

Product datasheet for **RG237804**

AIPL1 (NM_001285399) Human Tagged ORF Clone

Product data:

| | |
|---------------------------|--|
| Product Type: | Expression Plasmids |
| Product Name: | AIPL1 (NM_001285399) Human Tagged ORF Clone |
| Tag: | TurboGFP |
| Symbol: | AIPL1 |
| Synonyms: | AIPL2; LCA4 |
| Mammalian Cell Selection: | Neomycin |
| Vector: | pCMV6-AC-GFP (PS100010) |
| E. coli Selection: | Ampicillin (100 ug/mL) |
| ORF Nucleotide Sequence: | >RG237804 representing NM_001285399. Blue=ORF Red=Cloning site Green=Tag(s) |

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GCTCGTTTGTAGTAACCGTCAGAATTTGTAAATACGACTACTATAGGGCGCCGGGAATTCGTCGACTG
GATCCGGTACCGAGGAGATCTGCCGCCGCGATCGCC
ATGGATGCCGCTCTGCTCCTGAACGTGGAAGGGGTCAAGAAAACCATCTGCACGGGGGCACGGGCGAG
CTCCAAACTTCATACCGGATCCCGAGAGCGGACAGTCATTGACGACAGTCGGCAGGTGGGCCAGCCC
ATGCACATCATCATCGGAAACATGTTCAAGCTCGAGGTCTGGGAGATCCTGCTTACCTCCATGCGGGTG
CACGAGGTGGCCGAGTTCTGGTGCACACCATCCACACGGGGTCTACCCATCCTATCCCGAGCCTG
AGGCAGATGGCCAGGGCAAGGACCCACAGAGTGGCACGTGCACACGTGCGGGCTGGCCAAACATGTTT
GCCTACCACACGCTGGGCTACGAGGACCTGGACGAGCTGCAGAAGGAGCCTCAGCCTCTGGTCTTTGTG
ATCGAGCTGCTGCAGTTGATGCCCGAGTGATTACCAGAGGGAGACCTGGAACCTGAGCAATCATGAG
AAGATGAAGGCGGTGCCCGTCTCCACGGAGAGGAAATCGGCTCTTCAAGCTGGCCGCTACGAGGAG
GCCTCTTCCAAGTACCAGGAGGCCATCATCTGCCTAAGGAACCTGCAGACCAAGGAGAAGCCATGGGAG
GTGCAAGTGGTGAAGCTGGAGAAGATGATCAATACTCTGATCCTCAACTACTGCCAGTGCCTGCTGAAG
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GCCTACTACGTGCGTGCCCGGCTCACGCAGAGGTGTGGAATGAGGCCGAGGCCAAGGCCGACCTCCAG
AAAGTGTGGAGCTGGAGCCGTCATGCAGAAGGCGGTGCGCAGGGAGCTGAGGCTGCTGGAGAACCAG
ATGGCGGAGAAGCAGGAGGAGGAGCGGCTGCGCTGCCGGAACATGCTGAGCCAGGTGCCACGCAGCCT
CCCGCAGAGCCACCCACAGAGCCACCCGACAGTCATCCACAGAGCCACCTGCAGAGCCACCCACAGCA
CCATCTGCAGAGCTGTCCGACGGCCCCCTGCAGAGCCAGCCACAGAGCCACCCCGTCCCAGGGCAC
TCGCTGCAGCAC
ACGCGTACGCGGCCGCTCGAG - GFP Tag - GTTTAAAC
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Protein Sequence: >Peptide sequence encoded by RG237804
 Blue=ORF Red=Cloning site Green=Tag(s)

MDAALLLNVEGVKKTILHGGTGELPNFITGSRERTVIDDSRQVGGQPMHIIIGNMFKLEVWEILLTSMRV
 HEVAEFWCDTIHTGVYPIILSRSLRQMAQKDPTEWHVHTCGLANMFAYHTLGYEDLDELQKEPQLV
 IELLQVDAPSDYQRETWNLSNHEKMKAVPVLHGEGRNLFKLGRYEEASSKYQEAIIICLRNLQTKKPWE
 VQWLKLEKMINTLILNYCQCLLKKEEYVEVLEHTSDILRHHPGIVKAYVVRARAHAEVWNEAEAKADLQ
 KVLELEPSMQKAVRRELRLLENRMAEKQEERLRCRNMLSQGATQPPAEPPEPPAQSSTEPPAEPPTA
 PSAELSAGPPAEPATEPPPPSPGHSLQH
 TRTRPLEMESDESGLPAMEIECRITGTLNGVEFELVGGEGTPEQGRMTNKMSTKGALTFSPYLLSHV
 MGYGFYHFGTYPSTYENPFLHAINNGGYNTRIEKYEDGGVLHVSFSYRYEAGRVIGDFKVMGTGFPED
 SVIFTDKIIRSNATVEHLHPMGDNDLDGSFTRTFSLRDGGYSSVVDSHMHFKSAIHPSILQNGGPMFA
 FRRVEEDHSNTELGIVEYQHAFKTPDADAGEERV

Restriction Sites: SgfI-MluI

Cloning Scheme:



ACCN: NM_001285399

ORF Size: 1116 bp

OTI Disclaimer: The molecular sequence of this clone aligns with the gene accession number as a point of reference only. However, individual transcript sequences of the same gene can differ through naturally occurring variations (e.g. polymorphisms), each with its own valid existence. This clone is substantially in agreement with the reference, but a complete review of all prevailing variants is recommended prior to use. [More info](#)

OTI Annotation: This clone was engineered to express the complete ORF with an expression tag. Expression varies depending on the nature of the gene.

Components: 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).

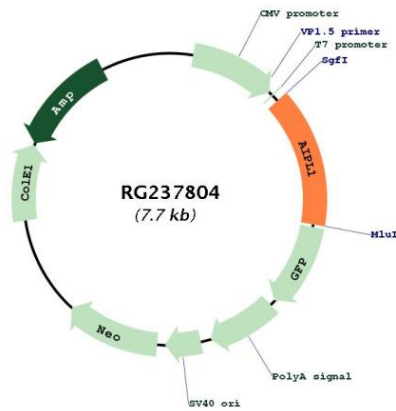
RefSeq: [NM_001285399.3](#)

RefSeq Size: 2954 bp

RefSeq ORF: 1119 bp
Locus ID: 23746
UniProt ID: [Q9NZN9](#)
Cytogenetics: 17p13.2
Protein Families: Druggable Genome
MW: 42.8 kDa
Gene Summary:

Leber congenital amaurosis (LCA) is the most severe inherited retinopathy with the earliest age of onset and accounts for at least 5% of all inherited retinal diseases. Affected individuals are diagnosed at birth or in the first few months of life with nystagmus, severely impaired vision or blindness and an abnormal or flat electroretinogram. The photoreceptor/pineal-expressed gene, AIPL1, encoding aryl-hydrocarbon interacting protein-like 1, is located within the LCA4 candidate region. The encoded protein contains three tetratricopeptide motifs, consistent with chaperone or nuclear transport activity. Mutations in this gene may cause approximately 20% of recessive LCA. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Jan 2014]

Product images:



Circular map for RG237804