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Tumor necrosis factor receptor-associated factor 6 (TRAF6) from mud crab participates in antilipopolysaccharide factors (ALFs) gene expression

umor necrosis factor receptor-associated factor 6 (TRAF6) is a cytoplasm key signal adapter protein that mediates signals activated by tumor necrosis factor receptor (TNFR) superfamily and the Interleukin-1 receptor/Toll-like receptor (IL-1/ TLR) superfamily. The full-length 2492 bp Scylla paramamosain TRAF6 (Sp-TRAF6) contains a 1800 bp of open reading frame (ORF) encoding 598 amino acids, including an N-terminal RING-type zinc finger, two TRAF-type zinc fingers and a conserved C-terminal meprin and TRAF homology (MATH) domain. Multiple alignment analysis shows that the putative amino acid sequence of Sp-TRAF6 has highest identity with Pt-TRAF6 (KP341006) from Portunus trituberculatus at 88%, while the similarity of other crustacea sequences was 54-55%. RT-PCR results indicated that the Sp-TRAF6 transcripts were predominantly expressed in the hepatopancreas and stomach, whereas it was barely detected in the heart and hemocytes in our study. Further, Sp-TRAF6 transcritripts were significantly up-regulated after immune challenge with Vibrio parahemolyticus or LPS. Our previous study had characterized two novel anti-lipopolysaccharide factor isoforms from S.paramamosain (SpALF5 and SpALF6). Both of them contain a conserved LPS-binding domain with two conservative cysteine residues, which is critical for their antimicrobial function. The vitro binding and antimicrobial activity assays indicated that the recombinant SpALF5 and SpALF6 protein generated from prokaryotic expression system showed a varying degree of binding activity towards bacteria and fungus, and exhibited a broad spectrum of antimicrobial activities against Gram-positive, Gram-negative bacterium and fungi. Therefore, six ALF isoforms from mud crab had been reported up to now. To investigate Sp-TRAF6 activating SpALFs gene expression, RNA interference assay was carrried out to examine the mRNA level of six SpALFs after silencing Sp-TRAF6 gene. The results showed that silencing Sp-TRAF6 gene could inhibit SpALF1, SpALF2, SpALF5 and SpALF6 expression in hemocytes, while SpALF1, SpALF3, SpALF4, SpALF5 and SpALF6 in hepatopancreas. Taken together, the acute-phase response to immune challenges and the inhibition of SpALFs gene expression indicate that Sp-TRAF6 plays an important role in host defense against pathogen invasion via regulation of ALF gene expression in S. Paramamosain.

Biography

Shengkang Li has completed his PhD from Sun Yat-sen University and Post-doctoral studies from IFREMER Centre de Nantes, France. He is the Principle Investigator of marine micro-organisms research group in Marine Biology Institute, Shantou University. He has published more than 30 papers in reputed journals and has been serving as a reviewer for many reputed journals.

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