



## 53BP1

Expressed:  
**mCherry 53BP1**

Plasmid:  
**pcDNA5-FRT/TO-mCherry 53BP1**

Parent Plasmid:  
**pcDNA5 FRT/TO mCherry**

DU Number:  
**DU33919**

Genbank:  
**NM\_001141980.1**

Species:  
**Human**

Synonyms:  
**TP53BP1, TDRD30, TP53, p202, p53BP1**

Sequence of Insert:

```
GGTACCGCCACCATGGACCCTACTGGAAGTCAGTTGGATTTCAGATTTCTCTCAGCAAGATACTC  
CTTGCCTGATAATTGAAGATTCTCAGCCTGAAAGCCAGGTTCTAGAGGATGATTCTGGTTCTCAC  
TTCAGTATGCTATCTCGACACCTTCTAATCTCCAGACGCACAAAGAAAATCCTGTGTTGGATGT  
TGTGTCCAATCCTGAACAAACAGCTGGAGAAGAACGAGGAGACGGTAATAGTGGGTTCAATGA  
ACATTTGAAAGAAAACAAGGTTGCAGACCCTGTGGATTCTTCTAACTTGGACACATGTGGTTCCA  
TCAGTCAGGTCATTGAGCAGTTACCTCAGCCAAACAGGACAAGCAGTGTCTGGGAATGTCAGT  
GGAATCTGCTCCTGCTGTGGAGGAAGAGAAGGGAGAAGAGTTGGAACAGAAGGAGAAAGAGA  
AGGAAGAAGATACTTCAGGCAATACTACACATTCCCTTGGTGCTGAAGATACTGCCTCATCACA  
GTTGGGTTTTGGGGTTCTGGAAGTCTCCAGAGCCAGGATGTTGAGGAAAATACTGTGCCATAT  
GAAGTGGACAAAGAGCAGCTACAATCAGTAACCACCAACTCTGGTTATAACCAGGCTGTCTGATG  
TGGATGCTAATACTGCAATTAAGCATGAAGAACAGTCCAACGAAGATATCCCCATAGCAGAACA  
GTCCAGCAAGGACATCCCTGTGACAGCACAGCCCAGTAAGGATGTACATGTTGTAAAAGAGCA  
AAATCCACCACCTGCAAGGTCAGAGGACATGCCTTTTAGCCCCAAAGCATCTGTTGCTGCTATG  
GAAGCAAAGAACAGTTGTCTGCACAAGAACTTATGGAAAGTGGACTGCAGATTCAGAAGTCAC  
CAGAGCCTGAGGTTTTGTCAACTCAGGAAGACTTGTGTTGACCAGAGCAATAAAACAGTATCTTCT  
GATGGTTGCTCTACTCCTTCAAGGGAGGAAGGTGGGTGTTCTTTGGCTTCCACTCCTGCCACCAC  
TCTGCATCTCCTGCAGCTCTCTGGTCAGAGGTCCCTTGTTCAGGACAGTCTTTCCACGAATTCTT  
CAGATCTTGTGCTCCTTCTCCTGATGCTTTCCGATCTACTCCTTTTATCGTTCCTAGCAGTCCCA  
CAGAGCAAGAAGGGAGACAAGATAAGCCAATGGACACGTCAGTGTATCTGAAGAAGGAGGAG  
AGCCTTTTCAGAAGAACTTCAAAGTGGTGAACCAGTAGAGTTAGAAAACCCCTCCTCCTGCC  
TGAGTCCACTGTATCACCACAAGCCTCAACACCAATATCTCAGAGCACACCAGTCTTCCCTCCTG  
GGTCACTTCTATCCCATCCCAGCCTCAGTTTTCTCATGACATTTTTATTCTTCCCCAAGTCTGG  
AAGAACAATCAAATGATGGGAAGAAAGATGGAGATATGCATAGTTCATCTTTGACAGTTGAGTG  
TTCTAAAACCTCAGAGATTGAACCAAAGAATTCCCCTGAGGATCTTGGGCTATCTTTGACAGGGG
```

ATTCTTGCAAGTTGATGCTTTCTACAAGTGAATATAGTCAGTCCCCAAAGATGGAGAGCTTGAGT  
TCTCACAGAATTGATGAAGATGGAGAAAACACACAGATTGAGGATACGGAACCCATGTCTCCAG  
TTCTCAATTCTAAATTTGTTCTGCTGAAAATGATAGTATCCTGATGAATCCAGCACAGGATGGT  
GAAGTACAAGTGAAGTGAAGTGAAGTGAAGTGAAGTGAAGTGAAGTGAAGTGAAGTGAAGTGAAGT  
ATTAGTATTTTAGCCACTGGTTGCAAGGGCAGAGAAGAAACGGTAGCAGAAGATGTTTGTATTG  
ATCTCACTTGTGATTCGGGGAGTCAGGCAGTTCCGTCAACCAGCTACTCGATCTGAGGCACTTTCT  
AGTGTGTTAGATCAGGAGGAAGCTATGGAAATTAAGAACACCATCCAGAGGAGGGGTCTTCA  
GGGTCTGAGGTGGAAGAAATCCCTGAGACACCTTGTGAAAGTCAAGGAGAGGAACTCAAAGAA  
GAAAATATGGAGAGTGTCCGTGTCACCTTTCTCTGACTGAAACTCAGTCCCAAGGGTTGTGTCT  
TCAAAGGAAATGCCAAAAAAGAATGCTCAGAAGCTATGGAAGTTGAAACCAGTGTGATTAGT  
ATTGATTCCCCTCAAAGTTGGCAATACTTGACCAAGAATTGGAACATAAGGAACAGGAAGCTT  
GGGAAGAAGCTACTTCAGAGGACTCCAGTGTGTCATTGTAGATGTGAAAGAGCCATCTCCAG  
AGTTGATGTTTCTTGTGAACCTTTGGAGGGAGTGGAGAAGTGTCTCAGATTCCCAGTCATGGGAG  
GATATTGCTCCAGAAATAGAACCATGTGCTGAGAATAGATTAGACACCAAGGAAGAAAAGAGTG  
TAGAATATGAAGGAGATCTGAAATCAGGGACTGCAGAAACAGAACCTGTAGAGCAAGATTCTTC  
ACAGCCTTCTTACCTTTAGTGAGAGCAGATGATCCTTTAAGACTTGACCAGGAGTTGCAGCAG  
CCCCAAACTCAGGAGAAAACAAGTAATTCATTAACAGAAGACTCAAAAATGGCTAATGCAAAGC  
AGCTAAGCTCAGATGCAGAGGCCAGAAGCTGGGGAAGCCCTCTGCCATGCCTCACAAAGCT  
TCTGTGAAAGTTCTAGTGAAACCCATTTCACTTTGCTAAAGAAGGTGATATCATCCCA  
CCATTGACTGGTGCAACCCACCTCTATTGGGCACCTAAAATTGGAGCCCAAGAGACACAGTA  
CTCCTATTGGTATTAGCAACTATCCAGAAAGCACCATAGCAACCAAGTGTGTCATGTCTGAAAGC  
ATGGTGGAGACCCATGATCCCATACTTGGGAGTGGAAAAGGGGATTCTGGGGCTGCCCCAGAC  
GTGGATGATAAATTATGTCTAAGAATGAAACTGGTTAGTCTGAGACTGAGGCGAGTGAAGAGT  
CTTTGCAGTTCAACCTGGAAAAGCCTGCAACTGGTGAAAGAAAAAATGGATCTACTGCTGTTGC  
TGAGTCTGTTGCCAGTCCCCAGAAGACCATGTCTGTGTTGAGCTGTATCTGTGAAGCCAGGCAA  
GAGAATGAGGCTCGAAGTGAAGATCCCCCACCACACCCATCAGGGGGAAGTTGCTCCACTTTC  
CAAGTTCTCAAGGAGAAGAGGAGAAGAAAAAATTGGAGGGTGACCATACAATCAGGCAGAGTC  
AACAGCCTATGAAGCCCATTAGTCTGTCAAGGACCCTGTTTCTCCTGCTTCCAGAAAGATGGTC  
ATACAAGGGCCATCCAGTCTCAAGGAGAGGCAATGGTGACAGATGTGCTAGAAGACCAGAAA  
GAAGGACGGAGTACTAATAAGGAAAATCCTAGTAAGGCCTTGATTGAAAGGCCCAGCCAAAAT  
AACATAGGAATCCAAACCATGGAGTGTTCCTTGAAGGGTCCAGAAACTGTTTCAGCAGCAACCC  
AGACTATAAGAATGTGTGTGAGCAGGGGACCAGTACAGTGGACCAGAAGTTTGGAAAGCAAG  
ATGCCACAGTTCAGACTGAGAGGGGGAGTGGTGAGAAACCAGTCAGTGTCTCCTGGGGATGATA  
CAGAGTCGCTCCATAGCCAGGGAGAAGAAGAGTTTGATATGCCTCAGCCTCCACATGGCCATGT  
CTTACATCGTCACATGAGAACAAATCCGGGAAGTACGCACACTTGTCACTCGTGTCAATTACAGAT  
GTGTATTATGTGGATGGAACAGAAGTAGAAAGAAAAGTAACTGAGGAGACTGAAGAGCCAATT  
GTAGAGTGTGAGGAGTGTGAAACTGAAGTTTCCCCTTACAGACTGGGGGCTCCTCAGGTGACC  
TGGGGGATATCAGCTCCTTCTCCTCCAAGGCATCCAGCTTACACCGCACATCAAGTGGGACAAG  
TCTCTCAGCTATGCACAGCAGTGAAGCTCAGGGAAAGGAGCCGGACCACTCAGAGGGAAAAC  
CAGCGGGACAGAACCCGCAGATTTTGCCTTACCAGCTCCCGAGGAGGCCAGGAAAACCTGAG  
TCCTAGAAAAGGGGTGAGTCAAGACAGGGACGCCAGTGTGTGAGGAGGATGGTGTGAGGCCT  
TGGCATCAGACAGGGAGGGAAGGCTCCAGTCACGCCTCGTGGGCGTGGGCGAAGGGGCCGCC  
CACCTTCTCGGACCACTGGAACCAGAGAAACAGCTGTGCCTGGCCCCTTGGGCATAGAGGACA  
TTTACCTAACTTGTCAACCAGATGATAAATCCTTACAGCCGTGTGCTGCCCCGAGTGCAGACTCC  
ACCAGACGAACAGATGTGGGTGCTGGTGTCTTTCGCTCGTAGTGACTCTCCAGAAATTCCTTTC  
AGGCTGCTGCTGGCCCTTCTGATGGCTTAGATGCCTCCTCTCCAGGAAATAGCTTTGTAGGGCTC  
CGTGTGTTAGCCAAGTGGTCATCCAATGGCTACTTTACTCTGGGAAAATCACACGAGATGTGCG  
GAGCTGGGAAGTATAAATTGCTCTTTGATGATGGGTACGAATGTGATGTGTTGGGCAAAGACAT  
TCTGTTATGTGACCCATCCCGCTGGACACTGAAGTGACGGCCCTCTCGGAGGATGAGTATTTTC  
AGTGCAGGAGTGGTGAAGGACATAGGAAGGAGTCTGGGGAAGTACTACAGCATTGAAAAA  
GAAGGCCAAAGAAAGTGGTATAAGCGAATGGCTGTATCCTGTCTTGGAGCAAGGAAACAGA  
CTGAGAGAGCAGTATGGGCTTGGCCCCTATGAAGCAGTAACACCTCTTACAAAGGCAGCAGATA  
TCAGCTTAGACAATTTGGTGAAGGGGAAGCGGAAACGGCGCAGTAACGTGAGCTCCCCAGCCA  
CCCCTACTGCCTCAGTAGCAGCAGCACAACCCCTACCCGAAAGATCACAGAAAGTCTCCTGTC

CTCCATGGGAGTTCTCTCAGGCCAAAAGAAAACCTTATCACTTCTGAAGAGGAACGGTCCCCTGCC  
AAGCGAGGTCGCAAGTCTGCCACAGTAAAACCTGGTGCAAGTAGGGGCAGGAGAGTTTGTGAGC  
CCCTGTGAGAGTGGAGACAACACCGGTGAACCCTCTGCCCTGGAAGAGCAGAGAGGGCCTTTG  
CCTCTCAACAAGACCTTGTCTGGGCTACGCATTTCTCCTTACCATGGCCACAACCAAGTGACAA  
GTTGGCCAGCCGCTCCAAACTGCCAGATGGTCTACAGGAAGCAGTGAAGAAGAGGAGGAATT  
TTTGAAATTCTCCTTTCAACAAGCAGTATACAGAATCCCAGCTTCGAGCAGGAGCTGGCTATA  
TCCTTGAAGATTTCAATGAAGCCCAGTGTAAACACAGCTTACCAGTGTCTTCTAATTGCGGATCAG  
CATTGTGCAACCCGGAAGTACTTCTGTGCCTTGCCAGTGGGATTCCTTGTGTGTCTCATGTCTG  
GGTCCATGATAGTTGCCATGCCAACCAGCTCCAGAACTACCGTAATTATCTGTTGCCAGCTGGG  
TACAGCCTTGAGGAGCAAAGAATTCTGGACTGGCAACCCCGTGAAAATCCTTTCCAGAATCTGA  
AGGTACTCTTGGTATCAGACCAACAGCAGAACTTCTGGAGCTCTGGTCTGAGATCCTCATGAC  
TGGTGGTGCAGCCTCTGTGAAGCAGCACCATTCAAGTGCCATAACAAAGATATTGCTTTAGGG  
GTATTTGATGTGGTGGTGACGGACCCCTCATGCCAGCCTCGGTGCTGAAGTGTGCTGAAGCAT  
TGCAGCTGCCTGTGGTGTCAACAAGAGTGGGTGATCCAGTGCCTCATTGTTGGGGAGAGAATTGG  
ATCAAGCAGCATCCAAAATATAAACACGATTATGTTTCTCACTAAGCGGCCG

Amino Acid Sequence:

MVSKGEEDNMAIIKEFMRFKVMHEGSVNGHEFEIEGEGEGRPYEGTQTAKLKVTKGGPLPFAWDILS  
PQFMYGSKAYVKHPADIPDYLKLSFPEGFKWER VMNFEDGGVVTVTQDSSLQDGEFIYKVKLRGTN  
FPSDGPVMQKKTMGWEASSERMYPEDGALKGEIKQRLKLDGGHYDAEVKTTYKAKKPVQLPGAY  
NV NIKLDITSHNEDYTIVEQYERAEGRHSTGGMDELYKSGLGSSRKGTATMDPTGSQLDSDFSQQDT  
PCLIEDSQPESQVLEDDSGSHFSMLSRHLPNLQT HKENPVLDVVSNPEQTAGEERGDNNSGFNEHL  
KENKVADPVDSSNLDTCGSISQVIEQLPQPNRTSSVLGMSVESAPAVEEEEKGEELEQKEKEKEEDTS  
G NTHSLGAEDTASSQLGFGVLELSQSQDVEENTVPYEVDKEQLQSVTTNSGYTRLSDVDANTAIAKH  
EEQSNEDIPIAEQSSKDIPVTAQPSKDVHVVEQ NPPPARSEDMPFSPKASVAAMEAKEQLSAQEL  
MESGLQIQKSPEPEVLSTQEDLFDQSNKTVSSDGCSTPSREEGGCSLASTPATTLLHLLQLSGQRSLV  
Q DSLSTNSSDLVAPSPDAFRSTPFIVPSSPTEQEGRQDKPMDTSVLSEEGGEPFQKQLQSGEPVELE  
NPPLLPESTVSPQASTPISQSTPVFPFGLPIPS QPQFSHDIFIPSPSLEEQSNKGKDGDMHSSSLTV  
ECSKTSEIEPKNSPEDLGLSLTGDSCKLMLSTSEYSQSPKMESLSSHRIDEDGENTQIEDTEPMS PVL  
NSKFVPAENDSILMNPAQDGEVQLSQNDDKTKGDDTDTRDDISILATGCKGREETVAEDVCIDLTCDS  
GSQAVPSPATRSEALSSVLDQEEAMEIKE HHPEEGSSGSEVEEIPETPCESQGEELKEENMESVPLH  
LSLTETQSQGLCLQKEMPKECSEAMEVETSVISIDSPQKLAILDQELEHKEQEAWEATSE DSSVVI  
VDVKEPSRVDVSCEPLEGVEKCSQSWEDIAPEIEPCAENRLDTKEEKSVEYEGDLKSGTAETEP  
VEQDSSQPSLPLVRADDPLRLDQELQQ PQTQEKTSNSLTEDSKMANAKQLSSDAEAQKLGKPSAH  
ASQSFCESSSETPFHFTLPKEGDIIPPLTGATPPLIGHLKLEPKRHSTPIGISNYPESTIAT SDVMSESM  
VETHDPILGSGKGDGAAPDVDDKLCRLMMLVSPETEASEESLQFNLEKPATGERKNGSTAVAESV  
ASPQKTMVLSLSCICEARQENEARSED PPTPIRGNLLHFPSSQGEEEEEKLEGDHTIRQSQQPMKPI  
SPVKDPVSPASQKMVIQGPSSPQGEAMVTDVLEDQKEGRSTNKENPSKALIERPSQNNI GIQTMECS  
LRVPETVSAATQTIKNVCEQGTSTVDQNFQKQDATVQTERGSGEKPVSAAGDDTESLHSQGEEFFD  
MPQPPHGHVLRHMRTIREVRTLVR VITDVYYVDGTEVERKVTEETEEPIVECQECETEVSPTSQTGG  
SSGDLGDISSFSSKASSLHRTSSGTSLSAMHSSGSSGKGAGPLRGKTSGETPADFALP SSRGGPGK  
LSPRKGVSQTGTPVCEEDGDAGLGIRQGGKAPVTTPRGRGRRRGRPPSRTTGTRETAVPGPLGIEDISP  
NLSPDDKSFVSRVPRVPDSTRRTDV GAGALRRSDSPEIPFQAAAGPSDGLDASSPGNSFVGLRVVA  
KWSSNGYFYSGKITRDVGAGKYKLLFDDGYECDVLGKDILLCDPIPLDTEVTALSEDEY FSAGVVKG  
HRKESGELYYSIEKEGQRKWKYKRMVILSLEQGNRLREQYGLGPYEAVTPLTKAADISLDNLVEGKR  
KRRSNVSSPATPTASSSSSTTPTRK ITESPRASMGVLSGKRKLITSEEERSPAKRGRKSATVKPGAV  
GAGEFVSPCESGDNTGEPSEALEEQRGPLPLNKTFLGYAFLLTMATTSDKLASRSKLP DGPTGSSE  
EEEEFLEIPPFNKQYTESQLRAGAGYILEDNFNAQCNTAYQCLLIADQHCRTRKYFLCLASGIPCUSHV  
WVHDSCHANQLQNYRNYLLPAGY SLEEQRILDWQPRENPFQNLKVVLLVSDQQQNFLELWSEILMTG  
GAASVKQHHSSAHNKDIALGVFDVVVTDPSCPASVLKCAEALQLPVVSQEWVIQCLI  
VGERIGFKQHPKYKHDIYVSH\*

Antibiotic:

## Amp

Comments:

**Contains two silent nucleotide changes as compared to the ref sequence; G>A 1245bp and G>A 3153 bp.**

Price per aliquot:

**£110.00**

