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>gaattactatgacagatacttataaccattgtggtttttacgtattttaaatacgttatacctataat
tagtcacacgaggaaatcaaatgctaaagtatgatatgtttttatgttttgttttttcttgcttagGGG
ATCATGGA CATTGAAGCATATTTTGAAAGAATTGGCTATAAGAACTCTAGGAACAAATGGACTTGGAA
AACATTAAC TGACATTCTTGAGCACCAGATCCGGGCTGTTCCCTTTGAGAACCTTAACATGCATTGTG
GGCAAGCCATGGAGTTGGGCTTAGAGGCTATTTTTGATCACATTGTAAGAAGAAA CCGGGTGGGTGG
TGTCTCCAGGTCAATCAACTTCTGTACTGGGCTCTGACCACAATCGGTTTT CAGACCACAATGTTAGG
AGGGTATTTTTACATCCCTCCAGTTAA CAAATACAGCACTGGCATGGTTCACCTTCTCCTGCAGGTGA
CCATTGACGGCAGGAATTACATTGTCTGATGCTGGGTCTGGAAGCTCCTCCAGATGTGGCAGCCTCTA
GAATTAATTTCTGGGAAGGATCAGCC TCAGGTGCCTTGCATTTTCTGCTTGACAGAAGAGAGAGGAAT
CTGGTACCTGGACCAAATCAGGAGAGAGCAGTATATTACAAACAAAGAATTTCTTAATTTCTCATCTCC
TGCCAAAGAAGAAACACCAAAAAATATACTTATTTACGCTTGAACCTCGAACAATTGAAGATTTTGAG
TCTATGAATACATACCTGCAGACGTCTCCAACATCTTCATTTATAACCACATCATTTTGTTCCTTGCA
GACCCAGAAGGGGTTTACTGTTTGGTGGCTTCATCCTCACCTATAGAAAATTC AATTATAAAGACAA
TACAGATCTGGTCGAGTTTAAACTCTCACTGAGGAAGAGGTTGAAGAAGTGCTGAGAAATATATTTA
AGATTTCTTTGGGAGAAATCTCGTGCCCAAACCTGGTGATGGATCCCTTACTATT TAGAATAAGGAA
CAAATAAACCTTGTGTATGTATCACCCAACCTCACTAATTATCAACTTATGTGCTATCAGATATCCT
CTCTACCCTCACGTTATTTTGAAGAAAATCCTAAACATCAAATACTTTTCATCCATAAAAATGTCAGCA
TTTATTA AAAACAATAACTTTTTAAAGAACATAAGGACACATTTTCAAATTAATAAAAATAAAGGC
ATTTTAAGGATGGCCTGTGATTATCTTGGGAAGCAGAGTGATTCATGCTAGAAAACATTTAATATTGA
TTTATTGTTGAATTCATAGTAAATTTTTACTGGTAAAT

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Figure 1 Annotated sequence of *NAT2* gene. The coding region of the sequence is shown in capital letters while the 5' and 3' untranslated region are shown in small letters. The start codon (ATG) is highlighted in grey while the stop codon (TAG) is highlighted in purple. The area where the forward and reverse primers anneal to the sequence are highlighted in green. Reported SNPs are underlined and annotated with blue font, while novel SNPs are underlined and annotated with red font.

Table A Novel and reported SNPs of *NAT2*

Novel SNPs
c.85G>C, c.87G>A, c.99G>A, c.126G>C, c.136C>G, c.147G>C, c.150G>A, c.162T>C, c.175A>G, c.180A>G, c.189C>T, c.219A>T, c.253C>G, c.261A>G, c.267A>G, c.277T>G, c.284T>G, c.286C>A, c.293T>C, c.294T>C, c.297C>A, c.316G>A, c.374C>T, c.379A>C, c.384C>A, c.386C>A, c.405A>G, c.408A>G
Reported SNPs
c.42C>G, c.98G>A, c.138A>T, c.191G>A, c.198G>A, c.243A>T, c.246C>T, c.282C>T, c.341T>C, c.345C>T, c.354T>C, c.357C>T