

## T7 RNA Polymerase Nucleotide Sequence

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ATGAACACGA  TTAACATCGC  TAAGAACGAC  TTCTCTGACA  TCGAACTGGC  TGCTATCCCG
TTCAACACTC  TGGCTGACCA  TTACGGTGAG  CGTTTAGCTC  GCGAACAGTT  GGCCCTTGAG
CATGAGTCTT  ACGAGATGGG  TGAAGCACGC  TTCCGCAAGA  TGTTTGAGCG  TCAACTTAAA
CTGGTGAGG  TTGCGGATAA  CGCTGCCGCC  AAGCCTCTCA  TCACTACCCCT  ACTCCCTAAG
ATGATTGCAC  GCATCAACGA  CTGGTTTGAG  GAAGTGAAG  CTAAGCGCGG  CAAGCGCCCG
ACAGCCTTCC  AGTTCCTGCA  AGAAATCAAG  CCGGAAGCCG  TAGCGTACAT  CACCATTAAG
ACCACTCTGG  CTTGCCTAAC  CAGTGCTGAC  AATACAACCG  TTCAGGCTGT  AGCAAGCGCA
ATCGGTCGGG  CCATTGAGGA  CGAGGCTCGC  TTCGGTCGTA  TCCGTGACCT  TGAAGCTAAG
CACTTCAAGA  AAAACGTTGA  GGAACAATC  AACAAGCGCG  TAGGGCACGT  CTACAAGAAA
GCATTTATGC  AAGTTGTCTA  GGCTGACATG  CTCTCTAAGG  GTCTACTCGG  TGGCGAGGCG
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TCTGAGACTA  TCGAACTCGC  ACCTGAATAC  GCTGAGGCTA  TCGCAACCCG  TGCAGGTGCG
CTGGCTGGCA  TCTCTCCGAT  GTTCCAACCT  TGCGTAGTTC  CTCCTAAGCC  GTGGACTGGC
ATTACTGGTG  GTGGCTATTG  GGCTAACGGT  CGTCTCCTC  TGGCGCTGGT  GCGTACTCAC
AGTAAGAAAG  CACTGATCGG  ACTAGAAGAC  GCTTACATGC  CTGAGGTGTA  CAAAGCGATT
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CCGATGAAAC  CGGAAGACAT  CGACATGAAT  CCTGAGGCTC  TCACCGCGTG  GAAACGTGCT
GCCGCTGCTG  TGTACCGCAA  GGACAGGCT  CGCAAGTCTC  GCCGTATCAG  CCTTGAGTTC
ATGCTTGAGC  AAGCCAATAA  GTTTGCTAAC  CATAAGGCCA  TCTGGTTCCC  TTACAACATG
GACTGGCGCG  GTCGTGTTTA  CGCTGTGTC  CGTGTCAACC  CGCAAGGTAA  CGATATGACC
AAAGGACTGC  TTACGCTGGC  GAAAGTAAA  CCAATCGGTA  AGGAAGGTTA  CTACTGGCTG
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AACTTTGTAC  ACAGCCAAGA  CCGTAGCCAC  CTTTCGTAAGA  CTGTAGTGTG  GGCACACGAG
AAGTACGGAA  TCGAATCTTT  TGCCTGATT  CACGACTCCT  TCGGTACCAT  TCCGGCTGAC
GCTGCGAAC  TGTTCAAAGC  AGTGC GCGAA  ACTATGGTTG  ACACATATGA  GTCTTGATGAT
GTACTGGCTG  ATTTCTACGA  CCAGTTCGCT  GACCAGTTGC  ACGAGTCTCA  ATTTGGACAAA
ATGCCAGCAC  TTCCGGCTAA  AGGTAAC'TTG  AACCTCCGTC  ACATCTTAGA  GTCGGACTTC
CGTTCCGCGT  AA

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A to G results in amino acid change Asn823 to Asp. Amino acid 823 is on surface of protein and is not found in any mutagenesis studies of T7 RNAP

A to G

## Protein Sequence 883 AA 98855 MW (translated from above)

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MNTINIAKND  FSDIELAAIP  FNTLADHYGE  RLAREQLALE  HESYEMGEAR  FRKMFERQLK
AGEVADNAAA  KPLITLLPK  MIARINDWFE  EVKAKRGKRP  TAFQFLQEIK  PEAVAYITIK
TTLACLTSAD  NTTVQAVASA  IGRAIEDEAR  FGRIRDLEAK  HFKKNVEEQL  NKRVGHVYKK
AFMQVVEADM  LSKGLLGGEA  WSSWHKEDSI  HVGVRCIEM  IESTGMVSLH  RQNAGVVGQD
SETIELAPEY  AEAIATRAGA  LAGISPMFQP  CVVPPKPWTG  ITGGGYWANG  RRPLALVRTH
SKKALMRYED  VYMPEVYKAI  NIAQNTAWKI  NKKVLAVANV  ITKWKHCPVE  DIPAIEREEL
PMKPEDIDMN  PEALTAWKRA  AAAYVRKDKA  RKSRRISLEF  MLEQANKFAN  HKAIWFPYNM
DWRGRVYAVS  MFNPQGNMT  KGLLTLAKGK  PIGKEGYWL  KIHGANCAV  DKVPPPERIK
FIEENHENIM  ACAKSPLENT  WWAEQDSPFC  FLAFCFEYAG  VQHHGLSYNC  SLPLAFDGSC
SGIQHFSAML  RDEVGGRAVN  LLPSETVQDI  YGIVAKKVN  ILQADAINGT  DNEVTVTDE
NTGELISEKVK  LGTKALAGQW  LAYGVTRSVT  KRSVMTLAYG  SKEFGFRQOV  LEDTIQPAID
SGKGLMFTQP  NQAAGYMAKL  IWESVSVTVV  AAVEAMNWLK  SAAKLLAAEV  KDKKTGEILR
KRCVHVWVTP  DGFVWQEQYK  KPIQTRLNLM  FLGQFRLQPT  INTNKDSEID  AHKQESGIAP
NFVHSQDGSH  LRKTVVWAHE  KYGIESFALI  HDSFGTIPAD  AANLFKAVRE  TMVDTYESCD
VLADFYDQFA  DQLHESQLDK  MPALPAKGNL  NLRDILESDF  AFA

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T7 RNA Polymerase with flanking sequences

ctttgctatgc catagcattt ttatccataa gattagcgga tcctacctga cgctttttat cgcaactctc  
 tactgtttctc catacccggt tttttgggct agcgaattcg agctcAAGAG GATACCAT...

Sac I site

ATGAACACGA	TTAACATCGC	TAAGAACGAC	TTCTCTGACA	TCGAACTGGC	TGCTATCCCG
TTCAAACTC	TGGCTGACCA	TTACGGTGAG	CGTTTAGCTC	GCGAACAGTT	GGCCCTTGAG
CATGAGTCTT	ACGAGATGGG	TGAAGCACGC	TTCCGCAAGA	TGTTTGAGCG	TCAACTTAAA
GCTGGTGAGG	TTGCGGATAA	CGCTGCCGCC	AAGCCTCTCA	TCACTACCCCT	ACTCCCTAAG
ATGATTGAC	GCATCAACGA	CTGGTTTGAG	GAAGTGAAAG	CTAAGCGCGG	CAAGCGCCCG
ACAGCCTTCC	AGTTCCTGCA	AGAAATCAAG	CCGGAAGCCG	TAGCGTACAT	CACCATTAAG
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ATCGGTCCGG	CCATTGAGGA	CGAGGCTCGC	TTCGGTTCGTA	TCCGTGACCT	TGAAGCTAAG
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ATCACC AAGT	GGAAGCATTG	TCCGGTCGAG	GACATCCCTG	CGATTGAGCG	TGAAGA ACTC
CCGATGAAAC	CGGAAGACAT	CGACATGAAT	CCTGAGGCTC	TCACCGCGTG	GAAACGTGCT
GCCCGTGCTG	TGTACCGCAA	GGACAAGGCT	CGCAAGTCTC	GCCGTATCAG	CCTTGAGTTC
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GACTGGCGCG	GTCGTGTTTA	CGCTGTGTCA	ATGTTCAACC	CGCAAGGTAA	CGATATGACC
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AAGCCTATTC	AGACGCGCTT	GAACCTGATG	TTCCTCGGTC	AGTTCGGCTT	ACAGCCTACC
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AAC'TTTGTAC	ACAGCCAAGA	CGGTAGCCAC	CTTCGTAAGA	CTGTAGTGTG	GGCACACGAG
AAGTACGGAA	TCGAATCTTT	TGCACTGATT	CACGACTCCT	TCGGTACCAT	TCCGGCTGAC
GCTGCGAACC	TGTTCAAAGC	AGTGCGCGAA	ACTATGGTTG	ACACATATGA	GTCTTGTGAT
GTACTGGCTG	ATTTCTACGA	CCAGTTCGCT	GACCAGTTGC	ACGAGTCTCA	ATTGGACAAA
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GCGTTCGCGT	AA...				

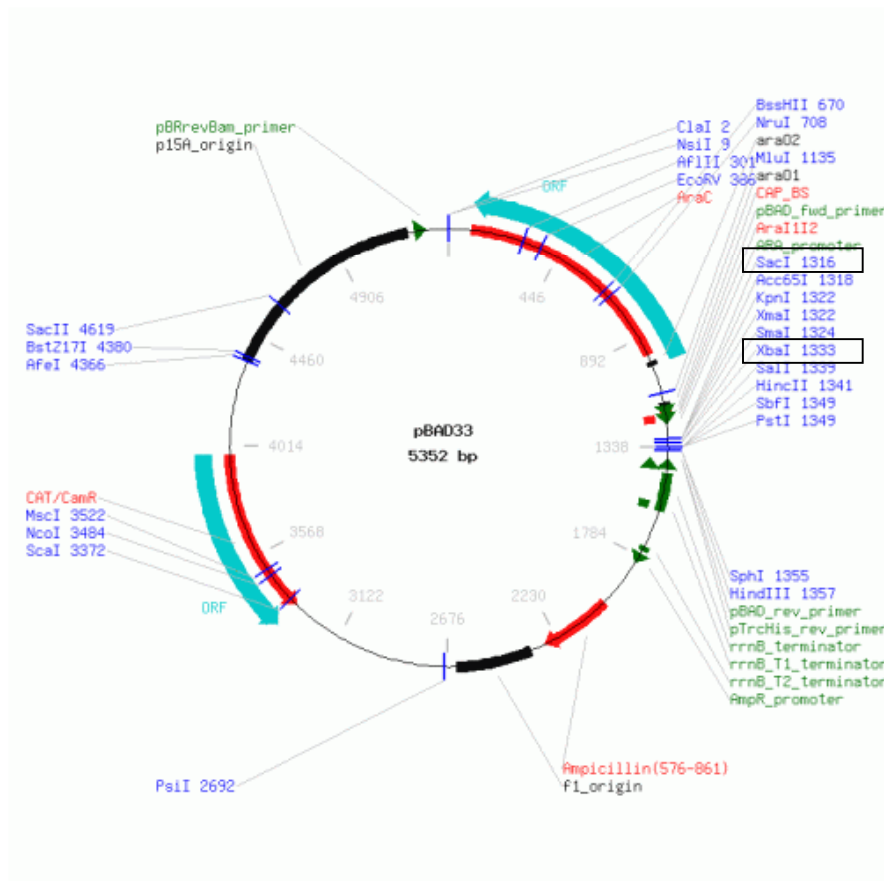
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 XbaI site

## pBAD

Sequence can be found at

<https://www.lablife.org/g?a=seqa&id=vdb%5fg2%2ejPxWBK9aksPdtGbxXRrF06TxA%2e4%2d%5fsequence%5f549a5591ef126b8ee61fb291e3498cd6c3147325%5f10>

Plasmid Name	pBAD33
Source/Vendor	Beckwith Lab
Plasmid Type	Bacterial Expression
Plasmid Size	5352
Sequencing Primer	5' ctgtttctccataccggtt 3' ctcatccgcaaaacag
Bacterial Resistance	Chloramphenicol
Notes	Vector backbone: pACYC-184 Cloning site 5': SacI Cloning site 3': HindIII Article: Tight regulation, modulation, and high-level expression by vectors containing the arabinose PBAD promoter. Guzman LM et al. (J Bacteriol. 1995 Jul . 177(14):4121-30. Pubmed)
Plasmid Sequence	<a href="#">View Sequence</a>



T7RNAP sequence was amplified with PCR from pAR1217 with SacI and XbaI sites at the ends and cloned into pPAD33 to create pTARA, with T7 RNA polymerase under the control of AraPBAD promoter and therefore AraC/arabinose control.

# Schematic for pTARA construction

