

Gene Synthesis QA Report

Gene Name	CYP78C5 cs	Order ID	SG140999-1
Lot#	GB151124-35412	Cloning Vector	pBluescript II SK(+)
Length (bp)	1668	Cloning Sites	EcoR I/BamHI

Quality Control

Test Items	Specifications	Results
Sequencing Alignment	Sequencing data consistent with target	[X] Pass
Vector Sequence	Flanking sequence of cloning sites are correct	[X] Pass
Restriction Digest	Insert size is correct and no contaminated bands	[X] Pass
ORF Across Junction	Correct and consistent with target	N/A
PCR Amplification	Correct and no contaminated bands	[X] Pass
Endotoxin Level	Verified, <0.1 EU/μg (Endo-Free Preps Only)	N/A
Appearance	Clear, no foreign particles	[X] Pass
DNA Purity	Purity (A 260/A280 = 1.8 - 2.0)	[X] Pass
DNA Quantity	Actual yield (by A 260)	5ug

Comments

Delivery form: lyophilized plasmid (TE lyophilized) containing the gene insert. It is stable at room temperature for extended period of time during shipping. The lyophilized plasmid can be dissolved in sterile TE buffer or nuclease-free water (neutral pH) depending on the established laboratory practice. After reconstitution, store the stock solution at -20°C or -80°C for long term storage. The lyophilized plasmid dissolved in TE buffer is stable for at least 6 months at 4°C while the lyophilized DNA dissolved in water is NOT STABLE at 4°C.

- Before opening the tube containing the plasmid, please briefly centrifuge the tube. Lyophilized plasmid could attach to the wall of the tube. Opening without centrifugation could cause DNA loss.
- Stock Solution: Reconstitute lyophilized plasmid (4ug or 10ug) in 40ul or 100ul of TE buffer or nuclease-free water (final concentration - 100ng/ul). To accurately determine the quantity of DNA present, please measure OD value of original stock at OD_{260nm} after reconstitution.
- Working Solution: make a 1:10 dilution of stock solution using TE buffer or nuclease-free water (final concentration - 10ng/ul)

Transformation and Replating:

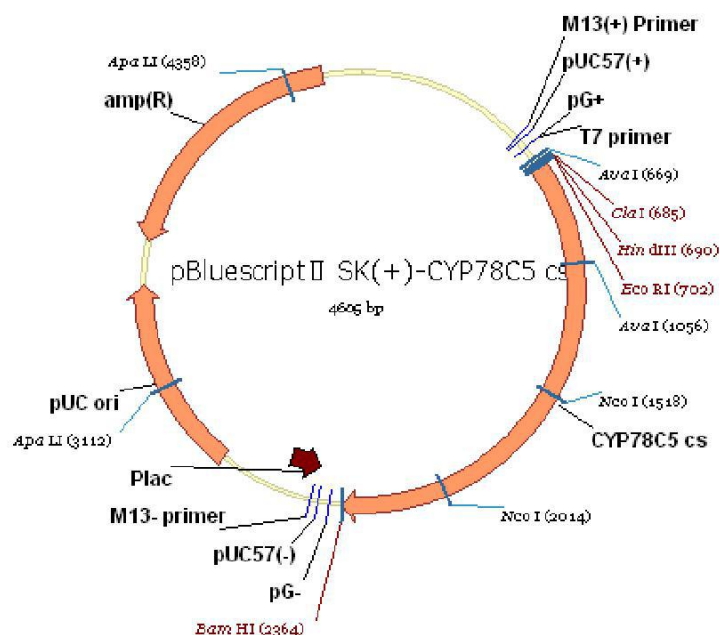
Transform 2ul of stock solution into appropriate *E. coli* competent cells according to standard laboratory protocol. Plate the mixture on LB agar (with desired antibiotic selection) and incubate at 37°C for overnight. Select a well separated, SINGLE colony and inoculate in LB medium with desired antibiotic selection for overnight culture. Purify Plasmid DNA from overnight cultures, verify sequences and continue with project of interest. *It is important to select only a SINGLE colony for overnight culture.

Restriction Digestion



Gene name: CYP78C5 cs
 Clone ID#: V9081-2
 RES: XbaI/HindIII

Construct Map:



Detailed Sequence of the Whole Construct:

1	CTAAATTGTA	AGCGTTAATA	TTTTGTAAAA	ATTCGCGTTA	AATTTTTGTT	AAATCAGCTC
61	ATTTTTTAAAC	CAATAGGCCG	AAATCGGCAA	AATCCCTTAT	AAATCAAAAG	AATAGACCGA
121	GATAGGGTTG	AGTGTGTGTC	CAGTTTGGAA	CAAGAGTCCA	CTATTAAAGA	ACGTGGACTC
181	CAACGTCAAA	GGGCGAAAAA	CCGTCTATCA	GGGCGATGGC	CCACTACGTG	AACCATCACC
241	CTAATCAAGT	TTTTTGGGGT	CGAGGTGCCG	TAAAGCACTA	AATCGGAACC	CTAAAGGGAG
301	CCCCCGATTT	AGAGCTTGAC	GGGGAAAGCC	GGCGAACGTG	GCGAGAAAGG	AAGGGAAGAA
361	AGCGAAAGGA	GCGGGCGCTA	GGGCGCTGGC	AAGTGTAGCG	GTCACGCTGC	GCGTAACAC
421	CACACCCGCC	GCGCTTAATG	CGCCGCTACA	GGGCGCGTCC	CATTCGCCAT	TCAGGCTGCG
481	CAACTGTTGG	GAAGGGCGAT	CGGTGCGGGC	CTCTTCGCTA	TTACGCCAGC	TGGCGAAAGG
541	GGGATGTGCT	GCAAGGCGAT	TAAGTTGGGT	AACGCCAGGG	TTTTCCAGT	CACGACGTTG
601	TAAAACGACG	GCCAGTGAGC	GCGCGTAATA	CGACTCACTA	TAGGGCGAAT	TGGGTACCGG
661	GCCCCCCTC	GAGGTCGACG	GTATCGATAA	GCTTGATATC	GAATTCATGG	ACATGGACTC
721	GTCGCCGTCG	ACACAGGACT	GTGGCGGCTG	GCTGCTGTAC	GTCTCCCTCG	CTGCCAAATG
781	CGGCGGGCGAC	CCTTGCCGCG	TCGTCGGCTT	CGTCGCCGTT	GCCGTCGTCG	CCTTCGCCGT
841	CACGTCGCTC	CTGCACTGGC	TGTCGCCCCG	TGGCCCCGGC	TGGGGGAGGT	ATTGGTGGAA
901	CAGGAGGGGT	GGTCTGGGCA	TTGCTGCCGC	CATTCTGCGG	CCCCGTGGGT	TGCCCGTGCT
961	CGGCAGCATG	TCGCTCATGG	CGGGACTCGC	GCACCGGAAG	CTCGCCGCGG	CGGCGGGGGG
1021	CTCGCCGGCG	AGGCGGCGCC	TCATGGCGCT	GTCTCTCGGG	GAGACACGGG	TGGTGGTCAC
1081	CGCCGACCCC	GGCGTCGCGC	GGGAGCTCCT	CGCCAGCGCG	GCGTTCGCGG	ACGGGCCGGT
1141	GAAGGAGTCC	GCGTACGGGA	TGCTGTTCCA	CCGCGCCATC	GGGTTGCGCG	CCTACGGCAC
1201	GTAATGGGCG	GCGCTCCGCC	GCGTCGCGTC	CACGCACCTC	TTCTCGCCGA	GGCAGGTGTC
1261	CGCCTCCGCC	GCGCAGCGCG	CGGTGATCGC	GCGCCAGATG	GTGGAGGCCA	TGAGGTCCGC
1321	CGCCGCCGCC	GCCGCCGGTG	GCGGCGTGGC	GGCGAGGCCG	TTCTGAAGC	GCGCGTCGCT
1381	GCACAACGTG	ATGTGGTCGG	TGATCGGGAG	GAAGTACGAG	CTGGCGGCGC	CGGAGAGCGA
1441	GGAGACGGCG	GAGCTGAGGA	GACTGGTGGA	CGAAGGCTAC	GACCTCCTCG	GCCAGCTCAA
1501	CTGGTCCGAC	CACCTCCCAT	GGCTCGCACC	CTTTGACCTC	AAGAAGACGC	GGTCAAGGTG
1561	CTCGTCCCTT	GTCCCCCGCG	TCAACCGCTT	CGTCACCCGC	ATCATCGACG	AGCACCCTGC
1621	TCGCCTCAGC	CTCGCCGTCG	ACGCCGCCGT	CGACTTCACC	GACGTCCTTC	TCTCCCTCCA
1681	CGGCGGCGAC	AAGCTCTCCG	ACGCCGACAT	GGTCGCCGTC	CTCTGGGAGA	TGATCTTTTCG

1741	AGGGACGGAC	ACGGTGGCGG	TCCTGATCGA	GTGGGTGGCG	GCGAGGCTGG	TGCTGCACCA
1801	GGACGTGCAG	GCCAGGGTCC	ATGACGAGCT	GGACCGAGTG	GTCGGGTCGG	ACCGGGCAGT
1861	GACCGAGTCG	GACGCGTCCA	AGCTGGTCTA	CCTCCAAGCG	GTGATCAAAG	AGGTCTCTGCG
1921	CCTCCACCCG	CCGGGCCCCAC	TGCTCTCGTG	GGCACGCCTC	GCCACGTCGG	ATGTACACGT
1981	CGGCGGGTTC	CTCATACCCT	CTGGGACCAC	CGCCATGGTG	AACATGTGGG	CCATAACCCA
2041	TGACCCTGCC	GTTTGGCCCC	ACCCGAACGA	GTTCAAACCA	GAGAGGTTTC	TCGCAGGGCC
2101	CTCGTCGGAC	CAGGCCACGG	AGTTTCCGAT	AATGGGGTCG	GATCTCAGGC	TCGCGCCGTT
2161	GGGTTCAGGA	AGCGGAAGCT	GCCCCGGCAA	GTCGCTCGCC	ATCGCCACTG	TCGGATTCTG
2221	GGTTGCCACG	TTGCTACACG	AGTTCGATTG	GCTTCCCTTG	TCAGATAAGT	CGCGCGGCGT
2281	CGATCTGTCT	GAGGTGCTGA	AGCTGTCTGT	CGAGATGGCA	ACCCCGCTGG	AGGCAAGGCT
2341	AAGGCCGCGA	CGCAAGGTGT	GAGGATCCAC	TAGTTCTAGA	GCGGCCGCCA	CCGCGGTGGA
2401	GCTCCAGCTT	TTGTTCCCTT	TAGTGAGGGT	TAATTGCGCG	CTTGGCGTAA	TCATGGTCAT
2461	AGCTGTTTCC	TGTGTGAAAT	TGTTATCCGC	TCACAATTCC	ACACAACATA	CGAGCCGGAA
2521	GCATAAAGTG	TAAAGCCTGG	GGTGCCTAAT	GAGTGAGCTA	ACTCACATTA	ATTGCGTTGC
2581	GCTCACTGCC	CGCTTTCCAG	TCGGGAAACC	TGTCGTGCCA	GCTGCATTAA	TGAATCGGCC
2641	AACGCGCGGG	GAGAGGCGGT	TTGCGTATTG	GGCGCTCTTC	CGCTTCCTCG	CTCACTGACT
2701	CGCTGCGCTC	GGTCGTTCGG	CTGCGGCGAG	CGGTATCAGC	TCACTCAAAG	GCGGTAATAC
2761	GGTTATCCAC	AGAATCAGGG	GATAACGCAG	GAAAGAACAT	GTGAGCAAAA	GGCCAGCAAA
2821	AGGCCAGGAA	CCGTAAAAAG	GCCGCGTTGC	TGGCGTTTTT	CCATAGGCTC	CGCCCCCCTG
2881	ACGAGCATCA	CAAAAATCGA	CGCTCAAGTC	AGAGGTGGCG	AAACCCGACA	GGACTATAAA
2941	GATACCAGGC	GTTTCCCCCT	GGAAGCTCCC	TCGTGCGCTC	TCTGTTCCTG	ACCCTGCCGC
3001	TTACCGGATA	CCTGTCCGCC	TTTCTCCCTT	CGGGAAGCGT	GGCGCTTTCT	CATAGCTCAC
3061	GCTGTAGGTA	TCTCAGTTCT	GTGTAGGTCG	TTCTGCTCAA	GCTGGGCTGT	GTGCACGAAC
3121	CCCCCGTTCA	GCCCCACCGC	TGCGCCTTAT	CCGGTAACCT	TCGTCTTGAG	TCCAACCCGG
3181	TAAGACACGA	CTTATCGCCA	CTGGCAGCAG	CCACTGGTAA	CAGGATTAGC	AGAGCGAGGT
3241	ATGTAGGCGG	TGCTACAGAG	TTCTTGAAGT	GGTGGCCTAA	CTACGGCTAC	ACTAGAAGGA
3301	CAGTATTTGG	TATCTGCGCT	CTGCTGAAGC	CAGTTACCTT	CGGAAAAAGA	GTTGGTAGCT
3361	CTTGATCCGG	CAAACAAACC	ACCGCTGGTA	GCGGTGGTTT	TTTTGTTTGC	AAGCAGCAGA
3421	TTACGCGCAG	AAAAAAAGGA	TCTCAAGAAG	ATCCTTTGAT	CTTTTCTACG	GGGTCTGACG
3481	CTCAGTGGAA	CGAAAACTCA	CGTTAAGGGA	TTTTTGGTCAT	GAGATTATCA	AAAAGGATCT
3541	TCACCTAGAT	CCTTTTAAAT	TAAAAATGAA	GTTTTAAATC	AATCTAAAGT	ATATATGAGT
3601	AAACTTGGTC	TGACAGTTAC	CAATGCTTAA	TCAGTGAGGC	ACCTATCTCA	GCGATCTGTC
3661	TATTTTCGTT	ATCCATAGTT	GCCTGACTCC	CCGTCGTGTA	GATAACTACG	ATACGGGAGG
3721	GCTTACCATC	TGGCCCCAGT	GCTGCAATGA	TACCGCGAGA	CCCACGCTCA	CCGGCTCCAG
3781	ATTTATCAGC	AATAAACCCAG	CCAGCCGGAA	GGGCCGAGCG	CAGAAGTGGT	CCTGCAACTT
3841	TATCCGCCCT	CATCCAGTCT	ATTAATTGTT	GCCGGGAAGC	TAGAGTAAGT	AGTTCGCCAG
3901	TTAATAGTTT	GCGCAACGTT	GTTGCCATTG	CTACAGGCAT	CGTGGTGTCA	CGCTCGTCGT
3961	TTGGTATGGC	TTCATTTCAGC	TCCGGTTCCC	AACGATCAAG	GCGAGTTACA	TGATCCCCCA
4021	TGTTGTGCAA	AAAAGCGGTT	AGCTCCTTCG	GTCTCTCCGAT	CGTTGTGAGA	AGTAAGTTGG
4081	CCGCAGTGTT	ATCACTCATG	GTTATGGCAG	CACTGCATAA	TTCTCTTACT	GTCATGCCAT
4141	CCGTAAGATG	CTTTTCTGTG	ACTGGTGAGT	ACTCAACCAA	GTCATTCTGA	GAATAGTGTA
4201	TGCGGCGACC	GAGTTGCTCT	TGCCCCGGCT	CAATACGGGA	TAATACCGCG	CCACATAGCA
4261	GAACTTTAAA	AGTGCTCATC	ATTGGAAAAAC	GTTCTTCGGG	GCGAAAACTC	TCAAGGATCT
4321	TACCGCTGTT	GAGATCCAGT	TCGATGTAAC	CCACTCGTGC	ACCCAACCTA	TCTTCAGCAT
4381	CTTTTACTTT	CACCAGCGTT	TCTGGGTGAG	CAAAAACAGG	AAGGCAAAAT	GCCGCAAAAA
4441	AGGGAATAAG	GGCGACACGG	AAATGTTGAA	TACTCATACT	CTTCTTTTTT	CAATATTATT
4501	GAAGCATTTA	TCAGGGTTAT	TGTCTCATGA	GCGGATACAT	ATTTGAATGT	ATTTAGAAAA
4561	ATAAACAAAT	AGGGGTTCCT	CGCACATTTC	CCCAGAAAGT	GCCAC	

Certified by: Patrick Barretto

Date: 10/25/2019