**سیویلیکا - ناشر تخصصی مقالات کنفرانس ها و ژورنال ها** گواهی ثبت مقاله در سیویلیکا CIVILICA.com

## عنوان مقاله:

3D structure analysis of VhhP2 Protein in Vibrio harveyi

محل انتشار: چهارمین کنفرانس بین المللی علوم و مهندسی (سال: 1395)

تعداد صفحات اصل مقاله: 15

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### خلاصه مقاله:

Vibrio harveyi is one of the significant aquatic pathogens founded predominately in marinehabitats. The gramnegative bacterium can cause infection in a broad range of marine vertebratesand invertebrates, involving fish, shrimp, lobster, and mollusk. Until now, the traditional formsof vaccines against V. harveyi have been produced including killed whole bacterial cells (mostly)and extracellular products (in part). In addition to these vaccines, currently a number of proteinbased(in fact recombinant subunit) vaccines have been manufactures and are shown to beeffective against some V. harveyi starins. The gene of vhhP2 belonged a pathogenic V. harveyistrain isolated from diseased fish has been identified and discovered that vhhP2 is widely distributed in V. harveyi strains of various geographical locations and sources. The present studywas designed to in silico resolving the major obstacles in the control or in prevention of Vibrioharveyi infections. We exploited bioinformatic tools to better understanding and characterizingthe vhhP2 3Dstructure and select appropriate regions as effective B cell epitops

**کلمات کلیدی:** Vibrio harveyi, vaccine, vhhP2, 3Dstructure

# لینک ثابت مقاله در پایگاه سیویلیکا:

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