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Evaluation of Multivariate Biomarker Indexes Application in Ecotoxicity Tests with Marine Diatoms Exposed to Emerging Contaminants

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Featured Application: Evaluation of oxidative stress indexes for ecotoxicological evaluation of marine diatoms exposed to emerging contaminants.

Abstract: Worldwide anthropogenic activities result in the production and release of potentially damaging toxic pollutants into ecosystems, thereby jeopardizing their health and continuity. Research studies and biomonitoring programs attend to this emerging problematic by applying and developing statistically relevant indexes that integrate complex biomarker response data to provide an holistic approach reflecting toxically induced alterations at the organism's or population level. Ultimately, indexes allow simple result communication, enhancing policy makers understanding, so contributing for better resource and environmental managing policies. In this study three indexes, the Integrated Biomarker Response index (IBR), the Bioeffects Assessment Index (BAI) and the Principal Components Analysis (PCA), were evaluated for their sensitivity in revealing toxically induced stress patterns in cells of the diatom *Phaeodactylum tricornerutum* under contaminant exposure. The set of biomarkers selected for index construction comprise the anti-oxidant enzymes APX, CAT and SOD, and the lipid peroxidation marker TBARS. Several significant inverse correlations with the concentration gradients applied, was noticed for all indexes, though, the IBR excels for its reliability in delivering statistically significant dose-response patterns for four out of the five compounds tested.

Keywords: antioxidant enzymes; pharmaceutical residues; pesticides; detergents; integrative indexes

1. Introduction

Daily worldwide spread use of pharmaceuticals, pesticides and detergents, that endangering the health of natural aquatic ecosystems, entails major concern [1,2]. These emerging contaminants have serious implications on the organisms living in aquatic environments, at different biological levels, leading to biochemical, physiological, metabolic, individual and population-level effects[3,4]. These impacts are often addressed from the biochemical point of view, as a starting point for understanding the early effects of these contaminants in the biota, using a collection of biomarkers [5–7]. These biomarker responses are used as indicators of health status, which are already considered at international environmental policies [8]. The World Health Organization (WHO) defines biomarkers as biological measurements at the biochemical, cellular or molecular level that

result from the interaction between an organism and environmental chemical, physical or biological agents [9].

The applicability of subcellular biomarkers to address pollutants in aquatic ecosystems is currently widely studied. Biomarkers of exposure such as antioxidant systems, for their sensitivity, are widely used within the scope of toxicological studies [10,11]. As chemical pollutants induce oxidative damages to biological systems, the activity levels of antioxidant enzymes, like superoxide dismutase (SOD) and catalase (CAT) - which are vitally intricated in the detoxification of reactive oxygen species (ROS) and reducing oxidative stress - can be evaluated and construed [7,10]. Likewise, biomarkers of lipid peroxidation (LPO) are vastly investigated [10], especially through the thiobarbituric acid reactive substances (TBARS) test, that measures the levels of secondary lipid oxidation products, such as malondialdehyde (MDA) formation [12]. This set of biomarkers provides a standing of the overall oxidative stress state of the cells under a certain compound exposure at a specific dose [13].

While from the scientific point of view, the mode of action of these antioxidant processes and how they are affected by different emerging contaminants is of utmost importance, these biochemical results are often difficult to communicate to the non-scientific community such as stakeholders and decision makers [14,15]. The pressing need for methodologies and information that can effectively connect researchers and environmental managers, linked to international environmental frameworks, led to the development of a variety of indexes that can easily communicate scientific results to stakeholders, as applicable tools for decision making in environmental managing [16–18]. An array of scoring indexes emerged in the past years [16], thus adopting an ecologically transversal approach is of greater importance in order to develop meaningful community management tools towards addressing contaminants of emerging concern [5,19,20]. Therefore, following biochemical evaluations, a new index development must reflect accurately the biological status, follow suitable validation and consider results communication, so that management entities can approach similarly new environmental matters within the scope of policy development [16,17]. As above mentioned several approaches have been developed to translate vast arrays of biochemical data into numeric indexes, such as the Integrated Biomarker Response index (IBR) [5], the Bioeffect Assessment Index (BAI) [19] and the Principal Component Analysis Scoring based index (denoted as PCA-index hereafter) [20,21]. These indexes are based on different approaches to the same biochemical data. The IBR index uses a normalization of the biochemical data to overcome the differences of magnitude between the different biochemical traits it integrates, and requires the expert judgment or previous experience from the operator for defining how a certain biochemical trait is altered (enhanced or inhibited) by a given compound [5]. The BAI approach divides each biochemical trait at each exposure level into statistical cohorts, considering its quartile distribution, and scoring each biochemical trait accordingly [19]. The PCA-index uses the values obtained from the PCA plot of the biochemical traits, to normalize the values and weight each variable according to its importance for the PCA-model [20,21]. Thus, these three indexes range from univariate (BAI) to multivariate approaches (IBR and PCA-based indexes), with or without operator intervention (IBR and PCA-based respectively), and a comparative analysis allows to assess which approach is more suitable to be applied in emerging contaminants ecotoxicological studies.

In the present work, the suitability of three different multivariate indexes (BAI, IBR, and PCA-based indexes) was tested through the application of previously obtained oxidative stress biomarker data, using marine diatoms as study model. The impacts of pharmaceuticals (propranolol [22], fluoxetine [23], ibuprofen [24], detergents (SDS), and herbicides (glyphosate [15]) on primary production and physiological fitness of the marine diatom *Phaeodactylum tricornerutum* were used to understand the suitability of multivariate indexes in depicting dose responses and contaminant type, as well as to evaluate the performance and accuracy of each approach for future ecotoxicological trials. Summarily, we intended to produce an index that allows to better communicate toxicity results, providing an easy-to-understand framework, accessible to the managers and the general public.

2. Materials and Methods

2.1. Diatom exposure trials and biomarker analysis

Data regarding growth and biomarker activity/concentration was collected from previous published works [15,23–25, 43]. *Phaeodactylum tricornutum* Bohlin (Bacillariophyceae; strain IO 108–01, Instituto Português do Mar e da Atmosfera (IPMA) axenic cell cultures (kept under asexual reproduction circumstances) were placed to grow in f/2 medium [25], under continuous aeration in a phytoclimatic cabinet, at 18 °C, set with a 14/10 h day/night photoperiod (RGB 1:1:1, maximum PAR 80 $\mu\text{mol photons m}^{-2} \text{s}^{-1}$), a sine function to mimic sunrise and sunset, and maximum light intensity at noon, set to reproduce natural conditions [26]. Exposure tests were performed according to the Organization for Economic Cooperation and Development (OECD) guidelines for microalgae assays [27], as described in [15,23–25, 43]. Briefly cultures were exposed for 48 h. Final concentrations were chosen targeting to cover a concentration range not only environmentally relevant concentrations found in the literature, but also concentrations recognized to have substantial biologic consequences in *P. tricornutum* [28,29]. Growth inhibition concentration (IC50) was computed according to the OECD recommendations for microalgae inhibition test [27]. Biomarker analysis was performed according to standard spectrophotometric procedures as described in [15,23–25, 43]. Briefly catalase (CAT), ascorbate peroxidase (APx) and superoxide dismutase (SOD) activities were assessed by spectrophotometric methodologies employing specific substrates as previously described [30–32]. Lipid peroxidation products were evaluated spectrophotometrically [33], using trichloroacetic acid extraction prior to the reaction with thiobarbituric acid.

2.3. Multivariate index calculation

2.3.1. Integrated Biomarker Response (IBR) Index

The IBR index was calculated for each tested compound according to Beliaeff and Burgeot (2002), posteriorly adapted by Broeg and Lehtonen (2006). Briefly, it was calculated by summing up triangular star plot areas calculated for each two neighbouring variables. To calculate IBR integrating all biomarkers, the general mean (m) and the standard deviation (s) of all data (including concentrations) regarding a given biomarker was calculated, followed by standardization to obtain Y :

$$Y = \frac{X - m}{s}$$

where X is the mean value for the biomarker at a given concentration. Then Z was calculated using $Z = -Y$ or $Z = Y$, in the case of a biological effect corresponding respectively to an inhibition or a stimulation and it was determined using the slope of the biomarker activity/concentration against the exogenous concentration applied. Positive slopes corresponded to a stimulation, while negative slopes corresponded to an inhibition. Regarding the biological effect, biomarkers can either increase or decrease depending on the type and concentration of the compound exposure, but also vary among organisms.

Subsequently, the score (S) was calculated as

$$S = Z + |\text{Min}|$$

where, $S \geq 0$ and $|\text{Min}|$ is the absolute value for the minimum value for all calculated Y for a given biomarker at all measurements made (again all concentrations considered). Star plots were then used to display Score results (S) and to calculate the IBR index as:

$$IBR = \sum_{i=15}^n A_i$$

being A_i the area between two consecutive clockwise scores in a given star plot:

$$A_i = \frac{S_i}{2} \sin \beta (S_i \cos \beta + S_{i+1} \sin \beta)$$

$$\beta = \tan^{-1} \frac{S_{i+1} \sin \alpha}{S_i - S_{i+1} \cos \alpha}$$

where S_i and S_{i+1} are two consecutive clockwise scores of a given star plot; n the number of biomarkers used and α :

$$\alpha = \frac{2\pi}{n}$$

2.3.2. Principal Component Analysis (PCA) -based Index

To assess which variables were more suitable for the index elaboration and implementation, a previously successfully tested statistical approach was used [20,21,35]. This was performed independently for each compound. Principal Component Analysis (PCA) was performed in order to select the appropriate measured parameters to integrate the index. For this selection, the five variables with the higher weighing factors from the factor axis with the higher explanatory value percentage were chosen [21]. PCA was performed after data normalization and transformation, using PRIMER 6 [36]. The PCA based index is defined as:

$$PCA - based\ index = \sum W_i E_i$$

Where W is the PCA weighing factor of the PCA selected variable and E its respective score. The scores were normalized using a sigmoidal equation limited from 1 to 0, as follows [21,35,37]:

$$E = \frac{a}{1 + x/x_0^b}$$

Where x is the variable verified value, a is the maximum score of the variable (in this case 0.535 so that the final index has 1 as its maximum value), x_0 is the average value of the variable and b is the value of the slope of the equation. The slope was defined as -2.5 for the better-adjusted curve tending to 1 for all the variables proposed.

2.3.3. Bioeffect Assessment Index (BAI)

Index variables are assigned numerical values based on the degree or severity of damage of an organ or tissue caused by environmental stressors [19]. The BAI was designed as an index for the assessment of the multifactorial contamination situation of coastal areas. Thus, it includes only biomarkers of general toxicity [19]. To assess the individual values of each biomarker within the BAI stages, biomarker population quartils were determined. The index value of biomarker ($Biomarker_{index}$) was attributed depending on the quartil interval of each value:

- Values > 3rd quartil: 40
- Values between 3rd and 2nd quartils: 30

- Values between 2nd and 1st quartils: 20
- Values < 1st quartil: 10

The BAI value results from:

$$BAI = \sum Biomarker_{index}^{1/n}$$

Where n is the number of biomarkers used for index calculation.

2.4. Statistical analysis

Due to the lack of normality and homogeneity of variances of the attained data, Kruskal–Wallis pairwise comparisons between different sample groups were used to disclose univariate statistical differences. Spearman correlation tests were computed to assess possible dose–response relationships between the external concentrations applied and the evaluated biomarkers. Both Kruskal–Wallis and Spearman tests were computed using Statistica software (StataSoft, version 12.5.192.7). Statistical significance was considered at $p < 0.05$. A multivariate approach was additionally used to test for dissimilarities in the whole oxidative stress data package [23,38,39]. Canonical analysis of principle (CAP) coordinates, using Euclidean distances, were assessed in a canonical plot using the dissimilarities concerning the biomarker considered traits while performing a cross-validation step and evaluating the distribution effectiveness into the different treatment sets. This multivariate approach is unsensitive to heterogeneous data and commonly employed to compare different sample groups using the intrinsic characteristics of each sample group (metabolic traits) [38–41]. Multivariate statistical analyses were accomplished using Primer 6 software (version 6.1.13, Plymouth, UK) [36].

3. Results

3.1. Diatom growth and ecotoxicological parameters (IC₅₀)

Data on the exposure concentration and toxicity of five contaminants that inhibit the growth of the marine diatom *P. tricornutum* were obtained from previous studies [15,23–25, 43] and are presented at Table 1. According to the IC₅₀ values, which refer to the exposure concentration capable to inhibit half the maximum growth in this marine diatom, SDS and fluoxetine are the most toxic compounds tested in *P. tricornutum*, with the lowest IC₅₀ values of the tested cultures (< 50 µg L⁻¹). In opposition, ibuprofen appears as the less toxic compound tested, with the highest dose (ca 350 µg L⁻¹) needed to reduce in half the growth of *P. tricornutum*.

Table 1. Exposure compounds and respective concentrations and *Phaeodactylum tricornutum* relative growth inhibition and calculated IC₅₀.

Compound	Concentrations tested (µg L ⁻¹)	Inhibition (%)	IC ₅₀ (µg L ⁻¹)	Reference
Propranolol	0.3	0.15	194.6	[22]
	8	4.11		
	80	41.11		
	150	77.08		
	300	154.16		
Fluoxetine	0.3	0.63	47.3	[23]
	0.6	1.27		
	20	42.26		
	40	84.52		
	80	169.04		
Ibuprofen	0.8	0.23	350.6	[24]
	3	0.86		
	40	11.41		
	100	28.52		
	300	85.57		

Glyphosate	10	4.43	225.9	[15]
	50	22.13		
	100	44.27		
	250	110.67		
	500	221.34		
SDS	0.1	15.5	11.4	[42]
	1	21.1		
	3	24.6		
	10	43.8		

3.2. Oxidative stress biomarkers

To assess possible cell damages owing to oxidative stress circumstances induced by xenobiotic exposure, various oxidative stress biomarkers were assessed in the diatom cells submitted to the different pollutant concentrations. Oxidative stress biomarker values (average \pm std deviation), linear regression slope and the statistical relation between each biomarker and the external concentration of the different contaminants applied are presented in the supplementary material Table S1. Further discussion on the biological considerations of these results can be found in the studies contributing for the present work [15,22–24,42]. To study the dose-response relationship of each of the biomarkers analysed, correlations between the biomarker response and the exogenous dose applied were also determined (supplementary material Table S1). Propranolol gradient concentrations showed a positive-dose relationship for SOD antioxidant enzyme ($r^2 = 0.87$, $p < 0.05$) and TBARS ($r^2 = 0.72$, $p < 0.05$), whereas CAT and APX showed no significant relation to this compound. A significant rise in lipid peroxidation products (TBARS) was observed in *P. tricornutum* cells exposed to fluoxetine concentration gradient, revealing a positive and significant correlation ($r^2 = 0.73$, $p < 0.05$). Among the antioxidant enzymes tested, SOD and APX followed a similar trend to lipid peroxidation, though not significant, but showing the highest activity levels under the highest concentration applied ($80 \mu\text{g L}^{-1}$), whereas CAT activity did not respond to any of the exogenous fluoxetine concentrations. A positive correlation with the exogenous ibuprofen dose applied is shown for TBARS ($r^2 = 0.77$, $p < 0.05$), and APX activities ($r^2 = 0.62$, $p < 0.05$). On the other hand, CAT and SOD antioxidant enzymes were irresponsive to this contaminant concentration variation. Glyphosate exposure revealed a positive correlation to APX ($r^2 = 0.60$, $p < 0.05$) and TBARS ($r^2 = 0.92$, $p < 0.05$) activities, where TBARS activity levels showed an increasing trend up to the maximum glyphosate exogenous concentration ($500 \mu\text{g L}^{-1}$). CAT antioxidant enzyme revealed no correlation to glyphosate exposure but a great increase in activity under the highest concentration applied ($500 \mu\text{g L}^{-1}$). SOD activity decreased as compound concentration increased, revealing a negative but significant dose-response correlation ($r^2 = 0.64$, $p < 0.05$). The applied set of oxidative stress biomarkers showed no correlation to the exogenous SDS exposure concentrations applied to the *P. tricornutum* cell culture.

To determine the efficiency of the considered biomarkers in classifying the samples exposure, a multivariate canonical analysis was conducted for each of the studied compounds (Figure 1). Fluoxetine (Figure 1B) and glyphosate (Figure 1D) multivariate analysis revealed the lowest classification efficiency of the samples (38.9%) using the considered biomarkers. This was attributed to a high degree of misclassification of the samples exposed to intermediate concentrations, indicative of a low impact of these compounds at these concentrations on the considered biomarkers. On the other hand, the assessed biomarkers in the cells exposed to propranolol (Figure 1A) and SDS (Figure 1E) were able to provide a correct classification of more than half of the evaluated samples (55.6% and 53.3 %, respectively). Once again, in intermediate concentrations the considered biomarkers had lower efficiency in classifying the samples, due to possible reduced impacts of these compounds in intermediate concentrations.

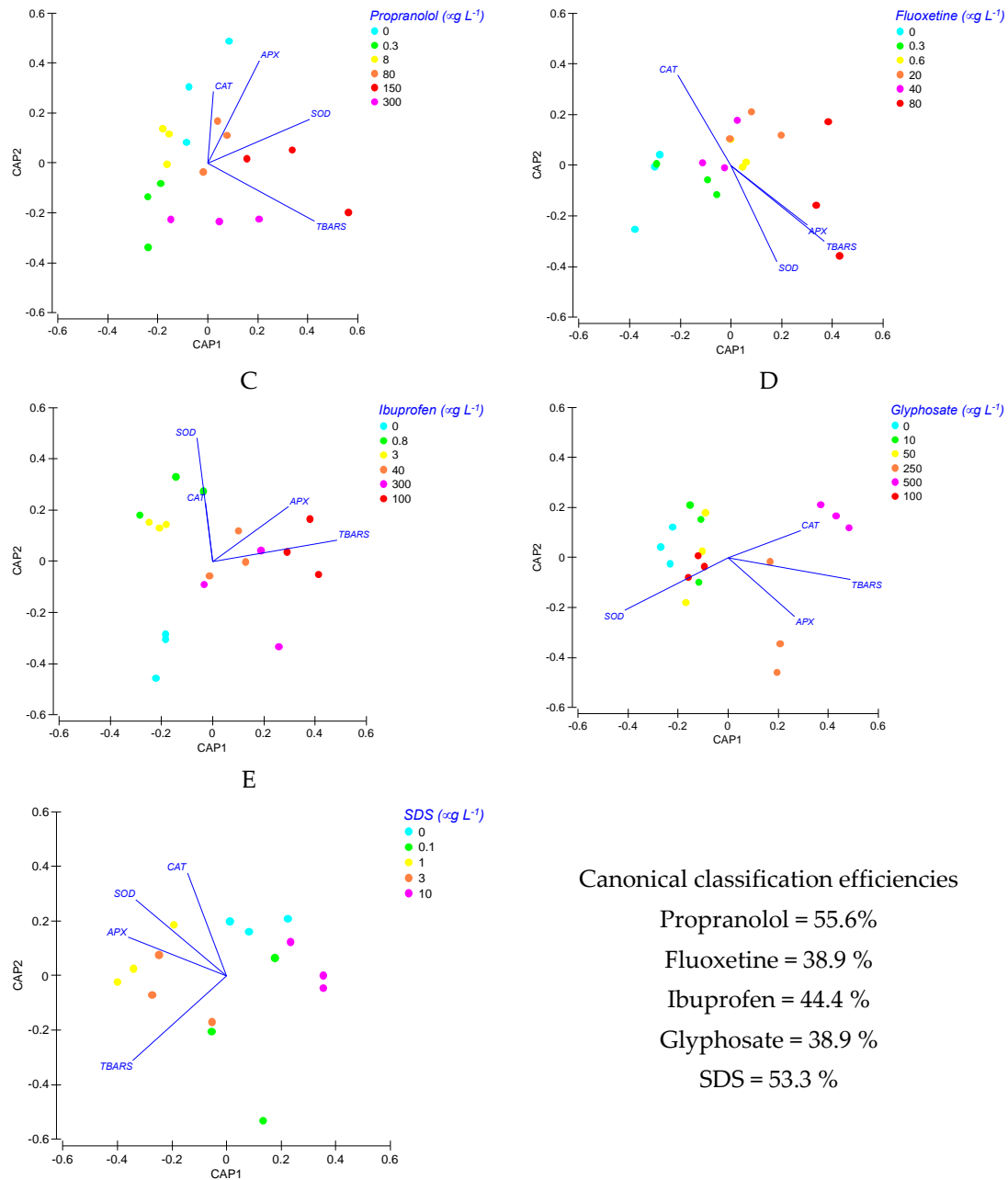


Figure 1. Canonical analysis of principal components plots of the diatom cells exposed to propranolol (A), fluoxetine (B), Ibuprofen (c), glyphosate (D) and SDS (e) at different exogenous concentrations, having the biomarker values as classifying variables. Data sources are available in Table 1.

3.3. Oxidative stress biomarkers indexes

To evaluate the applicability of these biomarkers in multivariate indexes, towards a better communication of the overall effect of each tested compound, three index approaches were undertaken: the Integrated biomarker response (IBR) index; the Principal component analysis (PCA) based index; and the Bioeffect Assessment Index (BAI)

The IBR index applied to biomarker results on propranolol and SDS contaminant exposure gradient, showed no significant variance between the exogenous concentrations tested (Figure 2A and 2D respectively). Nevertheless, a trend can be noticed for both pollutants, as the highest index values reflected higher concentrations. For the glyphosate

exogenous concentration gradient, IBR scores showed significant increases on index values at the 250 $\mu\text{g L}^{-1}$ and 500 $\mu\text{g L}^{-1}$ exposure concentrations (Figure 2B), akin for fluoxetine (Figure 2C) delivering a significantly higher index value at the 80 $\mu\text{g L}^{-1}$ concentration, suggesting a significant dose-response tendency for both pollutants. Concerning exposure to ibuprofen (Figure 2E), the IBR index revealed a significantly higher score at the maximum concentration tested of 300 $\mu\text{g L}^{-1}$, and in fact, this increase reflects a highly significant positive correlation together with the external ibuprofen gradient.

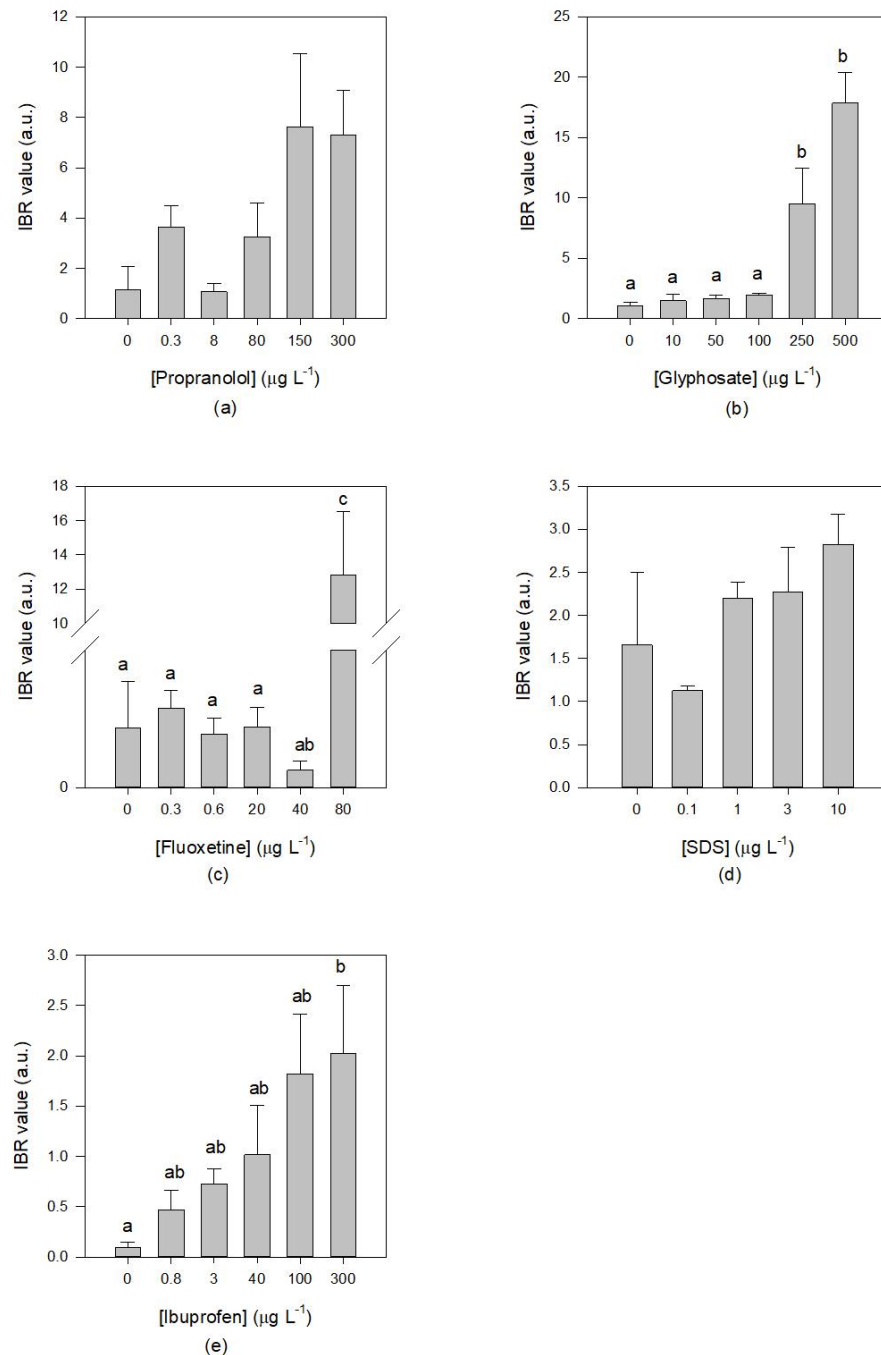


Figure 2. Integrated biomarker response (IBR) index of *Phaeodactylum tricoratum* cells exposed to different contaminants and concentrations. (A) Propranolol. (B) Glyphosate. (C) Fluoxetine. (D) SDS. (E) Ibuprofen. Average \pm standard deviation, N = 3, different letters indicate significant differences at $p < 0.05$.

Regarding PCA-based index results for propranolol exposure results, a significantly lower value was found in cultures exposed to $0.3 \mu\text{g L}^{-1}$, whilst the highest significant value is verified at the $150 \mu\text{g L}^{-1}$ propranolol concentration (Figure 3A). For glyphosate and fluoxetine exposure (Figure 3B and 3C respectively), PCA-based index attributes a significantly higher score at the maximum concentrations of $500 \mu\text{g L}^{-1}$ and $80 \mu\text{g L}^{-1}$, respectively, evidencing a positive correlation with both exogenous concentrations of these contaminants. Concerning SDS detergent (Figure 3D), PCA-based index scores did not show any significant variances under the increasing contaminant exposure gradient. On the other hand, in relation to ibuprofen (Figure 3E), significant and approximate high index values are presented for cultures exposed to $0.8 \mu\text{g L}^{-1}$ ibuprofen and all the concentrations above.

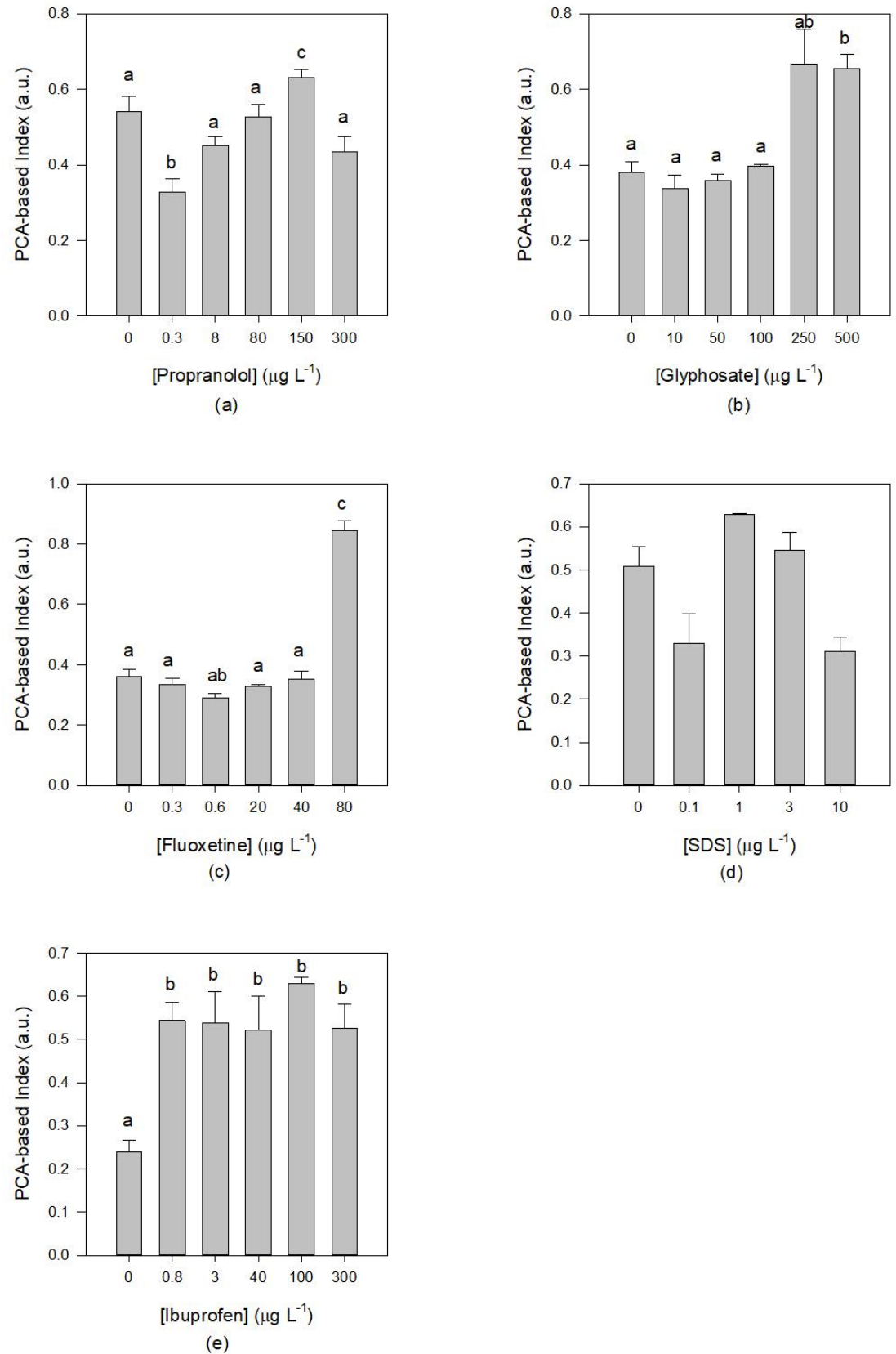


Figure 3. Principal component analysis (PCA) based index of *Phaeodactylum tricoratum* cells exposed to different contaminants and concentrations. (A) Propranolol. (B) Glyphosate. (C) Fluoxetine. (D) SDS. (E) Ibuprofen. Average \pm standard deviation, N = 3, different letters indicate significant differences at $p < 0.05$.

For the propranolol exposure gradient, BAI index provides similar results as the abovementioned PCA-based index, where a significant low index value is attributed to the minimum exogenous concentration of $0.3 \mu\text{g L}^{-1}$, and a significant index value increase is registered under the application of $150 \mu\text{g L}^{-1}$ propranolol, underlining a positive trend for dose-response (Figure 4A). Regarding glyphosate, BAI was unsuccessful to provide significant values for the evaluation of antioxidant stress response under this contaminant exposure gradient (Figure 4B). However, a slight increment in the index values follow contaminant concentration increase. Considering fluoxetine, BAI index values were only found to be significantly higher in the cells exposed to $80 \mu\text{g L}^{-1}$, the maximum contaminant concentration applied (Figure 4C). For the SDS detergent, a significantly higher index value is obtained in the cells exposed to $1 \mu\text{g L}^{-1}$ SDS, whereas an inverse trend is noticed for the maximum exogenous concentration, with significantly lower index scores, suggesting a significant inhibition of the activity levels of the antioxidant enzymes in the cells exposed to $10 \mu\text{g L}^{-1}$ SDS (Figure 4D). Applied to ibuprofen exposure data, the BAI index classifies the response to contaminant concentration gradient similarly to the PCA-based index, attributing significant and approximately high scores for cells exposed to $0.8 \mu\text{g L}^{-1}$ ibuprofen and all the concentrations above, without the suggestion of a dose-response trend (Figure 4E).

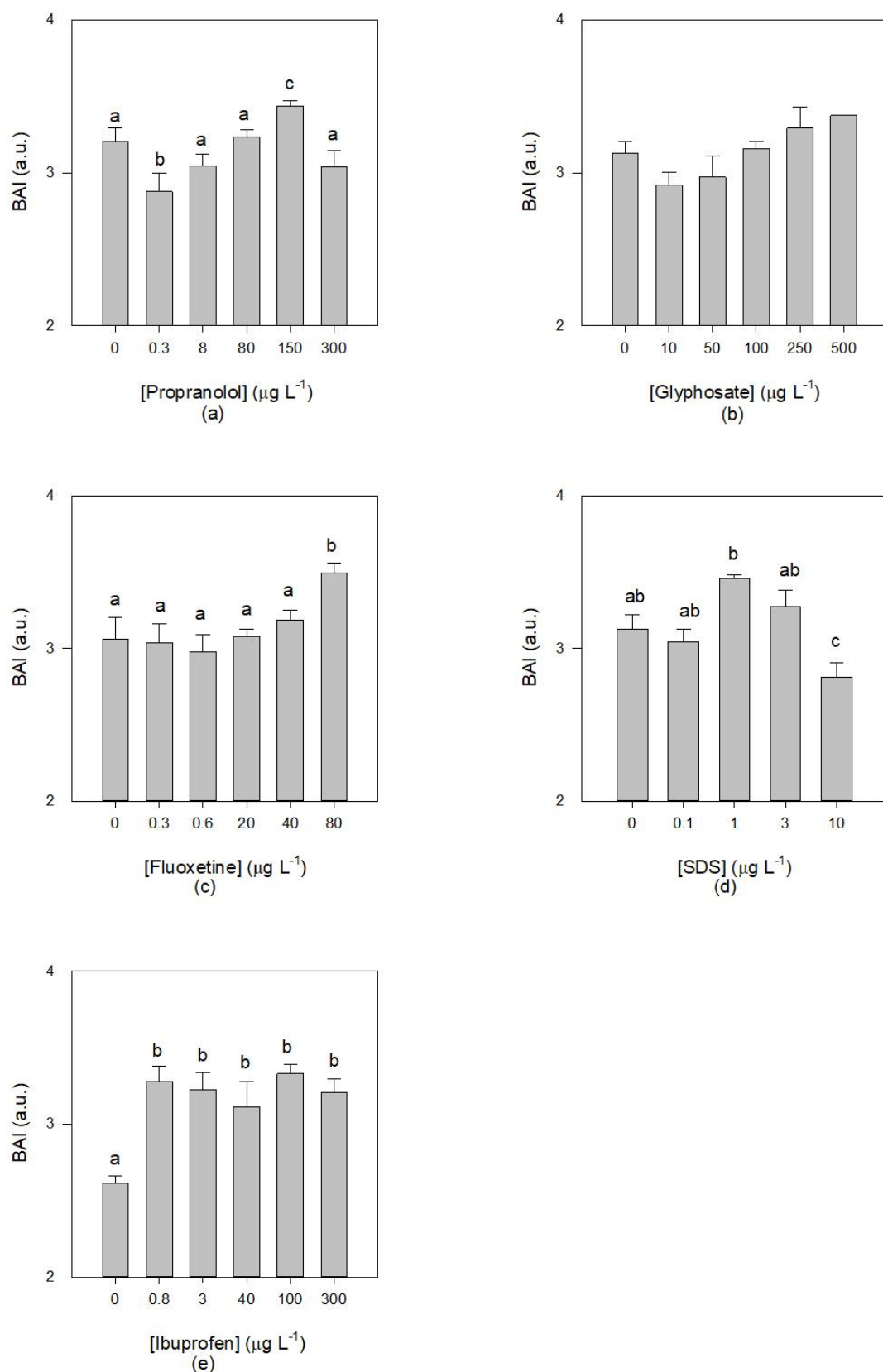


Figure 4. Bioeffect Assessment Index (BAI) of *Phaeodactylum tricornutum* cells exposed to different contaminants and concentrations. (A) Propranolol. (B) Glyphosate. (C) Fluoxetine. (D) SDS. (E) Ibuprofen. Average \pm standard deviation, N = 3, different letters indicate significant differences at $p < 0.05$.

In order to provide a comparison of the obtained index results, a visual exploration of the correlations between the exogenous contaminant concentrations and the suitability of the subjected indexes, is presented in Figure 5 via cluster heat maps. These clusters display the statistical significances of the associations between the considered variables and allow the evaluation of dataset quality. When comparing the IBR values with the exogenous concentration a significant positive dose-response correlation was found in the cells exposed to propranolol, ibuprofen, glyphosate and SDS, while for fluoxetine the inverse trend could be observed (Figure 5). If the same approach is performed for the PCA-based index it is possible to observe that the values of this index in the cells exposed to ibuprofen and glyphosate, present an inverse significant dose-response relationship with the exogenous dose for these compounds. As for the BAI values, these showed only significant correlations with the exogenous dose applied in the cells exposed to fluoxetine and glyphosate. It is also noticeable that in most cases, the majority of the indexes also present significant correlations between them..

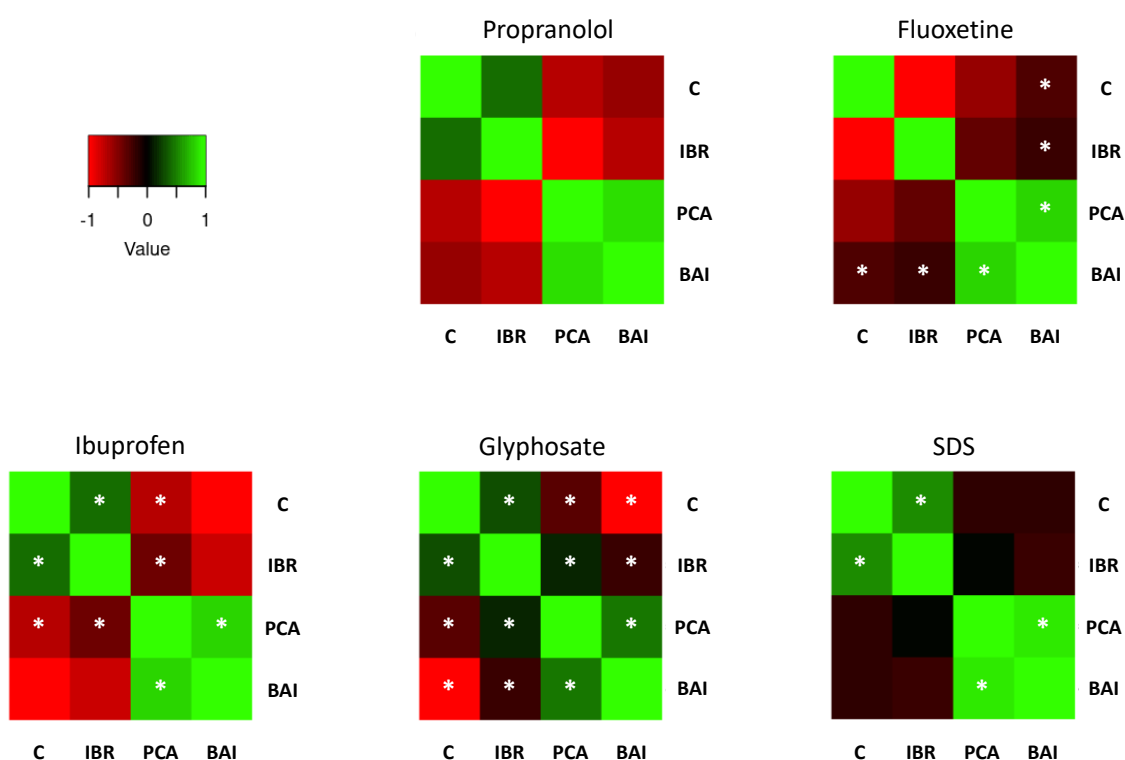


Figure 5. Spearman correlations heatmaps between the index values calculated and the exogenous concentrations applied for each compound tested. (*) denote significant differences at $p < 0.05$) and for all compounds tested. Indexes include the Integrated biomarker response (IBR) index; the Principal component analysis (PCA) based index; and the Bioeffect Assessment Index (BAI), whilst (C) refers to the gradient of exposure concentration.

4. Discussion

In the present work, selected biomarker data from five ecotoxicological contributions on emerging contaminants (three pharmaceuticals, one detergent and one herbicide), were subjected to integrated response analyses using three methods, the Integrated Biomarker Response (IBR) [5], the Principal Components Analysis -based index (PCA) and the Bioeffect Assessment Index (BAI) [19], to devise which could be more suitable delivering an accurate measure on the physiological stress response of diatom cell cultures under pollutant exposure.

Greater emphasis on evaluating the suitability of the already existing indexes prior to developing new ones is essential [16]. The set of biomarkers should be selected with flexibility considering cell function, biological level and pollutant affinity, in order to calculate relevant integrated responses of organisms to different contamination compounds, given that certain marker responses could emphasize certain functions or systems, while other levels of sensitivity would be at stake. Additionally, attention on the proper interpretation of biomarker responses integrating the index construction as parameters, and later, interpreting the derived index scores, is paramount [34]. In sum, a representative index is able to expose toxically induced response trends in natural systems which, therefore, are relevant to policy resolutions.

Previous contributions concerning aquatic environmental contamination, based their analysis on biomarker responses, proving that the adverse effects resulting from a wide variety of pollutants can be evaluated based on an array of specific and general exposure biomarkers, such as oxidative stress biomarkers, exhibiting sensitivity and accuracy in developing valid results [6,43], which are essential for the correct computation of the indexes [34]. Here, the efficiency of the applied set of biomarkers was determined by CAP (Canonical Analysis of Principal coordinates) multivariate statistical approach. The antioxidant enzymes CAT, SOD and APX, and the non-enzymatic TBARS oxidative stress biomarkers, involved in the stress response physiological processes of a cosmopolitan marine diatom species, *P. haedactylum tricornutum*, revealed an overall suitability for characterizing the different exogenous contaminants and the applied concentration gradients. This set of biomarkers produced statistically significant dose response relations for four out of the five pollutants fitting the experiment purpose, as figured by CAP efficiency results.

The IBR index is one of the most widely used and proficient in evaluating site good ecological status and contamination levels and in fact, available literature on biological effects following chemical exposure, integrate the IBR index as an important tool for contamination assessment [5–7,34,44–47]. IBR appears as a useful tool to mark pollution in field assessments, as it reflects contaminant levels in sites, despite the variability of biomarkers used for its calculation. Field and biomonitoring studies, concerning environmental pollution, rely on IBR for result support [48–52]. Fewer laboratory studies testify IBR's applicability within the scope of treatment comparison [47,53].

One of the premises of the IBR index is the 'user's judgment step', as one should decide the weight of each biomarker for the index construction *a priori*, as it may influence the index positively or negatively, depending on an induction or an inhibition in terms of physiological response to contamination. In the present work, the decision was based on the statistical relation of each biomarker alongside a gradient of contaminant concentrations [5]. This is the only empirical step of the IBR index, and can also be its weakness, putting at stake the index's reliability, as it is highly dependent on the user expert judgment and on data availability. As it was previously observed, the same biomarker can have different responses depending on the species and on the tested compound and thus this attribution of a positive or negative influence has to be greatly based on experimental data [56,57]. As the IBR index integrates multiple biomarker scores through standardization of diverse biological responses, it enables direct comparisons among contaminants and performs suitable scores for the set of biomarkers selected. In the end, a sum-up of the complex collected data is presented within a scoring framework, where high values represent environmental stress and low values represent a lack of stress or inhibition. The results of IBR data analysis in this study were consistent and reflected the overall oxidative stress conditions induced in *P. tricornutum* cells by the exogenous contaminants, translating the expected response of the cell physiological mechanisms to overcome stressful contaminated conditions. Higher index values were attributed to higher pollutant concentrations, which conforms with the general assumption of IBR, revealing statistically significant dose-response trends for four out of the five harmful compounds examined. Our considerations on IBR's advantages and disadvantages must regard some conditioning factors: i) a relatively large set of biomarkers is recommended as the higher the

number of biomarkers, the more significant the index becomes [34,44]; ii) it is recommended to classify which markers can be combined for IBR, differentiating effect and defence responses [34]; iii) to compare spatial and temporal series of datasets, new data should be normalized and incorporated together with previously obtained numerical values, since the numerical values of IBR are comparable only within each dataset, [34]; iv) IBR values must be interpreted in combination with individual biomarker scores [7,47]; v) IBR is appropriate for qualitative assessments [50]. In this context, IBR constitutes a robust and practical tool to assess the susceptibility of *P.tricornutum* cells to contaminants using multiple biomarker responses.

The BAI index came into sight as a modification of the 'Health Assessment Index' (HAI) [58], intending to assess contamination levels for coastal areas. It integrates only biomarkers of general toxicity, leading to a more holistic approach since the use of biomarkers that are non-specific to toxic effects allow the index to respond to a wider set of chemicals [19]. This is a valuable characteristic for an environmental health index, as the anthropogenic sources of contamination comprise a complex mixture of pollutants and tend to show an exponential worldwide increase [16]. The BAI index has confirmed suitability to address alterations at different bio-organizational levels, reflecting general toxicity patterns and revealing health status under polluted conditions, capturing degradative and restorative information on routine monitoring programs and allowing comparison between geographically distinct areas with differing contamination conditions [6,19,34,45]. The BAI index, an univariate statistical approach, transforms the complex alterations of biomarkers into single classes of values, losing some of the original data variability, so the combination of BAI analysis with other integrative index approaches is frequently applied for assessment protocols [6,45]. The BAI index revealed low sensitivity against our data set, confirming its scope of applicability as more adequate for general toxicity biomarkers and parameters of non-specific stress, as suggested in the literature [34], while present results were constructed from specific exposure oxidative stress biomarkers. Comparing, the BAI index exhibited a significant correlation to the PCA-based index, both indexes suggesting similar stress patterns under the application of all contaminant gradients. On the other hand, comparisons between the BAI index and the IBR index reflected the dissimilarities on these indexes productions, as generally, IBR will be based on parameters of general health biomarkers by itself or complemented with specific biomarkers, whereas the BAI takes only general toxicity markers [34]. Still, the BAI results support an overall significantly negative relation between index scores and pollutant concentration gradients, confirmed particularly for fluoxetine and glyphosate.

The PCA is a multivariate statistical technique, familiar in the identification of environmental pollution patterns and in the discrimination of the main variables responsible for the variance of chemical accumulation in organisms and consequential biological effects, thus, largely applied for the development of interpretative models of pollution data for decision-makers [48,59–61]. PCA is an effective tool when applied to high dimensional data sets, reducing their dimensionality while identifying the patterns between the inter-related variables, retaining as much as possible the variability of the original data set [62,63]. Also, PCA demonstrates applicability for index construction, selecting the appropriate measured parameters to integrate the index and identifying the main axes of variance within a data set [62,64,65]. Improved interpretation on other indexes results, such as the IBR index, may be supported by the appliance of the PCA analysis [48]. The PCA correct procedure assumes: i) large complex data sets; ii) the normalization and/or standardization of the original data should be considered; iii) biomarker weigh and influence are set solely based on statistical results; iv) patterns are suggested without reference to prior knowledge about the samples, and so it is also recommended for scaling not to be adjusted to match prior knowledge of the data. Nevertheless, limitations on PCA applicability should be considered upon output interpretation, which are, according to Lever et al. (2017): a) the assumption of a linear underlying structure of the data, b) highly correlated patterns may be undetermined due to the resultant uncorrelated principal components (axes of variance), and, c) maximizing variance is the objective rather than clustering

results. When applied to our data set, PCA is satisfactorily delivering a significant negative correlation for the overall experiment relative to compound concentration and index scores, particularly for ibuprofen and glyphosate gradients. In fact, the observed trend was identical by using either PCA or BAI index, which, in their hand, were statistically positively correlated. PCA relies on data variance to create next step assumptions, and a large data set delivers better results with this statistical method.

5. Conclusions

This study provides information about the suitability and reliability of three biological indexes in pollution monitoring research. An adequate index provides an integrative approach, constructed with the integration of multi-biomarker response data, to better clarify the links between contaminant exposure and biological effects in aquatic systems. The potential use of stress biomarkers in biological effects studies is here highlighted.

Overall, the three indexes addressed in the present study (IBR, PCA and BAI) could be successfully applied for the evaluation of environmental quality given that all of them depicted the response triggered by the tested pollutants under varying concentration gradients, reflecting disruption in the cell physiological processes by contamination, and providing a simplified output of the biomarker data despite their varying sensitivity and resolution. IBR performed the best, successfully reflecting the expected level of exposure, thus revealing the stress patterns exposed by the set of specific oxidative stress markers here selected, under a contaminant gradient, corroborating it as a suitable tool for the assessment of the biological effects of pollution in marine organisms. IBR combines multi-biomarker responses, delivers a qualitative approach and provides temporal assessment of contamination patterns.

Open access publications examining the applicability of the already available tools for pollution assessment, such as the present, improve scientific communication and efficiency within the research community, sustaining result communication to the general public and bringing great advantages for management deliberations.

Supplementary Materials: The following are available online at www.mdpi.com/xxx/s1, Table S1.

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