

SUPPLEMENTARY TABLES

Supplementary Table 1. Abbreviations of bone parameters for histomorphometrical and μ CT analysis.

Type of index	Parameter	Abbreviation	Unit	Analyzed bone
Structural (Trabecular)	Bone volume/tissue volume	BV/TV	%	Distal femur, L5 vertebra
	Number	Tb.N	/mm	
	Thickness	Tb.Th	mm	
	Separation	Tb.Sp	mm	
Structural (Cortical)	Thickness	Cort.Th	mm	Midshaft femur
	Density	Cort.Dens	mmHA/ccm	
	Area	Cort.A	mm ²	
	Medullary area	MA	mm ²	
	Porosity	Cort.Por	%	
Dynamic	Polar moment of inertia	pMOI	mm ⁴	Distal and midshaft femur
	Mineralizing surface/bone surface	MS/BS	%	
	Mineral apposition rate	MAR	μ m/day	
	Bone formation rate/bone surface	BFR/BS		
Formation	Bone formation rate/bone volume	BFR/BV (%/year)	%/year	Distal femur
	Osteoid volume/bone volume	OV/BV	%	
	Osteoid surface/bone surface	OS/BS	%	
	Osteoblast surface/bone surface	Ob.S/BS	%	
	Osteoblast number/tissue area	N.Ob/T.Ar		
Resorption	Osteoblast number/bone perimeter	N.Ob/B.Pm	/mm	Distal femur
	Erosion surface/bone surface	ES/BS	%	
	Osteoclast surface/bone surface	Oc.S/BS	%	
Osteocyte	Osteoclast number/bone perimeter	N.Oc/B.Pm	/mm	Distal and midshaft femur
	Osteocyte number/bone volume	N.Ot/BV	/mm ²	
	Ot density (/mm ²)		/mm ²	

Supplementary Table 2. Primer sequences for real time qPCR.

Gene Symbol	Forward (5'–3')	Reverse (5'–3')
<i>18S</i>	TCAAGAACGAAAGTCGGAGG	GGACATCTAAGGGCATCAC
<i>Actb</i> (β -actin)	GGCTGTATCCCCTCCATCG	CCAGTTGGTAACAATGCCATGT
<i>B2m</i>	TTCACCCCCACTGAGACTGAT	GTCTTGGGCTCGGCCATA
<i>Becn1</i>	GAGGCTAACTCAGGAGAGGAGC	TGCCTCCCCGATCAGAGTGA
<i>Bglap</i> (<i>Osteocalcin</i>)	CCGGGAGCAGTGTGAGCTTA	TAGATGCGTTTGTAGGCGGTC
<i>Cdkn1a</i> (<i>p21^{Cip1}</i>)	CCTGGTGATGTCCGACCTG	CCATGAGCGCATCGCAATC
<i>Cdkn2a</i> (<i>p16^{Ink4a}</i>)	GAACTCTTTCGGTCGTACCC	AGTTCGAATCTGCACCGTAGT
<i>Csf1</i> (<i>M-CSF</i>)	GACCCTCGAGTCAACAGAGC	TGTCAGTCTCTGCCTGGATG
<i>Ctsk</i>	GAAGAAGACTCACCAGAAGCAG	TCCAGGTTATGGGCAGAGATT
<i>Dmp1</i>	AAGCTAGCCCAGAGGGACAGGCAA	TTATCGGCGCCGGTCCCCGTAC
<i>Foxo1</i>	CCCAGGCCGGAGTTTAACC	GTTGCTCATAAAGTCGGTGCT

<i>Gapdh</i>	AGGTCGGTGTGAACGGATTTG	TGTAGACCATGTAGTTGAGGTCA
<i>Hprt</i>	TCAGTCAACGGGGGACATAAA	GGGGCTGTACTGCTTAACCAG
<i>Pth1r (PPR)</i>	CAGATTTTCCTGCTGCACCG	CTGCTGTGTGCAGAACTTCC
<i>Pthlh (PTHrP)</i>	CAGTTAGAGGCGCTGATTCC	AGCTCTGATTTCCGGCTGTGT
<i>Rlp13a</i>	GGGCAGGTTCTGGTATTGGAT	GGCTCGGAAATGGTAGGGG
<i>Sdha</i>	GGAACACTCCAAAAACAGACCT	CCACCACTGGGTATTGAGTAGAA
<i>Sirt1</i>	GCTGACGACTTCGACGACG	TCGGTCAACAGGAGGTTGTCT
<i>Sost (Sclerostin)</i>	CTTCAGGAATGATGCCACAGAGGT	ATCTTTGGCGTCATAGGGATGGTG
<i>Sphk1</i>	GAACCATAACTCTGTGCCTTTGTCT	AGCAATGGGGAGTGTCTTCTATATG
<i>Tnf (TNFα)</i>	AGACCCTCACACTCAGATCATCTTC	CCACTTGGTGGTTTGCTACGA
<i>Tnfsf11 (RANKL)</i>	CACAGCGCTTCTCAGGAGCTC	GAGATCTTGGCCCAGCCTCGA
<i>Tnfrsf11b (OPG)</i>	ACCCAGAAACTGGTCATCAGC	CTGCAATACACACACTCATCACT
<i>Trp53 (p53)</i>	CTCTCCCCCGAAAAGAAAAA	CGGAACATCTCGAAGCGTTTA
<i>Ywhaz</i>	GAAAAGTTCTTGATCCCCAATGC	TGTGACTGGTCCACAATTCCTT
