

Supplemental Information for:

The collection of NFATc1-dependent transcripts in the osteoclast includes numerous genes non-essential to physiologic bone resorption

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Supplemental Figure Legends

Supplemental Fig. 1 – Supplemental analysis of *Nhedc2*^{GT/GT} mice. (A) Genotyping PCR reactions for the *Nhedc2*^{WT} (left gel) and *Nhedc2*^{GT} (right gel) alleles on tail DNA from *Nhedc2*^{+/+} (lane 2), *Nhedc2*^{GT/+} (lane 3) and *Nhedc2*^{GT/GT} (lane 4) mice. A 100 bp DNA ladder is shown in lane 1. **(B)** Micro-CT quantification of femoral metaphyseal trabecular parameters in 12-14 week old female and male *Nhedc2*^{+/+} (n=9, female; n=11, male) and *Nhedc2*^{GT/GT} (n=9, female; n=11, male) mice. (n.s., not significant).

Supplemental Fig. 2 – Supplemental analysis of *Serpind1*^{-/-} mice. (A) Micro-CT quantification of femoral metaphyseal trabecular parameters in 12-week-old female and male *Serpind1*^{+/+} (female, n=7; male, n=6) and *Serpind1*^{-/-} (female, n=4; male, n=3). **(B)** Real time PCR analysis for osteoclast marker genes in samples from *Serpind1*^{+/+} (n=2) and *Serpind1*^{-/-} (n=4) OCPs stimulated with MCSF and RANKL to induce osteoclast differentiation. **(C, D)** Real time PCR analysis using primers targeting *Serpind1* exons 2 (forward) and 3 (reverse) on **(C)** *Serpind1*^{+/+} and *Serpind1*^{-/-} liver or **(D)** WT BMMs cultured with MCSF alone or MCSF and RANKL to induce osteoclast differentiation. **(E)** Real time PCR analysis for *Serpin* homologs in samples from *Serpind1*^{+/+} (n=2) and *Serpind1*^{-/-} (n=4) OCPs stimulated with MCSF and RANKL to induce osteoclast differentiation. (n.s., not significant; N.D., not detectable).

Supplemental Fig. 3 – Supplemental analysis of *Adcy3*^{-/-} mice. (A) Micro-CT quantification of femoral metaphyseal trabecular parameters in 8-week-old female and male *Adcy3*^{+/+} (female, n=6; male, n=5) and *Adcy3*^{-/-} (female, n=7; male, n=8) mice. **(B)** Body weight of 8-week old female and male *Adcy3*^{+/+} and *Adcy3*^{-/-} mice used for micro-CT analysis in Fig. 4 A,B and Supplemental Fig. 3A. **(C)** Real time PCR analysis for osteoclast marker genes expression in *Adcy3*^{+/+} (n=3) and *Adcy3*^{-/-} (n=4) OCPs stimulated with MCSF and RANKL to induce osteoclast differentiation. Data in (C) is normalized to WT littermates. **(D)** Real time PCR analysis for *Adcy3* expression in *Adcy3*^{+/+} (n=2) and *Adcy3*^{-/-} (n=3) OCPs stimulated with MCSF and RANKL to induce osteoclast differentiation. (n.s., not significant).

Supplemental Fig. 4 – Supplemental analysis of *Rhoc*^{-/-} mice. (A) Micro-CT quantification of femoral metaphyseal trabecular parameters in 8-10 week old female

and male *Rhoc*^{2/+} (female, n=7; male, n=2) and *Rhoc*^{-/-} (female, n=5; male, n=2). **(B)** Real time PCR analysis for *Rhoc* expression in *Rhoc*^{2/+} (n=2) and *Rhoc*^{-/-} (n=2) OCPs stimulated with MCSF and RANKL to induce osteoclast differentiation.

Supplemental Fig. 5 – Supplemental analysis of *Rab38*^{cht/cht} mice. **(A)** Micro-CT quantification of femoral metaphyseal trabecular parameters in 8-week-old female and male *Rab38*^{+/+} (female, n=2; male, n=4) and *Rab38*^{cht/cht} (female, n=3; male, n=3) mice. **(B)** Real time PCR analysis for osteoclast marker genes expression in *Rab38*^{+/+} (n=2) and *Rab38*^{cht/cht} (n=1) OCPs stimulated with MCSF alone or MCSF and RANKL to induce osteoclast differentiation. (n.s., not significant; *, p<0.05)

Supplemental Table 1 – Quantitative Real-time PCR primers used in this study

| Primer Set | Forward Primers (5'-3') | Reverse Primers (5'-3') | Source |
|-------------------------------------|-------------------------|-------------------------|--------|
| <i>Adcy3</i> | CCTGAGTCCTTGGAGAACCT | CCACGTAGCAGTCAAAGAGG | A |
| <i>Hmbs</i> | ATGAGGGTGATTTCGAGTGGG | CAAACGTATGCCAGGGTACAA | B |
| <i>Hprt</i> | GTTAAGCAGTACAGCCCCAAA | AGGGCATATCCAACAACAACTT | C |
| <i>Nfatc1</i> | TGCCTTTTGCAGCAGTATCT | CAGGCAAGGATGGGCTCATAT | D |
| <i>Nhedc1</i> | CCACTGGGCAGATGGTATTT | GCAGCATGCCAAGAAGA | E |
| <i>Nhedc2</i> | TTGTTCCATCACCGGAGGTAA | TTGGTGTTGAGTGCTTGCC | E |
| <i>Rhoc</i> | ACTCCATGCTGCCAGTTTCT | AGGCCAGAGGGACTAAGAGC | E |
| <i>Rab38</i> | CCAAGGGAAGGATGTGCTTA | GTGAGATGGGGCTTCACAAT | E |
| <i>Serpind1</i> (<i>ex2-3</i>) | GGCAAACAACCACATTCTG | TCTCTCATTAGCCGGAAGT | E |
| <i>Serpind1</i> (<i>ex3-4</i>) | GAGTACGTAGGGGGCATCAG | GGACCTCCACCAGGTTGTAA | E |
| <i>Serpinb1a</i> | ACATCCATTACGCTTCCAAA | GGCCAAGTCAGCACCATACAT | C |
| <i>Serpinb2</i> | GTGCTGGGGGTAACACTGAAC | GCGAAATCACAGCCACTGAAG | C |
| <i>Serpine2</i> | CACATGGGATCGCGTCCATC | CAGCACTTTACCAACTCCGTTTA | C |
| <i>Serpinh1</i> | GCCGAGGTGAAGAAACCCC | CATCGCCTGATATAGGCTGAAG | C |

A – Designed with the on-line GenScript Real Time PCR (TaqMan) Primer Design program.

B – Gift of Dr. Kevin McHugh, College of Dentistry, University of Florida

C - The Center for Comparative and Integrative Biology Primer Bank, Harvard Medical School. Primerbank ID: *Hprt* (7305155a2), *Serpinb1a* (13384828a1), *Serpinb2* (6755098a1), *Serpine2* (6678099a1), *Serpinh1* (6753304a1).

D - A.O. Aliprantis, Y. Ueki, R. Sulyanto, A. Park, K.S. Sigrist, S.M. Sharma, M.C. Ostrowski, B.R. Olsen, L.H. Glimcher, NFATc1 in mice represses osteoprotegerin during osteoclastogenesis and dissociates systemic osteopenia from inflammation in cherubism, *J Clin Invest.* 118 (2008) 3775-89.

E - Designed with the on-line Primer 3 tool. Steve Rozen and Helen J. Skaletsky (2000) Primer3 on the WWW for general users and for biologist programmers. In: Krawetz S, Misener S (eds) Bioinformatics Methods and Protocols: Methods in Molecular Biology. Humana Press, Totowa, NJ, pp 365-386.

Supplemental Table 2 – Raw microarray data for the genes depicted on the heat map in Fig. 1A.

| Heatmap.order | Probeset | Gene Symbol | p-value | FDR | log2(FC) KOvWT |
|----------------------|-----------------|--------------------|----------------|------------|-----------------------|
| 98 | 1417700_at | Rab38 | 0.005108 | 0.063074 | -3.71 |
| 97 | 1420575_at | Mt3 | 0.004283 | 0.060491 | -4.51 |
| 96 | 1437308_s_at | F2r | 0.003823 | 0.059292 | -4.82 |
| 95 | 1451710_at | Oscar | 0.002111 | 0.052424 | -4.88 |
| 94 | 1433474_at | Edil3 | 0.004097 | 0.060422 | -4.28 |
| 93 | 1448605_at | Rhoc | 0.022969 | 0.110736 | -2.87 |
| 92 | 1426951_at | Crim1 | 0.020935 | 0.106308 | -3.20 |
| 91 | 1460197_a_at | Steap4 | 0.015549 | 0.093885 | -6.51 |
| 90 | 1420461_at | Mst1r | 0.011264 | 0.082114 | -4.90 |
| 89 | 1431711_a_at | Kazn | 0.013887 | 0.089020 | -3.22 |
| 88 | 1421959_s_at | Adcy3 | 0.009205 | 0.076730 | -2.66 |
| 87 | 1425829_a_at | Steap4 | 0.009882 | 0.078992 | -3.83 |
| 86 | 1452646_at | Trp53inp2 | 0.011529 | 0.082658 | -4.30 |
| 85 | 1426869_at | Boc | 0.007521 | 0.071389 | -2.92 |
| 84 | 1450625_at | Col5a2 | 0.016968 | 0.097514 | -2.83 |
| 83 | 1422809_at | Rims2 | 0.015804 | 0.094740 | -3.29 |
| 82 | 1437226_x_at | Marcksl1 | 0.000028 | 0.044454 | -2.82 |
| 81 | 1422619_at | Ppap2a | 0.000004 | 0.044454 | -4.07 |
| 80 | 1424133_at | Tmem98 | 0.000042 | 0.044454 | -5.57 |
| 79 | 1418569_at | Fblim1 | 0.000068 | 0.044454 | -3.46 |
| 78 | 1417089_a_at | Ckmt1 | 0.000133 | 0.044454 | -2.98 |
| 77 | 1450429_at | Capn6 | 0.000069 | 0.044454 | -2.70 |
| 76 | 1450843_a_at | Serpinh1 | 0.000033 | 0.044454 | -4.39 |

| | | | | | |
|----|--------------|----------|----------|----------|-------|
| 75 | 1434588_x_at | Tbca | 0.001101 | 0.046468 | -2.68 |
| 74 | 1421791_at | | 0.000757 | 0.044454 | -3.07 |
| 73 | 1425702_a_at | Enpp5 | 0.000312 | 0.044454 | -2.90 |
| 72 | 1425339_at | Plcb4 | 0.000146 | 0.044454 | -4.77 |
| 71 | 1455235_x_at | Ldhb | 0.000055 | 0.044454 | -3.62 |
| 70 | 1425622_at | Edil3 | 0.000145 | 0.044454 | -4.18 |
| 69 | 1421511_at | Itgb3 | 0.000092 | 0.044454 | -6.33 |
| 68 | 1448237_x_at | Ldhb | 0.000107 | 0.044454 | -3.06 |
| 67 | 1422620_s_at | Ppap2a | 0.000103 | 0.044454 | -4.05 |
| 66 | 1416666_at | Serpine2 | 0.000070 | 0.044454 | -5.04 |
| 65 | 1422967_a_at | Tfrc | 0.000281 | 0.044454 | -2.79 |
| 64 | 1452257_at | Bdh1 | 0.000490 | 0.044454 | -2.71 |
| 63 | 1448942_at | Gng11 | 0.000355 | 0.044454 | -2.83 |
| 62 | 1419136_at | Akr1c18 | 0.000436 | 0.044454 | -4.28 |
| 61 | 1417730_at | Ext1 | 0.000294 | 0.044454 | -3.06 |
| 60 | 1422515_at | Pate4 | 0.000105 | 0.044454 | -3.43 |
| 59 | 1415874_at | Spry1 | 0.000112 | 0.044454 | -2.74 |
| 58 | 1430233_a_at | Nhedc1 | 0.000064 | 0.044454 | -2.86 |
| 57 | 1451596_a_at | Sphk1 | 0.000103 | 0.044454 | -2.90 |
| 56 | 1439995_at | Nhedc2 | 0.002112 | 0.052424 | -4.95 |
| 55 | 1435463_s_at | Myo1d | 0.002468 | 0.053783 | -4.18 |
| 54 | 1423892_at | Apbb1 | 0.001599 | 0.049710 | -3.05 |
| 53 | 1425442_at | Oscar | 0.001743 | 0.050429 | -4.25 |
| 52 | 1418680_at | Serpind1 | 0.001835 | 0.051020 | -3.35 |
| 51 | 1454114_a_at | Nhedc1 | 0.001183 | 0.046674 | -4.84 |
| 50 | 1449141_at | Fblim1 | 0.001226 | 0.047183 | -2.89 |

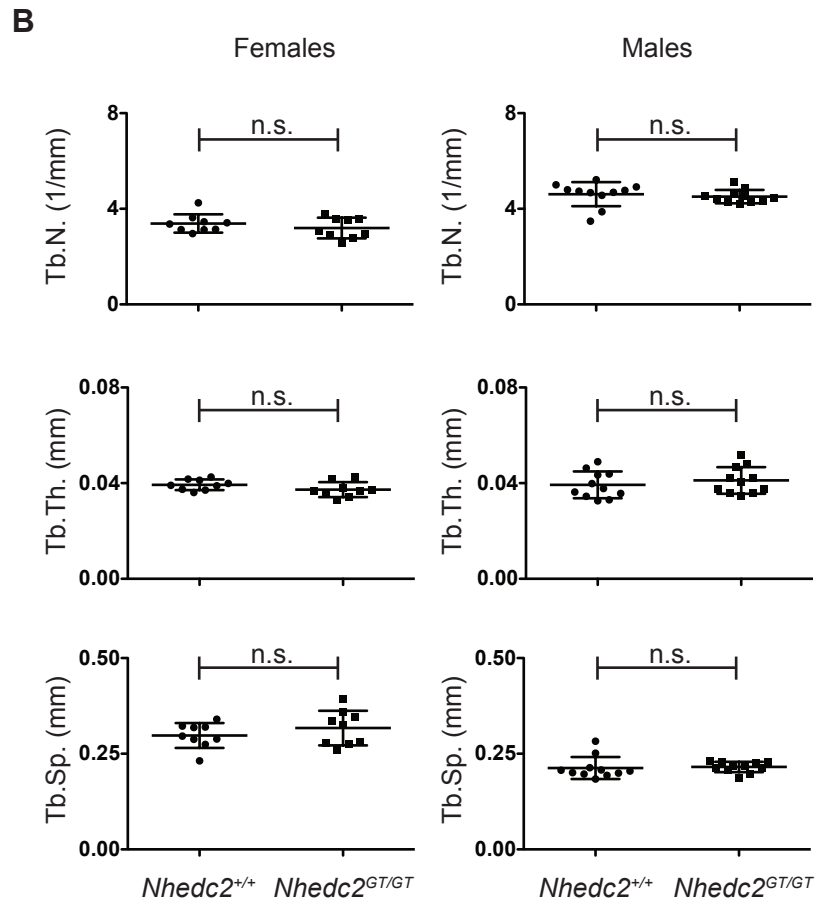
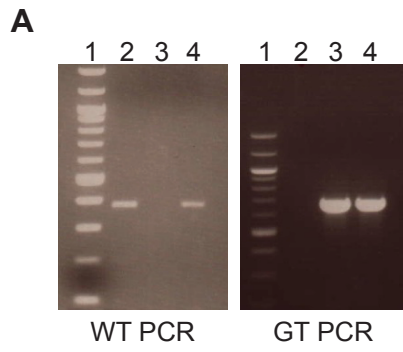
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|----|--------------|----------|----------|----------|-------|
| 49 | 1450852_s_at | F2r | 0.000737 | 0.044454 | -4.72 |
| 48 | 1451601_a_at | Spns2 | 0.000813 | 0.045119 | -3.20 |
| 47 | 1427891_at | Gimap6 | 0.000683 | 0.044454 | -5.31 |
| 46 | 1448502_at | Slc16a7 | 0.001020 | 0.046468 | -3.79 |
| 45 | 1434499_a_at | Ldhb | 0.000500 | 0.044454 | -4.13 |
| 44 | 1418688_at | Calcr | 0.000394 | 0.044454 | -4.80 |
| 43 | 1426959_at | Bdh1 | 0.000220 | 0.044454 | -3.55 |
| 42 | 1421425_a_at | Rcan2 | 0.000292 | 0.044454 | -3.13 |
| 41 | 1425250_a_at | Slc14a2 | 0.000940 | 0.046404 | -2.94 |
| 40 | 1449945_at | Ppargc1b | 0.001283 | 0.047286 | -2.64 |
| 39 | 1423893_x_at | Apbb1 | 0.000328 | 0.044454 | -2.62 |
| 38 | 1416601_a_at | Rcan1 | 0.000481 | 0.044454 | -2.63 |
| 37 | 1425842_at | Edil3 | 0.000503 | 0.044454 | -3.72 |
| 36 | 1422090_a_at | Pfkfb2 | 0.000861 | 0.045901 | -4.12 |
| 35 | 1452492_a_at | Slc37a2 | 0.001084 | 0.046468 | -2.61 |
| 34 | 1419082_at | Serpib2 | 0.037980 | 0.143389 | -2.59 |
| 33 | 1436929_x_at | Adcy3 | 0.031294 | 0.129404 | -3.09 |
| 32 | 1449943_at | Lfng | 0.023573 | 0.112131 | -2.72 |
| 31 | 1428895_at | Rftn2 | 0.005551 | 0.065208 | -4.22 |
| 30 | 1450932_s_at | Dock9 | 0.007990 | 0.073392 | -3.91 |
| 29 | 1452405_x_at | Gm10889 | 0.004160 | 0.060422 | -3.33 |
| 28 | 1426168_a_at | | 0.003382 | 0.057404 | -3.54 |
| 27 | 1423952_a_at | Krt7 | 0.002774 | 0.054990 | -3.65 |
| 26 | 1422875_at | Cd84 | 0.000035 | 0.044454 | -2.91 |
| 25 | 1434624_x_at | Rps9 | 0.000177 | 0.044454 | -2.85 |
| 24 | 1416007_at | Satb1 | 0.000082 | 0.044454 | -2.94 |

| | | | | | |
|----|--------------------|-----------|----------|----------|-------|
| 23 | 1428372_at | St5 | 0.001109 | 0.046468 | -4.79 |
| 22 | 1419621_at | Ankrd2 | 0.000473 | 0.044454 | -4.66 |
| 21 | 1425444_a_at | Tgfbr2 | 0.000379 | 0.044454 | -4.60 |
| 20 | 1425934_a_at | B4galt4 | 0.000027 | 0.044454 | 2.79 |
| 19 | 1448301_s_at | Serpinb1a | 0.000049 | 0.044454 | 3.10 |
| 18 | 1453223_s_at | Dppa2 | 0.000044 | 0.044454 | 3.87 |
| 17 | 1416444_at | Elovl2 | 0.000086 | 0.044454 | 3.65 |
| 16 | 1416318_at | Serpinb1a | 0.000336 | 0.044454 | 3.03 |
| 15 | 1427760_s_at | | 0.000177 | 0.044454 | 4.90 |
| 14 | 1437693_at | D1Pas1 | 0.000583 | 0.044454 | 4.02 |
| 13 | 1434334_at | Prkd2 | 0.000568 | 0.044454 | 3.79 |
| 12 | 1449033_at | Tnfrsf11b | 0.000033 | 0.044454 | 5.44 |
| 11 | 1427231_at | Robo1 | 0.001623 | 0.050016 | 3.15 |
| 10 | 1422368_at | Vmn1r43 | 0.001813 | 0.051020 | 3.47 |
| 9 | 1437275_at | Ctnna1 | 0.000322 | 0.044454 | 4.40 |
| 8 | 1452426_x_at | | 0.015491 | 0.093772 | 4.93 |
| 7 | 1439243_x_at | Cops5 | 0.016630 | 0.096553 | 4.22 |
| 6 | 1450922_a_at | Tgfb2 | 0.009969 | 0.079513 | 3.45 |
| 5 | 1415780_a_at | Armcx2 | 0.020081 | 0.103541 | 3.34 |
| 4 | AFFX-DapX- 5_at | | 0.009520 | 0.078134 | 2.70 |
| 3 | 1449782_at | AA517650 | 0.010090 | 0.079789 | 2.79 |
| 2 | 1419521_at | Zfp94 | 0.038739 | 0.144595 | 3.22 |
| 1 | 1450930_at | Hpca | 0.040907 | 0.148765 | 2.76 |

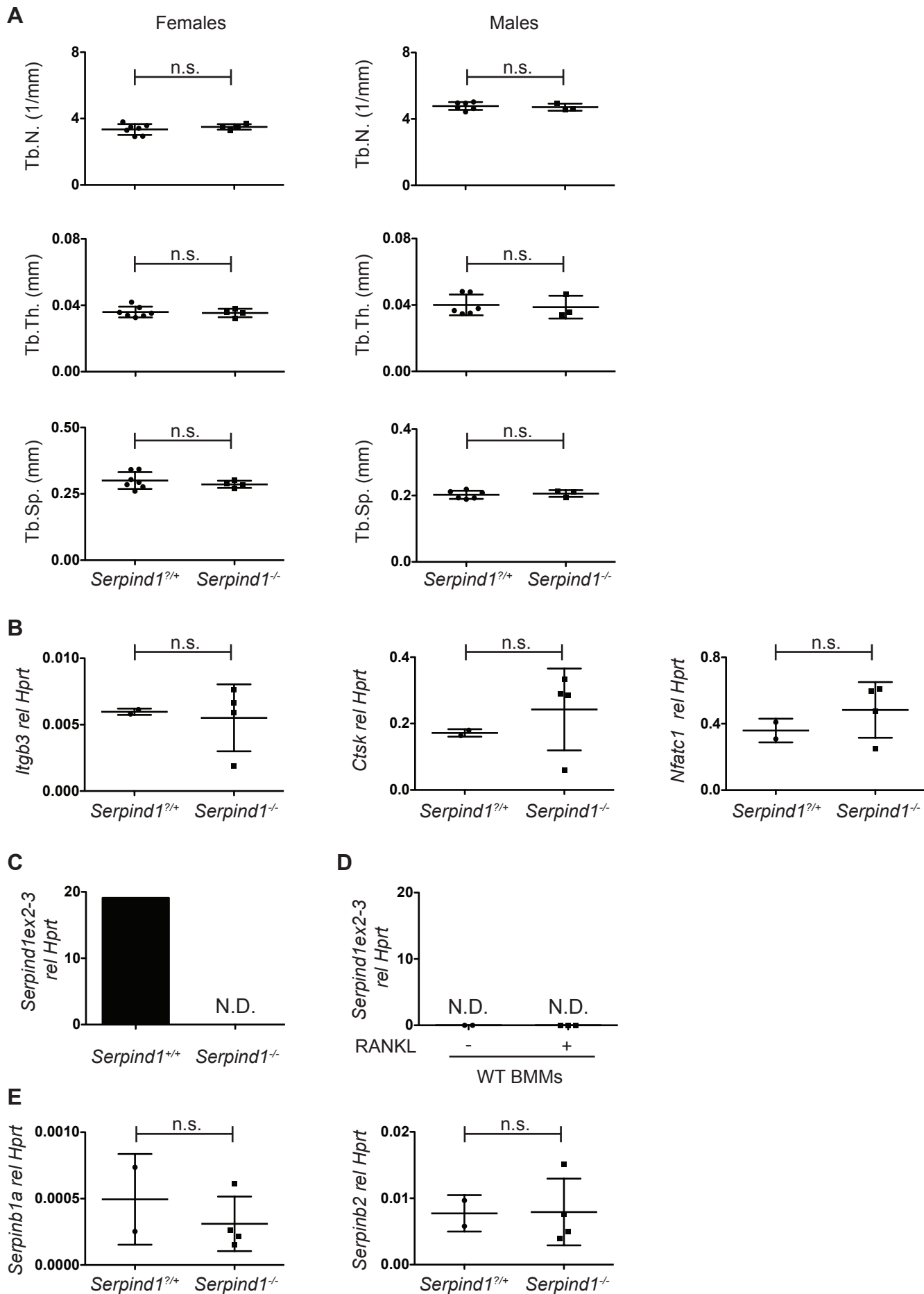
Supplemental Table 3 – Histomorphometric analysis of *Rhoc*^{-/-} mice.

Histomorphometric analysis of the tibia of 8 week of *Rhoc*^{+/+} or *Rhoc*^{-/-} female mice. p-value determined by 2-tailed student's t-test.

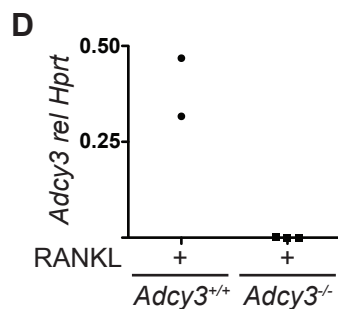
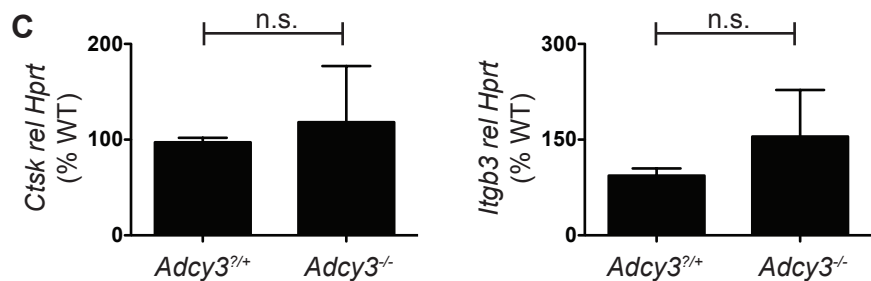
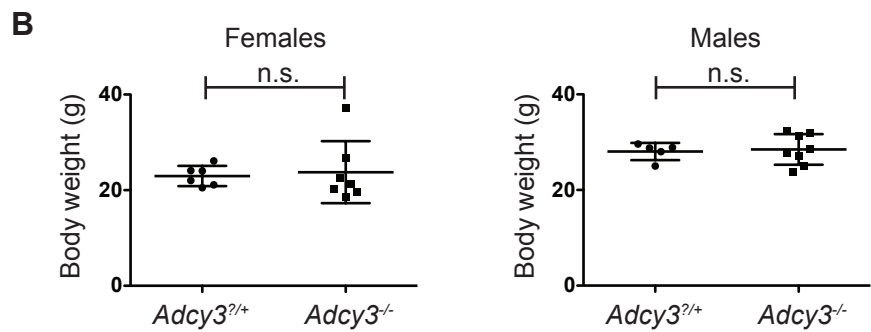
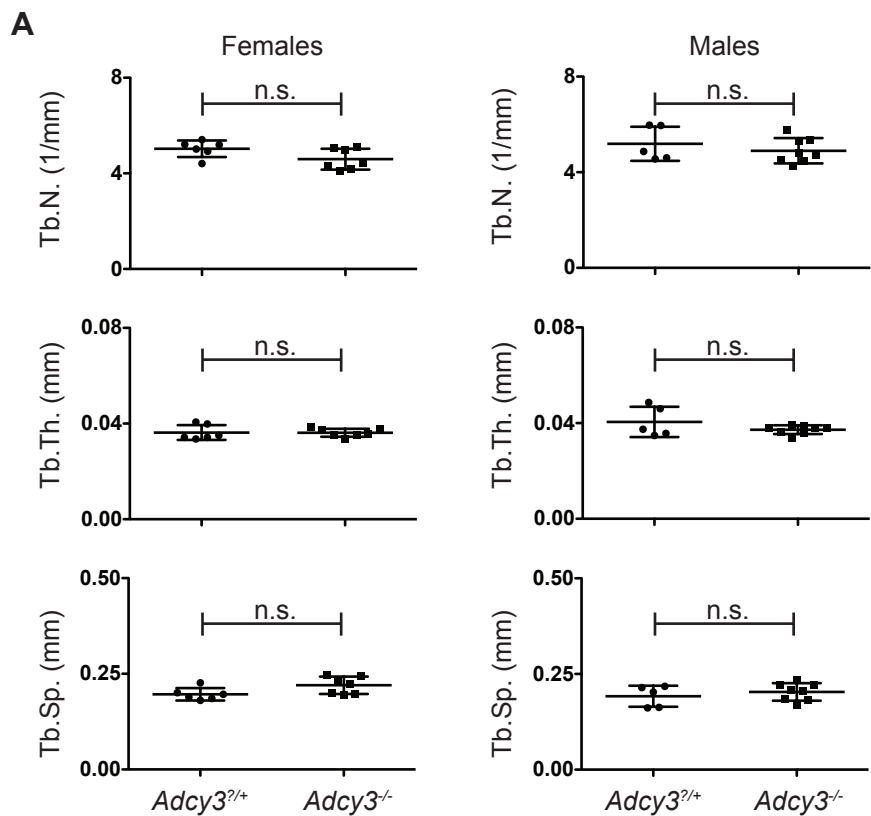
| | <i>Rhoc</i> ^{+/+} (n=5) | | <i>Rhoc</i> ^{-/-} (n=5) | | p-value |
|-----------------|----------------------------------|---------|----------------------------------|--------|---------|
| | Avg. | St.Dev. | Avg. | St.Dev | |
| BV/TV | 20.5 | 3.9 | 17.2 | 2.3 | 0.157 |
| TbTh | 29.2 | 1.1 | 31.6 | 4.4 | 0.355 |
| TbN | 7.0 | 1.3 | 5.5 | 0.6 | 0.057 |
| TbSp | 117.5 | 28.5 | 153.2 | 16.8 | 0.054 |
| BS/BV | 68.6 | 2.6 | 64.1 | 8.4 | 0.369 |
| OTh | 2.7 | 0.3 | 2.8 | 0.2 | 0.748 |
| OV/TV | 1.7 | 0.4 | 1.4 | 0.2 | 0.188 |
| OV/BV | 8.4 | 2.3 | 8.2 | 2.0 | 0.901 |
| OS/BS | 41.9 | 7.1 | 43.5 | 5.5 | 0.715 |
| NOb/TAR | 339.7 | 45.2 | 296.0 | 51.5 | 0.230 |
| NOb/BPm | 31.8 | 7.2 | 35.1 | 8.4 | 0.545 |
| NOb/Opm | 75.1 | 6.3 | 79.8 | 10.0 | 0.454 |
| Nob/Obpm | 111.0 | 6.9 | 110.5 | 3.2 | 0.893 |
| ObS/BS | 28.5 | 6.0 | 31.8 | 7.3 | 0.503 |
| ObS/OS | 67.7 | 3.2 | 72.2 | 8.6 | 0.379 |
| NOc/TAR | 26.9 | 4.6 | 24.4 | 3.6 | 0.386 |
| NOc/BPm | 2.5 | 0.6 | 2.9 | 0.6 | 0.413 |
| OcS/BS | 7.6 | 1.8 | 8.9 | 2.5 | 0.418 |



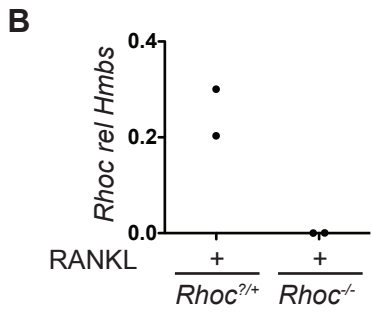
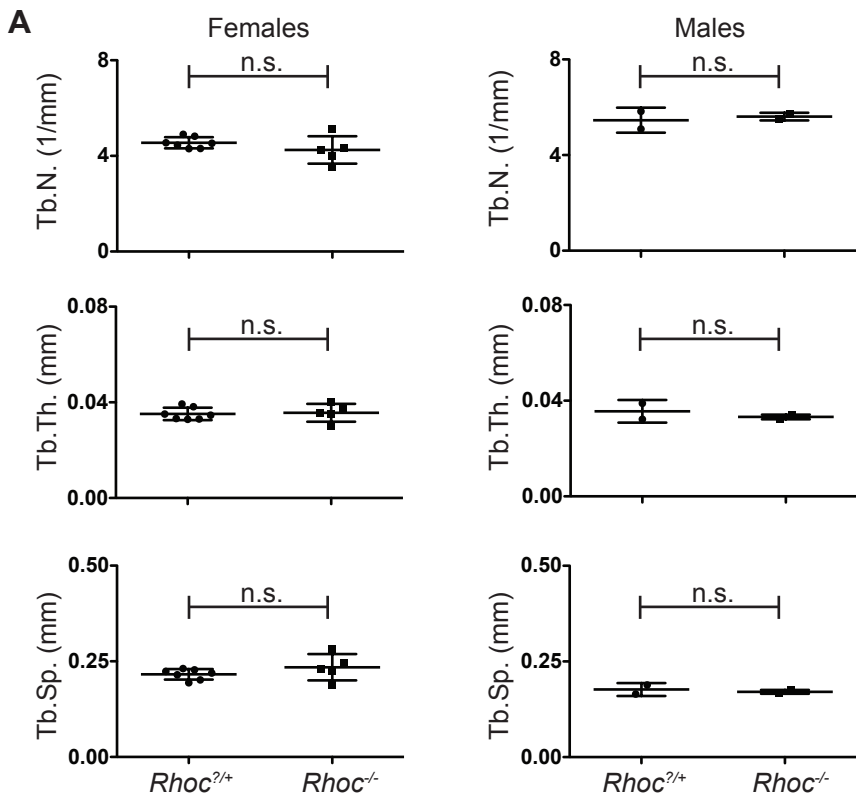
Supplemental Figure 1



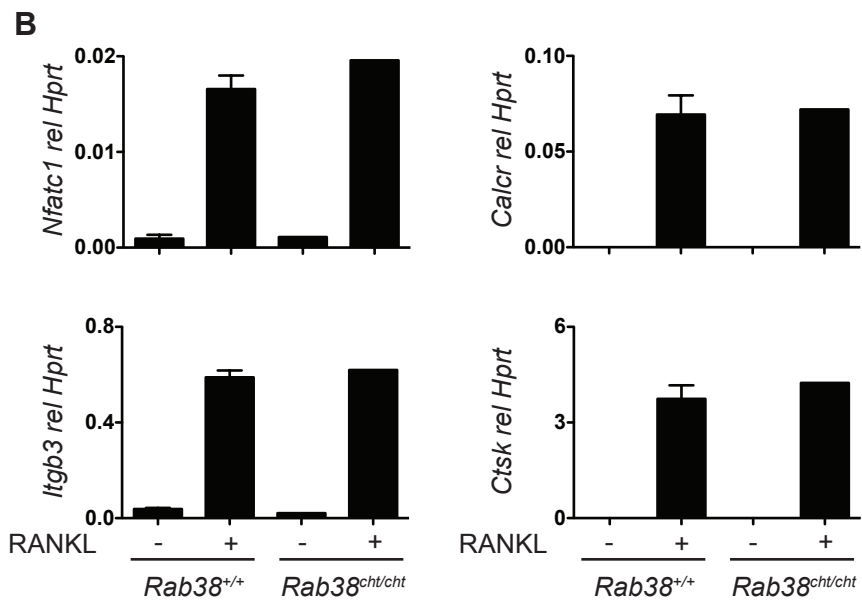
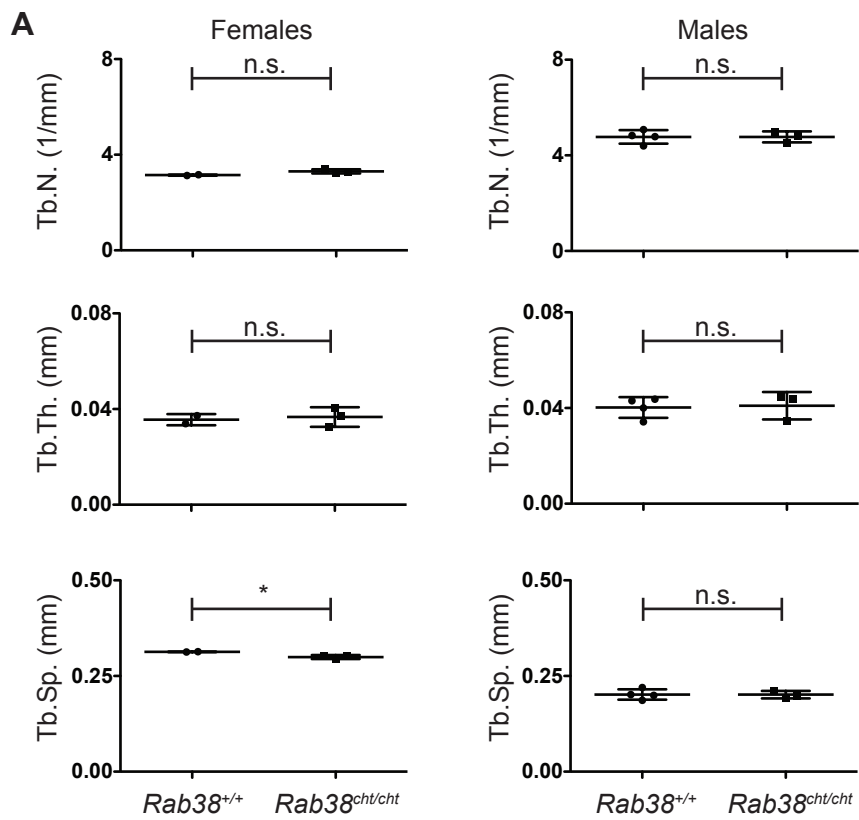
Supplemental Figure 2



Supplemental Figure 3



Supplemental Figure 4



Supplemental Figure 5