

Product datasheet for RC220707L4V

OriGene Technologies, Inc.

9620 Medical Center Drive, Ste 200 Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com CN: techsupport@origene.cn

Factor H (CFH) (NM 000186) Human Tagged ORF Clone Lentiviral Particle

Product data:

Product Type: Lentiviral Particles

Product Name: Factor H (CFH) (NM_000186) Human Tagged ORF Clone Lentiviral Particle

Symbol: Factor H

Synonyms: AHUS1; AMBP1; ARMD4; ARMS1; CFHL3; FH; FHL1; HF; HF1; HF2; HUS

Mammalian Cell

Selection:

Puromycin

Vector: pLenti-C-mGFP-P2A-Puro (PS100093)

Tag: mGFP

ACCN: NM_000186 **ORF Size:** 3693 bp

ORF Nucleotide

The ORF insert of this clone is exactly the same as(RC220707).

Sequence:

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.

RefSeg: NM 000186.2

 RefSeq Size:
 4004 bp

 RefSeq ORF:
 3696 bp

 Locus ID:
 3075

 UniProt ID:
 P08603

 Cytogenetics:
 1q31.3

Domains: CCP

Protein Families: Druggable Genome, Secreted Protein





Factor H (CFH) (NM_000186) Human Tagged ORF Clone Lentiviral Particle - RC220707L4V

Protein Pathways: Complement and coagulation cascades

MW: 139.07 kDa

Gene Summary: This gene is a member of the Regulator of Complement Activation (RCA) gene cluster and

encodes a protein with twenty short consensus repeat (SCR) domains. This protein is secreted into the bloodstream and has an essential role in the regulation of complement activation, restricting this innate defense mechanism to microbial infections. Mutations in this gene have been associated with hemolytic-uremic syndrome (HUS) and chronic hypocomplementemic nephropathy. Alternate transcriptional splice variants, encoding different isoforms, have

been characterized. [provided by RefSeq, Oct 2011]