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Decidual RANKL/RANK interaction promotes the residence and polarization of TGF- β 1-producing regulatory $\gamma\delta$ T cells

Rui-Qi Chang¹, Jun Shao², Yu-Han Meng³, Jian Wang¹, Da-Jin Li¹ and Ming-Qing Li^{1,4} 

Abstract

Decidual $\gamma\delta$ T (d $\gamma\delta$ T) cells play an essential role during successful pregnancy; however, the residence and polarization of $\gamma\delta$ T cells in decidua remain unclear. In this study, we observed higher levels of receptor activator for nuclear factor- κ B ligand (RANKL) on decidual stromal cells (DSCs), and its receptor RANK on d $\gamma\delta$ T cells in decidua from normal pregnancy compared with patients with recurrent spontaneous abortion (RSA). RANKL expressed by DSCs can induce the polarization of peripheral blood $\gamma\delta$ T (p $\gamma\delta$ T) and d $\gamma\delta$ T cells to Foxp3 + $\gamma\delta$ T cells, and upregulate the expression of transforming growth factor (TGF)- β 1. This process is mediated through activation of nuclear factor kappa-light-chain-enhancer of activated B cells (NF- κ B). In addition, RANKL promotes the adhesion of d $\gamma\delta$ T cells to DSCs in vitro, which is associated with the upregulation of ICAM-1 and VCAM-1 on DSCs and integrins on d $\gamma\delta$ T cells. RANKL knockout leads to the decreased numbers of uterus total $\gamma\delta$ T cells, Foxp3+ $\gamma\delta$ T cells and the expression of TGF- β 1, and the increased pregnancy loss in mice. These results suggest that RANKL is a pivotal regulator of maternal-fetal tolerance by triggering the polarization and residence of TGF- β 1-producing Foxp3+ $\gamma\delta$ T cells in early pregnancy. The abnormal low level of RANKL/RANK results in pregnancy loss because of the dialogue disorder between DSCs and d $\gamma\delta$ T cells. This observation provides a scientific basis on which a potential marker can be detected to early warning of pregnancy loss.

Introduction

Decidual immune cell (DIC), one of the major components at the maternal-fetal interface, is critical in the induction of maternal immune tolerance to fetal alloantigen during pregnancy^{1–3}. Abnormality of DIC is related to several pathological pregnancies, including recurrent spontaneous abortion (RSA), unexplained infertility, preeclampsia, and intrauterine growth restriction (IUGR)^{4,5}. Decidual $\gamma\delta$ T (d $\gamma\delta$ T) cells, accounted for over 60% of T cells in human decidua, participate in maintenance of pregnancy by recognizing

alloantigen without MHC restriction, producing cytokines and linking the innate and adaptive immune responses as a bridge^{6–8}.

Similar to CD4 helper T (Th) cells, $\gamma\delta$ T cells can be polarized toward six distinct subgroups upon activation based on their functional and developmental features^{9,10}. $\gamma\delta$ 1, 2, 17, 22, follicular helper (FH), and regulatory (reg) cells are characterized by its capacity to produce interferon (IFN)- γ , interleukin (IL)-4, IL-17, IL-22, Th2-cell-associated cytokines (including IL-4 and IL-10), and transforming growth factor (TGF)- β , respectively. Moreover, T-bet, GATA-binding protein 3 (GATA3), RORC, Bcl-6, and Foxp3 are the master transcription factors for the polarization of $\gamma\delta$ 1, 2, 17, FH, and reg, respectively^{11–15}. Accumulating evidence showed that d $\gamma\delta$ T cells have a tendency to secrete immunosuppressive cytokines, especially TGF- β and IL-10 at maternal-fetal interface^{7,16,17}. These results implicate that the polarization of d $\gamma\delta$ T cells may play an important role in regulation of immune response at the

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maternal-fetal interface. However, the related mechanism remains unclear.

Receptor activator for nuclear factor- κ B (RANK) and its only known ligand tumor necrosis factor ligand superfamily member 11 (TNFSF11, also known as RANKL) have dual roles in immune regulation. On the one hand, they promote adaptive immune response by inducing the production of IL-12 in mature dendritic cells and polarization of CD4⁺ T cells into Th1 cells¹⁸. On the other hand, they exert their immunosuppression through inducing the polarization of regulatory T cells and participating in the establishment of central as well as peripheral tolerance¹⁹. In our previous studies, RANKL/RANK has been identified and functionally described at the maternal-fetal interface where it involved in the maintenance of pregnancy by promoting the growth of decidual stromal cells (DSCs) and inducing decidual M2 macrophage polarization^{20,21}. However, to date there have no studies about the effects of RANKL/RANK interaction on $\text{d}\gamma\delta$ T cells.

In this article, we focus on the interaction between DSCs-derived RANKL and RANK expressed on $\text{d}\gamma\delta$ T cells and reveal their role in the maintenance of early pregnancy and RSA.

Results

The abnormal low level of RANKL/RANK at the maternal-fetal interface in RSA patients

To investigate the interaction between DSC-derived RANKL and RANK expressed on $\text{d}\gamma\delta$ T, we first analyzed the expression of RANKL and RANK in decidua during early pregnancy. As shown, the strong positive staining of RANKL and RANK located in the cytoplasm and cell membrane of DSCs was observed by immunohistochemistry (Fig. 1a). RANKL and RANK expression in decidua from normal pregnancy were significantly higher than that in control endometrium from non-pregnant women (Fig. 1a). Further analysis showed that DSCs from normal pregnancy had a higher level of membrane RANK (Fig. 1b, c). Flow cytometry analysis revealed high levels of RANK expression on $\text{d}\gamma\delta$ T cells, as the percentage of RANK⁺ $\gamma\delta$ T cells (CD45⁺CD3⁺ $\gamma\delta$ TCR⁺) was over 90% at the maternal-fetal interface, while less than 10% of peripheral blood (Fig. 1d, e). The tissue-specific high expression level of RANK on $\text{d}\gamma\delta$ T suggests the possible role of RANK in the regulation of $\text{d}\gamma\delta$ T and maternal-fetal immunotolerance.

Meanwhile, we investigated the relationship between RANKL/RANK signaling in decidua and the occurrence of RSA. Interestingly, by comparison with normal pregnant women, RSA patients showed weaker levels in decidua for the RANKL and RANK staining (Fig. 1a) and membrane RANKL expression (Fig. 1b, c). In accordance with this, a significant decrease in the percentage of RANK⁺ $\gamma\delta$ T cells was observed in patients with RSA

(53.79% \pm 9.647) in first trimester compared with normal pregnancy (92.65% \pm 1.517) (Fig. 1d, e). These findings indicate that the suppression of RANKL/RANK signaling at the maternal-fetal interface may be related to RSA.

DSCs upregulate the expression of RANK on $\text{d}\gamma\delta$ T cells

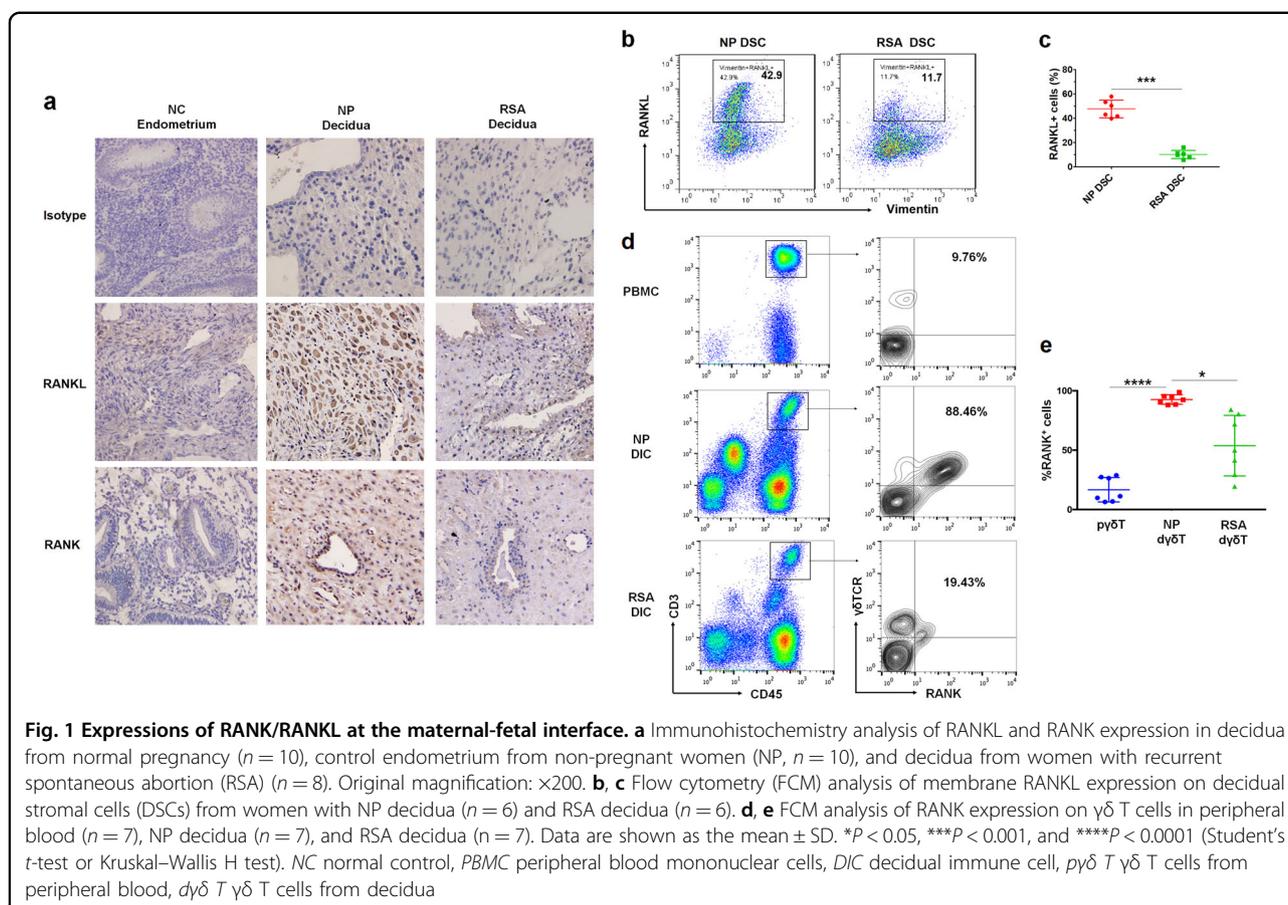
To investigate the potential relationship between high level of RANKL/RANK, and the interaction of DSCs and $\text{d}\gamma\delta$ T cells at the maternal-fetal interface, we constructed the co-culture model with DSCs and $\text{d}\gamma\delta$ T cells for imitating local microenvironment of decidua during early pregnancy. FCM analysis revealed markedly increased expression of RANKL in DSCs (Fig. 2a, b) and RANK on $\text{d}\gamma\delta$ T cells (Fig. 2c, d) in the co-culture unit. After culture alone for 48 h, the percentage of RANKL⁺ DSCs were reduced to 8.62% and increased to about two times with the presence of $\text{d}\gamma\delta$ T cells (Fig. 2a, b). Meanwhile, the presence of DSCs changed the frequency of RANK⁺ $\text{d}\gamma\delta$ T cells from 28.60 to 45.63% after culture for 48 h (Fig. 2c, d). Taken together, the interaction of DSCs and $\text{d}\gamma\delta$ T cells maintains the high levels of RANKL/RANK at the maternal-fetal interface.

RANK⁺ $\text{d}\gamma\delta$ T cells present the activated phenotype

Our further phenotypic analysis showed that both RANK⁺ $\text{p}\gamma\delta$ T and RANK⁺ $\text{d}\gamma\delta$ T cells expressed higher levels of a series of costimulatory molecules, including CTLA-4, OX-40, GITR, ICOS, and CD40, compared with RANK⁻ cells (Fig. 3a, b, Figure S1). On the contrary, the expression level of NKG2D, one of the cytotoxicity markers of $\gamma\delta$ T²², was decreased in RANK⁺ cells as compared with RANK⁻ cells (Fig. 3a, b, Figure S1). Moreover, cytokine pattern was different between RANK⁺ $\gamma\delta$ T cells and RANK⁻ $\gamma\delta$ T cells. RANK⁺ $\text{p}\gamma\delta$ T as well as RANK⁺ $\text{d}\gamma\delta$ T produce higher levels of IL-4, TGF- β 1, and IL-17A compared with RANK⁻ cells (Fig. 3a–c, Figure S1). In decidua, RANK⁺ $\gamma\delta$ T produce higher level of IL-10 compared with RANK⁻ $\gamma\delta$ T cells (Fig. 3a–c). However, such difference was not found in peripheral blood. Meanwhile, no significant difference in the production of IFN- γ between RANK⁺ and RANK⁻ $\gamma\delta$ T was detected (Figure S1). Taken together, RANK⁺ $\gamma\delta$ T are more functionally activated than RANK⁻ $\gamma\delta$ T.

RANKL expressed on DSCs induce polarization of Foxp3⁺ regulatory $\gamma\delta$ T cells and TGF- β 1 production

Different $\gamma\delta$ T subsets are characterized by the secretion of cytokines and expression of transcription factors⁹. By analyzing their profiles of cytokines and transcription factors, we explored whether RANKL/RANK signaling involved in the polarization of $\text{d}\gamma\delta$ T cells both in vitro and in vivo. To gain more insights into how RANKL overexpression affects the cytokine production of $\text{d}\gamma\delta$ T cells, we directly co-cultured the $\gamma\delta$ TCR⁺ cells isolated from human decidua tissues with RANKL-overexpressed



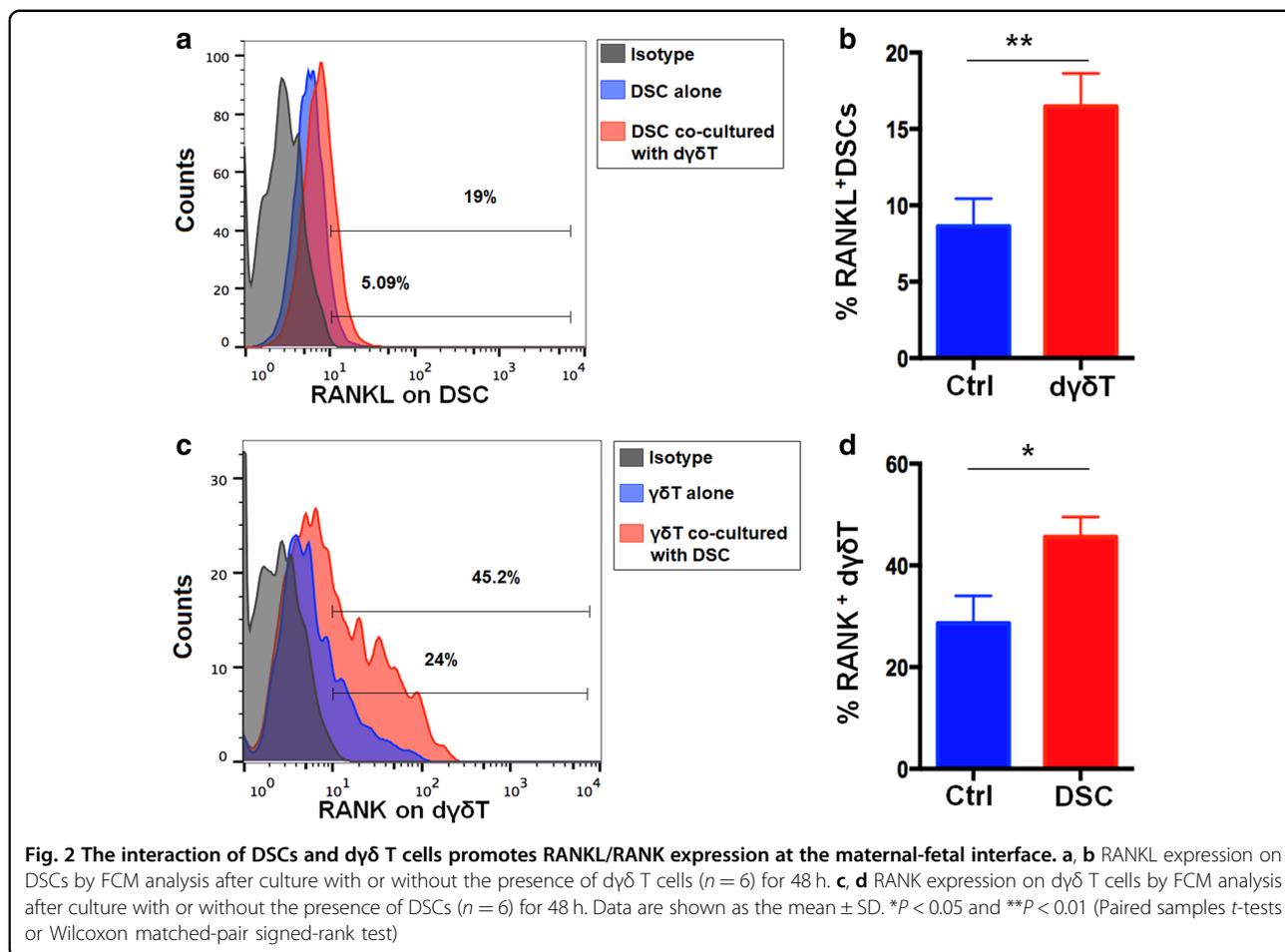
or control DSCs, and collected supernatant from co-culture system and subjected them to cytokines microarray analysis²³ (Fig. 4a). In the co-culture system with RANKL overexpression, cytokines were more highly produced including several TGF- β superfamily members TGF- β 1, 2, 3, as well as growth/differentiation factor (GDF) 9, 11, 15, and angiogenic factors vascular endothelial growth factor (VEGF) (Fig. 4b). Subsequently, FCM analysis showed that the secretion of cytokine TGF- β 1 and expression of transcription factor Foxp3 were significantly increased in $d\gamma\delta$ T cells after treated by RANKL-overexpressed DSCs as compared with control group (Fig. 4c, d). By contrast, there were no significant differences in other cytokines, including IFN- γ , TNF- α , IL-4, IL-17, IL-22, and IL-10, or transcription factors which include T-bet, GATA-3, RORC, AHR, and Bcl-6 (data not shown). Meanwhile, similar results were obtained when $p\gamma\delta$ T cells co-cultured with RANKL-overexpressed or control DSCs (Figure S2).

By using RANKL knockout mice, we further verified the effect of RANKL/RANK interaction on $\gamma\delta$ T cells at the maternal-fetal interface in vivo. Uterine $\gamma\delta$ T cells ($CD45^+CD3^+\gamma\delta TCR^+$) from pregnant RANKL^{-/-} mice

exhibited decreased Foxp3 expression and produced decreased amounts of TGF- β 1 compared to cells from pregnant wild-type (WT) mice (Fig. 4e). Thus, decidual RANKL/RANK interaction has little impacts on the polarization of $\gamma\delta$ T cells toward other subgroups, except for TGF- β 1-producing regulatory $\gamma\delta$ T cells.

Polarization of regulatory $\gamma\delta$ T cells and TGF- β 1 production induced by RANKL are dependent on the NF- κ B pathway

Recently, several studies discovered that nuclear factor kappa-light-chain-enhancer of activated B cells (NF- κ B), the major downstream pathway of RANKL/RANK interaction, is critical to the expression of Foxp3, as well as the development and immune suppression of regulatory T cells^{24,25}. As is known, when activated, NF- κ B is phosphorylated for further nuclear entry²⁶. To test whether RANKL/RANK signaling exerts effects on $\gamma\delta$ T cells by activating NF- κ B pathway, we assessed the phosphorylation status of NF- κ B in $\gamma\delta$ T cells. To this end, expression level of phospho-NF- κ B p65 were analyzed after gating on NF- κ B p65⁺ $\gamma\delta$ T cells. As shown in Fig. 5a, b, at day 8.5 of gestation, uterine NF- κ B p65⁺ $\gamma\delta$ T cells expressed less phospho-NF- κ B p65 in RANKL^{-/-}



mice compared to WT controls. Thus, RANKL-deficiency reduces NF- κ B activation in uterine $\gamma\delta$ T cells.

We further proved that the activation of NF- κ B is crucial for RANKL/RANK signaling to regulate $\text{d}\gamma\delta$ T polarization with the help of PDTC, a selective inhibitor of NF- κ B. As shown in Fig. 5c, d, increased levels of TGF- β 1 production and Foxp3 expression in $\text{d}\gamma\delta$ T, which was co-cultured with RANKL-overexpressed DSCs for 48 h, were almost completely aborted by PDTC. The pretreatment of human $\text{d}\gamma\delta$ T with PDTC for 1 h before the co-culture, was capable of aborting the effects of RANKL/RANK interaction on $\text{d}\gamma\delta$ T. These data suggest that NF- κ B signaling pathway is involved in the RANKL-induced polarization of Foxp3 + $\text{d}\gamma\delta$ T cells and TGF- β 1 production.

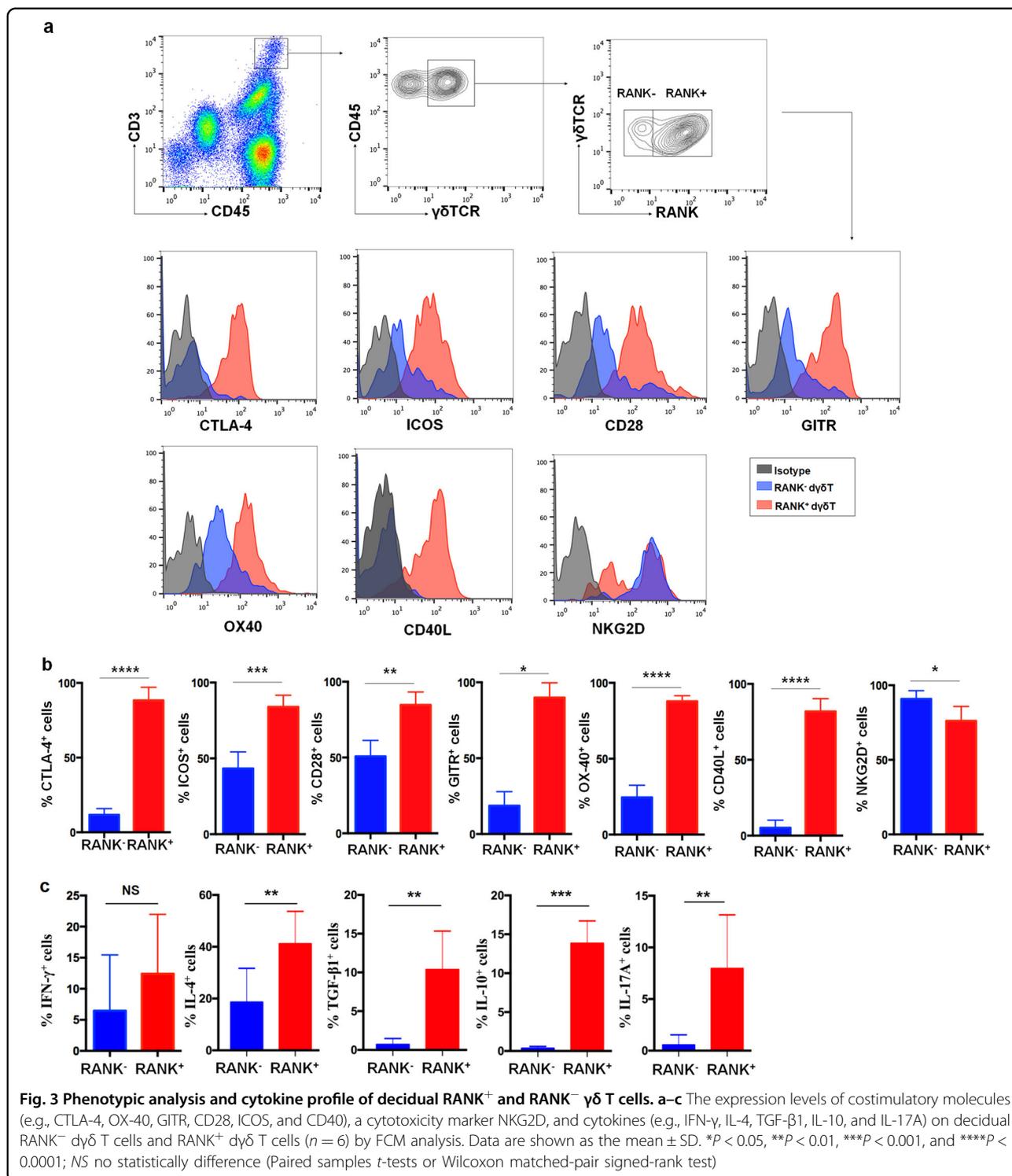
RANKL upregulates the expression of ICAM-1/VCAM-1 on DSCs and integrins on $\text{d}\gamma\delta$ T cells

Since RANKL enhances leukocyte residence by inducing adhesion molecules, including intercellular adhesion molecule 1 (ICAM-1) and vascular cell-adhesion molecule

1 (VCAM-1), in endothelial cells²⁷, we hypothesized that DSC adhesion molecule expression and $\gamma\delta$ T cell residence may also be regulated by RANKL. Importantly, in skin-draining lymph nodes, RANKL also upregulates cell-adhesion molecules in stromal cells²⁸. As shown, overexpression of RANKL significantly upregulated ICAM-1 and VCAM-1 in human DSCs (Fig. 6a, b).

In contrast, RANKL deficiency led to the decreased ICAM-1 and VCAM-1 expression in murine uterine stromal cells at day 8.5 of gestation (Fig. 6c). Thus, RANKL promotes the expression of cell-adhesion molecules in stromal cells at the maternal-fetal interface both in vitro and in vivo.

As is known, binding to integrin is a prerequisite of cell-adhesion molecule to mediate cell migration. ICAM-1 binds to integrins of lymphocyte function-associated antigen 1 (LFA-1, also known as CD11a/CD18), or macrophage-1 antigen (Mac-1, also known as CD11b/CD18), and VCAM-1 binds to integrins of very late antigen-4 (VLA-4, also known as integrin α 4 β 1, CD49d/CD29), or integrin α 4 β 7. As shown, all integrins



of ICMA-1 and VCAM-1 were expressed in γδ T cells of human decidua during early pregnancy (Fig. 6d). Of note, RANK⁺ dγδ T expressed higher levels of integrins, including LFA-1, CD11b, integrin α4β1, and integrin α4β7, compared with RANK⁻ cells (Fig. 6d). The results

of co-culture system in vitro showed that human dγδ T cells cultured with RANKL-overexpressed DSCs exhibited the increased integrins expression compared to cells cultured with control DSCs (Fig. 6e). Furthermore, significant reductions in integrins expression

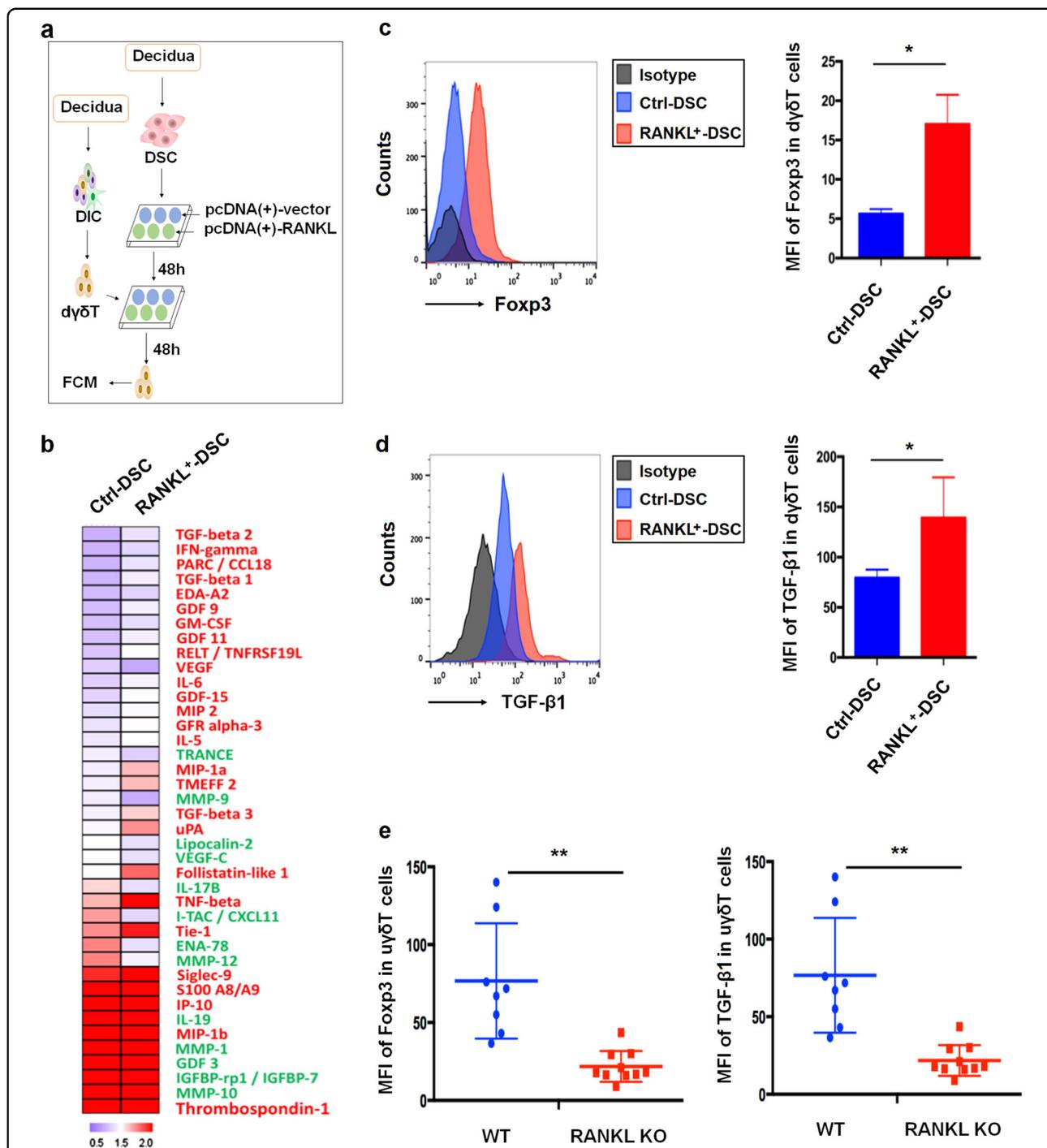
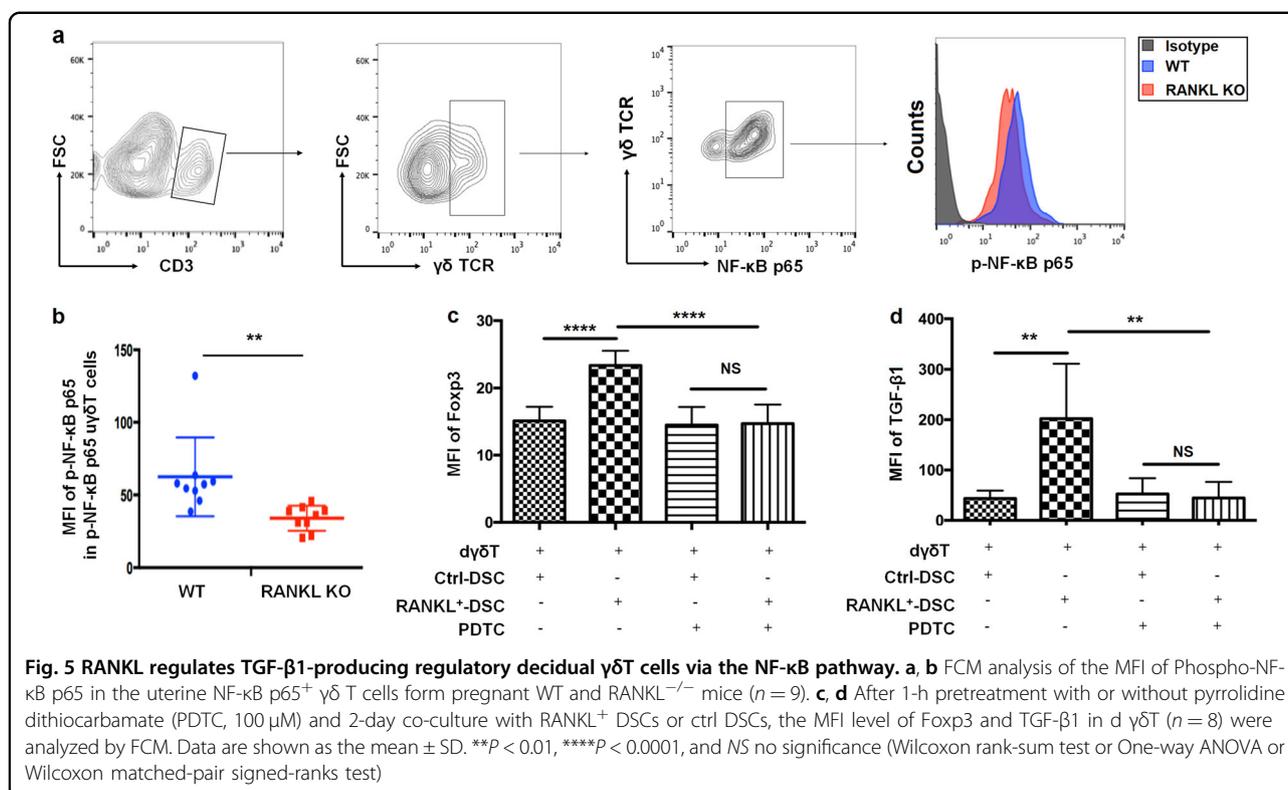


Fig. 4 RANKL from DSCs induce polarization of Foxp3⁺ regulatory $\gamma\delta$ T cells and TGF- β 1 production. **a** After co-culture with RANKL-overexpressed (RANKL⁺) or control (ctrl) DSCs at a 1:1 ratio for 48 h, the supernatants and decidual $\gamma\delta$ T (d $\gamma\delta$ T) cells were collected for protein array and flow cytometry analysis. **b** Heatmap of selected proteins differentially expressed in culture supernatants. **c, d** The median fluorescence intensity (MFI) of Foxp3 (**c**) and TGF- β 1 (**d**) in d $\gamma\delta$ T cells. **e** FCM analysis of MFI of Foxp3 and TGF- β 1 in uterine $\gamma\delta$ T cells from WT ($n = 8$) pregnant and RANKL^{-/-} ($n = 10$). WT wild-type, KO knockout. Data are shown as the mean \pm SD. * $P < 0.05$ and ** $P < 0.01$ (Wilcoxon matched-pair signed-ranks test or Student's t -test with correction)

were observed in uterine $\gamma\delta$ T cells of pregnant RANKL-deficient mice (Fig. 6f). Thus, these results indicate that DSC-derived RANKL upregulates cell-adhesion

molecule expression on DSCs, as well as integrin expression on $\gamma\delta$ T cells, and may further enhance $\gamma\delta$ T cell residence into decidua.



The RANKL/RANK interaction enhances the residence of $\gamma\delta$ T cell into decidua

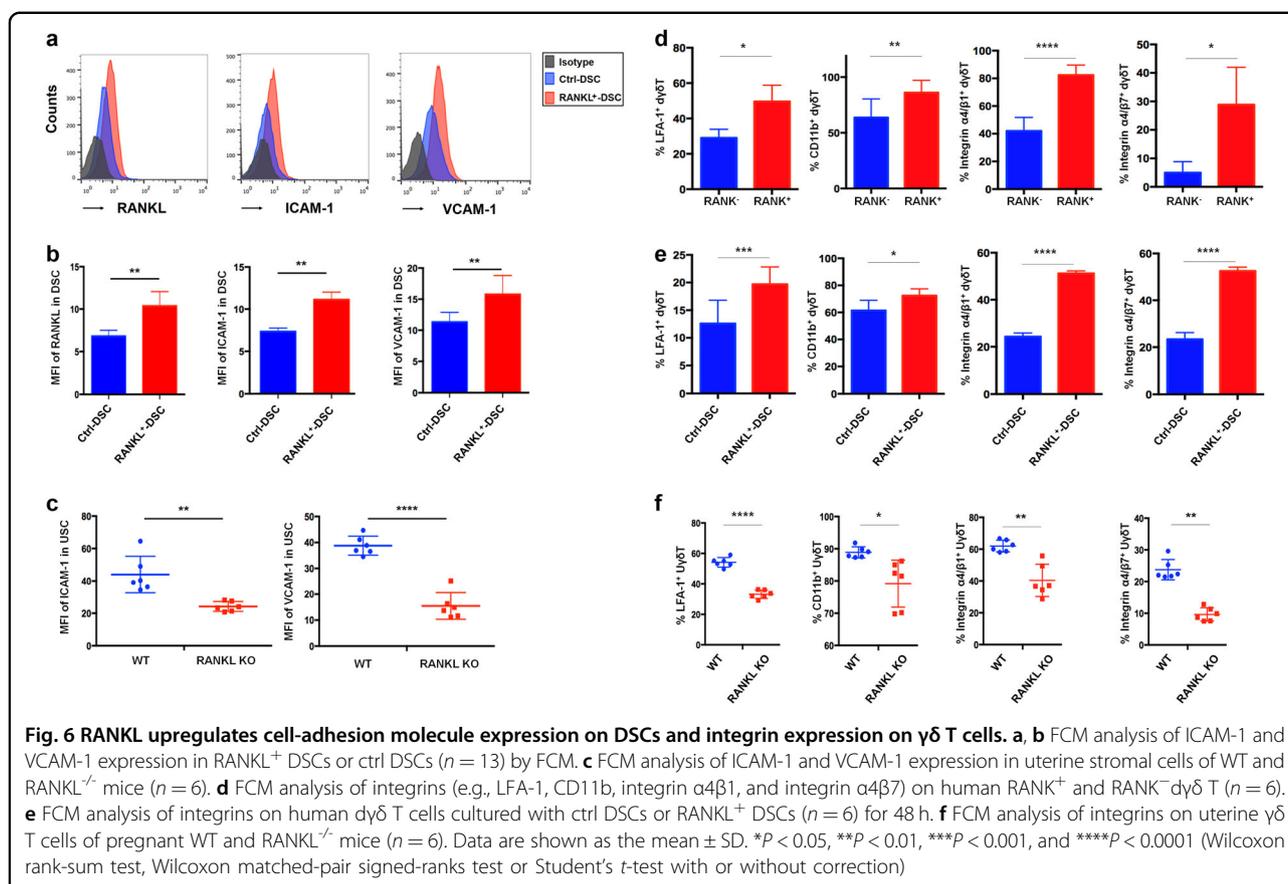
Considering ICAM-1 and VCAM-1 are important in leukocyte migration and residence, firstly we assessed the effect of RANKL on cell adhesion between DSCs and $\gamma\delta$ T cells by the adhesion experiments in vitro. As shown, overexpression of RANKL significantly increased the ability of DSCs to attract PKH-26-labeled $\gamma\delta$ T cells isolated from human decidua tissues (Fig. 7a, b). In support of these data, uterine immune cell populations in mice at day 8.5 of gestation were observed. Compared to WT mice, RANKL^{-/-} mice had reduced numbers and frequencies of uterine $\gamma\delta$ T cells, while the total number of T cells remained unaltered (Fig. 7c). Meanwhile, normal frequencies of other immune cell populations, including natural killer cells, natural killer T cells and macrophages, were present in the uterine of RANKL^{-/-} mice during early pregnancy (data not shown). Of note, the lack of RANKL resulted in the increase of mouse embryo absorption (Fig. 7d). These findings indicate that the RANKL/RANK interaction increases the residence of $\gamma\delta$ T cell into decidua in early pregnancy, and contributes to the establishment and maintenance of normal pregnancy.

Discussion

Here we demonstrated that, at the maternal-fetal interface, DSC-derived RANKL promotes the polarization of $\gamma\delta$ T cells into Foxp3⁺ regulatory $\gamma\delta$ T cells and

elevates TGF- β 1 production by activating NF- κ B pathway. Moreover, RANKL enhances the aggregation and residence of $\gamma\delta$ T cells into decidua by upregulating the expression of cell-adhesion molecules. Taken together, RANKL/RANK interaction contributes to the maternal-fetal tolerance by strengthening the crosstalk between DSCs and $\gamma\delta$ T cells.

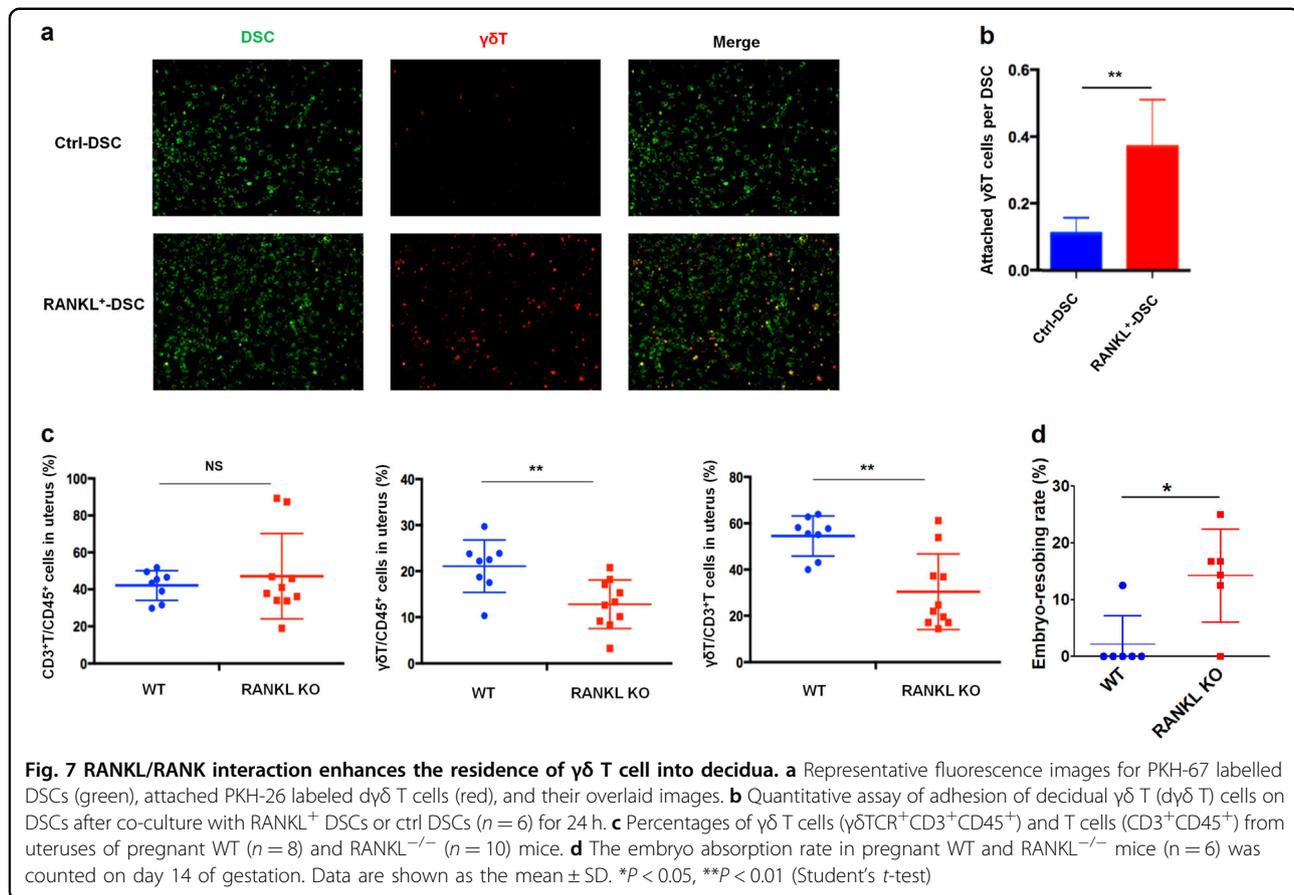
RANKL and RANK have been recently identified as key determinants of osteoclastogenesis. Hormones, cytokines, and other humoral factors influence osteoclastogenesis within the tumor/bone microenvironment primarily through increases in RANKL²⁹. In breast cancer, tumor-derived factors, including parathyroid hormone-releasing protein (PTHrP), prostaglandin E₂ (PGE₂), TNF α , and interleukins (e.g. IL-1, IL-6, IL-11), have been demonstrated to enhance RANKL expression by osteoblasts and other bone stromal cells present, which may directly contribute to osteoclastogenesis^{29,30}. In this study, the co-culture of DSCs and $\gamma\delta$ T cells led to the high levels of RANKL on DSCs and RANK on $\gamma\delta$ T cells, suggesting the cytokines secreted by DSCs and $\gamma\delta$ T cells may contribute to this regulatory process. Owing to the important role of PGE₂, IL-1 β , and IL-11 in the decidualization³¹⁻³⁵, it can be speculated that local microenvironment at maternal-fetal interface, including hormones, DSC, and DIC-derived cytokines, is bound to play an important role in maintaining high levels of RANKL; however, the specific mechanism remains to be studied.



In comparison with RANK⁻ $\gamma\delta$ T cells, RANK⁺ cells tend to be more active based on their higher expression level of costimulatory molecules and increased secretion of cytokines. Remarkably, among all seven tested costimulatory molecules, only NKG2D decreased in RANK⁺ $\gamma\delta$ T cells compared to that in RANK⁻ cells. As the most widely studied NK receptor (NKR) in $\gamma\delta$ T cells, NKG2D has been proved to mediate cytotoxicity of $\gamma\delta$ T cells and tissue damage^{36,37}. It is possible that, during early pregnancy, these highly activated RANK⁺ $\gamma\delta$ T cells play a role mainly in cytokine production instead of cell-killing at the maternal-fetal interface. Moreover, the increased expression level of CTLA-4 in RANK⁺ $\gamma\delta$ T cells is crucial in immunosuppression mediated by T cell³⁸.

We found that, during early pregnancy, RANKL plays a two-pronged strategy, which involves inducing polarization of Foxp3⁺ regulatory $\gamma\delta$ T cells through the highly expressed receptor RANK on $\gamma\delta$ T cells (over 90% are RANK⁺) and enhancing $\gamma\delta$ T cell residence into decidua. However, stimulation with recombinant human RANKL protein did not change the differentiation of regulatory $\gamma\delta$ T cells (data not shown), suggesting that the effect of DSCs on regulatory $\gamma\delta$ T cells polarization mainly depends on membrane RANKL. Previously, Foxp3⁺ T cells have been demonstrated to play a role in inducing

early maternal tolerance³⁹. In addition, the RANKL/RANK interaction promoted TGF- β 1 production of $\gamma\delta$ T cells. These processes are possibly dependent on the NF- κ B signaling pathway. TGF- β can directly inhibit the cytotoxicity and activity of effector cells at the maternal-fetal interface, including promotes the conversion of CD16⁺ NK cells into CD16⁻ NK cells⁴⁰, downregulates the expression level of CD80/CD86 on macrophages⁴¹, and limits IL-17 production in Th17 cells as well as T-bet expression and IFN- γ production in Th1 cells⁴². At the same time, TGF- β can in turn induces the polarization of Foxp3⁺ regulatory T cells⁴³, as well as protect them from apoptosis⁴⁴, and thus forms a positive feedback to suppress anti-fetal responses. Moreover, uterine $\gamma\delta$ T cells have been shown to prevent fetal rejection through TGF- β production at the maternal-fetal interface⁴⁵, which also consists with our results. In our previous study, we found that the depletion of macrophage in pregnant mice led to a significant decrease in RANKL on uterine DSCs, and adoptive transfer of RANK⁺ macrophages relieves murine embryo absorption induced by macrophage depletion²¹. These results suggest that the fetal loss in RANKL^{-/-} mice is closely related to the abnormal polarization of decidual macrophages. Further, in this study, we found that, except for participating in the polarization of



macrophages, RANKL also promotes the residence and polarization of TGF- β 1-producing regulatory $\gamma\delta$ T cells at the maternal-fetal interface. To sum up, RANKL/RANK interaction regulates the polarization, proliferation, immunoregulatory function, and residence of $\gamma\delta$ T cells, and may contribute to the suppression of fetal rejection as well as the maintenance of pregnancy during the first trimester, which needs further study.

At the maternal-fetal interface, the accumulation of $\gamma\delta$ T cells is quite obvious during early pregnancy. In peripheral blood, $\gamma\delta$ T cells are a minor subset of T cells (5–10%), while turns the major subset in decidua (over 60%)⁶. It has been reported that RANKL increases migration of human lung cancer cells through NF- κ B-upregulated ICAM-1⁴⁶. Our finding indicates that RANKL may also participate in $\gamma\delta$ T cells accumulation regulation by enhancing the adhesion of $\gamma\delta$ T cells to DSCs, and this effect should be dependent on the interaction of ICAM-1/VCAM-1 and integrins. However, the regulatory mechanism of RANKL on ICAM-1/VCAM-1 needs further research.

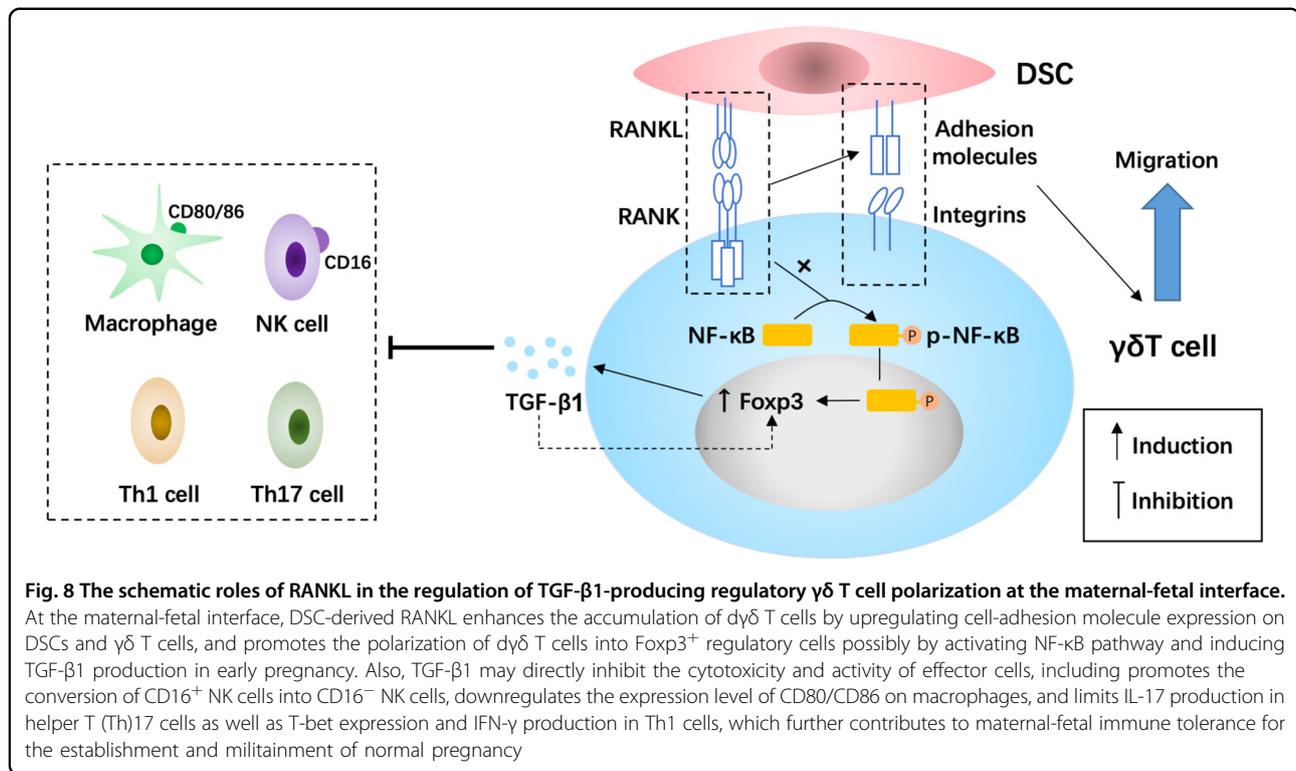
In summary, as shown in Fig. 8, our study demonstrated that during early pregnancy RANKL from DSCs, on the one hand, can induce the polarization of Foxp3⁺

regulatory $\gamma\delta$ T cells and TGF- β 1 production possibly by NF- κ B signaling pathway, on the other hand, can enhance the adhesion and residence of $\gamma\delta$ T cell into decidua. These integral effects will promote the resident and polarization of TGF- β 1-producing regulatory $\gamma\delta$ T cells, and contribute to the maternal-fetal immune tolerance, and establishment and maintenance of pregnancy. In contrast, an abnormal low expression of RANKL and RANK at the maternal-fetal interface leads to the deficiency of regulatory $\gamma\delta$ T cells and TGF- β 1 production and decidual macrophage dysfunction, and enhances the risk of RSA occurrence.

Materials and methods

Human samples and cell isolation

Decidua samples were obtained from 72 women with clinically normal pregnancy requiring pregnancy termination (age: 28.13 ± 4.37 years; gestational age at sampling: 50.92 ± 5.26 days; mean \pm SD) and 21 women with recurrent spontaneous abortions (age: 29.12 ± 3.09 years; gestational age at sampling: 61.04 ± 8.12 days; mean \pm SD). Abortions caused by genetic, endocrine, or anatomical abnormalities, as well as infection, were excluded. Normal endometrium was obtained at hysterectomy from patients



with uterine leiomyoma ($n = 10$) as healthy controls. None of the included patients had experienced complications related to pelvic inflammatory disease and none took any medications or received hormonal therapy within 6 months before surgery. The Ethics Committee of Obstetrics and Gynecology Hospital, Fudan University approved the study, and tissues were obtained with informed consent from each patient before surgery.

Fresh decidua tissues were washed in phosphate buffer saline (PBS) and cut into pieces within 30 min after operation. DICs and DSCs were then isolated from the shredded tissue as previously described⁴⁷. DICs and DSCs used in this study were isolated from the tissues of normal pregnancy recurrent spontaneous abortions.

Peripheral blood was collected from 26 healthy donors. Peripheral blood mononuclear cells (PBMCs) were isolated from peripheral blood by Ficoll-Paque (GE Healthcare) centrifugation.

Mice

Male and female RANKL heterozygote mice were purchased from the Jackson Laboratories (Sacramento, CA, USA). RANKL^{-/-} mice and WT littermates were generated by mating RANKL heterozygote. All mice were kept in the Laboratory Animal Facility of Fudan University (Shanghai, China). Age-matched (8 weeks) pregnancy mice were used in this study. The day of appearance of a copulatory plug was arbitrarily designated as day 0 of

gestation. Uteruses were collected from RANKL^{-/-} or WT littermates at day 8.5 of gestation. All procedures involving animals were approved by the Animal Care and Use Committee of Fudan University.

Primary cell cultures

Primary human and mice immune cells were cultured in RPMI 1640 (Gibco, Grand Island, NY, USA) with 10% fetal bovine serum (FBS; Gibco, Grand Island, NY, USA), 100 IU/mL penicillin, 100 μ g/mL streptomycin, 250 ng/mL amphotericin B, and 20 ng/mL recombinant human IL-2. Primary human and mice stromal cells were cultured in Dulbecco's Modified Eagle Media: Nutrient Mixture F-12 (DMEM/F-12; Gibco, Grand Island, NY, USA) with 10% fetal bovine serum (FBS; Gibco, Grand Island, NY, USA), 100 IU/mL penicillin, 100 μ g/mL streptomycin, and 250 ng/mL amphotericin B.

Flow cytometry

For flow cytometry (FCM) analyses, cells were incubated with monoclonal antibodies (Table S1) according to the manufacturer's instructions. Intracellular cytokines were detected after stimulating with Cell Activation Cocktail (Biolegend) for 6 h. Intracellular proteins or intranuclear transcription factors were stained using Fixation/Permeabilization Solution Kit (BD Pharmingen) or Transcription Factor Buffer Set (BD Pharmingen), respectively. All samples were run on a Beckman-Coulter

CyAN ADP Analyzer (Beckman-Coulter, Brea, CA, USA), and data were analyzed by FlowJo software (TreeStar, Ashland, OR, USA).

Immunohistochemistry

Anti-RANKL antibody (R&D Systems, Abingdon, UK) was used for human decidua and endometrium immunohistological staining according to a previously described method⁴⁷.

Purification of $\gamma\delta$ T cells

Human $\gamma\delta$ T cells were isolated from DICs or PBMCs using Anti- $\gamma\delta$ TCR MicroBead Kit (Miltenyi Biotec, Bergisch Gladbach, Germany) according to the manufacturer's instructions, and a 95% purity of CD45⁺CD3⁺ $\gamma\delta$ TCR cells was confirmed by flow cytometry.

Co-culture of DSCs and $\gamma\delta$ T cells

DSCs and $\gamma\delta$ T cells isolated from decidual of normal pregnancy were used to construct the co-culture system in vitro. More concretely, DSCs were cultured in a 6-well plate (5×10^5 cells per well) for overnight, and then co-cultured with or without freshly isolated $\gamma\delta$ T cells at a 1:1 ratio. In the meantime, part of the isolated $\gamma\delta$ T cells were cultured in a 6-well plate (5×10^5 cells per well) without DSCs. After 48 h, DSCs and $\gamma\delta$ T cells were collected for further flow cytometry analyses.

After 48 h of transfection with pcDNA(+)-RANKL or pcDNA(+)-vector plasmid (GeneChem Co., Ltd, Shanghai, China), DSCs were directly co-cultured with freshly isolated decidual/peripheral $\gamma\delta$ T cells at a 1:1 ratio. In addition, in some groups, $\gamma\delta$ T cells were pretreated with or without 100 μ M pyrrolidine dithiocarbamate (PDTC; MedChemExpress, USA) for 1 h, and co-cultured with RANKL-overexpressed or control DSCs. After 48 h of co-culture, $\gamma\delta$ T cells were collected for further flow cytometry analyses.

Protein array

After cultured in 5% FBS medium for an extra day, supernatants were collected from the DSC (from three women with normal pregnancy, mixed in the ratio of 1:1:1) $\gamma\delta$ T co-culture system. The expression levels of over 500 proteins (including cytokines, chemokines, growth factors, angiogenic factors, etc.) were tested using L-Series Antibody Array 507 (RayBiotech) by the H-Wayen Biotechnologies in Shanghai.

Adhesion experiment

After 2 days of transfection, RANKL-overexpressed or control DSCs were collected, washed, then stained with PKH-67 (Sigma), and cultured in a 96-well plate (3×10^4 cells per well) for overnight. After that, PKH-26- (Sigma)

labeled $\gamma\delta$ T cells were co-cultured with the treaded DSCs for 24 h. Then, to remove unattached $\gamma\delta$ T cells, the media was removed, and cells were washed gently with PBS twice. The results were obtained under a fluorescent microscope (Olympus, Tokyo, Japan) at five random fields of view and showed as the ratio of $\gamma\delta$ T cells to DSCs.

Statistical analysis

The continuous variable is shown as the mean \pm S.D. Variables between two dependent groups were analyzed by paired or unpaired samples *t*-test when the data are normally distributed and by Wilcoxon matched-pair signed-ranks test when the data are not normally distributed. As for independent samples, Student's *t*-test with or without correction, Wilcoxon rank-sum test, one-way ANOVA, or Kruskal–Wallis H test was used according to the analysis of normal distribution and variance homogeneity. All analyses were conducted with Graphpad Prism software (La Jolla, CA, USA). *P* < 0.05 (*) was considered statistically significant.

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Conflict of interest

The authors declare that they have no conflict of interest.

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