

Specifications:

Gene:	hCFHR1
Accession:	NP_002104
Insert size:	1006bp
Concentration:	10µg at 0.2µg/µL

hCFHR1 cDNA Plasmid

**CFHR1 complement factor H
related 1 [*Homo sapiens*
(human)]**

Also known as: CFHL; FHR1; HFL1;
HFL2; CFHL1; H36-1; H36-2; CFHL1P;
CFHR1P

Summary:

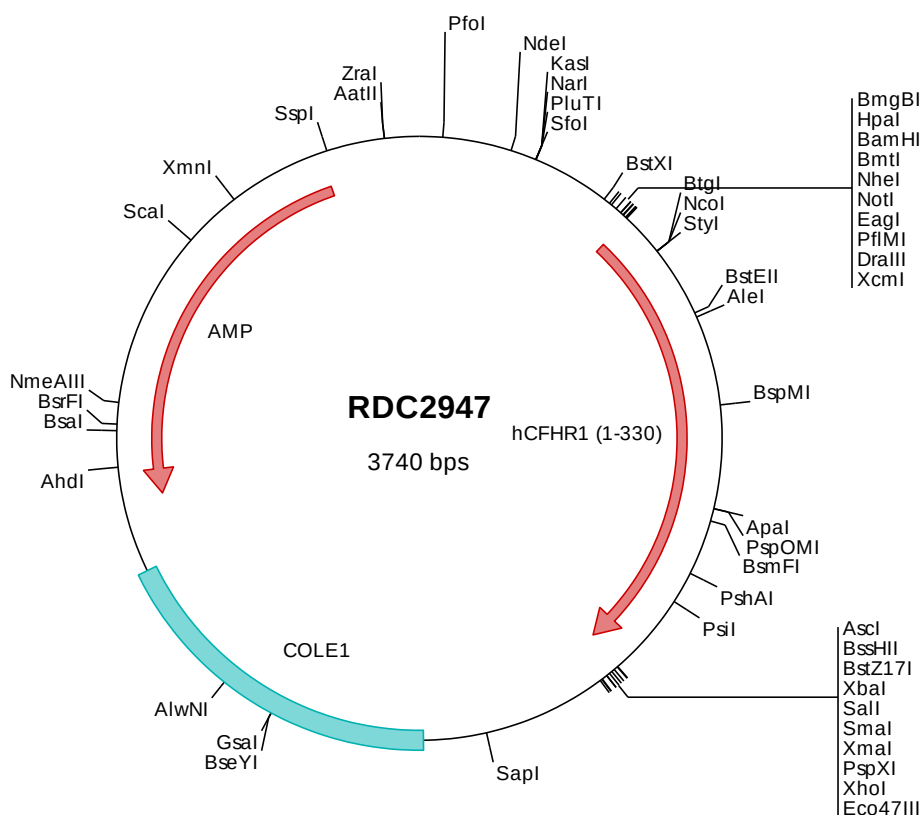
CFHR1 is a secreted protein belonging to the complement factor H protein family. It binds to *Pseudomonas aeruginosa* elongation factor Tuf together with plasminogen, which is proteolytically activated. It is proposed that Tuf acts as a virulence factor by acquiring host proteins to the pathogen surface, controlling complement, and facilitating tissue invasion. Mutations in CFHR1 are associated with an increased risk of atypical hemolytic-uremic syndrome.

Description

This shuttle vector contains the complete ORF for the gene of interest, along with a Kozak consensus sequence for optimal translation initiation. It is inserted NotI to AscI. The gene insert is flanked with convenient multiple cloning sites which can be used to easily cut and transfer the gene cassette into your desired expression vector.

Preparation and Storage

Formulation	cDNA is provided in 10 mM Tris-Cl, pH 8.5
Shipping	Ships at ambient temperature
Stability	1 year from date of receipt when stored at -20°C to -80°C
Storage	Use a manual defrost freezer and avoid repeated freeze-thaw cycles.



> RDC2947 Plasmid DNA Sequence

```

1   tcgcgcgctt  cggatgatgac  ggtgaaaacc  tctgacacat  gcagctcccc  gagacggtca  cagcttgtct  gtaagcggat  gccgggagca  gacaagcccg
101  tcagggcgcg  tcagcgggtg  ttggcgggtg  tcggggctgg  cttactatg  cggcatcaga  gcagattgta  ctgagagtgc  accatatgcg  gtgtgaaata
201  ccgcacagat  gcgtaaggag  aaaataccgc  atcaggcgcc  attcgccatt  caggctcgcg  aactgttggg  aagggcgatc  ggtgcgggcc  tcttcgctat
301  tacgccagct  ggcgaaagg  ggaatgtctg  caaggcgatt  aagttgggta  acgccagggt  tttcccgctc  acgacgttgt  aaaacgacgg  ccagtgaatt
401  ggagacgtgt  taacaagctt  ggatccgata  tcgctagcgc  ggccggccacc  atgtggctcc  tggtcagtgt  aattctaato  tcacggatat  cctctgttgg
501  gggagaagca  acattttgtg  attttccaaa  aataaaccat  ggaattctat  atgatgaaga  aaaatataag  ccattttccc  aggttcctac  aggggaagtt
601  ttctattact  cctgtgaata  taattttgtg  tctcttcaa  aatcattttg  gactgcata  acatgcacag  aagaaggatg  gtcaccaaca  ccaaagtgtc
701  tcagactgtg  tttctttcct  ttgttggaag  atggctatto  tgaatcttca  ggacaaacac  atctggaagg  tgatactgtg  caaattatct  gcaacacagg
801  atacagactt  caaaacaatg  agaacaacat  ttcagtgtga  gaacggggct  ggtccacccc  tcccaaatgc  aggtccactg  acacttctgt  tgtgaatccg
901  cccacagtac  aaaatgctca  tatactgtcg  agacagatga  gtaaataatc  atctggtgag  agagtacgtt  atgaatgtag  gagcccttat  gaaatgtttg
1001  agggatgaaga  agtgaatgtt  ttaaatggaa  actggacaga  accacctcaa  tgcaaaagatt  ctacgggaaa  atgtggggcc  cctccaccta  ttgacaatgg
1101  ggacattact  toattccogt  tgcagatata  tgcctcagct  toatcagttg  agtaccaatg  ccagaacttg  tatcaacttg  agggtaacaa  gcgaataaca
1201  tgtagaaatg  gacaatggtc  agaaccacca  aaatgcttac  atcctgtgtg  aatatcccg  gaaattatgg  aaaattataa  catagcatta  aggtggacag
1301  ccaaacagaa  cgtttatttg  agaacagggt  aatcagctga  atttgtgtgt  aaacggggat  atcgtcttct  atcagcttct  cacacattgc  gaacaacatg
1401  ttgggatggg  aaactggagt  atcccaactg  tgcaaaaaga  taaaggcgcg  ccagtatact  ctagagtcca  caccggggga  attcctcgag  cgctcgtctc
1501  tagcttggcg  taatcatggt  catagctgtt  tctctgttga  aattgtttat  cgctcacaat  tccacacaac  atacgagccg  gaagcataaa  gtgtaaagcc
1601  tggggtgccct  aatgagtgag  ctaactcaca  ttaatttcgt  tgcctcact  gcccgcttcc  cagtcgggaa  acctgtcgtg  ccagctgcat  taatgaatcg
1701  gccaacgcgc  ggggagaggc  ggtttgcgta  ttgggcgctc  ttccgcttcc  tcgctcactg  actcgcctcg  ctcggtcggt  cggctgcggc  gagcggatat
1801  agctcactca  aaggcggtaa  tacggttatc  cacagaatca  ggggataacg  caggaaagaa  catgtgagca  aaaggccagc  aaaaggccag  gaaccgtaaa
1901  aagggccgct  tgcgtggcgt  ttcccatagg  ctccgcccc  ctgacgagca  tcacaaaaat  cgacgctcaa  gtcagaggtg  gcgaaacccc  acaggaactat
2001  aaagatacca  ggcgtttccc  cctggaagct  ccctcgtcgc  ctctcctgtt  ccgacctcgc  cgcttaccgg  atacctgtcc  gcctttctcc  cttcggaagc
2101  cgtggcgctt  tctcaatgct  cacgctgtag  gtatctcagt  tcggtgtagg  tcggttcgct  caagctgggc  tgtgtgcacg  aacccccctg  tcagccccgac
2201  cgctgcgctt  tatcaggtaa  ctactcgtct  gagtccaaac  cggttaagaca  cgacttatcg  ccaactggcg  cagccactgg  taacaggatt  agcagagcga
2301  ggtatgtagg  cgggtgctca  gagttcctga  agtgggtggc  taactacggc  tacactagaa  ggacagtatt  tggtatctgc  gctctgtgta  agccagttac
2401  cttcgaaaaa  agagtgggta  gctcttgatc  cggcaaacaa  accaccgctg  gtacgggtgg  ttttttggtt  tgcaagcagc  agattacgcg  cagaaaaaaa
2501  ggatctcaag  aagatccttt  gatcttttct  acgggggtct  acgctcagtg  gaacgaaaac  tcaactgtaa  ggatttttgt  catgagatta  tcaaaaagga
2601  tcttcaccta  gatcctttta  aattaaaaat  gaagttttaa  atcaatctaa  agtatataat  agtaaaactg  gtctgacagt  taccaatgct  taatcagtga
2701  ggcacctatc  tcagcgatct  gtctatttct  ttcattccata  gttgctgac  tccccgtcgt  gtatataact  acgatacggg  agggcttacc  atctggcccc
2801  agtgcgtcaa  tgataccgag  agaccacgc  tcaccggctc  cagatttatc  agcaataaac  cagccagccg  gaagggccga  gcgcagaagt  ggtcctgcaa
2901  ctttatccgc  ctccatccag  tctattaatt  gttgccggga  agctagagta  agtagttcgc  cagttaatag  tttgcgcaac  gttgttgcca  ttgctacagg
3001  catcgtgggt  tcacgctcgt  cgtttgggat  ggcttcattc  agctccggtt  cccaacgata  aaggcgagtt  acatgatccc  ccatgttggt  caaaaaagcg
3101  gttagctcct  tcggtcctcc  gatcgttgtc  agaagtaagt  tggccgcagt  gttatcactc  atggttatgg  cagcactgca  taattctctt  actgtcatgc
3201  catccgtaag  atgcttttct  gtgactgggt  agtactcaac  caagtcattc  tgagaatagt  gtatgcggcg  accgagttgc  tcttgcccg  cgtcaatacg
3301  ggataatacc  gcgccaata  gcgaactttt  aaaagtgtct  atcattggaa  aacgttcttc  ggggcgaaaa  ctctcaagga  tcttaccgct  gttgagatcc
3401  agttcgtatg  aacccactcg  tgcacccaac  tgatcttcag  catcttttac  tttcaccagc  gtttctgggt  gagcaaaaac  aggaaggcaa  aatgccgcaa
3501  aaaaggggat  aaggcgac  cggaaatgtt  gaatactcat  actcttctct  tttcaatatt  attgaagcat  ttatcagggt  tattgtctca  tgagcggata
3601  catatttgaa  tgtatttaga  aaaataaaca  aataggggtt  ccgcgcacat  ttccccgaaa  agtggccact  gacgtctaag  aaaccattat  tatcatgaca
3701  ttaacctata  aaaataggcg  tatcacgagg  ccctttcgtc

```

> RDC2947 Translated Insert Sequence

```

1   mwllsvili  srissvggea  tfcdfpkinh  gilydeekyk  pfsqvptgev  fyysceynfv  spsksfwtri  tcteegwspt  pkclrlcfff  fvenghsess
101  gqthlegdtv  qlicntgyrl  qnnenniscv  ergwstppkc  rstdtscvnp  ptvqnahils  rqmskypsge  rvryecrspe  emfgdeevmc  lngnwtpeppq
201  ckdstgkcp  pppidngdit  sfplsvyapa  ssveyqcqnl  yqlegnkrit  crnqgwsepp  kclhpcvisr  eimenynial  rwtakqklyl  rtgesaefvc
301  krgyrlssrs  htlrttcwdg  kleyptcagr

```