

Supplementary Table S1. Homologous protein alignment sequence

| Name | Gene ID | Sequence |
|---------------------|-------------------------|--|
| AtNST1 (ANAC043) | AT2G46770 | MNLGVNGQSQVPPGFRFHPTEELLHYYLKKK VANQKI DLDVIQDIDLNLKEPWDIQEKC KIGSTPQNDWYFFSHKD KKYPTGTRTNRATTAGFWKATGRDKAICGSSRRIGMRK TLVFYKGRAPHGEKSDWIMHEYRLDDNIHELNVSNL SG DRDLGGDEGVVCRVFRKKNYVKALHNPTNNRNSTH MMIRSKKHELDQDGFMHLPQLDSPSTNGVHEPIQQMNQ DGSDSMFNYEDDHNQEAPNSTVNWL VELDRLVAYQLNG QASEASESTTNHLSCFNNHHLNHDSDITPFCLIPAGDYDR SNNNDRTTTAATTQDLLRSQSSRVFDNDSIDDHFWSF TKQSSSPSSLDPISHFSV |
| AtNST2 | AT3G61910 | MNISVNGQSQVPPGFRFHPTEELLKYYLRKKISNIKIDL DVIPDIDLNLKEPWDIQEEMCKIGTPQNDWYFYSHKDK KYPTGTRTNRATTVGFWKATGRDKTIYTNGDRIGMRKT LVFYKGRAPHGQKSDWIMHEYRLDESVLISSCGDH DVN VETCDVIGSDEGVVCRVFKNNLCKNMISSSPASSVK TPSFNEETIEQLLEV MGQSCKG EIVLDPFLKLPNLECHNN TTITSYQWLIDDVN NCHVSKVMDPSFITSWAALDRLV ASQLNGPNNSYSIPAVNETSQSPYHGLNRSGCNTGLTPDY YIPEIDLWNEADFARTTCHLLNGSG |
| AtNST3 | AT1G32770 | MADNKVNLSINGQSKVPPGFRFHPTEELLHYYLRKKV NSQKIDLDVIREVDLNKLEPWDIQEECRIGSTPQNDWYF FSHKDKKYPTGTRTNRATVAGFWKATGRDKIICSCVRRI GLRKTLVFYKGRAPHGQKSDWIMHEYRLDDTPMSNGY ADVVTEDPMSYNEEGWVVCRVFRKKNYQKIDDCPKITL SSLPDDTEEKGPTFHNTQNV TGLDHVLLYMDRTGSNIC MPESQTTQHQDDVLFMQLPSLETPKSESPVDQSFLTPS KLDfspvQEKITERPVCSNWASLDRLVAWQLNNGHNP CHRKSFDEEEENGDTMMQRWDLHWNNDDNVDLWSSF TESSSSLDPLLHLSV |
| FtNAC16 | FtPinG00003 81200.01 | MNLGVNGQSQVPPGFRFHPTEELLHYYLKKK VANQKI DLDVIQDIDLNLKEPWDIQEKC KIGSTPQNDWYFFSHKD KKYPTGTRTNRATTAGFWKATGRDKAICGSSRRIGMRK TLVFYKGRAPHGEKSDWIMHEYRLDDNIHELNVSNL SG DRDLGGDEGVVCRVFRKKNYVKALHNPTNNRNSTH MMIRSKKHELDQDGFMHLPQLDSPSTNGVHEPIQQMNQ DGSDSMFNYEDDHNQEAPNSTVNWL VELDRLVAYQLNG QASEASESTTNHLSCFNNHHLNHDSDITPFCLIPAGDYDR SNNNDRTTTAATTQDLLRSQSSRVFDNDSIDDHFWSF TKQSSSPSSLDPISHFSV |
| GaNAC43 | XP_0176296 | MPENMSISVNGQSQVPPGFRFHPTEELLQYYLRKKVSY |

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| | 05.1 | EKIDLDVIRDVDLNKLEPWDIQCESCKIGTPQNDWYFFS HKDKKYPTGTRTNRATAAGFWKATGRDKVIYSNCRRIG MRKTLVFYKGRAPHGQKSDWIMHEYRLDDNIVETTVSN AMGEGTQEEGWVVCRIFKKKNHHKTLDNPNSSLGSES RNHMLSACNEGELEQILEHMRRNCNEEGVANNSWRLM RPIETAISNSYPDSFMKLPSESPNSTSSQNCYQPMIVDNE GSITNQMSGDPNSRLTNWAALDRLVASQLNGQTETSRQ LACFNDHSISMGYSNPSADHHHDLQSPALRYNRSYNG TQDYNSEMDLWSLTRSSSSDPLCHVVNASV |
| SmNAC43 | QBZ39066.1 | SQVPPGFRFHPTEEELLQYYLKKKVASEKIDLDVIQDVD LNKLEPWDIQCCTIGSTPQNDWYFFSHKDVKYPTGTRT NRATASGFWKATGRDKVIYGNNSRRIGMRKTLVFYKGRA PRGQKSDWIMHEYRLDDNTTANITHNNHGSNAMGEG MQEEGWVVCRVFKKKNHSHKTLDTTPISSWSMLSTIN SCKEGTLEQMLQSIGTPFRRNHFPDSNNTRFLTLSNLH DSPNSSRTQNCYQSLHVQ |
