**SKIL ENSG00000136603**

**2055 nt**

ATGGAAAACCTCCAGACAAATTTCTCCTTGGTTCAGGGCTCAACTAAAAAACTGAATGGGATGGGAGATGATGGCAGCCCCCCAGCGAAAAAAATGATAACGGACATTCATGCAAATGGAAAAACGATAAACAAGGTGCCAACAGTTAAGAAGGAACACTTGGATGACTATGGAGAAGCACCAGTGGAAACTGATGGAGAGCATGTTAAGCGAACCTGTACTTCTGTTCCTGAAACTTTGCATTTAAATCCCAGTTTGAAACACACATTGGCACAATTCCATTTAAGTAGTCAGAGCTCGCTGGGTGGACCAGCAGCATTTTCTGCTCGGCATTCCCAAGAAAGCATGTCGCCTACTGTATTTCTGCCTCTTCCATCACCTCAGGTTCTTCCTGGCCCATTGCTCATCCCTTCAGATAGCTCCACAGAACTCACTCAGACTGTGTTGGAAGGGGAATCTATTTCTTGTTTTCAAGTTGGAGGAGAAAAGAGACTCTGTTTGCCCCAAGTCTTAAATTCTGTTCTCCGAGAATTTACACTCCAGCAAATAAATACAGTGTGTGATGAACTGTACATATATTGTTCAAGGTGTACTTCAGACCAGCTTCATATCTTAAAGGTACTGGGCATACTTCCATTCAATGCCCCATCCTGTGGGCTGATTACATTAACTGATGCACAAAGATTATGTAATGCTTTATTGCGGCCACGAACTTTTCCTCAAAATGGTAGCGTACTTCCTGCTAAAAGCTCATTGGCCCAGTTAAAGGAAACTGGCAGTGCCTTTGAAGTGGAGCATGAATGCCTAGGCAAATGTCAGGGTTTATTTGCACCCCAGTTTTATGTTCAGCCTGATGCTCCGTGTATTCAATGTCTGGAGTGTTGTGGAATGTTTGCACCCCAGACGTTTGTGATGCATTCTCACAGATCACCTGACAAAAGAACTTGCCACTGGGGCTTTGAATCAGCTAAATGGCATTGCTATCTTCATGTGAACCAAAAATACTTAGGAACACCTGAAGAAAAGAAACTGAAGATAATTTTAGAAGAAATGAAGGAGAAGTTTAGCATGAGAAGTGGAAAGAGAAATCAATCCAAGACAGATGCACCATCAGGAATGGAATTACAGTCATGGTATCCTGTTATAAAGCAGGAAGGTGACCATGTTTCTCAGACACATTCATTTTTACACCCCAGCTACTACTTATACATGTGTGATAAAGTGGTTGCCCCAAATGTGTCACTTACTTCTGCTGTATCCCAGTCTAAAGAGCTCACAAAGACAGAGGCAAGTAAGTCCATATCAAGACAGTCAGAGAAGGCTCACAGTAGTGGTAAACTTCAAAAAACAGTGTCTTATCCAGATGTCTCACTTGAGGAACAGGAGAAAATGGATTTAAAAACAAGTAGAGAATTATGTAGCCGTTTAGATGCATCAATCTCAAATAATTCTACAAGTAAAAGGAAATCTGAGTCTGCCACTTGCAACTTAGTCAGAGACATAAACAAAGTGGGAATTGGCCTTGTTGCTGCCGCTTCATCTCCGCTTCTTGTGAAAGATGTCATTTGTGAGGATGATAAGGGAAAAATCATGGAAGAAGTAATGAGAACTTATTTAAAACAACAGGAAAAACTAAACTTGATTTTGCAAAAGAAGCAACAACTTCAGATGGAAGTAAAAATGTTGAGTAGTTCAAAATCTATGAAGGAACTCACTGAAGAACAGCAGAATTTACAGAAAGAGCTTGAATCTTTGCAGAATGAACATGCTCAAAGAATGGAAGAATTTTATGTTGAACAGAAAGACTTAGAGAAAAAATTGGAGCAGATAATGAAGCAAAAATGTACCTGTGACTCAAATTTAGAAAAAGACAAAGAGGCTGAATATGCAGGACAGTTGGCAGAACTGAGGCAGAGATTGGACCATGCTGAGGCCGATAGGCAAGAACTCCAAGATGAACTCAGACAGGAACGGGAAGCAAGACAGAAGTTAGAGATGATGATAAAAGAGCTAAAGCTGCAAATTCTGAAATCATCAAAGACTGCTAAAGAATAG

Forward (108-128: 20 bp) Tm: 60.05

TCATGCAAATGGAAAAACGASKIL\_100SKIL

Reverse（331-351: 20 bp）Tm: 60.07

CGACATGCTTTCTTGGGAAT

Product: 244 bp

produced by Primer3 (version 0.4.0)

GRCh38.p13 (Ensemble Genome Browser: release 100)