

# Notch Signaling Promotes Osteoclast Maturation and Resorptive Activity

Jason W. Ashley,<sup>1</sup> Jaimo Ahn,<sup>1\*</sup> and Kurt D. Hankenson<sup>2,3\*\*</sup>

<sup>1</sup>Department of Orthopaedic Surgery, Perelman School of Medicine, University of Pennsylvania, Philadelphia, Pennsylvania

<sup>2</sup>Department of Small Animal Clinical Sciences, College of Veterinary Medicine, Michigan State University, East Lansing, Michigan

<sup>3</sup>Department of Physiology, Colleges of Natural Sciences and Osteopathic Medicine, Michigan State University, East Lansing, Michigan

## ABSTRACT

The role of Notch signaling in osteoclast differentiation is controversial with conflicting experimental evidence indicating both stimulatory and inhibitory roles. Differences in experimental protocols and in vivo versus in vitro models may explain the discrepancies between studies. In this study, we investigated cell autonomous roles of Notch signaling in osteoclast differentiation and function by altering Notch signaling during osteoclast differentiation using stimulation with immobilized ligands Jagged1 or Delta-like1 or by suppression with  $\gamma$ -secretase inhibitor DAPT or transcriptional inhibitor SAHM1. Stimulation of Notch signaling in committed osteoclast precursors resulted in larger osteoclasts with a greater number of nuclei and resorptive activity whereas suppression resulted in smaller osteoclasts with fewer nuclei and suppressed resorptive activity. Conversely, stimulation of Notch signaling in osteoclast precursors prior to induction of osteoclastogenesis resulted in fewer osteoclasts. Our data support a mechanism of context-specific Notch signaling effects wherein Notch stimulation inhibits commitment to osteoclast differentiation, but enhances the maturation and function of committed precursors. *J. Cell. Biochem.* 116: 2598–2609, 2015. © 2015 Wiley Periodicals, Inc.

**KEY WORDS:** Osteoclasts; Notch signaling; Bone resorption; Osteoclast fusion; Osteoclast commitment

Molecular signaling for osteoclast differentiation is initiated by monocyte/macrophage colony-stimulating factor (M-CSF) and Receptor Activator of NF- $\kappa$ B Ligand (RANKL) [Feng, 2005; Teitelbaum, 2007]. While indispensable for proper osteoclast formation, these pathways are subject to modulation [Izawa et al., 2012; Jules et al., 2012; Hong et al., 2013; McCoy et al., 2013]. Signaling pathways regulating osteoclast precursor proliferation, osteoclast fusion, function, and survival potentially impact the number and activity of osteoclasts differentiated under M-CSF/RANKL stimulation [Feng, 2005; Li et al., 2011; Maeda et al., 2012; Park et al., 2013]. Notch signaling is an important, yet incompletely understood, pathway that has been suggested to have multiple roles

in the differentiation of osteoclasts [Duan et al., 2008]. Understanding the seemingly variegated roles of Notch signaling in osteoclasts is critical to the rational design of bone-preserving therapies targeting this pathway.

Signaling from mammalian homologues of *Drosophila* Notch receptor is an essential component of embryonic development, tissue patterning, and differentiation of stem cell populations [Artavanis-Tsakonas et al., 1999]. In mammals, there are four Notch receptors (Notch1, 2, 3, and 4), and multiple ligands of the Delta-like (Delta-like1, Delta-like3, and Delta-like4) and Jagged (Jagged1 and Jagged2) families [Chen et al., 2014]. Notch signaling has two distinguishing characteristics. First, Notch signaling can only be

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\*Correspondence to: Jaimo Ahn, MD, PhD, Department of Orthopaedic Surgery, University of Pennsylvania, 3737 Market St, Suite 600, Philadelphia, PA 19104.

E-mail: jaimo\_ahn@alumni.stanford.edu

\*\*Correspondence to: Kurt D. Hankenson DVM, PhD, Department of Small Animal Clinical Sciences, G334 Veterinary Medical Center, 736 Wilson Road, East Lansing MI 48824.

E-mail: kdhan@msu.edu

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properly initiated in a target cell via receptor binding by a ligand on the plasma membrane of another cell (*trans*-activation); interactions between ligand and receptor on the same cell suppress Notch signaling (*cis*-inhibition)[Cordle et al., 2008; Wang et al., 2011]. In addition, activation of Notch signaling requires tensile force between receptor and ligand, and, because of this, Notch signaling can only be efficiently activated in culture through the use of immobilized ligands [Kopan and Ilagan, 2009; Wang and Ha, 2013]. Second, upon ligand stimulation, proteolytic cleavage events catalyzed by Tumor Necrosis Factor  $\alpha$ -converting enzyme (TACE) and  $\gamma$ -secretase release the intracellular domain of Notch (NICD) which translocates to the nucleus where it forms a complex with mastermind-like (MAML), CSL, and general transcription factors to activate expression of target genes [Groot et al., 2014]. Notch signaling can be inhibited with  $\gamma$ -secretase inhibitors, such as N-[N-(3,5-difluorophenacetyl-L-alanyl)]-(S)-phenylglycine t-butyl ester (DAPT). SAHM1, a peptide mimetic of a dominant negative form of MAML, inhibits canonical Notch transcription complex formation [Kornilova et al., 2003; Morohashi et al., 2006; Jones, 2009; Moellering et al., 2009].

The controversial findings of previous studies regarding Notch signaling in osteoclasts may be summarized as follows: (1) Notch signaling suppresses osteoclast differentiation, (2) Notch signaling enhances osteoclast differentiation, and (3) Notch1 signaling stimulated by Jagged1 (JAG1) suppresses osteoclast differentiation and Notch2 signaling stimulated by Delta-like1 (DLL1) enhances osteoclast differentiation [Yamada et al., 2003; Bai et al., 2008; Fukushima et al., 2008; Schwarzer et al., 2008; Sethi et al., 2011; Sekine et al., 2012; Zhou et al., 2014; Zhou et al., 2015]. The disparate findings of these studies likely stem from differences in experimental approach and may be evidence for context-dependent roles for Notch signaling in osteoclasts. Resolution of these conflicting findings is critical for informed design of mechanistic studies of Notch signaling in osteoclast function.

This study investigates osteoclast differentiation and function in the context of Notch stimulation and suppression in a cell-autonomous fashion using cultured osteoclast precursors, Notch ligands, and Notch signaling inhibitors. This study utilizes multiple specific methodologies in order to isolate Notch signaling effects at distinct points during differentiation and clarifies the context-dependent roles of Notch signaling in osteoclastogenesis.

## MATERIALS AND METHODS

### ANIMAL CARE AND USE

Mice were maintained according to Institutional Animal Care and Use Committee guidelines of the University of Pennsylvania.

### CELL CULTURE REAGENTS

Osteoclast/macrophage medium ( $\alpha$ -10) was prepared by dissolving one vial of Eagle's minimum essential medium powder (Sigma-Aldrich, M0894-10  $\times$  1L) and 2 g sodium bicarbonate (Fisher Scientific, BP328-500) in 890 mL deionized water and adding 100 mL heat-inactivated fetal bovine serum, 10 mL 10000 units/mL penicillin/10000  $\mu$ g/mL streptomycin (Life Technologies, 15140122),

and 10 mL 200 mM L-glutamine (Life Technologies, 25030-081). Medium was sterile filtered using two 500 mL 0.22  $\mu$ m filter bottles (Corning, 431097) and stored at 4°C. 10% L929 cell-conditioned medium was added to  $\alpha$ -10 as a source of monocyte/macrophage colony-stimulating factor (MCSF)

Recombinant mouse Receptor Activator of NF- $\kappa$ B Ligand (RANKL) (Shenandoah Biotechnology, 200-04) was reconstituted in sterile phosphate buffered saline (PBS) (CellGro, 21-031-CV) at 100  $\mu$ g/mL and stored at  $-80^{\circ}\text{C}$ . Recombinant human Jagged1-Fc (R&D Systems, 1277-JG-050) and Delta-like1-Fc (Adipogen, AG-40A-0116Y-C050) were reconstituted at 200  $\mu$ g/mL in sterile PBS and stored at  $-80^{\circ}\text{C}$ . Fc fragment-specific goat anti-human IgG (Jackson ImmunoResearch, 109-001-008) was reconstituted at 200  $\mu$ g/mL in sterile water and stored at  $-80^{\circ}\text{C}$ . DAPT (Sigma-Aldrich, D5942-5MG) and SAHM1 (EMD Millipore, 491002-1MG) were dissolved in DMSO (Fisher Scientific, BP231-100) at 10 mM and stored at  $-80^{\circ}\text{C}$ .

### MODULATION OF NOTCH SIGNALING WITH IMMOBILIZED LIGANDS AND SOLUBLE INHIBITORS

Notch signaling was induced by seeding cells onto IgG- (control), JAG1-, or DLL1-coated surfaces. Notch ligand-coated plates were prepared as previously described [Zhu et al., 2013]. 250  $\mu$ L (for 24-well plates) of 10  $\mu$ g/mL goat anti-human IgG in PBS were added to each well to be coated and incubated for 1 h at room temperature. Wells were washed 3  $\times$  with PBS, and 250  $\mu$ L of PBS (IgG wells), 10  $\mu$ g/mL Jagged1-Fc, or 10  $\mu$ g/mL Delta-like1-Fc were added and incubated for 2 h at room temperature. Wells were washed 3  $\times$  with PBS. Wells were kept filled with PBS until cell seeding to prevent drying. Notch signaling was inhibited by culturing cells in medium containing either 10  $\mu$ M DAPT, which inhibits  $\gamma$ -secretase, or 10  $\mu$ M SAHM1, which inhibits NICD-MAML-CSL transcription complex formation. Figure 1 summarizes the general mechanism of Notch signaling and points of experimental manipulation.

### OSTEOCLAST PRECURSOR CULTURE AND DIFFERENTIATION

Osteoclast precursors were obtained according to an established protocol [Ashley et al., 2011]. Mice were sacrificed by CO<sub>2</sub> inhalation followed by cervical dislocation. Tibiae and femora from each mouse were dissected and each was flushed with 2.5 mL  $\alpha$ -10 via a 10 mL syringe (BD 14-823-2A) with a 25-gauge needle (BD, 305125) into a single 15 mL conical tube which was mixed by inversion. Tissue debris were allowed to settle and the cell suspension was transferred to a new 15 mL tube and pelleted at 300g for 5 min. Pellets were resuspended in 1 mL ACK lysing buffer (Life Technologies, A10492-01) and incubated at 37°C for 2 min to lyse red blood cells. 8 mL PBS was then added to the tube and mixed by inversion. Cells were pelleted at 300g for 5 min and resuspended in  $\alpha$ -10 without MCSF, plated in a 100 mm tissue culture treated plate (Corning, 430167), and incubated in a humidified 37°C, 5% CO<sub>2</sub> incubator overnight. The next morning, non-adherent cells were collected for differentiation into osteoclast using one of these protocols.

(1) RANK signaling stimulation prior to Notch signaling stimulation [Ashley et al., 2011]. Non-adherent bone marrow cells

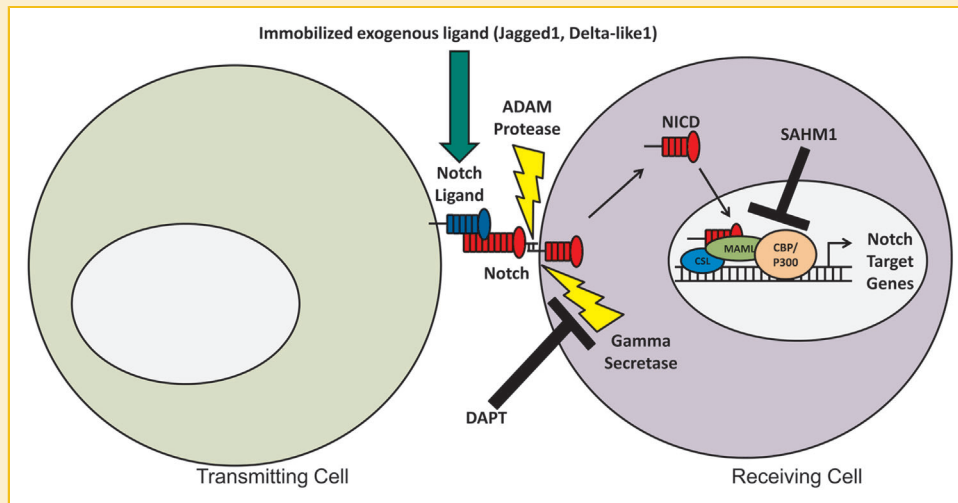


Fig. 1. Summary of experimental manipulation of Notch signaling. Notch signaling was stimulated by plating cells on antibody-immobilized Fc-fusion proteins of either Jagged1 or Delta-like1; antibody only was used as the control for ligand-stimulated cells. Notch signaling was inhibited with either the  $\gamma$ -secretase inhibitor, DAPT, which inhibits Notch signaling at the point of NICD release, or the dominant-negative MAM1 peptide mimetic, SAHM1, which inhibits Notch signaling at the transcriptional level; DMSO, the vehicle for both inhibitors, was used as the control for Notch inhibition.

TABLE 1. Primer Name

	Sequence (5' to 3')	Measured Transcript
18S_RTQ_F	GGTAACCCGTTGAACCCCAT	18S - Loading Control
18S_RTQ_R	CAACGCAAGCATTATGACCCG	
Notch1_RTQ_F	GCTCCGAGGAGATCAACGAG	Notch1
Notch1_RTQ_R	TTGACATCACCTCACACCG	
Notch2_RTQ_F	ATATCGACGACTGCCCAAC	Notch2
Notch2_RTQ_R	CCATAGCCTCCGTTTCGGTT	
Notch3_RTQ_F	ATGTGGCCGTGGCTATACTG	Notch3
Notch3_RTQ_R	AGTGCTTGACACTCATCCA	
Notch4_RTQ_F	GTCTGAGGGTAATTCCTGC	Notch4
Notch4_RTQ_R	CTTGGCAGGTTCCTTGTTTC	
DLL1_RTQ_F	CAATCTGTCTGCGAGGGTGT	Delta-like1
DLL1_RTQ_R	CGGATGCACTCATCGCAGTA	
DLL4_RTQ_F	GATGAGGGATGGGAGGTCT	Delta-like4
DLL4_RTQ_R	CGCGCAGGTCAAGGTACTAT	
JAG1_RTQ_F	GCAACGACCGTAATCGCATC	Jagged1
JAG1_RTQ_R	CCATTGCCGGTAGGGTTTA	
JAG2_RTQ_F	GGGTGGCAACTCTTCTACC	Jagged2
JAG2_RTQ_R	GTCAATGTGCCAGTCCAGG	
HES1_RTQ_F	GAGGCTGCCAAGGTTTTTGG	HES1
HES1_RTQ_R	ACTTTACGGGTAGCAGTGGC	
HEY1_RTQ_F	GCTCACCCAGACTACAGCTC	HEY1
HEY1_RTQ_R	CGCTTCTCGATGATGCCTCT	
ACP5_RTQ_F	CAGCTCAGTTGGGTAGCACA	TRAP
ACP5_RTQ_R	AGCCACAAATCTCAGGGTGG	
MMP9_RTQ_F	CGCTCATGTACCCCGCTGTAT	MMP9
MMP9_RTQ_R	CCGTGGGAGGTATAGTGGGA	
CTSK_RTQ_F	CAGTGTGGTGGTGGGCTAT	Cathepsin K
CTSK_RTQ_R	CATGTTGGTAAATGCCGAGG	
CTR_RTQ_F	GGTGCGCGGGATCCTATAA	Calcitonin Receptor
CTR_RTQ_R	CACGAGTGTAGTGGCGTGGATA	
CD200_RTQ_F	GGGGTGAATCATCACAGGGG	CD200
CD200_RTQ_R	CAAATCCCTCACAGGCTCGT	
DCSTAMP_RTQ_F	CATGTGGGTGCTGTTTGGC	DC-STAMP
DCSTAMP_RTQ_R	GACTCCTTGGGTTCCTTGCTT	
CD9_RTQ_F	TTCTGTCCAGTCGTTCTGTG	CD9
CD9_RTQ_R	CTGAGAGTGAATCGGAGCC	
CD81_RTQ_F	GAACTGGGAAACAAACCGGC	CD81
CD81_RTQ_R	ATAGCACCCAGGAAGCCTA	
SIRPA_RTQ_F	GAAACCATACCGTCTGGGA	Signal-regulatory protein alpha
SIRPA_RTQ_R	CTGGGTTATTTCCCTGGCGT	
ATP6V <sub>0</sub> D <sub>2</sub> _RTQ_F	ATGCAAAGCCAGCTCTCTAA	v-ATPase V <sub>0</sub> subunit D <sub>2</sub>
ATP6V <sub>0</sub> D <sub>2</sub> _RTQ_R	TTGCCATAGTCCGTGGTCTG	

were evenly distributed between five 60 mm non tissue culture-treated suspension plates (Corning, 08-772-31) in  $\alpha$ -10 with MCSF. Three days after seeding, adherent macrophages were washed once with PBS and incubated with 0.5 mL 5 mM EDTA (Fisher Scientific, S312-500) in PBS per plate for 3–5 min at 37°C. 1.5 mL  $\alpha$ -10 was added to each plate and lifted cells were pooled in a single 50 mL conical tube and counted using a hemocytometer. Cells were diluted to  $2 \times 10^5$  cells/mL in  $\alpha$ -10 with MCSF, and 250  $\mu$ L of the cell suspension was added to each well of a 24-well plate to achieve  $5 \times 10^4$  cells/well ( $2.6 \times 10^4$  cells/cm<sup>2</sup>). To each well, 250  $\mu$ L  $\alpha$ -10 with MCSF and double concentrations of each well's particular treatment (e.g., 200 ng/mL RANKL, 20  $\mu$ M DAPT) was added to achieve correct final concentrations. Fourty eight hours after seeding, media, and treatments in wells were refreshed. For wells treated with SAHM1, additional SAHM1 to a final concentration of 10  $\mu$ M was added to the wells daily. Seventy two hours after seeding, cells were washed with PBS and either fixed and stained for TRAP expression using a leukocyte acid phosphatase staining kit (Sigma-Aldrich, 387A) or lysed in 1 mL TRIzol (Life Technologies, 15596-026) for RNA extraction.

- (2) Near-simultaneous stimulation of Notch and RANK signaling [Bradley and Oursler, 2008]. Non-adherent bone marrow cells were transferred to a conical tube and counted. Variable numbers of non-adherent cells ( $2-10 \times 10^5$ ) were seeded into a 24-well plate with MCSF and 100 ng/mL RANKL. Differentiation medium was refreshed on day 3 and day 4 of the differentiation. Cells were TRAP stained after 5 days of differentiation.
- (3) RANK signaling stimulation after Notch signaling stimulation [Tevlin et al., 2014]. Non-adherent bone marrow cells were transferred to a conical tube and counted. Variable numbers of non-adherent cells ( $1-8 \times 10^5$ ) were seeded into a 24-well plate with MCSF and cultured for 3 days. On the third day of culture medium was replaced with  $\alpha$ -10 containing MCSF and 100 ng/mL RANKL. Medium was refreshed daily, and differentiated cells were TRAP stained 3 days later.

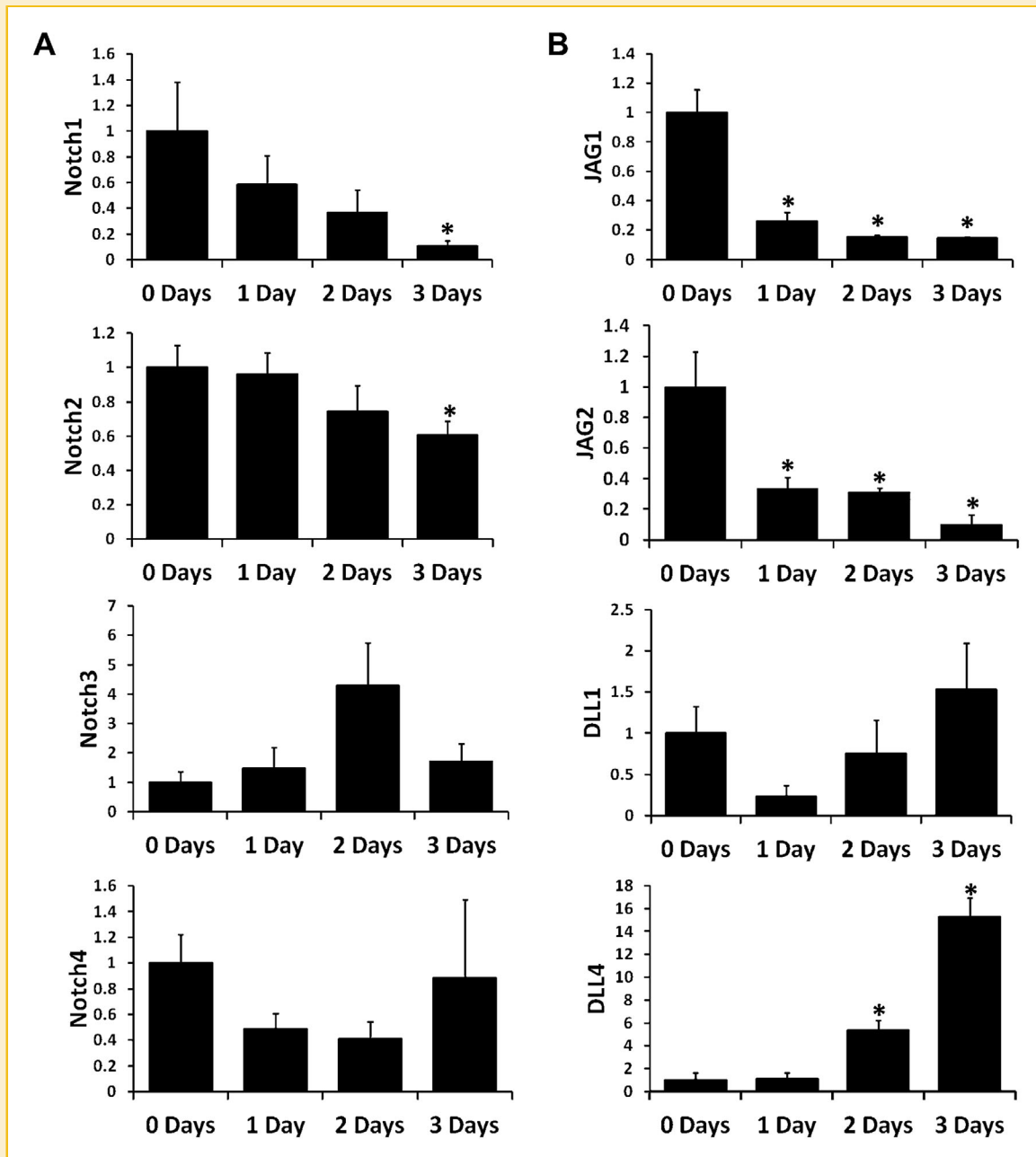
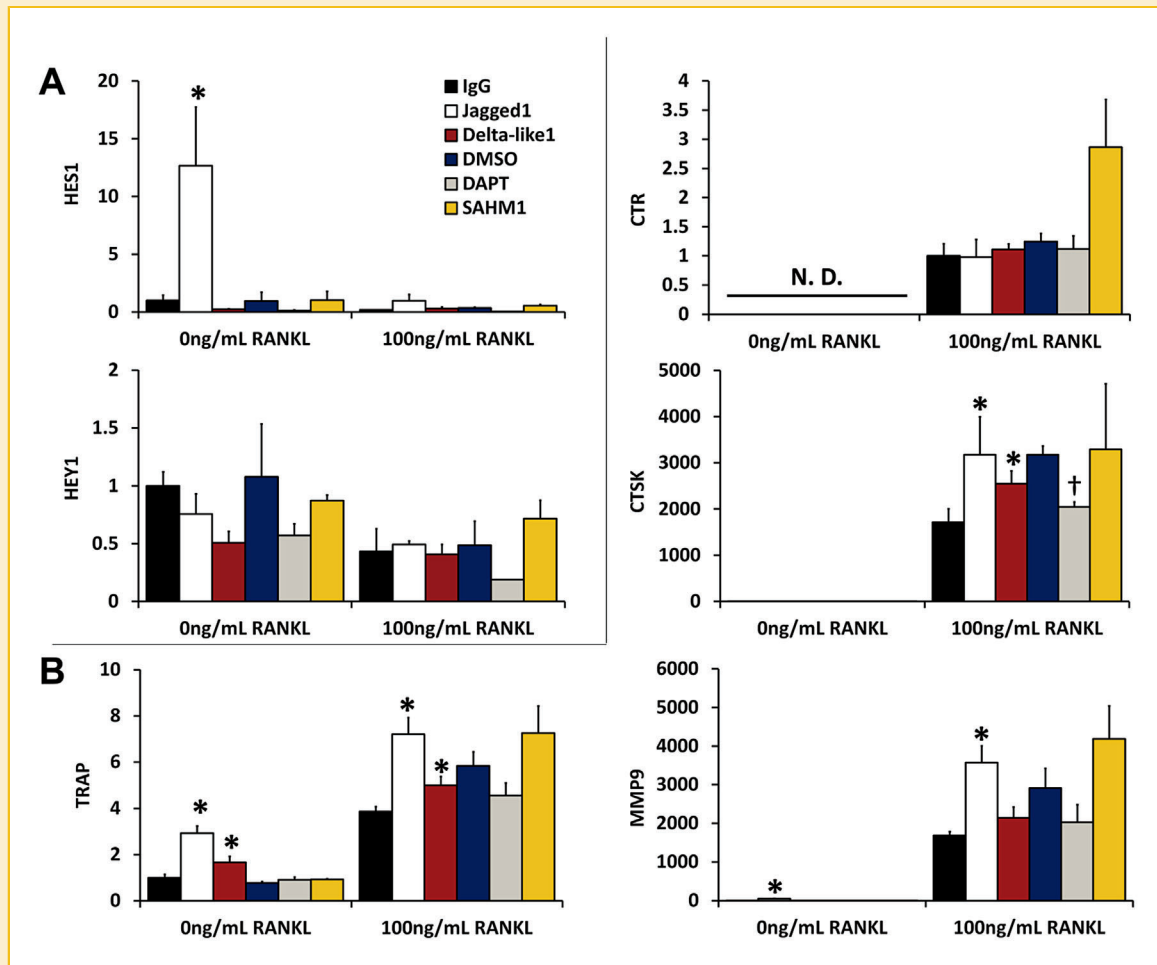


Fig. 2. Notch receptor and ligand expression during osteoclastogenesis. RNA was extracted from osteoclast precursors cultured with MCSF alone (0 Days) or with MCSF and 100ng/mL RANKL for 1, 2, or 3 days. Expression was measured via real-time RT-PCR. (A) Expression of Notch receptors (Notch1, Notch2, Notch3, and Notch4) relative to expression at 0 Days. (B) Expression of Notch ligands (Jagged1, Jagged2, Delta-like1, and Delta-like4) relative to expression at 0 Days. \* $P < 0.05$  versus 0 Days. Data are derived from three biological replicates.

#### QUANTIFICATION OF STAINED AREA, OSTEOCLAST NUMBER, OSTEOCLAST SIZE, AND NUCLEAR NUMBER

Following TRAP staining, wells were dried and plates were scanned at 4800DPI using a flatbed scanner. Using ImageJ (<http://rsb.info.nih.gov/ij/>), well images were converted to grayscale and black/white thresholded at brightness 0-110 of a 0-255 histogram. Using equal sized ellipse selections, the stained area fraction of each well was measured to generate stained area values.

Two fields (at 20 $\times$  magnification) per well showing the highest density of osteoclasts were imaged in color. Prior to analysis, images were automatically white-balanced using the adjustment layer>levels function of Adobe Photoshop. Osteoclasts with more than 3 nuclei were counted manually. Using ImageJ, images were converted to grayscale, automatically contrast enhanced, and black/white thresholded at brightness 0-190 of a 0-255 histogram. Using the Analyze Particles function, the median size of osteoclasts were



**Fig. 3.** Expression of Notch target and osteoclast genes under Notch signaling manipulation. RNA was extracted from osteoclast precursors cultured under Notch stimulation or inhibition with either MCSF only or MCSF and 100 ng/mL RANKL for 3 days. (A) Expression of Notch target genes (HES1 and HEY1). (B) Expression of osteoclast functional genes (Tartrate-resistant acid phosphatase, TRAP; Calcitonin receptor, CTR; Cathepsin K, CTSK; Matrix metalloproteinase 9, MMP9). \* $P < 0.05$  versus IgG; †,  $P < 0.05$  versus DMSO. Data are derived from three biological replicates. N.D., not detected.

determined in pixels and converted to square microliter using a microscope calibration standard. After all measurements were complete median osteoclast sizes for each group were averaged.

To quantify nuclear number, TRAP-stained wells were observed at 40 $\times$  magnification. Nuclei within the 20 largest osteoclasts per well were manually counted.

### RNA ISOLATION AND QUANTITATIVE RT-PCR

To isolate RNA from TRIzol lysates, 200  $\mu$ L chloroform (Fisher Scientific, BP1145-1) was added to each sample and shaken for 15 s. Samples were incubated at room temperature for 5 min and centrifuged at 12000g at 4°C for 15 min. Five hundred microliter of the top, aqueous layer was transferred to a new micro centrifuge tube and mixed with 500  $\mu$ L 100% ethanol. This mixture was applied to the columns of an RNeasy mini kit (Qiagen, 74106) and the remainder of the isolation was carried out according to the manufacturer's protocol. RNA concentration was determined using a NanoDrop spectrophotometer (Thermo Scientific).

To generate cDNA, 2  $\mu$ g RNA was used as template in a reverse transcriptase reaction using the SuperScript VILO cDNA synthesis kit (Life Technologies, 11754-050). The resulting cDNA solutions were diluted to an effective starting RNA mass of 100 ng (20-fold dilution when starting with 2  $\mu$ g).

Primers for real time PCR listed in Table 1 were designed using Primer-BLAST (<http://www.ncbi.nlm.nih.gov/tools/primer-blast/>) and synthesized by Integrated DNA Technologies (Table 1). For each reaction primer mix tubes and cDNA mix tubes were prepared. Primer tubes contained 5  $\mu$ L 2 $\times$  SYBR Select master mix (Life Technologies, 4472908), 2  $\mu$ L 2  $\mu$ M primer mix, and 3  $\mu$ L deionized water per reaction. Complementary DNA mixtures contained 5  $\mu$ L 2 $\times$  SYBR Select master mix, 2  $\mu$ L cDNA, and 3  $\mu$ L deionized water per reaction. Ten microliter each of primer mix and cDNA mix were added to the wells of a MicroAmp Fast Optical 96-well reaction plate (Life Technologies, 4346907). Plates were sealed with optical adhesive film (Life Technologies, 4311971) and run in an ABI 7500 real time PCR system using

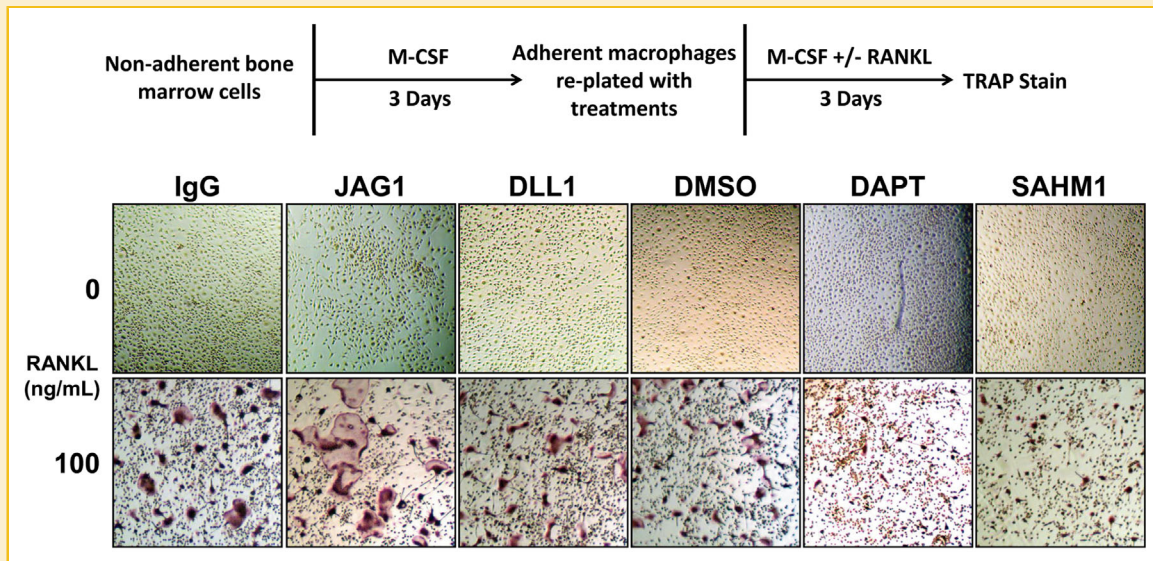


Fig. 4. Osteoclastogenic differentiation of adherent BMMs under Notch stimulation and inhibition. Non-adherent bone marrow cells were cultured with M-CSF for 3 days to generate adherent bone marrow-derived macrophages, which were lifted and re-plated into 24-well plates with indicated treatments. Osteoclast precursors were cultured in either MCSF alone or MCSF and 100ng/mL RANKL for 3 days under either Notch ligand stimulation or Notch signaling inhibition. At the conclusion of differentiation, cells were TRAP stained. Red/purple staining indicates TRAP activity.

the following program: (1) 50°C for 2 min, (2) 95°C for 2 min, (3) 95°C for 15 sec, (4) 60°C for 1 min, (5) repeat 3–4 40×. Single products were confirmed by determining melting curves at the conclusion of the reaction. Relative expression was calculated using the  $2^{-\Delta\Delta Ct}$  method normalized to 18S (endogenous loading control).

#### IN VITRO RESORPTION

Osteoclasts were differentiated from adherent bone marrow macrophages according to protocol (1) for 3 days in 24-well Osteo Assay plates (Corning, 3987) which contain a layer of mineral hydroxyapatite. Medium was refreshed every other day and osteoclasts were cultured for an additional 1 or 3 days to allow resorption to occur. Osteoclasts were removed from the plate at the conclusion of the resorption period by washing wells once with PBS and incubating for 5 min with 10% bleach followed by 2 washes with deionized water. To enhance contrast between non-resorbed and resorbed areas, wells were stained using a modified von Kossa protocol. Three microliter 5% silver nitrate (Midland Scientific Inc., 6828-16) was added to each well and incubated for 30 min at room temperature in the dark. Plates were then soaked in deionized water for 5 min. Stain was developed by adding 300  $\mu$ L 5% sodium carbonate (Fisher Scientific, S263-500) in formalin and incubating at room temperature for 4 min. Sodium carbonate solution was discarded and plates were dried at 50°C for 1 h.

Plates were scanned at 600DPI using a flatbed scanner. Using ImageJ, well images were converted to grayscale and black/white thresholded at brightness 0–110 of a 0–255 histogram. Using equal sized ellipse selections, the stained area fraction (stained area/measured area) of each well was measured to generate stained area values. Resorbed area was calculated as 100%–% stained area.

#### STATISTICAL ANALYSIS

Statistical significance was evaluated using 1-tailed student's *t*-tests for real time RT-PCR data and 2-tailed student's *t*-tests for all other data. *P*-values less than 0.05 were considered statistically significant.

## RESULTS

#### NOTCH RECEPTORS AND LIGANDS ARE EXPRESSED BY OSTEOCLAST PRECURSORS AND OSTEOCLASTS

Notch receptor (Fig. 2A) and ligand (Fig. 2B) gene expression at baseline and during osteoclastogenesis were measured via quantitative RT-PCR. Osteoclasts were differentiated from adherent bone marrow-derived macrophages (BMMs) with MCSF and RANKL, and RNA was collected after 1, 2, and 3 days of treatment. RNA samples from cells treated with MCSF alone for 3 days (0 days RANKL treatment) were also collected. Expression of all four Notch receptors was detectable throughout differentiation though expression of both Notch1 and Notch2 had decreased significantly by the end of the treatment (Fig. 2A). Among Notch ligands, expression of both Jagged1 (JAG1), and Jagged2 (JAG2) significantly decreased after 1 day of RANKL treatment and remained low for the remainder of the differentiation period (Fig. 2B). By contrast, Delta-like1 (DLL1) expression remained constant and Delta-like4 (DLL4) expression significantly increased after 2 and 3 days of RANKL treatment. Delta-like3 (DLL3) expression could not be detected at any time point (data not shown).

#### NOTCH SIGNALING MANIPULATION ALTERS EXPRESSION OF OSTEOCLAST FUNCTION-RELATED GENES

Following Notch stimulation and inhibition, mRNA levels of Notch-responsive genes and osteoclast functional genes were quantified via

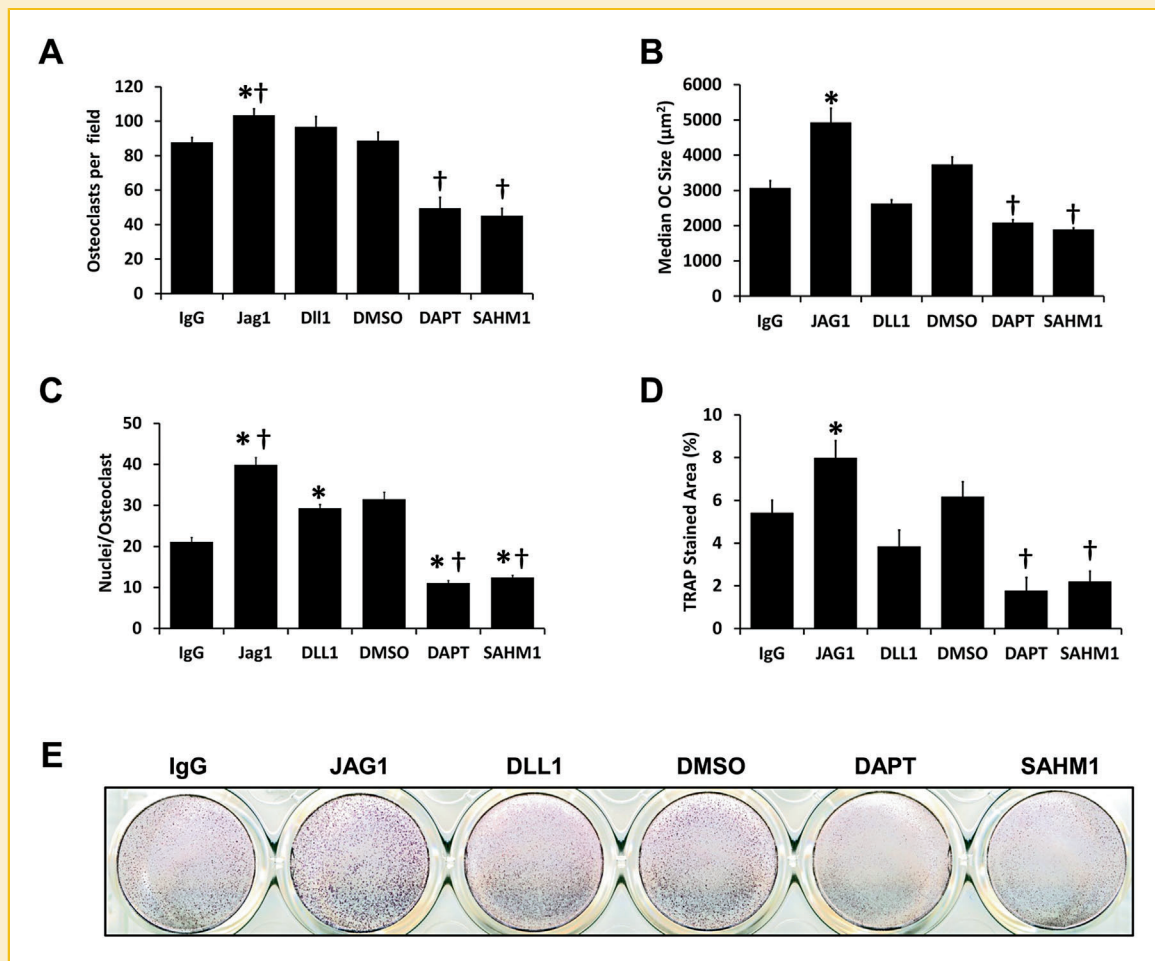


Fig. 5. Quantification of in vitro osteoclastogenesis parameters. (A) Mean number of osteoclasts per microscopic field. (B) Average of median osteoclast size in each visual field. (C) Mean number of nuclei per osteoclast. (D) Mean TRAP-stained areas. \* $P < 0.05$  versus IgG; †,  $P < 0.05$  versus DMSO. Data are aggregate of three independent experiments. (E) Representative image of TRAP-stained wells.

real-time quantitative RT-PCR. The Notch-responsive gene *Hes1*, but not *Hey1*, was upregulated by stimulation with JAG1, but not DLL1, in osteoclast precursors (Fig. 3A). Both JAG1 and DLL1 stimulation enhanced TRAP expression even in the absence of RANKL (Fig. 3B). JAG1 also significantly increased cathepsin K and MMP9 expression, where DLL1 increased only cathepsin K. With the exception of cathepsin K, which was significantly reduced by DAPT, there was no significant change in osteoclast functional genes under Notch signaling inhibition. No treatment altered expression of calcitonin receptor.

#### NOTCH SIGNALING ENHANCES OSTEOCLASTOGENESIS OF OSTEOCLAST PRECURSORS PRE-STIMULATED WITH RANKL

Adherent BMMs were differentiated to osteoclasts by seeding with MCSF and RANKL either onto JAG1- or DLL1-coated wells (control: IgG-coated wells) or with DAPT or SAHM1 (control: DMSO) (Fig. 4). Culture of cells on JAG1, but not DLL1, resulted in increased numbers of multinuclear osteoclasts; in contrast, Notch signaling inhibition with either DAPT or SAHM1 significantly reduced

osteoclast numbers (Figs. 4 and 5A). JAG1 stimulation resulted in larger osteoclasts, where Notch inhibition by both DAPT and SAHM1 resulted in smaller osteoclasts (Figs. 4 and 5B). In addition, both JAG1 and DLL1-stimulated cells had significantly increased numbers of nuclei where nuclear numbers were significantly reduced in DAPT and SAHM1 treated cells (Fig. 5C). Alterations in osteoclast size and number were reflected in measurements of TRAP-stained areas; JAG1-stimulated cells demonstrated significantly higher and DAPT- and SAHM1-treated cells demonstrated significantly lower stained areas compared to their respective controls (Fig. 5D and E). These results suggest that activation of Notch signaling in RANKL-stimulated osteoclast precursors enhances osteoclast differentiation.

#### NOTCH SIGNALING INHIBITS OSTEOCLASTOGENESIS OF NON-COMMITTED OSTEOCLAST PRECURSORS

To investigate context-dependent effects of Notch signaling on osteoclastogenesis, osteoclast precursors were differentiated under two additional conditions (Fig. 6). First, varying numbers of non-

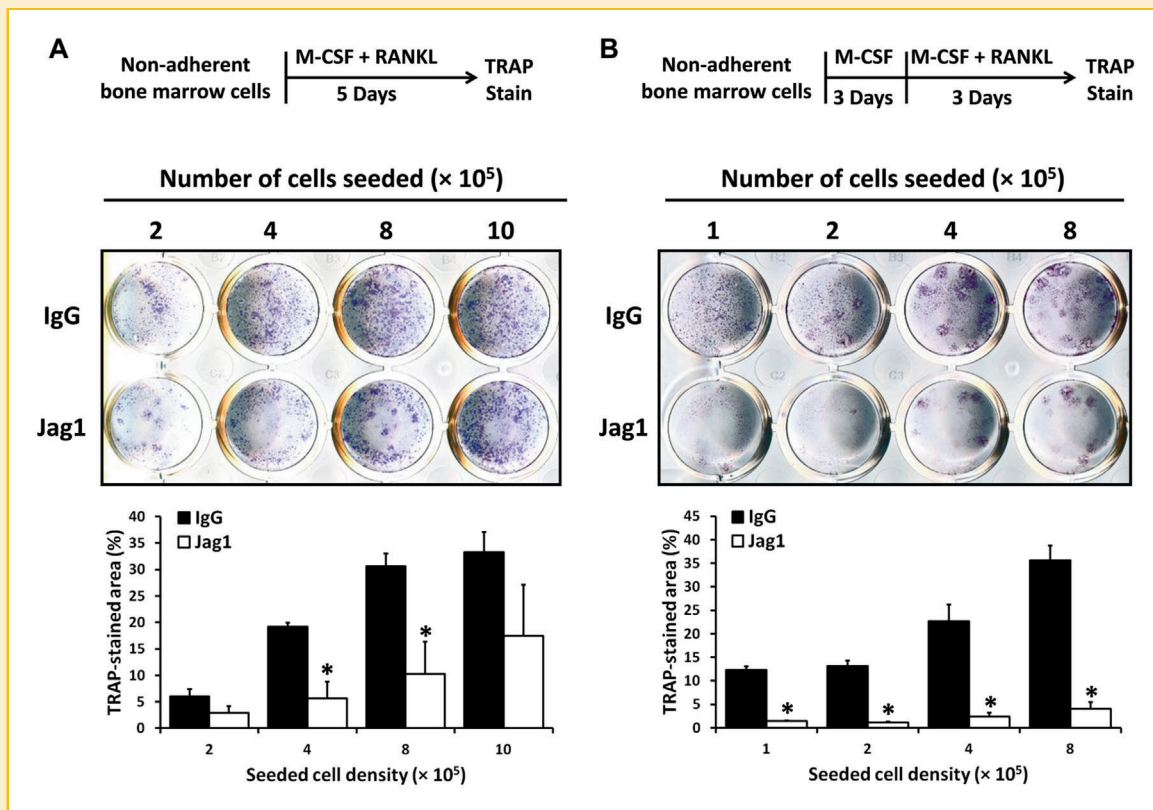


Fig. 6. Differentiation of osteoclasts from non-adherent bone marrow cells. (A) Osteoclasts were generated from varying seeding densities of non-adherent marrow cells on IgG- or JAG1-coated culture surfaces by culturing for 5 days with MCSF and RANKL. Following differentiation, cells were TRAP stained and TRAP-stained area was quantified. \* $P < 0.05$  versus IgG. (B) Osteoclasts were generated from varying seeding densities of non-adherent marrow cells on IgG- or JAG1-coated culture surfaces by first culturing 3 days with MCSF only followed by 3 days of MCSF and RANKL. Following differentiation, cells were TRAP stained and TRAP-stained area was quantified. \* $P < 0.05$  versus IgG. Each treatment was performed in duplicate. Images are representative and data are aggregate of 2 independent experiments.

adherent bone marrow cells were seeded with MCSF and RANKL into IgG- (control) or JAG1-coated wells. At the lowest density ( $1 \times 10^5$  cells), there was no significant difference in TRAP-stained areas between precursors cultured in IgG- or JAG1-coated wells (Fig. 6A). However, at intermediate densities (4 and  $8 \times 10^5$  cells) osteoclast differentiation was significantly higher in IgG-coated wells. At the highest density ( $10 \times 10^5$  cells), there were similar levels of osteoclastogenesis in IgG- and JAG1-coated wells.

Second, varying numbers of non-adherent bone marrow cells were seeded into IgG- or JAG1-coated wells with MCSF only and allowed to adhere and proliferate for 3 days prior to RANKL stimulation. Under this method, cells in IgG-coated wells demonstrated a greater amount of osteoclastogenesis regardless of seeding cell density (Fig. 6B). These results suggest that early activation of Notch signaling in osteoclast precursors suppresses osteoclastogenesis.

#### NOTCH SIGNALING ENHANCES OSTEOCLASTIC RESORPTION

Osteoclastic resorption of mineral surfaces was assessed under Notch signaling stimulation and suppression to determine whether alterations in osteoclast maturation translate to altered function.

Osteoclast precursors were cultured with and without RANKL on mineral-coated OsteoAssay surfaces under Notch stimulation with immobilized JAG1 or DLL1 or Notch inhibition with DAPT or SAHM1 for 4 (Fig. 7A) or 6 (Fig. 7B) days. After 4 days of culture, significant increases in resorption were evident in both JAG1 and DLL1-stimulated groups compared to IgG-coated wells, but there was not yet sufficient resorption in controls to assess effects of Notch inhibition (Fig. 7A). After 6 days of culture, resorption remained significantly higher in JAG1- and DLL1-coated wells compared to IgG control, and resorption under Notch-inhibition with both DAPT and SAHM1 was significantly reduced compared to DMSO control wells (Fig. 7B).

#### NOTCH SIGNALING MANIPULATION ALTERS EXPRESSION OF OSTEOCLAST FUSION GENES

The increases and decreases in nuclear number seen under Notch stimulation and inhibition, respectively, suggest that Notch signaling may contribute to the fusion of osteoclast precursors. To investigate whether Notch signaling might regulate fusion at the mRNA level, expression of established osteoclast fusion genes were measured in the context of Notch signaling stimulation (JAG1- and



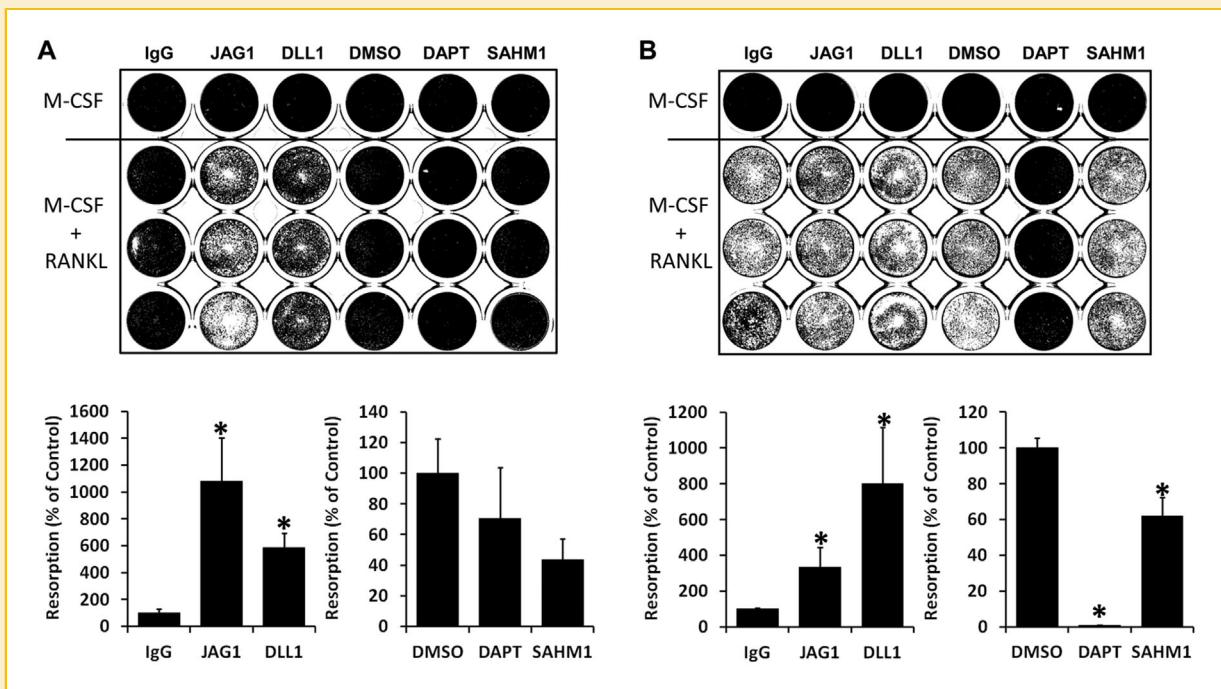


Fig. 7. Osteoclastic hydroxyapatite resorption under Notch signaling manipulation. Osteoclast precursors were cultured under Notch stimulation or inhibition with either MCSF only or MCSF and 100 ng/mL RANKL for either 4 or 6 days. At the conclusion of the culture period, cells were removed and remaining mineral was darkened via von Kossa stain. (A) Representative von Kossa-stained plate following 4 days of culture. (B) Quantification of hydroxyapatite resorption area following 4 days of culture. \* $P < 0.05$  versus IgG. (C) Representative von Kossa-stained plate following 6 days of culture. (D) Quantification of hydroxyapatite resorption area following 6 days of culture. \* $P < 0.05$  versus IgG or DMSO, respectively. Images are representative and data are aggregate of two independent experiments.

DLL1-coated wells; IgG-coated wells, control) and inhibition (DAPT and SAHM1 treatment; DMSO, control) (Fig. 8). Both CD200 and DC-STAMP were significantly upregulated by both JAG1 (64% and 41% increases over IgG) and DLL1 (85% and 13% increases over IgG) stimulation; similarly, Notch inhibition with DAPT significantly decreased expression of these genes where SAHM1 decreased only CD200 expression. v-ATPase V0 subunit d2 (ATP6V<sub>0</sub>D<sub>2</sub>), CD9, CD81, and signal regulatory protein  $\alpha$  (SIRPA) expression were not altered with Notch signaling manipulation.

## DISCUSSION

The role of Notch signaling in osteoclastogenesis has been controversial with evidence supporting both inhibitory and stimulatory roles. The findings of this study suggest that Notch signaling is necessary for the maturation of osteoclasts, but it suppresses the differentiation of macrophages that have not yet been stimulated with RANKL.

Genetic studies by Bai et al. suggested an inhibitory role for Notch signaling in osteoclastogenesis as compound knockout of Notch1, 2, and 3 resulted in enhanced osteoclastogenesis due to increased osteoclast precursor proliferation and enhanced response to RANKL [Bai et al., 2008]. In the present study, however, we have found that osteoclasts and their precursors also express Notch4; thus, the enhanced osteoclastogenesis seen in this model may be attributable

to signaling through Notch4 (Fig 2A). Yamada et al. similarly suggest an inhibitory role for Notch signaling in osteoclastogenesis in their experiments with immobilized DLL1 which demonstrated an inhibition of osteoclast formation and osteoclast precursor proliferation [Yamada et al., 2003]. However, the inhibitory effect of DLL1 has been challenged by the work of Sekine et al. who, using immobilized antibodies against specific Notch receptors and blocking antibodies against specific Notch ligands, demonstrated that Notch signaling initiated by DLL1, and Notch2 promote osteoclastogenesis and Notch signaling initiated by JAG1 and Notch1 suppress osteoclastogenesis [Sekine et al., 2012]. Our data agrees with that of Sekine et al. in that DLL1 increased osteoclast nuclear number and resorptive activity (Figs. 5C and 7).

In the above-described studies, while the exact details of the osteoclastogenesis protocols are not provided, a common aspect is that the osteoclast differentiation processes were concluded after 6 days of culture. In the present study, Notch signaling stimulation via immobilized JAG1 was carried out under three different osteoclastogenesis protocols: (1) seeding of adherent bone marrow-derived macrophages with MCSF and RANKL into coated plates wherein RANKL in the media stimulated RANK signaling prior to attachment to the ligand-coated surface (RANK first, Notch second), (2) seeding of non-adherent bone marrow cells with MCSF and RANKL into coated plates wherein cells become responsive to RANK and Notch signaling after they have differentiated into adherent macrophages (near-simultaneous stimulation of RANK and Notch), and (3) seeding of

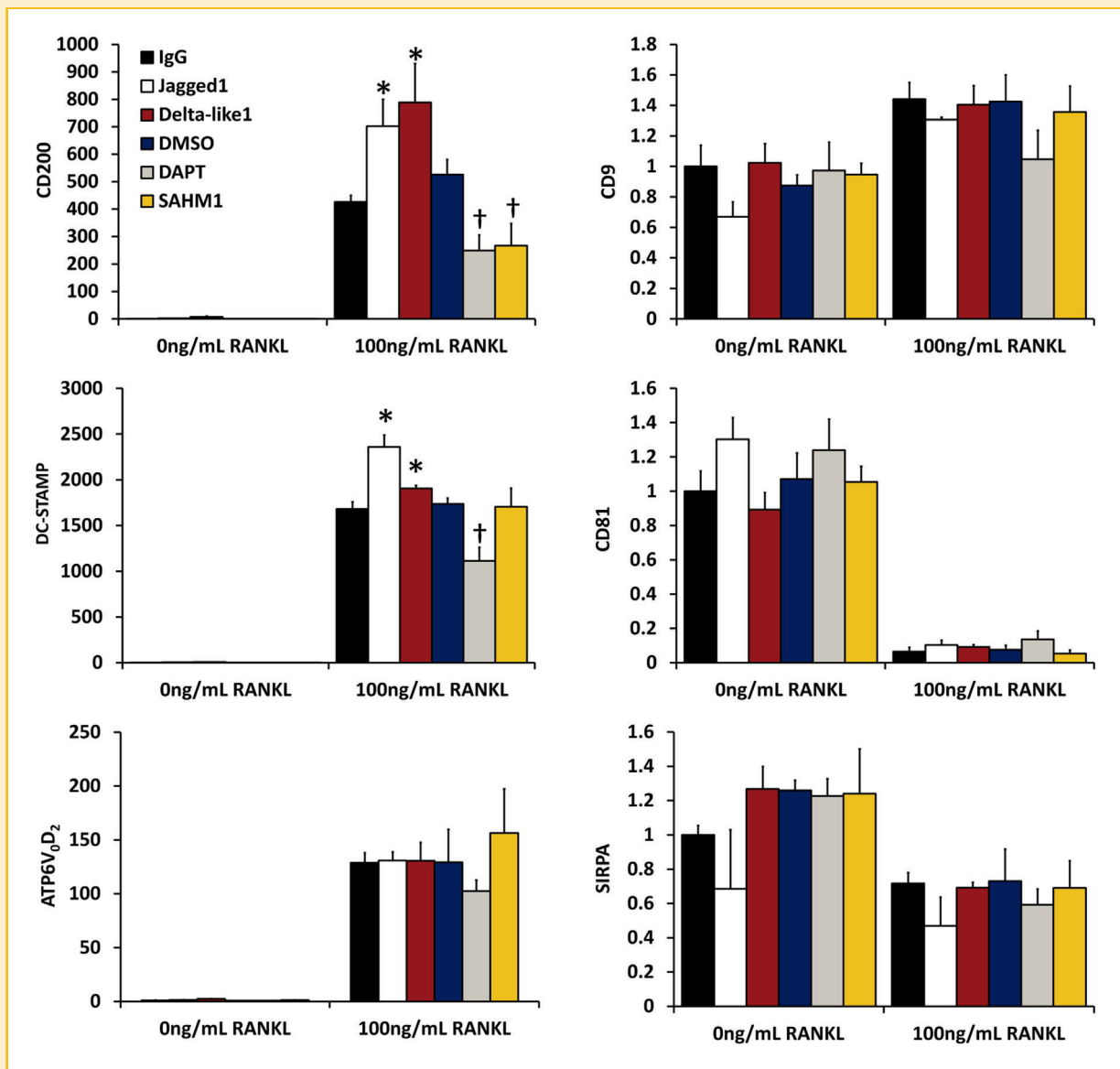


Fig. 8. Expression of osteoclast fusion genes under Notch signaling manipulation. RNA was extracted from osteoclast precursors cultured under Notch stimulation or inhibition with either MCSF only or MCSF and 100ng/mL RANKL for 3 days. Expression of osteoclast fusion genes CD200, DC-STAMP, v-ATPase  $V_0$  subunit  $d_2$  (ATP6V $_0$ D $_2$ ), CD9, CD81, Signal-regulatory protein alpha (SIRPA) were measured via qRT-PCR. \* $P < 0.05$  versus IgG; †,  $P < 0.05$  versus DMSO. Data are derived from three biological replicates.

non-adherent bone marrow cells with MCSF only into coated plates followed by 3 days of culture prior to the addition of RANKL (Notch first, RANK second). Under the first protocol, Notch stimulation resulted in a clear enhancement of osteoclastogenesis evidenced not only by increased number of nuclei per osteoclast and increased osteoclast size (in the case of JAG1) (Figs. 4 and 5), but also by increased mineral resorption (Fig. 7). The importance of Notch signaling in osteoclast maturation is further evidenced by suppression of osteoclast fusion and resorption by Notch inhibition with either DAPT or SAHM1.

Data from both Bai and Yamada suggest that the inhibitory effect of Notch signaling is due to suppression of osteoclast precursor proliferation [Yamada et al., 2003; Bai et al., 2008]. This is likely to be the case, as, when osteoclasts were differentiated according to the

second protocol, JAG1 appeared to decrease osteoclastogenesis, but this defect was rescued by seeding a higher density of cells (Fig. 6A). By contrast, JAG1 inhibited osteoclastogenesis when osteoclast precursors were cultured on coated surfaces for 3 days prior to RANKL stimulation regardless of cell density (Fig. 6B). These data are not only consistent with findings by others, but also provide evidence for divergent roles for Notch signaling in osteoclastogenesis that are governed by whether osteoclast precursors are exposed to RANKL before or after Notch activation. Such effects are not unusual as TNF exerts a similar behavior wherein it suppresses osteoclast differentiation from uncommitted osteoclast precursors, but enhances osteoclastogenesis if precursors are pre-stimulated with RANKL [Jules et al., 2010].

While there were significant alterations in the expression of some osteoclast genes under Notch signaling manipulation (Fig. 3), the magnitude of gene expression alteration was not great relative to the difference between cells treated with MCSF alone and those treated with both MCSF and RANKL. Furthermore, while osteoclasts were smaller and had fewer nuclei under Notch signaling inhibition, they were, nevertheless, TRAP-positive (Figs. 3 and 4). These findings suggest that Notch signaling does not largely contribute to the initial differentiation of osteoclasts, but rather osteoclast maturation. A role for Notch signaling in osteoclast maturation is further supported by increased expression of osteoclast fusion genes CD200 and DC-STAMP under Notch stimulation, and decreased expression of these genes under Notch inhibition (Fig. 8). From a clinical perspective, inhibition of osteoclast maturation without inhibition of initial osteoclast differentiation could be a useful and powerful approach to restore bone mass in osteoporosis or other disorders of bone loss, as small osteoclasts are deficient in resorption, but are nevertheless capable of stimulating osteogenesis [Henriksen et al., 2014].

Our final observation, which suggests future mechanistic investigative directions, stems from the distinct effects of DAPT and SAHM1. DAPT blocks Notch cleavage and liberation of NICD, and SAHM1 interferes with canonical transcription complex formation [Kornilova et al., 2003; Moellering et al., 2009]. In the case of osteoclast genes, fusion genes, and hydroxyapatite resorption, DAPT demonstrated a more robust inhibitory effect than SAHM1 (Figs., 7, and 8 5). The differences in intensity suggest that there are (1) targets of DAPT other than Notch signaling, such as ErbB4, involved in osteoclastogenesis and/or (2) elements of Notch signaling that are MAML-independent involved in osteoclastogenesis [Sardi et al., 2006; Besse et al., 2007]. It has been demonstrated that NICD1 and NICD2 interact with Calcium/calmodulin-dependent protein kinase IV and NF- $\kappa$ B subunit p65 respectively, and these interactions support osteoclastogenesis. These may be examples of MAML-independent Notch signaling that cannot be observed with Notch receptor knockout [Fukushima et al., 2008; Choi et al., 2013]. Future studies are necessary to understand these potentially important additional pathways that may interact with Notch signaling and what roles they play in osteoclast differentiation and function.

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## AUTHOR CONTRIBUTIONS

J.W. Ashley—designed and performed experiments, analyzed and interpreted data, and drafted manuscript; J. Ahn—designed

experiments, interpreted data, and edited manuscript; K.D. Hankenson—designed experiments, interpreted data, and edited manuscript.

## DISCLOSURES

J.W. Ashley: Nothing to disclose. J. Ahn: Unrelated research funding from Synthes. Consultant for Synthes and Merck. Part owner of Skelegen. K.D. Hankenson: Co-founder of Skelegen.

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