gene abbreviation	amplified sequence	length		efficiency %	R^2
Ano1 (TMEM16A)	ACAACTCCACCATGAGAACCTTCTCCTTGTGGATCTGCTGGCTGATGTCTTTGGGGATATCAGGGATCACCCAGTCCACAAAGTCACTCA TGAACATCACCAGGTTCTGGAAGACAATGACGAATGCCAGCCGGGCGGCCAGGA	114	intron_spanning	100	0,9986
Ano2	GGCAAGTTCTCAGTCATCATCAATGCTTTTGTCATTGCTGTCACCTCCGACTTTATCCCCCGCCTTGTATACCAGTACTCCTATAGCCACA ACGGGACGCTACATGGATTCGTGAACCA	119	intron_spanning	99	0.9996
Ano3	CAGGAATCAGCATTCCAGTATACCATCCCAGCCAGGCAAAGTATAAGCCAATCTTCTCACCAAAGTACAGCCTGATTAAGTCCAGAGGTTG ATGCTTGTACCACATTCC	109	intron_spanning	102	0.9998
Ano4	CCTTCGAGAATTCCATACCAAATTCCTATGTCCTTGGCCCTCGAAGCCAGAGGCCTTCTCCACTGAGTCACAAACTTGTAAGCATCAAGT CGGATTTCAATGATGTTATTCAGTAAGGCC	120	intron_spanning	99	0.9997
Ano6	AAGTGATGAACAACGTTAACAAATTTGGGATTAATAGACTGGTCAGTTCTGGCATCTACAAAGCAGCGTTCCCTCTGCACGACTGCAAATTC AACTATAAGTCGGAAGACC	111	intron_spanning	94	0.9993
Ano7	GCTGTGGTGATTATGTGTCTTGTATCTATCATCCTGTACCGGGCCGTGATGGCTATCATCGTGTCCAAGTCGAACAACGCCTTC	84	exonic	91	0.9916
Ano8	CGAGGACCTGTTATGTCTCTTCTGTCTCCCGGCCGCATCGCCAGAGCCAGTCTCCGGATCTCGGTCTCCATCAGATC	77	exonic	103	0.9976
Ano9	AGTCTCTTCTTGCTGTTCACGACAAAGTTCACAATTCGGATTCTTGTGGTAGCTATGTTGAAGTTCTGCTTGGCTCTGGACCTAGGATCTT CAGGCTTCATGAGGA	106	intron_spanning	96	0.9994
Ano10	GGAACAACTCCAAGTAATCATCAAAGGTCCCCAGGTAAGTGCCCATTTCCTTCTCCAGGAGGACTTGCTCATACAAGGTCGTGTCAACAT CCGACTTCAATGC	103	intron_spanning	101	0.9985
Best1	TGCTGATGATTCGAGAGAACCATTCTGTAGAGTCCACGGATGCAATAGTAGAGGAATATGAAGACAAGGAATTCTCCG	78	intron_spanning	96	0.9993
Clcn1	TCCATGTTCTCTGGTGTCTATGAGCAGCCATACTATTACACTGACATCCTGACAGTGGGCTGTGCCGTGGGGGTCGGTTGCTGTTTTGG AACACCACTTGGAG	103	intron_spanning	98	0.9996
Clcn2	CAGGACGGGTGCGGGCGCGGGCTCAGCGAGCCGGGAGCTGAGTCAAGGCCAGAGCCCGAGAAGCAAGAGGAGGCAAGAGGACAGTG CACCGAGATG	66	exonic	99	0.9997
Clcn3	GACTTCCGAAGACGACAATTTGTTAGACGGTGACACAGCAGCTGGAACTCATTATACAATGACAAATGGAGGCAGCATTAACAGCTCCACA CACT	95	intron_spanning	100	0.9990
Clcn4	TGGATCATCTTCAGGAACTGGTGCAGGCCTGGAGCGGACTGTGTCACACCAGGGCTTTATGCGATGGTGGGAGCTGCAGCCTGTCTAG GTGGAGTGACTAGGATGACAGTGTCTCTGG	118	intron_spanning	96	0.9994
Clcn5	CTTCTGTAAAAGCGCCTCAACCTTAATGAGCAAATGCGGGCTGCCAAGGTTGGTTCTTCCTTGAGGCAGAAAAAAAGAAAACTGTGAACGA AGGGAACAAGTCATACAATGGTGGTGGAATAGGCTCTTCAAATAGA	107	intron_spanning	101	0.9993
Clcn6	CATGTCGTTGTAGGTGTCATTGGGACAGAAAAATGTCTTGATAGTTGAATTCACATCTTCCGATGTGACCTGGAGCTGGAATGAGCCATTA CCCGTTTGACTCGTGGAAGACATCT	116	intron_spanning	102	0.9992
Clcn7	CCTAAGTGACACAGCGTCTAATCACAATGGGTTCCCTGTGGTGGAGGATGTAGGAGACACCCAGCCAGCCAGACTCCAAGGCCTAATCC TGCGTTCCCAGCTCATCGTGCTCCTGAAGCACAAGGTGTTTGTGGAGAGGTC	111	intron-spanning	97	0.9974
Clcnka	TCGGAATCTGGTCTTGTAGATGGAGGTGATGGTCTCCTGTTCACTGTTGAAGACCCCTAGGAGACGGAACATGAAGG	77	intron_spanning	99	0.9987
Clcnkb	CTCCCTGTACAGCCACTTATGTGCTCTGACCACACGTCCAATAGCAAAGTTCATGGCATAGCTGATCAGAGCCATGAGCACCCCGAGAG CCACCAGGAAGTACCAGTCCTCACCCACACGGAACAGCCGCTCTTTCAGCCACTCTAGGCTCCCCTGGATGTTTCTGCGGATT	172	intron_spanning	97	0.9994
Cftr	GATGTTGGCTATTACTGGATCTACTGGAGCAGGAAAGACATCACTCCTGATGCTGATACTGGGAGAACTGGAAGCGTCAGAGGGAATTATT AAGCACAGTGGAAG	105	intron_spanning	99	0.9998
Gadd45a	CTTAAGGCAGGATCCTTCCATTGTGATGAATGTGGGTTCGTCACCAGCACACAGTGTAAGTCCGGGGTCTGCGCAGCGCCCCCGCTCT CAGCGGGGCTCTTGTCGTTCTCCAGTAGCAACAGCTCTGCCAGCCGACC	107	intron_spanning	100	0.9998
Lrrc8a	AAGACCATCGAGGAGATCATCAGCTTCCAGCATCTGCACCGCCTCACCTGCCTTAAGCTGTGGTACAACCAC	72	exonic	97	0.9997
Lrrc8c	GAACTCGGTCACTGGAATCATGTTTCTTATATGTATATGTTACTTCTCTGAACATCTGTGGTCCCAGAATTCTTAGTCATAAGCAGCCAC	90	intron_spanning	91	0.9993
Lrrc8b	CCAAGAGTAGACACTGAAGTTTCTTGCACTGGAACAGCCCGTCTGGTAGCATCTCGATGTTATTGTTGGTCACGGCGAAGTACTGCAGGT TGGTCAGGTACTGG	104	intron_spanning	101	0.9996
Lrrc8d	TGCAGTGTAGCAGAGGATAAAAATGAACTTGGCGGTCTTGATGAGGGTCTGGACCACGTAGAGCTTGTAGATCAAGTCACTGTCTTCCAC	90	exonic	96	0.9997
Lrrc8e	CTGATGAGTTCTCACGAGATTCATGGCTGGGCAGGCAGATGATCTTGTCCTGTGTCACCTGGAGGGTGCACCCGAACACCCCAATCATG AGCATAGCCACAG	102	intron_spanning	97	0.9996
Sema3g	AGACTCAACAACCTCTCTCTCCCACCCGGCTAGGAAAAGGAGTCCAGGACGCCTGATTACCTCTTAGAGAAAATCTCTACCACCTACAGG AA	92	exonic	99	0.9995
Slc12a1	TGGATGGGTGAAAGGTGTGCTGGTGAGGTGCATGCTGAACATCTGGGGAGTCATGCTCTTCATTCGCCTCTCCTGGATTGTAGGAGAAG CGGGAATTGGTCTTGGAGaTCCTTATAATTCTTCTTTCCACCATGGTAACCTCTATCAC	117	intron_spanning	100	0.9999
Slc12a2	TGTAAGATCCGAGTATTCATTGGTGGAAAGATAAACAGAATAGACCATGACCGGAGAGCGATGGCTACTTTACTCAGTAAATTCCGAATAGA CTTCTCCGATATCATGGT	110	intron_spanning	94	0.9984
Slc12a4	GCTTGACATGTCGTGTGTGCCCGATGGGTAAAAGATGGCAGCTGGTGGAGCAATGTAGGTCAGCAAGATCTCAATGGCTCCTAGGAT	87	intron_spanning	99	0.9999
Slc12a5	CCGAGCCCGGCGCCGCGCAGCCACCATGCTCAACAACCTGACGGACTGCGAGGACGGCGATGGGGGAGCCAACCCGGGTGACGGC AATCCCAAGGAGAGCAGCCCCTTCATCAACAGCACGGACACGGAGAAGGGGAGAGAGTATGATGGCAGGAACATGG	113	intron_spanning	96	0.9998
Slc12a6	CATGCTATTACCATTCCTACTGAAACAGCACAAGGTTTGGCGAAAATGCAGCATACGGATCTTCACAGTAGCCCAACTAGAAGACAACAGT ATCCAGATGA	101	intron_spanning	96	0.9999
Slc12a7	GTGTTCCTGTCGTGAATCAGCTGGGCCTCTCTCTCCCGCTCATTCTTTGACAACTGCATCTGTTTCAGCATCTGTGACCTCTGCTCCAT CATTAGCGTCTTCTCATAGGTGA	112	intron_spanning	99	0.9992
Slc26a9	TTCGTTACCTTCCACACTCTCATCCTTGACATGAGTGGAGTCAGCTTTGTGGACCTGATGGGCATCAAAGCTCTAGCCAAGCTAAGCTCC ACCTATGAGAAGA	103	intron_spanning	95	0.9994
Tmem206	GAGCTGGTCCAAGTTCAAGGTCCAGGTGTTGTGCCAGGTGTGGACAATGAGTCTGCATCCAG	62	intron_spanning	99	0.9996
Ttyh1	CCTTGACCTCACAGAGAACAGCCTGCCTCTGCCTCCCTAGTACTGAGATTAAAGCATGTATCACCATATCACCAGC	76	exonic	101	0.9990
Ttyh2	GAGAACGTGCCACTCATCGGGAGAGGTTCCCCTCCGCCCACGTACTCTCCCAGCATGAGAGCCACCTACATGTCCGTGGCGGATGAA CACCTGAGACACTAC	102	Intron-spanning	96	0.9995
Ttyh3	CACTCTCGTCTTCGCAATGTCCGGCGCCTGCCTTCCCTTATCCACCCAGCTTACTTGGCTCTTCCTGTC	69	exonic	99	0.9999
Hprt1	GTAGATGGCCACAGGACTAGAACGTCTGCTAGTTCTTTACTGGCCACATCAACAGGACTCTTGTAGATTCAACTTGCCG	79	Exonic	98	0.9989
Sdha	CCTGAGCATTGCAGAATCTTGCAGGCCTGGAGATAAAGTTCCTCCGATTAAGGCAAATGCTGGAGAAGAGTCGGTTATGAATCTTGACAAG TTGAGATTTGCTGATGGAAGTGTAAGAACATCAGAGCTGCGCCTAAGCATGCAGAAG	118	Intron-spanning	97	0.998
Ywhaz	GTTGGAAGGCCGGTTAATTTTCCCCTCCTTCTCCCGCTTCTGCTTCGTCTCCTTGGGTATCCGATGTCCACAATGTCAAGTTGTCTCTCA GTAACTGCA	99	Intron-spanning	92	0.9998