

Functional Roles of Netrin-1 in Osteoblast Differentiation

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Abstract. *Aim: Recent studies have demonstrated that netrin-1 plays a vital role in bone metabolism. Previous studies have shown that osteoblasts produce netrin-1 which affects osteoclast differentiation. However, the role of netrin-1 in osteoblast differentiation is not well understood. In this study, we explored the roles of netrin-1 in osteoblasts. Materials and Methods: Quantitative reverse-transcriptase polymerase chain reaction (qPCR), RNA interference for netrin receptors, the generation of netrin-1 plasmid, transfection of plasmids, and cell proliferation assay were performed. Results: During osteoblast differentiation by ascorbic acid, netrin-1 expression was significantly decreased. Gene expression related with osteoblast differentiation was down-regulated by netrin-1 treatment. We also found that osteoblast differentiation by bone morphogenetic protein-4 (BMP-4) was inhibited in the presence of recombinant netrin-1. Forced expression of both BMP-4 and netrin-1 significantly decreased alkaline phosphatase expression. On the other hand, Unc5b, neogenin, and A2b which belong to netrin receptors were expressed by osteoblasts. Moreover, alkaline phosphatase expression was significantly decreased by knockdown for the combination of two receptors among these receptors. Conclusion: Netrin-1 is involved in the regulation of osteoblast differentiation.*

Bone tissue is constantly being remodeled to maintain its homeostasis by osteoblasts and the resorption by osteoclasts

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(1). The function of bone tissue is regulated by several factors which other organs produce to communicate with bones. Indeed, the vessels and nerves inside bone are essential for bone development and remodeling (2, 3). Previous studies have demonstrated that neural and vascular regulation factors, such as semaphorins and netrins, orchestrate bone metabolism. We have demonstrated that semaphorin 3A, which is a diffusible axonal chemorepellent that has an important role in axon guidance, regulates bone remodeling indirectly by modulating sensory nerve development and also found that it enhances the elongation of dendrite in osteocytes (4, 5). Semaphorin 3A exerts an osteoprotective effect by both suppressing bone resorption and increasing bone formation (6).

Mammalian netrins including the secreted proteins (netrin-1, netrin-3, netrin-4 and netrin-5) and membrane-bound proteins (netrin-G1 and netrin-G2) are involved in the regulation of both axon guidance and angiogenesis (7). The netrin system comprises of seven receptors including neogenin (Neo1), deleted in colorectal carcinoma (DCC), Unc5A, Unc5B, Unc5C, Unc5D, adenosine 2b (A2b), and Down syndrome cell adhesion molecule (DSCAM) (7). Among these netrins, netrin-1 (Ntn1) has involved in cytoskeleton reorganization and in epithelial cell adhesion and migration in lung, mammary gland, and pancreas (8-10). Recent studies showed that Ntn1 plays a crucial role in osteoclast differentiation (11, 12). As the function of osteoclasts is regulated by Ntn1, the question whether Ntn1 affects cellular function in osteoblasts has been raised. However, little is known about the role of Ntn1 in osteoblasts. In this study, we explore whether Ntn1 affects the function of osteoblasts.

Materials and Methods

Cell culture, reagents and transfection. The mouse pre-osteoblastic cell line MC3T3-E1 cells which were kindly provided from Dr. Toru Ogasawara (Department of Oral and Maxillofacial Surgery, The University of Tokyo, Japan) were maintained in the medium,

consisting of α -modified minimum essential medium (α -MEM) (WAKO, Osaka, Japan) with 10% fetal bovine serum (FBS) (BioWest, Nuaille, France). For osteoblastic differentiation assay, cells were cultured in the medium containing ascorbic acid (50 μ g/ml) and β -glycerophosphate (10 mM). The medium was changed every 3 days. All cultures were maintained at 37°C in humidified air including 5% CO₂. Recombinant mouse Ntn1 (1109-N1-025) was purchased from R&D systems (Minneapolis, MN, USA). Cells were transfected with plasmids using Lipofectamine 2000 (Thermo Scientific, Wilmington, DE, USA).

Cell proliferation assay. The cells were incubated with recombinant Ntn1 at the indicated concentration for 3 days. The sample cells were quantified using a Cell Titer 96 Aqueous One Solution Cell Proliferation Assay (Promega, Madison, WI, USA), according to the manufacturer's instructions. The measurements are represented by the means of at least three independent experiments, with each data point based upon six replicates.

Measurement of alkaline phosphatase (ALP) activity. Cells were plated at a density of 2×10⁴ cells in 24-well plates. After reaching confluence, cells were incubated with ascorbic acid (50 μ g/ml) and β -glycerophosphate (10 mM) in the presence or absence of recombinant Ntn1 at the indicated concentration for 7 days. ALP activity was assayed (WAKO, Tokyo, Japan) as previously described. The measurements are expressed as the means of three independent experiments, with each data point based on four replicates.

Quantitative reverse-transcriptase polymerase chain reaction (qPCR). To validate changes in gene expression, quantitative real time RT-PCR (qPCR) analysis was performed using Applied Biosystems Prism 7900HT Sequence Detection System according to the manufacturer's instructions (Thermo Scientific, Wilmington, DE, USA) as previously described (13). Total RNA was extracted from cells using ISOGEN (Nippon Gene, Tokyo, Japan). We used SYBR Green-based detection. Reverse-transcriptase reaction was performed with High Capacity cDNA Reverse Transcription kit (Thermo Scientific, Wilmington, DE, USA). The cDNA was amplified by PCR using murine specific primers for *alkaline phosphatase (Alp)*, *bone gamma-carboxyglutamic acid-containing protein (Bglap)*, *Ntn1*, *BMP-4*, *collagen 1 alpha (Col1a)*, *DCC*, *Unc5a*, *Unc5b*, *Unc5c*, *Unc5d*, *A2b*, *Neol*, *DSCAM*, and *β -actin (Actb)* as follows: *Alp* sense, 5'-GCTTTAAACCCAGACACAAG-3' and *Alp* antisense, 5'-GCAGTAACCACAGTCAAGGT-3'; *Bglap* sense, 5'-TGCTTGTGACGAGCTATCAG-3' and *Bglap* antisense, 5'-GAGGACAGGGAGGATCAAGT-3'; *Ntn1* sense, 5'-GCAAGCTGAAGATGAACAGA-3' and *Ntn1* antisense, 5'-CTTTGTCTGGCCTTCAGGA-3'; *BMP-4* sense, 5'-GAGGAGTTTCCATCACGAAGA-3' and *BMP-4* antisense, 5'-GCTCTGCCGAGGAGATCA-3'; *Col1a* sense, 5'-CATGTTTCAGCTTTGTGGACCT-3' and *Col1a* antisense, 5'-GCACTGACTTCAGGGATGT-3'; *DCC* sense, 5'-GCTTTTGTCTCA GCCAGGAC-3' and *DCC* antisense, 5'-CGCTCAAGTCATCC TGTTCA-3'; *Unc5a* sense, 5'-ACCTCTGAGGCTGAGGACTTC-3' and *Unc5a* antisense, 5'-GGCTTGTGCAGAGTGAGGTAG-3'; *Unc5b* sense, 5'-CCATGGACCGGTACCTAAACT-3' and *Unc5b* antisense, 5'-TGGCTACCAGCATCTCACTCT-3'; *Unc5c* sense, 5'-AACTCTCTGGGGGTCACCT-3' and *Unc5c* antisense, 5'-TGCAGAGTGAGGATGACAGG-3'; *Unc5d* sense, 5'-CCTGAGTCTCTGA AGTACCTG-3' and *Unc5d* antisense, 5'-ATGTGGACTCATC CTCCACTG-3'; *A2b* sense, 5'-TCCCGCTCAGGTATAAAGGTT-3'

and *A2b* antisense, 5'-GGACACACCCAAAGAAGTTGA-3'; *Neol* sense, 5'-AGAGGGCATGAGTCAGAGGA-3' and *Neol* antisense, 5'-GCTGGAGTGGAAATGATGGT-3'; *DSCAM* sense, 5'-ATATCA GGCGCGAATAAACG-3' and *DSCAM* antisense, 5'-CCGTGATG TTCTCATTGC-3'; *Actb* sense, 5'-AGAAGGACTCCTATGTGG GTGA-3' and *Actb* antisense, 5'-CATGATCTGGGTTCAT CTTTTC-3'. SYBR green-based qPCR was performed using THUNDERBIRD SYBR qPCR Mix (TOYOBO, Osaka, Japan). Thermal cycling conditions were conducted as previously described (13). Values were normalized to *Actb* using the 2^{- $\Delta\Delta$ Ct} method. All samples were examined in triplicate assays.

Plasmids. Mouse Ntn1 (accession number NM_008744.2) and BMP-4 (accession number NP_031580.2) was obtained by a standard RT-PCR technique using PrimeSTAR HS DNA polymerase (TaKaRa, Ohtsu, Japan) and cloned into pcDNA3.1 expression vector (Thermo Scientific, Wilmington, DE, USA) as previously described (14).

Small interfering RNA (siRNA) transfection. RNA interference was performed as previously described (13). Stealth Select RNAi siRNA for murine Ntn1 (MSS207200, MSS207201, MSS207202) and Stealth RNAi Negative Control Duplex (Low GC, Medium GC) as si-control (si-CTL) were purchased from Thermo Scientific (Wilmington, DE, USA). Briefly, they were all transfected with Lipofectamine RNAi MAX (Thermo Scientific, Wilmington, DE, USA) according to the manufacturer's instructions. Results were representative of more than three individual experiments.

Statistical analysis. Comparisons between two groups were analyzed using Student's *t*-tests and comparisons among three groups were analyzed using One-Way Analysis of Variance and Bonferroni/Dunn methods ([#]*p*<0.05, ^{##}*p*<0.01). All values are represented as the mean± S.E.M. Results are representative examples of more than three independent sets of experiments.

Results

Ntn1 inhibits osteoblast differentiation in MC3T3-E1. First, we assessed Ntn1 expression during osteoblast differentiation by qPCR analysis. As shown in Figure 1A, *Ntn1* mRNA expression was decreased at 7 and 14 days after induction of ascorbic acid and β -glycerophosphate. To assess whether treatment of Ntn1 affects osteoblast differentiation, we added recombinant mouse Ntn1 to the culture of conditioned medium. collagen 1 alpha (Col1a), an early marker of osteogenic differentiation, was down-regulated in a dose-dependent manner in the presence of Ntn1 (Figure 1B). Ntn1 inhibited ALP activity of MC3T3-E1 (Figure 1C). To confirm that the decrease of gene expression related with osteogenic differentiation is not due to the inhibition of cell proliferation, we assessed whether Ntn1 affects cell proliferation in MC3T3-E1 cells. Ntn1 did not affect cell proliferation at all in MC3T3-E1 cells (Figure 1D).

To estimate the function of Ntn1 in osteogenic differentiation by induction of bone morphogenetic proteins (BMP), we overexpressed Ntn1 and/or BMP-4 in MC3T3-E1 cells (Figure 2A). Addition of recombinant Ntn1 significantly

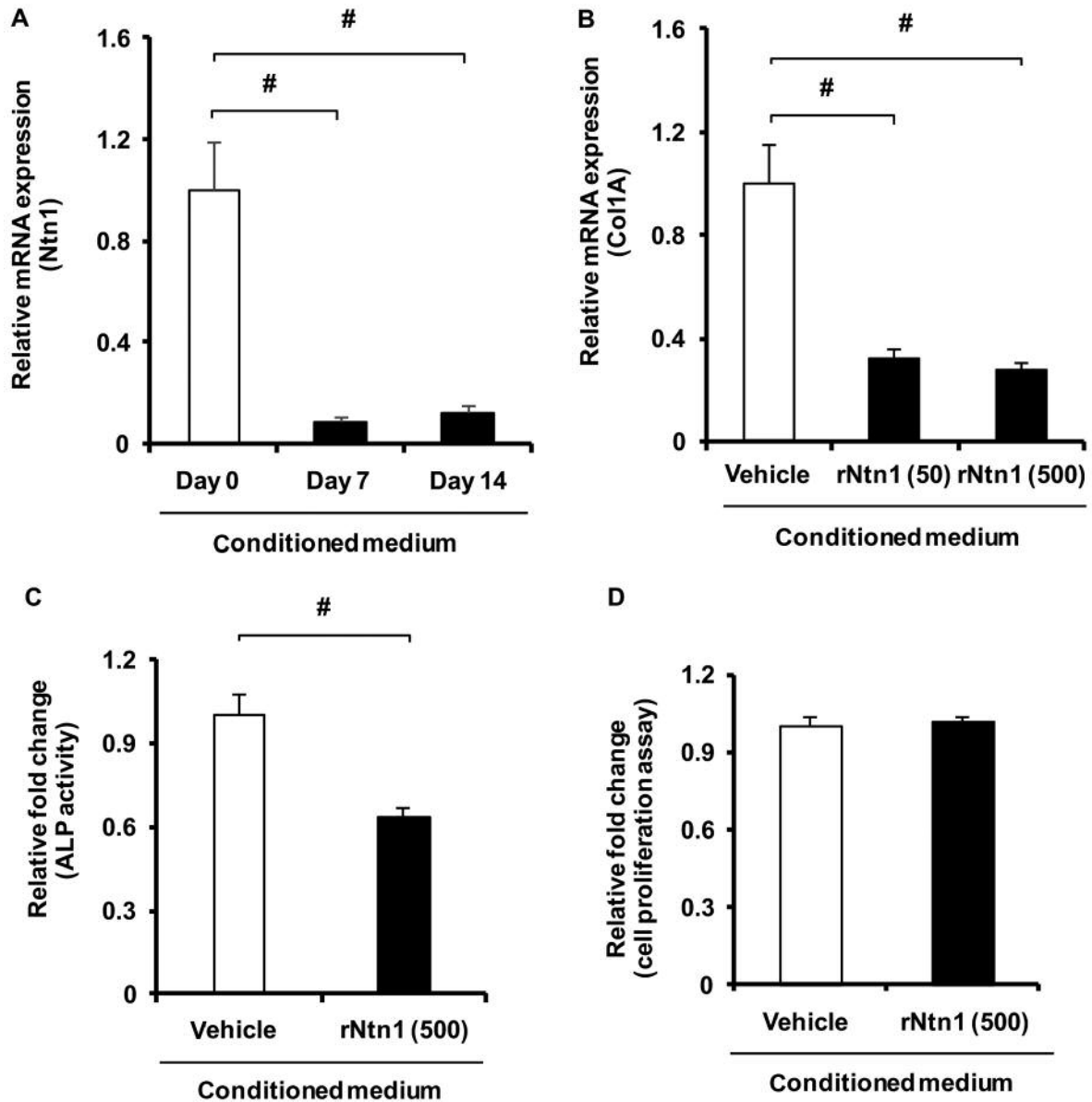


Figure 1. *Ntn1* inhibits osteoblast differentiation. A. Expression of *Ntn1* mRNA in MC3T3-E1 during osteoblast differentiation in the presence of ascorbic acid and β -glycerophosphate at 7 and 14 days estimated by qPCR analysis. B. Expression of *Col1a* mRNA in MC3T3-E1 with recombinant *Ntn1* (50 and 500 ng/ml) for 3 days estimated by qPCR analysis. C. Effect of recombinant *Ntn1* (500 ng/ml) on ALP activity of MC3T3-E1 at 7 days estimated by ALP activity assay. D. Effect of recombinant *Ntn1* (500 ng/ml) on cell proliferation of osteoblastic cells at 3 days estimated by cell proliferation assay. Data are calculated from three repeated experiments. Recombinant *Ntn1*, rNtn1. # $p < 0.05$. Data are expressed as the means \pm S.E.M.

decreased Alp mRNA level in BMP-4 transfected MC3T3-E1 in a dose dependent manner (Figure 2B). Forced expression of *Ntn1* significantly decreased both Alp and *Col1a* mRNA level (Figure 2C). Furthermore, simultaneous overexpression of both BMP-4 and *Ntn1* also significantly inhibited osteoblast differentiation (Figure 2C). These results suggest that *Ntn1* inhibits osteoblast differentiation.

Three netrin receptors are involved in the regulation of osteoblast differentiation by *Ntn1*. Next we assessed whether MC3T3-E1 cells express netrin receptors. As shown in Figure 3A-H, *Neo1*, *A2b*, and *Unc5b* were strikingly expressed by MC3T3-E1 compared to murine brain. Then we validated whether knockdown of these three receptors affect osteoblast differentiation in the presence of *Ntn1*. We confirmed that

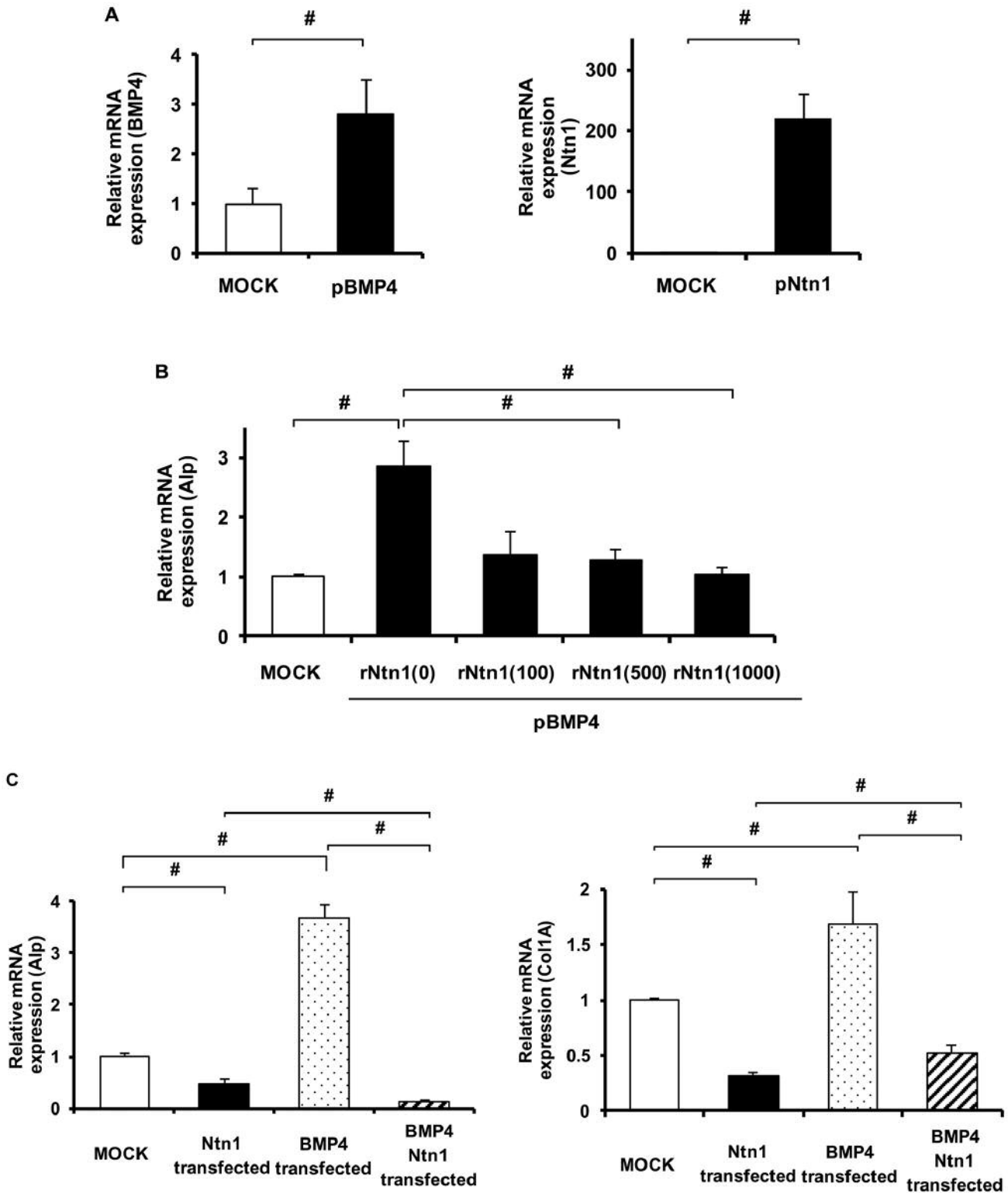


Figure 2. *Ntn1* overexpression inhibits osteoblast differentiation. A. MC3T3-E1 cells were transfected with *Ntn1* plasmid, BMP-4 plasmid, or an empty vector at 3 days estimated by qPCR analysis: (left panel) pBMP4, (right panel) pNtn4. B. Expression of *Alp* mRNA in BMP-4 transfected MC3T3-E1 with recombinant *Ntn1* (0, 100, 500 and 1,000 ng/ml) for 3 days estimated by qPCR analysis. C. Expression of *Alp* and *Col1a* mRNA in BMP-4 and/or *Ntn1* transfected MC3T3-E1: (upper panel) *Alp*, (lower panel) *Col1a*. Plasmid *Ntn1*, pNtn1; plasmid BMP-4, pBMP4; empty vector, MOCK; Recombinant *Ntn1*, rNtn1. #*p*<0.05. Data are expressed as the means±S.E.M.

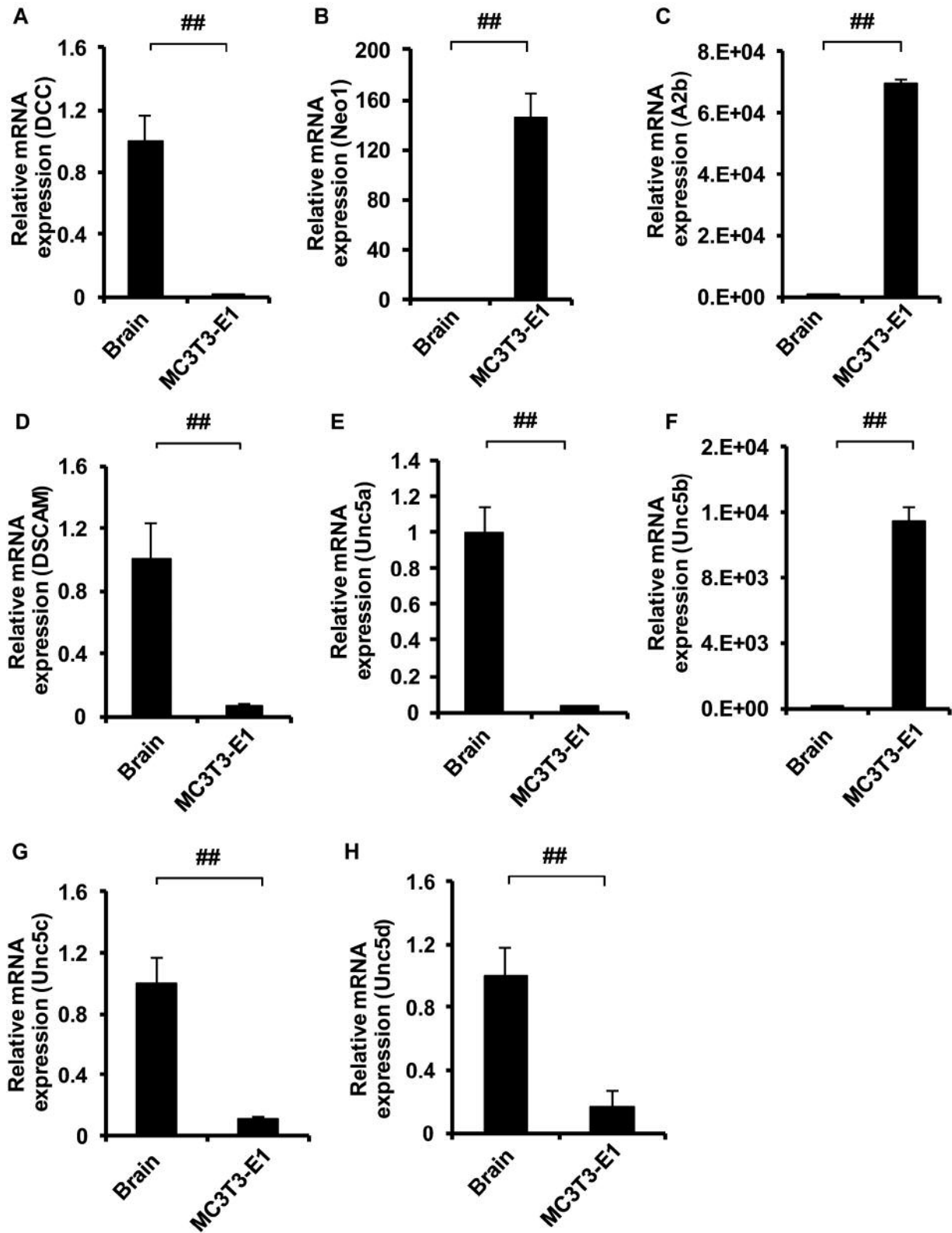


Figure 3. Netrin receptors mRNA expression in MC3T3-E1. Expression of netrin receptors in MC3T3-E1 and murine brain estimated by qPCR analysis A. DCC mRNA. B. Neo1 mRNA. C. A2b mRNA. D. DSCAM mRNA. E. Unc5a mRNA. F. Unc5b mRNA. G. Unc5c mRNA. H. Unc5d mRNA. # $p < 0.05$, ## $p < 0.01$. Data are expressed as the means \pm S.E.M.

single knockdown of each receptor (*i.e.* siRNA for A2b, Unc5b, Neo1) was successfully achieved (Figure 4A-C). Ntn1 inhibited *Alp* mRNA levels in siRNA for control transfected cells (Figure 4D). RNA interference of one receptor (A2b, Unc5b, Neo1) did not rescue the inhibition of differentiation by Ntn1 (Figure 4E-G). However, the combination of two among these three receptors recovers the inhibition of differentiation by Ntn1 (Figure 4H-J). These results suggest that the combination of two receptors is essential for regulating osteoblast differentiation by Ntn1.

Discussion

The present study demonstrated that Ntn1 plays a vital role for differentiation in osteoblasts. Osteoblast differentiation is partly controlled by bone morphogenetic protein (BMP) (15). Recent studies show that BMP is the key molecule to connect netrin with osteoblastic differentiation. Neogenin which belongs to one of the netrin receptors is involved in the regulation of BMP-induced Smad signaling and endochondral bone formation (16). Their studies provide evidence that neogenin may regulate chondrocyte maturation by promoting BMP induced BMP receptor association with lipid rafts, thus enhances effective BMP receptor concentration or BMP binding affinity and increases Smad phosphorylation and Runx2 induction by using neogenin-deficient mice, suggesting that neogenin promotes chondrogenesis *in vitro* and *in vivo*. On the other hand, this controversial result has been already reported. Hagihara *et al.* reported that neogenin negatively regulates the functions of BMP and that this effect of neogenin is mediated by the activation of RhoA (17). However, it is conjectured that the inhibition of osteoblast differentiation by BMP via the activation of neogenin may not occur under the physiological condition.

Togari *et al.* showed that Ntn1 is expressed by osteoblasts (18). Yagami *et al.* found that the expression of Ntn1 was decreased in the process whereby mesenchymal cells differentiate into osteoblasts by BMP (19). It has been reported that Ntn1 is involved in regulating bone remodeling. Mediero *et al.* demonstrated that Ntn1, which is produced by osteoclasts enhanced osteoclast differentiation by an autocrine/paracrine manner and Ntn1 deficient mice have markedly diminished osteoclasts, as well as increased cortical and trabecular bone density and volume compared to wild-type mice, suggesting that Ntn1 is a negative regulator in bone metabolism (11). In contrast, Maruyama *et al.* reported that Ntn1 suppresses osteoclast multinucleation, but not osteoclast differentiation and protected the mice against autoimmune bone destruction *in vivo*, indicating that Ntn1 is a positive regulator in bone metabolism (12). These researchers reported that Ntn1 does not affect osteoblast differentiation. As they use primary cells derived from murine calvaria which consist of a heterogeneous cell population, their results are different from our findings.

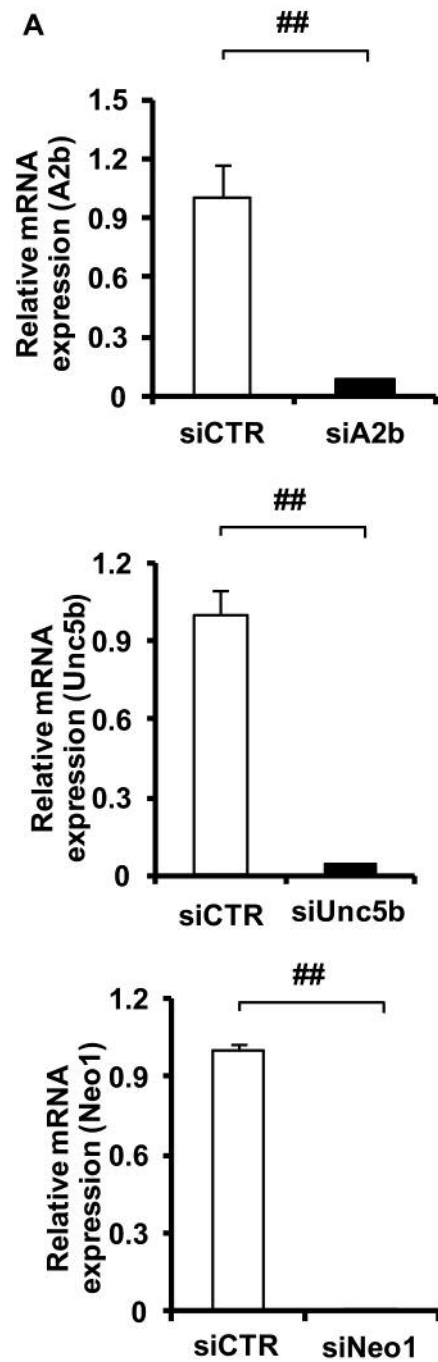


Figure 4. Continued

Ntn1 and its receptors are involved in regulating differentiation of various organs. Ntn1 is effective in regulating the expression of markers of neuroectodermal differentiation in human embryonal carcinoma cells (20). In mammary gland, Ntn1 facilitates functional differentiation of mammary epithelial cells (21). In terms of differentiation

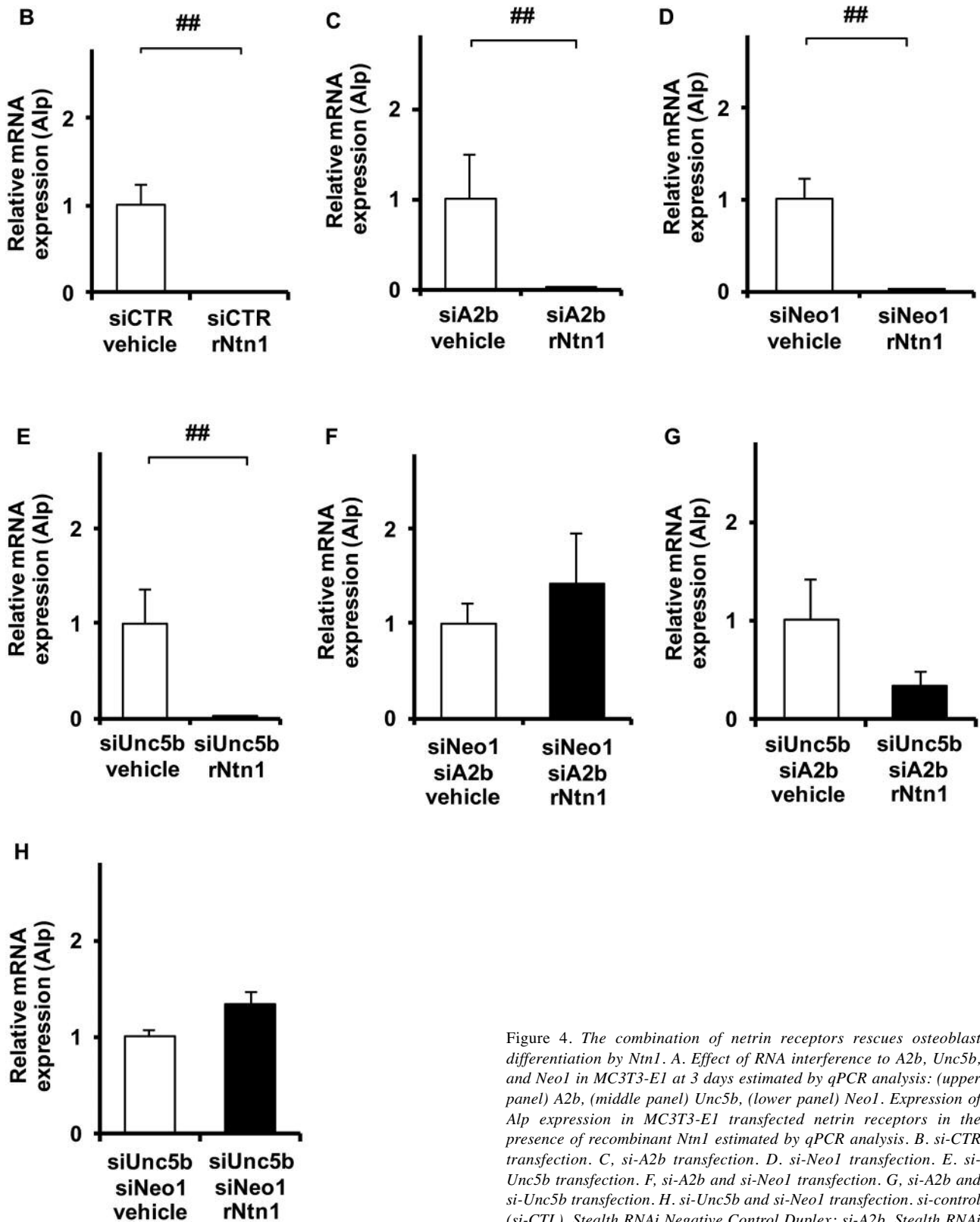


Figure 4. The combination of netrin receptors rescues osteoblast differentiation by Ntn1. A. Effect of RNA interference to A2b, Unc5b, and Neo1 in MC3T3-E1 at 3 days estimated by qPCR analysis: (upper panel) A2b, (middle panel) Unc5b, (lower panel) Neo1. Expression of Alp expression in MC3T3-E1 transfected netrin receptors in the presence of recombinant Ntn1 estimated by qPCR analysis. B. si-CTR transfection. C. si-A2b transfection. D. si-Neo1 transfection. E. si-Unc5b transfection. F. si-A2b and si-Neo1 transfection. G. si-A2b and si-Unc5b transfection. H. si-Unc5b and si-Neo1 transfection. si-control (si-CTL), Stealth RNAi Negative Control Duplex; si-A2b, Stealth RNAi to A2b; si-Neo1, Stealth RNAi to Neo1; si-Unc5b, Stealth RNAi to Unc5b # $p < 0.05$, ## $p < 0.01$. Data are expressed as the means \pm S.E.M.

potential, Ozmadenci *et al.* showed that murine induced pluripotent stem cells which derived with recombinant Ntn1 turn off the endogenous Oct4/GFP promoter activity (22). Neogenin overexpression favors inner cell mass differentiation (23). In placenta, a discrete pattern of distribution of Ntn1 and its two receptors, Unc5B and DCC were observed (24). Moreover, Xu *et al.* demonstrated that negative regulation of N- α -acetyltransferase 10 towards Ntn1 and its receptor Unc5B were also detected upon treatment of retinoid acid that is used to induce morphological differentiation (25).

We have previously demonstrated that Ntn4, derived from vascular endothelial cells inhibits osteoclast differentiation (13). We speculate that netrin family may play a vital role for bone metabolism. Further investigations are required to clarify the role of netrins in bone metabolism.

In conclusion, we found that Ntn1 inhibits osteoblast differentiation and the combination of two netrin receptors is essential for regulating osteoblast differentiation by Ntn1.

Acknowledgements

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