**2B chimera PcrA**

**Plasmid**

pET22B\_PcrA\_2Bchimera

**Protein**

2B chimera PcrA: *B. stearothermophilus* sequence with *S. aureus* 2B domain

**Depositor**

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**Protein sequence**

MNFLSEQLLA HLNKEQQEAV RTTEGPLLIM AGAGSGKTRV LTHRIAYLMA EKHVAPWNIL

AITFTNKAAR EMRERVQSLL GGAAEDVWIS TFHSMCVRIL RRDIDRIGIN RNFSILDPTD

QLSVMKTILK EKNIDPKKFE PRTILGTISA AKNELLPPEQ FAKRASTYYE KVVSDVYQEY

QQRLLRNHSL DFDDLIMTTI QLFDRVPDVL HYYQYKFQYI HIDEYQDTNR AQYTLVKKLA

ERFQNICAVG DADQSIYRWR GADIQNILSF ERDYPNAKVI LLEQNYRSTK RILQAANEVI

EHNVNRKPKR LWTENPEGKP ILYYEAMNEA DEAQFVAGRI REAVERGERR YRDFAVLYRT

NAQSRVMEEM LLKANIPYQI VGGQKFYDRK EIKDLLSYLR IIANSNDDIS LQRIINVPKR

GVGPSSVEKV QNYALQNNIS MFDALGEADF IGLSKKVTQE CLNFYELIQS LIKEQEFLEI

HEIVDEVLQK SGYREMLERE NTLESRSRLE NIDEFMSVPK DYEENTPLEE QSLINFLTDL

SLVADIDEAN GTEQAAEGDA VMLMTLHAAK GLEFPVVFLI GMEEGIFPHN RSLEDDDEME

EERRLAYVGI TRAEEELVLT SAQMRTLFGN IQMNPPSRFL NEIPAHLLET ASRRQAGASR

PAVSRPQASG AVGSWKVGDR ANHRKWGIGT VVSVRGGGDD QELDIAFPSP IGIKRLLAKF

APIEKV

**Gene sequence**

GAAATAATTTTGTTTAACTTTAAGAAGGAGATATACATATGAACTTTCTGAGCGAACAGCTGCTGGCACATCTGAATAAAGAACAGCAAGAAGCAGTTCGTACCACCGAAGGTCCGCTGCTGATTATGGCAGGCGCAGGTAGCGGTAAAACCCGTGTTCTGACCCATCGTATTGCATATCTGATGGCAGAAAAACATGTTGCACCGTGGAATATTCTGGCAATTACCTTTACCAATAAAGCCGCACGTGAAATGCGTGAACGTGTTCAGAGCCTGCTGGGTGGTGCAGCAGAAGATGTTTGGATTAGCACCTTTCATAGCATGTGTGTTCGTATTCTGCGTCGTGATATTGATCGCATTGGTATTAATCGCAACTTTAGCATTCTGGACCCGACCGATCAGCTGAGCGTTATGAAAACCATTCTGAAAGAAAAAAACATCGATCCGAAAAAATTCGAGCCTCGTACCATTCTGGGCACCATTAGCGCAGCAAAAAATGAACTGCTGCCTCCGGAACAGTTTGCAAAACGTGCAAGCACCTATTATGAAAAAGTTGTGTCCGATGTGTACCAAGAATATCAGCAACGTCTGCTGCGTAATCATAGCCTGGATTTTGATGATCTGATTATGACCACCATCCAGCTGTTTGATCGTGTTCCGGATGTTCTGCATTATTATCAGTACAAATTCCAGTACATCCACATCGATGAATATCAGGATACCAATCGTGCACAGTATACCCTGGTTAAAAAACTGGCAGAACGCTTTCAGAACATTTGTGCAGTTGGTGATGCAGATCAGAGCATTTATCGTTGGCGTGGTGCAGATATTCAAAACATTCTGAGCTTTGAACGCGATTATCCGAATGCCAAAGTTATTCTGCTGGAACAGAATTATCGTAGCACCAAACGCATTCTGCAGGCAGCAAATGAAGTGATTGAACATAATGTGAACCGTAAACCGAAACGTCTGTGGACCGAAAATCCGGAAGGTAAACCGATTCTGTATTATGAAGCAATGAACGAAGCAGATGAAGCACAGTTTGTTGCAGGTCGTATTCGTGAAGCAGTTGAACGTGGTGAACGTCGTTATCGTGATTTTGCAGTTCTGTATCGTACCAATGCACAGAGCCGTGTTATGGAAGAAATGCTGCTGAAAGCAAATATCCCGTATCAGATTGTTGGTGGCCAGAAATTCTATGATCGCAAAGAAATTAAAGATCTGCTGAGCTATCTGCGCATTATTGCCAATAGCAATGATGATATTAGCCTGCAGCGTATTATCAATGTTCCGAAACGTGGTGTTGGTCCGAGCAGCGTTGAAAAAGTGCAGAATTATGCCCTGCAGAACAACATCAGCATGTTTGATGCACTGGGTGAAGCAGATTTCATTGGTCTGAGCAAAAAAGTTACCCAAGAATGCCTGAACTTCTATGAACTGATTCAGTCCCTGATCAAAGAACAAGAATTCCTGGAAATCCATGAGATCGTTGATGAAGTGCTGCAGAAAAGCGGTTATCGCGAAATGCTGGAACGTGAAAATACCCTGGAAAGCCGTAGCCGTCTGGAAAATATTGATGAATTTATGAGCGTGCCGAAAGATTATGAAGAAAATACACCGCTGGAAGAACAGAGCCTGATCAATTTTCTGACCGATCTGAGCCTGGTTGCAGATATCGATGAAGCAAATGGCACCGAACAGGCAGCCGAAGGTGATGCCGTTATGCTGATGACCCTGCATGCCGCAAAAGGTCTGGAATTTCCGGTTGTGTTTCTGATTGGCATGGAAGAAGGTATTTTTCCGCATAATCGTTCCCTGGAAGATGATGATGAAATGGAAGAGGAACGTCGTCTGGCCTATGTTGGTATTACCCGTGCAGAAGAAGAACTGGTTCTGACCAGCGCACAGATGCGTACCCTGTTTGGTAACATTCAGATGAATCCGCCTAGCCGTTTTCTGAATGAAATTCCTGCACATCTGCTGGAAACCGCAAGCCGTCGTCAGGCTGGTGCCAGCCGTCCGGCAGTTAGTCGTCCGCAGGCAAGCGGTGCCGTTGGTAGCTGGAAAGTGGGTGATCGTGCAAATCATCGTAAATGGGGTATTGGCACCGTTGTTAGCGTTCGTGGTGGCGGTGATGATCAAGAACTGGATATTGCATTTCCGAGCCCGATTGGCATTAAACGCCTGCTGGCAAAATTTGCACCGATCGAAAAAGTGTAAGGATCCGAATTCGAGCTCCGTCGACAAGCTTGCGGCCGCAC

**Gene source**

*B. stearothermophilus* , with the amino acids in positions between 381 and 547( 2B domain) substituted with *S. aureus* 2B domain amino acids.

Synthesised by GeneArt

**Plasmid**

pET22B

**Insert position:** PcrA 199-2384 insertion using Nde and BamH1

**Plasmid sequence**

**Pet22b vector sequence**

**T7 primer sequence (MWG)** TAA TAC GAC TCA CTA TAG GG

**Reverse compliment T7** CCC TAT AGT GAG TCG TAT TA

**T7 ter sequence** CTA GTT ATT GCT CAG CGG T

1 atccggatat agttcctcct ttcagcaaaa aacccctcaa gacccgttta gaggccccaa

61 ggggttatg**c tagttattgc** **tcagcggt**gg cagcagccaa ctcagcttcc tttcgggctt

121 tgttagcagc cggatctcag tggtggtggt ggtggtgctc gagtgcggcc gcaagcttgt

181 cgacggagct cgaattcgga tccgaattaa ttccgatatc catggccatc gccggctggg

241 cagcgaggag cagcagacca gcagcagcgg tcggcagcag gtatttcata tgtatatctc

301 cttcttaaag ttaaacaaaa ttatttctag aggggaattg ttatccgctc acaattc**ccc**

361 **tatagtgagt cgtatta**att tcgcgggatc gagatctcga tcctctacgc cggacgcatc

421 gtggccggca tcaccggcgc cacaggtgcg gttgctggcg cctatatcgc cgacatcacc

481 gatggggaag atcgggctcg ccacttcggg ctcatgagcg cttgtttcgg cgtgggtatg

541 gtggcaggcc ccgtggccgg gggactgttg ggcgccatct ccttgcatgc accattcctt

601 gcggcggcgg tgctcaacgg cctcaaccta ctactgggct gcttcctaat gcaggagtcg

661 cataagggag agcgtcgaga tcccggacac catcgaatgg cgcaaaacct ttcgcggtat

721 ggcatgatag cgcccggaag agagtcaatt cagggtggtg aatgtgaaac cagtaacgtt

781 atacgatgtc gcagagtatg ccggtgtctc ttatcagacc gtttcccgcg tggtgaacca

841 ggccagccac gtttctgcga aaacgcggga aaaagtggaa gcggcgatgg cggagctgaa

901 ttacattccc aaccgcgtgg cacaacaact ggcgggcaaa cagtcgttgc tgattggcgt

961 tgccacctcc agtctggccc tgcacgcgcc gtcgcaaatt gtcgcggcga ttaaatctcg

1021 cgccgatcaa ctgggtgcca gcgtggtggt gtcgatggta gaacgaagcg gcgtcgaagc

1081 ctgtaaagcg gcggtgcaca atcttctcgc gcaacgcgtc agtgggctga tcattaacta

1141 tccgctggat gaccaggatg ccattgctgt ggaagctgcc tgcactaatg ttccggcgtt

1201 atttcttgat gtctctgacc agacacccat caacagtatt attttctccc atgaagacgg

1261 tacgcgactg ggcgtggagc atctggtcgc attgggtcac cagcaaatcg cgctgttagc

1321 gggcccatta agttctgtct cggcgcgtct gcgtctggct ggctggcata aatatctcac

1381 tcgcaatcaa attcagccga tagcggaacg ggaaggcgac tggagtgcca tgtccggttt

1441 tcaacaaacc atgcaaatgc tgaatgaggg catcgttccc actgcgatgc tggttgccaa

1501 cgatcagatg gcgctgggcg caatgcgcgc cattaccgag tccgggctgc gcgttggtgc

1561 ggatatctcg gtagtgggat acgacgatac cgaagacagc tcatgttata tcccgccgtt

1621 aaccaccatc aaacaggatt ttcgcctgct ggggcaaacc agcgtggacc gcttgctgca

1681 actctctcag ggccaggcgg tgaagggcaa tcagctgttg cccgtctcac tggtgaaaag

1741 aaaaaccacc ctggcgccca atacgcaaac cgcctctccc cgcgcgttgg ccgattcatt

1801 aatgcagctg gcacgacagg tttcccgact ggaaagcggg cagtgagcgc aacgcaatta

1861 atgtaagtta gctcactcat taggcaccgg gatctcgacc gatgcccttg agagccttca

1921 acccagtcag ctccttccgg tgggcgcggg gcatgactat cgtcgccgca cttatgactg

1981 tcttctttat catgcaactc gtaggacagg tgccggcagc gctctgggtc attttcggcg

2041 aggaccgctt tcgctggagc gcgacgatga tcggcctgtc gcttgcggta ttcggaatct

2101 tgcacgccct cgctcaagcc ttcgtcactg gtcccgccac caaacgtttc ggcgagaagc

2161 aggccattat cgccggcatg gcggccccac gggtgcgcat gatcgtgctc ctgtcgttga

2221 ggacccggct aggctggcgg ggttgcctta ctggttagca gaatgaatca ccgatacgcg

2281 agcgaacgtg aagcgactgc tgctgcaaaa cgtctgcgac ctgagcaaca acatgaatgg

2341 tcttcggttt ccgtgtttcg taaagtctgg aaacgcggaa gtcagcgccc tgcaccatta

2401 tgttccggat ctgcatcgca ggatgctgct ggctaccctg tggaacacct acatctgtat

2461 taacgaagcg ctggcattga ccctgagtga tttttctctg gtcccgccgc atccataccg

2521 ccagttgttt accctcacaa cgttccagta accgggcatg ttcatcatca gtaacccgta

2581 tcgtgagcat cctctctcgt ttcatcggta tcattacccc catgaacaga aatccccctt

2641 acacggaggc atcagtgacc aaacaggaaa aaaccgccct taacatggcc cgctttatca

2701 gaagccagac attaacgctt ctggagaaac tcaacgagct ggacgcggat gaacaggcag

2761 acatctgtga atcgcttcac gaccacgctg atgagcttta ccgcagctgc ctcgcgcgtt

2821 tcggtgatga cggtgaaaac ctctgacaca tgcagctccc ggagacggtc acagcttgtc

2881 tgtaagcgga tgccgggagc agacaagccc gtcagggcgc gtcagcgggt gttggcgggt

2941 gtcggggcgc agccatgacc cagtcacgta gcgatagcgg agtgtatact ggcttaacta

3001 tgcggcatca gagcagattg tactgagagt gcaccatata tgcggtgtga aataccgcac

3061 agatgcgtaa ggagaaaata ccgcatcagg cgctcttccg cttcctcgct cactgactcg

3121 ctgcgctcgg tcgttcggct gcggcgagcg gtatcagctc actcaaaggc ggtaatacgg

3181 ttatccacag aatcagggga taacgcagga aagaacatgt gagcaaaagg ccagcaaaag

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3301 gagcatcaca aaaatcgacg ctcaagtcag aggtggcgaa acccgacagg actataaaga

3361 taccaggcgt ttccccctgg aagctccctc gtgcgctctc ctgttccgac cctgccgctt

3421 accggatacc tgtccgcctt tctcccttcg ggaagcgtgg cgctttctca tagctcacgc

3481 tgtaggtatc tcagttcggt gtaggtcgtt cgctccaagc tgggctgtgt gcacgaaccc

3541 cccgttcagc ccgaccgctg cgccttatcc ggtaactatc gtcttgagtc caacccggta

3601 agacacgact tatcgccact ggcagcagcc actggtaaca ggattagcag agcgaggtat

3661 gtaggcggtg ctacagagtt cttgaagtgg tggcctaact acggctacac tagaaggaca

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4021 acttggtctg acagttacca atgcttaatc agtgaggcac ctatctcagc gatctgtcta

4081 tttcgttcat ccatagttgc ctgactcccc gtcgtgtaga taactacgat acgggagggc

4141 ttaccatctg gccccagtgc tgcaatgata ccgcgagacc cacgctcacc ggctccagat

4201 ttatcagcaa taaaccagcc agccggaagg gccgagcgca gaagtggtcc tgcaacttta

4261 tccgcctcca tccagtctat taattgttgc cgggaagcta gagtaagtag ttcgccagtt

4321 aatagtttgc gcaacgttgt tgccattgct gcaggcatcg tggtgtcacg ctcgtcgttt

4381 ggtatggctt cattcagctc cggttcccaa cgatcaaggc gagttacatg atcccccatg

4441 ttgtgcaaaa aagcggttag ctccttcggt cctccgatcg ttgtcagaag taagttggcc

4501 gcagtgttat cactcatggt tatggcagca ctgcataatt ctcttactgt catgccatcc

4561 gtaagatgct tttctgtgac tggtgagtac tcaaccaagt cattctgaga atagtgtatg

4621 cggcgaccga gttgctcttg cccggcgtca atacgggata ataccgcgcc acatagcaga

4681 actttaaaag tgctcatcat tggaaaacgt tcttcggggc gaaaactctc aaggatctta

4741 ccgctgttga gatccagttc gatgtaaccc actcgtgcac ccaactgatc ttcagcatct

4801 tttactttca ccagcgtttc tgggtgagca aaaacaggaa ggcaaaatgc cgcaaaaaag

4861 ggaataaggg cgacacggaa atgttgaata ctcatactct tcctttttca atattattga

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4981 aaacaaatag gggttccgcg cacatttccc cgaaaagtgc cacctgaaat tgtaaacgtt

5041 aatattttgt taaaattcgc gttaaatttt tgttaaatca gctcattttt taaccaatag

5101 gccgaaatcg gcaaaatccc ttataaatca aaagaataga ccgagatagg gttgagtgtt

5161 gttccagttt ggaacaagag tccactatta aagaacgtgg actccaacgt caaagggcga

5221 aaaaccgtct atcagggcga tggcccacta cgtgaaccat caccctaatc aagttttttg

5281 gggtcgaggt gccgtaaagc actaaatcgg aaccctaaag ggagcccccg atttagagct

5341 tgacggggaa agccggcgaa cgtggcgaga aaggaaggga agaaagcgaa aggagcgggc

5401 gctagggcgc tggcaagtgt agcggtcacg ctgcgcgtaa ccaccacacc cgccgcgctt

5461 aatgcgccgc tacagggcgc gtcccattcg cca

**Publication**

To be published

**Protocol for prep of protein**

1. The transformed B834 (DE3) cells, containing the pET22b-2BPcrA, are used to inoculate a starter culture, of 100 ml of Luria-Bertani broth (LB), with ampicillin (100 µg/ml). These were grown overnight, at 37 °C, on a shaker, at 225 rpm. The protein expression used 8 X 500 ml cultures. A 1/100 dilution of the overnight culture was made up in each 500 ml of LB, with ampicillin, in 2 L flasks. The cells were grown until they reached an OD595 0.5. PcrA expression was induced with 1 mM IPTG, and grown for 5 h, at 25 ° C, shaking at 225 rpm. The culture was transferred into 4 x L centrifuge bottles, after incubation, and cells were centrifuged, at 4 °C, at 4000 rpm, in JS 4.2 swing bucket rotor (Beckman Coulter), for 30 min. The culture pellets were resuspended, into 20 ml of PcrA resuspension buffer containing; 50 mM Tris·HCl (pH 7.5), 2 mM, EDTA, 1 mM DTT, 200 mM NaCl, and 10 % (w/v) sucrose. Then the cells were stored at -80 °C.
2. The resuspended cells were thawed and a dissolved protease inhibitor tablet (Complete, Santa Cruz biotechnology) added. The cells were lysed, using a probe sonicator, with burst of 4 x 20 s (80 Joules), on ice, then the cells were spun, at 13200 rpm, at 4 °C, for 20 min, using 45 Ti rotor (Beckman Coulter). The pellet was discarded. Saturated (NH4)2SO4, at a volume of 0.7 times that of the supernatant, was added gradually to the supernatant form centrifugation, with stirring at room temperature, to precipitate the PcrA. This was centrifuged at 13 200 rpm, at 4 °C, for 20 min, this time discarding the supernatant, the pellet was resuspended into 20 ml, of PcrA low salt pellet buffer containing; 50 mM Tris·HCl (pH 7.5), 2 mM, EDTA, 1 mM DTT and 100 mM NaCl. Its conductivity was adjusted to match the re-suspension buffer, by addition of PcrA adjustment buffer containing; 50 mM Tris·HCl (pH 7.5), 2 mM EDTA, 1 mM DTT.
3. The conductivity adjusted sample was loaded onto a 20 ml heparin sepharose column (GE healthcare), equilibrated in PcrA column buffer; 50 mM Tris·HCl (pH 7.5), 2 mM EDTA, 1 mM DTT, and 100 mM NaCl, using an AKTA FPLC (GE healthcare), at 4 ˚C, with a flow rate of 1 ml/min. The PcrA was eluted, with a NaCl gradient, from 100 mM NaCl to 700 mM NaCl, over 150 ml. PcrA eluted at approximately, 300-400 mM NaCl, across four fractions. The purity of the protein was confirmed by SDS-PAGE gel and those fractions with just a single protein band were pooled and concentrated, using a 20 ml, 10000 MWCO, Vivaspin concentrator (Millipore), by centrifugation, at 4000 RPM, at 4 ˚C, in a JS 4.2 swing bucket rotor. The absorbance ratio of 260 nm:280 nm showed a large amount of DNA present in the final pool of protein, therefore the protein was then gel filtered to remove the DNA.
4. A Superdex 200 column was washed overnight, in a gel filtration buffer containing; 500 mM NaCl, with 1 mM NaN3, at a flow rate of 0.2 ml/min, for 12 hours. A 2 ml aliquot of the pooled protein was loaded onto the gel filtration column; the column was then washed, with 40 ml, of buffer at a flow rate of 0.5 ml/min. The sample was eluted at a flow of 0.5 ml/min, with 2 ml fractions being collected over 120 ml. The DNA eluted first, at 45 – 64 ml, the protein eluted at 75 – 84 ml, this was confirmed on agarose and SDS-PAGE gels, the 260:280 nm spectrometry showed that there was little DNA contamination remaining in the sample. The molar extinction coefficient of 71865 M-1cm-1 at 280 nm (from amino acid sequence), was used to calculate the concentration of the PcrA protein. PcrA was stored in aliquots at -80 ˚C.