

## Product datasheet for **SC202639**

### **POLR1C (NM\_203290) Human 3' UTR Clone**

#### **Product data:**

<b>Product Type:</b>	3' UTR Clones
<b>Product Name:</b>	POLR1C (NM_203290) Human 3' UTR Clone
<b>Vector:</b>	pMirTarget (PS100062)
<b>Symbol:</b>	POLR1C
<b>Synonyms:</b>	AC40; HLD11; RPA5; RPA39; RPA40; RPAC1; RPC40; TCS3
<b>ACCN:</b>	NM_203290
<b>Insert Size:</b>	243 bp
<b>Insert Sequence:</b>	>SC202639 3'UTR clone of NM_203290 The sequence shown below is from the reference sequence of NM_203290. The complete sequence of this clone may contain minor differences, such as SNPs. <b>Blue</b> =Stop Codon <b>Red</b> =Cloning site  GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAAGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC GATGAAGTAGATGCGGTTTCAGATGGACTGAGCTTGGATGCTTCTGAGGCAAGCTGAAGCTTTGGGTTCT GACTGACCCACCCTACAGGACTGCTGAACAGAGAGCCAGTGTGACTAGGGATCCTGAGTTTTCTGGGA CAATTCCAGCTTTAATCAATACATTTTGTAAATGTGCCATAAAATGAGACTTTTTACGCCTTTATAAG GCCTTAGATGTAATAAACTCACCCAAACAAAAA <b>ACGCGT</b> AAGCGGCCGCGCATCTAGATTGAAAGAAATGACCGACCAAGCGACGCCAACCTGCCATCA CGAGATTCGATTCCACCGCCCTTCTATGAAAGG
<b>Restriction Sites:</b>	SgfI-MluI
<b>OTI Disclaimer:</b>	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).
<b>Components:</b>	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.
<b>RefSeq:</b>	<u><a href="#">NM_203290.4</a></u>



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**Summary:** The protein encoded by this gene is a subunit of both RNA polymerase I and RNA polymerase III complexes. The encoded protein is part of the Pol core element. Mutations in this gene have been associated with Treacher Collins syndrome (TCS) and hypomyelinating leukodystrophy 11. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Jan 2016]

**Locus ID:** 9533

**MW:** 9.2