Supplementary Material

**Table S1. The accession number and source organnisms of the 35 SNAT that was used to construct the phylogenetic tree, SNAT proteins from bacteria, algae, mosses, ferns, gymnosperms and angiosperms**

|  |  |
| --- | --- |
| Accession number | Source organisms |
| AK659369  | *Oryza sativa* |
| XP\_015648698.1  | *Oryza sativa* |
| At1g32070  | *Arabidopsis thaliana* |
| At1g26220 | *Arabidopsis thaliana* |
| NP\_442603  | *Cyanobacterium synechocystis* |
| XM\_002983152  | *Selaginella moellendorffii* |
| NP\_001143827  | *Zea mays* |
| XP\_003568235.1  | *Brachypodium distachyon* |
| XP\_002266361.1  | *Vitis vinifera* |
| XP\_002323094.1  | *Populus trichocarpa* |
| XM\_001782439  | *Physcomitrella patens*  |
| XM\_002955986  | *Volvox carterif.nagariensis* |
| NC\_007932 XP\_002439969.1  | *Pyropia yezoensis**Setaria italica* |
| PTQ28360  | *Marchantia polymorpha* |
| PTQ28361  | *Marchantia polymorpha* |
| WP\_011388274  | *Rhodospirillum rubrum* |
| WP\_011389455 CO176218 XP\_009780441.1 XP\_009774327.1 XP\_009614145.1XP\_009611002.1  | *Rhodospirillum rubrum**Pinus taeda**Nicotiana sylvestris**Nicotiana sylvestris**Nicotiana tomentosiformis**Nicotiana tomentosiformis* |
| XP\_016455489.1 | *NtSNAT1* |
| XP\_016498446.1 | *NtSNAT2* |
| XP\_016474869.1 | *NtSNAT3* |
| XP\_016446764.1 | *NtSNAT4* |
| XP\_016447514.1 | *NtSNAT5* |
| XP\_016450139.1 | *NtSNAT6* |
| XP\_016468822.1 | *NtSNAT7* |
| XP\_016481220.1 | *NtSNAT8* |
| XP\_016501830.1 | *NtSNAT9* |
| XP\_016506039.1 | *NtSNAT10* |
| XP\_016509944.1 | *NtSNAT11* |
| XP\_016510132.1 | *NtSNAT12* |

|  |  |
| --- | --- |
| **Primers Name** | **Primer sequence** |
| NtSNAT1-FNtSNAT1-RNtSNAT2-FNtSNAT2-RNtSNAT3-FNtSNAT3-RNtSNAT4-FNtSNAT4-RNtSNAT5-FNtSNAT5-RNtSNAT6-FNtSNAT6-RNtSNAT7-FNtSNAT7-FNtSNAT8-FNtSNAT8-RNtSNAT9-FNtSNAT9-RNtSNAT10-FNtSNAT10-RNtSNAT11-FNtSNAT11-RNtSNAT12-FNtSNAT12-RActin-FActin-R |

|  |
| --- |
| GACTCAACCAGATGGAACAGTCG |
| GCTTCTTTTCTCCGTTCCCCT |
| TCCGGGTTTGTTAAGAATAACACTG |
| CTTCGAGGCCATCCAACCTT |
| ATGTGACTGGAATATTGACCGTGGA |
| TGTGCCTCCGCTTTAGTTATGA |
| TCCTGGGGAAAGGGTGAATG |
| CACGAACACCTTTTCTGCACC |
| GAGCATCAAATACGCCAGCC |
| GCCGCTTAAGGGGGATGATA |
| AGGTTATCAGCTGTCTTGGCA |
| TTGAAACCTTGTAAGGCGGC |
| CTTAATTCCGACTGCCCACC |
| GGTAAAGGTAGTGAAGGGAGGC |
| CTCAACAACTTTCCCAACAAAACC |
| CGATACGTTCCGTCAATGCT |
| ATGGCAATCCTCATTACACCAT |
| GGTGAAGAAATTTGTTGGGAAAGT |
| GTGGACCTGTTTGTGAGAGC |
| CAATTCATCCGGAGTTACTGGC |
| TCGACGTCAGGGTATTGCTT |
| AGAGACGACGAAGTTTCTTCCA |
| CGACGTTCCCCATATCCACAA |
| ACTTCGAGGAGGAAGATGGTGAACTGGGACGATATGGAGAACACTGGCGTATAGGGACAAC |

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**Table S2. Primers used in this paper.**

**Table S3. The full amino acid sequences of the twelve NtSNAT.**

**NtSNAT1**

MQMQTLHLLSTSPVTASSSLSNFVSLNCCRCQFSNPLPFPCKTNLDFVKVKRQSKVSNLKAGFWESIRSGFVKNNTIQVIESPSSEEEEEEEPLPEEFVLIEKTQPDGTVEQIIFSSGGDVDVYDLQALCDKVGWPRRPLSKLAAALKNSYIVATLHSRKFSSGEEGNGEKKLIGMARATSDHAFNATIWDVLVDPSYQGQGLGKALIEKLIRTLLQRDIGNISLFADSQVVEFYKNLGFEPDPEGIKGMFWYPMY

**NtSNAT2**

MQMQTLHLLSTSTSSSSSLSTFVSLNCCRCQFSNQLPFPCKTNLGFVKVKRQSKVSNLKAGFWESIRSGFVKNNTVQVIESPSNEEEEEEEPLPEEFVLIEKTQPDGTVEQIIFSSGGDVDVYDLQALCDKVGWPRRPLSKLAAALKNSYIVATLHSRKFSSGEEGNGEKKLIGMARATSDHAFNATIWDVLVDPSYQGQGLGKALIEKLIKTLLQRDIGNISLFADSQVVEFYKNLGFEPDPEGIKGMFWYPMY

**NtSNAT3**

MWTVPLGRTTCIIGANNSLNLFLNPGSINIPFHFPASDFKASPKSFSSLPRRSGLCRASQIAELFPTTSPEVFVREARVEDCWEVAETHCSSFFPEYAFPLDFVLRIDRLIAMLFGFSIPNGCKRTCLVAVVGSRDEEACLIGTEELKLGGFDGRLSLNKGYVTGILTVDTVADFLPRKGPLRQRRKGIAYISNVAVRERYRRKGIAKKLITKAEAQARSWGCRAIALHCDTSNPGAIKLYIGEGFRIIKVPEGANWPQPKTSPNMQFNLLMKLLDI

**NtSNAT4**

MQLPDEVLHQENRLEFGQFMAREAMFDEEYWTAAWLRAESHWEDRQNDRYINNYKKQYAEQEFNALKRRCKAQIGQRCTCIVTVRNEEKNNRHTVLKSVVGTLDLMIGHLSHGEDFPGERVNAQVFCNIERRSSNRYGYIANLCVAKSARRQGVARNMLHYAIRSAKANGAEKVFVHVHTNNGPAQKLYQKVGFEVVQVPNLKLSEEQPHLLLLAA

**NtSNAT5**

MTTIRRFSCNDLLRFASVNLDHLTETFNMSFYMTYMARWPDYFHVAEAPGGKIMGYIMGKVEGQGESWHGHVTAVTVAPEYRRQQLAKKLMNLLEDVSDKIDKAYFVDLFVRASNTPAIKMYEKLDYVIYRRVLRYYSGEEDGLDMRKALSRDVERKSIIPLKRPVTPDELEYD

**NtSNAT6**

MELRSKFLPQFKIQQPELTWVFSKQGKNKPLFVLNIFSREAFPVSYDRWKNIEVHCNNDQSIRQTPLSKQDNAKLPELSFNRLQQTDDGYCGLQKRNFGRFIAREAMLDEEYWTAAWLRAEAHWESVSYMRHVDAYKRKYAEQEFYALKRRCSGQDGNCLKCFCFVAVKKEEKNVRRTVLNSVVGTLDLTIRQFVQRERYPGEIKRLSAVLACQDPFDSHKYAYIANVCVAKFARRQGIASNMIHLAADAAALQGFKQLFVHVNADNIPGQELYKKTGFTIVEEASSSLSKEQRLLMSLEL

**NtSNAT7**

MAAAAPPPSPTPAPAVIREDLIPTAHQVFSRIRLATNADVPHIHKLIHQMAVFERLTHLFSATESSLSTTLFPENSPPPFTTFTVFLLEVSQNPFLPIDNQNCTNFSPIHKTINLDLPVSDAEAEMFKSGGNDAVVAGFVLFFPNYSSFLAKPGFYIEDIFVRECYRRKGFGRLLLSTVAAQAAKMGYGRVEWVVLDWNVNAIKFYEEMGAQILQEWRVCRLTGGALEAFANVNI

**NtSNAT8**

MAILITPFSYSPQASSLYLSSKLHNTNISSTYGYNSSTPLRSFVICSSQQLSQQNQQISPPTPQPILIDKSILSISEAKSENELWAASCLRVRTFYDFQHDTLNTEDHTKYLTEREFEALTERIAGKRVGFGRVSCVNATLPFSKVSNVAYDLSTSCKFSQDNVELVVVGTLDINQCIRLPDEITGMKPKGIGADFARGYVSNVCVAKEMQRNGLGCALISKAKMVAKDMGISDLYVHVAIDNEPAKKLYMKCGFVYENEEPAWQARFLDRPRRLLLWTDLSSS

**NtSNAT9**

MAILITPFSYSPQASSLYLSSKLHNTNIIYNTYGCRSSTPLRSFVLCSSQQLSQQISSPTPHPILIDKSFLCISEAKSENELWAASSLRVRIFYDFQHDTLNTEDHTKYLTEREFEALTERIAGKRVGFGRVSCINATLPFSEVSNVAYDLSTSCKFSQDNVELVVVGTLDINQCIRLPDEITGMKPKGIGADFARGYVSNVCVAKEMQRNGLGCALISKAKTVAKDMGISDLYVHVAIDNEPAKKLYMKCGFVYENEEPAWQARFLDRPRRLLLWTDLSNS

**NtSNAT10**

MTTIRRFSCNDLLRFASVNLDHLTETFNMSFYMTYMARWPDYFHVAEAPGGRIMGYIMGKVEGQGESWHGHVTAVTVAPEYRRQQLAKKLMNLLEDVSDKIDKAYFVDLFVRASNTPAIKMYEKLDYVIYRRVLRYYSGEEDGLDMRKALSRDIEKKSIIPLKRPVTPDELEYD

**NtSNAT11**

MELNSKFLPQFKIQQPQFTWVFSKQGKNKPLFVVNIFSREAFPVSYDRWKNIEVHCNNDQSIRQTPLSKQDNAKLPELSFNRLQQTDDGYCGLQKRNFGRFIAREAVLDEEYWTAAWLRAEAHWESVSYMRHVDAYKRKYAEQEFYALKRRCSGQDGNCLKCFCFVAVKKEEKNVRRTVLNSVVGTLDLTVRQFVQRERYPGEIKKLSTVLACQDPFDSHKYAYIANVCVAKFARRQGIASNMIHLAADAASLQGFKQLFVHVNADNIPGQELYKKTGFKIVEETSSSLSKEQRLLMSLEL

**NtSNAT12**

MAAAAPPPSPTPAPAVIREDLVPTGHQVFSRIRLATNADVPHIHKFIHQMAVFERLTHLFSATESSLSATLFPENSPPPFTTFTIFLLEVSQNPFLPIDNQNCTNFSPIHKTINLDLPISDPEAEMFKSGGNDEVVAGFVLFFPNYSSFLAKPGFYIEDIFVRECYRRKGFGRLLLSAVAAQAAKMGYGRVEWVVLDWNVNAIKFYEEMGAQILQEWRVCRLTGGALEAFANVNI

**NsSNAT1**

MQMQTLHLLSTSTSSSSSLSTFVSLNCCRCQFSNQLPFPCKTNLGFVKVKRQSKVSNLKAGFWESIRSGFVKNNTVQVIESPSNEEEEEEEPLPEEFVLIEKTQPDGTVEQIIFSSGGDVDVYDLQALCDKVGWPRRPLSKLAAALKNSYIVATLHSRKFSSGEEGNGEKKLIGMARATSDHAFNATIWDVLVDPSYQGQGLGKALIEKLIKTLLQRDIGNISLFADSQVVEFYKNLGFEPDPEGIKGMFWYPMY

**NsSNAT2** MLLYNPISTHLPPTPTALSLKPTTIHHRNVIVSSQYQPIPTTVNISISDESLKSKGFNLHRSITNLNLDH

LNSVFVAVGFPRRDTTKIQLALENTDSLMWIEYEKTKRPVAFARATGDGVFNAIIWDVVVDPNFQGIGLG

KAVMERLVTELLGKGINNIALYSEPRVLGFYRPLGFVADPDGIRGMVYSRKKKKNR

**NtoSNAT1**

MQMQTLYLLSTSPVTASSSLSNFVSLNCCRCQFSNPLPFPCKTNLDFVKVKRQSKVSNLKAGFWESIRSGFVKNNTIQVIESPSSEEEEEEEPLPEEFVLIEKTQPDGTVEQIIFSSGGDVDVYDLQALCDKVGWPRRPLSKLAAALKNSYIVATLHSRKFSSGEEGNGEKKLIGMARATSDHAFNATIWDVLVDPSYQGQGLGKALIEKLIRTLLQRDIGNISLFADSQVVEFYKNLGFEPDPEGIKGMFWYPMY

**NtoSNAT2**

MLLYNPISTHLPPTPLTLKPTTHHHHQNVTVSSQYQPIPTTVNISISDESLKSKGFNLHRSITNLNLDHLNSVFVAVGFPRRDTTKIQLALENTDSLAWIEYEKTKRPVAFARATGDGVFNAIIWDVVVDPNFQGIGLGKAVMERLVTELLEKGISNIALYSEPRVLGFYRPLGFVADPDGIRGMVYSRKKKKNR

**Table S4**. ***NtSNAT* and *NtSNAT-*like genes and their related Gene Ontology (GO) terms.**

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
|  | ID | Terms | Genes | Numberof genes |
| Molecular Function | GO:0008080GO:0016740 | *N*-acetyltransferase activitytransferase activity | *NtSNAT1, NtSNAT2, NtSNAT-*like*3, NtSNAT-*like*4, NtSNAT-*like*5, NtSNAT-*like*6, NtSNAT-*like*7, NtSNAT-*like*8, NtSNAT-*like*9, NtSNAT-*like*10,**NtSNAT-*like*11, NtSNAT-*like*12* | 12 |
| GO:0004596 | peptide alpha-*N*-acetyltransferase activity | *NtSNAT-*like*10* | 1 |
| GO:0016747 | Transferase activity, transferring acyl groups other than amino-acyl groups | *NtSNAT-*like*7, NtSNAT-*like*12* | 2 |
| Celluar Component | GO:0009507 | chloroplast | *NtSNAT-*like*8, NtSNAT-*like*9* | 2 |
| GO:0031416 | NatB complex | *NtSNAT-*like*10* | 1 |
| Biological Process | GO:0017196 | *N*-terminal peptidyl-methionine acetylation | *NtSNAT-*like*10* | 1 |



**Supplementary Figure 1. Conserved motif map**