Friedman et al. supplementary data



Figure S1. Wnt11 and Rspo2 expression in control and retrovirally infected cells.

MC3T3 control (GFP), Wnt11+, or Rspo2+ cells were placed in osteogenic conditions with or without BMP. The cells were harvested for RNA after nine days of osteo-induction. Gene expression was subsequently analyzed by Q-RT-PCR. **A.** Wnt11. **B.** Rspo2. Expression values represent fold change in gene expression relative to non-BMP treated controls, normalized to beta-actin.

-1.5 kb	cacttttCAAAgttggg
-1 kb	ggaaaaaaCAAAgccaag
9 kb	gtgctaaCAAAgggccc

Figure S2. Subsequence analysis of the Rspo2 promoter. Genomatix program was used to analyze a 2 Kb region of the Rspo2 promoter including part of the first exon. Three Tcf/Lef consensus sites were identified.



Figure S3. Model of Wnt11/Rspo2 mediated osteoblast differentiation. In this model BMP increases the expression of TCF7, DMP, and BSP and inhibits Rspo2 expression. Wnt11 increases nuclear accumulation of beta-catenin, which is required for increased Rspo2 expression. Coordinated BMP-Wnt signaling, possibly through Tcf7-beta-catenin in addition to other BMP regulated pathways (Smad/MAPK), is required for increased DMP. Rspo2 on the other hand can directly increase DMP, BSP, and Phex gene expression and increase beta-catenin nuclear accumulation. This beta-catenin signaling may partially influence Phex expression, which requires BMP (Tcf7). These results suggest that Phex and BSP are two factors that could be required for the enhanced mineralization promoted by Wnt11 through Rspo2.

Table S1A. BMP Regulated Transcription Factors, ECM, Proteoglycans				
	o T	-		
Name	Gene Litle	Description		
Transcriptio	on Factors Increas	sed		
Id1	U43884	inhibitor of DNA binding 1		
Junb	NM 008416	Jun-B oncogene		
Tcf7	AI323642	transcription factor 7, T-cell specific		
klf4	BG069413	Kruppel-like factor 4 (gut)		
DIx3	U79738	distal-less homeobox 3		
Tcf7	NM_009331	transcription factor 7, T-cell specific		
Klf10	NM_013692	Kruppel-like factor 10		
Sp7	NM_130458	trans-acting transcription factor 7,		
	-	Osterix		
Tbx2	NM_009324	T-box 2		
Rcor2	NM_054048	REST corepressor 2		
Transcriptio	on Factors Decrea	sed		
ld4	BB121406	inhibitor of DNA binding 4		
		ů.		
ECM, Prote	eoglycans Increas	ed		
Lepre1	NM 019783	leprecan 1		
HapIn1	NM_013500	hyaluronan and proteoglycan link protein		
	-	1		
Cspg4	BB377873	chondroitin sulfate proteoglycan 4		
lbsp	L20232	integrin binding sialoprotein		
Col2a1	NM_031163	procollagen, type II, alpha 1		
Bglap1	NM_007541	bone gamma carboxyglutamate protein 1		
Gpc1	NM_016696	glypican 1		
P4ha2	1417149_at	procollagen-proline, 2-oxoglutarate 4-		
		dioxygenase (proline 4-hydroxylase),		
		alpha II polypeptide		
Loxl4	NM_053083	lysyl oxidase-like 4		
Col11a2	AV242706	procollagen, type XI, alpha 2		
ECM, Prote	eoglycans Decrea	sed		
Otor	AF243504	otoraplin, cartilage-derived retinoic acid		
		sensitive protein		
Col8a1	NM_007739	procollagen, type VIII, alpha 1		

Table S1A. BMP regulated expression of transcription factors, ECM, and proteoglycans in pre-osteoblasts. RNA was harvested at days 3, 6, and 9 after the start of osteo-induction. cRNA samples were hybridized to an Affymetrix 420A 2.0 microarray chip. Only genes showing an expression level \geq 400 with an interquartile range >4 were evaluated. The genes represented in this table showed increased or decreased expression at day 9 in BMP treated cells relative to non-BMP treated GFP controls.

Name	Gene Title	Description		
Receptors I	ncreased			
Ly6a	BC002070	lymphocyte antigen 6 complex, locus A,		
-		Sca-1		
Unc5b	BG065285	unc-5 homolog B (C. elegans)		
Tnfrsf11b	AB013898	tumor necrosis factor receptor		
		superfamily, member 11b		
		(osteoprotegerin)		
Lifr	D17444	leukemia inhibitory factor receptor		
Bambi	AF153440	BMP and activin membrane-bound		
		inhibitor, homolog (Xenopus laevis)		
Vldlr	NM_013703	very low density lipoprotein receptor		
Pthr1	BC013446	parathyroid hormone receptor 1		
Vdr	AV290079	vitamin D receptor		
		phosphate regulating gene with		
Dhav	NIM 011077	nomologies to endopeptidases on the X		
Priex		chromosome		
Decenteral	Deerseed			
Receptors I		CD24 entiren		
Cu34	NIM_133034	CD34 antigen		
Signal Tran	sduction Increased			
Rasa3	NM 009025	RAS p21 protein activator 3		
Rras2	NM 025846	related RAS viral (r-ras) oncogene		
		homolog 2		
Rassf5	NM 018750	Ras association (RalGDS/AF-6) domain		
	-	family 5		
Ramp1	NM_016894	receptor (calcitonin) activity modifying protein		
Plcd1	NM_019676	phospholipase C, delta 1		
Rab27b	BB121269	RAB27b, member RAS oncogene family		
Jak2	NM_008413	Janus kinase 2		
Gpr133	BB548889	G protein-coupled receptor 133		
Gpr35	NM_022320	G protein-coupled receptor 35		
Trib3	BB508622	tribbles homolog 3 (Drosophila)		
Signal I ran	A FOODERA	u		
KNOJ Rao2	AF309564	ras nomolog gene family, member J		
rtysz	AF210000	regulator of G-protein signaling 2		

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Table S1B. BMP Regulated Receptors and Signal Transduction Molecules

Table S1B. BMP regulated expression of receptors and signal transduction

molecules in pre-osteoblasts. RNA was harvested at days 3, 6, and 9 after the start of osteo-induction. cRNA samples were hybridized to an Affymetrix 420A 2.0 microarray chip. Only genes showing an expression level \geq 400 with an interquartile range >4 were evaluated. The genes represented in this table showed increased or decreased expression at day 9 in BMP treated cells relative to non-BMP treated GFP controls.

Table S2. Wnt-11 Regulated Transcription Factors, ECM, Proteoglycans				
Name	Gene Title	Description		
Transcription Factors Increased				
Sox6	AJ010605	SRY-box containing gene 6		
Ebf1	BB125261	early B-cell factor 1		
ECM, Prote	oglycans Increased			
Pcolce2	AF352788	procollagen C-endopeptidase enhancer 2		
HapIn1	NM_013500	hyaluronan and proteoglycan link protein 1		
Otor	AF243504	otoraplin, cartilage-derived retinoic acid sensitive protein		
Dmp1	U65020	dentin matrix protein 1		
Phex	U73910	phosphate regulating gene with		
		homologies to endopeptidases on the X		
		chromosome (hypophosphatemia,		
		vitamin D resistant fickets)		
Secreted Fa	actors Increased			
Wif1	BC004048	Wnt inhibitory factor 1		
lgf1	NM_010512	insulin-like growth factor 1		
Rspo2	BG067392	R-spondin 2 homolog (Xenopus laevis)		
Lect1	NM_010701	leukocyte cell derived chemotaxin 1		
Receptors I	ncreased			
Ly6a	BC002070	lymphocyte antigen 6 complex, locus A, Sca-1		
Vdr	AV290079	vitamin D receptor		
Ly6c	NM_010741	lymphocyte antigen 6 complex, locus C		
Itga6	BM935811	integrin alpha 6		
		phosphate regulating gene with		
Phex	NM_011077	homologies to endopeptidases on the X chromosome		
Signal Transduction Increased				
Camk2b	BG862223	calcium/calmodulin-dependent protein kinase II, beta		

Table S2. Wnt11 regulated gene expression. RNA was harvested at days 3, 6, and 9 after the start of osteo-induction. cRNA samples were hybridized to an Affymetrix 420A 2.0 microarray chip. Only genes showing an expression level \geq 400 with an interquartile range >4 were evaluated. The genes represented in this table showed increased or decreased expression at day 9 in non-BMP treated Wnt11+ cells relative to non-BMP treated controls.

Table S3 A. BMP+Wnt-11 Regulated Transcription Factors, ECM,		Table S3 B. BMP+Wnt-11 Regulated Receptors and Signal			
Proteoglycans, and Secreted Factors		Transduction Molecules			
Name	Gene Title	Description	Name	Gene Title	Description
Transcription Factors Increased		Receptors Increased			
Hmga1	NM_016660	high mobility group AT-hook 1	Ank	NM_020332	progressive ankylosis
Egr2	X06746	early growth response 2			phosphate regulating gene with
Junb	NM_008416	Jun-B oncogene	Phex	NM_011077	homologies to endopeptidases on the X
Foxc2	NM_013519	forkhead box C2			chromosome
Fosl1	U34245	fos-like antigen 1			
Nfatc1	NM_016791	nuclear factor of activated T-cells,	Receptors Decreased		
			Ptgfr	NM 008966	prostaglandin F receptor
ECM, Prote	eoglycans Increas	ed	U U	-	
Dmp1	U65020	dentin matrix protein 1	Signal Transduction Increased		
Tnfaip6	NM_009398	tumor necrosis factor alpha induced	Ras4	NM 009062	regulator of G-protein signaling 4
		protein 6	Sprv2	BB529691	sprouty homolog 2 (Drosophila)
Ereg	NM_007950	epiregulin	Rab27b	BB121269	RAB27b member RAS oncogene family
Otor	AF243504	otoraplin	Ptas2	M94967	prostaglandin-endoperoxide synthase 2
		F	Gprc5b	BC020004	G protein-coupled receptor family C
ECM. Proteoglycans Decreased		00.000	2002000	group 5. member B	
Aspn	NM_025711	asporin	Rgs16	U94828	regulator of G-protein signaling 16
Secreted Factors Increased		Signal Transduction Decreased			
Ctqf	NM 010217	connective tissue growth factor	Ddit4	AK017926	DNA-damage-inducible transcript
ll12a	AF128210	interleukin 12a			4/REDD1
Adm	NM 009627	adrenomedullin	Plekha5	1425543 s at	pleckstrin homology domain containing.
Bmp8a	NM_007558	bone morphogenetic protein 8a			family A member 5
Rspo2	BG067392	R-spondin 2 homolog (Xenopus laevis)	Osr1	NM_011859	odd-skipped related 1 (Drosophila)
Secreted F	actors Decreased				
lgfbp5	BF225802	insulin-like growth factor binding protein 5			
•••		5 01			

Table S3. Genes showing additive or synergistic changes in expression withWnt11 and BMPMicroarray analysis was performed and analyzed as previouslymentioned. The genes represented in this table showed increased or decreasedexpression at day 9 in BMP treated Wnt11+ cells relative to BMP treated controls.