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Induction of an Osteocyte-like Phenotype by Fibroblast Growth Factor-2

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Abstract

The purpose of this study was to characterize the molecular phenotype that occurs during the profound morphological shift of cultured osteogenic cells upon treatment with fibroblast growth factor-2 (FGF2). A time course of treatment with FGF2 was performed on an osteoblast cell line, primary bone marrow stromal cells and an osteocyte-like cell line. Morphologic changes were recorded, and gene profiling was carried out by real time PCR. By 8 hours of FGF2 treatment, there is a striking morphological shift of osteoblast and stromal cells to an elongated dendritic-like morphology that is reminiscent of osteocytes. In osteoblasts treated with FGF2, this morphologic shift is preceded by an induction of several osteocyte markers, including dentin matrix protein 1 (>20-fold) and E11 (>5-fold). There is a transient increase in the gene expression of sclerostin (3.5-fold) and PHEX (2.5-fold). Sclerostin regulation by FGF2 is complex, as gene expression becomes markedly inhibited by FGF2 at times points after 8 hours of treatment before rebounding at day 12. Analogous modulation of osteocyte markers is seen in bone marrow stromal cells and MLO-Y4 osteocyte-like cells. In conclusion, this study shows that FGF2 can regulate the transition of osteogenic cells towards the osteocyte lineage, as well as, regulate the expression of critical genes in osteocytes.

Keywords (<6)

Osteoblast; Bone Marrow Stromal Cell; Osteocyte; Dentin Matrix Protein 1; Sclerostin; Connexin43

1. Introduction

Fibroblast growth factor-2 (FGF2) plays an important role in bone formation and osteoblast function (Reviewed in [1;2]). Deletion of the FGF2 gene in mice leads to reduced bone mass as well as an impaired ability to respond to the osteoanabolic effects of PTH [3;4;5;6]. In a transgenic mouse model, overexpression of the 18kDa-isoform of FGF2 in cells of the osteogenic lineage markedly enhances bone mineral density [7]. The short-term, systemic administration of FGF2 has been shown to increase bone formation in rats [8;9]. FGF2 has been shown to restore bone mass in ovariectomized rats [10;11]. Moreover, FGF2 accelerates fracture repair in multiple animal models [12;13;14]. The mechanism by which FGF2 enhances bone formation has long been hypothesized to be via the recruitment and

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expansion of osteoprogenitor cell populations rather than via enhancement of osteoblast differentiation. Indeed, for the most part, *in vitro* analysis of the effects of FGF2 on osteoblasts and osteoprogenitors has led to the conclusion that FGF2 is antagonistic to osteoblast differentiation. FGF2 has potent mitogenic effects on osteoprogenitors and markedly downregulates the expression of markers of osteoblasts, such as alkaline phosphatase, type I collagen and osteocalcin [15;16]. However, the effects of FGF2 are stage specific, as treatment of mature osteoblasts with FGF2 can have opposing outcomes from those arising from FGF2 treatment of immature osteoblasts [17].

The terminally differentiated cell of the osteoblast lineage is the osteocyte (Reviewed in [18;19;20]). These cells are the most abundant cell type in bone and are found in the newly deposited osteoid or embedded in the mineralized skeleton. Osteocytes have a dendritic-like morphology and lack many of the key molecular markers of their preceding osteoblasts, including greatly reduced expression of type I collagen and alkaline phosphatase (ALP). Further, these cells are unlikely to be able to mineralize. Osteocytes are of great importance to skeletal physiology, as data continues to accumulate to implicate these cells in the mechanical and hormonal responsiveness of bone, as well as, phosphate homeostasis and the regulation of Wnt signaling. Osteocytes are thought to be vital to the orchestration of bone remodeling. However, osteocytes have proven to be difficult to study due to the fact that they are embedded in the mineralized skeleton and have low mitogenic potential. Importantly, little is known about the factors that regulate the transition of osteoblasts to osteocytes. In this study, we explore the hypothesis that FGF2 can induce an osteocyte-like phenotype in osteogenic cells, and regulate the gene expression patterns of osteocytes.

2. Materials and methods

2.1. Cell culture

MC3T3-E1 clone 4 (MC4) cells were obtained from the American Type Culture Collection (Manassas, VA). These cells were maintained in α MEM media (Cellgro, Herndon, VA), supplemented with 10% fetal bovine serum (Hyclone, Logan, UT), penicillin (50 IU/ml) and streptomycin (50 μ g/ml). Mouse primary bone marrow stromal cells were prepared as described previously [21]. MLO-Y4 cells were kindly provided by Lynda Bonewald (University of Missouri, Kansas City) and were cultured as described [22]. For FGF2 treatments, cells were cultured in α MEM containing 1% fetal bovine serum, 50 μ g/ml ascorbic acid and 3 mM sodium phosphate for 24 hours prior to the addition of FGF2 (10 ng/ml). FGF2 was purchased from Millipore (Temecula, CA). The vehicle diluent for FGF2 (phosphate buffered saline, 0.1% bovine serum albumin, 1mM dithiothreitol) was used as a negative control for FGF2 treatments. For long-term cultures, media with or without FGF2 were replaced every 2–3 days. Cell viability was routinely assessed as described previously [23] and was statistically unaffected among treatment groups.

2.2. Crystal violet staining and microscopy

Following the indicated treatments, cells were fixed in 4% paraformaldehyde for 15 minutes, prior to staining with crystal violet (0.1% crystal violet in 10% ethanol) for 20 minutes. Cells were observed on a Nikon Eclipse 50i microscope (Melville, NY) and images captured with a Photometrics Coolsnap CCD camera.

2.3. Proliferation assay

MC4 cells were seeded in 24 well plates at 20,000 cells/well. Cells were treated with or without FGF2 for 24 hours. Cell numbers were assessed with the Cell Counting Kit-8 assay (Dojindo, Rockville, MD) or Cell Titer-Glo assay (Promega, Madison, WI). The results of three experiments performed in triplicate wells were averaged for analysis.

2.4. Migration assay

MC4 cells (6,000 cells/well) were seeded into the upper chamber of a Corning transwell chamber insert (8.0µm pore size; Corning, NY). FGF2 (10ng/ml) containing medium was added to the lower chamber. Medium lacking FGF2 added to the bottom chamber was used for comparison of migration. The cells were allowed to migrate for 4 or 24 hours. The upper side of the transwell filter was cleaned of cells with a cotton swab. The remaining cells were stained with crystal violet, as above. The number of cells migrating to the underside of the filter was counted. The results of three experiments performed in triplicate wells were averaged together.

2.5. RNA isolation and real time PCR

RNA was extracted from cultured cells at the indicated time points, as we have previously reported [24]. Quantitation of gene expression by real time PCR was performed as described previously [25]. The primers used for PCR are listed in Table I.

2.6. ALP activity assay

ALP activity was determined from MC4 cells treated with or without FGF2 for 4, 8 and 12 days by monitoring the conversion of p-nitrophenol phosphate to p-nitrophenol as described previously [26]. ALP activity was normalized to protein content for each well.

2.7. Statistical analysis

All experiments were performed in triplicate wells and repeated at least three times. Results are expressed as means \pm standard deviations. Data were assessed for statistical significance by ANOVA followed by a Tukey's post hoc test. P-values less than 0.05 are considered statistically significant.

3. Results and discussion

We observed that treatment of MC4 osteoblasts with FGF2 (10 ng/ml) induces a distinct change in cellular morphology (Fig. 1A). Within 8 hours of FGF2 treatment, MC4 cells begin to take on a dendritic-like morphology with development of long cell processes that are reminiscent of osteocytes. By 24 hours, long cellular processes are seen to extend from the cell body of FGF2 treated MC4 cells. While the occasional cell with this osteocyte-like morphology is seen in MC4 cells that have not been treated with FGF2, nearly every cell in the FGF2 treated culture has adopted this morphology by 24 hours. Further, the osteocyte-like morphology remains persistent for at least 12 days, as long as FGF2 remains in the culture media. The cells maintain this osteocyte-like morphology even at high densities. Notably, in the MC4 cells used in this study, FGF2 has only a modest effect on cell proliferation (Fig. 1B) and no significant impact on cell migration (Fig. 1C). The limited impact of FGF2 on the proliferation of these cells may not be too surprising. MC4 cells are considered a differentiated osteoblast with a high mineralizing capacity [27], and FGF2 has diverse effects on osteoblasts dependent upon their differentiation stage [17].

Given the striking morphological changes that were reminiscent of osteocytes, we examined the expression of osteocyte marker genes in these cells. We wanted to test the hypothesis that FGF2 was not functioning to inhibit osteoblast differentiation but rather that it was enhancing progression towards an osteocyte-like phenotype. In fact, others and we have shown that treatment of MC4 osteoblasts with FGF2 can enhance osteocalcin gene expression, a marker of mature osteoblasts and osteocytes [23;28]. To our knowledge, no one has looked to see if FGF2 could induce the expression of osteocyte markers. Indeed, osteocytes, like osteoprogenitor cells, are characterized by their low expression of both ALP and type I collagen relative to osteoblasts [22;29;30;31]. Consistent with previous studies in

osteoblasts [15;16], we observe a rapid and sustained decrease in the expression of COL1A1 and ALP gene expression caused by FGF2 treatment of MC4 cells as assessed by real time PCR (Fig. 2A and B). Likewise, ALP activity assays reveal an FGF2-dependent decrease in alkaline phosphatase enzymatic activity (Fig. 2C). We observe a rapid (within 2 hours) and sustained (up to 12 days) elevation in dentin matrix protein 1 (DMP1), with expression levels as high as 22-fold greater than those observed in MC4 cells cultured in the absence of FGF2 (Fig. 2D). DMP1, the acid phosphoprotein of the SIBLING (small, integrin binding ligand, N-linked glycoprotein) family, is expressed in osteocytes and pre-osteocytes, but not osteoblasts, *in vivo* [32]. DMP1 is present in the pericellular bone matrix surrounding osteocytes as well as along their cellular processes. Underscoring its physiologic relevance in osteocyte and skeletal biology, genetic ablation of DMP1 in mice results in osteomalacia and hypophosphatemia, as well as, defective osteocyte differentiation and a disorganized osteocyte lacuno-canalicular network [33].

Similarly, another important osteocyte marker, E11, is increased 2- to 5-fold in MC4 osteoblasts upon FGF2 treatment (Fig. 2E). This increase is statistically significant by 4 hours and remains elevated for the first two days. Like DMP1, E11 is associated with the dendritic processes of osteocytes and short interfering RNA mediated knockdown of E11 has been shown to block the growth of these cell processes [34]. Consistent with the notion that E11 (and perhaps DMP1) may be involved in dendrite formation, the expression of these genes precedes the formation of the dendritic cell processes, which become prominent only at ~8 hours of exposure to FGF2. The FGF2-induced expression of E11 is no longer statistically significant at 8 and 12 days of culture. As E11, is known to be an early marker of osteocyte differentiation, this may suggest that these cells continue to progress towards terminal differentiation while exposed to FGF2.

While not specifically a marker of osteocytes, the importance of the gap junction protein connexin43 in osteocytes is well documented [35;36;37]. Connexin43 gene expression is modestly upregulated following FGF2 treatment in MC4 cells (Fig. 2F). Similarly, there is a transient increase in the expression of the osteocyte-associated gene PHEX (phosphate-regulating gene with homologies to endopeptidases on the X chromosome) at 4 and 8 hours after FGF2 treatment (Fig. 2G). Like DMP1, PHEX plays an important role in the skeletal control of phosphate homeostasis [38].

Next, we examined the late osteocyte marker, sclerostin (SOST). SOST is a Wnt antagonist that serves as a negative regulator of bone formation [39]. Following FGF2 treatment of MC4 cells, SOST gene expression is complex (Fig. 2H). At 2 hours, there is a 3.5-fold induction of SOST gene expression. By 8 hours, SOST expression is markedly diminished (~ 4-fold reduction) by FGF2 treatment and remains so at 8 days (10-fold reduction). However, by 12 days in culture with FGF2, MC4 cells again demonstrate an increase in SOST gene expression (2.4-fold induction) relative to vehicle-treated controls.

Of interest, the effects of FGF2 on MC4 osteoblast gene expression/osteocyte differentiation are reversible. Within 24 hours of the removal of FGF2 from the culture media, the cells revert to an osteoblast-like spindle-shaped/cuboidal morphology (data not shown). Likewise, removal of FGF2 from the culture media results in the reversion of the gene expression to the same levels as cultured MC4 cells without FGF2 within 24 hours (data not shown). This data demonstrates that FGF2 is not sufficient to permanently alter the cell phenotype towards the osteocyte lineage. It does support the notion that FGF2 is, at minimum, capable of “pushing” osteoblastic cells towards the osteocyte lineage. At present, it is unknown whether additional factors may be required to terminally and irreversibly differentiate these cells into the osteocyte lineage. Regardless, this data is the first to demonstrate that FGF2 may be a factor influencing the transition of osteoblast to an osteocyte lineage.

To see if the effects of FGF2 on the development of an osteocyte-like phenotype are specific to MC4 cells, we examined the morphology and gene expression patterns of primary murine bone marrow stromal cells (BMSCs) that had been treated with FGF2. Unlike the response in MC4 cells, there is a robust stimulation of cell proliferation among the BMSCs upon FGF2 treatment. However, we observe a subset of cells that develop an osteocyte-like appearance (not shown). Further, we observe a similar increase in the expression of osteocyte markers, including DMP1, E11, Cx43 and a decrease in COL1A1 and ALP expression at 24, 28 and 72 hours of treatment with FGF2 (Fig. 3). In contrast to the observed effect in MC4 cells, PHEX was noticeably diminished by FGF2 treatment in BMSCs. SOST expression was undetected in the BMSCs. At time points later than 72 hours, the population of cells that proliferate in response to FGF2 overgrows the plate. It is likely that this population of cells blunts the apparent changes in gene expression in the subset of cells that have taken on a osteocyte-like morphology. Future studies will need to be undertaken to separate the sub-populations of cells in the BMSC cultures to fully appreciate the effect of FGF2 on this subset of responsive cells. Regardless, it is apparent that FGF2 can rapidly induce osteocyte-like changes in the gene expression and morphology in BMSCs, and thus is not solely specific to MC4 osteoblasts.

Finally, we examined whether FGF2 could impact the phenotype of osteocytes directly. Accordingly, we used the MLO-Y4 osteocyte-like cell line, which was derived from osteocytes isolated from transgenic mice overexpressing the SV40 large T-antigen under control of the osteocalcin promoter [22]. In contrast to MC4 osteoblasts and BMSCs, DMP1 is the only tested gene induced during the early time points, with FGF2 increasing DMP1 expression 3.7-fold at 8 hours of treatment in the MLO-Y4 cells (data not shown). However, at the 24, 48 and 72 hour time points, FGF2 induced the expression of DMP1 (Fig. 4A) and E11 (Fig. 4B) in MLO-Y4 osteocyte-like cells. The expression of connexin43 was increased ~2 fold at the 48 and 72 hours in FGF2 treated MLO-Y4 cells (Fig. 4C). The expression of PHEX (Fig. 4D) and SOST (Fig. 4E) were both inhibited by FGF2 treatment at the 24, 48 and 72 hour time points. The inhibition of SOST expression by FGF2 in osteocytes is particularly intriguing. SOST is a critical inhibitor of bone formation by antagonizing Wnt signaling [39]. Paradoxically, FGF2 has been shown to be osteoanabolic *in vivo* despite being largely seen as antagonistic towards osteoblast differentiation and mineralization. However, the inhibition of SOST expression by FGF2 may suggest that the osteoanabolic actions of FGF2 are not solely mediated by regulating osteoblasts/osteoprogenitor proliferation, but may also extend to regulation of mineralization signals (e.g., SOST) by osteocytes.

4. Conclusions

The current study shows that FGF2 induces changes in morphology and gene expression in an osteoblast cell line and a subset of BMSCs that are consistent with progression towards an osteocyte like phenotype. Importantly, these data support the hypothesis that FGF2 influences the transition of osteoblasts to osteocytes. Further, this study presents a complimentary model to study osteocyte biology *in vitro*. Additionally, the data reveal that FGF2 regulates the expression of genes by osteocytes that are involved in the development of the osteocyte cellular process, the formation of the elaborate canalicular network, Wnt signaling and phosphate homeostasis. Lastly, this study also reveals that FGF2 may inhibit expression of SOST, an important negative regulator of bone formation, by osteocytes and may add to the current understanding of the potent osteoanabolic effects of FGF2 *in vivo*.

Acknowledgments

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Abbreviations

ALP	alkaline phosphatase
BMSCs	bone marrow derived stromal cells
COL1A1	collagen I α 1
DMP1	dentin matrix protein 1
FGF2	fibroblast growth factor 2
MC4	MC3T3 clone 4
PHEX	phosphate-regulating gene with homologies to endopeptidases on the X chromosome
SIBLING	small, integrin binding ligand, N-linked glycoprotein
SOST	sclerostin

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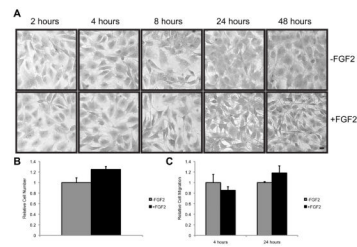


Figure 1.

FGF2 induces an osteocyte-like morphology in MC4 osteoblasts. (A) Photomicrographs of crystal violet stained MC4 cells treated with or without FGF2 (10ng/ml) for the indicated time. Scale bar, 10 μ m. (B) The relative number of viable cells were determined in MC4 cells cultured in the presence or absence of FGF2 (10ng/ml) for 16 hours. (C) The relative number of MC4 cells migrating through an 8 μ m transwell chamber in response to the presence or absence of FGF2 (10ng/ml) for the indicated time was assessed by and counting crystal violet stained cells under a light microscope.

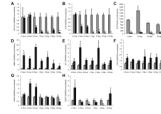


Figure 2.

FGF2 induces the expression of several osteocyte markers in MC4 osteoblasts. Real time PCR was performed on cDNA prepared from MC4 cells cultured in the presence (black bars) or absence (gray bars) of FGF2 (10ng/ml) for the indicated time periods. Gene specific primers were used to determine the expression of (A) COL1A1, (B) ALP, (D) DMP1, (E) E11, (F) Cx43, (G) PHEX, and (H) SOST. Gene expression data are normalized to 18S rRNA. ALP enzymatic activity (C) is shown for cultures treated with (black bars) or without (gray bars) FGF2 for the indicated time periods. Each time point is normalized relative to the respective -FGF2 control. Asterisks (*) indicate a p-value <0.05 relative to the -FGF2 control.

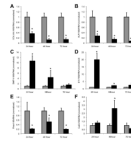


Figure 3.

FGF2 induces the expression of several osteocyte markers in BMSCs. Real time PCR was performed on cDNA prepared from mouse BMSCs cultured in the presence (black bars) or absence (gray bars) of FGF2 (10ng/ml) for the indicated time periods. Gene specific primers were used to determine the expression of (A) COL1A1, (B) ALP, (C) DMP1, (D) E11, (E) PHEX, and (F) Cx43. Gene expression data are normalized to 18S rRNA. Each time point is normalized relative to the respective -FGF2 control. *, p-value <0.05 relative to the -FGF2 control.

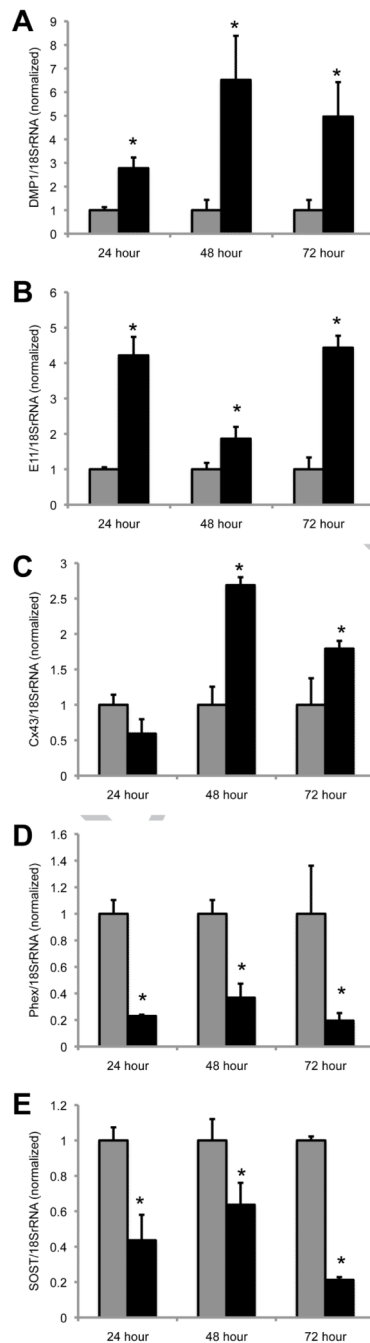


Figure 4. FGF2 modulates gene expression in MLO-Y4 osteocytes

Real time PCR was performed on cDNA prepared from MLO-Y4 cells cultured in the presence (black bars) or absence (gray bars) of FGF2 (10ng/ml) for the indicated time periods. Gene specific primers were used to determine the expression of (A) DMP1, (B) E11, (C) Cx43, (D) PHEX, and (E) SOST. Gene expression data are normalized to 18S rRNA. Each time point is normalized relative to the respective -FGF2 control. *, p-value <0.05 relative to the -FGF2 control.

TABLE I

Mouse Real Time PCR Primers

TARGET	FORWARD	REVERSE
COL1A1	CTTCACCTACAGCACCTTGTG	GATGACTGTCTTGCCCAAGTT
ALP	CCTCCGGATCCTGACCAAA	GTCAATCCTGCCTCCTCCA
DMP1	TGTCATTCTCCTTGTGTTCTTTG	AGAGCTTCAGATTCAGTATTGTGGTAT
E11	TGGCAAGGCACCTCTGGTA	TGAGGTGGACAGTTCCTCTAAGG
CX43	CAGGCCGGAAGCACCAT	GCTGTCGTCAGGGAAATCAA
PHEX	GGAAGAAAACCATTGCCAATTATT	CGCCTGCTGAGGTTTGGA
SOST	GGAATGATGCCACAGAGGTCAT	CCCGGTTTCATGGTCTGGTT
18S rRNA	CATTAAATCAGTTATGGTTCCTTTGG	TCGGCATGTATTAGCTCTAGAATTACC