### **RESEARCH ARTICLE**

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# A mutation that blocks integrin $\alpha_4\beta_7$ activation prevents adaptive immunemediated colitis without increasing susceptibility to innate colitis



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### **Abstract**

**Background:**  $β_7$  integrins are responsible for the efficient recruitment of lymphocytes from the blood and their retention in gut-associated lymphoid tissues. Integrin  $α_4β_7$  binds MAdCAM-1, mediating rolling adhesion of lymphocytes on blood vessel walls when inactive and firm adhesion when activated, thereby controlling two critical steps of lymphocyte homing to the gut. By contrast, integrin  $α_Eβ_7$  mediates the adhesion of lymphocytes to gut epithelial cells by interacting with E-cadherin. Integrin  $β_7$  blocking antibodies have shown efficacy in clinical management of inflammatory bowel disease (IBD); however, fully blocking  $β_7$  function leads to the depletion of colonic regulatory T (Treg) cells and exacerbates dextran sulfate sodium (DSS)-induced colitis by evoking aberrant innate immunity, implying its potential adverse effect for IBD management. Thus, a better therapeutic strategy targeting integrin  $β_7$  is required to avoid this adverse effect.

**Results:** Herein, we inhibited integrin  $\alpha_4\beta_7$  activation in vivo by creating mice that carry in their integrin  $\beta_7$  gene a mutation (F185A) which from structural studies is known to lock  $\alpha_4\beta_7$  in its resting state. Lymphocytes from  $\beta_7$ -F185A knock-in (KI) mice expressed  $\alpha_4\beta_7$  integrins that could not be activated by chemokines and showed significantly impaired homing to the gut. The  $\beta_7$ -F185A mutation did not inhibit  $\alpha_E\beta_7$  activation, but led to the depletion of  $\alpha_E\beta_7^+$  lymphocytes in the spleen and a significantly reduced population of  $\alpha_E\beta_7^+$  lymphocytes in the gut of KI mice.  $\beta_7$ -F185A KI mice were resistant to T cell transfer-induced chronic colitis, but did not show an increased susceptibility to DSS-induced innate colitis, the adverse effect of fully blocking  $\beta_7$  function.

**Conclusions:** Our findings demonstrate that specific inhibition of integrin  $\alpha_4\beta_7$  activation is a potentially better strategy than fully blocking  $\alpha_4\beta_7$  function for IBD treatment.

**Keywords:** Inflammatory bowel disease, Integrin  $\alpha_4\beta_7$ , Activation, Lymphocyte

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### **Background**

Inflammatory bowel disease (IBD), including ulcerative colitis (UC) and Crohn's disease (CD), is an idiopathic intestinal disorder caused by an inappropriate inflammatory response to intestinal microbes in a genetically susceptible host [1, 2]. The pronounced infiltration of both innate and adaptive immune cells is observed in the gut of UC and CD patients [3, 4], suggesting that both types of immunity are involved in the progression of IBD [5]. The migration of leukocytes into inflamed intestinal tissue is tightly regulated by specific cell adhesion molecules [6]. The cell-surface glycoprotein  $\beta_7$  integrins regulate the homing and retention of lymphocytes in the gut-associated lymphoid tissues (GALT) [7]. Integrin  $\alpha_4\beta_7$  mediates the homing of lymphocytes into the GALT via interaction with mucosal addressin cell adhesion molecule-1 (MAdCAM-1) on the intestinal vasculature [8, 9], whereas integrin  $\alpha_E \beta_7$  facilitates the retention of lymphocytes in the gut epithelium through binding to E-cadherin [10]. The highest expression of  $\alpha_4\beta_7$  can be found on memory gut-homing CD4+ T cells, and it is also expressed on other T cell subsets (T<sub>H</sub>2 and T<sub>H</sub>17) and B cells as well as some other leucocytes [11-13].  $\alpha_E \beta_7$  mainly expresses in CD8<sup>+</sup> intraepithelial T cells,  $T_H9$ ,  $CD69^+\alpha_E^+$  intestinal tissue-resident memory T(T<sub>RM</sub>) cells, Treg cells, and mucosal dendritic cell subsets [14–16]. Of note, integrin  $\beta_7$  has emerged as a promising drug target for the treatment of IBD since blocking  $\beta_7$  function suppresses the migration of inflammatory lymphocytes to the GALT [17, 18] and consequently inhibits adaptive immune-mediated colitis [19].

Clinical studies have reported that two humanized monoclonal antibodies that can block integrin  $\beta_7$  function, vedolizumab (anti- $\alpha_4\beta_7$  antibody) and etrolizumab (anti- $\beta_7$  antibody), effectively maintain clinical remission in IBD patients [20, 21]. Vedolizumab specifically blocks  $\alpha_4\beta_7$  interaction with MAdCAM-1 by binding to the specificity-determining loop (SDL) that projects from the  $\beta_7$  I domain at the interface with the  $\alpha_4$   $\beta$ -propeller domain [22]. This epitope is only accessible in the heterodimer of  $\alpha_4$  and  $\beta_7$  subunits. Etrolizumab binds to the  $\beta_7$  subunit of both  $\alpha_4\beta_7$  and  $\alpha_E\beta_7$  integrins and blocks the interaction of  $\beta_7$  integrins with their ligands MAdCAM-1 and E-cadherin, respectively [23]. Interestingly, UC patients who received 100 mg etrolizumab show a significantly higher remission rate than patients receiving 300 mg etrolizumab [21]. Moreover, aggravated colitis is observed in a small percentage of UC patients treated with a high dose of vedolizumab [20]. These reports suggest that excessive inhibition of  $\beta_7$  function may have adverse effects on the management of IBD under certain conditions. Indeed, total loss of  $\alpha_4\beta_7$  function has been shown to lead to the depletion of colonic Treg cells and consequently exacerbate dextran sulfate sodium (DSS)-induced colitis by evoking aberrant innate immunity [24]. Moreover, vedolizumab treatment significantly suppresses homing of Treg cells to the gut in UC patients [25]. Therefore, a more specific approach to suppress adaptive immune response without evoking aberrant innate immunity in colitis could be more efficacious for IBD treatment.

Integrin  $\alpha_4\beta_7$  mediates rolling and firm adhesion of lymphocytes when inactive and activated, respectively [22, 26], controlling two critical steps in tissue-specific homing of lymphocytes [27, 28]. Integrin activation can be dynamically regulated via a cluster of three metal ion-binding sites in the  $\beta$  I domain [26, 29]. A cation- $\pi$  interaction in the human  $\beta_7$  I domain between F185 in the SDL and the synergistic metal ion-binding site (SyMBS) cation has been shown to be essential for the activation of  $\alpha_4\beta_7$  [30]. Disruption of the cation- $\pi$  interaction through mutation of F185 to Ala inhibits  $\alpha_4\beta_7$  activation and  $\alpha_4\beta_7$ -mediated cell migration.

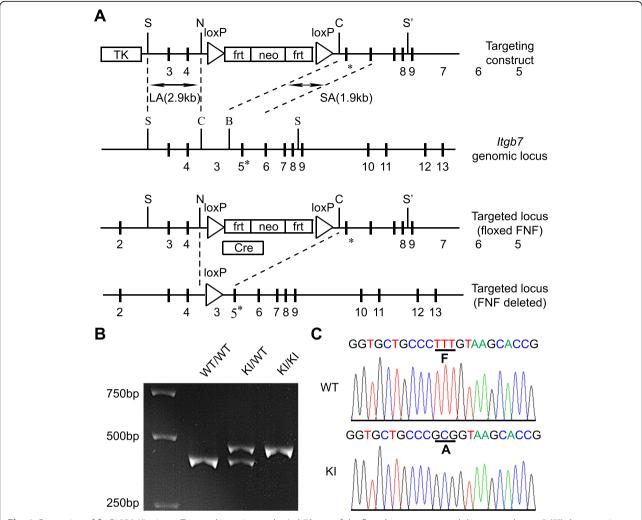
Herein, we generated knock-in (KI) mice bearing the integrin  $\beta_7$ -F185A mutation which blocked  $\alpha_4\beta_7$  activation. Moreover, this mutation did not inhibit the chemokine-induced activation of  $\alpha_E \beta_7$ , but induced the significant reduction of  $\alpha_E \beta_7$  expression in KI mice. Similar to lymphocytes from integrin  $\beta_7$  knock-out (KO) mice, lymphocytes from KI mice showed significantly deficient homing to the GALT, and CD4+CD45RBhigh T cells from KI mice could not induce colitis in a T cell transfer model. Notably, KI mice exhibited sufficient colonic Treg cells, in stark contrast to the depletion of colonic Treg cells in  $\beta_7$  KO mice, and thus avoided the aberrant innate immunity in DSS-induced colitis. Taken together, specific blockade of integrin  $\alpha_4\beta_7$  activation is sufficient to prevent adaptive immune-mediated colitis without increasing susceptibility to innate colitis and therefore is a potentially better treatment for IBD than complete blockade of  $\alpha_4\beta_7$  function.

### **Results**

### Generation of KI mice bearing the β<sub>7</sub>-F185A mutation

To block the activation of integrin  $\alpha_4\beta_7$  in vivo, we generated a mouse with the F185A point mutation in the mouse  $\beta_7$  gene, thereby disrupting the synergistic metal-binding site (SyMBS) cation-F185 interaction within  $\beta_7$  I domain and locking  $\alpha_4\beta_7$  in a resting state, as shown previously for human  $\beta_7$  [30]. Specifically, using a replacement-type gene-targeting strategy [31], a phenylalanine-to-alanine mutation (F185A) was introduced into the mouse integrin  $\beta_7$  gene, *Itgb7*. Embryonic stem cells were electroporated with the targeting construct encoding mutant  $\beta_7$ -F185A and were then injected into blastocysts. Chimeric mice with germline transmission of the targeted allele were crossed with an EIIa-cre mouse to excise the *neo* cassette (Fig. 1a). The correct integration of the mutant  $\beta_7$  gene

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**Fig. 1** Generation of  $β_7$ -F185A Kl mice. **a** Targeted insertion to the *ltgb7* locus of the floxed *neo* cassette, and the mutated exon 5 (5\*) that contains  $β_7$ -F185A. The targeting vector, the WT *ltgb7* locus, the targeted *ltgb7* allele containing floxed *neo* cassette, and the mutated *ltgb7* (F185A) allele are shown. Exons are shown as thick lines, as well as long arm (LA) and short arm (SA) of homology are also shown. The *loxP*-flanked *neo* cassette is deleted by intercrossing the mutant mouse strains with an Ella-Cre strain, leaving 1 loxP site. S, Sacll; N, Not; C, Clal; S', Sall. **b** Genotyping and confirmation of deleted *neo* cassette by PCR. Genomic DNA isolated from tails was used for PCR analyses. PCR bands are shown for WT (WT/WT, 360 bp), heterozygote (KI/WT, 380 and 360 bp), and homozygote (KI/KI, 380 bp) samples. **c** Sequencing analysis of WT and Kl mice. DNA sequencing confirmed a phenylalanine-to-alanine substitution at position 185 of the mouse  $β_7$  integrin gene in Kl mice

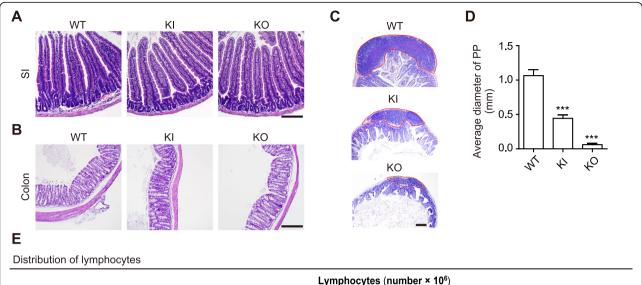
 $Itgb\mathcal{T}^{F185A}$  was confirmed by PCR and DNA sequencing (Fig. 1b, c).  $Itgb\mathcal{T}^{F185A/F185A}$  ( $\beta_7$ -F185A KI) mice showed no abnormalities by visual inspection and had a normal life span. In addition to the  $\beta_7$ -F185A KI mice, we also included  $\beta_7$  KO ( $Itgb\mathcal{T}^{-/-}$ ) mice in all experiments to compare the effects of blockade of  $\alpha_4\beta_7$  activation with total loss of  $\beta_7$  function.

### Reduced lymphocytes in the gut of β<sub>7</sub>-F185A KI mice

The small intestine (SI) and colon of KI and KO mice exhibited basically normal architectures (Fig. 2a, b); however, Peyer's patches (PP) with decreased cellularity and rudimentary follicles were observed in KI and KO mice compared with wild-type (WT) mice (Fig. 2c, d).

The spleen (SP), peripheral lymph nodes (PLN), and mesenteric lymph nodes (MLN) were indistinguishable among WT, KI, and KO mice (Additional file 1: Figure S1). We next analyzed the distribution of lymphocytes in the lymphoid organs of these mice. Flow cytometric analyses showed that compared with WT mice, KI mice contained significantly fewer lymphocytes in the gut including fewer intraepithelial lymphocyte (IEL) and lamina propria lymphocyte (LPL) in the SI and fewer T and B cells in the PP and colon (Fig. 2e). Moreover, KO mice showed a greater decrease in CD3<sup>+</sup> T cells in the gut than did KI mice. Thus, both integrin  $\beta_7$ -F185A mutation and  $\beta_7$  KO can specifically inhibit lymphocyte recruitment to the GALT. It is noteworthy that  $\beta_7$  KO

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	Lymphocytes (number × 10°)					
Organ	CD3 <sup>+</sup>			CD19 <sup>+</sup>		
	WT	KI	ко	WT	KI	ко
PP (per mouse)	0.8±0.01	0.045±0.009 <sup>AA</sup>	0.01±0.006 <sup>AA</sup>	2.6±0.05	0.03±0.004 <sup>AA</sup>	0.02±0.001 <sup>AA</sup>
SI-IEL (per mouse)	3.4±0.01	1.4±0.07 <sup>A</sup>	1.1±0.01 <sup>A</sup>	ND	ND	ND
SI-LPL (per mouse)	1.2±0.03	0.40±0.02 <sup>A</sup>	0.35±0.01 <sup>A</sup>	0.63±0.007	0.25±0.003 <sup>A</sup>	0.22±0.005 <sup>A</sup>
Colon (per mouse) <sup>B</sup>	0.20±0.006	0.1±0.006 <sup>A</sup>	0.06±0.009 <sup>A</sup>	0.10±0.001	0.03±0.001 <sup>A</sup>	0.03±0.002 <sup>A</sup>
Spleen (per mouse)	28.0±3.3	28.6±0.3	30.2±3.6	35.2±4.1	36.0±0.2	38.1±4.1
Thymus (per mouse)	10.5±1.0	9.1±0.1	8.9±0.4	ND	ND	ND
PB (per ml)	0.8±0.04	0.74±0.01	0.66±0.05	0.4±0.02	0.5±0.05	0.4±0.04
BM (per 1 femur)	1.14±0.12	1.32±0.2	1.36±0.5	7.3±0.7	7.8±0.3	8.7±0.7
PLN (per node)	0.20±0.02	0.23±0.03	0.25±0.01	0.09±0.004	0.12±0.001	0.12±0.002
MLN (per mouse)	3.0±0.6	2.8±0.1	2.4±0.5	1.6±0.3	1.4±0.3	1.3±0.2

**Fig. 2** Reduced lymphocytes in the GALT of  $β_7$ -F185A KI mice. Representative histological sections of the small intestine (SI) (**a**), colon (**b**), and Peyer's patch (PP) (**c**) of WT,  $β_7$ -F185A KI (KI), and  $β_7$ -KO (KO) mice were analyzed by hematoxylin and eosin staining. Scale bars, 100 μm. **d** Quantification of the average diameter of PP in the individual group of mice (n = 9 per group). \*\*\*\*P < 0.001 (Student's t test). **e** Flow cytometry enumeration of lymphocyte distribution in lymphoid organs from the individual group of mice (n = 9 per group). PB, peripheral blood; PLN, peripheral lymph node; MLN, mesenteric lymph node; BM, bone marrow; IEL, intraepithelial lymphocyte; LPL, lamina propria lymphocyte. Results are presented as cell number × 10<sup>6</sup>,  $^{A}P < 0.01$ ;  $^{AA}P < 0.005$  (Student's t test).  $^{B}$ The cecum was excluded. ND, not detected. Data are mean ± s.d. of at least 3 independent experiments (**d, e**)

results in a greater inhibition of T cell recruitment to the gut.

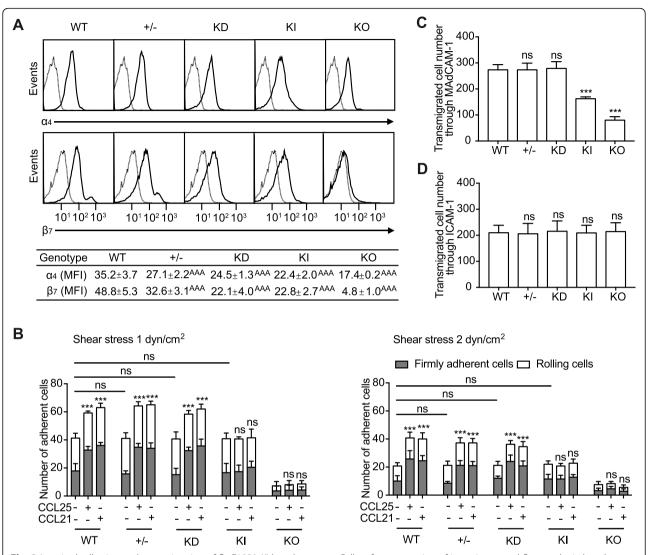
# Chemokine fails to promote $\alpha_4\beta_7$ -mediated adhesion of $\beta_7$ -F185A KI lymphocytes

We found that splenic lymphocytes from KI mice showed an approximately 50% reduction in  $\beta_7$  integrin cell surface expression compared with cells from WT mice (Fig. 3a). Decreased expression of  $\alpha_4$  integrin was also observed in KI and KO mice, likely resulting from the reduction in  $\beta_7$  expression (Fig. 3a). Although quantitative reverse transcription polymerase chain reaction (qRT-PCR) showed that  $\beta_7$  mRNA level was comparable between WT and KI splenic lymphocytes (Additional file 1: Figure S2A), flow

cytometric analysis of permeabilized cells indicated that the total expression of  $\beta_7$  integrin, including cell surface and intracellular expression, was decreased in KI lymphocytes (Additional file 1: Figure S2B).

Next, we examined  $\alpha_4\beta_7$ -mediated splenic lymphocyte adhesion on MAdCAM-1 substrates using a parallel wall flow chamber. Considering the reduced expression of  $\alpha_4\beta_7$  in KI lymphocytes, we included the  $Itgb7^{+/-}$  (+/-) and  $\beta_7$  knock-down (KD) splenic lymphocytes as controls because those cells also express reduced levels of  $\beta_7$  integrins (Fig. 3a). In the presence of the physiologic cations 1 mM  $Ca^{2+}/Mg^{2+}$ , which maintain  $\alpha_4\beta_7$  predominantly in the resting state [26, 30], WT, +/-, KD and KI lymphocytes showed similar adhesive behavior on

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**Fig. 3** Impaired adhesion and transmigration of  $β_7$ -F185A KI lymphocytes. **a** Cell surface expression of integrins  $α_4$  and  $β_7$  on splenic lymphocytes from WT,  $ltgb7^{+/-}$  (+/-),  $β_7$  knock-down (KD), KI, and KO mice. All viable lymphocytes were gated using a combination of forward angle and side scatter to exclude dead cells and debris. And the results were presented as histograms for  $α_4$  and  $β_7$  expression. The numbers within the table show the specific mean fluorescence intensities of FlB504 (anti- $β_7$ ) and GK1.5 (anti- $α_4$ ) mAbs. **b** Adhesion of WT, +/-, KD, KI, and KO splenic lymphocytes to MAdCAM-1 at 1 dyn/cm² or 2 dyn/cm² before and after chemokine stimulation. **c**, **d** Transmigration of WT, +/-, KD, KI, and KO splenic lymphocytes toward a serum gradient through MAdCAM-1-coated (**c**) or ICAM-1-coated (**d**) permeable insert was examined using a modified Boyden chamber assay with a transwell tissue culture system. \*\*\*P < 0.001; ns, not significant. AAAP < 0.001 (Student's t test in **a-d**). Data are mean ± s.d. of at least 3 independent experiments (**a-d**). The asterisk in **b** indicates the changes of total adherent cells

MAdCAM-1 substrates in flow (Fig. 3b), indicating that the F185A mutation and the reduction in  $\alpha_4\beta_7$  expression have no effect on the adhesion of unstimulated lymphocytes to MAdCAM-1. As expected,  $\beta_7$  KO lymphocytes rarely adhered to MAdCAM-1 due to the loss of  $\alpha_4\beta_7$  expression (Fig. 3b).

Activation of  $\alpha_4\beta_7$  by chemokines promotes the firm adhesion of lymphocytes to MAdCAM-1, which is a critical step during lymphocyte homing to the gut [32, 33]. The flow chamber results showed that chemokine (C-C

motif) ligand 25 (CCL25) and chemokine (C-C motif) ligand 21 (CCL21) significantly increased the number of firmly adherent WT, +/-, and KD splenic lymphocytes on MAdCAM-1 substrates, whereas these chemokines were not able to increase the adhesion of KI cells to MAdCAM-1 (Fig. 3b), indicating that chemokines cannot activate the  $\alpha_4\beta_7$ -F185A mutant, thus failing to promote  $\alpha_4\beta_7$ -mediated cell adhesion. Despite the reduced  $\alpha_4\beta_7$  expression on +/- and KD splenic lymphocytes, these cells showed adhesive behavior similar to WT cells

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upon chemokine treatment, suggesting that the impaired adhesion of  $\beta_7$ -F185A cells is not due to the reduction in  $\alpha_4\beta_7$  expression.

# $\beta_7\text{-F185A}$ KI lymphocytes show impaired $\alpha_4\beta_7\text{-mediated}$ transmigration

Next, we studied the impact of defective  $\beta_7$  integrin activation on  $\alpha_4\beta_7$ -mediated lymphocyte transmigration using a transwell assay. Compared with WT lymphocytes, significantly fewer KI cells transmigrated through the MAdCAM-1-coated insert (Fig. 3c), indicating that blockade of  $\beta_7$  integrin activation perturbed  $\alpha_4\beta_7$ -mediated cell transmigration. Moreover, knockout of  $\beta_7$  led to a further decrease in lymphocyte transmigration across MAdCAM-1 substrates (Fig. 3c). Similar numbers of WT, +/-, and KD splenic lymphocytes transmigrated through a MAdCAM-1-coated insert, suggesting that the decreased expression of  $\beta_7$  integrin does not affect  $\alpha_4\beta_7$ -mediated cell transmigration. As a control, WT, +/-, KD, KI, and KO splenic lymphocytes showed the intact ability of  $\beta_2$  integrin to mediate cell transmigration through an ICAM-1-coated insert (Fig. 3d).

To confirm the effects of  $\beta_7$ -F185A mutation on the adhesion and migration of human lymphocytes, we established Jurkat T- $\beta_7$  WT and Jurkat T- $\beta_7$  F185A, which stably expressed similar level of WT or F185A β<sub>7</sub> (Additional file 1: Figure S3A). Because Jurkat T cells lack CCL25 chemokine receptor 9 (CCR9), CCR9 was also co-transfected into those cells. Consistent with the results of mouse lymphocytes, CCL25 and CCL21 only promoted the adhesion of Jurkat  $T-\beta_7$  WT but not Jurkat T-β<sub>7</sub> F185A to MAdCAM-1 (Additional file 1: Figure S3B). Compared with Jurkat T- $\beta_7$  WT cells, significantly fewer Jurkat T-β<sub>7</sub> F185A cells transmigrated through the MAdCAM-1-coated insert (Additional file 1: Figure S3C). Taken together, inhibition of integrin  $\beta_7$  activation has the same effects on the adhesion and migration of mouse and human lymphocytes.

### $\alpha_E \beta_7^+$ lymphocytes are reduced in $\beta_7$ -F185A KI mice

In addition to integrin  $\alpha_4\beta_7$ , the  $\beta_7$  subunit also forms a heterodimer with integrin  $\alpha_E$  subunit. The  $\alpha_E\beta_7$  integrin facilitates the retention of lymphocytes in the gut epithelial layer via interactions with E-cadherin. In contrast to the inhibition of  $\alpha_4\beta_7$  activation by  $\beta_7$ -F185A, CCL21 and CCL25 promoted the adhesion of both Jurkat T- $\alpha_E\beta_7$  WT and Jurkat T- $\alpha_E\beta_7$  F185A to E-cadherin (Additional file 1: Figure S4), indicating that  $\beta_7$ -F185A mutation does not inhibit chemokine-induced activation of  $\alpha_E\beta_7$ . Unlike integrin  $\alpha_4\beta_7$  which binds to ligand via the metal ion-dependent adhesion site (MIDAS) in  $\beta_7$  subunit,  $\alpha_E\beta_7$  has an I domain in the  $\alpha_E$  subunit and binds to E-cadherin through MIDAS in  $\alpha_E$  I domain [26, 34]. Thus it is not surprising that the  $\beta_7$ -F185A mutation

inhibits the cytokine-activated β<sub>7</sub>-mediated MAdCAM-1, but not the  $\alpha_E$ -mediated adhesion to sion to E-cadherin. We observed, however, a significant decrease of  $\alpha_E \beta_7^+$  splenic lymphocytes in KI mice and the depletion of  $\alpha_E \beta_7^+$  splenic lymphocytes in KO mice (Additional file 1: Figure S5A). Furthermore, flow cytometry showed that the percentages of  $\alpha_E^+$  IEL and  $\alpha_E^+$ LPL significantly decreased in the gut of KI mice and almost disappeared in KO mice (Additional file 1: Figure S5B), these percentage changes being seen in the context of the reduced total IEL and LPL populations in these mice (Fig. 2e). In addition, the expression level of  $\alpha_E$  is significantly decreased in  $\alpha_E^+$  IEL and LPL populations in KI and KO mice (Additional file 1: Figure S5C). Because  $\alpha_E \beta_7$  is critical for the retention of intestinal lymphocytes and the  $\alpha_E$  KO has been shown to induce the reduction of intestinal lymphocytes in mice [35], it is tempting to speculate that  $\beta_7$ -F185A mutation-induced downregulation of  $\alpha_E \beta_7$  expression impaired the retention of intestinal lymphocytes in KI mice. To summarize, although the  $\beta_7$ -F185A mutation did not inhibit the chemokine-induced activation of  $\alpha_E \beta_7$ , it led to the significant reduction of  $\alpha_E \beta_7$ -expressing lymphocytes and reduced the  $\alpha_E$  expression level in  $\alpha_E \beta_7^+$  IELs and LPLs in KI mice, which would be expected to impair lymphocyte retention in the gut.

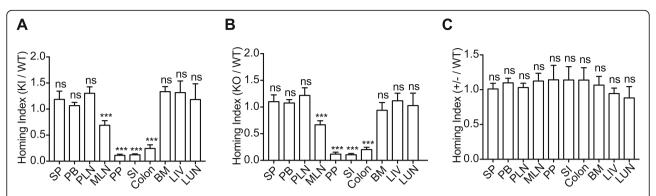
### Suppressed β<sub>7</sub>-F185A lymphocyte homing to the GALT

Next, we investigated whether lymphocyte homing to the GALT was affected by the defective activation of  $\beta_7$  integrin using a competitive homing assay [36]. Splenic lymphocytes freshly isolated from WT and KI mice were fluorescently labeled with CellTrace Violet and CellTrace Yellow, respectively. Equal numbers of WT and KI lymphocytes were mixed and then intravenously administered into C57BL/6J recipient mice. Organs were harvested 18 h after administration, and the homing indices were then determined. WT and KI lymphocytes homed equally well to the SP, PLN, bone marrow (BM), liver (LIV), and lung (LUN), whereas homing of KI lymphocytes to MLN, PP, SI, and the colon was severely decreased in comparison (Fig. 4a). This indicates that blockade of  $\beta_7$  integrin activation specifically suppresses lymphocyte homing to the GALT. In addition, KO lymphocytes showed decreased homing to the GALT similar to that seen with KI lymphocytes (Fig. 4b), while the homing of +/- lymphocytes was not affected (Fig. 4c). Collectively, these results demonstrate that either integrin  $\beta_7$ -F185A mutation or  $\beta_7$  KO can efficiently inhibit gut-specific homing of lymphocytes.

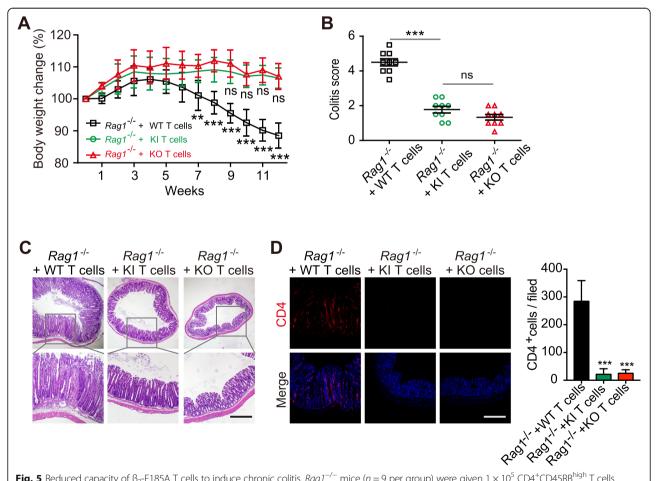
# Reduced capacity of $\beta_7$ -F185A T cells to induce chronic colitis

Integrin  $\beta_7$ -mediated T lymphocyte recruitment to the gut has been implicated in the pathogenesis of chronic

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**Fig. 4** Reduced in vivo homing of  $β_7$ -F185A lymphocytes to the GALT. In vivo competitive homing of splenic lymphocytes from WT mice and KI (a) or KO (b) or +/- mice (c). The ratio of KI, KO, or +/- lymphocytes over WT lymphocytes that homed to specific tissues was determined 18 h after injection. SP, spleen; PB, peripheral blood; PLN, peripheral lymph node; MLN, mesenteric lymph node; PP, Peyer's patch; SI, small intestine; BM, bone marrow; LIV, liver; LUN, lung. \*\*\*\*P < 0.001; ns, not significant (Student's t test). Data are mean t s.d. of at least 3 independent experiments



**Fig. 5** Reduced capacity of  $β_7$ -F185A T cells to induce chronic colitis.  $Rag 1^{-/-}$  mice (n = 9 per group) were given  $1 \times 10^5 \text{ CD4}^+\text{CD45RB}^{high}$  T cells isolated from WT, KI, or KO splenic lymphocytes and monitored for 12 weeks. **a** Body weight change after T cell transfer. **b** Quantitative histopathologic grading of colitis severity. **c** Hematoxylin and eosin staining of colon sections at week 12 after T cell transfer. **d** Immunofluorescence staining and CD4<sup>+</sup> T cell quantification of colon sections with anti-CD4 (red) and DAPI (blue) as in **d**. Scale bars, 100 μm (**c**, **d**). \*\*P < 0.005, \*\*\*P < 0.001; ns, not significant (two-way ANOVA in **a**, Student's P < 0.001; ns, not are mean P < 0.001 at least 3 independent experiments (**a**, **b**)

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colitis [19, 20]. Therefore, we studied the capacity of  $\beta_7$ -F185A T cells to induce intestinal inflammation using a T cell transfer model of chronic colitis [37]. As previously described [37],  $Rag1^{-/-}$  mice transferred with 1 × 10<sup>5</sup> WT CD4<sup>+</sup>CD45RB<sup>high</sup> T cells presented progressive loss of body weight and higher colitis activity score (Fig. 5a, b) and showed clinical symptoms of severe colitis including massive infiltration of mononuclear cells in colonic lamina propria, disruption of epithelial boundaries, and disappearance of goblet cells (Fig. 5c). On the contrary, Rag1<sup>-/-</sup> mice reconstituted with the same number of KI or KO CD4+CD45RBhigh T cells appeared healthy, showed no progressive body weight loss (Fig. 5a, b), and maintained normal colonic mucosal architecture (Fig. 5c). Consistent with the above results, markedly fewer KI and KO CD4+ T cells were present in the colonic lamina propria compared with WT CD4<sup>+</sup> T cells (Fig. 5d). Thus, blocking  $\alpha_4\beta_7$  activation is equivalent to fully inhibiting  $\alpha_4\beta_7$  function with regard to suppressing the recruitment of inflammatory T cell to the gut and subsequently eliminating adaptive immune-mediated chronic colitis.

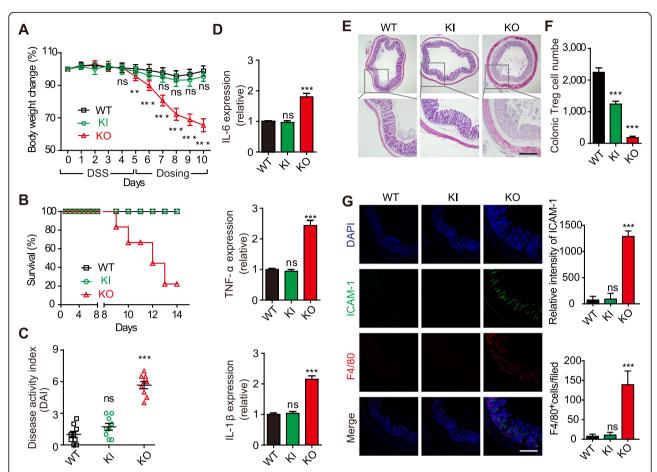
# $\beta_{7}\text{-F185A KI}$ mice are resistant to DSS-induced acute colitis

In addition to adaptive immunity, innate immunity also contributes to the pathogenesis of IBD [38]. In both IBD patients and the DSS-induced colitis model, macrophages were considered a critical factor involved in disease progression [39, 40]. Moreover, total loss of  $\alpha_4\beta_7$ function has been shown to lead to colonic Treg depletion, which promotes ICAM-1 expression and macrophage infiltration into the colon and exacerbates DSSinduced colitis [24]. Therefore, we next evaluated the effect of blocking  $\alpha_4\beta_7$  activation on the susceptibility to innate immune-mediated acute colitis using the DSS colitis model [41]. Instead of using high concentration of DSS (3.5-5%) to induce severe acute colitis [42, 43], mice were treated with low concentration of DSS (2%) to induce impairment of mucosal barrier function in the gut and moderate acute colitis [24, 39, 44]. Compared with the rapid body weight loss of DSS-treated  $\beta_7$  KO mice, DSS-treated WT and KI mice showed similar mild body weight changes throughout the observation period (Fig. 6a). Notably, WT and KI mice exhibited a 100% survival rate at day 14 after initial DSS treatment compared with the 20% survival rate of KO mice (Fig. 6b). In contrast to the severe clinical symptoms of colitis found in KO mice, WT and KI mice developed more mild clinical symptoms of colitis, including lower colitis activity score (Fig. 6c), decreased colonic expression of proinflammatory cytokines (IL-6, TNF- $\alpha$ , and IL-1 $\beta$ ) (Fig. 6d), fewer inflammatory infiltrates, and moderate disruption of mucosal structures (Fig. 6e). Of note, KI mice retained around 55% of Treg cells in the colonic lamina propria compared with the approximately 92% depletion of colonic Treg cells in KO mice (Fig. 6f). In contrast to the increased ICAM-1 expression and macrophage infiltration in the colon of KO mice, KI mice exhibited no visible ICAM-1 expression or only mild macrophage infiltration into the colon (Fig. 6g). Because the colonic Treg cells have been shown to inhibit ICAM-1 expression and macrophage infiltration in the colon in DSS colitis [24], it is tempting to speculate the retained colonic Treg cells in KI mice may be related to the susceptibility to DSS-induced innate colitis.

Taken together, specific blockade of integrin  $\alpha_4\beta_7$  activation is sufficient to prevent adaptive immunemediated colitis without increasing susceptibility to innate colitis, suggesting the efficacy of this approach in treating IBD. Of note, specifically blocking  $\alpha_4\beta_7$  activation retains colonic Treg cells, thus appears to be advantageous over fully blocking  $\alpha_4\beta_7$  function by avoiding aberrant innate immune response. Therefore, the targeting of integrin  $\alpha_4\beta_7$  activation for the treatment of IBD has the potential to induce fewer adverse effects than complete blockade of  $\alpha_4\beta_7$  function.

### **Discussion**

Integrin  $\alpha_4\beta_7$  mediates rolling adhesion and firm adhesion of lymphocytes pre- and post-activation respectively [22, 26], controlling two critical steps of lymphocyte homing to the gut [27, 28]. Blocking the function of  $\alpha_4\beta_7$ efficiently suppresses the recruitment of inflammatory lymphocytes to the gut and consequently inhibits adaptive immune-mediated colitis [19, 45]. The US Food and Drug Administration has approved the integrin  $\alpha_4\beta_7$ blocking antibody vedolizumab for the treatment of adults with moderately to severely active UC or CD. Clinical studies report that vedolizumab effectively maintains clinical remission in some IBD patients; however, aggravated colitis is observed in a small percentage of UC patients treated with high dose of vedolizumab [20]. One possible reason could be that the excessive loss of  $\alpha_4\beta_7$  function leads to colonic Treg depletion, which exacerbates colitis by evoking aberrant innate immune response in a DSS colitis model [24]. Indeed, a study has reported that vedolizumab treatment significantly suppresses Treg cells homing to the gut in UC patients [25]. Instead of completely blocking the function of integrin, an appropriate modulation of integrin function might be required not only to suppress aberrant immune response-induced inflammation but also to maintain immune homeostasis [46, 47]. In this study, we demonstrate that specific inhibition of integrin  $\alpha_4\beta_7$  activation is sufficient to prevent adaptive immunemediated colitis without increasing susceptibility to innate colitis, suggesting a potentially better treatment Zhang et al. BMC Biology (2020) 18:64 Page 9 of 15



**Fig. 6** β<sub>7</sub>-F185A KI mice are resistant to DSS-induced acute colitis. Eight-week-old WT, KI, and KO mice were treated with low concentration of DSS (2%) for 5 days, followed by regular drinking water (dosing). **a–c** Body weight change (**a**), survival ratio (**b**), and disease activity index (**c**) were obtained from individual groups of mice (n = 6 per group). **d** Quantitative PCR analysis of IL-6, TNF-α, and IL-1β expression in distal colon tissue from individual groups of mice (n = 6 per group) at day 10 after DSS treatment. Results are normalized to *GAPDH*. **e** Hematoxylin and eosin staining of distal colon sections at day 10 after DSS treatment. **f** Flow cytometric analysis of Treg cell number in colonic tissues from individual groups of mice (n = 6 per group) at day 4 after DSS treatment. **g** Immunofluorescence staining and quantification of F4/80<sup>+</sup> macrophages and ICAM-1 expression in distal colon sections with anti-ICAM-1 (green), anti-F4/80 (red), and DAPI (blue) at day 4 after DSS treatment. Scale bars, 100 μm (**e**, **g**). \*\*P < 0.005; \*\*\*P < 0.001; ns, not significant (two-way ANOVA in **a**, Student's t test in **c**, **d**, **f**, and **g**). Data are mean  $\pm$  s.d. of at least 3 independent experiments (**a**, **c**, **d**, **f**, **g**)

for IBD than complete blockade of  $\alpha_4\beta_7$  function. It provides novel mechanism of action for the development of new IBD drugs.

Leukocyte recruitment to the gut is dynamically regulated by adhesion cascades including cell capture, slow rolling, adhesion strengthening, intravascular crawling, and transcellular transmigration [48]. Previous in vitro studies have shown that resting  $\alpha_4\beta_7$  mediates robust cell rolling on immobilized MAdCAM-1 substrates and firm cell adhesion upon activation [26, 30]. Full blockade of  $\alpha_4\beta_7$  function disables the ability of this integrin to mediate both rolling and firm cell adhesion, whereas specifically blocking  $\alpha_4\beta_7$  activation only partially inhibits  $\alpha_4\beta_7$ -mediated cell adhesion, which still allows this integrin to support cell rolling [30]. Consistent with this, our in vitro data showed

that there were significantly more  $\beta_7$ -F185A KI cells than  $\beta_7$  KO cells that adhered to (mostly rolling events) and transmigrated through MAdCAM-1 substrates (Fig. 3b, c). Interestingly, the in vivo shortterm competitive homing assay showed that there was no significant difference between the suppression of recruitment of KI and KO lymphocytes to the GALT (Fig. 4), which is consistent with the data that CD4+CD45RBhigh T cells from KI and KO mice are similarly impaired in their ability to induce colitis in a T cell transfer model. Thus, in a manner similar to full inhibition of  $\alpha_4\beta_7$  function, blocking  $\alpha_4\beta_7$ -mediated firm cell adhesion through specific inhibition of  $\alpha_4\beta_7$  activation suppresses lymphocyte homing into the gut. This indicates both approaches are capable of inhibiting adaptive immune-mediated colitis.

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As shown by clinical studies that high doses of etrolizumab reduce the remission rate of IBD patients, it is possible that homing of Treg cells is reduced due to the excessive inhibition of  $\beta_7$  function, as is the KO phenotype. It is noteworthy that  $\beta_7$  KO mice did show a greater decrease in CD3+ T cells in the gut compared with  $\beta_7$ -F185A KI mice (Fig. 2e). Importantly, there were significantly more colonic Treg cells in β<sub>7</sub>-F185A KI mice than in KO mice after DSS treatment, suggesting the  $\alpha_4\beta_7$ -mediated rolling adhesion may contribute to the maintenance of the colonic Treg cell population. Although the number of colonic Treg cells in β<sub>7</sub>-F185A KI mice was less than that in WT mice, there appear to be enough colonic Treg cells present to be resistant to the aberrant innate immune response that exacerbates gut inflammation in the DSS colitis model (Fig. 6). Thus, specifically blocking  $\alpha_4\beta_7$  activation avoids excessive depletion of colonic Treg cells, which is important for suppressing aberrant innate immune responses under certain conditions.

In addition to integrin  $\alpha_4\beta_7$ , the  $\beta_7$  subunit also forms a heterodimer with the  $\alpha_E$  subunit. Through an interaction with E-cadherin,  $\alpha_E \beta_7$  mediates cell-to-cell interactions between T cells and intestinal epithelial cells, thereby playing an important role in the retention of lymphocytes in mucosal tissue [49]. Integrin  $\alpha_E$  KO mice display a reduction in the number of lymphocytes residing in the intestinal epithelium and lamina propria [35]. The reduced expression of integrin  $\alpha_E$  mainly impairs the retention of lymphocytes in the gut, but has little effect on the gut homing of lymphocytes [50, 51]. In this study, we did observe significantly reduced expression of  $\alpha_E \beta_7$  in KI lymphocytes, the depletion of the  $\alpha_E \beta_7^+$  lymphocytes in the spleen, and significantly reduced the population of  $\alpha_E \beta_7^+$  IEL and LPL in the gut of KI mice. β<sub>7</sub>-F185A mutation did not inhibit the chemokineinduced activation of  $\alpha_E \beta_7$ , but downregulated the expression of  $\alpha_E\beta_7$  in lymphocytes, thereby impairing lymphocyte retention in the gut.

It is also noteworthy that a small population of  $\beta_7$ -high splenic lymphocytes disappeared in KI mice (Fig. 3a). Further analysis showed that this group of cells mainly consisted of CD8<sup>+</sup> T cells (79.05%), CD4<sup>+</sup> T cells (8.74%), CD4<sup>+</sup>CD25<sup>+</sup> Treg cells (1.42%), and CD19<sup>+</sup> B cells (3.90%) (Additional file 2: Table S1). Whether the depletion of this  $\beta_7$ -high lymphocyte population is related to colitis needs to be further studied.

Etrolizumab is an anti- $\beta_7$  monoclonal antibody that is able to inhibit the functions of both  $\alpha_4\beta_7$  and  $\alpha_E\beta_7$  integrins, whereas vedolizumab is an anti- $\alpha_4\beta_7$  antibody which only inhibits the function of  $\alpha_4\beta_7$ . It is reported that vedolizumab is more effective in UC than in CD [52]. One potential explanation is that UC is related to  $T_H2$  and  $T_H17$  cells, which mainly express  $\alpha_4\beta_7$ . By

contrast, etrolizumab is more effective in inflammatory environments rich in CD69<sup>+</sup> $\alpha_E$  <sup>+</sup> intestinal  $T_{RM}$  cells and  $T_{H}9$  cells because these cells express high level of  $\alpha_E\beta_7$  [14, 15]. In addition, a phase II study reports that patients with elevated intestinal expression of  $\alpha_E\beta_7$  may have an increased chance of clinical remission in response to etrolizumab treatment [53]. Thus, a personalized selection of vedolizumab or etrolizumab for the treatment of different IBD patients might benefit the clinical remission. It is noteworthy that blocking  $\beta_7$  activation leads to the loss of  $\alpha_E\beta_7$  expression in lymphocytes as shown in this study, which should be able to suppress the functions of both  $\alpha_E\beta_7$  and  $\alpha_4\beta_7$  integrins.

### **Conclusions**

Collectively,  $\beta_7\text{-F}185A$  mutation not only inhibited integrin  $\alpha_4\beta_7$  activation thus reducing the  $\alpha_4\beta_7\text{-mediated}$  homing of lymphocytes into the gut, but also induced the downregulation of integrin  $\alpha_E\beta_7$  expression in lymphocytes and impaired the retention of lymphocytes in the gut. Importantly, rather than completely inhibiting  $\beta_7$  function, inhibiting the activation of  $\alpha_4\beta_7$  could retain enough Treg cells in the gut, thus could prevent adaptive immune-mediated colitis without increasing the susceptibility to innate colitis, suggesting this is potentially a better strategy than fully blocking  $\beta_7$  function for IBD treatment. Therefore, the development of antibodies and small molecule antagonists that specifically block  $\alpha_4\beta_7$  activation may provide a new approach for clinical management of IBD.

### Methods

### Generation of β<sub>7</sub>-F185A KI mice

For the construction of Itgb7 gene-targeting vector, a 2.9kb SacII/NotI fragment containing exon 3, 4 and a 1.3-kb ClaI/SaII fragment containing exon 5, 6 were subcloned into the upstream regions and downstream of the loxPflanked neo gene, respectively. F185A was introduced in exon 5 by site-directed mutagenesis. The exonic sequences in the targeting vector were confirmed by DNA sequencing. The targeting vector was transfected into ES cells of the 129/SvJ mouse strain. G418-resistant ES cell clones were screened for homologous recombination. The correctly targeted ES clones were injected into C57BL/6J blastocysts, and the resulting chimeric males were mated to C57BL/6J females for germline transmission. The neo cassette was deleted after crossed with EIIa-cre mice. The offspring were screened for heterozygosity by PCR (Primers, Additional file 2: Table S2) to confirm in vivo deletion of the neo cassette. The mice that were backcrossed into the C57BL/6J background for ten generations were used in all experiments.

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### Mice

Itgb7<sup>-/-</sup> mice, Rag1<sup>-/-</sup> mice, and EIIa-cre mice with C57BL/6J background were from Jackson Laboratory (Bar Harbor, ME). All mice were maintained under specific pathogen-free conditions. Age- and sex-matched mice were used at 8–10 weeks of age. All animal studies were approved by the Institutional Animal Care and Use Committee of the CAS Center for Excellence in Molecular Cell Science, Shanghai Institute of Biochemistry and Cell Biology, Chinese Academy of Sciences.

### Antibodies and reagents

Monoclonal antibodies (mAbs) specific for mouse CD3 (17A2), CD4 (GK1.5), CD8 (53-6.7), CD19 (1D3), CD45RB (C303.16A), and integrin  $\alpha_4$  (9C10) were all from BD Bioscience (San Jose, CA); mAb to mouse CD3 (17A2) was from Invitrogen (Carlsbad, CA); mAb to human/mouse E-Cadherin (24E10) was from Cell Signaling; mAbs to mouse F4/80 (BM8), CD25 (PC61.5), and Foxp3 (FJK-16s) were from eBioscience (San Diego, CA); mAb to mouse ICAM-1 (M-19) was from Santa Cruz (Santa Cruz, CA); mAb to integrin  $\beta_7$  (N1N3) was from GeneTex (Irvine, CA); rat mAb FIB504 against human/ mouse  $\beta_7$  and rat mAb M290 against human/mouse  $\alpha_E$ were prepared by using hybridomas (Developmental Studies Hybridoma Bank, University of Iowa); mAbs to mouse CD19 (1D3), human CCR9 (L053E8), FITC goat anti-rat IgG, and FITC goat anti-mouse IgG were from BioLegend; 4,6-diamidino-2-phenylindole (DAPI) was from Sigma (St. Louis, MO); CellTrace Violet Cell Proliferation Kit and CellTrace Yellow Cell Proliferation Kit were from Thermo Fisher Scientific (Waltham, MA). Human and mouse chemokines CCL21 and CCL25 were purchased from R&D Systems (Minneapolis, MN). Recombinant mouse extracellular maturation peptide Ecadherin (Asp157-Val709) fused with Fc tag were expressed in 293T cells and purified by protein A (Pierce) affinity chromatography.

### **Cell lines**

Jurkat T cell line expressing WT or mutant human  $\alpha_4\beta_7$  was generated by infection with the recombinant lentivirus, and human CCR9 was infected at the same time. Similarly, Jurkat T cell line expressing WT or mutant mouse  $\alpha_E\beta_7$  was generated by infection with the recombinant lentivirus. Integrins and CCR9 expressing efficiency were assessed by flow cytometry 72 h after transfection using a FACS Celesta flow cytometer (BD Biosciences).

### Mononuclear cell isolation and flow cytometry

Lymphocyte cells were isolated from the spleen, thymus, PB, BM, PLN, MLN, PP, and colon as previously described [25, 54–56]. Briefly, lymphocytes were isolated

from the spleen, thymus, PLN, MLN, and PP by dispersion through a fine wire mesh. To harvest the IEL and LPL, the intestinal tube was opened laterally and cut into pieces measuring 0.5 cm. Intestinal pieces were incubated for 3 or 5 consecutive times in HBSS/EDTA (1 mmol/L DTT) for 20 min at 37 °C with stirring. For the isolation of LPL, tissue fragments were further subjected to 2 rounds of treatment in RPMI-1640 with 5% FBS, collagenase D, DNase I, and dispase for 30 min at 37 °C. The resulting suspensions were layered on a discontinuous 40%/70% Percoll gradient, centrifuged at 650g for 20 min, and IEL and LPL were recovered from the interphase. Aliquots of  $1 \times 10^6$  viable cells were stained with 50 µl of optimally titrated antibody cocktail in PBS containing 5% FBS. For intracellular staining of Foxp3, cells were fixed and permeabilized using the mouse Treg cell staining kit (eBioscience) before staining with anti-Foxp3 mAb. Cells were maintained on ice throughout the staining procedure. Samples were analyzed by using the FACS Celesta flow cytometer (BD Biosciences). All viable lymphocytes were gated using a combination of forward angle and side scatter to exclude dead cells and debris. Acquisition and analysis were performed on FlowJo (LLC, Ashland, Oregon, USA).

### Histology and immunofluorescence microscopy

Formalin-fixed, paraffin-embedded small intestine and distal colon sections of 4- $\mu$ m thickness were mounted on glass slides and followed by hematoxylin and eosin staining. For immunostaining analysis, distal colons were collected at individual days after the initial DSS treatment or T cell transfer. Cryostat sections were made permeable with cold acetone and blocked with 1% (wt/vol) BSA. Samples were incubated with fluorochrome-conjugated anti-F4/80 (5  $\mu$ g/ml), anti-ICAM-1 (10  $\mu$ g/ml), or anti-CD4 (5  $\mu$ g/ml), and counterstaining of nuclei was with DAPI (1  $\mu$ g/ml). Images were acquired with a Leica TCS SP8 confocal microscope (Leica, Mannheim, Germany).

### RNA isolation and real-time quantitative PCR

Total RNA was extracted from mouse splenic lymphocytes or distal colonic tissues with TRIzol reagent according to the manufacturer's instructions (Invitrogen, Carlsbad, CA). For cDNA synthesis, RNA was reverse-transcribed with an M-MLV reverse transcriptase (Promega, Madison, WI). Then, cDNA was amplified by real-time PCR (primers, Additional file 2: Table S3) with a SYBR Premix ExTaq kit (TaKaRa, Otsu, Japan) on an AbiPrism 7500 sequence detector (Applied Biosystems, Foster City, CA). The expression of target genes was normalized to the expression of the housekeeping gene GAPDH.

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### Silencing of β<sub>7</sub> in splenic lymphocytes

Silencing of  $\beta_7$  expression in WT mouse splenic lymphocytes was achieved by shRNA. Cells with  $\beta_7$  were generated by infection with the recombinant lentivirus, which harbors packaging plasmid psPAX2 (7.5 µg), envelope plasmid pMD2.G (3 µg), and transfer plasmid pLKO.1-shItgb7 (10 µg). The oligonucleotide targeting sequence for mouse  $\beta_7$  was 5′-CCCGTCTTCTAGTGTTCACTT-3′. Knockdown of  $\beta_7$  was confirmed by flow cytometry 72 h after transfection using a FACS Celesta flow cytometer (BD Biosciences).

### Flow chamber assay

Cell adhesion under physiologic shear stress was studied as previously described [26]. Briefly, a polystyrene petri dish was coated with 20 µl of MAdCAM-1/Fc (10 µg/ml) alone or with chemokines (2 µg/ml) in coating buffer (PBS, 10 mM NaHCO<sub>3</sub>, pH 9.0) for 1 h at 37 °C followed by blocking with 2% BSA in coating buffer for 1 h at 37 °C. Cells were diluted to  $1 \times 10^6$ cells/ml in HBSS (10 mM HEPES, 1 mM Ca<sup>2+</sup>/Mg<sup>2+</sup>) and immediately infused in the flow chamber using a syringe pump through the flow chamber at a constant flow of 1 dyn/cm<sup>2</sup> or 2 dyn/cm<sup>2</sup>. All adhesive interactions between the flowing cells and the coated substrates were determined by manually tracking the motions of individual cells as previously described [57]. The motion of each adherent cell was monitored for 10 s following the initial adhesion point, and two categories of cell adhesion were defined. Adhesion was defined as rolling adhesion if the adherent cells were followed by rolling motions  $\geq 5$  s with a velocity of at least 1 μm/s for splenic lymphocytes or 2 μm/s for Jurkat T cells due to the larger cell size, whereas a firmly adherent cell was defined as a cell that remained adherent and stationary for at least 10 s. For integrin  $\alpha_E \beta_7$ -mediated adhesion, a polystyrene petri dish was coated with 20 µl of mouse E-cadherin-Fc (40 µg/ml) alone or with chemokines (2 µg/ml) in coating buffer. Cells were diluted to  $1 \times 10^6$  cells/ml in HBSS and immediately infused in the flow chamber using a syringe pump through the flow chamber at a constant flow of 1 dyn/cm<sup>2</sup>. Then, cells were allowed to adhere for 2 min, and the number of adherent cells was determined.

### Transwell migration assay

Transwell migration was performed using Millicell inserts with 5- $\mu$ m pore size for splenic lymphocytes and 8- $\mu$ m pore size for Jurkat T cells (Millipore, Billerica, MA). Ten micrograms per milliliter of MAdCAM-1/Fc or 10  $\mu$ g/ml ICAM-1/Fc was coated on the upper surface of inserts, and the lower chamber was filled with 1 ml RPMI 1640 medium with 10% FBS.  $2 \times 10^5$  cells in 0.2-

ml serum-free RPMI 1640 medium were added into the upper chamber. Cells were incubated for 6 h at 37  $^{\circ}$ C in 5% CO<sub>2</sub>. Cells remaining on the upper surface of the inserts were scraped with a cotton swab, and cells migrating to the bottom surface were counted after fixation with 3.7% formaldehyde and staining with DAPI.

### Competitive in vivo homing assay

A competitive homing assay was conducted as previously described [36].  $2\times10^7$  splenic lymphocytes from heterozygote (+/-), KI, or KO mice labeled with Cell-Trace Yellow were mixed with the same number of WT cells labeled with CellTrace Violet and injected intravenously into C57BL/6J mice. An aliquot was saved to assess the input ratio (calculated as [Yellow<sup>+</sup>][13, 15]<sub>in-put</sub>/[Violet<sup>+</sup>]<sub>input</sub>). Recipient mice were scarified 18 h after injection, and lymphocytes from tissues were harvested. The homing index was calculated as the [Yellow<sup>+</sup>]<sub>tissue</sub>/[Violet<sup>+</sup>]<sub>tissue</sub> ratio to the input ratio.

### T cell transfer colitis

CD4<sup>+</sup> CD45RB<sup>high</sup> naive T cells were isolated from mouse splenic lymphocytes using LSRII (BD Bioscience) cell sorting with FITC-conjugated anti-CD45RB and PEconjugated anti-CD4 mAbs.  $1\times10^5$  WT, KI, or KO CD4<sup>+</sup>CD45RB<sup>high</sup> cells in 0.2 ml PBS were injected intravenously into  $Rag1^{-/-}$  recipient mice, respectively. Mice were weighted weekly and observed for signs of illness as reported previously [58]. At week 12, mice were scarified, and histological tissue was taken and analyzed.

### **DSS-induced acute colitis**

Acute colitis was induced by the provision of 2% (wt/ vol) DSS with molecular mass of 36-50 kDa (MP Biomedicals, Irvine, CA) in drinking water for a total of 5 days (days 0-5), followed by regulatory drinking water (days 6-15). Mice were assessed daily for body weight, diarrhea, and bloody stool. The disease activity index (DAI) and histological damage were assessed by trained individuals blinded to the treatment groups, as reported previously [58, 59]. DAI was calculated as the combined score of stool consistency (0, normal; 1, moist/sticky; 2, soft; 3, diarrhea), presence of blood in the stool (0, no blood; 1, blood in the stool or around the anus; 2, for severe bleeding), and mouse appearance (0, normal; 1, ruffled fur or altered gait; 2, lethargic or moribund). The severity of colitis was assessed on stained colonic sections in a blinded fashion using established criteria based on crypt damage and ulceration. Crypt damage was scored as follows: 0, intact crypts; 1, loss of the basal one-third; 2, loss of the basal two-thirds; 3, entire crypt loss; 4, change of epithelial surface with erosion; and 5, confluent erosion. Ulceration was scored as follows: 0, Zhang et al. BMC Biology (2020) 18:64 Page 13 of 15

absence of ulcer; 1, one or two foci of ulceration; 2, two to four foci of ulcerations; and 3, confluent or extensive ulceration. On day 10, mice were sacrificed and colons removed and analyzed. For interim analyses, experiments were done as described above but terminated on day 4. Colon tissues were used for histological and quantitative PCR analysis.

### Statistical analysis

Statistical analyses were performed with GraphPad PRISM software 5.0 (GraphPad Software, La Jolla, CA). Significances were determined by two-tailed Student's t test or two-way ANOVA as indicated. \*P<0.01, \*\*P<0.005, and \*\*\*P<0.001 were considered statistically significant in all figures.

### **Supplementary information**

**Supplementary information** accompanies this paper at https://doi.org/10. 1186/s12915-020-00784-6.

Additional file 1: Figure S1. The secondary lymphoid tissues other than GALT appear normal in  $\beta$ 7-F185A KI and  $\beta$ 7-KO mice. Representative histological sections of the peripheral lymph node (PLN), mesenteric lymph node (MLN) and spleen (SP) of WT,  $\beta$ 7-F185A KI (KI) and  $\beta$ 7-KO (KO) mice were analyzed by hematoxylin and eosin staining. Scale bars, 500μm. Figure S2. Expression of integrins β7 in splenic lymphocytes of WT and β7-F185A mice. (A) Quantitative PCR analysis of integrin β7 expression in splenic lymphocytes from WT and KI mice. Results are normalized to GAPDH. (B) Total (cell surface plus intracellular) protein expression of integrin  $\beta 7$  in splenic lymphocytes was determined by flow cytometry using permeabilized cells. Data are mean  $\pm$  s.d. of at least 3 independent experiments (A-B). Figure S3. Impaired adhesion and transmigration of Jurkat T-β7 F185A cells. (A) Expession of β7 and CCR9 in Jurkat T-β7 WT and Jurkat T-β7 F185A cell lines were determined by flow cytometry. (B) Adhesion of Jurkat T-β7 WT, Jurkat T-β7 F185A and Jurkat T cells to MAdCAM-1 substrates at 1 dyn/cm<sup>2</sup> or 2 dyn/cm<sup>2</sup> before and after chemokine stimulation. (C) Transmigration of Jurkat T-β7 WT, Jurkat T-F185A and Jurkat T cells through MAdCAM-1-coated insert. Data are mean  $\pm$  s.d. of at least 3 independent experiments (**B C**). \*\*\* P < 0.001; ns, not significant (Student's t-test). Asterisk in **B** indicates the changes of total adherent cells. Figure S4.  $\beta$ 7-F185A mutation does not affect  $\alpha$ E $\beta$ 7-mediated cell adhesion to E-cadherin substrates. (A) Expession of  $\beta7$  and  $\alpha E$  in Jurkat T-αΕβ7 WT and Jurkat T-αΕβ7 F185A cell lines were determined by flow cytometry. The numbers within the panels show the specific mean fluorescence intensities of FIB504 (anti- $\beta$ 7) and M290 (anti- $\alpha$ E) mAbs. (**B**) Adhesion of Jurkat T-αΕβ7 WT, Jurkat T-αΕβ7 F185A and Jurkat T cells to the immobilized E-cadherin substrates (40 µg/ml) at 1dyn/cm<sup>2</sup> before and after chemokine stimulation. αΕβ7-E-cadherin binding was inhibited by pre-treatment of cells with 10  $\mu$ g/ml  $\alpha$ E blocking antibody M290. Data are mean  $\pm$  s.d. of at least 3 independent experiments (**A-B**).\*\*\* P <0.001; ns, not significant (Student's t-test). Figure S5. Integrin αE<sup>+</sup> lymphocytes in spleen, SI and colon. (A) Expression of  $\alpha E$  in WT, +/-, KI and KO splenic lymphocytes was determined by flow cytometry. (**B-C**) Expression of  $\alpha E$  in intestinal IEL and LPL was detected by flow cytometry. The numbers within the panels (**A-B**) show the percentage of  $\alpha E^+$  lymphocytes. The numbers within the table (C) show the specific mean fluorescence intensities of M290 (anti-αE) mAb in αE<sup>+</sup> lymphocytes. Data are mean  $\pm$  s.d. of at least 3 independent experiments (A-C). ^AAP < 0.05; ^AAAP < 0.001 (Student's t-test in C). SP, spleen; SI, small intestine; IEL, intraepithelial lymphocyte; LPL, lamina propria lymphocyte.

**Additional file 2: Table S1.** The analysis of integrin  $\beta$ 7-high splenic lymphocytes subsets in WT mice. **Table S2.** The list of primers used for genotyping. **Table S3.** The list of primers used for real-time quantitative PCR analyses

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### Authors' contributions

HLZ, YJZ, and JFC designed the experiments; HLZ, YJZ, YDP, CDL, SHW, ZJY, and LL performed the experiments and analyzed the data; HLZ, YJZ, YDP, and JFC interpreted the results; JFC, GXG, JSL, and YAZ supervised the study; the manuscript was drafted by HLZ, YJZ, and CDL and edited by JFC. The authors read and approved the final manuscript.

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### Availability of data and materials

All data generated or analyzed during this study are included in this published article and its additional information files.

### Ethics approval and consent to participate

The research of mouse experiments in the manuscript has been conducted under the guidance of international ethical standards. All research protocols were conducted with the approval of the Institutional Animal Care and Use Committee of the Shanghai Institute of Biochemistry and Cell Biology, Center for Excellence in Molecular Cell Science.

### Consent for publication

Not applicable.

### Competing interests

The authors declare that they have no competing interests.

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