Effect of glucosamine, a therapeutic agent for osteoarthritis, on osteoblastic cell differentiation

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Received February 4, 2011; Accepted March 22, 2011

DOI: 10.3892/ijmm.2011.686

Abstract. Osteoarthritis (OA) is characterized by qualitative and quantitative changes in the architecture and composition of all the joint structures. Glucosamine (GlcN) has been used to treat OA in humans, because GlcN is present in the cartilage tissues as a component of glycosaminoglycans, and exhibits the symptom-modifying effect on OA by normalizing cartilage metabolism. On the other hand, the pathological change of subchondral bone is associated with the initiation and progression of cartilage damage in OA. However, the effect of GlcN on bone metabolism remains unsolved. In the present study, we determined the effect of GlcN on bone metabolism (osteoblastic cell differentiation) using mouse osteoblastic MC3T3-E1 cells by evaluating the expression of early (type I collagen and alkaline phosphatase), middle (osteopontin) and late (osteocalcin and mineralization) stage differentiation markers, and further compared its effects to those of N-acetyl-D-glucosamine (GlcNAc), a derivative of GlcN. The results indicated that the mineralization of mature osteoblasts was increased by treatment with GlcN and GlcNAc. Furthermore, reverse transcription-polymerase chain reaction (RT-PCR) analyses revealed that GlcN and GlcNAc substantially increased the expression of a middle stage marker and a late stage marker, although they did not essentially affect the expression of early stage markers. In addition, GlcN and GlcNAc suppressed the expression of receptor activator of NF-κB ligand (RANKL), a key factor involved in the osteoclastic cell differentiation and activation. Together these observations suggest that both GlcN and GlcNAc may have a potential not only to induce osteoblastic cell differentiation especially at middle-late stages, but also to suppress the osteoclastic cell differentiation, thereby possibly increasing bone matrix deposition and decreasing bone resorption, and eventually modulating bone metabolism in OA.

Introduction

Osteoarthritis (OA) is the most common joint disorder with an immense socioeconomic impact. OA is characterized by qualitative and quantitative changes in the architecture and composition of all the joint structures (1,2). An altered imbalance between the biosynthesis and the degradation of matrix components leads to a progressive destruction of the tissue (3). Currently, several treatments are available for OA ranging from conservative measures to more surgical extremes. Conservative measures involve lifestyle modifications, physical therapy and pharmacological treatment with nonsteroidal anti-inflammatory drugs and intra-articular injection of hyaluronic acid (4). Treatments for early OA are primarily based on symptomatic relief, whereas irreversible joint disability in advanced OA usually requires surgical intervention to relieve pain and improve joint function (5). However, current treatments are mostly targeting the symptoms and not addressing the destructed structure of articular cartilage in OA.

D-Glucosamine (GlcN), a naturally occurring amino monosaccharide, is present in the connective and cartilage tissues as a component of glycosaminoglycans, and contributes to maintaining the strength, flexibility and elasticity of these tissues. According to the biochemical and pharmacological findings, the administration of GlcN normalizes cartilage metabolism, so as to inhibit the degradation (6) and stimulate the synthesis of proteoglycans (7,8), and to restore articular function. Thus, GlcN has been widely used to treat OA in humans for more than two decades (9-12). In this context, several short- and long-term clinical trials in OA have shown the significant symptom-modifying effect of GlcN (13-15). Moreover, the updated Osteoarthritis Research Society International (OARSI) recommendations for management of hip and knee OA have suggested that GlcN has symptom-relieving and structure-modifying effects in knee OA (16). In addition, we have previously demonstrated that GlcN is able to suppress cytokine-induced synovial cell activation (such as nitric oxide-, PGE2- and IL-8-production), thereby possibly exhibiting anti-inflammatory actions in arthritis (17).

On the other hand, it is known that the pathological change of subchondral bone is associated with the initiation...
and progression of cartilage damage in OA (18). Due to non-physiological load-bearing to the subchondral bone, the bone resorption increases, thereby enhancing the progression of cartilage damage in OA (19). However, the effect of GlcN, a therapeutic agent for OA, on bone metabolism is not fully understood. To address this, in the present study, we examined the effect of GlcN on the bone metabolism using mouse osteoblastic MC3T3-E1 cells by examining osteoblastic cell differentiation, and compared it with that of N-acetyl-D-glucosamine (GlcNAc), a derivative of GlcN.

Materials and methods

Reagents. GlcN, GlcNAc, alizarin red-S (AR-S) and paraformaldehyde were purchased from Wako Pure Chemical Industries Ltd. (Osaka, Japan). Dexamethasone and ascorbic acid were purchased from Sigma-Aldrich (MO, USA). β-glycerophosphate was purchased from Calbiochem (CA, USA).

Cells. Mouse osteoblastic MC3T3-E1 cells, which were established from newborn mouse calvaria, were purchased from RIKEN Cell Bank (RCB 1126, Tsukuba, Japan). The MC3T3-E1 cells were maintained in α-minimum essential medium (Gibco Invitrogen, Carlsbad, CA, USA) containing 10% heat-inactivated fetal bovine serum (FBS), penicillin and streptomycin at 37°C in 5% CO₂. To induce osteoblastic differentiation of MC3T3-E1 cells, the cells were fed with differentiation medium (DM: normal medium supplemented with 500 nM dexamethasone, 10 mM β-glycerophosphate and 50 mg/ml ascorbic acid). The medium was changed every third day (20,21).

AR-S staining. MC3T3-E1 cells (1x10⁵ cells/well) were seeded in 6-well plates overnight. After changing the medium to DM, the cells were incubated in the absence or presence of GlcN (0.1 or 1 mM) or GlcNAc (1 mM) for 21 days. The incubation media were changed every third day with further addition of GlcN or GlcNAc. After the incubation, the cells were washed twice with ice-cold phosphate-buffered saline (PBS; 137 mM NaCl, 2.7 mM KCl, 8.1 mM Na₂HPO₄, 1.5 mM KH₂PO₄, pH 7.4), and fixed with 2% paraformaldehyde for 15 min at 4°C. The cells were then washed twice with ice-cold PBS and once with ice-cold distilled water. After the washing, the AR-S staining solution (40 mM AR-S, pH 4.2 adjusted by 1% KOH solution) was added to each well (1 ml/well), followed by incubation for 15 min at room temperature with gentle rocking. The cells were then washed five times with distilled water, followed by rinsing with PBS for 15 min with gentle rocking to reduce non-specific AR-S staining. Stained cells were photographed, and then lysed in 10% hexadecylpyridinium chloride (Nacalai Tesque, Kyoto, Japan)-10 mM sodium phosphate-buffer (pH 7.0) for 1 h at room temperature with gentle rocking. For quantification, the concentration of AR-S in each sample was determined by absorbance at 562 nm (20-22).

Semi-quantitative reverse transcription-polymerase chain reaction (RT-PCR). MC3T3-E1 cells (2x10⁶ cells/dish) were seeded in 6 cm plates overnight. After changing the medium to DM, the cells were incubated in the absence or presence of GlcN (0.1 or 1 mM) or GlcNAc (1 mM) up to 21 days. The incubation media were changed every third day with further addition of GlcN or GlcNAc. After the incubation, the cells were collected by a cell scraper (Sumitomo Bakelite Co., Ltd., Tokyo, Japan). Then, total-RNA was purified using an RNasea plus mini kit (Qiagen, CA, USA) and QIAshredder (Qiagen) to remove contaminating DNA, according to the manufacturer's protocol, and stored at -80°C. RT-PCR was performed using a ReverTra-Plus RT-PCR kit (Toyobo, Osaka, Japan), and PCR amplification was performed with the GoTaq Master mix (Promega, WI, USA) in a thermal cycler (GeneAmp PCR System 9700, Applied Biosystems, CA, USA) for type I collagen (COLI), alkaline phosphatase (ALP), osteopontin (OPN), osteocalcin (OCN), receptor activator of NF-κB ligand (RANKL) and for glyceraldehyde-3-phosphate dehydrogenase (GAPDH), according to the manufacturer's protocol. In brief, cDNA was synthesized by reverse transcription of total RNA (500 ng) using ReverTra Ace reverse transcriptase and oligo(dt)20. To discriminate mRNA-derived PCR products from genomic DNA-derived products, the intron-spanning PCR primers were used with the annealing temperature and cycle number indicated in Table I. PCR products were resolved by 2% agarose gel electrophoresis in 1X Tris-acetate-EDTA buffer and stained with ethidium bromide. In our preliminary experiments, we tried to semi-quantitatively detect mRNA by using different numbers of PCR cycles. The results revealed that the amounts of RT-PCR products increased as the cycle number increased. Thus, we decided to measure the mRNA levels by RT-PCR with the number of cycles indicated in Table I.

<table>
<thead>
<tr>
<th>Genes</th>
<th>Forward primer</th>
<th>Reverse primer</th>
<th>Annealing temperature (°C)</th>
<th>Number of cycles</th>
</tr>
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<tr>
<td>COL1a</td>
<td>5'-acctatggcatcatcagc-3'</td>
<td>5'-gcaacacagtggcaatgag-3'</td>
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<td>27</td>
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<tr>
<td>ALPb</td>
<td>5'-gactgtcatctggagattg-3'</td>
<td>5'-gcttcatccttgggtaatg-3'</td>
<td>55</td>
<td>31</td>
</tr>
<tr>
<td>OPNc</td>
<td>5'-ctgctgctctgctggc-3'</td>
<td>5'-ctgtgctctgctgctggc-3'</td>
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<td>28</td>
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<td>5'-ctgtgctctgctgctggc-3'</td>
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<td>26</td>
</tr>
<tr>
<td>RANKL</td>
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<td>5'-ctgtgctctgctgctggc-3'</td>
<td>58</td>
<td>37</td>
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<td>60</td>
<td>20</td>
</tr>
</tbody>
</table>

aSaito et al (20); bAli et al (23); cToyobo.

Table I. Gene specific PCR primers, annealing temperature (°C) and number of PCR cycles.
Table I. The detected bands were quantified using MultiGauge (Version 3.0, FujiFilm, Tokyo, Japan).

**Statistical analysis.** Data are expressed as the mean ± SD, and analyzed for significant differences by a Student's t-test using Excel. Differences were considered statistically significant at P<0.05.

**Results**

**Effects of GlcN and GlcNAc on the mineralization of osteoblastic MC3T3-E1 cells.** Mineralization of differentiated osteoblastic MC3T3-E1 cells was measured after the incubation with GlcN or GlcNAc for 21 days. GlcN significantly increased the mineralization at 0.1 (P<0.05) and 1 mM (P<0.01) compared with control without GlcN and GlcNAc (Fig. 1). Moreover, GlcNAc significantly increased the mineralization at 1 mM (P<0.01) (Fig. 1). Notably, GlcN was more potent than GlcNAc in inducing mineralization at the same concentration (Fig. 1).

**Effects of GlcN and GlcNAc on the osteoblastic differentiation of MC3T3-E1 cells.** Osteoblastic cells express differentiation markers, such as COL1 and ALP which are markers for the early stage of osteoblastic differentiation, OPN, a marker for middle stage differentiation, and OCN, a marker for late stage differentiation (24-29). Thus, the effect of GlcN on osteoblastic cell differentiation was investigated by measuring the expression of differentiation markers by RT-PCR. First, the expression of the markers for the early stage of osteoblastic differentiation (COL1 and ALP) was evaluated. The expression of COL1 was not changed by GlcN (0.1 and 1 mM) after incubation for 1-3 days, compared to the control without GlcN and GlcNAc (Fig. 2). In contrast, the expression of COL1 was slightly, but significantly, decreased by GlcNAc (1 mM) after 2 days of incubation (Fig. 2). Furthermore, the expression of ALP was not substantially changed by either GlcN or GlcNAc (Fig. 2).

The expression of OPN, a marker for middle stage differentiation, was also evaluated. The expression of OPN was not changed by GlcN nor by GlcNAc after incubation for 7 days. However, the expression was apparently increased by both GlcN and GlcNAc (1 mM) after the incubation for 21 days (Fig. 4), although the change was not statistically significant. Furthermore, the expression of a marker for late stage of OCN was evaluated. Of importance, the expression of OCN was significantly (P<0.05) increased by GlcN (1 mM) after the incubation for 21 days (Fig. 5). The expression of OCN was also increased by GlcNAc after 21 days, although the change was not significant.

**Effects of GlcN and GlcNAc on the expression of RANKL by MC3T3-E1 cells.** Bone tissue is continuously remodeled through the combined activities of two cell types, osteoblasts and osteoclasts. Osteoblasts deposit bone matrix, whereas osteoclasts resorb bone matrix. The bone deposition and resorption are tightly coupled, and their balance defines both the mass as well as the quality of the bone tissue (30-32). RANKL, which is expressed and secreted by mature osteoblasts, is a key factor of osteoclastogenesis (33,34). Thus, the expression of RANKL was evaluated during osteoblastic differentiation of MC3T3-E1 cells by RT-PCR. Of note, the expression of RANKL was significantly (P<0.05) suppressed...
Figure 3. Effects of GlcN and GlcNAc on the mRNA expression of alkaline phosphatase (ALP). Osteoblastic cells (MC3T3-E1) were differentiated in differentiation medium in the absence (Control) or presence of GlcN (0.1 and 1 mM) or GlcNAc (1 mM) for 3 days. After the incubation, total RNA was purified, and the expression of mRNA for ALP was evaluated by RT-PCR. (A) PCR products were resolved by 2% agarose gel electrophoresis, and stained with ethidium bromide. (B) ALP mRNA expression is expressed as a ratio relative to Control. Data represent the mean ± SD of three independent experiments. Values were compared between the absence (Control) and presence of GlcN or GlcNAc.

Figure 4. Effects of GlcN and GlcNAc on the mRNA expression of osteopontin (OPN). Osteoblastic cells (MC3T3-E1) were differentiated in differentiation medium in the absence (Control) or presence of GlcN (0.1 and 1 mM) or GlcNAc (1 mM) for 21 days. After the incubation, total RNA was purified, and the OPN mRNA expression was evaluated by RT-PCR. (A) PCR products were resolved by 2% agarose gel electrophoresis, and stained with ethidium bromide. (B) OPN mRNA expression is expressed as a ratio relative to Control. Data represent the mean ± SD of three independent experiments. Values were compared between the absence (Control) and presence of GlcN or GlcNAc.

Figure 5. Effects of GlcN and GlcNAc on the mRNA expression of osteocalcin (OCN). Osteoblastic cells (MC3T3-E1) were differentiated in differentiation medium in the absence (Control) or presence of GlcN (0.1 and 1 mM) or GlcNAc (1 mM) for 21 days. After the incubation, total RNA was purified, and the OCN mRNA expression was evaluated by RT-PCR. (A) PCR products were resolved by 2% agarose gel electrophoresis, and stained with ethidium bromide. (B) OCN mRNA expression is expressed as a ratio relative to Control. Data represent the mean ± SD of three independent experiments. Values were compared between the absence (Control) and presence of GlcN or GlcNAc. *P<0.05.

Figure 6. Effects of GlcN and GlcNAc on the mRNA expression of RANKL. Osteoblastic cells (MC3T3-E1) were differentiated in differentiation medium in the absence (Control) or presence of GlcN (0.1 and 1 mM) or GlcNAc (1 mM) for 21 days. After the incubation, total RNA was purified, and the RANKL mRNA expression was evaluated by RT-PCR. (A) PCR products were resolved by 2% agarose gel electrophoresis, and stained with ethidium bromide. (B) RANKL mRNA expression is expressed as a ratio relative to Control. Data represent the mean ± SD of three independent experiments. Values were compared between the absence (Control) and presence of GlcN or GlcNAc. *P<0.05.
by both GlcN and GlcNAc (1 mM) after incubation for 21 days (Fig. 6).

**Discussion**

OA is characterized by qualitative and quantitative changes in the architecture and composition of all the joint structures (1,2). GlcN has been used to treat OA in humans, because it is present in the cartilage tissues as a component of glycosaminoglycans (35), and exhibits symptom-modifying effects on OA by normalizing cartilage metabolism (6-8,13-15).

On the other hand, the pathological change of subchondral bone is implicated in the initiation and progression of cartilage damage in OA (18). Due to non-physiological load-bearing to the subchondral bone, the bone resorption increases, thereby enhancing the progression of OA (19). However, the effect of GlcN on bone metabolism remains unsolved.

There are a few reports on the effects of GlcN on bone metabolism. First, a GlcN-containing diet has been shown to increase the bone mineral density of the femur in mice (36); however, the involved mechanism was not clarified. Second, GlcN treatment was reported to increase the ALP activity in NOS-1 cells, a human osteoblastic cell line (37). Third, GlcN treatment was found to increase OCN secretion from MG-63 cells, a human osteoblastic cell line, whereas collagen synthesis was not affected by GlcN (38). However, in these studies, only a few markers were used for evaluating the effect of GlcN on bone metabolism. Moreover, the effect of GlcNAc, a derivative of GlcN, has never been assessed.

In this study, we determined the effect of GlcN on bone metabolism (osteoblastic cell differentiation) using mouse MC3T3-E1 osteoblasts by evaluating the expression of differentiation markers of the early (COLI and ALP), middle (OPN) and late (OCN and mineralization) stages, and further compared the GlcN effects with those of GlcNAc. The results indicate that mineralization of mature osteoblasts is increased by treatment with GlcN after incubation for 21 days (Fig. 1). Furthermore, RT-PCR analyses revealed that GlcN treatment substantially increased the expression of a middle stage marker (OPN) and a late stage marker (OCN) after incubation for 21 days, whereas it did not essentially affect the expression of early stage markers (COLI and ALP) (Figs. 2-5). In addition, GlcN treatment suppressed the expression of RANKL, which regulates the differentiation and activation of osteoclasts, cells involved in the resorption of bone matrix (Fig. 6). Together these observations likely suggest that GlcN not only induces osteoblastic cell differentiation especially at the middle-late stages, but also suppresses osteoclastic cell differentiation, thereby increasing bone matrix deposition and decreasing bone resorption to modulate the bone formation.

Furthermore, the present results demonstrate that GlcNAc, a derivative of GlcN, as well as GlcN is able to increase the mineralization of mature osteoblasts and the expression of a middle stage marker (OPN) and a late stage marker (OCN) for osteoblastic cell differentiation, and to suppress the expression of RANKL, a key factor involved in the osteoclastic cell differentiation and activation (Figs. 1-6). These observations likely suggest that not only GlcN, but also GlcNAc, has a potential to induce osteoblastic cell differentiation and to suppress osteoclastic cell differentiation, thereby promoting bone formation (Fig. 7). In support of this finding, it has been recently reported that oral administration of GlcNAc increases bone mineral density in C57BL/6J mice, possibly by promoting bone formation and suppressing bone resorption (49).

In conclusion, the present study revealed that GlcN and GlcNAc increase the mineralization of mature osteoblasts and the expression of middle and late stage markers (OPN and OCN, respectively) during osteoblastic differentiation, and reduce the expression of RANKL, a differentiation and activation factor of osteoclasts. Together these observations likely suggest that both GlcN and GlcNAc may have a potential to induce osteoblastic cell differentiation, and to suppress osteoclastic cell differentiation, thereby increasing bone matrix deposition and decreasing bone resorption to modulate bone metabolism in OA. However, the effect of GlcN and GlcNAc on bone metabolism in OA should be carefully evaluated in the future.
Acknowledgements
This study was partially supported by the Research Institute for Diseases of Old Age, Juntendo University School of Medicine.

References