

**Seattle Genova** 

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## **Laronidase In Vitro Transcribed mRNA-LNP**

Catalog Number:SG-MRNA-LNP-1905

Gene Name  Source  The OR prepare mRNA PEG:SM  Alternative names  Laronic  SPECIFICATIONS  Cap  5'-UTR  ORF  Laronic  3'-UTR  3' UTR  aminot encode  Poly(A) Tail  A 110-residue adenos  Modifications  N1-met  Neutral Lipid  Cholesterol  Lonizable Lipid  PEG-lipid  Heptad  8-oxool  Storage  PBS, ph	dase In Vitro Transcribed mRNA-LNP  L-iduronidase
Source The OR prepare mRNA PEG:SM  Alternative names  Laronic  SPECIFICATIONS  Cap  5'-untr optimiz  ORF  3'-UTR  3' UTR aminot encode  Poly(A) Tail  A 110-r residue adenos  Modifications  Neutral Lipid  Cholesterol  Lonizable Lipid  PEG-lipid  PEG-lipid  Storage  PBS, ph	
Alternative names  SPECIFICATIONS  Cap m7Gpp 5'-UTR 5'-untroptimiz  ORF Laronic  3'-UTR 3' UTR aminot encode  Poly(A) Tail A 110-residue adenos  Modifications N1-met  Neutral Lipid 1,2-dist  Cholesterol Cholest  Lonizable Lipid 1,2-dim (PEG20)  PEG-lipid Heptad  8-oxood  Storage -20 °C  Buffer PBS, ph	
SPECIFICATIONS  Cap m7Gpp 5'-UTR 5'-untroptimiz  ORF Laronic 3'-UTR 3' UTR aminot encode  Poly(A) Tail A 110-residue adenos  Modifications N1-met  Neutral Lipid 1,2-dist  Cholesterol Cholest  Lonizable Lipid 1,2-dim (PEG20  PEG-lipid Heptad 8-oxoor  Storage -20 °C  Buffer PBS, ph	RF of Laronidase was cloned in our IVT vector and mRNA was ed through in vitro transcription and purification. The purified was further encapsulated with LNP(DSPC:Cholesterol:DMG-M102).
Cap m7Gpp 5'-UTR 5'-untroptimiz ORF Laronic 3'-UTR 3' UTR aminot encode Poly(A) Tail A 110-r residue adenos Modifications N1-met Neutral Lipid 1,2-dist Cholesterol Cholest Lonizable Lipid 1,2-dim (PEG20 PEG-lipid Heptad 8-oxoor Storage -20 °C Buffer PBS, ph	lase
5'-UTR  5'-untroptimiz  ORF  Laronic  3'-UTR  3' UTR  aminot encode  Poly(A) Tail  A 110-r residue adenos  Modifications  N1-met  Cholesterol  Lonizable Lipid  1,2-dim (PEG20  PEG-lipid  Heptad 8-oxoor  Storage  -20 °C  Buffer  PBS, ph	
ORF Laronic 3'-UTR 3'-UTR 3'-UTR Poly(A) Tail A 110-r residue adenos Modifications N1-met Neutral Lipid Cholesterol Lonizable Lipid PEG-lipid Heptad 8-oxoor Storage PBS, ph	PN
3'-UTR aminot encode  Poly(A) Tail  A 110-r residue adenos  Modifications  N1-met  Neutral Lipid  Cholesterol  Lonizable Lipid  PEG-lipid  Heptad 8-oxoor  Storage  -20 °C  Buffer  PBS, ph	ranslated region derived from human alpha-globin RNA with an zed Kozak sequence
aminot encode  Poly(A) Tail  A 110-r residue adenos  Modifications  N1-met  Neutral Lipid  Cholesterol  Lonizable Lipid  PEG-lipid  Heptad 8-oxoor  Storage  PBS, ph	łase
residue adenos  Modifications N1-met  Neutral Lipid Cholesterol Cholest  Lonizable Lipid PEG-lipid Heptad 8-oxood Storage -20 °C  Buffer PBS, ph	comprising two sequence elements derived from the terminal enhancer of split (AES) mRNA and the mitochondrial ed 12S ribosomal RNA
Neutral Lipid 1,2-dist Cholesterol Cholest Lonizable Lipid 1,2-dim (PEG20 PEG-lipid Heptad 8-oxool Storage -20 °C Buffer PBS, ph	nucleotide poly(A)-tail consisting of a stretch of 30 adenosine es, followed by a 10-nucleotide linker sequence and another 70 sine residues.
Cholesterol Cholest Lonizable Lipid 1,2-dim (PEG20 PEG-lipid Heptad 8-oxool Storage -20 °C Buffer PBS, ph	thyl-pseudouridine
Lonizable Lipid  1,2-dim (PEG20  PEG-lipid  Heptad 8-oxoo  Storage  -20 °C  Buffer  PBS, ph	tearoyl-sn-glycero-3-phosphocholine (DSPC)
PEG-lipid Heptad 8-oxoo Storage -20 °C Buffer PBS, ph	terol
Storage -20 °C  Buffer PBS, ph	nyristoyl-rac-glycero-3-methoxypolyethylene glycol-2000 000-DMG)
Buffer PBS, ph	lecan-9-yl 8-((2-hydroxyethyl)(8-(nonyloxy)— ctyl)amino)octanoate)(SM-102)
Cryoprotectant Trehald	ose
BACKGROUND	
Gene Accession	
Gene Alias Laronio	lase



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polymorphic form of human a-L-iduronidase. It contains 6 N-linked oligosaccharide modification sites. Laronidase catalyses the hydrolysis of terminal alpha-L-iduronic acid residues of dermatan sulfate and heparin sulfate.

## Background

Human recombinant alpha-L-iduronidase, 628 residues (mature form). Laronidase is a glycoprotein with a molecular weight of approximately 83 kD. The predicted amino acid sequence of the recombinant form, as well as the nucleotide sequence that encodes it, are identical to a