



Cloning and sequence analysis of the LOC339524 gene in Sprague-Dawley rats

Z.H. Long¹, H. Li¹, F. Chen¹ and L.Y. Zou²

¹Department of Anesthesiology, Xinqiao Hospital, Third Military Medical University, Chongqing, China

²College of Basic Medical Sciences, Third Military Medical University, Chongqing, China

Corresponding author: H. Li

E-mail: Lh78553@163.com

Genet. Mol. Res. 14 (4): 16577-16584 (2015)

Received August 5, 2015

Accepted October 25, 2015

Published December 11, 2015

DOI <http://dx.doi.org/10.4238/2015.December.11.4>

ABSTRACT. We cloned the LOC339524 gene in Sprague-Dawley (SD) rats and analyzed the structure and function of the protein encoded by it. Based on the known human LOC339524 gene sequences, the full-length coding sequence of the LOC339524 gene in SD rats was cloned and amplified by the polymerase chain reaction using the complementary DNA of SD rats as a template. Bioinformatics analysis showed that the length of the cloned LOC339524 gene (GenBank accession No. KM224520) was 831 bp and it encoded a deduced protein of 276 amino acids. Sequence analysis revealed that the coded protein was identical to that produced in humans and its functional domain was located in the 138-236 amino acid fragments, a proline-rich region. Our results suggest that the encoded protein may be a significant regulator of the inflammatory response and may provide sufficient information to justify an in-depth investigation of the role of the LOC339524 gene.

Key words: Sprague-Dawley rat (*Rattus norvegicus*); LOC339524 gene; Clone; Bioinformatics