



Biomarkers in marine ecosystems monitoring

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An increasing awareness of the importance of protecting, conserving or enhancing marine ecosystems has risen over the past decades as a consequence of the growing human impacts on marine areas and the possible detrimental effects for the environment and human health.

The European Union Marine Strategy Framework Directive (MSFD) (Directive 2008/56/EC) (1) aiming to achieve or maintain 'Good Environmental Status' (GES) in EU waters by 2020 evidences the need to assess the status of marine areas and to develop new monitoring strategies and approaches to quantify impacts of contaminants in marine ecosystems (2-4).

Recent marine environmental monitoring programmes aim to assess the level of contamination in marine areas evaluating the effects of different stressors, including contaminants, on marine organisms, thus linking contaminants with the health of the ecosystem in term of biological effect, and biomarkers, defined as measurable alterations occurring at molecular, cellular, and physiological levels in response to environmental stressors, have been recognized as remarkable tools for the biological effect assessment in marine environmental monitoring. Several biomarkers have been evaluated in various sentinel organisms in response to different contaminants and have also been adopted in the framework of international biomonitoring programs (5-7) to study the quality of the aquatic environment.

Among the sentinel organisms, mussels (*Mytilus* spp.) have long been utilized as a bioindicator for biomonitoring of marine pollution (8-10) and currently the investigations on pollutant responsive biomarkers in mussels count 10% of all reported biomarker studies (11).

Through the years, the studies mainly conducted in strictly controlled laboratory conditions, to assess the impacts of different environmental stressors on the biological response in mussels, have provided suitable sets of biomarkers with a great potential for biomonitoring of marine ecosystems.

Since the advancements in genomics have been applied in marine sciences a boost promoting biomarker discovery has come from our knowledge of the molecu-

lar mechanisms underlying the physiological response of marine organisms to environmental stressors, including chemicals and emerging contaminants, thereby giving new valuable tools for assessing marine health status (12). Moreover, it is becoming increasingly evident that distantly related species share common response pathways to the same environmental stimuli and biomarkers early identified in response to one stressor are actually involved in the response to multiple environmental stressors (12,13).

Transcriptomic approaches contributed to the identification of unknown hypoxia-responsive biomarkers in mussels *Mytilus galloprovincialis* experimentally subjected to air exposure (14) that have effectively applied in ecotoxicological studies of natural marine areas (15) and are likely to be used in future environmental monitoring studies of hypoxic areas that are expected to increase worldwide.

Multi-biomarker approaches have been successfully applied for assessing the impacts of climate change and anthropogenic contaminants on aquatic organisms and demonstrated to be valuable tools not only to reveal the presence of stressor(s) but also to recognize regions of reduced ecosystem health. In particular, the application of a multi-biomarker panel resulted effective for assessing the biological effects of petrochemical contamination on the health status of mussels *Mytilus galloprovincialis* (11,15).

Despite the relationships between exposure to contaminant(s) and biological responses in mussels have been widely addressed, at least for persistent organic pollutants (POPs) and metals (Cu, Hg, Pb, and Zn) pollution, great attention has recently focused on the contaminants of emerging concern including pharmaceuticals and personal care products, microplastics and nanoparticles that are currently recognized as environmental threats in the aquatic environment (16-20). The few investigations on their effects in marine organisms, evidenced several responses at molecular level, physiological performance and organisms health depending on the species under study, and both of kind and concentration of contaminant.

Noteworthy, the rapid development of various omic

methodologies, namely genomics, transcriptomics, metabolomics and proteomics together with the more recent epigenomics, and their successful application in pollution biomonitoring supported the discovery of new biomarkers, thereby providing novel approaches integrating monitoring of chemicals and their biological effects on sentinel species (21).

Future studies should focus on multidisciplinary approach (genomics, proteomics and metabolomics) to understand the synergetic effects of multiple environmental stressors, especially emerging pollutants under field conditions, on marine organisms and to reveal the complexity of anthropogenic impacts on aquatic environment.

Scientific evidences should therefore support the active role of the governments in rising public awareness on environmental concerns and in developing new and efficient strategies to achieve marine ecosystem health.

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