



Genetic Insights: Applications of HSP90 in Chlorella Biotechnology

Williams Anderson*

Department of Genetics and Microbiology, Keele University, Keele, United Kingdom

DESCRIPTION

Chlorella vulgaris, a single-celled green microalga, has emerged as a potential candidate for various biotechnological applications, including biofuel production, wastewater treatment, and as a source of valuable biomolecules. In recent scientific trial, researchers have explored the complexity of molecular machinery of *Chlorella vulgaris*, focusing on the cloning and expression of a cytosolic HSP90 gene. Heat Shock Protein 90 (HSP90) is a molecular chaperone with pivotal roles in cellular homeostasis, and understanding its function in *Chlorella vulgaris* facilitates an approach for exploiting the potential of this microalga.

Understanding the role of HSP90 gene

HSP90, a highly conserved molecular chaperone, plays a potential role in cellular proteostasis by aiding in the folding, stabilization, and degradation of client proteins. It is involved in diverse cellular processes, including signal transduction, cell cycle control, and response to environmental stress. In the context of microalgae like *Chlorella vulgaris*, the role of HSP90 becomes particularly intriguing as it adapts to varying environmental conditions, including temperature fluctuations and nutrient availability.

Cloning the cytosolic HSP90 gene

In the initial stage of understanding *Chlorella vulgaris*'s HSP90 involves cloning its cytosolic gene. This process involves isolation of gene of interest from the microalga's genomic DNA, amplifying it using Polymerase Chain Reaction (PCR), and then inserting it into a suitable vector for further analysis and manipulation. The cloned gene provides researchers with a specific target for studying the structure, regulation, and function of the cytosolic HSP90 in *Chlorella vulgaris*.

Expression of the cloned gene

Once the cytosolic HSP90 gene is successfully cloned, the next critical step is its expression within *Chlorella vulgaris* cells.

Expression involves translating the genetic information encoded in the gene into functional HSP90 proteins. This can be achieved through various expression systems, such as plasmids or viral vectors, introduced into the microalga. The expression of the cloned HSP90 gene allows researchers to investigate its dynamics, cellular localization, and response to different environmental stimuli.

Implications for cellular homeostasis

The presence and expression of a cytosolic HSP90 gene in *Chlorella vulgaris* suggest its involvement in maintaining cellular homeostasis under varying conditions. As a molecular chaperone, HSP90 assists in the correct folding of proteins, preventing the formation of non-functional or misfolded structures. This is particularly potential in microalgae, which experience fluctuations in temperature, light, and nutrient availability. Understanding how *Chlorella vulgaris* utilizes HSP90 for cellular stability provides valuable insights into its adaptability and resilience.

Environmental stress response

Chlorella vulgaris, like other microorganisms, challenged by environmental stressors such as heat, cold, and nutrient deprivation. The expression of the cytosolic HSP90 gene is expected to be complicatedly linked to the microalga's response to such stress conditions. Researchers can investigate the upregulation of HSP90 in response to heat shock or other environmental stressors, focusing on the adaptive mechanisms employed by *Chlorella vulgaris* to thrive in diverse ecological functions.

Biotechnological applications

Beyond the field of fundamental research, the cloning and expression of a cytosolic HSP90 gene in *Chlorella vulgaris* hold potential for biotechnological applications. Microalgae are increasingly recognized as sustainable platforms for the production of biofuels, high-value compounds, and pharmaceuticals. By understanding the role of HSP90, researchers

Correspondence to: Williams Anderson, Department of Genetics and Microbiology, Keele University, Keele, United Kingdom, E-mail: Wandersons@gmail.com

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can potentially manipulate its expression to enhance the microalga's productivity, stress tolerance, and overall performance in biotechnological processes.

Optimizing biofuel production

Chlorella vulgaris has gained attention as a potential feedstock for biofuel production due to its high lipid content. The efficient production of lipids, which can be converted into biofuels, relies on the optimal functioning of cellular processes. Manipulating the expression of *HSP90* may provide a means to enhance lipid biosynthesis and improve the overall efficiency of *Chlorella vulgaris* as a biofuel feedstock.

Enhancing biomolecule production

Chlorella vulgaris is also known for its ability to accumulate various biomolecules with potential industrial and pharmaceutical

applications, such as pigments, antioxidants, and proteins. The controlled expression of *HSP90* may be a key factor in optimizing the production of these biomolecules, ensuring their correct folding and functionality.

The cloning and expression of a cytosolic *HSP90* gene in *Chlorella vulgaris* represent a significant step in understanding the molecular mechanisms underlying this microalga's cellular processes. This research not only contributes to our knowledge of fundamental biology but also holds potential for applications in biofuel production, biomolecule synthesis, and environmental biotechnology. As scientists continue to explore the encoded genomes of microorganisms like *Chlorella vulgaris*, the potential for sustainable and innovative solutions to critical global challenges becomes increasingly apparent.