

SUPPLEMENTARY TABLE S3. PRIMERS USED IN QUANTITATIVE REVERSE
TRANSCRIPTION-POLYMERASE CHAIN REACTION

| <i>Gene</i> | <i>Accession</i> | <i>Forward</i> | <i>Reverse</i> |
|---------------|------------------|--------------------------|---------------------------|
| <i>Apc</i> | NM_007462.3 | GCTAGCTATGTCCAGCTCCC | CTCTTTACTGCCCCGGGAAT |
| <i>Bap1</i> | NM_027088.2 | ACATGGCAGCTTCCTGACTC | GCTTCTTCCCAACGAGGACA |
| <i>Bmp2</i> | NM_007553.3 | TGCTTCTTAGACGGACTGCG | CTGGGAAGCAGCAACTA |
| <i>Bmp4</i> | NM_007554.2 | CCCGGAAGCTAGGTGAGTTC | AATCCCATCAGGGACGGAGA |
| <i>Bnip3</i> | NM_009760.4 | AAATTAAGGGTGCGTGCGG | CAAAGTGGGGTTCGTGGGTA |
| <i>Cisd2</i> | NM_025902.3 | AAGCTCTTTAGGCCCGAGTG | TGATGATGCTTCCTGCCCTG |
| <i>Dab2</i> | NM_023118.5 | ACTCTGAAATCGACACCAATCA | CAGGCAAGAAGGATGCCCTGA |
| <i>Ep300</i> | NM_177821.6 | GACCCTGTCATGCAAAGCCT | GAAATGATACCTGTTCTGGTAACTG |
| <i>En2</i> | NM_010134.3 | TCCTACGCAGCCTCTCTTCT | CAGGGGTTCATCGACACAGAC |
| <i>Fgf5</i> | NM_010203.5 | AGTCAATGGCTCCCACGAAG | TACAATCCCCTGAGACACAGC |
| <i>Fzd5</i> | NM_022721.3 | CACAGGTACCTAGCTTGTCGT | TGATTAGGGCTCCGACTCCA |
| <i>Fosl1</i> | NM_010235.2 | TCATCTGGAGAGGTGGGTCC | CTGCGGTTCTGACTCACTCG |
| <i>Gata6</i> | NM_010258.3 | TCAGGGGTAGGGGCATCAG | GTAGAGGCCGTCTTGACCTG |
| <i>Hax1</i> | NM_011826.4 | AGCATCTTCAGCGAGATGGG | GCAGTCTCTCACCAGGTGTT |
| <i>Hnf4a</i> | NM_008261.2 | GGATATGGCCGACTACAGCG | AGATGGGGACGTGTCATTGC |
| <i>Jun</i> | NM_010591.2 | GCACATCACCCTACACCGA | GGGAAGCGTGTCTGGCTAT |
| <i>Lef1</i> | NM_010703.4 | CCTGGCATTCCGACATTCCC | GGTCAAAGCAAAGAGCTGCC |
| <i>Mcam</i> | NM_023061.2 | GTGTGTACCTTGGTGCTTGC | ATCACTCTTACGAGTCGGGG |
| <i>MKi67</i> | NM_001081117.2 | CCAAGGCCCAAGTTTGATGC | GACTTGGCCCCGAGATGTAG |
| <i>c-Myc</i> | NM_010849.4 | GTTGGAAACCCCGCAGACAG | ATAGGGGTGTACGGAGTCGT |
| <i>Ncam1</i> | NM_001081445.1 | AGAGGACGGGAACTCCATCA | GAGCGCTCTGTACTTGACCA |
| <i>Olig2</i> | NM_016967.2 | GCTTAGATCATCCCTGGGGC | AGATCATCGGGTCTGGGGA |
| <i>PCNA</i> | NM_011045.2 | TGGTAGTTGTCGCTGTAGGC | ATCAGGCGTGCCTCAAACAT |
| <i>Runx2</i> | NM_001146038.2 | AGTCCATGCAGGAATATTTAAGGC | AAGAAGCTTTGCTGACACGGT |
| <i>Sod1</i> | NM_011434.1 | GGAACCATCCACTTCGAGCA | CCCATGCTGGCCTTCAGTTA |
| <i>Sox7</i> | NM_011446.1 | AAAGTGTTGGGGTGTGCTGA | GCAGACATTGGGCAGAGAGT |
| <i>Sox17</i> | NM_011441.5 | CCCCAACACTCCTCCCAAAG | ACAGTAGCGGGTCCAGAATG |
| <i>Spp1</i> | NM_001204201.1 | CTGGCAGCTCAGAGGAGAAG | TTCTGTGGCGCAAGGAGATT |
| <i>T</i> | NM_009309.2 | CAAGAACGGCAGGAGGATGT | GTACATGGCATTGGGGTCCA |
| <i>Tcf7l1</i> | NM_001079822.2 | GTCGTCCCTGGTCAACGAAT | ATCCTGGGGCCTTCTCACTT |
| <i>Tbx3</i> | NM_011535.3 | CTACGGGGGAGCAATGGATG | CGCTTGGGAAGGCCAAAGTA |
| <i>Trp53</i> | NM_011640.3 | GGCGTAAACGCTTCGAGATG | CTTCAGGTAGCTGGAGTGAGC |