

Table S1. Oligonucleotides used for site-directed mutagenesis of AnxA2.

Mutants	Sense	Antisense
Δ25-36	5' - CCAAGTGCATATCGGGATGCTTTGAACATTGAAAC-3'	5' - CATCCCATATGCACTTGGGGGTGT-3'
Δ37-48	5' - GATGCTGAGAAAGGTGTGGATGAGGTCAC-3'	5' - CCACACCTTTCTCAGCATCAAAGTTAGTATAGGC-3'
Δ49-61	5' - CATCAAGACCAACCCGACCAATGCAC-3'	5' - CTGCGGTTGGTCTTGATGGCTGTTTCAATG-3'
Δ62-75	5' - CAACATTTTGACCAGAAGGACCAAAAAGGAACCTTGC-3'	5' - GGTCCCTCTGGTCAAAATGTTGACAATGGTG-3'
Δ37-66	5' - GATGCTGAGCAGAGACAGGATATTGCCTTC-3'	5' - CCTGTCTCTGCTCAGCATCAAAGTTAGTATAGGC-3'
Δ74-88	5' - GCCTTCTCAGCCTTATCTGGCCAC-3'	5' - CCAGATAAGGCTGAGAAGGCAATATCCTGTCTC-3'
Δ82-88	5' - GAAGGACCAAAAAGTCAGCCTTATCTGGCCAC-3'	5' - GATAAGGCTGACTTTTTGGTCTTCTCTGGTAGG-3'
Δ89-101	5' - CAGCACTGAAGCTATTGAAGACACCTGCTCAG-3'	5' - GTCTTCAATAGCTTCAGTGTGATGCAAGTTC-3'
Δ102-108	5' - GATTTTGGGCTATGACGCTTCTGAGCTAAAAG-3'	5' - GAAGCGTCATAGCCCAAAATCACCGTCTC-3'
> 25-27 (AnxA1)	5' - CTACACCCCAAGTGCATATAGCGCTGTC-3'	5' - GACAGCGCTATATGCACTTGGGGGTGTAG-3'
> 28-30 (AnxA1)	5' - GGGTCTGTCAGTCCCTATACTAACTTTG-3'	5' - CAAAGTTAGTATAGGGACTGACAGACCC-3'
> 31-33 (AnxA1)	5' - GTCAAAAGCCTATCCTACCTTTGATGCTGAG-3'	5' - CTCAGCATCAAAGGTAGGATAGGCTTTGAC-3'
> 34-36 (AnxA1)	5' - CTATACTAACTTTAATCCTAGTCGGGATGCTTTG-3'	5' - CAAAGCATCCCAGACTAGGATTAAGTTAGTATAG-3'
> 37-39 (AnxA1)	5' - CTTTGATGCTGAGAGCGATGTTTGAACATTG-3'	5' - CAATGTTCAAACATCGCTCTCAGCATCAAAG-3'
> 40-42 (AnxA1)	5' - CGGGATGCTGCGGCCCTTGAACAGCCATC-3'	5' - GATGGCTGTTTCAAGGCCCGCAGCATCCCG-3'
> 43-45 (AnxA1)	5' - CTTTGAACATTCACAAAGCCATCAAGACC-3'	5' - GGTCTTGATGGCTTTGTGAATGTTCAAAG-3'
> 46-48 (AnxA1)	5' - GAAACAGCCATCATGGTCAAAGGTGTGG-3'	5' - CCACACCTTTGACCATGATGGCTGTTTC-3'
> 49-75 (AnxA1)	5' - AACAAATGCACAGAGACAGCAAATTAAGCCGCTACCAGAGAAGGACCAAAAAGGAACCTTG-3'	5' - CTGGTAGGCGGCTTTAATTTGCTGTCTCTGTGCATGTTGCGTTTGGTCAAATGTGC-3'
> 49-81 (AnxA1)	5' - GGCCACCATTATCGACATTTTGACCAACCGCAGCAATGCACAGAG-3'	5' - CAAAATGTCGATAATGGTGGCCTCATCCACACCTTTGGTC-3'
> 62-75 (AnxA1)	5' - AAACGCAACAATGCACAGAGACAGCAGATTAAGGCCGCTACCAGAGAAGGACC-3'	5' - GGCTTAATCTGCTGTCTCTGTGCATGTTGCGTTTGGTCAAATGTTGACAATGGTG-3'
K28S+D34N+E36S	5' - TCAGCCTATACTAACTTTAATGCTTCGCGGGATGCTTTGAACATTG-3'	5' - CGAAGCATTAAGTTAGTATAGGCTGAGACAGACCCATATGCACTTG-3'
R37S+E43H+K47M	5' - CGGATGCTTTGAACATTCACACAGCCATCATGACCAAGGTGTGGATGAG-3'	5' - CATGATGGCTGTGTAATGTTCAAAGCATCCGACTCAGCATCAAAGTTAGTATAGGC-3'
V98L	5' - GGAGACGCTGATTTTGGCCTATTG-3'	5' - CAATAGGCCCAAAATCAGCGTCTCC-3'