

Supplementary Information Table 3

Supplementary Information Table 3A. Primers used on the Sequenom-MassArray platform

| SNP_ID | PCR primers | Extension primers |
|--------|-------------------------------------------------------------------|--------------------------|
| -80708 | ACGTTGGATGGTAGTTATGAGAGTGCTGCC ACGTTGGATGCCTTCCTGAGACCTAAGAAC | ATCTCTGTTGTGGAAGC |
| -68168 | ACGTTGGATGCTGCAACTAGCAAATCACAG ACGTTGGATGAAGGAACTGCGTTTATCAC | ATCACAGTTTGAGGAGG |
| -52070 | ACGTTGGATGAGAACAACCTCGCCCTTTTCC ACGTTGGATGGTACTTTGAAACCTGTCAGC | CAGAGATAGAAGCATTACATTT |
| -29144 | ACGTTGGATGGGCTAAAAGCAAGCCATGTG ACGTTGGATGGATCGCCCCAAGAAATCAAG | CACACAAGGGCCTGGAT |
| -23852 | ACGTTGGATGTACAGCACAAATGCCAGGCAC ACGTTGGATGGTCTCTGTAGATATAGAC | GCACAATGCCAGGCACATAGCAAA |
| -18304 | ACGTTGGATGTCAGGTTTCTCTGGTTAAGG ACGTTGGATGATCAGTTACTTGCCATGCTC | CCTGCAACTACATCTGTTA |
| -12534 | ACGTTGGATGATGGCACCTTATTTCTCCC ACGTTGGATGAACTAGGCAAGGAAGTGGAC | CTCCCACCACAGGGAACCTACA |
| -5289 | ACGTTGGATGTGCCAGCCATATGGGTGCAA ACGTTGGATGTTTGCTGCTTCCATCTTCCC | GCCATATGGGTGCAATGGCCT |
| -4570 | ACGTTGGATGGTGGTCCAAGGCCATTTTGTG ACGTTGGATGTTAAGAAGAGCTGTGTGCC | AAGGCCATTTTGTGCTGGCTGTAC |
| -1580 | ACGTTGGATGCTGGAGCCTGAAGAACCATC ACGTTGGATGTGTAAGAGCCTGTGCTCTG | GGCCTCGGGGAATTTTAC |
| -386 | ACGTTGGATGCCATCTCTCCAAGCTTCATG ACGTTGGATGCCTATTCGCCCATTTAATC | CTCATTTGTATAAAAAGGAAAAAG |
| -355 | ACGTTGGATGTATGAACACCCCCCTTGAG ACGTTGGATGACCTGGGTTTTGTTCTAGGC | ACCCCCCTTGAGGGTGTTAGGA |
| +177 | ACGTTGGATGAGAGTCCCGCCCCACCGA ACGTTGGATGAAAGGTGAAGAAGGGAAGCG | GGGGTGC GCGCCGCCTC |
| +3977 | ACGTTGGATGACCTGGGTAATCCAGGAATC ACGTTGGATGGTTACCTTCCGTGGCAAAAAG | GTTCTTAATCACAGCTGC |
| +6032 | ACGTTGGATGCAGATGTCTTCTCTGGTTCC ACGTTGGATGAACCATAGCAGCAAAGGAGG | CCTGTACCCATTCCAAC |
| +7685 | ACGTTGGATGGAATGAGAGCAGGGTAAGTG ACGTTGGATGGTCTTGACCTCATGGAGCAG | AGCCTTTGTGATTCTCTCTCTCT |
| R13W | ACGTTGGATGTCCACTCACCAGCCATCTTG ACGTTGGATGCTATGGCATCATCGACCTCC | CTAGAGGCTTCTTAGGCC |
| S55N | ACGTTGGATGAACTCCCCAGAAAGCACCTC ACGTTGGATGACGTGACTGAGCTGAGGCTT | GCCTACCTCACAGGACA |
| Q101Q | ACGTTGGATGAAGACTATGACGTCTGCGTG ACGTTGGATGCTGCCTTCCAAGTAGGAGAC | GGAAGACCTGGTGGCCGCCCA |
| S180L | ACGTTGGATGTCAGGTGGTAGGCAGCTCT ACGTTGGATGAAGTACCAGATGCTGCAGGC | CAGCTCTGCTGAGGCC |
| A186A | ACGTTGGATGCGACGTAGTACATGAATCGG ACGTTGGATGTGCACCATCCCCCTGCTGTC | ATCGGAGCTCAGGTGGGTA |
| V197I | ACGTTGGATGCACCTGAGCTCCGATTCATG ACGTTGGATGCGCATGACAGCTTCTTTGAC | AGCTCCGATTCATGTACTAC |
| +10122 | ACGTTGGATGTGGTCAGTGAGGAAACCAAG ACGTTGGATGGCTCCTGCACTTATTAACC | AGCGGGTGAAGTAACTTG |

| | | |
|--------|--------------------------------------------------------------------|------------------------|
| +10610 | ACGTTGGATGTGCAGACACTCAGTTGACAC ACGTTGGATGCTGCATGGGTTTTCTGTTTC | TCATGGGACCCCGGAAAT |
| +11347 | ACGTTGGATGATCCATGGGCTTGTGATCTC ACGTTGGATGAGGGAGCAAGTGAAGAGATG | GTGATCTCAGGTGAAGGC |
| +13466 | ACGTTGGATGGTGATTTCTGAGGTTTGCTTC ACGTTGGATGTTACCAACTCTGAGCTACTC | TGCTTCAAATGATCTTGTGG |
| +13955 | ACGTTGGATGCCTCCCCTACATCTGTTTTTC ACGTTGGATGAGAACAACAAAACCCAAAC | CATATTTAGGTCTTTAATCCAC |
| +14472 | ACGTTGGATGGAAAATATGAAATGCTAGGG ACGTTGGATGGTGTATTTTCAGTAAAGCTTTC | AAGAGGAGCTGACCAATGT |
| +21162 | ACGTTGGATGTGATTGGTTCACTGCTTCCC ACGTTGGATGGAGTCTGCGCGGACAAAAT | CTGGTACCTTCCCGTGTAG |
| +35464 | ACGTTGGATGCCAGGGATGGGTCAGATAAC ACGTTGGATGCACCAACCTCTCGTTCTTTC | TCAGATAACCGGACTTCGC |
| +44402 | ACGTTGGATGGAAAACACGTTGGGAAACAC ACGTTGGATGTGTAAGCGACTCGTGTTC | GGAAACACGATACACAG |
| +60339 | ACGTTGGATGTCAGGAACATCCTCCACCAG ACGTTGGATGGGAACACACTGGAATCTGAG | TTCACCAAACCACGTGGAA |
| +71446 | ACGTTGGATGAATGCCAACAGCTTCTGCC ACGTTGGATGAGTCTGCCTCCAAGTTGTTTC | CGTGCAAGGAATGATTTAG |

Supplementary Information Table 3B. Primers used for direct DNA sequencing

| Primer name | Sequence |
|---------------|--------------------------|
| TIRAP_5'_1F | CCATACTTGGCGCCGTC AAG |
| TIRAP_5'_1R | CAGCTCCGCTCTGCAAAACGC |
| TIRAP_5'_2F | GCGTTTTGCAGAGCGGAGCT G |
| TIRAP_5'_2R | GCTGCAGAGCCGCTGCAAAC |
| TIRAP_5'_3F | GTTTGCAGCGGCTCTGCAGC |
| TIRAP_5'_3R | CTCGGGCACAATCCGGCTGCG |
| TIRAP_ex1_F | CGCAGCCGGATTGTGCCCGAG |
| TIRAP_ex1_R | GATATCCACAACGCACTGAGC |
| TIRAP_ex2_F | GTGTGTCTAATAATACTTGC |
| TIRAP_ex2_R | CCTGCAGCCGTCAGCTCAG |
| TIRAP_ex3_F | CTGAGCTGACGGCTGCAGG |
| TIRAP_ex3_R | CACGACTGTGATTGGAGCTG |
| TIRAP_ex4_F | CAGCTCCAATCACAGTCGTG |
| TIRAP_ex4_R | ACAGCAGAGTCGCGAGTCCG |
| TIRAP_int4_1F | CGGACTCGCGACTCTGCTGT |
| TIRAP_int4_1R | GAGAAACACTGCTCAGGTCTGTGT |
| TIRAP_int4_2F | ACACAGACCTGAGCAGTGTTC |
| TIRAP_int4_2R | CTAGATTGTCGCTAAATG |
| TIRAP_int4_3F | CATTTAGCGACAATCTAG |
| TIRAP_int4_3R | GTTACCAGGTGCCCTCTCACTG |
| TIRAP_ex5_F | AGCACCTCCAGCGATGCTTC |
| TIRAP_ex5_R | TACTGTAGCTGAATCCCCTTCC |
| TIRAP_ex6_F | GGAACGGGATTCAGCTACAGTA |
| TIRAP_ex6_R | TCACTCCCTATACCTCTTGTAC |
| TIRAP_ex7_F | GTGACAAGAGGTATAGGGAGTGA |
| TIRAP_ex7_R | GTGTCTTTAATGGCACCTAG |
| TIRAP_ex8_F | TCATTGACTGACTAGTCCAT |
| TIRAP_ex8_R | GCTTTACTGAAATACAC |