Editorial

RBP-J and ITAM crosstalk

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Osteoclasts, multinucleated giant cells derived from the monocyte/macrophage lineage, are the exclusive specialized bone-resorbing cells that are indispensable for skeletal development, bone remodeling and repair and mineral homeostasis. Increased osteoclastogenesis leads to excessive bone resorption in osteoporosis and in inflammatory diseases with bone destruction, such as rheumatoid arthritis (RA), psoriatic arthritis (PsA), juvenile chronic arthritis (JCA) and periodontitis [1]. As bone is one of the major metastatic sites for tumorigenesis of the breast, prostate, lung as well as multiple myeloma, abnormal generation and activation of osteoclasts facilitate skeletal metastasis and tumor growth in bone tissue [2]. Thus, appropriate control of osteoclastogenesis will be a direct and highly effective approach in the prevention and treatment of osteolysis in pathological conditions.

The immunoreceptor tyrosine-based activation motif (ITAM)-mediated signaling pathways play important roles in various cellular activities, including immune response and cancer activation. The main ITAM-containing adaptors expressed by myeloid osteoclast precursors are DNAXactivating protein 12 (DAP12) and Fc receptor common γ subunit (FcR γ). These adaptors associate with and mediate signaling by various receptors, including DAP12associated triggering receptor expressed in myeloid cells 2 (TREM2) and signal-regulatory protein β 1 (SIRP β 1), FcRy-associated osteoclast-associated receptor (OSCAR), paired immunoglobulin-like receptor-A (PIR-A) and FcRs. Osteoclasts require ITAM-mediated costimulation of RANK signaling for their appropriate differentiation during bone homeostasis [3]. Osteoclastogenesis is delicately modulated at various levels, but little is known about the mechanisms that regulate ITAM-mediated costimulation. Recombinant recognition sequence binding protein at the J₂ site (RBP-J, also named RBP-J₂, CSL or CBF1) functions as a central transcription factor that receives inputs from several signaling pathways, such as the canonical Notch pathway, Wnt-β-catenin, NF-κB pathway, TLR and TNF signaling pathways. We recently identified RBP-J as a key negative regulator that imposes the requirement for ITAM-mediated costimulation of osteoclastogenesis, homeostatic bone remodeling and inflammatory bone resorption, and discovered the signaling pathways and mechanisms by which RBP-J regulates ITAM-mediated costimulation [4, 5].

Loss of DAP12 or both DAP12 and FcR γ in mice leads to severe osteopetrosis due to impaired

osteoclastogenesis and resorptive function. To investigate the role of RBP-J in ITAM-mediated bone remodeling, we generated DAP12 and RBP-J double knockout (KO) mice (R-DKO) and DAP12, FcRy and RBP-J triple KO mice (TKO) [5]. Strikingly, we found that RBP-J deficiency almost completely reversed the defects of osteoclast differentiation program and significantly rescued the osteopetrotic bone phenotype of Dap12 KO or Dap12/ *Fcrg* double knockout (DKO) mice by bypassing the requirement for costimulation of osteoclastogenesis during bone homeostasis. In inflammatory settings, RBP-J deficiency enabled TNF- α to induce osteoclast formation and bone resorption in Dap12 KO mice. These data show that RBP-J deficiency allows osteoclast differentiation to occur independently of ITAM-mediated costimulation during homeostatic bone remodeling and inflammatory bone destruction. Thus, RBP-J imposes the requirement for ITAM-mediated costimulation. During osteoclastogenesis, ITAM-mediated PLCy2-calcium signaling is constitutively active at basal level, which provides essential costimulatory signals for RANK signaling to induce osteoclast differentiation. We show that RBP-J restrains ITAM signaling and limits crosstalk between RANK and ITAM associated receptors by suppressing PLCy2 expression and keeping PLCy2-calcium signaling in a repressed basal state. This is the first study showing the negative regulation of ITAM signaling by RBP-J. The regulatory mechanisms of PLCy2 expression were unclear. Our results show that the mechanism of PLC γ 2 regulation is transcriptional in osteoclast differentiation. Furthermore, built upon RNAseq experiments, we identified a novel regulatory loop whereby RBP-J restrains the expression of TGFBR1 and thereby modulates cell responsiveness to TGF- β , which in turn regulates transcription of PLC γ 2 [5].

Our findings provide evidence for a cross talk between RBP-J and ITAM-mediated signaling pathways in osteoclast precursors and suggest a conceptually new model where ITAMs positively while RBP-J negatively regulates PLC γ 2-calcium signaling. The balance between these opposing pathways fine tunes osteoclastogenesis and determines the strength of activating signals required for osteoclastogenesis in physiological and various pathological settings. For example, we observed that RBPJ expression level was significantly suppressed in synovial fluid macrophages derived from RA patients, indicating that RBP-J expression can be altered in response to environmental cues in pathological states [5]. As RBPJ is a newly identified risk locus for RA, our data providing new evidence for a link between RBP-J and RA highlights the significance of understanding RBP-J mediated mechanisms in osteoclastogenesis and bone resorption. The cross talk between RBP-J and ITAMmediated signaling pathway as well as the signaling network identified by our study may also exist in other cell types. In cancers, Notch/RBP-J is directly involved in abnormal differentiation of myeloid cells, including dendritic cells and myeloid-derived suppressor cells (MDSC) that can function as osteoclast progenitors to enhance bone loss in tumors [6, 7]. Therefore, the newly described RBP-J/PLC γ 2/calcium pathway would also have therapeutic implications for abnormal osteoclastogenesis and associated skeletal destruction in the cancer setting.

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