

Product datasheet for **SC337127**

KIRREL1 (NM_001286349) Human Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	KIRREL1 (NM_001286349) Human Untagged Clone
Tag:	Tag Free
Symbol:	KIRREL1
Synonyms:	KIRREL; NEPH1
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)



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Fully Sequenced ORF: >SC337127 representing NM_001286349.
 Blue=Insert sequence Red=Cloning site Green=Tag(s)

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GCTCGTTTAGTGAACCGTCAGAATTTTGAATACGACTCACTATAGGGCGGCCGGGAATTCGTGCGACTG
GATCCGGTACCGAGGAGATCTGCCGCCGCGATCGCC
ATGCTGAGCCTCCTCGTCTGGATCCTCACTCTCTCCGATACTTTCTCCCAAGTCCCCCAGAGGACACC
AGGATTGACGGAGGCCCTGTGATTCTACTGCAGGCAGGCACCCCCACAACCTCACATGCCGGGCCCTTC
AATGCGAAGCCTGCTGCCACCATCATCTGGTTCCGGGACGGGACGCAGCAGGAGGGCGCTGTGGCCAGC
ACGGAATTGCTGAAGGATGGGAAGAGGGAGACCACCGTGAGCCAACTGCTTATTAAACCCACGGACCTG
GACATAGGGCGTGTCTTCACTTGCCGAAGCATGAACGAAGCCATCCCTAGTGGAAGGAGACTTCCATC
GAGCTGGATGTGACCAACCTCTACAGTGACCTGTCCATTGAGCCACAGACGGTGACGAGGGTGAG
CGTGTGTCTTTACCTGCCAGGCCACAGCAACCCGAGATCTTGGGCTACAGGTGGGCCAAAGGGGGT
TTCTTGATTGAAGACGCCACGAGAGTCGCTATGAGACAAATGTGGATTATTCCTTTTTCACGGAGCCT
GTGCTTGTGAGGTTACAACAAAGTGGGAAGCACCAATGTCAGCACTTTAGTAAATGTCCACTTTGCT
CCCCGATTGTAGTTGACCCAAACCCACAACCACAGACATTGGCTCTGATGTGACCTTACCTGTGTC
TGGGTTGGGAATCCCCCCTCACTCTACCTGGACCAAAAAGGACTCAAATATGGTCCTGAGTAACAGC
AACCAGCTGCTGCTGAAGTCGGTGACTCAGGCAGACGCTGGCACCTACACCTGCCGGGCCATCGTGCT
CGAATCGGAGTGGCTGAGCGGGAGGTGCCGCTCTATGTGAACGGGCCCCCATCATCTCCAGTGAGGCA
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CGCATAGCATGGGCTGGAAGGAGAAGTCTTGGAGGTGGGGACCTGGAACGCTATACAGTGGAGAGG
ACCAACTCAGGCAGTGGGGTGTATCCACGCTCACCATCAACAATGTGATGGAGGCCGACTTTCAGACT
CACTACAACTGCACCGCTGGAACAGCTTCGGGCCAGGCACAGCCATCATCCAGTGGAAGAGCGAGAG
GTGTTACCTGTGGGCATCATAGCTGGGGCCACCATCGGCGCGAGCATCCTGCTCATCTTCTTCTCATC
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GACCTGCGCTGCGACACCATCGACACCCGGGAGGAGTATGAGATGAAGGACCCCAACATGGCTACTAC
AACGTGCGTGCCCATGAAGACCGCCCGTCTTCCAGGGCAGTGCTCTATGCTGACTACCGTGCCCTGGC
CCTGCCCGCTTCGACGGCCGCCCTCATCCGTCTCTCCACTCCAGCGGCTATGCCAGCTCAACACC
TATAGCCGGGGCCCTGCCTCTGACTATGGCCCTGAGCCACACCCCTGGCCCTGCTGCCACAGTGGC
ACTGACACAACCAGCCAGCTGTCTACGAGAACTATGAGAAGTTCAACTCCCATCCCTTCCCTGGGGCA
GCTGGGTACCCACCTACCGACTGGGCTACCCCAAGGCCACCTCTGGCCTGGAGCGGACCCCATAT
GAGGCGTATGACCCATTGGCAAGTACGCCACAGCCACTCGATTCTCTACACCTCCAGCACTCGGAC
TACGGCCAGCGATTCCAGCAGCGCATGCAGACTCACGTGATG
ACGCGTACGCGGCCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGAT
TACAAGGATGACGACGATAAGGTTTAAACGGCCGGC
  
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Restriction Sites: SgfI-MluI

ACCN: NM_001286349

Insert Size: 1974 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:	<u>NM_001286349.1</u>
RefSeq Size:	7526 bp
RefSeq ORF:	1974 bp
Locus ID:	55243
UniProt ID:	<u>Q96J84</u>
Cytogenetics:	1q23.1
Protein Families:	Transmembrane
MW:	72.7 kDa
Gene Summary:	NEPH1 is a member of the nephrin-like protein family, which includes NEPH2 (MIM 607761) and NEPH3 (MIM 607762). The cytoplasmic domains of these proteins interact with the C terminus of podocin (NPHS2; MIM 604766), and the genes are expressed in kidney podocytes, cells involved in ensuring size- and charge-selective ultrafiltration (Sellin et al., 2003 [PubMed 12424224]).[supplied by OMIM, Mar 2008] Transcript Variant: This variant (2) lacks two in-frame exons in the 5' coding region compared to variant 1. The encoded isoform (2) is shorter than 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.