

Product datasheet for SC111686

RHD (NM_016124) Human Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	RHD (NM_016124) Human Untagged Clone
Tag:	Tag Free
Symbol:	RHD
Synonyms:	CD240D; DIIIc; RH; Rh4; RH30; RHCED; RhDCw; RHDel; RHDVA(TT); RhII; RhK562-II; RhPI; RHPII; RHXIII
Mammalian Cell Selection:	None
Vector:	<u>pCMV6-XL4</u>
E. coli Selection:	Ampicillin (100 ug/mL)
Fully Sequenced ORF:	>OriGene ORF within SC111686 sequence for NM_016124 edited (data generated by NextGen Sequencing)

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ATGAGCTCTAAGTACCCGCGGTCTGTCCGGCGCTGCCTGCCCTCTGGGCCCTAACACTG
GAAGCAGCTCTCATTCTCCTCTTCTATTTTTTACCCACTATGACGCTTCCTTAGAGGAT
CAAAAGGGGCTCGTGGCATCCTATCAAGTTGGCCAAGATCTGACCGTGATGGCGGCCATT
GGCTTGGGCTTCCCTCACCTCGAGTTTCCGGAGACACAGCTGGAGCAGTGTGCCTTCAAC
CTCTTCATGCTGGCGCTTGGTGTGACAGTGGCAATCCTGCTGGACGGCTTCTGAGCCAG
TTCCCTTCTGGGAAGGTGGTCATCACACTGTTCAAGTATTCGGCTGGCCACCATGAGTGCT
TTGTCCGGTCTGATCTCAGTGGATGCTGTCTTGGGGAAGGTCAACTTGGCGCAGTTGGT
GTGATGGTCTGGTGGAGGTGACAGCTTAGGCAACCTGAGGATGGTCATCAGTAATATC
TTCAACACAGACTACCACATGAACATGATGCACATCTACGTGTTCCGAGCCTATTTGGG
CTGTCTGTGGCCTGGTGCCTGCCAAAGCCTCTACCCGAGGGAACGGAGGATAAAGATCAG
ACAGCAACGATACCCAGTTTGTCTGCCATGCTGGGCGCCCTCTTCTTGTGGATGTTCTGG
CCAAGTTTCAACTCTGCTCTGCTGAGAAGTCCAATCGAAAGGAAGATGCCGTGTTCAAC
ACCTACTATGCTGTAGCAGTCAGCGTGGTACAGCCATCTCAGGGTCATCCTTGGCTCAC
CCCCAAGGGAAGATCAGCAAGACTTATGTGCACAGTGCAGTGGTGGCAGGAGGCGTGGCT
GTGGGTACCTCGTGTCACCTGATCCCTTCTCCGTGGCTTGCCATGGTGTGGTCTTGTG
GCTGGGCTGATCTCCGTCGGGGGAGCCAAGTACCTGCCGGGTGTTGTAACCGAGTGCTG
GGGATTTCCCACAGCTCCATCATGGGCTACAACCTCAGCTTGTGGTCTGCTTGGAGAG
ATCATCTACATTGTGCTGCTGGTGTGATACCGTCGGAGCCGGCAATGGCATGATTGGC
TTCCAGTCTCCTCAGCATTGGGGAACCTCAGCTTGGCCATCGTGATAGCTCTCAGCTCT
GGTCTCCTGACAGGTTTGTCTCCTAAATCTTAAAAATATGGAAGCACCTCATGAGGCTAAA
TATTTTGTGATACCAAGTTTTCTGGAAGTTTCTCATTGGCTGTTGGATTTTAA

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Clone variation with respect to NM_016124.3



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5' Read Nucleotide Sequence:

>OriGene 5' read for NM_016124 unedited
 GCATTTTGTAAATACGACTCACTATAGGGCGGCCGCAATTTCGGCAGGAGCTGGTCTGGT
 GGAACCCCTGCACAGAGACGGACACAGGATGAGCTCTAAGTACCCGCGGTCTGTCCGGCG
 CTGCCTGCCCCCTCTGGGCCCTAACACTGGAAGCAGCTCTCATTCTCTCTTATTTTTT
 TACCCACTATGACGCTTCTTAGAGGATCAAAAGGGGCTCGTGGCATCCTATCAAGTTGG
 CCAAGATCTGACCGTGATGGCGGCCATTGGCTTGGGCTTCCCTCACCTCGAGTTCCGGAG
 ACACAGCTGGAGCAGTGTGGCCTTCAACCTCTTCACTGCTGGCGCTTGGTGTGCAGTGGGC
 AATCCTGCTGGACGGCTTCTGAGCCAGTTCCTTCTGGGAAGGTGGTATCACACTGTT
 CAGTATTCGGCTGGCCACCATGAGTGCTTTGTCGGTGCTGATCTCAGTGGATGCTGTCTT
 GGGGAAGGTCAACTTGGCGCAGTTGGTGGTATGGTGGTGGTGGAGGTGACAGCTTAGG
 CAACCTGAGGATGGTATCAGTAATATCTTCAACACAGACTACCACATGAACATGATGCA
 CATCTACGTGTTTCGACGCTATTTTGGGCTGTCTGTGGCCTGGTGCCTGCCAAAGCCTCT
 ACCCGAGGGAACGGAGGATAAAGATCAGACAGCAACGATACCCAGTTTGTCTGCCATGCT
 GGGCGCCCTCTTCTGTGGATGTTCTGGCCAAGTTTCAACTCTGCTCTGCTGAGAAGTCC
 AATCGAAGGGAGAATGCCGTGTTTACACCTACTATGCTGTAGCAGTCAGCGTGGTACAG
 CCATCTCAGGGTATCCTTGGCTCACCCNACGGGNAAGATCAGCAGACTATGTGCACAGT
 GCGTGTGGCAAGAGCGTGGCTGTGGNTACCTCGTGTACCTGATCCCTTCTNCGTGGCT
 NGCCATGGNGCTGGGN

3' Read Nucleotide Sequence:

>OriGene 3' read for NM_016124 unedited
 AAACAAGGAAAACACCANACACATTCACCCCTTTTACTTGNACGCGGCCGCAATT
 NANGATCGGCTT
 TTTTTTTTTTATGGGAAACATGGCTGTATTTTATTGGTGAATTTACAGCGCATAAAAA
 ATGGGGAAATTTCTCAAAAAGGGCAAAAAAGGATTCAACTCCATTTTCTTTGACTCC
 AGGGCCTGGGCAACATTGAAACTTTGCTGTCTAAGCGTTTCTCACCTCCAAATGCCG
 GCAACAGTGAAAGGAATTTGCTTTTAAACAGGCCTGTTTTTTTTGGATGCTTTT
 GCTTAAATCCAACCTCCAAATGAGGAACTTCCAAAAAATTGGTCTCCTCAAAATTTA
 ACCTCATGAGGGCTTTCCATTTTTTAAATTTAGGAACAAACCTGTCCGGAGACCAAAC
 GTGAAATCTTACCACGGCCAAGCTGATTTCCCAATGTTGGGAGGCCCTTGAACCA
 ATCCTGCCATTGGCCGGCTCCAACGGGTTCAAACACCCCCCAACAATGTAACAAATCTT
 CTCAAGCACAACCAACAGGTTCAACTTGAACCCCTTTATGGAACCCGCGGGGAAATCCC
 CCGATTTTGGTTCCAAAACCCCGGAGAGACTTTGGGTTCCCGCGGGAAATCCCCCC
 ACCCCAAACACCCACCCCTTGGCCACCCCCCGTAAAAGGTGACCGGTTACAACAAGTA
 TCCTCCCAGTCCCCTTTTTGGAAAACCCGATTTGTGAAAAAAAAGCCTGGTGGATTT
 CCCTGGGGGGGAGCCACACGGAATCCTCTTAAATGGGTTTACCCACTCTATTTGTTAAAA
 TTAATAGGGGTTTAAACACGGCTTTTTCTTTTCTCATGGACCCCCCAAAAAATAATT
 CAACCCTGGCCAAATCCCCCACTAAACGGG

Restriction Sites:

NotI-NotI

ACCN:

NM_016124

Insert Size:

1550 bp

OTI Disclaimer:

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).

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).

Reconstitution Method:

1. Centrifuge at 5,000xg for 5min.
2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.
3. Close the tube and incubate for 10 minutes at room temperature.
4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.
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.

RefSeq: [NM_016124.2](#), [NP_057208.2](#)

RefSeq Size: 2709 bp

RefSeq ORF: 1254 bp

Locus ID: 6007

UniProt ID: [Q02161](#)

Cytogenetics: 1p36.11

Domains: Ammonium_transp

Protein Families: Transmembrane

Gene Summary: The Rh blood group system is the second most clinically significant of the blood groups, second only to ABO. It is also the most polymorphic of the blood groups, with variations due to deletions, gene conversions, and missense mutations. The Rh blood group includes this gene, which encodes the RhD protein, and a second gene that encodes both the RhC and RhE antigens on a single polypeptide. The two genes, and a third unrelated gene, are found in a cluster on chromosome 1. The classification of Rh-positive and Rh-negative individuals is determined by the presence or absence of the highly immunogenic RhD protein on the surface of erythrocytes. Multiple transcript variants encoding different isoforms have been found for this gene. [provided by RefSeq, Jul 2008]
Transcript Variant: This variant (1) represents the predominant transcript and encodes isoform 1. Sequence Note: This RefSeq record was created from transcript and genomic sequence data to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the transcript record were based on transcript alignments.