**Table S1. Primers used for obtaining templates of sgRNA with T7 promoter.**

|  |  |
| --- | --- |
| Name | Sequence (5’ to 3’) |
| T7-Sg1# | TAATACGACTCACTATAGGGGGTGAGCGTGAGCAGCGTGC |
| T7-Sg2# | TAATACGACTCACTATAGGGCTGCCTCATCATCCTCTTCT |
| T7-Down | AAAAAAAGCACCGACTCGGTGCCA |

**Table S2. 16 potential off-target sites examined by PCR and primers used for list.**

|  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
|  | **20** | **19** | **18** | **17** | **16** | **15** | **14** | **13** | **12** | **11** | **10** | **9** | **8** | **7** | **6** | **5** | **4** | **3** | **2** | **1** | **N** | **G** | **G** |
| **MC1R sg1#** | **G**  | **G**  | **T**  | **G**  | **A**  | **G**  | **C**  | **G**  | **T**  | **G**  | **A**  | **G** | **C**  | **A**  | **G** | **C**  | **G** | **T**  | **G**  | **C**  | **T**  | **G**  | **G**  |
| **OTS 1** | **G** | **G** | **T** | **G** | **A** | **G** | **C** | **C** | **T** | **G** | **T** | **G** | **C** | **A** | **G** | **C** | **G** | **T** | **G** | **C** | **T** | **G** | **G** |
| **OTS 2** | **C** | **G** | **T** | **G** | **G** | **A** | **C** | **G** | **T** | **G** | **A** | **G** | **C** | **A** | **G** | **C** | **G** | **T** | **G** | **C** | **G** | **G** | **G** |
| **OTS 3** | **G**  | **G**  | **A**  | **G**  | **G**  | **G**  | **C**  | **G**  | **T**  | **G**  | **A**  | **G**  | **C**  | **A**  | **G**  | **C**  | **G**  | **T**  | **G**  | **G**  | **G**  | **A**  | **G**  |
| **OTS 4** | **G**  | **G**  | **T**  | **G**  | **A**  | **G**  | **C**  | **T**  | **G**  | **C**  | **A**  | **G**  | **C**  | **A**  | **G**  | **C**  | **G**  | **T**  | **G**  | **C**  | **A**  | **A**  | **G**  |
| **OTS 5** | **G**  | **G**  | **C**  | **A**  | **A**  | **G**  | **C**  | **G**  | **T**  | **G**  | **A**  | **G**  | **G**  | **A**  | **G**  | **C**  | **G**  | **T**  | **G**  | **C**  | **T**  | **G**  | **G**  |
| **OTS 6** | **G**  | **G**  | **G**  | **G**  | **A**  | **G**  | **A**  | **G**  | **G**  | **G**  | **A**  | **G**  | **C**  | **A**  | **G**  | **C**  | **G**  | **T**  | **G**  | **C**  | **A**  | **G**  | **G**  |
| **OTS 7** | **G**  | **G**  | **T**  | **G**  | **A**  | **G**  | **C**  | **T**  | **T**  | **C**  | **A**  | **G**  | **C**  | **A**  | **G**  | **C**  | **C**  | **T**  | **G**  | **C**  | **G**  | **G**  | **G**  |
| **OTS 8** | **G**  | **G**  | **C**  | **C**  | **A**  | **G**  | **G**  | **C**  | **T**  | **G**  | **A**  | **G**  | **C**  | **A**  | **G**  | **C**  | **G**  | **T**  | **G**  | **C**  | **C**  | **A**  | **G** |

|  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- |
|  | Site | Potential Off Target Site | Position | No. of Mismatches | Gene | PCR Primer (5’ to 3’) | Product Size(bp) |
| Sg1# | OT1 | GGTGAGCCTGTGCAGCGTGCTGG | Chr2:+95042062 | 2 | - | F: TTTGCAACTCACTGCCCTCTR: CCTTTGTGTCTTTGCCTGGTG | 707 |
| OT2 | CGTGGACGTGAGCAGCGTGCGGG | Chr12:+33652570 | 3 | TMEM151B | F: ACTGCTACTTCCGCCTCTTTCR: CGGGACACCAGAGTAAGGAG | 772 |
| OT3 | GGAGGGCGTGAGCAGCGTGGGAG | chrUN0:+53260732 | 3 | STK10 | F: GACAGAGGGAGGTGAAAGGCR: GGGTCCTAGCTGTCATGTCG | 715 |
| OT4 | GGTGAGCTGCAGCAGCGTGCAAG | chrUN0:-951264 | 3 | - | F: CCTTTCGGTTCTCTGTGCTCTR: CTGTCCATGGCGCTTGTATG | 531 |
| OT5 | GGCAAGCGTGAGGAGCGTGCTGG | chr1:+103377309 | 3 | - | F: GCACCCTTTCCCCGTAACTR: GTGGTCCCTGTTGTGGGTTC | 511 |
| OT6 | GGGGAGAGGGAGCAGCGTGCAGG | chr14:+39467157 | 3 | - | F: TCGCTTTAGTTTAGGTGGAAGTGTR:CAGCACTGCAGGTTAGAGGTTTAG | 607 |
| OT7 | GGTGAGCTTCAGCAGCCTGCGGG | chrUN0:+143328002 | 3 | GPR153 | F: TCTTCCTTCCCACAGGCAGTR: TCTTGCTCTTGCTGGACCTTC | 738 |
| OT8 | GGCCAGGCTGAGCAGCGTGCCAG | chr17:-14779856 | 4 | - | F: TTGAGAAACTGCCAAGCTGAR: GCACAATACTCACCCCTAACC | 636 |

|  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
|  | **20** | **19** | **18** | **17** | **16** | **15** | **14** | **13** | **12** | **11** | **10** | **9** | **8** | **7** | **6** | **5** | **4** | **3** | **2** | **1** | **N** | **G** | **G** |
| **MC1R sg2#** | **C**  | **T**  | **G**  | **C**  | **C**  | **T**  | **C**  | **A**  | **T**  | **C**  | **A**  | **T**  | **C**  | **C**  | **T**  | **C**  | **T**  | **T**  | **C**  | **T**  | **T**  | **G**  | **G**  |
| **OTS 1** | **C**  | **T**  | **T**  | **C**  | **C**  | **T**  | **C**  | **A**  | **T**  | **C**  | **A**  | **T**  | **C**  | **C**  | **T**  | **C**  | **T**  | **T**  | **C**  | **T**  | **A**  | **A**  | **G**  |
| **OTS 2** | **C**  | **T**  | **G**  | **G**  | **C**  | **T**  | **C**  | **A**  | **T**  | **C**  | **T**  | **T**  | **C**  | **C**  | **T**  | **C** | **T** | **T** | **C** | **T**  | **T**  | **G**  | **G**  |
| **OTS 3** | **C**  | **T**  | **G**  | **T**  | **C**  | **T**  | **C**  | **A**  | **C**  | **C**  | **A**  | **T**  | **C**  | **C**  | **T**  | **C**  | **T**  | **T**  | **C**  | **T**  | **A**  | **G**  | **G**  |
| **OTS 4** | **C**  | **T**  | **G**  | **C**  | **C**  | **T**  | **C**  | **A**  | **T**  | **T**  | **C**  | **T**  | **C**  | **C**  | **T**  | **C**  | **T**  | **T**  | **C**  | **T**  | **G**  | **G**  | **G**  |
| **OTS 5** | **T** | **T** | **C**  | **T**  | **C**  | **T**  | **C**  | **A**  | **T**  | **C**  | **A**  | **T**  | **C**  | **C**  | **T**  | **C**  | **T**  | **T**  | **C**  | **T**  | **A**  | **G**  | **G**  |
| **OTS 6** | **C**  | **A**  | **T**  | **C**  | **A**  | **T**  | **C**  | **A**  | **T**  | **C**  | **A**  | **T**  | **C**  | **C**  | **T**  | **C**  | **T**  | **T**  | **C**  | **T**  | **G**  | **A**  | **G**  |
| **OTS 7** | **C**  | **T**  | **G**  | **C**  | **C**  | **T**  | **C**  | **A**  | **T**  | **C**  | **T**  | **T**  | **C**  | **C**  | **T**  | **C**  | **T**  | **T**  | **C**  | **C**  | **G**  | **A**  | **G**  |
| **OTS 8** | **C** | **T**  | **G**  | **C**  | **T**  | **T**  | **T**  | **C**  | **T**  | **C**  | **A**  | **T**  | **C**  | **C**  | **T**  | **C**  | **T**  | **T**  | **C**  | **T**  | **C**  | **A**  | **G** |

|  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- |
|  | Site | Potential Off Target Site | Position | No. of Mismatches | Gene | PCR Primer (5’ to 3’) | Product Size(bp) |
| Sg2# | OT1 | CTTCCTCATCATCCTCTTCTAAG | chr7:+142130706  | 1 | TRAK2 | F: TTTGCAACTCACTGCCCTCTR: CCTTTGTGTCTTTGCCTGGTG | 500 |
| OT2 | CTGGCTCATCTTCCTCTTCTTGG | chr3:-82297932  | 2 | RRS1 | F: ACTGCTACTTCCGCCTCTTTCR: CGGGACACCAGAGTAAGGAG | 486 |
| OT3 | CTGTCTCACCATCCTCTTCTAGG | chrUN0:-1694385  | 2 | - | F: GACAGAGGGAGGTGAAAGGCR: GGGTCCTAGCTGTCATGTCG | 605 |
| OT4 | CTGCCTCAGTCTCCTCTTCTGGG | chrUN0:+1712663  | 3 | - | F: CCTTTCGGTTCTCTGTGCTCTR: CTGTCCATGGCGCTTGTATG | 482 |
| OT5 | TTCTCTCATCATCCTCTTCTAGG | chr1:+72469871  | 3 | - | F: GCACCCTTTCCCCGTAACTR: GTGGTCCCTGTTGTGGGTTC | 527 |
| OT6 | CACCATCATCATCCTCTTCTGAG | chr14:-157875265  | 3 | - | F: TCGCTTTAGTTTAGGTGGAAGTGTR: CAGCACTGCAGGTTAGAGGTTTAG | 627 |
| OT7 | CTGCCTCATCTTCCTCTTCCGAG | chrUN0:-549895  | 2 | - | F: TCTTCCTTCCCACAGGCAGTR: TCTTGCTCTTGCTGGACCTTC | 605 |
| OT8 | CTGCTTTCTCATCCTCTTCTCAG | chr17:-23331789  | 3 | - | F: TTGAGAAACTGCCAAGCTGAR: GCACAATACTCACCCCTAACC | 613 |

**Table S3. Information regarding the specific primers used for the qPCR.**

|  |  |  |  |
| --- | --- | --- | --- |
| Gene | Primer Sequences (5’ to 3’) | GenBank accession number | Product size (bp) |
| MITF | F: CGTGTATGCAGATGGATGATGR: CAGGTAACGTATTTGCCATTTG | XM\_002713310.3 | 118 |
| TYR | F: CATCTTCGATCTGAGTGTCTCR: TGCCTACAGGGATGACATAG | NM\_001082077.1 | 98 |
| TYRP1 | F: CAGAGGATTCTTACAGTCAGGAGR: CTGGTGGCAATGACAAACTG | NM\_001297495.1 | 122 |
| DCT | F: TGTGCATGACCTTCGATAGCR: CCTGGTGTCAGTTTGTACCTC | NM\_001297493.1 | 125 |