

## Product datasheet for **SC125747**

### Bile salt activated lipase (CEL) (BC042510) Human Untagged Clone

#### Product data:

Product Type:	Expression Plasmids
Product Name:	Bile salt activated lipase (CEL) (BC042510) Human Untagged Clone
Tag:	Tag Free
Symbol:	Bile salt activated lipase
Synonyms:	BAL; bile-salt-activated lipase; bile salt-dependent lipase, oncofetal isoform; bile salt-stimulated carboxyl ester lipase; bile salt-stimulated lipase; BSDL; BSSL; carboxy; carboxyl ester hydrolase; CEase; CELL; FAP; FAPP; LIPA; MODY8; OTTHUMP00000022450
Mammalian Cell Selection:	None
Vector:	<u><a href="#">pCMV6-XL5</a></u>
E. coli Selection:	Ampicillin (100 ug/mL)



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**Fully Sequenced ORF:**

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>OriGene sequence for BC042510 edited
GATGCTACCATGGGCGCCTGCAACTGGTTGTGTTGGGCCTCACCTGCTGCTGGGCAGT
GGCGAGTGCCGCGAAGCTGGGCGCCGTGTACACAGAAGGTGGGTTCTGGAAGGCGTCAA
TAAGAAGCTCGGCCTCTGGGTGACTCTGTGGACATCTTCAAGGGCATCCCCTTCGCAGC
TCCCACCAAGGCCTGAAAAATCCTCAGCCACATCCTGGCTGGCAAGGGACCTGAAGGC
CAAGAACTTCAAGAAGAGATGCCTGCAGGCCACCATACCCAGGACAGCACCTACGGGGA
TGAAGACTGCCTGTACCTCAACATTTGGGTGCCCGAGGCAAGCAAGTCTCCCGGGA
CCTGCCCGTTATGATCTGGATCTATGGAGGCGCCTTCTCATGGGTCCGGCCATGGGGC
CAACTTCTCAACAACCTACCTGTATGACGGCGAGGAGATCGCCACACGCGAAACGTCAT
CGTGGTACCTTCAACTACCGTGTGCGGCCCTTGGGTTCTCAGCACTGGGACGCCAA
TCTGCCAGGTAATATGGCCTTCGGGATCAGCACATGGCCATTGCTTGGGTGAAGAGGAA
TATCGCGCCTTCGGGGGGACCCCAACAACATCACGCTCTTCGGGGAGTCTGCTGGAGG
TGCCAGCGTCTCTGCAGACCCTCTCCCCTACAACAAGGGCCTCATCCGGCGAGCCAT
CAGCCAGAGCGGCGTGGCCCTGAGTCCCTGGGTATCCAGAAAAACCACTCTTCTGGGC
CAAAAAGGTGGCTGAGAAGGTGGGTTGCCCTGTGGGTGATGCCGCGAGGATGGCCAGTG
TCTGAAGGTTACTGATCCCCGAGCCCTGACGCTGGCCTATAAGGTGCCGCTGGCAGGCT
GGAGTACCCCATGCTGCACTATGTGGGTTCTGTCCTGTGATTGATGGAGACTTCATCCC
CGCTGACCCGATCAACCTGTACGCCAACGCCGCGACATCGACTATATAGCAGGCACCAA
CAACATGGACGGCCACATCTTCGCCAGCATCGACATGCCTGCCATCAACAAGGGCAACAA
GAAAGTACGGAGGACTTCTACAAGCTGGTCAGTGAGTTCACAATCACCAAGGGGCTCAG
AGGCGCCAAGACGACCTTTGATGTCTACACCGAGTCTGGGCCAGGACCCATCCCAGGA
GAATAAGAAGAAGACTGTGGTGGACTTTGAGACCGATGTCCTCTTCTGGTGCCACCGA
GATTGCCCTAGCCAGCACAGACCAATGCCAAGAGTGCCAAGACCTACGCTACCTGTT
TTCCCATCCCTCTCGGATGCCCCGTCTACCCCAAATGGGTGGGGCCGACCATGCAGATGA
CATTACAGTACGTTTTTCGGGAAGCCCTTCGCCACCCCCACGGGCTACCGGCCCAAGACAG
GACAGTCTCTAAGGCATGATCGCTACTGGACCAACTTTGCCAAAACAGGGGACCCCAA
CATGGGCGACTCGGCTGTGCCACACACTGGGAACCCTACACTACGAAAAACAGCGGCTA
CCTGGAGATCACCAAGAAGATGGGCAGCAGCTCCATGAAGCGGAGCTGAGAACCAACTT
CCTGCGCTACTGGACCCTCACCTATCTGGCGTGCCACAGTGACCGACCAGGAGGCCAC
CCCTGTGCCCCCACAGGGGACTCCGAGGCCACTCCCGTGCCCCCACGGGTGACTCCGA
GACCGCCCCCGTCCGCCACGGGTGACTCCGGGGCCCCCCCCGTGCCGCCACGGGTGA
CTCCGGGGCCCCCCCCGTGCCGCCACGGGTGACTCCGGGGCCCCCCCCGTGCCGCCAC
GGGTGACTCCGGGGCCCCCCCCGTGCCGCCACGGGTGACTCCGGCGCCCCCCCCGTGCC
GCCACGGGTGACGCGGGCCCCCCCCGTGCCGCCACGGGTGACTCCGGCGCCCCCCCC
CGTGCCGCCACGGGTGACTCCGGGGCCCCCCCCGTGACCCACGGGTGACTCCGAGAC
CGCCCCGTGCCGCCACGGGTGACTCCGGGGCCCCCCCCGTGCCGCCACGGGTGACTC
TGAGGCTGCCCTGTGCCCCACAGATGACTCCAAGGAAGCTCAGATGCCTGCAGTCAT
TAGGTTTTAGCGTCCCATGAGCCTTGGTATCAAGAGGCCACAAGAGTGGGACCCAGGGG
CTCCCCCTCCATCTTGAGCTCTTCTGAATAAAGCCTCATACCCCTGAAAAAAAAAAAAA
AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA
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<b>5' Read Nucleotide Sequence:</b>	>OriGene 5' read for BC042510 unedited GTTTAGATATTTGTAATACGACTTTATATAGGGCGGCCGCGNATTCANATCTGGTACCGG TCCGGAATTCCTGGGATGATGCTCACCATGGGGCGCCTGCAACTGGTTGTGGGCTC ACCTGCTGCTGGGAGTGGCGAGTGCCGCGAAGCTGGGCGCCGTGTACACAGAAGGTGGG TTCGTGGAAGGCGTCAATAAGAAGCTCGGCCTCCTGGGTGACTCTGTGGACATCTTCAAG GGCATCCCCTTCGAGCTCCCACCAAGGCCCTGAAAAATCCTCAGCCACATCCTGGCTGG CAAGGGACCCTGAAGGCCAAGAAGCTTCAAGAAGAGATGCCTGCAGGCCACCATCAGCCAG GACAGCACCTACGGGGATGAAGACTGCCTGTACCTCAACATTTGGGTGCCCCAGGGCAGG AAGCAAGTCTCCCGGGACCTGCCCGTTATGATCTGGATCTATGGAGGCGCCTTCCTCATG GGGTCCGGCCATGGGGCCAACCTCCTCAACAACACTACCTGTATGACGGCGAGGAGATCGCC ACACGCGGAAACGTATCGTGGTACCTTCACTACCGTGTGGCCCTTGGGTTCTCA GCACTGGGAGCGCAATCTGCCAGGTAATATGGCCTTCGGGATCAGCACATGGCCATTG CTTTGGTTGAGGAAGAAATATCGCGGCCTCGGGGGGGGACCCCAACAACATCACGCTTC TCGGGGAGTCTGCTGGAAGTCAAGCGTCTTCTGCAAAACCCTTCCCCCTAAACAAGGG CCTCATCCGGCGAGCCATCACCCAAAGGGGCTGGCCCTGAATCCCTGGGTATCCAGAAA AACCCACTTTTTGGGCCAAAAGTGGCTGAAAGGTGGTTGCCTGGGGGGATCCCCGGATG GCCATGGTTTGAGTTCTGTTCC
<b>Restriction Sites:</b>	Please inquire
<b>ACCN:</b>	BC042510
<b>OTI Disclaimer:</b>	Our molecular clone sequence data has been matched to the reference identifier above as a point of reference. Note that the complete sequence of our molecular clones may differ from the sequence published for this corresponding reference, e.g., by representing an alternative RNA splicing form or single nucleotide polymorphism (SNP).
<b>Components:</b>	The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).
<b>Reconstitution Method:</b>	<ol style="list-style-type: none"> <li>1. Centrifuge at 5,000xg for 5min.</li> <li>2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.</li> <li>3. Close the tube and incubate for 10 minutes at room temperature.</li> <li>4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.</li> <li>5. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date of shipping when stored at -20°C.</li> </ol>
<b>RefSeq:</b>	<a href="#">BC042510.1</a> , <a href="#">AAH42510.1</a>
<b>RefSeq Size:</b>	2316 bp
<b>Locus ID:</b>	1056
<b>Cytogenetics:</b>	9q34.13
<b>Protein Families:</b>	Druggable Genome
<b>Protein Pathways:</b>	Glycerolipid metabolism, Metabolic pathways, Steroid biosynthesis

**Gene Summary:**

The protein encoded by this gene is a glycoprotein secreted from the pancreas into the digestive tract and from the lactating mammary gland into human milk. The physiological role of this protein is in cholesterol and lipid-soluble vitamin ester hydrolysis and absorption. This encoded protein promotes large chylomicron production in the intestine. Also its presence in plasma suggests its interactions with cholesterol and oxidized lipoproteins to modulate the progression of atherosclerosis. In pancreatic tumoral cells, this encoded protein is thought to be sequestered within the Golgi compartment and is probably not secreted. This gene contains a variable number of tandem repeat (VNTR) polymorphism in the coding region that may influence the function of the encoded protein. [provided by RefSeq, Jul 2008]