

53BP1 (mouse)

Plasmid:

pHR-SIN-CSGW HA (mouse) 53BP1 (lent)

Parent Plasmid:

pHR SIN CSGW

DU Number:

DU5731

Genbank:

NM_013735

Species:

Mouse

Synonyms:

TP53BP1

Sequence of Insert:

**ACTAGTGCCACCATGTACCCATACGATGTGCCAGATTACGCCATGGACCCTACTGGAAGTCAAT
TGGATTGAGATTCTCTCAGCAAGACACTCCTTGCCTGATAATAGAAGATTCTCAGCCTGAAAGC
CAGGTTCTGGAAGAAGATGCAGGCTCTCACTTCAGCGTGCTATCTCGACACCTTCTAATCTGCA
GATGCACAAAGAGAACCCCGTGTTGGATATTGTATCAAATCCGGAACAATCTGCTGTAGAACAA
GGAGACAGTAATAGCTCATTCAATGAACATCTGAAAGAAAAGAAAGCTTCAGATCCTGTGGAGT
CTTCTCATTGGGTACCAAGTGGTCCATCAGTCAGGTCATTGAACGGTTACCTCAGCCAAACAGG
ACAAGCAGTGCTCTGGCAGTGACAGTAGAAGCTGCTTCTCTCCAGAGGAGGAGAAGGAAGAA
GAGGAGTTAGAGGAGAAGGAAGGGGTGGGAGCAAACGCTCCCGGTGCTGACTCCCTTGCTGCT
GAAGATTCTGCTTCATCACAGTTGGGCTTTGGAGTTCTGGAAGTGTCCAGAGCCAGGATGTTG
AAGAACATACAGTGCCATATGATGTCAACCAAGAGCATCTGCAGTTGGTGACCACTAACTCGGG
TTCTAGCCCGCTATCTGATGTGGATGCGAGCACTGCAATTAATGTGAAGAACAGCCACTGAA
GATATTGCCATGATAGAACAGCCAGCAAAGACATCCCTGTTACAGTACAGCCCGGTAAGGTA
TCCATGTGGTAGAAGAACAAAATCTACCACTTGTAAAGTCCGAAGACCGGCGCTCCAGTCCTCA
AGTTTCTGTTGCTGCTGTGGAAACAAAGGAACAGGTACCTGCCCGGGAGCTGCTGGAAGAGGG
GCCGCAGGTTGAGCCGTCATCAGAGCCTGAGGTTTCTCAACCCAGGAGGACTTGTTTGACCAG
AGTAGTAAACAGCTTCTGATGGTTGTTCTACTCCTTCAAGGGAGGAAGGTGGGTGCTCTCCGG
TTTCCACACCTGCTACCACCCTGCAGCTCCTGCAGCTCTCTGGTCAGAAGCCCCTTGTTGAGGA
GAGTCTTCCACGAATTCCTCAGATCTTGTGCTCCTTCCCCTGATGCTTTCCGATCTACCCCTTT
TATCGTTCCTAGCAGTCCCACAGAGCAAGGAGGGAGAAAAGATGAGCCCATGGATATGTCAGT
GATACCTGCAGGAGGGGAGCCTTTCCAGAAGCTTCATGATGACGAAGCAATGGAGACAGAAAA
ACCCCTTCTCCCGTCTCAGCCTACTGTGTCACCGCAAGCATCAACACCAGTGTCTCGGAGCAGC
CCAGTTTTCACTCCTGGCTCTCTTCCATCCCGTCCCAGCCGAGTTCTCTCATGACATTTTCATT
CCATACCAAGTCTGGAAGAACCATCAGATGATGTGAAGAAAGGTGGAGGTTTACATAGTCTCAT
CTCTTACTGTTGAGTGTTCGAAGACTTCAGAGAGTGAACCAAGAATTTCACTGATGACCTTGGG
CTCTCCATGACAGGGGATTCTTGCAAACTGATGCTTTCTACAAGTGAGTATAGTCAGTCCTCAA
GATGGAGAGCTTGGGTTCTCCCAGGACTGAGGAAGACAGAGAGAACACACAGATTGACGATAC
TGAACCTTTGCTCCAGTTAGCAATTCTAACTTCTGCTGACAGTGAGAATGTCTGCTGACTC**

CATCGCAGGACGACCAGGTAGAAATGAGTCAGAATGTAGATAAAGCAAAAGAGGATGAAACGG
AGGACAGAGGTGACTGTAAAGGCAGAGAAGACGCGGTTGCTGAAGATGTTTGCATCGACCTCA
CTTGTGATTCTGGGAGTCAGGCAGTTCCTTCTCCAGCTACCCGTTCCGGAGGCACTTTCCAGTGT
TTAGATCAGGAGGAAGCTATGGACACTAAAGAACACCATCCAGAGGAAGGGTTTTCCGGGATCTG
AAGTAGAAGAAGTCCCTGAGACTCCCTGTGGAAGTCACAGAGAGGAGCCCAAAGAAGAAGCGA
TGGAAGTATCCCCTGACACTTTCTGACTGAAACCCAGTCTGAGGCATTGTGTCTGCAGAA
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CTGAGAATAGATTAGATACTCCAGAAGAAAAGCGTATAGAATGTGATGGA
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GGATTCTGAGTCTGCCCCAGAAATGGATGGAAAAGTGTCTCTGAAAATGA
AACTGGTTAGTCTGAGACAGAGGCCAGTGAAGAATCTTTGCAATTTAGC
CTGGAAAAGCCTACAAGTCTGAGAGAAAAAATGGATCTACTGCCATTGC
AGAGCCTGTTGCCAGTCTCCAGAAGCCGGTGCCTGTGTTTGGCTGCATCT
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CCTGTTTCGCAGCAGAGGGCCTCACAGGAGCCCTTCAGTCTGCAGAGGA
CGTGATGGAAACAGACCTGCTGGAAGGACTGGCTGCTAACCAGGACAGAC
CTAGTAAGATGTTGATGGACAGGCCACCCAGAGTAACATCGGGATCCAG
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GACTGTGAAGAGTGTGTGTGAGCAGGGGACCAGTACAGCGGAGCAGAACT
CTCGGAAACAAGATGCCACTGTGCAGACTGAGAGGGGGAGTGGAGAGAAA
CCTGCCAGTGTCCCGTGGACGACACAGAGTCCCTCCACAGCCAGGGGGA
AGAGGAATTTGAAATGCCCCAGCCTCCGCATGGCCATGTCTTGCATCGCC
ACATGAGAACCATTGAGAAGTCCGTACACTCGTCAACCCGCGTCATCACA
GATGTTTATTATGTGGATGGGACAGAAGTGGAAAGGAAAGTAACGGAGGA
GACTGAAGAACCAATCGTAGAGTGTGAGGAATGTGAAACAGAAGTTTCCC
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GAAAAGCCAGTGGGACAGAAGCTGCAGATTTTGCCTTACCCAGTTCCCGA
GGAGGCCCAGGAAAAGTGTGATCCTAGAAAAGGGATCAGTCAGACAGGGGC
ACCGGTGTGTGAGGAAGATGGTGTGATGCAGGCCTTGGCATCAGACAGGGAG
GGAAGGCTCCAGTTACACCTCGTGGGCGTGGGCGAAGGGGCCGCCACCT
TCTCGGACCACTGGAACCAGAGAAACAGTTGTCTCTGGTCCGTTGGGCGT
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TTATGCCTCGTGTGCCAGATTCTACCAAACGGACGGATGCCAGTTCTAGT
ACTTTGCGGCGGAGTGATTCTCCAGAGATTCTTTTACAGGCTGCTACTGG
TTCCTCTGATGGCTTGGATTCTCATCTTCAGGAAACAGCTTTGTGGGTC
TCCGTGTTGTAGCTAAGTGGTTCATCCAATGGCTACTTTTACTCTGGGAAG
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GTACGAATGTGACGTGCTGGGCAAAGACATTCTCCTGTGTGACCCCATCC

CCCTGGACACTGAAGTGACAGCCCTCTCAGAAGATGAATATTTTCAGTGCA
GGAGTGGTCAAAGGACATAGGAAGGAGTCTGGGGAGCTGTATTACAGCAT
TGAAAAAGAAGGCCAAAGGAAGTGGTATAAACGAATGGCGGTCATTCTGT
CCTTGGAGCAAGGAAACAGACTGAGAGAGCAATATGGGCTTGGCCCATAT
GAAGCTGTCACACCGCTCACGAAGGCAGCAGATATCAGCTTAGACAATTT
GGTGAAGGAAAGCGGAAACGGCGCAGTAACATCAGCTCCCCAGTCACCC
CCACTGCCGCCAGTAGCAGCAGCACAAACGCCACACGTAAAGCCACAGAG
AGTCCCCGTGCTTCCACGGGAGTCCCGTCAGGCAAAGGAAACTTCCGAC
TTCTGAAGAGGAACGCTCTCCTGCTAAGCGAGGTCGAAAGTCTGCCACCG
TGAAACCTGGAACAGTGGGGGCAGCAGAGTTTGTGAGTCCCTGTGAGACT
GGAGACAACATAGGTGAGCCTTCTGTCTAGAAAGAGCCAAGAGGGCCTTT
GCCCTCAACAAGACCTTGTCTGGGCTATGCCTTTCTCCTCACCATGG
CTACAAGTGTGACAAGCTGGCCAGTCGCTCTAAGCTGCTAGATGGTCCT
ACAGGAAGCAGTGAAGAAGAGGAGGAATTTTAGAAATTCCTCCTTTCAA
CAAGCAGTATACAGAATGCCAGCTTCGAGCAGGAGCTGGGTATATCCTTG
AAGACTTCAATGAAGCCCAGTGTAACACAGCCTACCAAGTGTCTCCTAATT
GCGGACCAGCACTGTGGAACCCGGAAGTACTTCTGTGCCTTGCCAGTGG
CATTCTTGTGTGTCTCATGTCTGGGTCCATGACAGTTGCCATGCCAACC
AACTTCAAACTATCGTAATTATCTGCTGCCTGCTGGGTATAGCCTTGAA
GAGCAACGAATTCTGGATTGGCAACCCCGTGAAAACCCTTTCCAGAATCT
GAAGGTCTCTTGGTGTGAGATCAACAACAGAACTTCTTGGAGCTCTGGT
CTGAGATCCTCATGACTGGAGGGGCAGCCTCTGTGAAGCAGCACCATTCA
AGTGCCCAACAACAAGACATTGCTTTAGGGGTATTTGATGTGGTGGTGAC
AGACCCCTCATGCCAGCCTCGGTGCTCAAGTGTGCTGAAGCCTTGCAAC
TGCCTGTGGTATCACAAGAATGGGTGATCCAGTGCCTCATTGTTGGGGAG
AGAATTGGATTCAAGCAGCATCCAAAATATAAACATGATTATGTTTCTCA CTAACGCGT

Amino Acid Sequence:

MYPYDVPDYAMDPTGSQLDSDFSQQDTPCLIEDSQPESQVLEEDAGSHF
SVLSRHLPLNMQMHKENPVLDIVSNPEQSAVEQGDSNSSFNEHLKEKKASD
PVESSLGTSISQVIERLPQPNRTSSALAVTVEAASLPEEEKEEELE
EKEGVGANAPGADSLAAEDSASSQLGFGVLELSQSQDVEEHTVPYDVNQE
HLQLVTTNSGSSPLSDVDASTAIKCEEQPTEDIAMIEQPSKDIPVTVQPG
KGIHVVEEQNLPLVRSEDRPSSPQVSVAAVETKEQVPARELLEEGPQVQP
SSEPEVSSTQEDLFDQSSKTASDGCSTPSREEGGCSPVSTPATTLLQLLQL
SGQKPLVQESLSTNSSDLVAPSPDAFRSTPFIVPSSPTEQGGRKDEPMDM
SVIPAGGEPFQKLHDDEAMETEKPLLPSQPTVSPQASTPVSRSSTPVFTPG
SLPIPSQPEFSDIFIPSPSLEEPSDDVKKGGGLHSSSLTVECSKTSESE
PKNFTDDLGLSMTGDCKLMLSTSEYSQSSKMESLGSPTREEDRENTQID
DTEPLSPVSNKLPADSENVLVTSPQDDQVEMSNVDKAKEDETEDRGDC
KGREDAVAEDVCIDLTCDSGSQAVPSPATRSEALSSVLDQEEAMDTKEHH
PEEGFSGSEVEVPETPCGSHREEPKEEAMESIPLHLSTTETQSEALCLQ
KEAPKEECPEAMEVETSVISIDSPQKLQVLDQELEHKDPDTWEEATSEDS
SVVIVDVKEPSPRADVSCPELEEVEKCSDSQSWEGVAPEEEPCAENRLDT
PEEKRIECDGDSKAETTEKDAVTEDSPQPPLPSVRDEPVRPDQETQQPQV
QEKESPVTVDAEVADDKQLGPEGACQQLKAPACASQSFCESSSGISNYP
ESTIATSDVTSESMVEINDPLLGNEKGDSESAPEMDGKLSLKMMLVSPET
EASEESLQFSLEKPTTAERKNGSTAIAEPVASLQKVPVFGCIYEAQKEK
EAQSEAPPSAPDRANLLHFPSAQEEDKERPDVTPKLRQSEQPVKPVGPVM
DDAAPEDSASPVSQQRASQEPFSPAEDVMETDLLEGLAANQDRPSKMLMD
RPTQSNIGIQTVDHSLCAPETVSAATQTVKSVCEQGTSTAEQNSRKQDAT
VQTERGSGEKPASAPVDDTESLHSQGEFEMPQPPHGHVLRHRMRTIRE
VRTLVTRVITDVYYVDGTEVERKVTEETEEPIVECQECETEVSPTSQGGG
SGDLGDISSFSSKASSSHHTSSGTSLSAIHSSGSSGRGAGPLKGKASGTE

AADFALPSSRGGPGKLSRKGISQTGAPVCEEDGDAGLGIRQGGKAPVTP
RGRGRRGRPPSRTTGTRETVVSGPLGVEDISPSMSPDDKSFTRIMPRVPD
STKRTDASSSTLRRSDSPEIPFQAATGSSDGLDSSSSGNSFVGLRVVAKW
SSNGYFYSGKITRDVGAGKYKLLFDDGYECDVLGKDILLCDPIPLDTEVT
ALSEDEYFSAGVVKGHRKESGELYYSIEKEGQRKWKYKRMVILSLEQGNR
LREQYGLGPYEAVTPLTKAADISLDNLVEGKRKRNRNISPVTPTAASSS
STTPTRKATESPRASTGVPSGKRKLPTSEEERSPAKRGRKSATVKPGTVG
AAEFVSPCETGDNIGEPSVLEEPRGPLPLNKTFLFLGYAFLLMATTSDKL
ASRSKLLDGPTGSSEEEEEEFLEIPPFNKQYTECQLRAGAGYILEDFNEAQ
CNTAYQCLLIADQHCRTRKYFLCLASGIPC VSHVWVHDSCHANQLQNYRN
YLLPAGYSLEEQRILDWQPRENPFQNLKVLLVSDQQNFLELWSEILMTG
GAASVKQHHSSAHNKDIALGVFDVVVTD PSCPASVLKCAEALQLPVVSQE
WVIQCLIVGERIGFKQHPKYKH DYVSH*

Antibiotic:

Amp

Comments:

Plasmid for lentiviral expression in mammalian cells (SFFV promoter) N.B. Nucleotide sequence differences compared with NM_013735 CDS: Delta nt 2654-2773 (delta aa 885-924). t3255c, c3441t, t3468g, a4947g (all silent).

Price per aliquot:

£110.00



University
of Dundee