

Product datasheet for **SC327504**

SLC14A1 (NM_001146036) Human Untagged Clone

Product data:

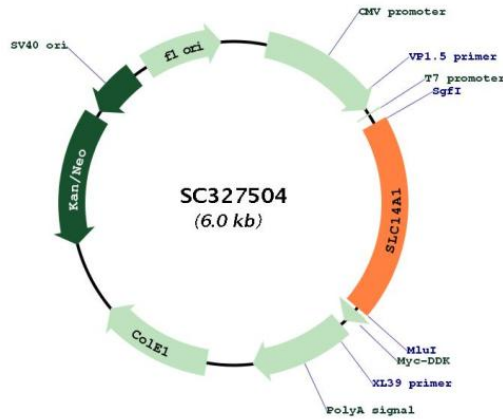
Product Type:	Expression Plasmids
Product Name:	SLC14A1 (NM_001146036) Human Untagged Clone
Tag:	Tag Free
Symbol:	SLC14A1
Synonyms:	HsT1341; HUT11; JK; Jk(b); RACH1; RACH2; UT-B1; UT1; UTE
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)
Fully Sequenced ORF:	>SC327504 representing NM_001146036. Blue=Insert sequence Red=Cloning site Green=Tag(s)

```
GCTCGTTT TAGTGAACCGTCAGAATTTTGT AATACGACTCACTATAGGGCGCCGGGAATTCGTCGACTG
GATCCGGTACCGAGGAGATCTGCCGCC GCGATCGCC
ATGGAGGACAGCCCACTATGGTTAGAGTGGACAGCCCACTATGGTTAGGGGTGAAAACAGGTTTCG
CCATGTCAAGGGAGAAGGTGCTTCCCCAAAGCTTTGGCTATGTCACCGGTGACATGAAAGAAGTTGCC
AACCAGCTTAAAGACAAACCCGTGGTGTCCAGTTCATTGACTGGATTCTCCGGGCATATCCCAAGTG
GTGTTTCGTCACAACCCCGTCAGTGGAACTCCTGATTCTGGTAGGACTTCTGTTCAGAACCCTGGTGG
GCTCTCACTGGCTGGCTGGGAACAGTGGTCTCCACTCTGATGGCCCTCTGCTCAGCCAGGACAGGTCA
TTAATAGCATCTGGGCTCTATGGCTACAATGCCACCCTGGTGGGAGTACTCATGGCTGTCTTTTCGGAC
AAGGGAGACTATTTCTGGTGGCTGTACTCCCTGTATGTGCTATGTCCATGACTTGCCCAATTTCTCA
AGTGCATTGAATTCATGCTCAGCAAATGGGACCTCCCGCTTTCACCCTCCCTTCAACATGGCGTTG
TCAATGTACCTTTCAGCCACAGGACATTACAATCCATTCTTTCAGCCAAACTGGTCATACCTATAACT
ACAGCTCCAAATATCTCCTGGTCTGACCTCAGTGCCTGGAGTTGTTGAAATCTATACCAGTGGGAGTT
GGTCAGATCTATGGCTGTGATAATCCATGGACAGGGGGCATTTCCTGGGAGCCATCCTACTCTCCTCC
CCACTCATGTGCCTGCATGCTGCCATAGGATCATTGCTGGGCATAGCAGCGGGACTCAGTCTTTCAGCC
CCATTTGAGGACATCTACTTTGGACTCTGGGGTTTCAACAGCTCTCTGGCCTGCATTGCAATGGGAGGA
ATGTTTCATGGCGCTCACCTGGCAAACCCACCTCCTGGCTTTGGCTGTGCCCTGATTCAGGCCTATCTT
GGAGTCGGCATGGCAAACCTTTATGGCTGAGGTTGGATTGCCAGCTTGACCTGGCCCTTCTGTTTGCC
ACGCTATTGTTCTCATCATGACCACAAAAAATCCAACATCTACAAGATGCCCTCAGTAAAGTTACT
TATCCTGAAGAAAACCGCATCTTCTACCTGCAAGCCAAGAAAAGAATGGTGGAAAGCCCTTTGTGA
ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGAT
TACAAGGATGACGACGATAAGGTTTAAACGGCCGGC
```

Restriction Sites: Sgfl-Mlul



[View online »](#)

Plasmid Map:


ACCN: NM_001146036

Insert Size: 1170 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).

OTI Annotation: This TrueClone is provided through our Custom Cloning Process that includes sub-cloning into OriGene's pCMV6 vector and full sequencing to provide a non-variant match to the expected reference without frameshifts, and is delivered as lyophilized plasmid DNA.

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_001146036.2](#)

RefSeq Size: 4058 bp

RefSeq ORF: 1170 bp

Locus ID: 6563

UniProt ID: [Q13336](#)

Cytogenetics: 18q12.3

Protein Families: Transmembrane

MW: 42.5 kDa

Gene Summary: The protein encoded by this gene is a membrane transporter that mediates urea transport in erythrocytes. This gene forms the basis for the Kidd blood group system. [provided by RefSeq, Mar 2009]
Transcript Variant: This variant (3) differs in the 5' UTR and lacks an in-frame portion of the 5' coding sequence compared to variant 4. The resulting isoform (2) has a shorter N-terminus compared to isoform 1. Variants 2 and 3 both encode the same isoform (2). 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.