

Cell line	Sequence of PCR product	Integration site according to BLAST alignment
A549 SIB01	CTCTACAAAATGCTTGTTATATGATGTAAGTCTCAAAA TGCTACCTGATTCTAAAGAATGCATTTGCTCATAAAACA ATGAGAGGTGGCAGGTGCATTGTTCTCTTGGATT CCAACAAACATTTAATATGCTTTTCTGAATTGTTACTTTC CTTGTCTATCTGT	Homo sapiens chromosome 12. NT_009755.19. Position 7510752 to 7510922. Features in this sequence: transmembrane protein 132D; TMEM132D. Intron 2.
AGS SIB02	TTGATGAGTCTGAGTAAAGGAGAAAAGAGCCCACTG CAGATTTTCAAGCCAATTCCATCCCTTCAAGCCCACCCC CACCTTTGCACACACGCAAACACACACGTGCACACACA CACACACATCACTGCTGCCACCACCACCCTGCCACCAC CACCATTTTCTCCTCTCATATAAAATAACTGACAATGTT CCCCTGAATATAAACACATGGACCCACTAAAGCTACAT GCAAATTGACAGAAGATTCCAATCCTGCATCATGCCAG AGAGCTCAAGGAAGAAAGTCCTATCCTGCGGTCTCAGT TGGCAGGAGCCTCTGTCCTACAATTTCTGACCCGGTCA AAGCAACCTTATAACCTCACGGGCAAAGACATGAGATG AATCAAGAGGGTACTCAAGACTCATC	Homo sapiens chromosome 2. NT_022184.15. Position 39155292 to 39155675. No features in this sequence. Features flanking this sequence: 856359 bp at 5' side: similar to hCG1815165. 345912 bp at 3' side: B-cell CLL/lymphoma 11A isoform 3.
AGS SIB02	ACCTATCTGAAAAAAAtAAACAATAATTaaaTAAATAAAT AAAGCCTTGGGAAAGCCCGTTGGTAAAAGACACATGTC TGATTTTATCTTTTACGAGCCCCCTTACCCTGTGAGTTTA TCAGCCTTACACGGCTAATGCACGGGGGGGGAATAGTT TCTTATTATGGTGTCTTTTTCCCCTAAGACTGAAAAAA CC	Homo sapiens chromosome 2. NT_022184.15. Position 21913858-21914014. No features at this site. Features flanking this sequence: 72295 bp at 5' side: 3-hydroxyanthranilate 3,4-dioxygenase. 359331 bp at 3' side: zinc finger protein 36, C3H type-like 2
L929 SIB01	ACATCACGTTGGGTAACATTGCTTTTCAAAAACATTTTG AGACTATACGTAATGATCtTATTgATTTTGTATATATGA GTACACCACCGTTGCTGaTCTTCAGACAGACTA	Mus musculus chromosome 17. NT_039649.7. Position 52357582-52357685. No features in this sequence. Features flanking this part of sequence: 41915 bp at 5' side: Rab31-like. 9702 bp at 3' side: protein phosphatase 4, regulatory subunit 1.
L929 SIB01	TGTGCGGTGGGGTGGGAAGTCCCTCTACTCGTATCAGA CACCTGGGAGATGATACAGTTTCTGGACTGGACAAGAG TGACTTTAAGTGAAATGCTCATTCTGGGGAGAGGTTG TTCTGTTGTTCTTTTACACGAGGTCAATTCTGGGTTAA AAATTTGACTGTGGGTTAATAACACCATCCCTCCTCTT AACGCCCTGTCTACCTCCTGGAAGTTAACTATGAGTT ACCTCTCCCCACTGTTGAGCATTTCACCTAAAGTCACTC CTGCTCAGTCTGAAACTCTCTCATCTCCAGGTCTCTG CTACTTTCTAGAGGGACTTCCCACCCACCCACACTTTCT GAGCCTGGGAGCTCTCTG	Mus musculus chromosome 19. Position 43272966-43273076. No features in this sequence. Features flanking this part of sequence: 55846 bp at 5' side: similar to rat ribosomal protein L13a. 64525 bp at 3' side: SORCS receptor 1.
L929 SIB01	GGCcAGGGGGCCTCTCCCCTCACCaCCTCATTGACCTC CCCAGTCCTTGGAACTAAcTtTTCTCATTGGCTTCcCT AGGATAC	Mus musculus chromosome 19. NT_039687.7. Position 38005522- 38005436. Feature in this sequence: Paired box gene 2. Intron 8. Features flanking this part of subject sequence: 119617 bp at 5' side: olfactory receptor 1505. 363523 bp at 3' side: hypothetical protein.