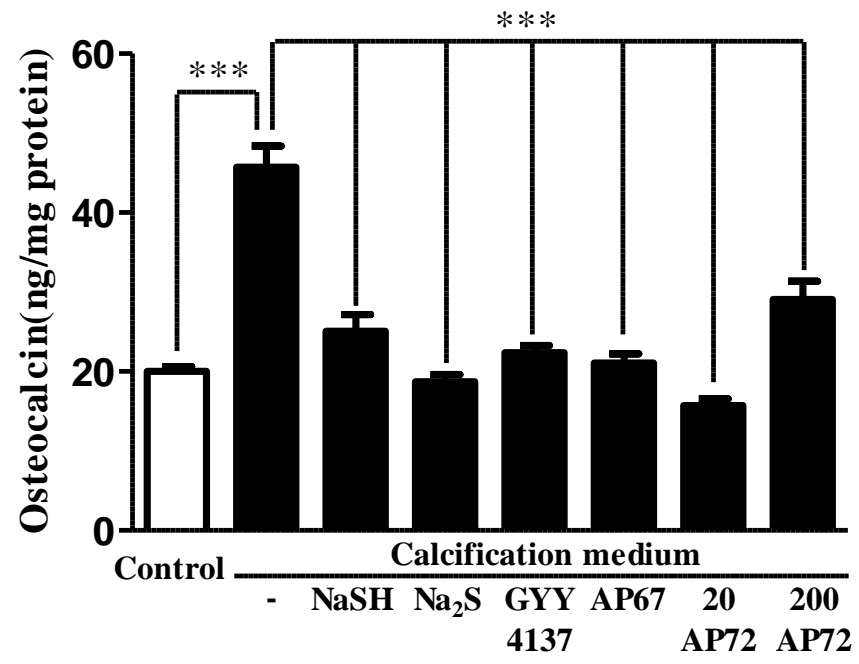
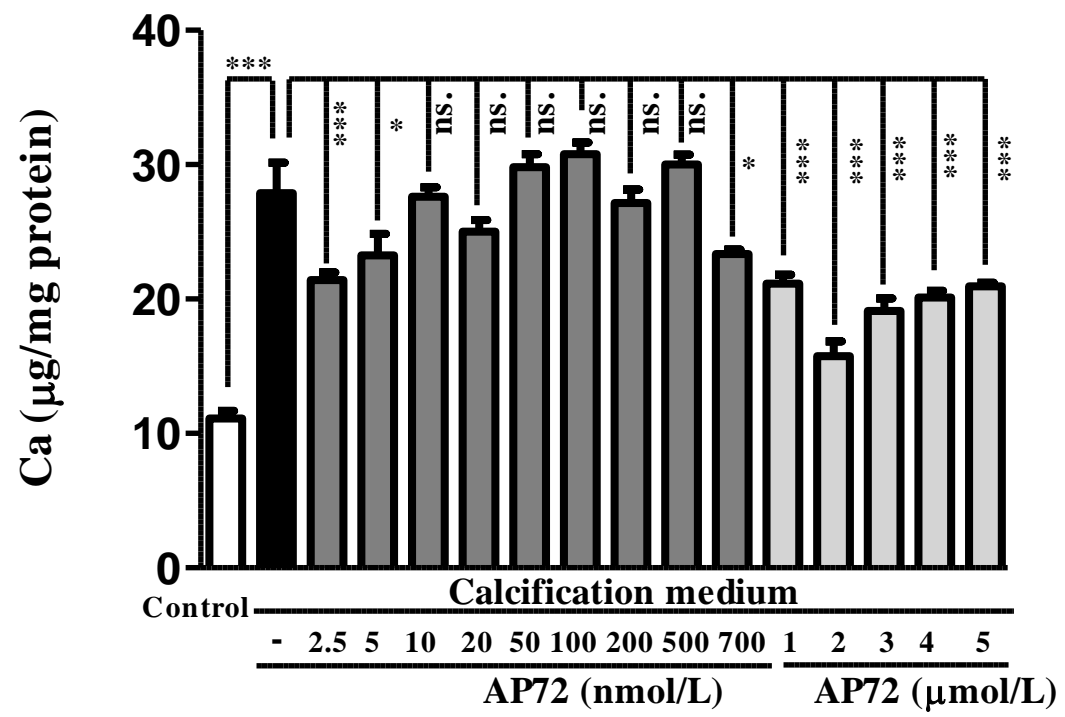


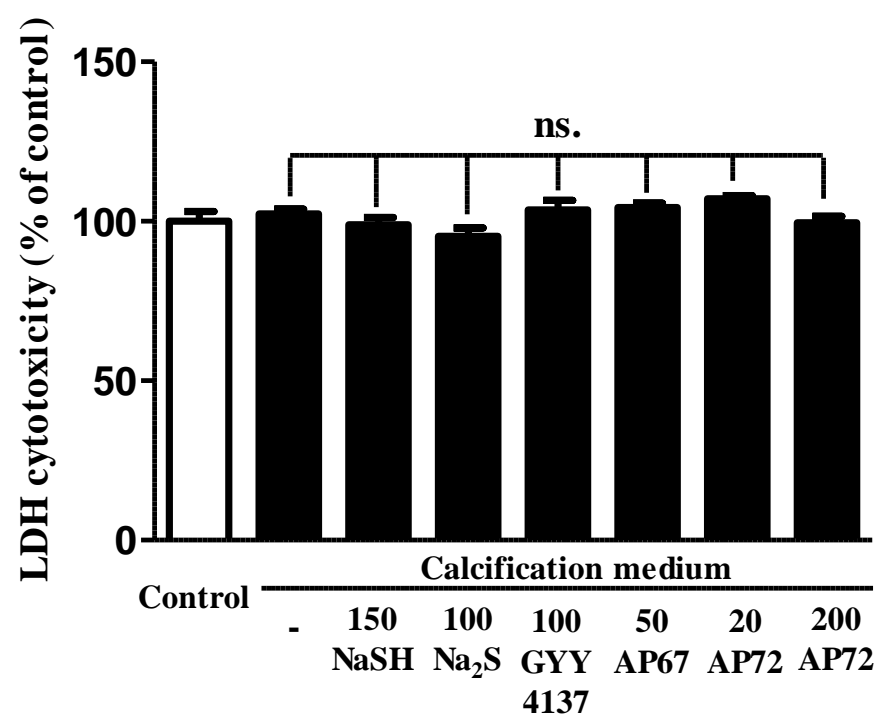
A

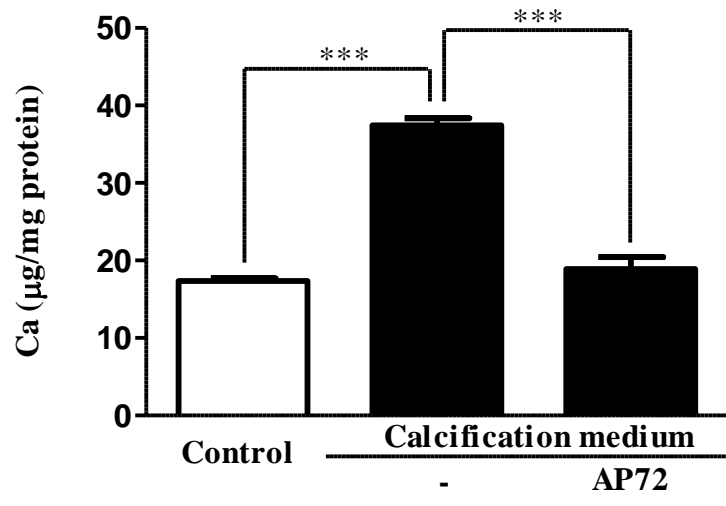
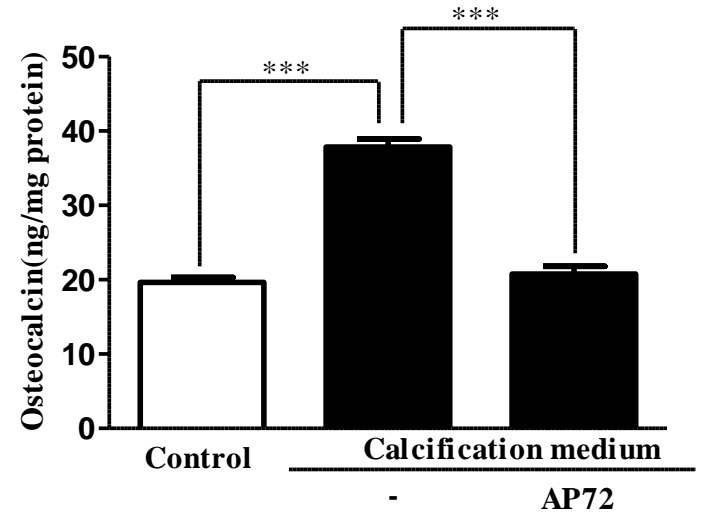
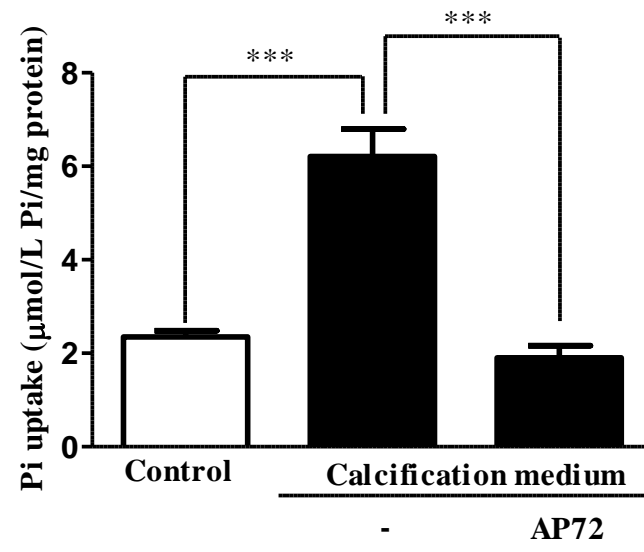
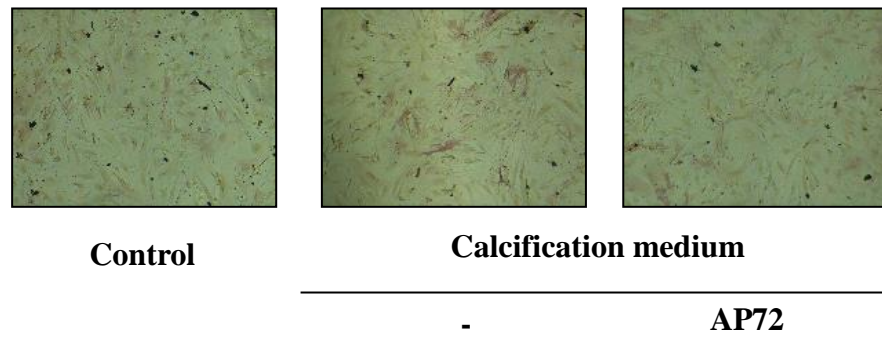
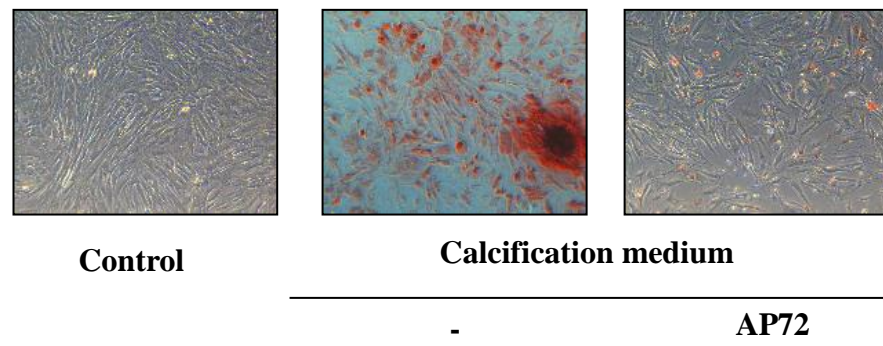
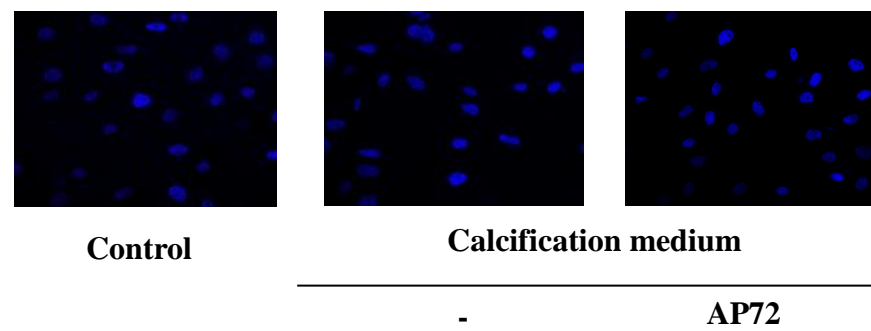


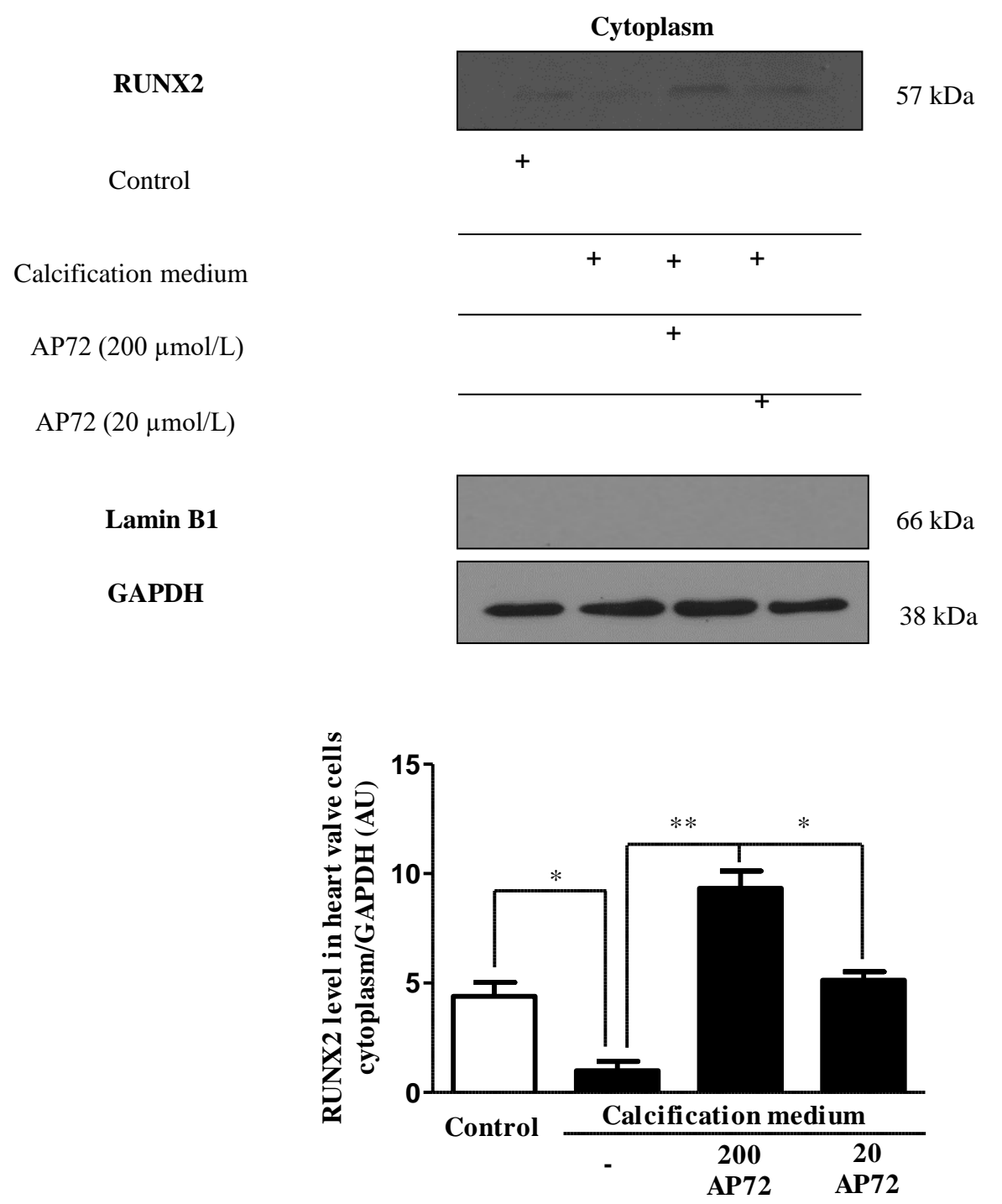
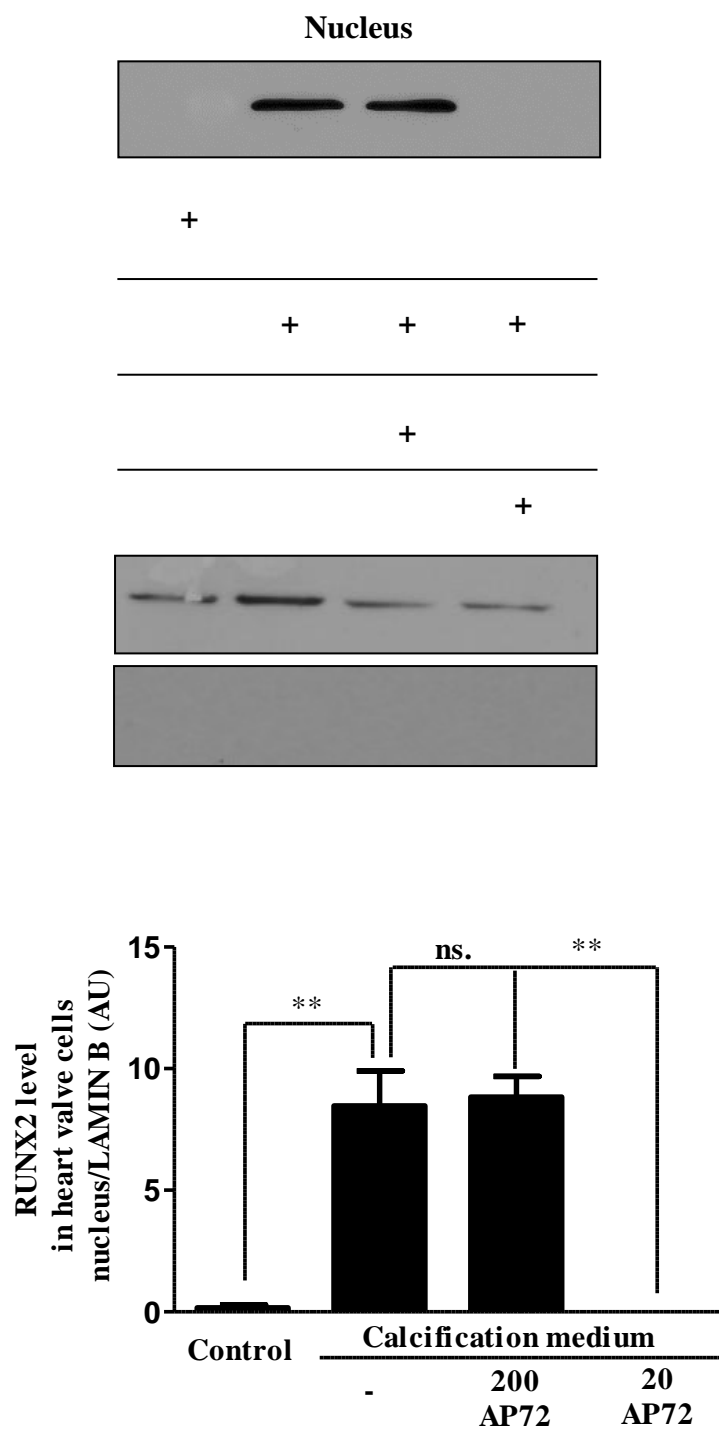
B

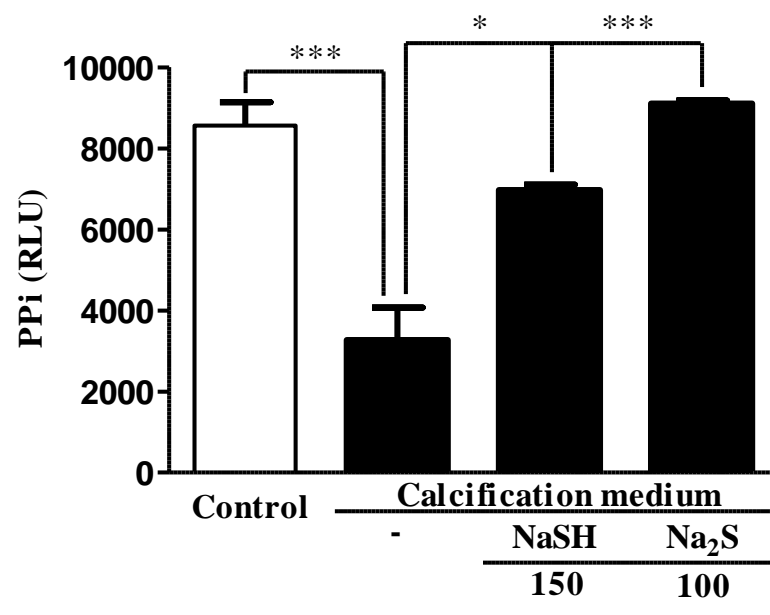


C

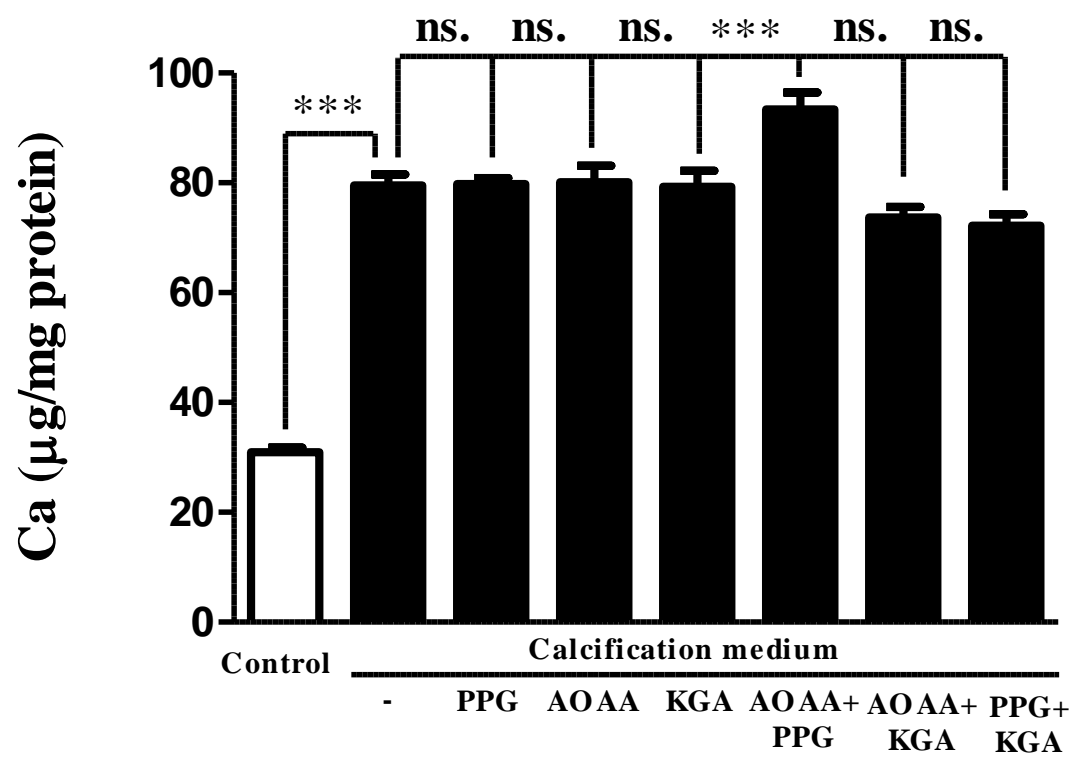


A**B****C****D****E****F**

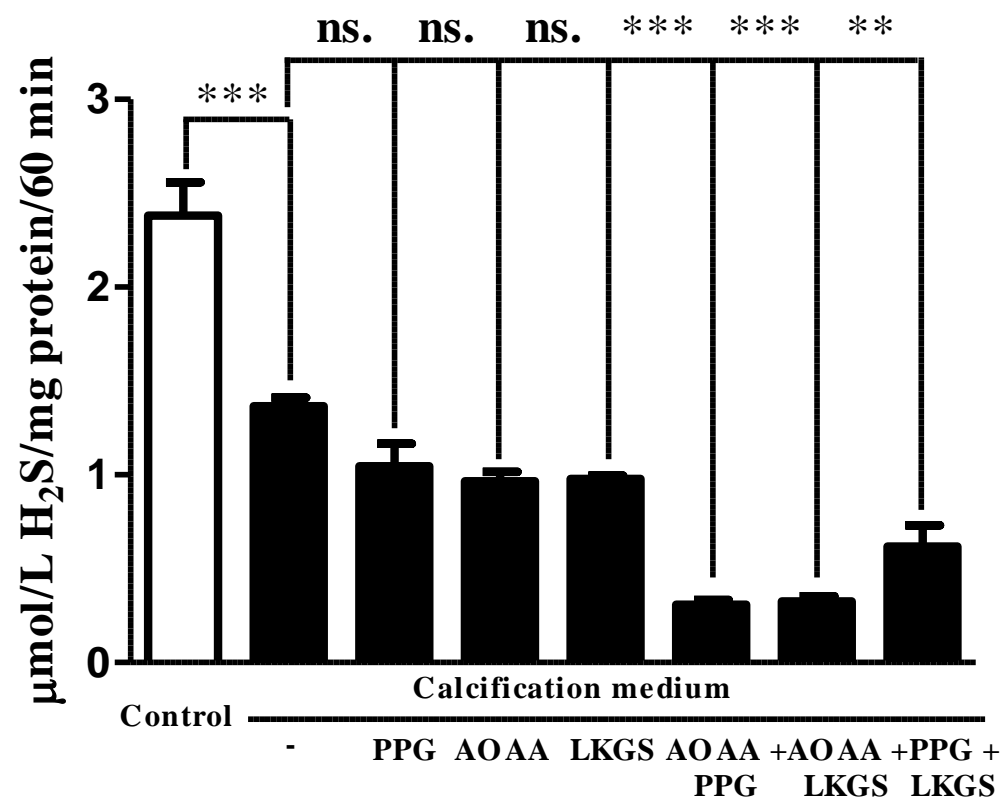




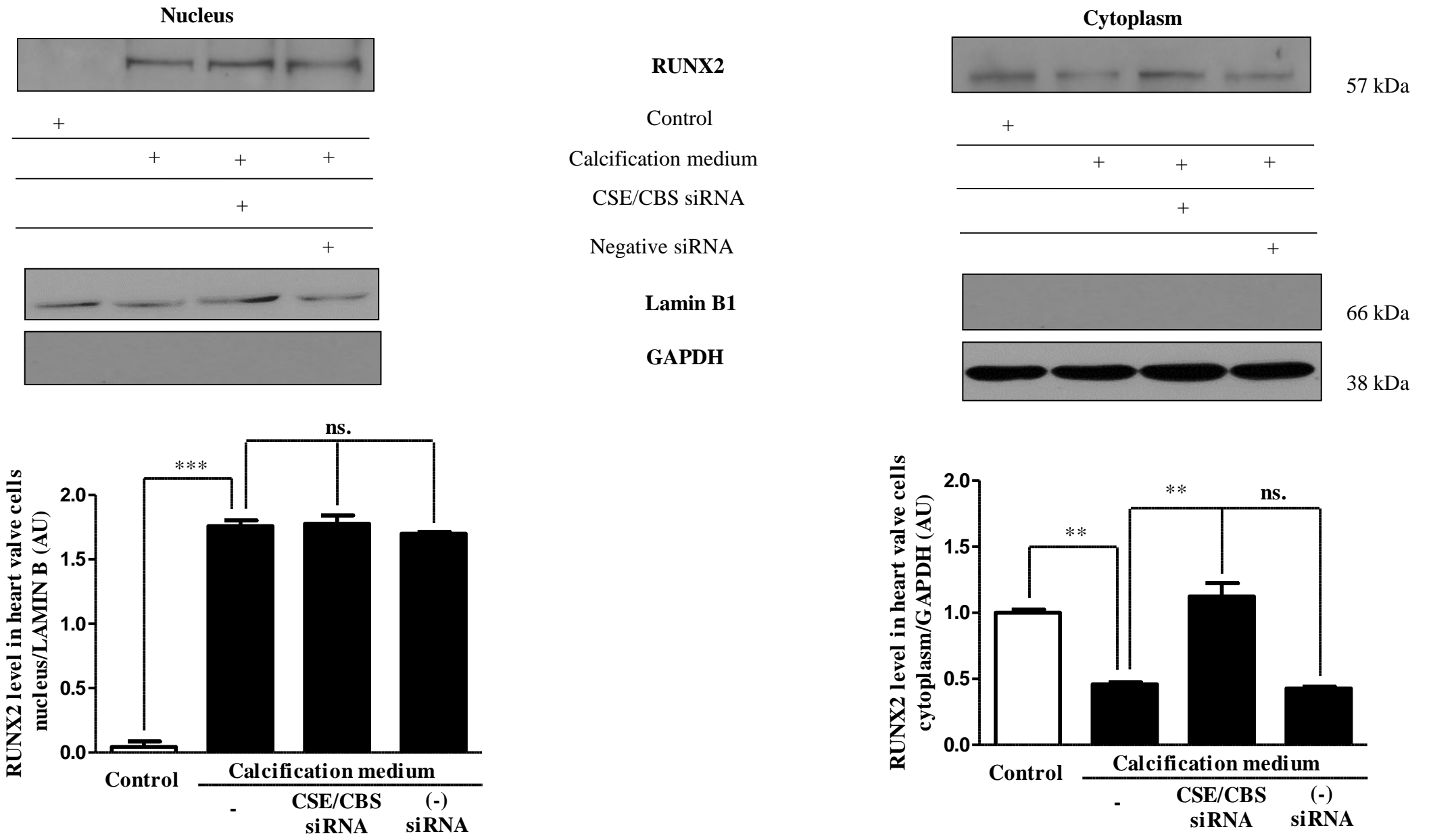
A



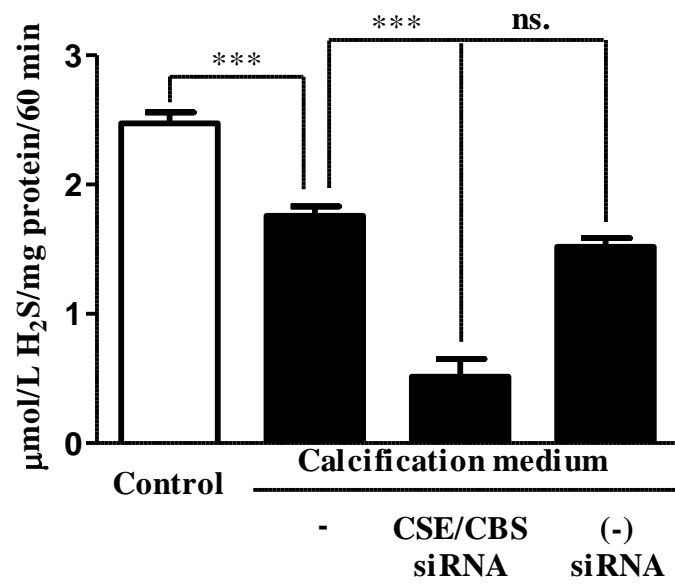
B



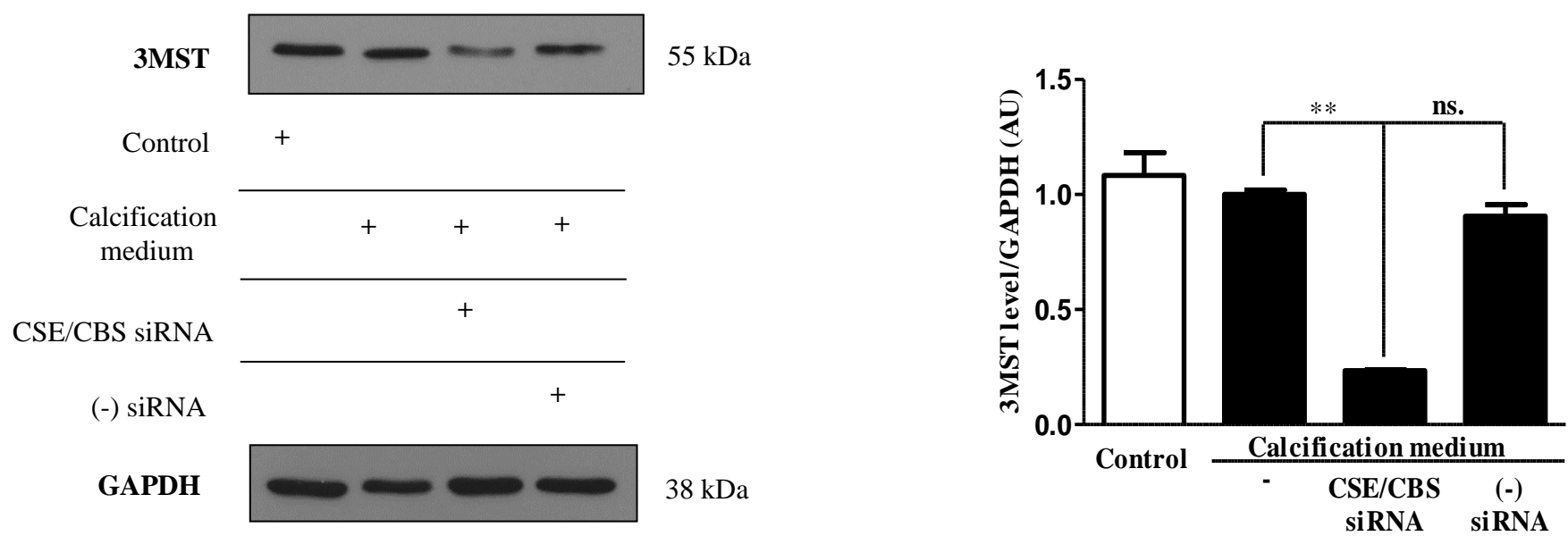
A

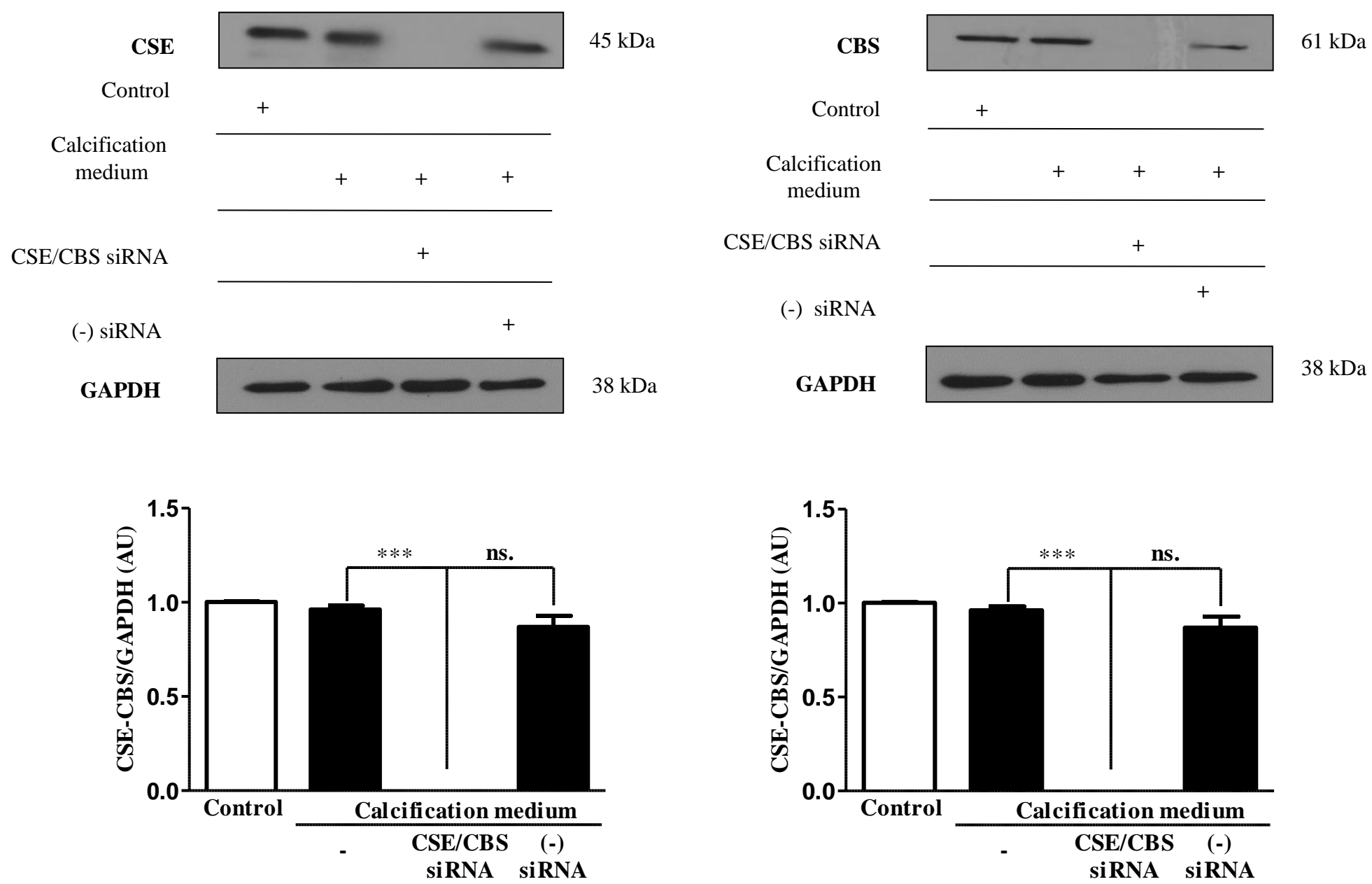
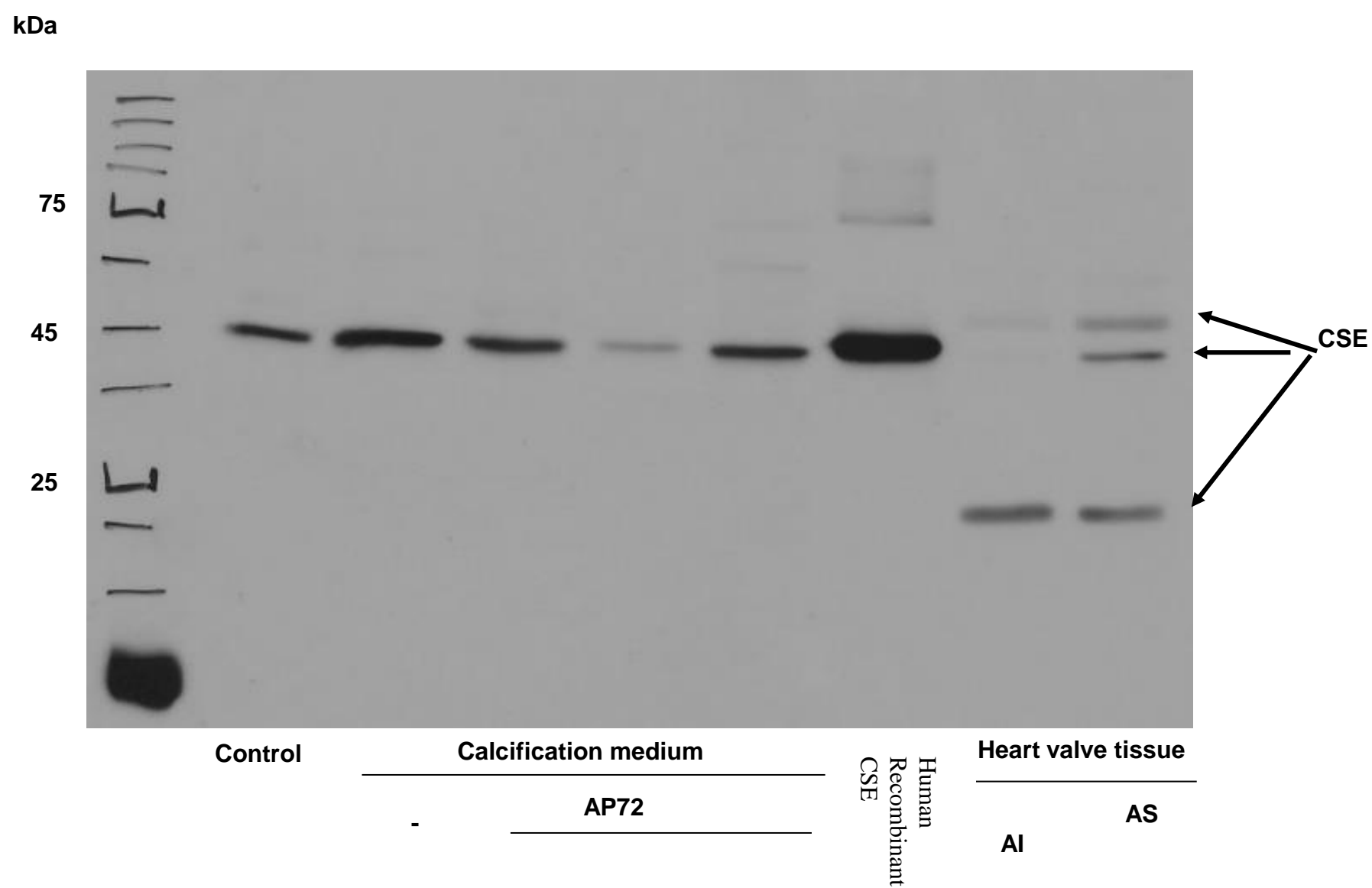


B



C



A**B**

SUPPLEMENTARY FIGURE LEGENDS

Supplementary Figure 1. Osteocalcin levels, calcium deposition and cytotoxicity in VIC

Cells were cultured on 24 well plates for 5 days in growth medium or calcification medium and supplemented with NaSH (150 μ mol/L); Na₂S (100 μ mol/L); GYY4137 (50 μ mol/L); AP67 (20 μ mol/L); AP72 (200 μ mol/L). A) Osteocalcin level of the cells were measured after 5 days and normalized to the protein content of the cells. Graph shows mean \pm SEM of five separate experiments. B) VIC was cultured in phenol red-free growth or calcifying conditions or supplemented with AP72 (2.5 nmol/L to 5 μ mol/L) and calcium content was measured. C) LDH cytotoxicity in VIC exposed to H₂S donors was assessed. All experiments were performed. Results were shown mean \pm SEM at least of six independent experiments. Ns.: not significant; *P < 0.05; ***P < 0.001.

Supplementary Figure 2. In phenol red-free condition AP72 inhibits valvular calcification at 2 μ mol/L

VIC were cultured in growth medium or calcification medium alone (without phenol red) or supplemented with AP72 (2 μ mol/L) for 5 days. A) Calcium content and B) osteocalcin level C) Phosphate uptake of the cells were measured after 5 days and normalized to the protein content of the cells. D) Alkaline phosphatase staining, E) Alizarin Red S staining provides representative microscopic images of extracellular calcium depositions. F) Fluorescence staining assay for cytotoxicity was assessed. Scale bar: 50 μ m. Graph shows mean \pm SEM of five separate experiments. ***P < 0.001.

Supplementary Figure 3. Inhibition of valvular calcification by AP72

VIC was cultured in growth medium or calcification medium alone or supplemented with AP72 (20 μ mol/L) for 5 days. A) Calcium content and B) osteocalcin level of the cells were measured after 5 days and normalized with the protein content of the cells. C) Alkaline phosphatase staining and D) Alizarin Red S staining provide representative microscopic images of extracellular calcium depositions. E) Differences in cellular cytotoxicity is visualized using a fluorescence staining assay (see Methods). Scale bar: 50 μ m. Graph shows mean \pm SEM of five separate experiments. ***P < 0.001.

Supplementary Figure 4. RUNX2 nuclear translocation in calcifying condition in the presence of 20 and 200 μ mol/L of AP72

VIC grown on coverslips were exposed to growth medium or calcification medium alone or supplemented with AP72 (20 and 200 μ mol/L) for 24 hours. A) RUNX2 levels of cytoplasm and nucleus were determined by Western blot analyses. The band intensities are normalized to Lamin B1 in case of nuclear extracts and for GAPDH in case of cytoplasm extracts. Representative staining of five independent experiments was shown. Ns.: not significant; **P < 0.01; ***P < 0.001.

Supplementary Figure 5. Changes of PPI levels by fast sulfide donor treatment on VIC

VIC were cultured in growth medium or calcification medium in the presence or absence of NaSH or Na₂S for 5 days. PPI levels of VIC were measured and normalized to protein content of the cells. Graph shows mean ±SEM of five separate experiments. *P < 0.05; ***P < 0.001.

Supplementary Figure 6. Inhibition of H₂S production by pharmacological inhibitors in VIC

A) Calcium accumulation of extracellular matrix of VIC maintained in growth medium or calcification medium after 5 days treatment with PPG; AOAA; KGA; AOAA+PPG; AOAA+KGA; PPG+KGA (20 μmol/L each inhibitor). Calcium levels of VIC were normalized to protein content of the cells. B) H₂S production of VIC treated with pharmacological inhibitors alone or in combination was detected. Generation of H₂S was normalized to the protein content of cells. Graph shows mean ±SEM of five separate experiments. Ns.: not significant; ***P < 0.001.

Supplementary Figure 7. CSE/CBS silencing enhanced the progression of calcification and decreased the expression of 3-MST

CSE and CBS double gene silencing using siRNAs were performed. Cells were transfected with siRNA against CSE/CBS. Next day cells were washed and treated with high inorganic phosphate and calcium for 3 days. A) RUNX2 expression in a nuclear and cytoplasmic fraction were shown. B) H₂S production of CSE/CBS silenced samples was measured. C) Western blot demonstrated expression of 3-MST in CSE/CBS silenced VIC. Graph shows mean ±SEM of five independent experiments. Ns.: not significant; **P < 0.01; ***P < 0.001.

Supplementary Figure 8. CSE and CBS expression of the double silenced VIC

A) VIC were cultured in growth or calcification medium and transfected with siRNA for CSE/CBS. Next day VIC were treated with calcification medium for 3 days. CSE (left panel) and CBS (right panel) protein level were measured. Samples were normalized to GAPDH. B) Western blot of the CSE protein of VIC lysate and AI/AS were shown. Human recombinant CSE protein (5 and 10 ng) was used as a positive control. Ns.: not significant; ***P < 0.001.

Sequences of the siRNA

CTH si RNA sequence:

>gnl|UG|Hs#S6091963 Homo sapiens cystathionase (cystathionine gamma-lyase) (CTH), transcript variant 2, mRNA /cds=p(199,1284) /gb=NM_153742 /gi=299473760 /ug=Hs.19904 /len=2008

TCGTGAGACTGCAGACCCGCTAATAAAAATCCACCCCAACAATCGCTGTGTGCCGCTTTAG
TGCGCTCGCCGTCGGCTCTACCTGCGTGCTTTAGCTCCTTCTCGCCTGATCCTTCTGTCT
CTCCCAACCCCGGACACCCGGCTTCGACTGGTTATATCTTCGGTGTTCCTTTCTCTCTT
CTTCTTTTCGCGGTTTCAGCATGCAGGAAAAAGACGCCTCCTCACAAGGTTTCCTGCCACAC
TTCCAACATTTTCGCCACGCAGGCGATCCATGTGGGCCAGGATCCAGAGCAATGGACCTCC
AGGGCTGTAGTGCCCCCATCTCACTGTCCACCACGTTCAAGCAAGGGGCGCCTGGCCAG
CACTCGGGTTTTGAATATAGCCGTTCTGGAAATCCCCTAGGAATTGCCTTGAAAAAGCA
GTGGCAGCACTGGATGGGGCTAAGTACTGTTTGGCCTTTGCTTCAGGTTTAGCAGCCACT
GTA ACTATTACCATCTTTTAAAAGCAGGAGACCAAATTATTTGTATGGATGATGTGTAT
GGAGGTACAAACAGGTACTTCAGGCAAGTGGCATCTGAATTTGGATTAAAGATTTCTTTT
GTTGATTGTTCCAAAATCAAATTA CTAGAGGCAGCAATTACACCAGAAACCAAGCGCCCT
TTGGCTCTGGGAGCTGATATTTCTATGTATTCTGCAACAAAATACATGAATGGCCACAGT
GATGTTGTAATGGGCCTGGTGTCTGTTAATTGTGAAAGCCTTCATAATAGACTTCGTTTC
TTGCAAAACTCTCTTGGAGCAGTTCATCTCCTATTGATTGTTACCTCTGCAATCGAGGT
CTGAAGACTCTACATGTCCGAATGGAAAAGCATTTCAAAAACGGAATGGCAGTTGCCAG
TTCTGGAATCTAATCCTTGGGTAGAAAAGGTTATTTATCCTGGGCTGCCCTCTCATCCA
CAGCATGAGTTGGTGAAGCGTCAGTGTACAGGTTGTACAGGGATGGTCACCTTTTATATT
AAGGGCACTCTTCAGCATGCTGAGATTTTCCTCAAGAACCTAAAGCTATTTACTCTGGCC
GAGAGCTTGGGAGGATTCGAAAGCCTTGCTGAGCTTCCGGCAATCATGACTCATGCATCA
GTTCTTAAGAATGACAGAGATGTCCTTGGAAATTAGTGACACACTGATTCGACTTTCTGTG
GGCTTAGAGGATGAGGAAGACCTACTGGAAGATCTAGATCAAGCTTTGAAGGCAGCACAC
CCTCCAAGTGGAAGTCACAGCTAGTATTCCAGAGCTGCTATTAGAAGCTGCTTCCTGTGA

AGATCAAATCTTCCTGAGTAATTAATGGACCAACAATGAGCCTTTGCAAATTTTCAAG
CGGAAATTTTAAGGCACCTCATTATCTTTCATAACTGTAATTTTCTTAGGGATCATCTCT
GTTAAAAAGTTTTCTGTATGTCATGTTATAATTACAGGTCAATTCTGTTAATATCTTTTT
GTTAATTTTGCTCTATGTTTGCCTCTGAAGGAGGTGAGATTTGTGCTACTTTGGGAGATT
ATGTTCTTTTTTCATGTCTAAGATTTATTTTGATCATGTTTATAATATAATGGTAATTCA
TTTTTGATGTTTTGTGAAGAATTTAAATTTAAACGAATGTTCTTAAATCAAGTGTGATTT
TTTTGCATATCATTGAAAAGAACATTAAGCAATGGTTTACACTTAGTTACCATAAGCC
GAAAATCAAATACTTGAAAAGTTTACTGTGAAATTCTACTGATTTAAGACTATACTTAAT
ATTTTTAAAAAATAAATCAGCTGGGCGCGGTGGCTCACGCATGTAATGCCAGCACTTTT
GGAGGATAAGGCGGGCGGATCACGAGGTCAGGAGATTGAGACCATCCTGGCTAGCGCAGT
GAAACCCCATCTCTACTAAAAATGCAAAAAAATTAGACGGACGTGGTGGCGGGTGCCT
GTAGTCCCAGCTACTTGGGAGGCTGAGG

CBS siRNA sequence

>gnl|UG|Hs#S55481398 Homo sapiens cystathionine-beta-synthase (CBS), transcript variant 2, mRNA
/cds=p(223,1878) /gb=NM_001178008 /gi=295821199 /ug=Hs.533013 /len=2586

CACGCCCTCGGGGTCGGTCCTCGAGGACGCGCAGGGCCCCCACCCACCAGGACGCACGT
TTCAAGCTCATCAGTAAAGGTTTCTTAAATTCCCGAAGGGCAAGAAGTTAACCAAGTAAA
ACAGCATCGGAACACCAGGATCCCATGACAGATTCTGTTGTCACGTCTCCTTACAGAGTT
TGAGCGGTGCTGAACTGTCAGCACCATCTGTCCGGTCCCAGCATGCCTTCTGAGACCCCC
CAGGCAGAAGTGGGGCCACAGGCTGCCCCACCGCTCAGGGCCACACTCGGCGAAGGGG
AGCCTGGAGAAGGGGTCCCCAGAGGATAAGGAAGCCAAGGAGCCCCTGTGGATCCGGCCC
GATGCTCCGAGCAGGTGCACCTGGCAGCTGGGCCGGCCTGCCTCCGAGTCCCCACATCAC
CACACTGCCCCGGCAAAATCTCCAAAAATCTTGCCAGATATTCTGAAGAAAATCGGGGAC
ACCCCTATGGTCAGAATCAACAAGATTGGGAAGAAGTTCGGCCTGAAGTGTGAGCTCTTG
GCCAAGTGTGAGTTCTTCAACGCGGGCGGGAGCGTGAAGGACCGCATCAGCCTGCGGATG
ATTGAGGATGCTGAGCGCGACGGGACGCTGAAGCCCGGGGACACGATTATCGAGCCGACA

TCCGGGAACACCGGGATCGGGCTGGCCCTGGCTGCGGCAGTGAGGGGCTATCGCTGCATC
ATCGTGATGCCAGAGAAGATGAGCTCCGAGAAGGTGGACGTGCTGCGGGCACTGGGGGCT
GAGATTGTGAGGACGCCACCAATGCCAGGTTGACTCCCCGGAGTCACACGTGGGGGTG
GCCTGGCGGCTGAAGAACGAAATCCCCAATTCTCACATCCTAGACCAGTACCGCAACGCC
AGCAACCCCTGGCTCACTACGACACCACCGCTGATGAGATCCTGCAGCAGTGTGATGGG
AAGCTGGACATGCTGGTGGCTTCAGTGGGCACGGGCGGCACCATCACGGGCATTGCCAGG
AAGCTGAAGGAGAAGTGTCTGGATGCAGGATCATTGGGGTGGATCCCGAAGGGTCCATC
CTCGCAGAGCCGGAGGAGCTGAACCAGACGGAGCAGACAACCTACGAGGTGGAAGGGATC
GGCTACGACTTCATCCCCACGGTGCTGGACAGGACGGTGGTGGACAAGTGGTTCAAGAGC
AACGATGAGGAGGCGTTCACCTTTGCCCGCATGCTGATCGCGCAAGAGGGGCTGCTGTGC
GGTGGCAGTGCTGGCAGCACGGTGGCGGTGGCCGTGAAGGCCGCGCAGGAGCTGCAGGAG
GGCCAGCGCTGCGTGGTCATTCTGCCCGACTCAGTGCGGAACTACATGACCAAGTTCCTG
AGCGACAGGTGGATGCTGCAGAAGGGCTTTCTGAAGGAGGAGGACCTCACGGAGAAGAAG
CCCTGGTGGTGGCACCTCCGTGTTTACAGGAGCTGGGCCTGTCAGCCCCGCTGACCGTGCTC
CCGACCATCACCTGTGGGCACACCATCGAGATCCTCCGGGAGAAGGGCTTCGACCAGGCG
CCCGTGGTGGATGAGGCGGGGGTAATCCTGGGAATGGTGACGCTTGGGAACATGCTCTCG
TCCCTGCTTGCCGGGAAGGTGCAGCCGTCAGACCAAGTTGGCAAAGTCATCTACAAGCAG
TTCAAACAGATCCGCCTCACGGACACGCTGGGCAGGCTCTCGCACATCCTGGAGATGGAC
CACTTCGCCCTGGTGGTGCACGAGCAGATCCAGTACCACAGCACCGGGAAGTCCAGTCAG
CGGCAGATGGTGTTCGGGGTGGTCACCGCCATTGACTTGCTGAACTTCGTGGCCGCCAG
GAGCGGGACCAGAAGTGAAGTCCGGAGCGCTGGGCGGTGCGGAGCGGGCCCCGCCACCCTT
GCCACTTCTCCTTCGCTTTCCTGAGCCCTAAACACACGCGTGATTGGTAACTGCCTGGC
CTGGCACCGTTATCCCTGCACACGGCACAGAGCATCCGTCTCCCCTCGTTAACACATGGC
TTCCTAAATGGCCCTGTTTACGGCCTATGAGATGAAATATGTGATTTTCTCTAATGTAAC
TTCCTTTAGGATGTTTACCAAGGAAATATTGAGAGAGAAGTCGGCCAGGTAGGATGAA
CACAGGCAATGACTGCGCAGAGTGGATTAAAGGCAAAAGAGAGAAGAGTCCAGGAAGGGG

CGGGGAGAAGCCTGGGTGGCTCAGCATCCTCCACGGGCTGCGCCGTCTGCTCGGGGCTGA
GCTGGCGGGAGCAGTTTGCGTGTTGGGTTTTTTAATTGAGATGAAATTCAAATAACCTA
AAAATCAATCACTTGAAAGTGAACAATCAGCGGCATTTAGTACATCCAGAAAGTTGTGTA
GGCACCACCTCTGTCACGTTCTGGAACATTCTGTCATCACCCCGTGAAGCAATCATTTC
CCTCCCGTCTTCCTCCTCCCCTGGCAACTGCTGATCGACTTTGTGTCTCTGTTGTCTAAA
ATAGGTTTTCCCTGTTCTGGACATTTTCATATAAATGGAATCACACAAAAAAAAAAAAAAAA
AAAAAA.