<u>**Research Article**</u>

Title: CTLA-4 antibody-drug conjugate reveals autologous destruction of B-lymphocytes
associated with regulatory T cell impairment

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Abstract: Germline CTLA-4 deficiency causes severe autoimmune diseases characterized by dysregulation of Foxp3$^+$ Tregs, hyper-activation of effector memory T cells, and variable forms autoimmune cytopenia including gradual loss of B cells. Cancer patients with severe immune-related adverse events (irAE) after receiving anti-CTLA-4/PD-1 combination immunotherapy also have markedly reduced peripheral B cells. The immunological basis for B cell loss remains unexplained. Here we probe the decline of B cells in human CTLA-4 knock-in mice by using anti-human CTLA-4 antibody Ipilimumab conjugated to a drug payload emtansine (Anti-CTLA-4 ADC). The anti-CTLA-4 ADC-treated mice have T cell hyper-proliferation and their differentiation into effector cells which results in B cell depletion. B cell depletion is mediated by both CD4 and CD8 T cells and at least partially rescued by anti-TNF-alpha antibody. These data revealed an unexpected antagonism between T and B cells and the importance of regulatory T cells in preserving B cells.

INTRODUCTION

Foxp3 is a master regulator of regulatory T cells (Tregs) and its mutations result in fatal autoimmune disease in mice and human.[1-5] Among many known functions, Foxp3 is a transcriptional factor for expression of CTLA-4, which is constitutively expressed in Tregs.[6,7] *Ctla4* deletion in mice phenocopies that of Foxp3. In humans, CTLA-4 deficiency caused by autosomal heterozygous mutation of the CTLA-4 gene[8,9] or mutations in recycling partner LPS-Responsive beige-like anchor (LRBA) protein[10] are associated with severe autoimmune diseases. Although CTLA-4 can be expressed at lower levels in other cell types and has been suggested as a negative regulator for naïve T cell activation, lineage-specific deletion of the *Ctla4* gene in Foxp3^{+} Tregs results in development of systemic lymphoproliferation, fatal autoimmune disease, and potent tumor immunity.[11] These data are consistent with the notion that the predominant function of CTLA-4 is Treg-intrinsic.

A largely overlooked area is the cross-regulation between B cells and regulatory T cells. CTLA-4 conditional null mice and those with depletion of Tregs showed increase in germinal center B cells and heightened antibody responses with some reports showing a decrease B cell percentage.[12-15] B cell loss is a common feature in genetic mutations of Foxp3, scurfy mice, Foxp3 knockout mice or Treg depletion.[16-21] Among cancer patients who received immunotherapy, early change of circulating B cells of patients who received combination immunotherapy anti-CTLA-4/PD-1 correlated to irAE.[22] The change in B cells included decline of circulating B cells, and increase in CD21 low B cells and plasmablasts.[22]

The correlation between B cell loss and immune activation associated with defective Foxp3-CTLA-4 function remains unexplained. Since both CTLA-4 and Foxp3 are largely expressed outside of B cell compartment, and since this pathway is the master regulatory of Treg

function, we hypothesized that B cell loss maybe associated with activation of T cells that are

autodestructive of B cells. This hypothesis is noteworthy as no autodestructive T cells for B cells

have been described. To test this hypothesis, we generated anti-CTLA-4 ADC and showed that

the ADC caused selective depletion of Tregs in mice and a marked reduction of B cells.

Remarkably, activation of CD4 and CD8 T cells is the underlying cause of B cell depletion. Our

data explains the B cell loss associated with Foxp3 and CTLA-4 dysfunction and suggested an

unexpected antagonism between T cells and B cells.

RESULTS

CTLA-4 antibody-drug conjugate depleted regulatory T cells and B cells

Anti-CTLA-4 antibody Ipilimumab (Ipi) or human IgGFc (hIgGFc) control were

conjugated to a well-known drug payload DM1, emtansine, to furnish the corresponding ADCs:

Ipilimumab-DM1 (Ipi-DM1) and hIgGFc-DM1 with corresponding size shifts in the SDS-gel

(Figure 1A). The drug to antibody ratio (DAR) was calculated based on experimentally

determined extinction coefficient A280 & A252 for each antibody and reported values for DM1

(Table S1). The DAR from various conjugation is ~3.2 for Ipi-DM1 and ~ 1.7 for hIgGFc-DM1

control (Table S1), as expected based on the sizes of the proteins. The binding for Ipi-DM1 ADC

was evaluated by ELISA binding to immobilized His-hCTLA-4 and by flow cytometry using

hCTLA-4 expressing CHO cells (CHO-hCTLA-4). Ipi-DM1 binding was found comparable to

the parent antibody Ipi (Figures 1B, 1C). Specific killing of CTLA-4-expressing cells by Ipi-

DM1 was assessed in human CTLA-4 expressing CHO (CHO-hCTLA-4) and wild type CHO

(CHO-WT) cells in vitro (Figure 1-figure supplement 1), which showed Ipi-DM1 reduced

viability of CLTA-4 expressing CHO but not wild type cells. As expected, there was no change

in viability for either CHO cell lines when treated with Ipi (Figure 1-figure supplement 1).

In order to evaluate the in vivo effects of CTLA-4 ADC on Tregs, we treated human

CTLA-4 knock-in (*Ctla4$^{h/h}$*) mice with control hIgGFc or Ipi-DM1 ADC (Figure 1D). Peripheral

blood was stained for flow cytometry and gated on CD45 and T-cell subset, including CD8,

CD4, CD4 Foxp3^{+} and CD4 Foxp3^{-} subsets (Figure 1-figure supplement 2A). We found that Ipi-

DM1 ADC significantly depleted Foxp3^{+} Tregs as indicated by the reduction in percentage and

cell number of Foxp3^{+} CD4 T cells when compared to hIgGFc control (Figure 1E). In addition,

total CTLA-4 levels and Ki67 staining in Tregs were decreased compared to control group

(Figures 1F, 1G). In contrast, Ipi-DM1 ADC did not alter either % of CD4 Foxp3^{-} cells among

CD45^{+} leukocytes or CTLA-4 levels on the subset, but did slightly decrease the cell number of

CD4 Foxp3^{-} cells when compared hIgGFc control (Figure 1-figure supplement 3A, 3B).

Unexpectedly, Ipi-DM1 ADC caused a dramatic decline of B cells as percentages of CD45 and

cell number in the periphery (Figure 1H). B cells from mice treated with Ipi-DM1 exhibited

higher proliferation than those from hIgGFc treated control mice (Figure 1I).

Since a dramatic loss of B cells and a slight decrease in CD4-nonTregs in Ipi-DM1 treated

mice were observed, we sought ensure that this phenomenon was not a result of downstream

release of payload drug DM1. As shown in Figure 1-figure supplement 4A, hIgGFc-DM1 ADC

(payload control) did not alter Treg percentage and cell number. A slightly higher level of

CTLA-4 was noted in CD4 subset, while Treg proliferation did not change significantly

compared to control (Figure 1-figure supplement 4B, 4C). No other effects on T cell subsets

were noted (Figure 1-figure supplement 4D, 4E). Importantly, B cell number and proliferation

were unaffected by hIgGFc-DM1 ADC treatment (Figure 1-figure supplement 4F, 4G).

Additionally, to ensure B cell depletion was not directly caused by Ipi-DM1 ADC, we stained

B220$^+$ B cells and Foxp3$^+$ Tregs for human CTLA-4 in both human CTLA-4 knock-in (Ctla4$^{h/h}$)

and WT mice to evaluate expression of CTLA-4 in B cells. As expected, CTLA-4 was detected

in Foxp3$^+$ Tregs of Ctla4$^{h/h}$ but not B cells (Figure 1-figure supplement 5).

We then investigated the impact of anti-CTLA-4 ADC in lymphoid organs. Treg

percentage and absolute numbers were not impacted by Ipi-DM1 treatment compared to control

in the spleen at day 9. The percent of B cells decreased significantly, although the reduction in

absolute number of B cells was not statistically significant (Figure 1-figure supplement 6A, 6B).

Flow analysis of lymph nodes after Ipi-DM1 treatment resulted in a significant increase in

absolute cell numbers of Tregs and B cells and B cell percentage. (Figure 1-figure supplement

6C, 6D).

We then asked how Ipi-DM1 ADC induced Treg impairment impacted B cell

lymphopoiesis. Bone marrow cells were stained for flow cytometry and gated on CD45 and T-

cell subsets were defined as CD8 (CD8$^+$B220$^-$), CD4 (CD4$^+$B220$^-$), Tregs (CD4$^+$B220$^-$ Foxp3$^+$)

and CD4-nonTregs (CD4$^+$B220$^-$ Foxp3$^-$) (Figure 1-figure supplement 2B). Ipi-DM1 depletion of

Tregs was confirmed in the bone marrow (Figure 1J). Additionally, total CTLA-4 level in bone

marrow Tregs decreased, however Ki67 staining did not change significantly compared to

control group (Figures 1K, 1L). In contrast, Ipi-DM1 ADC did not alter bone marrow CD4-

nonTreg cell percentage among CD45$^+$, absolute cell number, or CTLA-4 level compared

hIgGFc control (Figure 1-figure supplement 3C, 3D). FACS analysis of bone marrow B cells

revealed the loss of mature B cells but not immature transitional type1 or Pre-pro/Pro/Pre B cells

from Ipi-DM1 treated mice (Figure 1M-1Q, Figure 1-figure supplement 7). Corresponding to

mature B cell loss, the Pre-Pro/Pro/Pre B cell subtype percentage in B220 are enriched (Figure

1P, 1Q, Figure 1-figure supplement 7). Additionally, CD21/35 expression level in the remaining

mature B cells in Ipi-DM1 treated mice was lower than those from hIgGFc treated control mice

(Figure 1R).

We then investigated B cell apoptosis in peripheral blood and lymphoid organs (Figure 2,

Figure 2-figure supplement 1). Only mature bone marrow B220^{hi} B cells showed increase in

apoptosis from Ipi-DM1 treatment, while progenitor/immature bone marrow B220^{lo} B cells were

similar to control group. Blood and spleen B cell apoptosis were similar to hIgGFC control

group, while lymph node B cell apoptosis was lower than the control group. Thus, the loss of B

cells in the peripheral blood is likely due to loss of mature B cells in the bone marrow, and that

such loss correlates with Treg depletion.

Taken together, data in Figure 1 & 2 showed that Ipi-DM1 can impair Treg function by

depleting Tregs, preferentially the proliferating Tregs with higher CTLA-4 levels, and loss of B

cells in the blood and bone marrow while B cell progenitors are not impacted. B cell death

increase is observed in only B220^{hi} bone marrow B cells, which are predominantly mature B

cells.

B-cell depletion correlates with Treg impairment by Ipi-DM1 and increases

immunoglobulins

To investigate the kinetic relationship between the decline of B cells and Treg

impairment, human CTLA-4 knock-in (Ctla4$^{h/h}$) mice were treated with control hIgGFc or Ipi-

DM1 and bled according to schedule diagram in Figure 3A. We observed that peripheral blood

samples from mice treated with Ipi-DM1 had a gradual decrease in total leukocytes compared

hIgGFc control group followed by a full recovery by Day 25 (Figure 3B). The large decrease of

CD45 cells in Ipi-DM1 group is predominantly attributed to the decline of B cells, which

plateaus on days 9 and 13 and recovers by day 25 (Figures 3C, 3D) in correlation with the

kinetics of Tregs (Figure 3E, 3F). Similar to the bone marrow data in Figure 1, peripheral blood

mature B cells decreased and transitional type1 B cells were enriched in Ipi-DM1 group

compared hIgGFc control (Figure 3G-3I). Additionally, Treg impairment by Ipi-DM1 increases

immunoglobulins especially IgE (Figure 3-figure supplement 1). The lack of Treg depletion in

the spleen and lymph nodes at these time points remains to be explained, although significant

immune activation, as suggested by splenomegaly and adenopathy (Figure 3-figure supplement

2), may result in production of cytokines that drive Treg proliferation.

Depletion of T-cells but not macrophage rescues B-cells from ablation by Ipi-DM1

We then evaluated how Ipi-DM1-mediated impairment of Treg function affects total CD4

and CD8 T cells in peripheral blood. Mice treated with Ipi-DM1 ADC did not affect the total

percentage of CD4 or CD8 T cells among CD45$^+$ leukocytes, but the cell number of both cell

types decreased (Figure 4A, 4B). Analysis of lymphoid organ T cells showed differing T cell

absolute number phenotype where bone marrow, thymus and spleen T cells were similar to

control hIgGFc while lymph node T cells increased (Figure 4-figure supplement 1). Staining of

Ki67 in CD4 and CD8 revealed a greater percentage of T cells in hyper-proliferative state in

blood and lymphoid organs (Figure 4C, Figure 4-figure supplement 2). Alternatively, mice

treated with hIgGFc-DM1 payload control did not change CD4 or CD8 percentage of CD45, cell

numbers and Ki67 staining compared to hIgGFc control in the periphery (Figure 4-figure

supplement 3).

We then analyzed the functional subsets of T cells in mice that received control hIgGFc

or Ipi-DM1 at day 9. Using CD44 and CD62L markers, we observed an expansion of effector

memory T cells (CD44^{hi}CD62L^{low}) in Ipi-DM1 group for both CD4 and CD8 T cells (Figures

4D, 4E). Correspondingly, the frequency of naive T cells was reduced for Ipi-DM1 while central

memory T cells were unchanged (Figures 4D, 4E). These results show that CTLA-4 ADC can

impair Treg function thereby resulting an increase in effector memory T cells and a hyper-

proliferative state similar to animals that lack the expression of CTLA-4 or Foxp3.

Since B cells are devoid of CTLA-4, Ipi-DM1 must have induced other effector cells that

are directly responsible for B cell depletion. To understand which cell types are responsible for

the destruction of B cells, we depleted either T cells or macrophages using either anti-Thy1.2

mAb or chlondrosome. Ctla4$^{h/h}$ mice were treated with control hIgGFc, Ipi-DM1, Ipi-DM1 in

combination with anti-Thy1.2, or Ipi-DM1 in combination with chlondrosome and bled on day 9

(Figure 5A). As shown in the Figure 5B, top panel, anti-Thy1.2 mAb efficiently depleted total T

cells while macrophage depletion by chlondrosome had no effect (Figure 5B top panel, 5C, 5D).

Additionally, combination of Ipi-DM1 with T cell depleting antibody resulted in a remarkable

rescue of B cells as indicated by percentage B220 in CD45 and cell number while combination

with macrophage depletion did not rescue the B cells (Figures 5B bottom panel, 5E).

In order to understand which subsets of T cells played a critical role in the T cell

mediated destruction of B cells, we treated Ctla4$^{h/h}$ mice with hIgGFc control, Ipi-DM1, Ipi-

DM1 with CD4 depleting antibody, or Ipi-DM1 with CD8 depleting antibody (Figure 5A). Both

CD4 and CD8 depleting antibodies provide efficient depletion (Figure 5F, top panel) of their

respective targeted T cell. Depletion of either CD4 or CD8 T cells results in B cell increase

(Figures 5F bottom panel, 5I). Interestingly, the cell number of CD4 and CD8 T cells increase

reciprocally by depletion of the other subsets (Figures 5G, 5H).

Taken together our CTLA-4 antibody-drug conjugate Ipi-DM1 can impair Treg function,

which results in a T cell mediated destruction of B cells. Whole T cell depletion with anti-

Thy1.2, depletion of CD4 or CD8 T cells rescued B cells from abrogation by Ipi-DM1.

B-cell rescue by Belatacept suggest a role for B7-CD28 interaction in B-cell depletion

The process for T cell activation requires two signals. The first signal is the binding of T-

cell receptor (TCR) to antigen-bound major histocompatibility complex (MHC) and the second

signal is a costimulatory molecule CD28 binding with B7-1/2 (CD80/86) on the antigen-

presenting cell. Having shown that the destruction of B cell can be prevented by total depletion

of T cell, we sought to rescue the B cells by breaking the CD28-B7 signal with a soluble CTLA-

4 that can bind to B7 but not Ipilimumab or Ipi-DM1 ADC. As shown in Figure 6-figure

supplement 1, Abatacept can neutralize Ipilimumab/ADC while Belatacept does not.

Furthermore, we previously showed that human CTLA4-Ig or mutants can bind to murine B7-1/2

(CD80/86) and block their binding to murine CD28 efficiently. [23,24] Therefore, Belatacept can be

used to test if activation of T cells is required for B cell depletion.

We treated mice with Ipi-DM1 to impair Treg function and used Belatacept to break

CD28/B7 signal to block T cell activation. Briefly, *Ctla4$^{h/h}$* mice were treated with hIgGFc or

Ipi-DM1 with/out Belatacept and bled on day 9 for flow analysis. Foxp3^{+} Tregs percentage in

CD4 and cell number decreased similarly for Ipi-DM1 with/out Belatacept compared hIgGFc

control (Figures 6A, 6B). Additionally, Belatacept did not impact Ipi-DM1 to target CTLA-4

expressing Tregs, as Ipi-DM1 with/out Belactacept had similar total CTLA-4 level reduction

compared to control group (Figure 6C). However, Belatacept rescued B cells from Ipi-DM1-

mediated depletion (Figures 6D, 6E). Correspondingly, Belatacept reduced effector memory T

cells (Figures 6F, 6G) as well as granzyme B and IFN-γ expression in CD4 and CD8 T cells

compared to Ipi-DM1 treated group (Figures 6H-6K, Figure 6-figure supplement 2).

Collectively, the data presented here show that Ipi-DM1 can impair Treg function

resulting in T cell activation as indicated by increase in effector memory T cell markers,

GranzymB, and cytokine production thereby resulting in the loss of B cells. However, under Ipi-

DM1 induced Treg impairment the addition of mutant soluble CTLA-4-Ig, Belatacept, can

reduce T cell activation and thereby preserving B cells.

B-cell depletion is partially rescued by anti-TNF-alpha

Since T cells are the effector cells responsible for B cell abrogation, it is of great interest

to evaluate the molecular mechanism by which B cells are eliminated by T cells. To address this

issue, we tested if either FasL or TNF-alpha, which are produced by activated T cells are

responsible. To answer this question, Ctla4$^{h/h}$ mice were treated with hIgGFc control or Ipi-DM1

with/out Adalimumab a human anti-TNF-alpha that also binds to mouse TNF-alpha (Figure 7A,

7B). Flow analysis showed Tregs were decreased similarly between Ipi-DM1 with/out

Adalimumab compared hIgGFc control group (Figures 7C top panel, 7D). Remarkably, the anti-

TNF-alpha partially rescued B cells as shown by percentage difference of B cell marker B220

between Ipi-DM1 and in combination with Adalimumab as well as cell number (Figures 7C

bottom panel, 7E). Peripheral blood samples stimulated with Iononmycin/PMA increased

intracellular cytokines TNF-alpha and IFN-gamma for T cells from Ipi-DM1 treated mice

compared to hIgGFc control, while that with Ipi-DM1 in combination with Adalimumab had

slight increase but the change was insignificant except for IFN-gamma in CD8 T cells (Figure

7F, 7G). In contrast, blocking FAS-L with antibody did not result in B cell rescue (Figure 7-

figure supplement 1).

DISCUSSION

We have reported that pH-insensitive anti-CTLA-4 antibodies trafficked to, and were

degraded in the lysosomes.[25] Since lysosomal degradation is needed to release DM1 from the

antibody for cytotoxicity, the pH-insensitive Ipilimumab was chosen for preparation of antibody-

drug conjugate to be used to impact Treg function. We found that in Ctla4[h/h] mice anti-CTLA-4

ADC, Ipi-DM1, can recapitulate the phenotype of loss of B cells associated clinical deficiency of

CTLA-4[8,9]   /recycling partner LBRA[10]. Our analysis revealed B cell loss in both bone marrow

and peripheral blood, specifically mature B cells. Loss of mature peripheral B cells is consistent

with clinical data for some patients with CTLA-4 haploinsufficiency.[8,9]

B cell loss was previously reported in mice with either naturally occurred (in Scurfy

mice) or targeted mutation of Foxp3 and Treg depletion.[16-21] The kinetics of B cell loss and Treg

depletion in the Ipi-DM1 treated mice revealed that B cell loss was transient and correlated to

Treg impairment (proliferation, CTLA-4 level, cell number). Our data further showed that B cell

destruction under Treg impairment conditions is T cell-mediated and required T cell activation.

Scurfy mice have poor central and peripheral B lymphopoiesis, however neonatal WT

Treg adoptive transfer in scurfy mice resulted in robust population of mature B cells in the

spleen.[19] Genetic ablation of TCRα gene by crossing scurfy with TCRA$\alpha^{-/-}$ mice efficiently

support B cell lymphopoiesis and was sufficient to restore B cells in the bone marrow and

peripheral compartments.[19] Additionally, reconstitution of bone marrow chimera from scurfy or

wild type mixed with that from μMT mice that is deficient in B cells results in normal B cells

after 8 weeks from reconstitution.[18] This is in line with our data that Treg impairment results in

activated T-cell-mediated apoptosis of mature B cells. Mature B cell loss in bone marrow

explains the loss of B cell in the PBL and rules out the possibility that the loss of B cell in the

periphery is merely a consequence of defective circulation of B cells normally produced in bone

marrow. Others have implied that under Treg depletion, activated T cells are targeting

interleukin 7 (IL-7) secreting ICAM1+ perivascular stromal cells needed to progress B cell

progenitors.[21] IL-7 is essential for B lymphopoiesis transition from Pro-B cell into Pre-B cell.[26]

However, our model system is distinct where only mature B cells are depleted, while B cell

progenitor precursors, and specifically Pre-B cells are not impacted. The involvement of T cells

is demonstrated by T cell depletion and requirement for T cell activation is demonstrated by

requirement for B7-CD28 interaction for B cell loss. Furthermore, we found that anti-TNF-α can

partially rescue B-cells from loss imposed by Treg impairment. These data showed that TNF-α

is one of the mediators of B cell loss. Additional studies are needed to reveal other mechanism

for B cells loss in Treg-defective environment.

While in vitro studies by others showed B cell killing by activated $CD25^+CD4$ Tregs, but

not by $CD25^-$ CD4 T cells, [27] this is not the case in vivo where B cell loss is associated with

Foxp3 Treg loss or impairment. [16-21] Furthermore, our current study shows that both CD4 and

CD8 T cells contributed to B cell loss. The concept that B cells are actively eliminated by T cells

in vivo expands our understanding of T-B cell interaction by showing antagonism rather than just

"immunological help" from T cells to B cells.

MATERIALS and METHODS

Experimental animals

C57BL/6 mice that express the CTLA-4 protein with 100% identity to human CTLA-4 protein

under the control of endogenous mouse *Ctla4* locus, *Ctla4[h/h]*, have been previously described.[28]

All animals used in experiments were 7-9 weeks age (female mice were used unless male mice are

indicated). No blinding or randomization was used and mice were fairly distributed into different

treatment groups so that initial average weight of each group were similar. All mice were

maintained at the Research Animal Facility of the Institute of Human Virology at the University

of Maryland Baltimore School of Medicine. All animal studies were approved by the Institutional

Animal Care and Use Committee.

Cell culture and treatment

CHO cells that were stably transfected with human CTLA-4 have been reported.[23] CHO cells were

grown in DMEM (Dulbecco's Modified Eagle Medium, Gibco) supplemented with 10% FBS

(Hyclone), 100 units/mL of penicillin and 100 µg/mL of streptomycin (Gibco). Mice Peripheral

blood leukocytes were cultured in RPMI-1640 medium (containing 10% FBS and 2%

penicillin/streptomycin). All cell lines were incubated at 37 °C and were maintained in an

atmosphere containing 5% CO2.

Viability assay

Wild type CHO (CHO-WT) or human CTLA-4 expressing CHO (CHO-hCTLA-4) cells were

seeded at 1,000 cell/well in a flat 96-well plate at 37 °C for 24 hrs in cell culture incubator. The

medium was then replaced with fresh medium containing 1/4 serially diluted vehicle (PBS),

Ipilimumab or Ipilimumab-DM1 ADC and cell were incubated at 37 °C for additional 72 hrs.

Each treatment group was in duplicate or triplicates. CCK-8 viability dye was then added to each

well according to manufacturer's protcol and incubated for another (2-2.5 hrs) at 37 °C. Wells

were subsequently read for absorbance at 450nM on Spectramax ID3 Molecular Devices plate

reader. In case of testing whether soluble human CTLA-4-Ig or mutant can neutralize

Ipilimumab or Ipilimumab-DM1, the same conditions above were used except that Abetacept or

Belatacept concentrations were kept constant at 6 μg/mL. Data is normalized according to the

following equation (treatment$_{OD450}$ - background$_{averageOD450}$ /vehicle$_{averageOD450}$ - background$_{average}$)

x 100.

Cell surface CTLA-4 binding

Freshly trypsinized human CTLA-4 expressing CHO (CHO-hCTLA-4) cells were stained with

1/4 serially dilute anti-CTLA-4 Iplimiumab or Iplimiumab-DM1 ADC in FACS buffer (2% FBS

with 2 mM EDTA) for 30 minutes on ice. Cells were then washed twice with FACS buffer and

incubated with anti-human IgG AF488 secondary antibody for 20 minutes on ice. Cells were

washed twice with FACS buffer and processed on BD Canto II flow cytometer. In the case of

whether soluble human CTLA-4-Ig or mutant can neutralize Ipilimumab or Ipilimumab-DM1

and prevent them from binding cell to surface CTLA-4 the same conditions above were used

except that Abetacept or Belatacept concentrations were kept constant at 6 μg/mL.

Peripheral blood T cell stimulation

Peripheral blood samples 50 μL were treated with ACK Lysis buffer and washed with RPMI-1640

medium. Leukocytes were then stimulated with 1 µg/ml each of phorbol 12-myristate 13-acetate

(PMA) (Sigma-Aldrich, St. Louis, MO), ionomycin (Sigma Aldrich, St. Louis, MO) and BD

GolgiStop™ (BD Biosciences, cat. 51-2092KZ) and cultured in RPMI-1640 medium (containing

10% FBS and 2% penicillin/streptomycin) for 4 hours at 37°C in 96-well plate. Medium was

removed and cells were washed twice with FACS buffer (2% FBS with 2 mM EDTA) followed

by surface staining and fix/perm and intracellular staining.

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Flow Cytometry

Leukocytes from blood or bone marrow were FACS stained directly or after red blood cell lysis

with ACK buffer. Fc Receptor was blocked with anti-FCR clone 2.4G2 at 10 μg/mL in FACS

buffer for 10 minutes at room temperature and respective surface staining antibodies cocktails

were added to each sample and incubated on ice for additional 20 minutes. Cells were then washed

twice with 1xPBS and stained with 1x live/Dead Fixable dye Aqua in 1x PBS for 7 minutes at

room temperature. Cells were then washed twice with FACS buffer and fixed with eBioscience™

Foxp3/Transcription Factor Staining Buffer Set for 40 minutes. Samples were either washed twice

and resuspended in FACS buffer and processed for flow acquisition or further permeabilized for

intracellular staining using perm buffer from same kit (eBioscience™ Foxp3/Transcription Factor

Staining Buffer Set). In General, all intracellular staining of Foxp3, Ki67, hCTLA-4, GranzymeB,

or cytokines were done overnight at 4 degree. In the case of detecting intracellular cytokines,

samples were cultured and simulated according to the above protocol followed by surface then

intracellular cytokine staining. Apoptosis staining with Annexin V/Pi was performed according

manufacturer protocol after surface staining. Samples were acquired by the BD Canto II Flow

cytometer and data were analyzed by Flowjo software.

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ELISAs

96-well high-binding polystyrene plates were pre-coated with 50 µL of 1 µg/mL of His-hCTLA-

4, or mouse TNF-alpha in coating buffer (0.1M bicarbonate) at 4°C overnight. After washing away

the unbound protein/antibody thrice with 0.05% PBST, the plates were blocked with blocking

buffer (1% BSA in PBST) for 1hr at room temperature. All primary antibody incubation was done

in blocking buffer at 4°C overnight or room temperature for 1 hour. For coated His-hCTLA-4 a

given concentration of either Ipilimumab/DM1, or hIgGFc/DM1; For coated mouse TNF-alpha a

given concentration of Adalimumab or Ipilimumab negative control were used. Following primary

incubation plates were then washed with PBST for four times and incubated with a goat anti-

human IgGFc HRP conjugate secondary antibody at 1/20,000 dilution for detection in blocking

buffer for 1 hr at room temperature. Plates were then washed 4 times with PBST followed by

development with 1-Step™ Ultra TMB-ELISA Substrate Solution for 10 minutes and stopped with

2N sulfuric acid. For Immunoglobulin quantification, an ELISA kit was used following

manufacturer protocol. Wells were read at 450nM on Spectramax ID3 Molecular Devices plate

reader.

Antibody-drug conjugate preparation

Ipilimumab or hIgGFc were buffer exchanged using prepacked column PD-10 into 1X PBS.

Ipilimumab or hIgGFc 5 mg each at concentration 1.5 mg/mL were conjugated with SMCC-

DM1(15 eq) in 1x PBS in the presence of 10% DMSO at 37°C under mild shaking conditions

for 50 minutes to furnish Ipilimumab-DM1 or hIgGFc-DM1 ADC respectively. Reaction

mixture was cleaned up from excess SMCC-DM1 by buffer exchange with a in house ADC

buffer at pH 6.5 containing (20 mM Histidine, 8 % sucrose, and Polysorbate 80 0.02%) by

Econo-Pac 10DG prepacked column according to manufacturer protocol. Confirmed fraction

containing ADC by nano drop were combined and concentrated down to 1 mL using Pierce™

Protein Concentrator PES, (30K for hIgGFc-DM1) or (50K for Ipilimumab-DM1) MWCO

according manufacture protocol followed by sterile filtration. ADC concentration was

determined using BCA assay according to manufacturer protocol. ADC were diluted to 0.4 – 0.5

mg/mL in 1xPBS and A_{280} &A_{252} were recorded on Nanodrop One. The drug to antibody ratio

(DAR) was calculated following previous literature.[29]

Antibodies and fusion proteins used for in vivo studies

CTLA-4-Ig fusion proteins were synthesized by Sydlabs, Inc (Boston, MA). Recombinant

Ipilimumab with amino acid sequence disclosed in WC500109302 was produced by Sydlabs Inc.

(Boston, MA). Azide-free human IgG-Fc was purchased from Athens Research and Technology

(Athens, GA, USA). Antibody-drug conjugates Ipilimumab-DM1 and hIgGFc-DM1 were

prepared from parent antibodies in lab. Depletion antibodies anti-mouse Thy1.2, clone

30H12(BE0066); anti-mouse CD4, clone GK1.5 (BE0003-1); and anti-mouse CD8α, clone 2.43

(BE00-61) were purchased by Bioxcell Inc. (West Lebanon, NH, USA). Anti-mouse FASL,

clone MFL3 (BE00319) was purchased by Bioxcell Inc. (West Lebanon, NH, USA). Anti-TNF-

α, Adalimumab, clinical grade Humira™ was purchased from Premium Health Services

(Columbia, MD, USA).

Other reagents and material

SMCC-DM1 drug payload with cross linker (Cederlanelabs/Cayman Chemical Co, 23926-10)

Anti-mouse FcγR, clone 2.4G2 (Bioxcell Inc, BE0307). Clodrosome® (Encapsula Nano

Sciences, SKU# CLD-8909). Mouse anti-human IgGFc secondary antibody (Invitrogen/Thermo

Fisher Scientific, 05-42-00). Goat anti-human IgGFc (HRP) preadsorbed (Abcam, ab98624).

Polyhistidine-tagged human CTLA-4 (HIS-hCTLA-4) (Sino Biological Inc, 11159-H08H).

Mouse TNF-alpha (Sino Biological Inc, 50349-MNAE). IgE ELISA Kit (Thermo Fisher

Scientific/Invitrogen, 88-50460-86). IgG ELISA Kit (Thermo Fisher Scientific/Invitrogen, 88-

50400-86). IgM ELISA Kit (Thermo Fisher Scientific/Invitrogen, 88-50470-86). IgA ELISA Kit

(Thermo Fisher Scientific/Invitrogen, 88-50450-86). 123 eBeads^{TM} counting beads flow

(Thermo Fisher Scientific/Invitrogen, 01-1234-42). NuPage^{TM} BIS-TRIS gels 4-12% (Thermo

Fisher Scientific/Invitrogen, NP0335BOX). NuPAGE™ MOPS SDS Running Buffer (20X)

(Thermo Fisher Scientific/Invitrogen, NP0001). NuPage^{TM} LDS sample buffer (4x) (Thermo

Fisher Scientific/Invitrogen, NP0007). Protein Ladder (Fisher Scientific, BP3603500). Sucrose

(Sigma Aldrich, S0389-1KG). L-Histidine (Sigma Aldrich, H8000-10G). Dimethyl sulfoxide

(DMSO) (Santa Cruz Biotechnology, Sc-258801). Polysorbate 80 (Fisher Scientific, L13315).

Buffer exchange prepacked column PD-10 (GE health care/ Cytiva Life Sciences, 17085101).

Buffer exchange prepacked column Econo-Pac 10DG (Bio-Rad, 7322010). Cell Counting Kit-8

(CCK-8) (Bimake, B34304). 1-Step™ Ultra TMB-ELISA Substrate Solution (Thermo Fisher

Scientific, 34028). Gibco™ ACK Lysing buffer (Thermo Fisher scientific/Gibco, A1049201).

Live/Dead™ Fixable Aqua Dead Cell Stain (Thermo Fisher Scientific/Life Technologies,

L34966). eBioscience™ Foxp3 / Transcription Factor Staining Buffer Set (Thermo Fisher

Scientific/Invitrogen, 00-5523-00). Pierce™ Protein Concentrator PES, 30K MWCO, 2-6 mL

(Thermo Fisher Scientific/ Pierce, 88521) Pierce™ Protein Concentrators PES, 50K MWCO, 2–

6 mL (Thermo Fisher Scientific/ Pierce, 88538). Pierce™ BCA Protein Assay Kit (Thermo

Fisher Scientific/Pierce, 23227).

Flow antibodies

eBioscience/ Thermo Fisher Scientific (San Diego, CA): APC-eFlour780 anti- mouse CD45,

clone 30-F11 (47-0451-82);eFlour450 anti-mouse CD4, clone GK1.5 (48-0041-82); Pacific Blue

anti-mouse CD4, clone RM-4-4 (116008); PE-Cyanine7 anti-mouse CD8α, clone 53-6.7 (25-

0081-82); PE-Cyanine7 anti-mouse CD8β, clone H35-17.2 (25-0083-82);PerCP-Cy5.5 anti-

mouse B220, clone RA3-6B2 (45-0452-82); APC anti-mouse Foxp3, clone FJK-16s (17-5773-

82);FITC anti-mouse Ki67, clone SolA15 (11-5698-82);FITC Isotype rat IgG2ak, clone eBR2a

(11-4321-82);FITC anti- mouse CD44, clone IM7 (11-0441-85);PerCP-Cy5.5 anti-mouse

CD62L, clone MEL-14 (45-0621-82);PE-Cyanine7 anti-mouse B220, clone RA3-6B2 (25-0452-

82); APC anti-mouse IgM, clone II/41 (17-5790-82); eFlour 450 anti-mouse CD21/CD35, clone

4E3 (48-0212-82); AF488 anti-mouse TNF-α, clone MP6-XT22 (53-7321-82);AF488 Isotype rat

IgG1k, clone eBRG1 (53-4301-80); APC anti-mouse IFNγ, clone XMG1.2 (17-7311-82);APC

Isotype rat IgG1k, clone eBRG1 (17-4301-82);FITC anti- mouse GranzymeB, clone NGZB (11-

8898-82); Alexa Fluor 488-conjugated goat anti-human IgG (H+L) cross-adsorbed secondary

antibody (A-11013). BioLegend (San Diego, CA): PE anti-human CTLA-4, clone BNI3

(369604);PE Isotype mIgG2ak, clone MPC-173 (400212); PerCP-Cy5.5 anti-mouse IgD, clone

11-26c.21 (405710); BV421 anti-mouse CD21/CD35, clone 7E9 (12342); APC Annexin V

Apoptosis Detection Kit with PI (640932);

Statistical analysis

The specific tests used to analyze each set of experiments are indicated in the figure legends.

Data were analyzed using a two-tailed t-test to compare between two groups, one-way or two

way ANOVA (analysis of variance) for multiple comparisons. In the graphs, y-axis error bars

represent S.E.M. or S.D. as indicated. Statistical calculations were performed using GraphPad

Prism software (GraphPad Software, San Diego, California). Non-significant [ns], $*p < 0.05$, $**p$

< 0.01, $***p < 0.001$, $****p < 0.0001$. Reported CTLA-4 level and absolute cell numbers are

normalized by the average value of hIgGFc control group.

Acknowledgments

Funding: This study is supported by grants from the National Institutes of Health

(R01AI154722) and OncoC4, Inc.

Author contributions: M.M.M performed the study with technical assistance from X.D. M.L.,

X.W., W.W., and C.A.; M.M.M, Y.L, P.Z, and L.S. designed and/or supervised the research.

M.M.M and Y.L. wrote the paper with input from other coauthors.

Competing interests: Y.L. and P.Z are co-founders of OncoC4, Inc. Other authors declare no

competing interests**.**

Supplementary Materials

Table S1. Drug to antibody ratio (DAR)

Figure 1-figure supplement 1. CTLA-4 antibody-drug conjugate killing is CTLA-4 specific

Figure 1-figure supplement 2. Gating strategy for T cells

Figure 1-figure supplement 3. CD4-nonTregs and CTLA-4 expression

Figure 1-figure supplement 4. B-cell depletion is not mediated by DM1 payload off-target

release

Figure 1-figure supplement 5. B-cells do not express CTLA-4

Figure 1-figure supplement 6. Treg and B cells in spleen and lymph nodes

Figure 1-figure supplement 7. IgM negative B-cell subtypes in bone marrow

Figure 2-figure supplement 1. B cell apoptosis in blood and lymphoid organs

Figure 3-figure supplement 1. Treg impairment increases plasma immunoglobulins

Figure 3-figure supplement 2. CTLA-4 antibody-drug conjugate leads to enlarged spleen and

lymph nodes

Figure 4-figure supplement 1. CTLA-4 antibody-drug conjugate impact on T cells in lymphoid

organs

Figure 4-figure supplement 2. CTLA-4 antibody-drug conjugate increase T cell proliferation in

lymphoid organs

Figure 4-figure supplement 3. T-cell quantity and proliferation are not impacted by IgG-DM1

treatment

Figure 6-figure supplement 1. Mutant soluble CTLA-4-Ig does not neutralize Ipilimumab or its

drug conjugate

Figure 6-figure supplement 2. T-cell cytokine production

Figure 7-figure supplement 1. Anti-FASL fails to rescue B cells

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Figure 1. CTLA-4 antibody-drug conjugate impairs regulatory T-cell and leads to B-cell depletion.

(A) SDS-Reducing gel showing size shift after DM1 conjugation. Lanes (left to right) are 1 hIgGFc, 2 hIgGFc-DM1 antibody-drug conjugate (ADC), 3 Ipilimumab (Ipi), 4 Ipilimumab-DM1 (Ipi-DM1) antibody-drug conjugate (ADC), and 5 Ladder. (B) ELISA binding of hIgGFc,

hIgGFc-DM1, Ipi, and Ipi-DM1 to pre-coated 1 μg/mL His-hCTLA-4 and detected with anti-

hIgG-HRP (n=2). (C) Flow cytometry binding of Ipi and Ipi-DM1 to hCTLA-4 expressing CHO

cells (CHO-hCTLA-4) and detected with anti-hIgG-AF488. (D) Diagram of experimental

design, $Ctla4^{h/h}$ mice were treated intraperitoneally (i.p.) (100 μg/mouse) with hIgGFc or Ipi-

DM1 every three days and mice were bled or harvested for bone marrow extraction on day 9 for

downstream flow analysis. (E-I) flow data analysis of day 9 peripheral blood, Tregs defined

($CD45^+ CD4^+ Foxp3^+$) and B cells defined ($CD45^+ B220^+$). (E) % $Foxp3^+$ in CD4 and

normalized cell number. (F) Relative CTLA-4 Level in Tregs. (G) % $Ki67^+$ in Tregs. (H) FACS

profiles depicting gating strategy after gating on $CD45^+$ and data summaries of % $B220^+$ in

CD45 and normalized cell number. (I) % $Ki67^+$ in B cells. (J-R) Flow data analysis of day 9 bone

marrow, Tregs defined ($CD4^+B220^-Foxp3^+$) and $B220^+$ B cell subtypes defined mature (M)

($IgD^{hi} IgM^{low}$), immature transitional type 1 (T1) ($IgD^{low}IgM^+$), and Pre-pro/Pro/Pre B

($B220^+IgM^-$). (J) % $Foxp3^+$ in $CD4^+B220^-$ and absolute cell number. (K) Relative CTLA-4 Level

in Tregs. (L) % $Ki67^+$ in Tregs. (M) FACS profile gating of M and T1 B cells, (N) % of mature

B cells in B220 and absolute cell number summaries, (O) % of T1 immature B cells in B220 and

absolute cell number summaries. (P) FACS profile gating of Pre-pro/Pro/Pre B cells, (Q) % of

Pre-pro/Pro/Pre B cells in B220 and absolute cell number summaries. (R) CD21/35 expression in

mature B cells in bone marrow. (B, C) Representative data of two or more repeats. (E-I) Data

combined from two independent experiments (n=13-14). (J-R) Data representative of two

independent experiments (n=5). Data analyzed using an unpaired two-tailed Student's t test and

represented as mean ± SEM. *$p < 0.05$, **$p < 0.01$, ***$p < 0.001$, ****$p < 0.0001$.

Figure 2. CTLA-4 antibody-drug conjugate increases apoptosis in mature B cells in bone

marrow.

Flow analysis of B cell apoptosis on day 9 after treatment with hIgGFc or Ipi-DM1 ADC. (A)

FACS profile defining B220^{hi} and B220^{lo} B cell populations in bone marrow (B) Gating strategy

for B220^{hi} (mature B cells) and B220^{lo} (progenitor/immature B cells) for apoptosis analysis. (C,

D) FACS profiles and summaries depicting apoptosis of B cells in (C) B220^{hi} bone marrow

cells, (D) B220^{lo} bone marrow cells. Data representative of two independent experiments (n=5)

and analyzed using an unpaired two-tailed Student's t test and represented as mean ± SEM. Non-

significant (ns) *$p < 0.05$, **$p < 0.01$, ***$p < 0.001$, ****$p < 0.0001$.

Figure 3. B-cell depletion is transient and correlated with a decrease in regulatory T-cell.

(A) Diagram of experimental design, $Ctla4^{h/h}$ mice were treated intraperitoneal (i.p.)

(100 μg/mouse) with hIgGFc or Ipi-DM1 every three days and mice were bled on day 3,9,13, 25

for downstream flow analysis. (B-F) Flow data of time course of CD45, B, and Treg cells in

peripheral blood, (B) CD45 normalized cell number, (C) % of B-cells in CD45, (D) B-cell

normalized cell number, (E) % Foxp3^{+} Tregs in CD4, (F) Treg normalized cell number. (G-J)

Flow analysis of B-cell subtypes from day 13 peripheral blood. (G) FACS profile of Mature (M)

(IgD^{hi} IgM^{low}), and immature Transitional type 1 (T1) (IgD^{low} IgM^{+}) B cell subtypes after gating

on CD45^{+}B220^{+} B cells. (H) % of mature B cells in B220. (I) % of T1 B cells in B220. (B-F)

Data combined from two independent experiments (n=10). (G-I) Data representative of two

independent experiments (n=6). Data were analyzed by an unpaired two-tailed Student's t test

and represented as mean ± SEM. *p < 0.05, **p < 0.01, ***p < 0.001, ****p < 0.0001

Figure 4. CTLA-4 antibody-drug conjugate leads to T-cell activation.

Flow data analysis of peripheral blood from $Ctla4^{h/h}$ mice on day 9 after treatment with hIgGFc

or Ipi-DM1 ADC. (A) % CD4$^+$ in CD45 and normalized cell number. (B) % CD8$^+$ in CD45 and

normalized cell number. (C) % Ki67$^+$ in CD4 and CD8 T cells. (D, E) FACS profiles and

summaries depicting the increase in effector T-cells after Ipi-DM1 treatment, naïve (Q1: CD44^{low}

CD62L^{hi}), central memory (Q2: CD44^{hi} CD62L^{hi}) and effector (Q3: CD44^{hi} CD62L^{low}),

phenotype of CD4$^+$ T cells (D), and CD8$^+$ T cells (E). (B-C) Data combined from two

independent experiments (n=13-14). (D-E) Data representative of two independent experiments

(n=5). Data analyzed using an unpaired two-tailed Student's t test and represented as mean ±

SEM. Non-significant [ns], *p < 0.05, **p < 0.01, ***p < 0.001, ****p < 0.0001.

Figure 5. B-cell depletion depends on T-cells but not macrophage.

(A) Diagram of experimental design, male or female *Ctla4$^{h/h}$* mice were treated intraperitoneal

(i.p.) (100 μg/mouse) with hIgGFc or Ipi-DM1 with/out (T-cell depleting antibody 100 μg/mouse

of anti-Thy1.2 or 150μL/mouse of clondrosome or depleting antibody CD4 or CD8

100 μg/mouse) every three days and mice were bled on day 9 for downstream flow analysis. (B)

FACS profiles depicting gating strategy after gating on CD45 for CD8 & CD4 T (top panel) and

B cells (bottom panel). (C-E) Data summaries from FACS profile(panel B), (C) % CD4^{+} in CD45

and normalized cell number, (D) % CD8^{+} in CD45 and normalized cell number, and (E) % B220^{+}

in CD45 and normalized cell number. (F) FACS profiles depicting gating strategy after gating on

CD45 for CD8 & CD4 T (top panel) and B cells (bottom panel). (G-I) Data summaries from

FACS profile(panel F), (G) % $CD4^+$ in CD45 and normalized cell number, (H) % $CD8^+$ in CD45

and normalized cell number, and (I) % $B220^+$ in CD45 and normalized cell number. (B-E) Data

combined from two independent experiments (n=10) and analyzed by ordinary one-way ANOVA

with Tukey's multiple comparisons test and represented as mean ± SEM. Non-significant [ns],

*$p < 0.05$, **$p < 0.01$, ***$p < 0.001$, ****$p < 0.0001$. (F-I) Data combined from two independent

experiments (n=11) and analyzed by unpaired two-tailed Student's t test and represented as mean

± SEM. *$p < 0.05$, **$p < 0.01$, ***$p < 0.001$, ****$p < 0.0001$.

Figure 6. Mutant soluble CTLA-4-Ig rescues B-cell.

Ctla4[h/h] mice were treated intraperitoneal (i.p.) (100 μg/mouse) with hIgGFc or Ipi-DM1

with/out (100 μg/mouse of Belatacept) every three days for total of three doses and mice were

bled on day 9 for downstream flow analysis.(A, D) FACS profiles depicting gating strategy after

gating on CD45 for (A) Tregs (% Foxp3^{+} in CD4 T cells) and (D) B cells (% B220^{+} in CD45).

(B) % Foxp3^{+} (Tregs) in CD4 and normalized cell number. (C) Representative FACS profile of

CTLA-4 expression in Tregs (CD4^{+} Foxp3^{+}) and Relative CTLA-4 summary. (E) % B220^{+} in

CD45 and normalized cell number. (F-G) FACS gate and summaries showing mutant CTLA-4-Ig

can decrease effector memory T-cells associated with Ipi-DM1 treatment, naïve (Q1: CD44^{low}

CD62L^{hi}), central memory (Q2: CD44^{hi} CD62L^{hi}) and effector (Q3: CD44^{hi} CD62L^{low}),

phenotype of CD4^{+} T cells (F), and CD8^{+} T cells (G). (H) FACS of GranzymeB gating in CD45

(Top panel), CD4 (middle panel), and CD8 (bottom panel). (I-K) GranzymeB expression

summaries in (I) CD45, (J) CD4, and (K) CD8 cells. (A-E) Data combined from three

independent experiments (n=15). (F-K) Data representative of two independent experiments

(n=5). Data analyzed by ordinary one-way ANOVA with Tukey's multiple comparisons test and

represented as mean ± SEM. Non-significant [ns], $*p < 0.05$, $**p < 0.01$, $***p < 0.001$,

$****p < 0.0001$.

Figure 7. B-cell depletion is partially rescued by anti-TNF-alpha.

(A) ELISA binding of clinical grade drug Adalimumab (Humira) to pre-coated 1μg/mL mouse-

TNF-alpha and detected with anti-hIgG-HRP, Ipilimumab negative control. (B) Diagram of

experimental design, male $Ctla4^{h/h}$ mice were treated intraperitoneal (i.p.) (100 μg/mouse) with

hIgGFc or Ipi-DM1 with/out (100 μg/mouse of Adalimumab) every three days for total of three

doses and mice were bled on day 9 and 13. (C) FACS profiles depicting gating strategy after

gating on CD45 for Tregs (% Foxp3$^+$ in CD4) (top panel) and B cells (% B220$^+$ in CD45)

(bottom panel) from day 9 peripheral blood. (D) % Foxp3$^+$ (Tregs) in CD4 and normalized cell

number. (E) % B220$^+$ in CD45 and normalized cell number. (F, G) Day 13 peripheral blood post

red blood lysis by ACK buffer were cultured and stimulated in the presence of

Iononmycin/PMA, and GolgiPlug for 4 hrs followed by intracellular cytokine detection (TNF-

alpha, IFN-gamma) in CD4 and CD8 T cells, (F) TNF-alpha, and (G) IFN-gamma. (C-E) Data

693   combined from two independent experiments (n=11-12) and analyzed using an unpaired two-

694   tailed Student's t test and represented as mean ± SEM. $*p < 0.05$, $**p < 0.01$, $***p < 0.001$,

695    $****p < 0.0001$. (F, G) Data representative of two independent experiments (n=5-6) and analyzed

696   by ordinary one-way ANOVA with Tukey's multiple comparisons test and represented as mean ±

697   SEM. $*p < 0.05$, $**p < 0.01$, $***p < 0.001$, $****p < 0.0001$.

Table 1 Drug to antibody ratio (DAR)

Entry	A_{280}	A_{252}	CmAb (Moles)	CDM1(Moles)	DAR CDM1/CmAB
hIgGFC-DM1	0.51	0.48	6.92705E-06	1.1531E-05	1.7
Ipilimumab-DM1	0.62	0.42	2.30582E-06	7.32032E-06	3.2

Drug to antibody ratio (DAR)= CDM1/ CmAB. DM1 known extinction coefficients (A_{280}, 5700 $M^{-1}cm^{-1}$; A_{252}, 28084 $M^{-1}cm^{-1}$). Experimentally derived extinction coefficients for Ipilimumab (A_{280}, 242115 $M^{-1}cm^{-1}$; A_{252}, 92989 $M^{-1}cm^{-1}$) and hIgGFc (A_{280}, 64136 $M^{-1}cm^{-1}$; A_{252}, 22544 $M^{-1}cm^{-1}$)

708   **Table S1. Drug to antibody ratio (DAR)**

Figure 1-figure supplement 1. CTLA-4 antibody-drug conjugate killing is CTLA-4 specific.

Cell viability of CHO-hCTLA-4 cells and wild type CHO cells after 72 hours incubation with Ipi

or Ipi-DM1 as measured by MTT assay (n=2). Data representative of two or more repeats.

Figure 1-figure supplement 2. Gating strategy for T cells.

(A) Tregs (CD4$^+$ Foxp3$^+$) and CD4-nonTreg (CD4$^+$ Foxp3$^-$) in blood. (B) CD8-T (CD4$^+$B220$^-$),

CD4-T (CD4$^+$B220$^-$), Tregs (CD4$^+$B220$^-$Foxp3$^+$) and CD4-nonTreg (CD4$^+$B220$^-$Foxp3$^-$) in bone

marrow.

Figure 1-figure supplement 3. CD4-nonTregs and CTLA-4 expression.

(A, B) Blood, (A) % CD4$^+$Foxp3$^-$ in CD45 and normalized cell number, (B) relative CTLA-4

level in CD4-nonTregs. (C, D) Bone marrow, (C) % CD4$^+$B220$^-$Foxp3$^-$ in CD45 and absolute

cell number, (B) relative CTLA-4 level in CD4-nonTregs (CD4$^+$B220$^-$ Foxp3$^-$). (A, B) Data

combined from two independent experiments (n=13-14). (C, D) Data representative of two

independent experiments (n=5). Data analyzed using an unpaired two-tailed Student's t test and

represented as mean $\pm$ SEM. *p < 0.05, **p < 0.01, ***p < 0.001, ****p < 0.0001.

Figure 1-figure supplement 4. B-cell depletion is not mediated by DM1 payload off-target

release.

Mice treated with hIgGFc or hIgGFc-DM1, blood flow analysis on day 9. (A) % Foxp3^{+} in CD4

and normalized cell number. (B) Relative CTLA-4 level in Tregs. (C) % Ki67^{+} in Tregs. (D) %

CD4^{+}Foxp3^{-} in CD45 and normalized cell number. (E) Relative CTLA-4 level in CD4-nonTregs.

(F) FACS profiles depicting gating strategy after gating on CD45^{+} and data summaries of %

B220^{+} in CD45 and normalized cell number. (G) % Ki67^{+} in B cells. Data combined from two

independent experiments (n=11) and analyzed using an unpaired two-tailed Student's t test and

represented as mean ± SEM. *$p < 0.05$, **$p < 0.01$, ***$p < 0.001$, ****$p < 0.0001$.

749 **Figure 1-figure supplement 5. B-cells do not express CTLA-4.**

750 FACS profile of human CTLA-4 in wild-type and human CTLA-4 knock-in mice peripheral

751 blood, (A) % CTLA-4$^+$ in Tregs (CD45$^+$ CD4$^+$ Foxp3$^+$), and (B) % CTLA-4$^+$ in B cells (CD45$^+$

752 B220$^+$). Data representative of 2 mice.

Figure 1-figure supplement 6. Tregs and B cells in spleen and lymph nodes.

Analysis of spleen and lymph nodes from $Ctla4^{h/h}$ mice on day 9 after treatment with hIgGFc or

Ipi-DM1 ADC. (A, B) Spleen, (A) % Foxp3+ in CD4 and absolute cell number, (B) % B220+ in

CD45 and absolute cell number. (C, D) combined inguinal and cervical lymph nodes, (C) %

Foxp3+ in CD4 and absolute cell number, (D) % B220+ in CD45 and absolute cell number. Data

representative of two independent experiments (n=5-7) and analyzed using an unpaired two-

tailed Student's t test and represented as mean ± SEM. Non-significant [ns], *p < 0.05, **p <

0.01, ***p < 0.001, ****p < 0.0001.

Figure 1-figure supplement 7. IgM negative B-cell subtypes in bone marrow.

773 (A) FACS profile gating of Pre B (IgM⁻CD43⁻) and Pre-pro/Pro (IgM⁻CD43⁺) cells in B220. (B-

774 C) % of B cell subtype in B220 and absolute cell numbers, (B) Pre B cells, (C) Pre-pro/Pro B

775 cells. Data representative of two independent experiments (n=5) and analyzed using an unpaired

776 two-tailed Student's t test and represented as mean ± SEM. *p < 0.05, **p < 0.01, ***p < 0.001,

777 ****p < 0.0001.

Figure 2-figure supplement 1. B cell apoptosis in blood and lymphoid organs.

B cell apoptosis on day 9 after treatment with hIgGFc or Ipi-DM1 ADC. (A) B220^{+} blood cells.

(B) B220^{+} spleen cells. (C) B220[+] lymph node cells. Data representative of two independent

experiments (n=5) and analyzed using an unpaired two-tailed Student's t test and represented as

mean ± SEM. Non-significant [ns], *p < 0.05, **p < 0.01, ***p < 0.001, ****p < 0.0001.

Figure 3-figure supplement 1. Treg impairment increases plasma immunoglobulins.

Mice treatment with hIgGFc or Ipi-DM1 ADC and plasma collected on day 9, 13, 25 for ELISA

Ig level quantification. (A) IgG. (B) IgM. (C) IgA. (D) IgE. Data representative of two

independent experiments (n=5) and analyzed using ordinary two-way Anova with Bonferroni's

multiple comparisons test and as mean ± SEM. Non-significant [ns], *p < 0.0332, **p < 0.0021,

***p < 0.0002, ****p < 0.0001.

Figure 3-figure supplement 2. CTLA-4 antibody-drug conjugate leads to enlarged spleen

and lymph nodes.

Day 9 or 13 after treatment with hIgGFc or Ipi-DM1 ADC. (A) Spleen weight. (B)

Splenomegaly. (C) Lymphadenopathy. (A) Data representative of two independent experiments

(n=5-7) and analyzed using an unpaired two-tailed Student's t test and represented as mean ±

SEM. Non-significant [ns], *p < 0.05, **p < 0.01, ***p < 0.001, ****p < 0.0001.

Figure 4-figure supplement 1. CTLA-4 antibody-drug conjugate impact on T cells in lymphoid organs.

Flow analysis of T cells at day 9 after treatment with hIgGFc or Ipi-DM1 ADC. (A, B) Bone marrow, (C-G) thymus, (H, I) spleen, (J, K) combined inguinal and cervical lymph nodes. (A) % $CD4^+B220^-$ in CD45 and absolute cell number. (B) % $CD8^+B220^-$ in CD45 and absolute cell number. (C) Gating strategy defining T-cell types in $CD45^+$ cells in thymus. (D) Tregs (% $Foxp3^+$ in $CD4^+CD8^-$) and absolute cell number. (E) % $CD4^+CD8^-$ in CD45 and absolute cell number. (F) % $CD8^+CD4^-$ in CD45 and absolute cell number. (G) % $CD4^+CD8^+$ in CD45 and absolute cell number. (H, J) % $CD4^+$ in CD45 and absolute cell number. (I, K) % $CD8^+$ in CD45 and absolute cell number. Data representative of two independent experiments (n=5-7) and analyzed using an unpaired two-tailed Student's t test and represented as mean ± SEM. *$p < 0.05$, **$p < 0.01$, ***$p < 0.001$, ****$p < 0.0001$.

Figure 4-figure supplement 2. CTLA-4 antibody-drug conjugate increase T cell

proliferation in lymphoid organs.

816   Flow analysis at day 9 after treatment with hIgGFc or Ipi-DM1 ADC. (A) % Ki67$^+$ in

817   CD4$^+$B220$^-$ and CD8$^+$B220$^-$ T cells in bone marrow. (B) % Ki67$^+$ in CD4 and CD8 T cells in

818   spleen. (C) % Ki67$^+$ in CD4 and CD8 T cells in lymph nodes. Data representative of two

819   independent experiments (n=5-7) and analyzed using an unpaired two-tailed Student's t test and

820   represented as mean ± SEM. *p < 0.05, **p < 0.01, ***p < 0.001, ****p < 0.0001.

Figure 4-figure supplement 3. T-cell quantity and proliferation are not impacted by IgG-DM1 treatment.

Mice treated with hIgGFc or hIgGFc-DM1, blood flow analysis on day 9. (A) % CD4$^+$ in CD45 and normalized cell number. (B) % CD8$^+$ in CD45 and normalized cell number. (C) % Ki67$^+$ in CD4 and CD8 T cells. Data combined from two independent experiments (n=11) and analyzed using an unpaired two-tailed Student's t test and represented as mean ± SEM. *p < 0.05, **p < 0.01, ***p < 0.001, ****p < 0.0001.

Figure 6-figure supplement 1. Mutant soluble CTLA-4-Ig does not neutralize Ipilimumab

or its drug-conjugate.

(A) Detached CHO cells expressing hCTLA-4 were incubated with serial dilution of Ipi or Ipi-

DM1 in the presence of given dose of Abatacept or Belatacept on ice for 30 minutes followed

by FACS detection using mouse anti-human IgG AF488, mean fluorescence intensity (MFI) . (B)

MTT cell viability of CHO-hCTLA-4 cells after 72 hours incubation with Ipi or Ipi-DM1 in the

presence of given dose of Abatacept or Belatacept. Data representative of two independent

experiments (n=3).

Figure 6-figure supplement 2. T-cell cytokine production.

Day 13 peripheral blood samples from treated (hIgGFc or Ipi-DM1 with/out Belatacept) mice

post ACK buffer red blood lysis were cultured and stimulated in the presence of

Iononmycin/PMA, and GolgiPlug for 4 hrs followed by intracellular cytokine detection (TNF-

alpha, IFN-gamma) in CD4 and CD8 T cells. Data is representative of two independent

experiments (n=5) analyzed by ordinary one-way ANOVA with Tukey's multiple comparisons

test and represented as mean ± SEM. *$p < 0.05$, **$p < 0.01$, ***$p < 0.001$, ****$p < 0.0001$.

Figure 7-figure supplement 1. Anti-FASL fails to rescue B cell.

Ctla4$^{h/h}$ mice were treated intraperitoneally (i.p.) (100 µg/mouse) with hIgGFc or Ipi-DM1

with/out (100 µg/mouse of anti-FASL) every three days for a total of three doses and mice were

bled on day 9. (A) FACS profile after gating on CD45 for B cells. (B) % B220^{+} in CD45 and

absolute B cell number summaries. Data representative of one experiment (n=5) analyzed by

ordinary one-way ANOVA with Tukey's multiple comparisons test and represented as mean ±

SEM. *$p < 0.05$, **$p < 0.01$, ***$p < 0.001$, ****$p < 0.0001$.