

Product datasheet for MR216449

Tspoap1 (NM_172449) Mouse Tagged ORF Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	Tspoap1 (NM_172449) Mouse Tagged ORF Clone
Tag:	Myc-DDK
Symbol:	Tspoap1
Synonyms:	Bzrap1; D230016K05; mKIAA0612; PRAX-1
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry2 (PS100063)
E. coli Selection:	Kanamycin (25 ug/mL)
ORF Nucleotide Sequence:	>MR216449 representing NM_172449 Red=Cloning site Blue=ORF Green=Tags(s)

TTTTGTAATACGACTCACTATAGGGCGGCCGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
GCC**CGATCGCC**

ATGGAGCAACTGACAACCCTCCCACGGCTTGGAGACCTTGGAGCCATGGAGCCATGGGCACTGCCTGCCT
GGCAGCACTGGACTCAGGGCCAGGGGTGCAAACCTGGAGATGCATCTCCAAGCATTGCTGGTACTCCGAC
AGCTCTGCAGGTTAAAGGATTGAGGTTTGAAGAGAGTTCCAAGCCTGAGGGAGCTCATAGCCCTGGACCT
GTCGGAATACTGATCCTGAAGCAACAGAGACCGGGCTGCCTAAGCTGGGGCAGCAAGCAGAGAGCCCTG
GGTACAGCTGTTCCGGGCTGGAGGAGGAGGACAGGCTTATAAGGCCAAGTTCAACATAGGCTTCGG
GGACAGGCCTAATCTGGAGCTGCTAAGGGCCCTGGGAGAGCTGCAGCAGCGCTGTACCATCTGAAGGAG
GAAAACAGATGCTGAGAAAAAGCAGCTTCCCAGAGACGGAGGAGAAGGTACGGAGGCTGAAGCGGAAGA
ATGCTGAACTGGCAGTCATTGCCAAGCGCTGGAGGAGAGGGCACAGAAGCTGCAGGAAACGAACATGAG
GGTGGTGAAGTGCCTGTGCCCGACCCGGATCCAGTTTGGAGTTGTGCCGTAAGGCTCTAGCTCGCCAG
CGAGCCCAGACCTCAGTGAGACAGCCAGTGCCTGTTGGCAAGGACAAACAGATTGCTGCCTTGACAGC
GGGAGTGCAGGGAGCTGCAGGCCAGACTCTCTGGTGGCAAGGAAGTCCCAGTGGCTGCATATGCC
GGACTTCGACCGGTTGCTGCGCGAGTCCCAACGGGAGGTGCTGCGGCTGCAGAGGCAGATCGCCCTGCC
AACCAGCGGGAGCCGCTCCGGCTGCCCGGTCCCGGCTCTACTGCCCATCTAGAGTAGGGGCGCCGG
CCCCCGGGCCCCGGGAGAGGCCGTAATTCAGGATGATGTGGAGAGCCACAGGTAAGTCTAAGGGAACC
AGAGAAGCAGCAAAGGGTGCAGCAGCTGGAGTCTGAGCTCTGCAAGAAGCGAAAGAAATGCGAGAGTCTG
GAGCAGGAAGCCAGGAAAAAGCAGAGGCGATGTGAGGAGCTGGAGCTACAGCTGAGGGCAGCCCAGAATG
AAAATGCCCGCTGGTGGAGGAGAACTCTCGGCTCAGTGGGAGAGCCACAGAGAAGGAGCAGGTGGAATG
GGAGAATTCGGAGCTGAAGGGACAGCTCCTGGGGTGACACAAGAGAGGGACTCGGCCCTCTTAAGAGC
CAGGGCCTGCAAAGCAAGCTGGAGAGCTGGAGCAGGTGCTGAAGCATATGCGGGAGGTGGCCAGCGCC
GGCAGCAGCTGGAGGTGGAGCATGAGCAGGCTAGGCTCAGCCTGCAGGAGAAGCAGGAGGAGGTCCGGAG
GCTGCAGCAGGCCAGGCAAGCAAGAGGGAACATGAGGGAGCCGTTACAGCTGCTGGAGTCTACCTTG



[View online »](#)

GATTCCATGCAGGCCCGGGTTCGAGAGCTTGAGGGCCAGTGCCGAAGCCAGACTGAGCGTTTCAGCCTTC
 TGGCTCAGGAGCTCCAGGCCCTCCGTCTGCACCCAGGCCCTCTGGATCTGCTTACTTCAGCCTTGGGCTG
 TAGTGCCCTTGGGGACCACCCGCCCCCACTGCTGCTGCTCTATCCCTCAGCCCTGCCAGGGGTCCGGC
 CCCAAAGATCTTGACCTCCCACCGGGCTCCCAGGACGCTGCACCCCAAATCTTCTGAACCTGCTCTCA
 CCACCTTACTGGAATCCCTCGAAGGACAGCTAAGAAGGCCGAGTCTCTTTCTAATCTTCTCGCTCAGA
 GTCCATCCACAACAGCCCAAGTCATGTCCACACCAGAGGTGGACACAGCCAGTGAGGTGGAGGAACTG
 GAGGTAGACAGTGTTCCTGCTCCCAGCAGCTCCGGAGAGTCACTCGGGAGGAGCCAGAAATCCAGTCT
 TCCTAGCACGCTATAGCTACAACCCTTTGAGGGTCCCAATGAGAATCCAGAGGCCGAGCTTCTCTGAC
 AGCGGGCGAGTATATCTACATCTATGGTAACATGGACGAGGATGGCTTTTTTGAAGGGGAGCTCATGGAT
 GGCCGAAGGGGCTGGTCCCTTCCAACCTTGTAGAGCGCGTGTCTGATGATGATCTTCTGTCCACCTCC
 CTCGGGAGCTGGCTGATTCGTGCGACAGCTCAGGCCCGAGCTCAGTTTCTGAGTGGAGGCGGGGTGG
 TTGCAGTAGTGGGGCCAGAGCAGCGGGGACGTAGCCAGCCAGACCAGAGGAGGAGGCCCGAGGAGAT
 GAACTCAGTCTGAGCCCCCACCAGAGGGACTTGGCGAGCCTCTGGCTGTGCCTTACCCCGACACATCA
 CGGTTCTCAAGCAGTTAGCCACAGTGTGGTGTGGCCTGGGAGTTGCCTCCTGAGAGAGTAGATCTGCG
 TGGCTTCCATATCTTTGCAATGGGAACTCCGTAGGCCCTTGGGGCCTGGGGTCCCCCAAGCTGTG
 CTTGAAAACATGGACCTGCGGACTGGGCCTCTCCATGTGTCTGTCCAGGCCCTAACCCAGCAAGGGCAGCT
 CAGACCCTCTGCGCTGTTGCCTGGCTGTGGGTGCCGGGGCTGGGGTGGTACCCAGCCAGCTGCGGATCCA
 TCGGCTGACAGCCACATCTGCTGAGATCGCTGGGTGCCGGGAATAGCAACTTGGCCCATGCCATCTAC
 CTCAATGGAGAAGAGTGGCCACCTGCTCGCCCCAGCACATATTGGGCAACCTTTTGTAACTGAGACCTG
 GCACACTCTATCAGGCCCGAGTGGAGGCTCAGATCCCATCTCAGGGGCCCTGGGAACCAGGCTGGGAGAG
 GCCAGAGCAGAGAGCTGTACCTTGCAGTTCACCACACTTCCAGCAGGTCTGCCTGATGCCCCACTGGAT
 GTACAGGCTGAACCAGGACCCTCTCTGGAATCTTGATGATCAGTTGGCTCCCTGTCACCATTGATGCTG
 CTGGTACCTCCAATGGTGTCCGGTTACAGGCTATGCCATCTATGCTGATGGGCAGAAGATTAGGAGGT
 GGCTTCCCCACAGCAGGCAAGTGTGCTGGTGGAGGTGTCCAGCTGCAGCTGTTGCAGGCCCTGCCATGAG
 GTGACTGTACGCACTATGTCACCCCATGGCGAGTCCAGTGACTCCATCCCAGGCCCGGTTGCCCCAGCCC
 TGGCTTCTGCCTGCCAGCCAGCCAGGATGTCCTGTCTCTCACCACGACCAAGCCAGAGGTGAGAACACC
 CCTTGCTCAGTCTCCCCAGGGCTTGGGGATACCAGCTTTCCTCCGGCATCTGTCCCCCATGGAACCT
 CAAGATTTCTCTGCAAGTCTTCCATAGAGATGTCAAAGGACCCAGGAGGAACCTCCAGTCCCTTGTCT
 CTCAGGAAGAGGCTGGGGCAGCTGTGCGGAGCATCTCAGAAGAGAAGAGGGCTATCGAGCCAACCTGCG
 CCAGGAAGGCCCTGAGCCTGTGGCTCCTTCCCTGGCTAAGCAGGAAGTGGAGTGCATTCAGGAGATGCT
 GGCCTGTACCTGCTCCACCAAGGAGAGCTGACCCAGAAGAAGCAAGTATTGAAGCCTGCCATGGGG
 GAGATCTGGACTCCGGGCTGAAACTTAGATCTGAGAAAGAAGATATGTCAGAGCTTGGAGTTCACCTGGT
 GAATCCCTTGTGGATCACAGCCCAACTCAGACTTATCAGACATCCAGGAAGAAGAGGAAGAGGAGGAA
 GAAGAGGAAGAGGAACTGGGTTCCAGGCCCTTGTCTCCAGAAGCAGGTTGCTGGCAACAGCATCAGGG
 AGAATGGAGCCAAGCCCCAGCCAGACCCCTTTTGTGAGACTGACAGCGACGAGGAGATCTTGGAGCAGAT
 ACTGGAGTTGCCTCTCCAGCGGCTCTGCAGCAAGAAGCTGTTGAGCATCCCTGAGGAGGAAGAGGAGGAG
 GAAGAGGAGGAAGGGCTGGAGAAACCAGGGCCAGCCGCACTTCCAAGACCCTAGCCAGCCTGAACTTG
 CGTTGCTAGGGCCGGGCTGTGATAGCAGTACAGCCAGGGACCTGGCCTGTGCTCCCTGTCTCCTGAGCT
 CTCTGGGGTCAGGGAGCACCTGGAGGATGTGCTGGGAGTCGTTGGTGGAAACGGCAGGAGGAGAGGAGGT
 GGCTCCCCGAGAAACTCCAAACCAGCAAGCGACCTCAGGACCCCGAGAACATTGCAGCCGGCTTCTTG
 GCAATGGCGGGCCCCAGGCCTCTGCACGGCCGGTCCCTCCAGGGAGAGGGGAGCCTCCCTGTGATTGA
 GGGCACCAGGGTTGGACAGGAGCCCGTGGGAGAGGGCGCCGGGTCTTCCCGAGGTGTCCCCGTGGC
 CCTGCTCCAGAATCCAGCTTAGTCAGCTGCCTCTCTCAAAGTGTGGAGATCAGTATTGAATATGATT
 CTGAGGATGAGCAGGAGGCGGCAGCGGGGTGTGAGCATCAACAGCTCCTGCTACCCACAGATGGGGA
 GGCCTGGGGCACAGCGGCAGTAGGAAGGCCAGGGGACCTCCGAAGGTCAATCCAGGCCCAACGCCTAC
 CTGCGCCTCCCAGCCTGGGAGAAAGGGGAGCCAGAGCGGAGAGGCCGAGTGCATTGGCAGAACCAAGG
 AGCCACCTCCCCGGCAACAGAAACTGGGGAGTCCAGAGGGCAGGACAACCTTGGGCGGAGAGGACCCCA
 GAGGAGAGGGGCCGGGTGCCTAGGTCTGGTACCACTGAGCTGGCCCTCCAAGGAGCCCCAAGAAGCA
 CCACCTCATCAGGACCTGCCTGTGAGGGTCTTTGTGGCTCTGTTGACTATGACCCTGTATCAATGTCAC
 CGAACCTGATGCCGGAGAAGAGGAACTGCCCTTAAAGGAGGGCCAACCTCCTCAAGGTGTTTGGAGACAA
 GGACGCTGATGGTTTCTACCGGGGTGAGAGTGGGGCCGTACAGGCTACATCCCCGCAACATGGTGGCT
 GAGGTGGCTGTGGACAGTCCAGCAGGGAGACAACAGCTGCTCCAGCGGGTCTTGGCCCCAAATGTTT

TCACCGAGGCCTCAGGAAATGGTCCCTCTGTGTACTCCTCAGCACACACACCCGGGCCTCCCCCAAGCC
 TCGTCGGTCCAAGAAAGTGGAGCTGGAAGGTCCTACACAGCTCTGTCCAGGTCCTCTAAGCTGATTCAT
 TCTGCTGCCAGAAAACCTCCCGACCTATGGTGGCTGCATTTGACTATAATCCTCGGGAGAACTCCCCCA
 ATATGGATGTGGAGGCAGAGCTGCCCTTACAGAGCAGGGGATGTCATTACTGTGTTTGGGAACATGGACGA
 TGATGGTTTCTACTATGGGGAGCTGAATGGACAAAGGGGCCTGGTTCCATCCAACCTCTGGAAGGCCT
 GGGCCTGAGTCAGGCAGCCTAGAGTCTGGGACATCTCAAGCCGAGAGTCAGAGAACGAGGAGGAGAAGAG
 TCCAGTGC

ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCTGGATT
 ACAAGGATGACGACGATAAGGTTAA

Protein Sequence:

>MR216449 representing NM_172449

Red=Cloning site Green=Tags(s)

MEQLTTLPRGLDGAMEPWALPAWQHWTQGGCKPGDASPSIAGTPTALQVKGLRFEESKPEGASHPGP
 VGNTDPEATETGLPKLGQQAESPYSGLSEEEEAQAYKAKFNIGFDRPNLELLRALGELQQRCTILKE
 ENQMLRKSFPETEELKVRRLKRKNAELAVIAKRLAQAQKLEETNMRVVSAPVPRPGSSLELCKRALARQ
 RARDLSETASALLAKDKQIAALQRECRELQARLSLVGKEGPQWLHMRDFDRLRESQREVLRLQRQIALR
 NQREPLRPARSPGPTAPSRVGAPAPGAPGEAVLQDDVESPQVVLREPEKQQRVQLESELCKRKKCESL
 EQEARKKQRRCELELQLRAAQNENARLVEENSRLSGRATEKEQVEWENSELKGQLLGVTVQERDSALLKS
 QQLQSKLESLEQLKHMREVAQRRQQLVEHEQARLSLQEKQEEVRLQQAQAEAKREHEGAVQLLESTL
 DSMQARVRELEGGQCSQTERFSLLAQELQAFRLHPGPLDLLTSALGCSALGDHPPPHCCCSIPQPCQSGS
 PKDLDLPPGSPGRCTPKSSEPALTTLTGIPRRTAKKAESLSNSSRSESIHNSPKSCTPEVDTASEVEEL
 EVDSVSLPAAPESHSGGARIQVFLARYSNPFEGPNENPEAELPLTAGEYIYIYGNMDEDFEGELMD
 GRRGLVPSNFVERVSDDLLSTLPRELADSSHSPELSFLSGGGGGSSGGQSSGGRSQRPEEEAAGD
 ELSLSPPEGLGEPLAVPYPRHITVLKQLAHSVVLAWELPPERVDLRGFHIFVNGELRQALGPGVPPKAV
 LENMDLRTGPLHVSQALTSKGSDDLRCCLAVGAGAGVPSQLRIHRLTATSAEIAWVPGNSNLAHAIY
 LNGEPCPPARPSTYWATFCNLRPGTLYQARVEAQIPSGPWEPEWERPEQRAATLQFTTLPAGLPDAPLD
 VQAEPGSPGILMISWLPVTIDAAGTNGVVRVTGYAIYADGQKIMEVASPTAGSVLVEVSQLQLLQACHE
 QDFASLSIEMSKGPQEEPPVPCSQEEAGAAVRSISEEKRAIEPTLGQEGPEPVAPSLAKQEVECTSGDA
 GPVPCSTQGELTQKPSIEACHGGDLDSGLKLRSEKEDMSELGVHLVNSLVDHSRNSDLSDIQEEEEEE
 EEEEELGSRPCSSQKQVAGNSIRENGAKPQDPDFCETDSDEEILEQILELPLQLRCSKCLFSIPEEEEE
 EEEGLEKPGPSRTSQDPSQPELALLGPGCDSSQPQGPGLCPLSPELSGVREHLEDVLGVVGGNGRRRG
 GSPEKLPNRKRPQDPREHCSRLGNGGPQASARPVPPRERGSPLVIEGTRVQEPGGRGRPGLSRRCPRG
 PAPESSLVSLSPKCLEISIEYDSEDEQEAAGSGVSNSSCYPTDGEAWGTAAVGRPRGPPKVNPGPNAY
 LRLPAWEKGEPPERGRSAIGRTKEPPSRATETGESRGQDNSGRRGPQRRGARVPRSGTTELAPRSPQEA
 PPHQDLVVRVVALFDYDPVSMSPNDAGEEELPFKEGQLLKVFQDKDADGFYRGESEGGRTGYIPCNMVA
 EVAVDSPAGRQQLLQRGFLPPNVLTEASNGGPSYSSAHTPGPPKPRRSKKVLEEGPTQLCPGPPKLIH
 SAAQKTSRPMVAADFYNPRENSPMDVEAELPFRAAGDVITVFGNMDDDGFYYGELNGQRGLVPSNLFLEGP
 GPESGLESQESQRTRRRRVQC

TRTRPLEQKLISEEDLAANDILDYKDDDDKV

Chromatograms:

https://cdn.origene.com/chromatograms/jc1403_a01.zip

Restriction Sites:

Sgfl-MluI

RefSeq Size: 7633 bp
RefSeq ORF: 5541 bp
Locus ID: 207777
UniProt ID: [Q7TNF8](#)
Cytogenetics: 11 C
MW: 200.4 kDa