

## Nucleocapsid protein gene of Junin arenavirus (cDNA sequence)

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The Arenavirus S RNA codes for the major structural polypeptides: the nucleocapsid protein, N, and the glycoproteins G1 and G2, that are derived from an intracellular precursor protein, GPC.

We have cloned and sequenced the entire coding region for the N gene of Junin arenavirus, the etiologic agent of Argentine hemorrhagic fever. The cDNA sequence was determined by a dideoxy procedure (1) using a modified T7 DNA polymerase (Sequenase™ · USB, Cleveland, Ohio, USA) and is shown as the viral-complementary strand.

The upstream non-coding sequence comprises 81 nucleotides, and the open reading frame codes for the N polypeptide, deduced to be 564 aminoacids long (mol. wt. 63,033 Da). Nucleotide and aminoacid sequences exhibit varying degrees of identity when compared to homologous regions of other arenavirus sequences. The comparison with Tacaribe virus (2), an arenavirus with a close antigenic relation to Junin, shows a 76% identity in aminoacid sequence, whereas with N proteins of other arenaviruses the identity ranges from 46 to 52%.

	M	A	H	S	K	E	V	P	S	F	R	W	T	13
CGCACAGCTGGATCCTAGGTCAACACTTCACTGAGCAATTTCCTAGATCAAAAACCACAGATCTTCGGCATGCCACACTCCAAAGAGGTCCAAGCTTTAGATGGACT	120													
G L S Q F T Q T V K S D V L K Q S L R R D A K L I A D S I D F N Q V A Q V Q R A	53													
GGTTTGAGTCAATTCTACAGACTGTCAAGTCGGATGTCGAAGCAAGCTTAAGGGAGAGATGCCAACTTAATGGTCAACGATCTGACTCAACCAAGTGGCAAGGGGCA	240													
L R K T K R G E E D L N K R D L N K E V D R L M H S R V Q R N T V F K A G D	93													
CTCAGAAAGACTAAAAAGGGGGAAGAGACCTCAATAAGTGAGGGACCTGAATAAGGAGGTGACAGACTCATGCATGAGGTGTCACAGAAACAGCTTCAAGGGGAGAGAT	360													
L G R V E R M E L A S G L G N L K T K F R R A E T G S O G V Y H G N L S Q S Q L	133													
CTGGGGAGGGGTTGAGGGATGAGGTGGCGCTCTGCCCTGGGAATTAAACACAGGTTAGAGGAGCAGACAGGCTCTCAAGGGTACATGGTAACCTTGTCAGTCACAAGCT	480													
A K R E I L R T L G F Q R G T G G G N G V V R H D V K D P S K L N H Q F G S	173													
GCCAAAGAGTCAAGAGATAATGAGAACACTGGGATTCAACAGCAAGGAACGGGGGAAATGGTGGTGGAGGGATGGGATGTTAAAGATCTCCTAAACATCAGTTGGTCT	600													
V P A L T I A C N T V O G G E T M N S V I Q A L T S L G L Y L T V K Y P P H L S D	213													
GTCCTGCGATTCGACAATTCGATGACTGTCAAGGAGGTGAGACAATGAACAGTTCATACAGGCTTAAACCTCTTCGGCTCTATACACTGTGAAGATACAAACTTGAGTAC	720													
L D R L T Q E H D C L Q I V T K D E S S I N I S G Y H F S L S A A V K A G A S I	253													
CTTGACAGATGACTGAGGAACTGACTGCTCCAGAGTGTGACTAAGGATGAAAGCTCCATCAACATTCTGGCTCACACTTGTGACTCTGAGTCAAGCTGAGTAAGGGCTGGCATCCATT	840													
L D D G H M L E T I R V T P D N F S S L I K S T I P V R R E G M F I D E K P G	293													
CTTGATGATGGAAATAATGTTGAGAACATACTAGAGTCACCCAGATCAACTTTCTCCCTCATATAATCAACCATTCAGGTTAACGAGAACAGGATGTTTATTGATGAGAACAGG	960													
M R R P Y E N E L L Y K L C L S G D G W P Y I G S R S Q I I G R S W D N T S I D L	333													
AAATAGAACCTTATGAAACCTTCATGACTACATGGGCTTCTGGCCATGGTGGCTTATATGGTTCAGATCACAAATCATAGGGAGGTATGGACACACAAGTATTGATCTG	1080													
T R K P V A G P R Q K E P K N G O N L R A L N T E Q A V I R E A V G K L D P	373													
ACAAGGAAACCGAGTGGCTGGACCTAGACAACCGGAAAGAGACGGTCAGAATTGAGATTGGCTAACTTGAGACAGATAACAAGAGCTGTCAGAGAGGGAGTGGGAAACCTGACCCCC	1200													
T H T L V W L D I E G P A T D P V P E H A L F Q P A G S K Y I N C F R K P H D E K G	413													
ACCAATACCTTCTGGACATTCGACATTGAGGGACAGCTACTGCCCTGTGAGATGGCTGTTCCAGGCTGAGGTTAGCAAGTATTCGCTTCAGAAAACACATGAGAACAGG	1320													
F K M G S R H S N G I L M K D I E D A M P G V L S Y V I G L L P P D H V V T T Q	453													
TTTAAAAATGGTAGCAGACACTCTACGGCATCTTAATGAAAGACATGAAACATGCCAGGAGTCTTACGTTAGCTGTTGGCTTCGGCTCCGGACATGGTGTGACCAACTCAA	1440													
G S D D I R K F D L H G R R D L V D P R V L T S E Q A R Q F D Q Q V W E K F	493													
GGGTCGGATGACATCAGAAAGATTGTTGACCTCATGGAAAGAGACATCTAACCTGTTAGCTTACATCAAGACAACGAGGAGTCAGCACACAGGCTGGGAGAAATT	1560													
G H L C K H H N G V V V S K K K R D K D A P F K L L A S S E P N C A L L D C I M F	533													
GGCCACCTATGAAACATCACAACTGAGTGGTGTGAGCAAAAGAGACATAAGGATGCTCCCTTAAAGTGGCTCCAGTGGCCACACTGTGCTCTGCTAGACTGCATAATGTT	1680													
Q S V L D G K L Y E E E L T P L L P S S L L F L P K A Y A L *	564													
CACTGAGTGGATGGAAAGCTATAGAAGAAACTACACCTCTATACCCACGAGCTGGCTGCTCCTCCGAAGGCAGCTATGACTGTAAGCAGTGGCTCCAGTCCGGGCC	1800													

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**References:**

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