferation of splenic CD4⁺CD25⁻ Teff in the absence or presence of CD4⁺CD25⁺ Treg isolated from naive C57BL/6 mice. CD4⁺CD25⁻ Teff (1×10^5) from WT C57BL/6 mice (**a**), WT or OX40^{-/-} C57BL/6 mice (**b**), or OX40^{-/-} C57BL/6 mice (**c**) were labelled with CFSE and cocultured with mDCs or mSBDCs (1×10^4) plus anti-CD3 antibody (1µg ml⁻¹). Control medium, antibodies and/or unlabelled CD4⁺CD25⁺ Treg (1×10^5) were added as indicated. Sixty hours later, CFSE intensity of Teff was measured by FACS. Numbers indicate percent CFSE^{lo}/proliferated Teff. Results are representative of three independent experiments. (**d**) CD4⁺CD25⁺ Treg isolated from C57BL/6 mice were pretreated with medium, mDCs, mSBDCs or mSBDCs plus anti-OX40L antibody for 5 h. CFSE-labelled OT-II Teff (1×10^5) and peptide-pulsed mDCs (1×10^4) were cocultured in the presence of medium or with CD4⁺CD25⁺ Treg isolated from the pretreatment condition at the indicated Treg to OT-II Teff ratios. Seventy-two hours later, CFSE intensity of OT-II Teff was examined by FACS. Representative data from one of two repeated experiments are shown. *P* values were calculated with Student's *t*-test.

Supplementary Fig. 3). As Tregs constitutively express OX40 and activated Teff but not DCs express OX40 (refs 21,23-25), we next determined the contribution of cell expression of OX40 for the activation of Teff by mSBDCs in the presence of Treg. We first examined the role of OX40 expression on Teff. As shown in Fig. 2b, there was only a marginal decrease in the proliferation of $OX40^{-/-}$ CD4⁺ Teff, as compared with WT Teff, in response to mSBDCs or mDCs. However, when Tregs were present, the proliferation of the $OX40^{-/-}$ CD4⁺ Teff stimulated by mSBDCs decreased significantly as compared with those of WT Teff, indicating that loss of OX40 on CD4⁺ Teff rendered them more sensitive to Treg-mediated suppression as compared with their WT counterparts when stimulated by mSBDCs. In addition, we also investigated the role of Treg-expressed OX40 in Tregmediated suppressive function. Since $OX40^{-/-}$ Treg from $OX40^{-/-}$ mice showed impaired suppressive function²⁴, we could not use OX40^{-/-} Treg in our studies. We cocultured mSBDCs, OX40^{-/-} CD4⁺ Teff and Treg, in which OX40/ OX40L costimulation only occurs between mSBDCs and Treg, and found that the proliferation of $OX40^{-/-}$ CD4⁺ Teff induced by mSBDCs was significantly inhibited when OX40Lblocking antibody was added (Fig. 2c), suggesting that interaction

between mSBDCs and Treg via OX40/OX40L costimulation may lead to reduced inhibitory ability of Treg. Furthermore, mSBDCpretreated Treg became less effective in inhibiting the proliferation of OT-II Teff compared with control or mDCpretreated Treg in an OX40L-dependent manner (Fig. 2d). Similar results were also obtained from CD8⁺ T cells stimulated by mSBDCs in the presence or absence of Treg (Supplementary Fig. 5). Thus, these findings indicate the importance of OX40/ OX40L signalling in mSBDC-induced resistance to Treg inhibition of Teff proliferation.

OX40 costimulation by mSBDCs inhibits Teff conversion to Treg. We found a significantly smaller Treg population (CD4⁺ CD25⁺Foxp3⁺) among CD4⁺ T cells in the lymph nodes of mSBDC-immunized mice as compared with mDC-immunized mice (Fig. 3a), indicating that mSBDCs inhibited the expansion of preexisting Treg and/or conversion of CD4⁺CD25⁻ Teff to Treg. Since mSBDCs and mDCs had similar stimulatory effects on expansion of Treg (Supplementary Fig. 6a), we hypothesized that mSBDCs may inhibit the conversion of CD4⁺CD25⁻ Teff to Treg. As shown in Fig. 3b, mSBDCs generated significantly



Figure 3 | mSBDCs inhibit Teff to Treg conversion by OX40 costimulation. (a) FACS analysis of Treg population in pooled draining lymph nodes 10 days after the second immunization. C57BL/6 mice were immunized twice with 1×10^6 OT-II peptide-loaded mDCs or mSBDCs with or without anti-CD25 antibody treatment. Numbers represent the percentages of CD25⁺ Foxp3⁺ Treg among gated CD4⁺ T cells. (**b,c**) Conversion of Teff into Treg was suppressed by mSBDC stimulation. CD4⁺ CD25⁻ Foxp3⁻ Teff (1×10^5) isolated from WT or $OX40^{-/-}$ C57BL/6 mice were cocultured with mDCs or mSBDCs (1×10^4) in the presence of anti-CD3 antibody ($1 \mu g m I^{-1}$) and IL-2 ($20 U m I^{-1}$) with the addition of varying doses of TGF- β 1 (**b**) or TGF- β 1 ($2 n g m I^{-1}$) plus antibodies indicated (**c**). Numbers indicate the percentages of CD25⁺ Foxp3⁺ Treg gated on CD4⁺ T cells in 4-day cultures. (**d**) Schematic diagram of Smad3 binding site in Foxp3 enhancer I region and Stat3 binding site in Foxp3 enhancer II region. (**e**,**f**) CD4⁺ CD25⁻ Teff were cocultured with mDCs or mSBDCs in the presence of TGF- β 1 and IL-2 for 4 days. CD4⁺ T cells were purified for analyses. (**e**) ChIP assay was performed using Smad3 antibody or control IgG. Precipitated DNA was subjected to RT-PCR using primers targeting enhancer I region. Error bars represent s.d. (n = 3). (**f**) ChIP assay was performed using Stat3 antibody or control IgG. Precipitated DNA was subjected to RT-PCR using primers. *P* values were calculated with Student's *t*-test.

fewer CD4⁺CD25⁺Foxp3⁺ Treg than mDCs *in vitro* cultures of CD4⁺CD25⁻ Teff without exogenous TGF- β 1. Surprisingly, when TGF- β 1 induced a large number of Treg from Teff when stimulated with mDCs, mSB74DCs or Si-non-mDCs, mSBDCs, mSB80DCs and Si-p38 α -mDCs strongly inhibited the conversion of Teff to Treg (Fig. 3b and Supplementary Fig. 6b,c).

To shed light on the mechanisms driving these effects, we screened IFN- γ , IL-4, IL-12, IL-6, OX40L and other molecules in mSBDCs for their potential contribution to the inhibition of Teff conversion to Treg^{7,26}. Our results showed that mAbs blocking OX40L, but not other molecules (Supplementary Fig. 6d), almost completely abrogated the inhibitory effect of mSBDCs on Treg conversion (Fig. 3c), indicating that OX40/OX40L signalling may be involved. In addition, mSBDCs and mDCs had similar ability

in conversion of $OX40^{-/-}$ Teff into Treg in the presence or absence of exogenous TGF- β 1 (Fig. 3c). Moreover, *in vivo* blocking OX40/OX40L pathway by injecting OX40L-blocking antibodies restored the generation of Treg population in tumourdraining lymph nodes from mSBDC-immunized mice to the same level of mice immunized with mDCs (Supplementary Fig. 7). To elucidate the mechanism underlying mSBDC-inhibited conversion of Treg, we investigated the binding of phosphorylated (p)Smad3 to the enhancer region I of Foxp3 promoter, in light of a previous study showing that the binding of pSmad3 to the enhancer region I increases Foxp3 transcription, whereas the binding of pStat3 to the conserved enhancer region II works as a gene silencer during Foxp3 transcription (Fig. 3d)²⁷. Our results show that, although the level of pSmad3 and pStat3 did not change in CD4⁺ T cells cocultured with mDCs or mSBDCs in the presence of TGF- β 1 (Supplementary Fig. 6e), binding of pSmad3 to the enhancer region I and pStat3 to the enhancer region II were significantly suppressed or enhanced, respectively, in mSBDC-cocultured CD4⁺ T cells in an OX40L-dependent manner (Fig. 3e,f). These results suggest epigenetic inhibition of Treg generation by mSBDCs. Together, these findings show that OX40/OX40L signalling provided by mSBDCs may contribute to the reduced generation of immunosuppressive cells *in vitro* and *in vivo*.

Sustained p50 activation mediates OX40L expression in mSBDCs. Our results indicate that both enhanced stimulatory function of mSBDCs and the inhibition of Teff conversion into Treg by mSBDCs are mediated exclusively by OX40L. We next elucidated the molecular mechanism underlying p38 inhibitionmediated upregulation of OX40L in mSBDCs. Because nuclear factor- κB (NF- κB) pathway contributes to the immunogenicity of DCs and Ox40l promoter contains a potential NF-κB binding site (5'-GGGAAATTCA-3', -84 to -75, upstream of Ox40l), we hypothesized that the NF- κ B pathway has an important role in the increased expression of OX40L on mSBDCs. By comparing NF-kB molecules in the nuclear fractions of mDCs and mSBDCs, we found that inhibition of p38 MAPK resulted in slightly increased nuclear translocation of RelA and RelB and robust nuclear translocation of p50 (Fig. 4a and Supplementary Fig. 8). Similarly, as compared with untreated mDCs and Si-non-mDCs, Si-p38\alpha-mDCs also demonstrated a substantially upregulated nuclear translocation of p50 (Supplementary Fig. 9a). Next, by chromatin immunoprecipitation (ChIP) assay, we detected an increased physiological binding of RelA, RelB and especially p50 to the *Ox40l* promoter in mSBDCs (Fig. 4b). To examine whether these NF- κ B molecules could activate the *Ox40l* promoter, we performed luciferase reporter gene assays in HEK 293T cells. RelA, RelB or p52 alone did not activate the *Ox40l* promoter, whereas p50 together with RelB, and to a lesser extent, with RelA or c-Rel, did (Fig. 4c). Considering that these NF- κ B molecules were increased in the nuclear fraction of mSBDCs, these data indicate that nuclear translocation of p50, which formed transcriptionally active complexes with RelA, c-Rel and especially with RelB, was responsible for the upregulation of OX40L expression in mSBDCs.

To further verify the p50-dependent upregulation of surface expression of OX40L on mSBDCs, we generated BMDCs from BM cells of $p50^{-/-}$ B6-mice. The expression of OX40L on $p50^{-/-}$ iDCs and $p50^{-/-}$ mDCs was significantly reduced as compared with those generated from BM cells of WT-B6 mice (Fig. 4d,e). Furthermore, SB202190-treated $p50^{-/-}$ iDCs or $p50^{-/-}$ mDCs failed to upregulate the expression of surface OX40L. The surface OX40L expression level was similar between $p50^{-/-}$ mDCs and $p50^{-/-}$ mSBDCs, confirming a crucial role of p50 in modulating OX40L expression of mSBDCs.

Inhibition of PPARγ **contributes to p50 activation in mSBDCs**. Next, we examined the signalling pathways that may lead to activation of p50 following p38 inhibition. As $I\kappa B-\alpha$ did not



Figure 4 | Inhibition of p38 MAPK activates p50-dependent pathways that result in increased surface expression of OX40L on mSBDCs. (a) Western blot analysis of the cytoplasmic and nuclear localization of NF- κ B molecules in iDCs, mDCs, iSBDCs and mSBDCs. (b) ChIP assay of p50, ReIA and ReIB binding to the *Ox40l* promoter in mDCs and mSBDCs. Error bars represent s.d. (n=3). (c) Luciferase reporter assay showed NF- κ B-dependent activation of the *Ox40l* promoter in HEK 293T cells. HEK 293T cells in triplicated wells were transiently transfected with mouse *Ox40l* promoter-inserted vector mOx40I-PG04 or empty plasmid NEG-PG04, followed by transfection with vectors for NF- κ B molecules. Error bars represent s.d. (n=3). (d) Surface expression of OX40L on iDCs, iSBDCs, mDCs and mSBDCs generated from WT and $p50^{-/-}$ C57BL/6 mice analysed by FACS. Numbers showed the percentage of OX40L-positive DCs gated on CD11c⁺ cells. (e) Summarized data (n=4) from d. Error bars represent s.d. Representative data from one of two repeated experiments are shown. *P* values were calculated with Student's t-test.

decrease in mSBDCs (Supplementary Fig. 9b), we performed gene array analysis to investigate the potential molecules that may have contributed to the nuclear translocation of p50. We hypothesized that the decreased PPAR γ expression we observed in gene array analysis may have a role in activating p50 (ref. 22). First, we verified that the protein level of PPARy was sharply decreased in mSBDCs as compared with mDCs (Fig. 5a and Supplementary Fig. 10) and in Si-p38\alpha-mDCs as compared with Si-non-mDCs (Supplementary Fig. 9c). Also, the level of CCAAT/enhancerbinding protein (pC/EBPβ), a known downstream target of p38 MAPK²⁸, was reduced in mSBDCs and in Si-p38\alpha-mDCs (Fig. 5a and Supplementary Figs 9c and 10), which could potentially be responsible for decreased levels of PPARy, considering it is the master transcriptional factor of PPAR γ^{29} . As a confirmation for the observations, knockdown of C/EBPB by siRNA diminished PPARy level in mDCs (Fig. 5b and Supplementary Fig. 10).

Second, we measured PPAR γ activity by detection of mRNA levels of *Ap2*, which has been described as a direct PPAR γ target³⁰. As expected, inhibition of p38 MAPK reduced *Ap2* levels in mDCs as well as the PPAR γ inhibitor, whereas both

rosiglitazone (RGZ), a PPAR γ activator, and the p38 activator, anisomycin³¹, enhanced *Ap2* expression (Fig. 5c). In addition, inhibition of p38 MAPK failed to suppress PPAR γ activity directly induced by RGZ (Fig. 5c). By analysis of the nuclear translocation of p50, we found that p50 activity was retroactively correlated with PPAR γ activity in mDCs (Fig. 5c,d and Supplementary Fig. 10). Moreover, we detected the presence of p50 in PPAR γ immunoprecipitate from mDCs but not from mSBDCs (Fig. 5e and Supplementary Fig. 10), suggesting a possible mechanism of transrepression of p50 activity by PPAR γ by complex formation²², which antagonizes p50 signalling in DCs.

Finally, we also found that PPAR γ inhibitor-induced p50 activation led to upregulated expression of surface OX40L on mDCs, while p50 deficiency abrogated this effect (Fig. 5f,g). As inhibition of p38 MAPK could not reverse RGZ-mediated PPAR γ activation, RGZ-treated mSBDCs failed to express the high levels of OX40L observed on mSBDCs (Fig. 5f,g). These results suggest that inhibition of p38 MAPK suppresses PPAR γ expression and activity in mSBDCs by decreasing C/EBP β phosphorylation,



Figure 5 | Downregulation of PPAR γ **expression/activity in mSBDCs contributes to the p50-mediated increased expression of surface OX40L.** (a) Western blot analysis of the PPAR γ , pC/EBP β and C/EBP β in mDCs and mSBDCs treated with or without p38 inhibitor SB202190, PPAR γ inhibitor GW9662 and/or PPAR γ activator RGZ. (b) The protein level of PPAR γ , pC/EBP β and C/EBP β in mDCs treated with C/EBP β -specific siRNA. (c) RT-PCR analysis of *Ap2* mRNA expression in mDCs treated with or without p38 inhibitor SB202190, PPAR γ activator RGZ, and/or p38 activator anisomycin. Data shown (n = 3) were normalized to the *Gapdh* gene. Error bars represent s.d. (d) Western blot analysis of the cytoplasmic and nuclear localization of p50 molecule in treated mDCs and mSBDCs. (e) Coimmunoprecipitation of endogenous p50 and PPAR γ . mDCs and mSBDCs lysates were immunoprecipitated with PPAR γ antibody or isotype control IgG and immunoblotted with p50 antibody. (f) Surface expression of OX40L on treated mDCs and mSBDCs generated from WT and $p50^{-/-}$ mice analysed by FACS. Numbers showed the percentage of OX40L-positive DCs gated on CD11c⁺ cells. (g) Summarized data (n = 4) from **f**. Error bars represent s.d. Representative data from one of two performed experiments are shown. *P* values were calculated with Student's t-test. thereby leading to reduced functional inhibition of PPAR γ on p50 transcription factor activity and increased expression of OX40L on mSBDCs.

mSBDCs overcome Treg-mediated immunosuppression in vivo. The observed regulatory role of p38 MAPK in DC differentiation and immunogenicity promoted us to examine whether p38 inhibited-mDC can induce potent antigen-specific antitumour immunity and facilitate tumour rejection. Similar to other reports of the B16-OVA³² and B16 (ref. 33) tumour model, mice were immunized with OT-I peptide-pulsed mSBDCs (mSBDCs-OT-I), mDCs-OT-I, mSBDCs-OT-II, mDCs-OT-II, mSBDC-TRP2 or mDC-TRP2 at days 3 and 10 after tumour cell inoculation. Groups of mice were depleted of CD25⁺ Treg with an intraperitoneal injection of CD25-specific antibody or treated with combined Treg cell depletion (3 days before immunization) and DC immunization. We (Supplementary Fig. 11a) and others³⁴ have found that a single injection of CD25-specific antibody can deplete >90% of Treg for a week by examining CD25+Foxp3+ T cells in the spleens. Tumour growth was significantly inhibited by immunization with mSBDCs-OT-I, mSBDCs-OT-II or mSBDC-TRP2, whereas immunization with non-pulsed mSBDCs or Treg depletion had marginal effects, and mDCs-OT-I, mDCs-OT-II or mDC-TRP2 immunization had moderate antitumour effects (Fig. 6a-c and Supplementary Fig. 11b). Notably, mSBDCs immunization alone was at least as effective as the combined treatments of Treg depletion and mDC immunization in inhibiting the growth of pre-established tumour (Fig. 6a-c). In addition, the enhanced antitumour effects of antigen-pulsed mSBDCs were correlated with the significantly higher frequencies of antigen-specific $CD4^+$ or $CD8^+$ IFN- γ -secreting T cells in spleens and the significantly increased CTL activity in immunized mice (Fig. 6d-f and Supplementary Fig. 11c). Importantly, Si-p38a-mDCs-OT-I, mSBDCs-OT-I and mSB80DCs-OT-I, but not the Si-non-mDCs-OT-I, mDCs-OT-I and mSB74DCs-OT-I, could prevent the outgrowth of pre-established B16-OVA tumour (Supplementary Fig. 11d-g), further demonstrating the crucial role of inhibiting p38 MAPK signalling in DCs to induce superior antitumour response after immunization.

We also examined the contribution of OX40L signal delivered by mSBDCs in activating T-cell immunity. We found that OX40L antibody treatment could abrogate the proliferation of activated OT-II T cells induced by mSBDC-OT-II in tumour-draining lymph nodes (Supplementary Fig. 12), and substantially reduced the ability of mSBDCs-OT-II and mSBDCs-OT-I to induce tumour regression (Supplementary Fig. 13). Furthermore, $p50^{-/-}$ mSBDCs-OT-I expressed low levels of OX40L and failed to induce an effective antitumour response compared with mSBDCs-OT-I from WT mice (Supplementary Fig. 11f). These observations reveal the importance of OX40 costimulation initiated by mSBDCs in inducing antitumour immunity *in vivo*.

mSBDCs activate antitumour immunity in the effector phase. We then examined the *in vivo* effects of mSBDCs in the effector phase of antitumour immune response. Tumour-infiltrating lymphocytes (TILs) were isolated from B16-OVA tumour-bearing C57BL/6 mice receiving different treatments. The numbers of tumour-infiltrating, IFN- γ -secreting CD8⁺ or CD4⁺ T cells were significantly increased in mSBDCs-OT-I- or mSBDCs-OT-II-immunized mice, respectively, than the other groups (Fig. 7a). We observed a significantly upregulated expression of OX40 and OX40L on both CD8⁺ and CD4⁺ T feff that might amplify the OX40/OX40L signal on the effector phase of antitumour immune response (Fig. 7b). Tumour-infiltrating CD8⁺ or CD4⁺ CD25⁻

(Fig. 7c) Teff isolated from mice immunized with mSBDCs-OT-I or mSBDCs-OT-II, respectively, had significantly enhanced proliferation when restimulated in vitro with OT-I or OT-II peptideloaded mDCs than those from other groups, even in the presence of CD4⁺CD25⁺ Treg from naive mice. We also analysed tumour-infiltrating CD4+CD25+Foxp3+ Treg population, and to our surprise, mSBDCs-OT-I or mSBDCs-OT-II immunization resulted in significantly decreased percentages of Treg in gated CD4⁺ TILs (Fig. 7d). Moreover, CD4⁺CD25^{hi} Treg isolated from TILs in mSBDCs-OT-II-immunized mice had significantly decreased suppressive function as compared with those from mDC-immunized or phosphate buffered saline (PBS)-treated mice in inhibiting the proliferation of CD4⁺CD25⁻ OT-II cells stimulated with mDCs-OVA (Fig. 7e). In addition, as compared with mSBDCs-OT-I, mSB74DCs-OT-I and p50^{-/-} mSBDCs-OT-I displayed significantly decreased capacity to activate tumour-infiltrating Teff, reduce Treg cell population and compromise the suppressive function of tumour-infiltrating Treg (Supplementary Fig. 14). Taken together, these data indicate that the inhibited p38 MAPK signalling, as well as the subsequently activated p50 NF-KB pathway, are indispensable for mSBDCstimulated activation of tumour-infiltrating Teff during the effector phase.

The role of p38 MAPK in human monocyte-derived DCs. We further asked whether p38 MAPK inhibition could enhance the Teff-stimulatory capacity of human monocyte-derived DCs (h-MoDCs) in the presence of Treg. Inhibition of p38 MAPK during the differentiation of h-MoDCs by 1 µM SB202190 or SB203580, but not SB202474 or DMSO, significantly upregulated the surface expression of OX40L on these cells after maturation (Fig. 8a). In the absence of Treg, the OX40/OX40L signalling appears to only have a minimal effect on the proliferation of alloreactive CD4⁺ Teff induced by p38 MAPK inhibitor-treated h-mMoDCs (Fig. 8b). Importantly, similar to murine p38 MAPK-inhibited BMDCs, p38 MAPK inhibitor-treated h-mMoDCs induced a profound proliferation of alloreactive CD4⁺ Teff in the presence of autologous Treg, whereas the proliferation of CD4⁺ Teff stimulated by SB202474- or DMSO-treated h-mMoDCs was largely inhibited (Fig. 8c). OX40 costimulation is crucial to p38 MAPK inhibitor-treated h-mMoDCs-mediated activation of Teff in the presence of Treg, because OX40L blockage abrogated this effect (Fig. 8c). Overall, these results highlight the translational potential of inhibition of p38 MAPK during the differentiation of h-MoDCs for improving DC-based immunotherapy in human cancers.

Discussion

p38 MAPK activity is required for LPS-, TNF-α- and CD40Linduced maturation of iDCs^{35,36}. However, the present study shows that p38 MAPK activity in DC progenitor cells acts as an antigen presentation attenuator, and disabling this critical brake during DC differentiation endows DCs with superior activity in antigen presentation and induction of immune responses. Our results further demonstrate the crucial role of inhibition of p38 MAPK activity in DCs in activating CD8⁺ and CD4⁺ T cells during both priming and effector phases of the immune response in an antigen-specific manner, possibly owing to the improved maturation status³⁷ of mSBDCs as shown by the significantly enhanced expression of MHC class II, CD80, CD86, CD40 and OX40L, as well as IL-12 production. When p38 MAPK signalling in DC progenitor cells was inhibited, the remarkably improved maturation status of mSBDCs was not limited to maturation induced by TNF- α /IL-1 β , but could also extend to other stimuliinduced maturation, such as LPS, Poly I:C, CpG, sCD40L and



Figure 6 | Antitumour activity of DC immunization and Treg depletion using anti-CD25 antibody. (**a**-**c**) Comparison of antitumour activity against pre-established tumours. C57BL/6 mice (five per group) were inoculated subcutaneously with B16-OVA (**a**,**b**) or B16 (**c**) tumour cells (3×10^5), and 3 days later were immunized with 1×10^6 OT-I peptide-loaded (**a**), OT-II peptide-loaded (**b**) or TRP2 peptide-loaded (**c**) mDCs or mSBDCs twice at a weekly interval. Arrows represent the timing of DC therapy. Anti-CD25 antibodies (50 µg per mouse) were administered intraperitoneally into some groups of mice 3 days before immunization. Error bars represent s.d. Representative results from one of three repeated experiments are shown. (**d**) IFN- γ ICS of splenocytes from tumour-bearing mice 10 days after second immunization was performed after restimulation with the immunizing peptides overnight *in vitro*. Numbers represent percentages of IFN- γ secreting lymphocytes gated on CD4⁺ or CD8⁺ T cells from one representative experiment of three performed. (**e**) *In vivo* CTL assay. CFSE^{lo} (left population in gates, non-target) and CFSE^{hi} (right population in gates, target) cells were pulsed with control peptide and OT-I peptide, respectively, and were injected into mice 7 days after OT-I peptide-pulsed DC immunization. Splenocytes were collected 6 h later for detection of CFSE-labelled cells. Numbers represent percentages of specific killing. Results are representative of one of three independent experiments. (**f**) Pooled data (*n* = 3) for *in vivo* CTL assay. Splenocytes were collected 36 h later for detection of CFSE-labelled cells in OT-II and TRP2 peptide-loaded DC immunization. Error bars represent s.d. *P* values were calculated with Student's *t*-test.

TNF- α /IL-1 β /IL-6/prostaglandin E2 (PGE2; Supplementary Fig. 15). Additional inhibition of p38 MAPK during maturation had no obvious effect in improving SBDC maturation status when matured by TNF- α /IL-1 β , sCD40L or TNF- α /IL-1 β /IL-6/ PGE2, but to some extent, suppressed the maturation of SBDCs stimulated by LPS, Poly I:C or CpG (Supplementary Fig. 15), indicating the important role of p38 MAPK in toll-like receptor (TLR) agonist-induced maturation of iSBDCs. Hence, this study provides new insights into an intracellular negative feedback mechanism, and indicates that DC generation and activation is self-limiting through p38 MAPK activation in DC progenitor cells.

We also found that the self-limiting regulation of DC immunogenicity, controlled by p38 MAPK, is mediated through transcriptional repression of NF- κ B family members (Fig. 9). We showed that p38 MAPK phosphorylates C/EBP β and pC/EBP β induces the expression of PPAR γ , whose activity affects the maturation and function of DCs^{38,39}. PPAR γ has been reported to physically associate with many distinct families of transcription factors, prevent the transcription factors from binding to their response elements and thus suppress their ability to induce gene transcription²². In line with this finding, our results revealed a physical interaction between PPAR γ and NF- κ B p50, which may repress the transactivation of p50. Inhibition of p38 MAPK in



Figure 7 | **Effects of mSBDCs on TILs. (a)** IFN- γ ICS of pooled CD8⁺ T cells or CD4⁺ T cells isolated from TILs of mice 10 days after final immunization was performed after overnight *in vitro* restimulation with OT-I peptide- or OT-II peptide-loaded mDCs. Numbers represent percentages of IFN- γ -secreting lymphocytes gated on CD4⁺ or CD8⁺ T cells. (b) Expression (mean fluorescence intensity, MFI) of OX40 and OX40L on CD4⁺ or CD8⁺ T cells isolated from TILs of mice immunized with OT-II peptide- or OT-I peptide-loaded mDCs and mSBDCs, respectively. (c) Enhanced proliferation of TILs in the absence or presence of Treg. CFSE-labelled 1 × 10⁵ CD8⁺ Teff (left) or CD4⁺ CD25⁻ Teff (right) from TILs were cocultured with 1 × 10⁴ OT-I peptide- or OT-II peptide-loaded mDCs, respectively, in the absence or presence of 1 × 10⁵ unlabelled CD4⁺ CD25⁺ Treg from naive mice in triplicates. Error bars represent s.d. (*n* = 3). (d) FACS analysis of the frequency of tumour-infiltrating Treg. Numbers represent the percentages of CD25⁺ Foxp3⁺ Treg in gated CD4⁺ T cells. (e) Reduced suppressive activity of tumour-infiltrating Treg. Splenic CD4⁺ CD25⁻ Teff (1 × 10⁵) from OT-II mice were cocultured with 1 × 10⁴ OVA-loaded mDCs in the absence or presence of CD4⁺ CD25⁺ Treg (1 × 10⁵) from TILs. CFSE intensity of Teff was measured by FACS 60 h later. Numbers indicate percent CFSE^{lo}/proliferated Teff. Representative data from one of two (**c-e**) or three (**a,b**) repeated experiments are shown. *P* values were calculated with Student's *t*-test.

DCs abrogated this self-limiting regulation and resulted in a robust p50 nuclear translocation. This sustained activated p50 forms transcriptionally active complexes with RelA, c-Rel and especially with RelB, which bind the Ox40l promoter and induce the substantially increased expression of surface OX40L on mSBDCs. Our result that p50 deficiency abrogated the enhanced expression of OX40L on mSBDCs further supports this conclusion.

Tregs have a major role in tumour-related immune suppression, and have been considered as one of the most crucial hurdles to overcome for successful antitumour immunotherapy⁴⁰. A recent report has shown an attenuated IL-10-secreting Treg induced by TLR agonists through inhibition of p38 MAPK signalling in DCs during maturation, owing to the decreased IL-10 production in DCs stimulated with TLR agonists⁴¹. In addition, others also showed that constitutive activation of p38 MAPK is responsible for inducing a tolerogenic profile in DCs during melanoma progression in tumour-bearing mice, with DCs secreting significantly more IL-10 and less IL-12 (ref. 42). Indeed, IL-12 might be downregulated by p38 MAPK signalling in many different antigen-presenting cells, especially following TLR agonists stimulation⁴¹⁻⁴⁵. Our findings differ from these reports because we used cytokine cocktails, not TLR agonists, to mDCs; the former are more relevant to the cells that are being used in clinical trials. A surprising effect of mSBDCs in inhibiting conversion of Teff to CD4+CD25+Fop3+ Treg has been

described in this study. The enhanced OX40/OX40L signals provided by mSBDCs may be responsible for the significantly decreased Treg cell population in the tumour and lymph nodes. In line with these result, several groups have reported that the generation of induced Treg is affected by OX40 costimulation^{25,46}.

In addition to inhibition of Treg conversion, mSBDCs were more resistant to cancer- and Treg cell-derived suppressive factors (Supplementary Fig. 16). Notably, mSBDCs partially abrogated Treg-mediated immunosuppression by activating CD4⁺ or CD8⁺ Teff in the presence of Treg and by inhibiting the suppressive function of Treg, probably via the mechanism of enhanced OX40 signal on both Teff and Treg. Our results show that mSBDCs were as effective as combined treatments of mDC immunization and Treg depletion by anti-CD25 antibodies in controlling the growth of pre-established tumours. Importantly, we also evidenced profound effects of mSBDCs on the effector phase of antitumour immunity, as demonstrated by substantially increased antigen-specific, activated tumour-infiltrating CD4⁺ or CD8⁺ Teff that were more resistant to Treg-mediated suppression, and by the reduced suppressive function of tumourinfiltrating Treg that may be a result of the increased OX40 signalling delivered by mSBDCs and/or activated Teff within the tumours. OX40L expression is only observed on cells present at sites of inflammation and not in those invading tumours or on tumour cells themselves, and this insufficient ligation of OX40



Figure 8 | Effects of p38 MAPK inhibition on h-MoDCs. (a) Expression of OX40L on treated h-mMoDCs. Inhibitors were added during the differentiation of h-MoDCs. Numbers are percentages of OX40L-expressing CD11c⁺ DCs as determined by FACS analysis. (**b,c**) Proliferation of CFSE-labelled $CD4^+CD25^-$ Teff (1×10^5) in the absence (**b**) or presence (**c**) of unlabelled $CD4^+CD25^+$ Treg (1×10^5) induced by various h-mMoDCs (1×10^4). Teff and Treg were isolated from different donors of peripheral blood mononuclear cells. Treg and h-mMoDCs were derived from the same donors. Control IgG or anti-OX40L mAbs were added to the coculture. Six days later, CFSE intensity of Teff was measured by FACS. Numbers indicate percent CFSE^{Io}/ proliferated Teff. Representative results from one of two performed experiments are shown. *P* values were calculated with Student's *t*-test.



Figure 9 | The role of p38 MAPK signalling in the regulation of DC immunogenicity. p38 MAPK phosphorylates C/EBPβ, and pC/EBPβ induces the production of PPARγ. PPARγ can be activated by its ligand or agonist that can then repress the transactivation of p50 by physical interaction with p50 to prevent its binding to the response elements such as *Ox401* promoter. p38 MAPK inhibitor, such as SB202190, could abrogate this self-limitation break and result in a robust p50 nuclear translocation. p50, after forming transcriptionally active complexes with RelB, induces the strikingly increased surface OX40L expression, which is crucial to activation of Teff and inhibition of Treg.

could be a limiting factor for immune-mediated destruction of tumour cells⁴⁷. In support of our findings, OX40L expressed by mDCs has been shown to be important for induction of antitumour immunity⁴⁸, and other groups also demonstrated that OX40 signalling triggered by agonist antibodies facilitates tumour rejection by blocking the suppression of Treg⁴⁹. Taken together, in addition to its strong immunostimulatory effects on Teff, mSBDCs can induce efficient antitumour immune responses by counteracting Treg-mediated immune suppression. Hence, mSBDCs should be very useful as a tool for DC-based immuno-therapy to induce strong antitumour immune responses in cancer patients.

Methods

Mice. C57BL/6 mice were purchased from the National Cancer Institute (NCI), and $OX40^{-/-}$ (B6.129S4-Tnfrsf4^{tm1Nik}/J), OT-II (C57BL/6-Tg(TcraTcrb)425Cbn/J), OT-I (C57BL/6-Tg(TcraTcrb)1100Mjb/J) and $p50^{-/-}$ (C57BL/6-129P-Nfkb1^{tm1Bal}/J) mice were purchased from the Jackson Laboratory. Male, 6–8-week old mice were used in the beginning of each animal experiment. The studies were approved by the Institutional Animal Care and Use Committee of Cleveland Clinic Foundation.

Cell lines and reagents. The WT B16 and B16 transfected with OVA (B16-OVA) melanoma cell lines (American Type Culture Collection (ATCC)) were cultured in Iscove's modified Dulbecco's media (Invitrogen) supplemented with 10% heatinactivated fetal bovine serum (FBS; Thermo Scientific), 100 U ml-1 penicillinstreptomycin and 2 mM L-glutamine (both from Invitrogen). p38 MAPK inhibitor SB202190 and SB203580, and their inactive analogue SB202474 were purchased from EMD Biosciences. PGE2, p38 MAPK activator anisomycin, PPARy activator RGZ and PPARy inhibitor GW9662 were purchased from Sigma. MHC class IIrestricted TRP1 (SGHNCGTCRPGWRGAACNQKILTVR), H2-Kb-restricted TRP2 (SVYDFFVWL), H2-K^b-restricted OT-I (SIINFEKL) and MHC class IIrestricted OT-II (ISQAVHAAHAEINEAGR) peptides were purchased from Gen-Script. Murine OX40L-blocking antibody (RM134L), CD40-blocking antibody (16-0402), CD80-blocking antibody (16-10A1), CD86-blocking antibody (GL1), IL-12blocking antibody (C17.8) and anti-CD25 antibody (PC6.1) were purchased from eBioscience. Human OX40L-blocking antibody (159403) was purchased from R&D Systems. Mouse or human Granulocyte-macrophage colony-stimulating factor (GM-CSF), TNF-α, IL-1β, IL-4, IL-6, IL-2, sCD40L and human TNF-β1 were purchased from R&D Systems. LPS, CpG, Poly I:C were purchased from InvivoGen.

BMDC generation. Mouse BMDCs were prepared as described previously⁵⁰. In brief, BM cells were cultured with 2.5 ml of RPMI-1640 supplemented with 10% FBS and mGM-CSF (20 ng ml⁻¹). At day 4, 2.5 ml fresh medium containing mGM-CSF (20 ng ml⁻¹) was added. At day 7, semi-adherent cells were collected as iDCs. In some cultures, 1.5 μ M of SB202190, SB203580 or SB202474 was added to the cultures on day 0, and cultures with addition of DMSO (0.1%; Sigma) served as control. iDCs collected at day 7 were extensively washed and maintained in medium containing mGM-CSF (10 ng ml⁻¹) before adding TNF- α (10 ng ml⁻¹) and IL-1 β (10 ng ml⁻¹) to help mature the cells. After 48 h, semi-adherent cells were collected as mDCs. No additional DMSO, SB202190, SB203580 or SB202474 was added during the culture or during maturation. Flow cytometry analysis showed that both iDCs and mDCs contain >90% of CD11c⁺ cells, while mDCs express significantly upregulated surface expression of DC-related molecules than iDCs (Supplementary Fig. 1).

h-MoDC generation. h-MoDCs were generated as described previously⁵¹. Monocytes were positively selected from peripheral blood mononuclear cells of healthy donors using anti-CD14-conjugated magnetic microbeads (Miltenyi Biotec). Monocytes were cultured in AIM-V medium supplemented with 50 ng ml⁻¹ human GM-CSF and human Interleukin 4 (hIL-4). In some cultures, 1 μ M of SB202190, SB203580 (ref. 52) or SB202474 was added to the cultures on day 0. After 5 days, semi-adherent cells iDCs were extensively washed and then matured for 2 days by adding human tumor necrosis factor- α (hTNF- α) (10 ng ml⁻¹), hIL-1 β (10 ng ml⁻¹), hIL-6 (10 ng ml⁻¹) and 0.5 μ g ml⁻¹ PGE2. The study was approved by the Institutional Review Board Committee of Cleveland Clinic Foundation.

Flow cytometric analysis. Cells were stained with fluorescein isothiocyanate (FITC), phycoerythrin, (PE), allophycocyanin (APC)-conjugated mAbs (1:100 dilution) after pre-blocking FC γ receptors. Rat mAbs specific for mouse CD4 (GK1.5), CD8 (53-6.7), CD25 (PC6.1), CD11c (N418), CD40 (1C10), CD80 (16-10A1), CD86 (GL1), OX40L (RM134L), OX40 (OX86) and matched isotype controls were purchased from eBioscience. Intracellular IFN- γ staining for T cells

was performed using a Cytofix/Cytoperm kit (BD Biosciences), with anti-IFN- γ (XMG1.2) antibodies (eBioscience). Before intracellular IFN- γ staining, splenocytes were restimulated by peptides (5 µg ml⁻¹) or peptide-loaded mDCs at a 10:1 T cells to DCs ratio overnight, and for the final 6 h of culture, GolgiPlug (BD Biosciences–Pharmingen) was added. Foxp3 staining was performed using a Foxp3 staining kit (eBioscience) with anti-Foxp3 (FJK-16s) antibodies (eBioscience). For flow cytometric analysis, stained cells were analysed on a FACSCalibur (Becton Dickinson) and CELLQuest software.

In vivo CTL assay. In vivo CTL assay was performed as described previously⁵³ with some modifications. Spleen cells from C57BL/6 mice were pulsed with $5 \mu g m l^{-1}$ OT-1 peptide, OT-II peptide, TRP1 peptide or TRP2 peptide. Pulsed splenocytes were labelled with CFSE at a final concentration of $1 \mu M$ (CFSE^h) or at a final concentration of $0.1 \mu M$ (CFSE^h). Two cell populations were mixed at 1:1 ratio, and 2×10^7 total cells were injected intravenously into treated mice. The percentage of antigen-specific lysis was determined 6 h later after target cell injection in OT-I peptide-loaded DC-immunized mice (OT-I peptide-pulsed CFSE^h target + TRP2 peptide-pulsed CFSE^{lo} non-target), and 36 h later in TRP2 peptide-pulsed DC-immunized mice (OT-II peptide-pulsed DC-immunized mice (OT-II peptide-pulsed DC-immunized mice (OT-II peptide-pulsed CFSE^{lo} non-target), by analysing the relative proportion of CFSE^{hi} target cells and CFSE^{lo} non-target cells in splenocytes based on CFSE-labeling intensity. The percentage of specific lysis was calculated as follows: $(1 - R \times M2/M1) \times 100$, where M1 = non-target events, M2 = target events, R = M1 in untreated mice/ M2 in untreated mice.

Isolation of TILs. On different days after DC immunization, mice were killed and TILs were isolated from tumour cell suspensions after a Ficoll gradient centrifugation to eliminate dead cells. Subsets of TILs ($CD8^+$, $CD4^+CD25^-$ and $CD4^+CD25^+$ T cells) were sorted with flow cytometer after staining.

Enzyme-linked immunosorbent assay and western blot. Enzyme-linked immunosorbent assay for IFN- γ and IL-12p70 production was performed by ELISA kits (R&D Systems). For western blot, anti-mouse p38 α (cat. no. 9218), p38 γ (cat. no. 2307), p38 (cat. no. 8690), p-p38 (cat. no. 4631), MAPKAPK-2 (cat. no. 3042), pMAPKAPK-2 (cat. no. 3041), RelA (cat. no. 4764), RelB (cat. no. 4954), p52 (cat. no. 4882), GAPDH (cat. no. 2118), PARP (cat. no. 9532), C/EBP- β (cat. no. 3087), pC/EBP- β (cat. no. 3084), PPAR γ (cat. no. 2430), pStat3 (cat. no. 9145), Stat3 (cat. no. 12640), pSmad3 (cat. no. 9520), Smad3 (cat. no. 9523), IKB- α (cat. no. 4812) and β -actin (cat. no. 4970) from Cell Signaling (1:1,000 dilution), and anti-mouse p38 β , (cat. no. sc-718) and c-Rel (cat. no. sc-70) from Santa Cruz Biotechnology (1:250 dilution) were used.

T-cell proliferation and suppression assays. Splenic CD8⁺, CD4⁺CD25⁻ or CD4⁺CD25⁺ T cells from WT, OX40^{-/-}, OT-I- or OT-II-transgenic mice were isolated by using the MACS T-cell isolation kits (Miltenyi Biotec). T-cell proliferation was measured with CFSE dilution assay. CD8 + or CD4 + CD25 - T cells were incubated for $5\,\text{min}$ at $37\,^\circ\text{C}$ with $1\,\mu\text{M}$ CFSE in PBS and then washed extensively. For antigenic stimulation using TCR-transgenic OT-I or OT-II T cells and antigen-pulsed DCs, 1×10^5 purified T cells and 1×10^4 DCs pulsed with OT-I or OT-II peptides (1 µg ml⁻¹) were seeded in each well of a round-bottom 96-well microtiter plate in triplicate in 200 ml RPMI-1640 medium supplemented with 10% FBS. For stimulation using WT or OX40^{-/-} T cells, 1×10^5 purified T cells and 1×10^4 unpulsed DCs were cocultured in the presence of anti-CD3 antibodies $(1 \, \mu g \, ml^{-}$ ¹). For suppression assays, $1-2 \times 10^5$ unlabelled CD4⁺CD25⁺ Tregs were added in the cocultures. Isotype-matched control immunoglobulin G (IgG) and blocking mAbs were added at a concentration of $10 \,\mu g \,ml^{-1}$. We measured proliferation by relative CFSE dilution after 60 h of culture. For human T-cell proliferation, h-MoDCs were mixed with CFSE-labelled alloreactive CD4 + CD25 Teff in Aim-V medium in the presence or absence of autologous CD4 + CD25 + Treg. Human T-cell proliferation was assessed 6 days after coculture.

Chromatin immunoprecipitation. ChIP assay were performed with the ChIP assay kit (Millipore) according to the manufacturer's instructions. Chromatin was extracted from CD4 ⁺ Foxp3 ⁻ T cells cocultured for 4 days with mDCs or mSBDCs in the presence of TGF- β 1, mDCs and mSBDCs after fixation with formaldehyde. Anti-pSmad3, anti-Pstat3, anti-RelA, anti-RelB, anti-ps0 and iso-type-matched control antibodies (all from Cell Signaling, 1:20 dilution) were used for the immunoprecipitation of chromatin. The precipitated DNA was analysed by reverse transcription (RT)–PCR with the following primer sets: surrounding region of NF- κ B binding site at *0x401* promoter, 5'-CTTCAGAGCACACGTTTGTG AAT-3' and 5'-TGAAGG AGCAGAGCAGAGTCTCTT-3'; enhancer I region: 5'-caagctgacctcaaatcacaaag-3'; 5'-catacccacattttgacctcgc-3'; 5'-gcttctgtgtatgtttt gtgt-3'; 5'-atcacccacacattgagaga-3'; enhancer II region: 5'-GGAACGTCACCTACC ACATCC-3'; 5'-CTACCCACAGGTTTCGTT-3'. Values were subtracted with the amount from IgG control and were normalized to corresponding input control.

Luciferase reporter assays. HEK 293T cells were transiently transfected with a 1121-bp mouse *Ox401* promoter (Genecopoeia)-inserted luciferase reporter vector pEZX-PG04 (mOx401-PG40), or control vector (NEG-PG40) and expression vectors for NF-кB molecules (p50, p52, RelA, RelB and c-Rel, Addgene) by Lipofectamine 2000 (Invitrogen). Promoter activity was measured with Secrete-Pair Dual Luminescence Assay Kit (GeneCopoeia) according to the manufacturer's instructions. Values are expressed as the mean ± s.d. of relative luciferase units normalized to internal control.

DC immunization and tumour models. iDCs, iSBDCs, $p50^{-/-}$ iSBDCs, iSB74DCs, iSB80DCs and siRNA-treated iDCs were pulsed with OT-I (5 µg ml⁻¹), OT-II (15 µg ml⁻¹) or TRP2 peptides (30 µg ml⁻¹) for 8 h, and matured for additional 48 h in the presence of TNF- α and IL-1 β . After washing with PBS, mDCs or treated mDCs were prepared for immunization of mice (five per group). In the therapeutic model, 3×10^5 B16-OVA or B16 tumour cells were then randomly divided into groups and injected with 1×10^6 mDCs or treated mDCs subcutaneously in the left back 3 and 10 days later. Tumour volumes were measured two or three times per week with an electronic caliper.

Statistical analyses. For statistical analysis, Student's *t*-test was used. A *P* value < 0.05 was considered statistically significant. In some Figures, the actual *P* values were shown as superscript numbers. Results are typically presented as means \pm s.d.

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Author contributions

Y.L. and Q.Y. initiated the study, designed the experiments and wrote the paper; Y.L. performed the majority of experiments and statistical analyses; M.Z., B.H., S.W. and. H.L. helped with ChIP and luciferase assay; Y.Z., J.Q., J.H and J.Y. provided critical suggestions to this study; and Q.Y. and J.H. contributed to the conceptual idea and to the writing and editing of the paper.

Additional information

Accession codes: The microarray data were deposited in the NCBI Gene Expression Omnibus (GEO) database under accession numbers GSE57879.

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