

## Product datasheet for SC205816

## OriGene Technologies, Inc.

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## Argininosuccinate Lyase (ASL) (NM\_001024944) Human 3' UTR Clone

**Product data:** 

**Product Type:** 3' UTR Clones

Product Name: Argininosuccinate Lyase (ASL) (NM 001024944) Human 3' UTR Clone

**Vector:** pMirTarget (PS100062)

Symbol: ASL

Synonyms: ASAL

**ACCN:** NM\_001024944

**Insert Size:** 694 bp

Insert Sequence: >SC205816 3'UTR clone of NM\_001024944

The sequence shown below is from the reference sequence of NM\_001024944. The complete

sequence of this clone may contain minor differences, such as SNPs.

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

AAAA

**ACGCGT**AAGCGGCCGCGCATCTAGATTCGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

**Restriction Sites:** Sgfl-Mlul

**OTI Disclaimer:** Our molecular clone sequence data has been matched to the sequence identifier above as a

point of reference. Note that the complete sequence of this clone is largely the same as the

reference sequence but may contain minor differences, e.g., single nucleotide

polymorphisms (SNPs).





## Argininosuccinate Lyase (ASL) (NM\_001024944) Human 3' UTR Clone - SC205816

**Components:** The cDNA clone is shipped in a 2-D bar-coded Matrix tube as 10 ug dried plasmid DNA. The

package also includes 100 pmols of both the corresponding 5' and 3' vector primers in

separate vials.

RefSeq: <u>NM 001024944.2</u>

**Summary:** This gene encodes a member of the lyase 1 family. The encoded protein forms a cytosolic

homotetramer and primarily catalyzes the reversible hydrolytic cleavage of argininosuccinate into arginine and fumarate, an essential step in the liver in detoxifying ammonia via the urea cycle. Mutations in this gene result in the autosomal recessive disorder argininosuccinic aciduria, or argininosuccinic acid lyase deficiency. A nontranscribed pseudogene is also located on the long arm of chromosome 22. Alternatively spliced transcript variants encoding

different isoforms have been described. [provided by RefSeq, Jul 2008]

Locus ID: 435

MW: 25.1