

## **Product datasheet for SC200974**

## AKR1A1 (NM 153326) Human 3' UTR Clone

**Product data:** 

**Product Type:** 3' UTR Clones

**Product Name:** AKR1A1 (NM\_153326) Human 3' UTR Clone

Symbol: AKR1A1

Synonyms: ALDR1; ALR; ARM; DD3; HEL-S-6

Mammalian Cell

Selection:

Neomycin

**Vector:** pMirTarget (PS100062)

**ACCN:** NM\_153326

**Insert Size:** 123 bp

Insert Sequence: >SC200974 3'UTR clone of NM\_153326

The sequence shown below is from the reference sequence of NM\_153326. The complete

sequence of this clone may contain minor differences, such as SNPs.

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

CCTCTGTACCCCTTTAATGACCCGTACTGAGACCACAGCTTCTTGGCCTCCCTTCCAGCTCTGCAGCTA

ATGAGGTCCTGCCACAACGGAAAGAGGGAGTTAATAAAGCCATTGGAGCATCCA

**ACGCGT**AAGCGGCCGCGCATCTAGATTCGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

**Restriction Sites:** Sgfl-Mlul

**OTI Disclaimer:** Our molecular clone sequence data has been matched to the sequence identifier above as a

point of reference. Note that the complete sequence of this clone is largely the same as the

reference sequence but may contain minor differences, e.g., single nucleotide

polymorphisms (SNPs).

**Components:** The cDNA clone is shipped in a 2-D bar-coded Matrix tube as 10 ug dried plasmid DNA. The

package also includes 100 pmols of both the corresponding 5' and 3' vector primers in

separate vials.

**RefSeq:** <u>NM 153326.3</u>



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## AKR1A1 (NM\_153326) Human 3' UTR Clone - SC200974

Summary: This gene encodes a member of the aldo/keto reductase superfamily, which consists of more

than 40 known enzymes and proteins. This member, also known as aldehyde reductase, is involved in the reduction of biogenic and xenobiotic aldehydes and is present in virtually every tissue. Multiple alternatively spliced transcript variants of this gene exist, all encoding

the same protein. [provided by RefSeq, Jan 2011]

**Locus ID:** 10327

MW: 4.8