**Table S4. Primers and melting temperature (Tm) used for Q-MeDIP and pyrosequencing.**

|  |  |  |  |
| --- | --- | --- | --- |
| **Name** | **sequence 5'-3'** | **TM** | **Length** |
| **QMeDIP** |
| ZNF336\_qMeDIP.Fow | GGCCAAGCTCTCCCTTGCTC | 60 | 122 |
| ZNF336\_qMeDIP.Rev | CGGTGTTCCAGACAGGCAGG |   |   |
| FYN\_qMeDIP.Fow | TGGACCGTCTGTAGGCAGGTGT | 60 | 188 |
| FYN\_qMeDIP.Rev | GGCAGTGGCTGGGGAGATGG |   |   |
| TMCO7\_qMeDIP.Fow | TCCAGGGTGTGCTGGTGGAGA | 60 | 176 |
| TMCO7\_qMeDIP.Rev | ACTCCTCTGCGTTTCACACAAGTT |   |   |
| ANKRD50\_qMeDIP.Fow | TGCCAAAGGTGAAGTACGTGGA | 60 | 171 |
| ANKRD50\_qMeDIP.Rev | TCTTGGTTTTAAGACGTCCGGGT |   |   |
| ZNF808\_qMeDIP.Fow | ACGTTCAGTCAGAAGGCAACCCT | 60 | 180 |
| ZNF808\_qMeDIP.Rev | ACGCACGAAAGCCTTGTCACA |   |   |
| TGIF1\_qMeDIP.Fow | TTGCTTGGCCTTGAGGCAGC | 60 | 120 |
| TGIF1\_qMeDIP.Rev | CGTTGCTTGGTGCTGGCGGA |   |   |
| ACOT2\_qMeDIP.Fow | CTATTAGCCGGGCGCGGTGG | 60 | 187 |
| ACOT2\_qMeDIP.Rev | CCCACATCGAAGGGTGCTGGC |   |   |
| ITGB6\_qMeDIP.Fow | TCACGTACAAGGTAAGACCTGCGT | 60 | 184 |
| ITGB6\_qMeDIP.Rev | TCACTGTGGGAACTGCCCGGT |   |   |
| RNF219\_qMeDIP.Fow | TCACCCCATCCAATCCCCGAA | 60 | 163 |
| RNF219\_qMeDIP.Rev | GGGAGCTGCAAGATTCCTGCGA |   |   |
| GRM5\_qMeDIP.Fow | TGGTACATGGGGACAAAATGGTCCT | 60 | 131 |
| GRM5\_qMeDIP.Rev | GCATGGTGGGGAAAATTCAGGAGGG |   |   |
| OR1J4\_qMeDIP.Fow | ACATGAAGGCTCATGACTAGCCTGA | 60 | 171 |
| OR1J4\_qMeDIP.Rev | TCGCCAAGAGAGTAGAGGGTTTGA |   |   |
| AVPR1A\_qMeDIP.Fow | ATGGCATCCAGCACAACCAG | 60 | 237 |
| AVPR1A\_qMeDIP.Rev | TGTTGGTGGATACAGTTGGAAGA |   |   |
| IL1R2\_qMeDIP.Fow | TGGGGCCACAGTTGGGCAAAA | 60 | 230 |
| IL1R2\_qMeDIP.Rev | GGGTGGTGTCACCCGCTCAC |   |   |
| IL1RN\_qMeDIP.Fow | GCTTCTCGCAGTGGGGCAGG | 60 | 201 |
| IL1RN\_qMeDIP.Rev | CCCAGAGGGTCGGCAGATCGT |   |   |
| IL31\_qMeDIP.Fow | GAAGCCTTCGACCCGCCACA | 60 | 214 |
| IL31\_qMeDIP.Rev | AGCCCAGGGTTGACAGACACGA |   |   |
| IL33\_qMeDIP.Fow | ACTATGCATGATTGTATGTGCT | 60 | 204 |
| IL33\_qMeDIP.Rev | GGCTTTGGTCAATGGATTGC |   |   |
| JUNB\_qMeDIP.Fow | TGGGTCATGCAGCTCCACCCA | 60 | 268 |
| JUNB\_qMeDIP.Rev | CGCAGCAAGTGGGGTGAGGG |   |   |
| OR13C8\_qMeDIP.Fow | TCACAGAGTTCTGTTGGAGATCAGT | 60 | 152 |
| OR13C8\_qMeDIP.Rev | GCAGGATCATCAGGTACATCCACA |   |   |
| PCDHB8\_qMeDIP.Fow | TGCCCAAAACTCAGTTAGCATTGT | 60 | 131 |
| PCDHB8\_qMeDIP.Rev | TGCCTGTATGTACCCTGGGACT |   |   |
| SLC41A2\_qMeDIP.Fow | TGCGGTTCTGATTCATCTGCTGT | 60 | 247 |
| SLC41A2\_qMeDIP.Rev | TGCCTGTCTTCTTGGTTTTTCTGGT |   |   |
| **Pyrosequencing** |
| ITGB6\_bis\_out.Fow | GGATTGAATTGTTTTGTTTGTTTTT | 55 | 286 |
| ITGB6\_bis\_out.Rev | TCCTCATTAAAATACCTATACAACCTTTAT |   |   |
| ITGB6\_bis\_nest.Fow | TTGTTTGTTTTTTTTATTTTTAGGAAG | 60 | 224 |
| ITGB6\_bis\_nest.Rev | /5Biosg/AAAACATTCACTATAAAAACTACCC |   |   |
| ITGB6\_bis\_seq1.Fow | AGGAAGGAATGATTA |   |   |
| ITGB6\_bis\_seq2.Fow | ATTATAGATAAATAT |   |   |
| GRM5\_bis\_out.Fow | ATTATTTAATTAAATGGTATATGGGGATAA | 55 | 204 |
| GRM5\_bis\_out.Rev | AAACTAACAAAATCAACAAAAAAACC |   |   |
| GRM5\_bis\_nest.Fow | TATGGGGATAAAATGGTTTTTTAGA | 60 | 150 |
| GRM5\_bis\_nest.Rev | /5Biosg/AAAAAATTCAAACCAATAAAAATAACATAA |   |   |
| GRM5\_bis\_seq1.Fow | TAATTTTTTGATTTG |   |   |
| OR13C8\_bis\_nest.Fow | GTTGGAGATTAGTATATGTAATATG | 55 | 163 |
| OR13C8\_bis\_nest.Rev | /5Biosg/CTAAAATAAAAACTCCATTTCC |   |   |
| OR13C8\_bis\_out.Fow | TTTTGTTGGAGATTAGTATATGT | 60 | 300 |
| OR13C8\_bis\_out.Rev | ACCTTTTTCTTTACTACCAA |   |   |
| OR13C8\_bis\_seq1.Fow | TATGGAAAGGATTAA |   |   |
| OR13C8\_bis\_seq2.Fow | TTAGATAGTTTTTTT |   |   |
| IL31\_bis\_nest.Fow | ATTTGGAGGGTAATTTTAGT | 55 | 220 |
| IL31\_bis\_nest.Rev | /5Biosg/AAAAAAAACAAAACCAACTTC |   |   |
| IL31\_bis\_out.Fow | GTTTATTTGGAGGGTAATTTTAG | 60 | 298 |
| IL31\_bis\_out.Rev | CCTAATAAAATAAAATAATACC |   |   |
| IL31\_bis\_seq1.Fow | GTTTTTTTATTTAGT |   |   |
| IL31\_bis\_seq2.Fow | TGGTTTTGTTATGGG |   |   |