

TP53

<i>ext_probe_sequence</i>	<i>lig_probe_sequence</i>	<i>MIP_Name</i>
TCCCCACTTTTCCTCTT	TGGAAACTGTGAGTGGATCCATT	TP53_02_A1
GCAGCCAGACTGCCTTCGGG	GAAGGGCAGGCCACCACC	TP53_02_A2
TGCCCTTCCAATGGAT	GCAGTCTGGCTGCTGCAAGAGGAA	TP53_02_S1
GGTTGGGGTSGGGGTGGTGGG	ATGGCAGTGACCCGGAAGG	TP53_02_S3
CCCCAACCCAGCCCCCTAGCA	GGGGACCTGGAGGGCTGG	TP53_03_A1
CCCAGCCCTCCAGGTCCCAG	CTCTGCTAGGGGGCTGGGG	TP53_03_S5
AGGTCCCAGCCCTCCAGGTC	CTCTGCTAGGGGGCTGGGG	TP53_03_S6
CCAGGTCCCAGCCCAACCCTT	GCTAGGGGGCTGGGGTTG	TP53_03_S7
GACCAGGTCCTCAGCCC	CCCATGGAATTTTCGCTTCCCAC	TP53_03_S8
ACTGCTCTTTTACCCA	CTCCCAGAATGCCAGAGGCTGCT	TP53_04_A1
TGGTTCACTGAAGACCC	GGCCCTGTCATCTTCTGTCCCT	TP53_04_A2
GTGGCCCCTGCACCAGC	TGGGCTTCTTGCATTCTGGGACA	TP53_04_A3
GAAAACCTACCAGGGCAG	GCCTGGCCGTATCCCCCTGCAT	TP53_04_A4
CAGCATCAAATCATCC	CCCTCAGGTCCCAGCCCAACCC	TP53_04_S1
CACGSGGGGAGCAGCCTCT	GGGGGACTGTARATGGGTGAA	TP53_04_S2
GGTTTTCTGGGAAGGG	GGGAGCTTCATCTGGACCTGGGTC	TP53_04_S3
AGGGCAACTGACCGTGAAG	GAGGGGGCTGGTGCAGGGGC	TP53_04_S4
ACAAAAGAAATGCAGGGGG	CGGAAACCGTAGCTGCCCTGG	TP53_04_S5
ATCTGGACCTGGGTCTTCAGTG	AGGTCCTCAGCCCCCAG	TP53_04_S6
GTGAGGCGCTGCCCCACC	CTCTTAGGTCTGGCCCTCCT	TP53_05_A1
TGCCCTGTGCAGCTGTGGGT	AGCGCTGCTCAGATAGCGAT	TP53_05_A2
GTGCCCTGACTTTCAAC	ACCCCCGCCCGCACCCGCGTCC	TP53_05_A3
CTGTGCAGCTGTGGGTTGATT	TGCTCAGATAGCGATGGTG	TP53_05_A4
CCAGCTGCTCACCATC	GCGGGGGTGTGGAATCAACCACA	TP53_05_S1
GTCATGTGCTGTGACTGC	GTTGAGGGCAGGGGAGTACTGT	TP53_05_S2
ATAAGATGCTGAGGAGGGG	GAGCAGCGCTCATGGTGGGGG	TP53_05_S3
GCTGCACAGGGCAGGTCTTGG	TGGAAGACGGCAGCAAAGA	TP53_05_S4
CTTATCCGAGTGGAAGGA	TGGGAGGRGGGGTTAAGGGTGG	TP53_06_A1
TGGGGCTGGAGAGACGACAGG	GTGGAGTATTTGGATGACA	TP53_06_A2

AAAGCCCCCTACTGCTCA	CGAAAAGTGTTTCTGTCATCC	TP53_06_S1
GCTCATAGGGCACCAC	ATCAGTGAGGAATCAGAGGCCTGG	TP53_06_S2
CCACTACAACATACATGT	TGGCCTGCTGTGCCCCAGCCTCT	TP53_07_A1
TGCCACAGGTCTCCCCAAGGC	CATGAACCGGAGGCCCATC	TP53_07_A2
AGCAGGCCAGTGTGCAGGG	GTAGTGGATGGTGGTACAGTC	TP53_07_S1
TGGAGTCTTCCAGTGTG	GATAACACAGGCCCAAGATGAGG	TP53_07_S2
CGTGTTTGTGCTGTCCTGGG	AGAAGCGGTGGAGGAGACC	TP53_08_A1
AATGGGACAGGTAGGACCTGA	GAGACCGGCGCACAGAGGA	TP53_08_A2
GCTTCTTGTCTGCTT	GCACAAACACGCACCTCAAAGCTG	TP53_08_S1
AGATTCTTCTCTCTG	GCAGTAAGGAAATCAGGTCCTACC	TP53_08_S2
AGGAAAGAGGCAAGGAAAGG	GTGGTGAGGCTCCCCTTTCT	TP53_08_S3
CACTGCCCAACAACACCAGC	TAACACTCAAAATGCCGTTT	TP53_09_A1
GTAAGCAAGCAGGACAAGAAG	CCCAGCCAAAGAAGAAACC	TP53_09_A2
GAGTGTTAGACTGGAAACT	TGGGCAGTGCTAGGAAAGAGG	TP53_09_S1
TCTCCATCCAGTGGTTTCTTC	TCCTCCACCGCTTCTTGTC	TP53_09_S2
TAACTCAGGTACTGTG	TGCCCAGGCTGGGAAGGAGCCAGG	TP53_10_A1
TGGGCGTGAGCGCTTCGAGA	CCTTCCTGGCCCTWCTCCCC	TP53_10_A2
AGGAAGGGGCTGAGGT	GCTCACGCCACGGATCTGCAGCA	TP53_10_S1
AGCTCTCGAACATCTCG	GACTTTATGACGGTACAAAAGC	TP53_10_S2
TGAAGTCCAAAAGGGTC	CCCATCTCTCCCTCCCCTGCCA	TP53_11_A1
GGGAAAAGGGGCACAGACC	ATGTTCAAGACAGAAGGGCCT	TP53_11_A2
CATCTCTCCTCCCTGCTTCTG	CCACTTCTTGTTCCCCACT	TP53_11_A3
GACAGAAGGGCCTGACTCAG	CAATAGGTGTGCGTCAGAAG	TP53_11_A4
TGGGAGGCTGTCAGTGGGGAA	CTGTAGGAGACAGAAGCAG	TP53_11_S1