

Supplemental Table 1, Sartori et al

Gene	Forward	Reverse	Probe
Atf4	GGCTATGGATGATGGCTTG	TCGAAACAGAGCATCGAAGT	AGGCATCCTCCTTGCCGGTG
Grp94	GGCTCAAGGACAGATGATGA	CCTGTTCACCTTCAGCTTGGGA	TCTGTGATGCGTTTAACCCATCCAA
XBP (spliced)	CTGAGTCCGCAGCAGGT	TGTCAGAGTCCATGGGAAGA	GGCCCAGTTGTACCTCCCC
XBP (total)	CACCTTCTTGCCCTGCTGGAC	GGGAGCCCTCATATCCACAGT	N/A
Bip/Grp78	AGGAGACTGCTGAGGCAT	CAGCATCTTTGGTTGCTTGT	CCAGCTTACTTCAATGATGCCCAGC
Hspa1a	AAGCAGACGCAGACCTTCA	CTCGTACACCTGGATCAGCA	CTACTCGGACAACCAGCCCCGG
Hspa1L	GGATGCCAAGATGGATAAAG	CTTGTGAGATCCCGTCCAT	TCCACCCGCATCCCAAAGGT
Wsf1	GTCCCATGAAGGCAGATGT	GCTGAGCAGCTGTTGAGTTC	ACGCCTTGCCAACGATGCAG
Herpud2	GTCCTCCAGTTTCCAAAA	TTGAATCCGAATGGTCTGAA	CCAGCACCGATAGAGGAAGT
Edem1	TCTGGTTGATGCCTTGATA	GCTTCGAAGACCTGGACTGT	CATCCGAGTTCCAGAAGGCAGTCA
Hrd1	TGGTGGCTCATGCCTACTAC	AAGAACACCTTGCCCATCAC	TGGCAGTCTGTACATCCAGGCC
Chop	TGGGGGCACCTATATCTCAT	GATGTGCGTGTGACCTCTGT	TGGCCCTGGCTCCTCTGTCA
Bim isoform 1	CCAGGCCTTCAACCACTATC	GCTCCTGTGCAATCCGTATC	ACCTGAAGATCTGCGCCCCGG
Bim isoform 2	ACAAACCCCAAGTCCCTCTT	GCTCCTGTGCAATCCGTATC	ACCTGAAGATCTGCGCCCCGG
Bim isoform 3	ATCGGAGACGAGTTCAACGA	ACCATTTGAGGGTGGTCTTC	TTGCAAATGATTACCGCGAGGC
Bcl2	GCCCTGTGGATGACTGAGTA	CAGCCAGGAGAAATCAAACA	ACCGGCATCTGCACACCTGG
Bak	ATGGCATCTGGACAAGGAC	GCTTCGAAAAGACCTCCTCTG	CTCATCGCAGCCACCTTCG
Bad	GAGGAGGAGCTTAGCCCTTT	CTCTTTGGGCGAGGAAGTC	CAATCTCTGGGCAGCGCAGC
Bax	TTGGAGATGAACTGGACAGC	CCAGTTGAAGTTGCCATCAG	ACATGTCAGCTGCCACCCGG
Fas	GTGGATCTGGGCTGTCTT	TCCTTGATATAATCCTTCTGAGCA	TTCACGAACCCGCCTCCTCA
Insulin	CTTCAGACCTTGCCGTTGGA	ATGCTGGTGCAGCACTGATC	CCCCGGCAGAAGCGTGGCATT
Amylase	AAATCTGCACAAGGTCTGGA	CCACAGGTAAGTCTGTTTCC	CGGACACCAACATTGTTGCACCT CAACCCTGGAAGACACCCCAATCTC G
Gata4	GTCTGCACAGCCTGCCT	TGACACACTCTCTGCCTTCTGAG	
Gata6	CAAGATGAATGGCCTCAGCA	GTTGGCACAGGACAGTCCAA	AAGCGCGTGCCTTCATCACGG
Cyclophilin	CAGACGCCACTGTGCTTT	TGCTTTGGAACCTTGTCTGCAA	CCCTTGGGCCGCGTCTCCTT
Pnli1p1	CCTTGGATGCTTTTCTGATGCTG	GCACATGTCAACCACCCAGTTC	N/A
Pnli1p2	TTGCTATGGACACCTTGGATGC	CAGCAACCAGCCTTCTTCTCC	N/A
Clu	GAGATTGAGAACGCCGTCCA	GCTCCTGCAGACACGTGCATA	N/A
Nr5a2	GCCAAATTGACAAAACGCAGA	GAGGCAAGGCTACATGGCTCA	N/A
Zg16	CCATCACCGCCTTCCGTATC	CAGAGCCAGATCGCCACTAA	N/A
Prss3	CCCCTACCAGGTGTCCCTGA	GCCCCAGCCAGAGATGAGAC	N/A
Pla2gb1	CTGCTGCACACAGCATCAGC	CTCCCGGAGCATGAGTAGGAG	N/A
Gp2	GGCATTGTCAGCCACACAGC	CCTGCCTACAGACACAGGACCA	N/A
Homer2	GCTGCCAATGTGAAGAAGTGG	TGGGACCTCTGACTCCAGCTC	N/A
Tff2	GGCTGACACCCCAACAGAGA	CACCAGGGCACTTCAAAGATCA	N/A
Efna5	CCAGAGGGGTGACTACCACATTG	TCTTCCGTTGTCTGGGATTGC	N/A
Rgs3	CTGTAACCCTGCCCGGACTC	GGAGGTAGAGGGGATTTCTCAGG	N/A
Rnf4	CTGCTGCCAGACCATCAAA	TGCTGCCCACTGAACATCAC	N/A
Prss2	GACCAGTGGGTGGTGTCTGC	TGCCCCAGCCAGAGATGAG	N/A
Gata4 loxp	GAAGACACCCCAATCTCGGTGAGTTAC	GTGAAAACCCCTACAGGAAAGTCATCT	N/A
Gata6 loxp	ACCCTACTATGACCAATTCCAGAACGAT	GTGGAAATAGCTAGAAAAGGCAACCTGT	N/A
CHOP wt	AATTCTACGTGTTAGAAAAGGGACAAG	GAGACAGGGTCTTGTATAGGCTACATT	N/A
Beta gal	GTTGCAGTGCACGGCAGATACACTT	GCCACTGGTGTGGCCATAATTCAA	N/A
Ins-Cre ^{ERT}	GCATTACCGGTGCGATGCAACGAGTGATGAG	GAGTGAACGAACCTGGTCAAAATCAGTGC G	N/A
Ubc-Cre ^{ERT2}	CGCTCGGGGTTGGCGAGTGTGTTTTGTGAA	GCCTGGCGATCCCTGAACATGTCCATCAG G	N/A