| 1 | A revisited hemolytic assay for palytoxin detection: limitations for its quantitation in |
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26 Abstract

Palytoxin (PLTX) and its analogues have been detected as seafood contaminants and associated with a series of human foodborne poisonings. Due to some fatalities ascribed to the ingestion of PLTX-contaminated marine organisms, the development of methods for its detection in seafood is recommended by the European Food Safety Authority (EFSA). Due to its feasibility, the spectrophotometric hemolytic assay is widely used to detect PLTX in different matrices, even though a standardized protocol is still lacking.

Thus, on the basis of available assay procedures, a new standardized protocol was set up 33 using purified human erythrocytes exposed to PLTX (working range: 3.9×10^{-10} - 2.5×10^{-8} M) in 34 a K^+ -free phosphate buffered saline solution, for 5 h at 41°C. An intra-laboratory 35 characterization demonstrated its sensitivity (limit of detection, LOD=1.4x10⁻¹⁰ M and 36 quantitation, LOQ=3.4x10⁻¹⁰ M), accuracy (bias=-0.8%), repeatability (RSDr=15% and 6% 37 for intra- and inter-day repeatability, respectively) and specificity. However, the standardized 38 method seems not to be suitable for PLTX quantitation in complex matrices, such as mussels 39 (Mytilus galloprovincialis) extracts, below the limit suggested by EFSA (30 µg PLTXs/Kg 40 shellfish meat). Thus, the hemolytic assay for PLTX quantitation in seafood should be used 41 only after a careful evaluation of the specific matrix effects. 42

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44 Keywords

45 Palytoxin, hemolytic assay, matrix effect, Mytilus galloprovincialis

Abbreviations: PLTX, palytoxin; LOD, limit of detection; LOQ, limit of quantitation; RSDr, relative standard deviation of repeatability; 42-OH-PLTX, 42-hydroxy-palytoxin; Ost-D, Ostreocin-D; OVTX, ovatoxin; EFSA, European Food Safety Authority; SAGM, Saline-Adenine-Glucose-Mannitol solution; ADSOL, Adenine-Dextrose Solution; D-PBS, Dulbecco's Phosphate Buffered Saline; EDTA, ethylenediaminetetracetic acid; O.D., optical density.

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47 **1. Introduction**

Palytoxin (PLTX) is a highly toxic non-polymeric complex molecule, originally isolated from 48 zoanthids of the genus Palythoa (Moore and Scheuer, 1971), and later identified in benthic 49 dinoflagellates of the genus Ostreopsis and cyanobacteria of the genus Trichodesmium 50 (Ciminiello et al., 2008; Kerbrat et al., 2011). Over the years, several PLTX analogues have 51 been detected in different marine organisms, including: (i) Ostreocin-D (Ost-D) in Ostreopsis 52 siamensis (Usami et al., 1995); (ii) two stereoisomers of 42-hydroxy-PLTX (42-OH-PLTX), 53 differing only for the configurational inversion at C50, identified in *Palythoa toxica* and *P*. 54 tuberculosa (Ciminiello et al., 2014a); and (iii) ovatoxin-a (OVTX-a), the major toxin 55 produced by Ostreopsis cf. ovata in the Mediterranean Sea (Ciminiello et al., 2012). In 56 addition, a series of OVTX-a analogues have been identified, such as OVTX-b to -k and 57 isobaric palytoxin (Brissard et al., 2015; García-Altares et al., 2015; Tartaglione et al., 2016). 58 These toxins may enter the human food chain through their accumulation into different 59

marine edible organisms, such as fishes, crustaceans, bivalves, gastropods, cephalopods, echinoderms, sponges and polychaete worms (Aligizachi et al., 2011; Biré et al., 2013; Gleibs and Mebs, 1999). In particular, a series of human poisonings characterized by general malaise, myalgia, cardiac problems, respiratory distress, and sometimes death, have been ascribed to the ingestion of PLTX-contaminated fishes and crabs in tropical areas (Deeds and Schwartz, 2010; Tubaro et al., 2011b; Wu et al., 2014).

In recent years, microalgae belonging to the genus *Ostreopsis* have frequently bloomed in the temperate Mediterranean Sea and the relevant toxins (mainly OVTX-a) have been detected in microalgae, aerosolized seawater and some marine edible organisms. However, no foodborne poisonings associated with PLTXs have yet been documented in this area, so far (Biré et al., 2013; Ciminiello et al., 2014a; Del Favero et al., 2012).

Although there is no official regulation for PLTXs in seafood, the European Food Safety 71 72 Authority (EFSA) suggested a maximum limit of 30 µg PLTXs/Kg of shellfish meat, and recommended the development of specific, rapid, precise and accurate methods for PLTXs 73 quantitation in seafood during monitoring programs (EFSA, 2009). Despite several methods 74 for PLTXs detection have been already published (Riobò et al., 2011; Tubaro et al., 2014), 75 they are not completely validated to be officially accepted, frequently requiring expensive 76 equipments and highly qualified operators, and/or presenting limitations in terms of 77 sensitivity, specificity and matrix effects. 78

Among them, the delayed hemolytic assay is one of the most used screening methods for 79 PLTX quantitation, due to its simplicity, cheapness, rapidity and sensitivity. It is based on the 80 toxin ability to convert the Na^+/K^+ ATPase of mammal erythrocytes to a nonspecific cationic 81 channel, leading to a rapid loss of K⁺ ions from cells and a delayed hemoglobin release that 82 83 can be easily measured spectrophotometrically (Habermann et al., 1981). The hemolytic assay is usually carried out following the Bignami's protocol (1993) with or without modifications, 84 but a standardized and universally accepted procedure has not been defined, so far. Most 85 literature studies report the original assay with a series of modifications mainly concerning 86 the origin of erythrocytes, the time and temperature of their incubation with PLTX, and/or the 87 working buffer composition (Aligizaki et al., 2008; Biré et al., 2013; Brissard et al., 2014; 88 Gleibs et al., 1995; Kim et al., 2002; Lenoir et al., 2004; Malagoli, 2007; Onuma et al., 1999; 89 Pezzolesi et al., 2012; Riobò et al., 2006; Riobò et al., 2008; Taniyama et al., 2001; Taniyama 90 et al., 2003; Volpe et al., 2014; Wachi and Hokama, 2001). Due to these experimental 91 variables, a standardized protocol for the hemolytic assay to be considered as a reference 92 procedure in PLTX quantitation is still lacking. 93

Moreover, edible marine species could contain hemolytic compounds different from PLTXs, tentatively interfering with the assay giving false positive results. For instance, a recently

developed biosensor to quantify PLTX in mussels, based on lactate dehydrogenase release 96 from sheep erythrocytes, appeared to be influenced by a significant mussels matrix effect and 97 low PLTX recovery (Volpe et al., 2014). Thus, the accuracy and precision of the hemolytic 98 assay for spectrophotometric PLTX quantitation in shellfish needs to be properly evaluated. 99 Hence, this study was carried out to set up a novel standardized protocol for the hemolytic 100 assay useful as reference procedure and to characterize its suitability for PLTX quantitation in 101 mussels (Mytilus galloprovincialis), often heavily contaminated by PLTXs (Brissard et al., 102 2014; Aligizaki et al., 2008; Aligizaki et al., 2011; Amzil et al., 2012). 103

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105 **2. Materials and methods**

106 *2.1 Toxins and other materials*

PLTX was purchased from Wako Pure Chemicals Industries Ltd. (Osaka, Japan; purity > 90 107 %). 42S-hydroxy-50S-palytoxin and 42S-hydroxy-50R-palytoxin were isolated from Palythoa 108 tuburculosa and P. toxica, as previously reported (Ciminiello et al., 2009; Ciminiello et al., 109 2014b). Ovatoxin-a and the mixture of ovatoxin-a,-d,-e were kindly provided by Prof. P. 110 Ciminiello (Università di Napoli "Federico II", Naples, Italy). The mouse monoclonal anti-111 112 PLTX antibody 73D3 (mAb-PLTX) was produced by a hybridoma cell culture, as previously described (Bignami et al., 1992). All the other reagents were of analytical grade and 113 purchased from Sigma-Aldrich (Milan, Italy), if not otherwise specified. 114

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116 2.2 Red blood cell purification

Blood samples from healthy human volunteers were obtained from the Transfusion Center, Azienda Ospedaliera Universitaria, Trieste, Italy. All donors signed an approved consent form giving permission for the collection and use of blood for research purposes (WMA Declaration of Helsinki). Blood was drawn by venipuncture between 08.00 a.m. and 10.00 a.m. to minimize variability due to circadian rhythms, and immediately processed. Blood, collected into standard triple bag systems, was fractionated following standard procedures to obtain buffy-coats, used to purify red blood cells. Buffy coats (50 ml) were diluted 1:1 (v/v) in the erythrocytes preservation solution (specified in section 2.3) and then centrifuged at 2400 rpm for 10 min at 4 °C. Red blood cells pellet (2 ml), suspended in the red blood cells preservation solution (10 ml), was washed three times by centrifugation at 1500 rpm for 5 min at 15 °C. Then, the final pellet was re-suspended 1:10 (v/v) in the preservation solution.

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129 2.3 Red blood cells storage

Three storage solutions were used to evaluate red blood cells preservation at 4° C: Saline-130 Adenine-Glucose-Mannitol solution (SAGM, containing 5x10⁻² M dextrose, 0.15 M NaCl, 131 2.4x10⁻³ M adenine and 2.9x10⁻² M d-mannitol) (Zehnder et al., 2008), Adenine-Dextrose 132 Solution (ADSOL, containing 0.12 M dextrose, 0.15 M NaCl, 3.9x10⁻³ M adenine and 4.1x10⁻ 133 ² M d-mannitol) (Moore, 1991) and Dulbecco's Phosphate Buffered Saline (D-PBS) 134 containing 1mM ethylenediaminetetracetic acid (EDTA) and 5 mM glucose (Lowe et al., 135 1973). Red blood cells preservation was evaluated by means of erythrocyte resistance to 136 spontaneous lysis and erythrocyte concentration. For the first parameter, 0.4 ml of red blood 137 cells suspension was centrifuged at 1500 rpm for 5 min and the optical density of the 138 supernatant was then measured at 405/540 nm (Microplate autoreader; Bio-Tek Instruments; 139 Vinooski, VT). Red blood cell concentration was evaluated by cell counting following the 140 Trypan Blue Exclusion Test. Acceptable thresholds were constant cells concentration and 141 optical densities lower than 0.5, a value close to the average optical densities of the negative 142 controls (125 µl of red blood cell suspension incubated with K⁺-free D-PBS without toxin). 143 Both parameters were assessed daily for 36 days. 144

146 *2.4 Experimental design*

The hemolytic assay was standardized considering the following parameters: the use of purified human erythrocytes *vs* whole blood, the influence of selected ions (borates, calcium, sodium, and potassium) on PLTX-induced hemolysis, and the incubation temperature of erythrocytes exposed to PLTX. The standardized assay was subsequently characterized according to the international principles, as described by the Eurachem Guide (Magnusson and Örnemark, 2014).

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154 2.5 Standardized hemolytic assay

155 After washing, red blood cells were pelleted by centrifugation (1500 rpm for 5 min) and resuspended in K⁺-free D-PBS at the concentration of 1×10^8 cells/ml. In 96-wells plates, 125 µl 156 of PLTX solution and 125 µl of the erythrocytes suspension were added to each well and 157 incubated for 5 h at 41 °C (PLTX final concentrations: 1.22x10⁻¹¹-4.00x10⁻⁷ M). As negative 158 controls, 125 µl of red blood cell suspension were incubated with 125 µl K⁺-free D-PBS 159 without toxin. As a positive control, 100% hemolysis was achieved by incubating the 160 erythrocytes suspension with 125 µl of 0.1% Tween 20 (v/v) for 5 h at 41° C. After 161 incubation, the plate was centrifuged at 1500 rpm for 5 min at 15 °C and, using a 162 multichannel pipette, the supernatant was carefully transferred into a clear flat bottom 96-163 wells plate avoiding pellet braking up. Optical density (O.D.) of supernatant was then 164 measured at 405/540 nm and the percentage of hemolysis calculated with respect to the 165 positive control by the following formula: 166

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167 % of hemolysis = 100 x (O.D. PLTX exposed sample-O.D. negative control)/(O.D. positive
168 control - O.D. negative control).
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170 2.6 Evaluation of the matrix effect

To assess the hemolytic assay suitability to quantify PLTX in mussels at levels below the 171 suggested EFSA limit (30 µg PLTXs/kg edible parts, corresponding to about 11.2x10⁻⁹ 172 mol/kg; EFSA, 2009), different extracts of Mytilus galloprovincialis edible parts were 173 prepared. Each extract was analyzed by liquid chromatography high resolution mass 174 spectrometry (LC-HRMS) to verify the absence of PLTX before the matrix effect evaluation. 175 Mussels were collected in the Gulf of Trieste (Italy) and shucked meat (200 g) was 176 homogenized (14000 rpm, 3 min) using an Ultra-Turrax (Ika-Werk; Staufen, Germany). The 177 homogenate (1 g) was extracted three times with 3 ml of different solvents (80%, 50% or 20%) 178 aqueous ethanol or aqueous methanol). Each extractive solution was then centrifuged at 5500 179 rpm for 30 min, the corresponding supernatants were pooled and the volumes adjusted to 10 180 ml with the relevant extraction solvents to obtain six extracts at a final concentration of 0.1 g 181 mussels meat equivalents/ml. The hemolytic activity of each extract was then evaluated at 182 five dilutions (1:1, 1:10, 1:50, 1:100 and 1:1000, v/v) to assess background hemolysis. 183

At dilutions devoid of background hemolysis, extracts were spiked with different PLTX concentrations to prepare matrix matched-samples at PLTX levels ranging from 3.9×10^{-10} to 2.5×10^{-8} M. These matched samples were then analyzed using the hemolytic assay. The relevant hemolytic activity was compared to that induced by the same PLTX concentrations without matrix.

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190 *2.7 Statistical analyses*

Results of the hemolytic assay are presented as mean±SE of at least three independent experiments performed in triplicate. Linearity (r²) of the calibration curve was estimated by linear regression analysis, using the GraphPad Prism software version 5.0 (GraphPad Prism; GraphPad Software, Inc.; San Diego, CA). Concentration-effect curves were compared by two-way ANOVA statistical analysis and Bonferroni post test, and significant differences were considered at p < 0.05. EC₅₀ (effective concentration giving 50% hemolysis) was calculated by nonlinear regression using a four parameters curve-fitting algorithm of the GraphPad Prism software.

Hemolytic assay performance was characterized according to the international principles 199 described by the Eurachem Guide (Magnusson and Örnemark, 2014). Briefly, limit of 200 detection (LOD) and quantitation (LOQ) were expressed as PLTX concentration 201 corresponding to the average of 10 blank values plus 3 or 10 times the standard deviations, 202 respectively. Accuracy was measured as % Bias (n = 10), calculated as % difference between 203 PLTX concentration measured by the assay and the theoretical concentration in the sample 204 205 divided by PLTX theoretical concentration. Repeatability was expressed as relative standard deviation of repeatability (RSDr), measured as % ratio between the standard deviation of 206 independent results and their mean value. Both independent results obtained by the same 207 operator in one day (intra-assay RSDr; n = 10) and within a 6-month period by different 208 operators (inter-assay RSDr; n = 10) were considered. 209

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211 **3. Results and discussion**

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213 *3.1 Storage of purified human erythrocytes*

The human erythrocytes model was chosen due to the easy availability of human blood, rapid isolation of significant cell numbers, and low cost. The use of purified human erythrocytes poses the need of a medium suitable to preserve the cells. To this aim, three cells storage solutions were evaluated: Saline-Adenine-Glucose-Mannitol Solution (SAGM), Adenine-Dextrose Solution (ADSOL) and D-PBS containing 1 mM EDTA and 5 mM glucose (PBS-EDTA-glucose), as described in section 2.3 (Lowe et al., 1973; Moore, 1991; Zehnder et al., 2008). Erythrocytes preservation in these solutions was daily evaluated, monitoring two

parameters up to 36 days: the spontaneous hemolysis (measured spectrophotometrically as 221 hemoglobin release) and the erythrocytes concentration determined by visual cell counting. 222 Acceptable thresholds were constant cell concentration in the storage solution and optical 223 densities lower than 0.5. Only PBS-EDTA-glucose solution allowed mainteinance of red 224 blood cells suitable for the assay for up to 3 weeks (Fig. S1), probably due to the presence of 225 crucial constituents providing the proper energy source to erythrocytes (glucose) and 226 preventing coagulation (EDTA). Hence, these conditions allow to prepare a batch of human 227 erythrocytes suitable for the hemolytic assay up to 3 weeks, reducing the working time and 228 avoiding to purify the erythrocytes before each single experiment. 229

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231 *3.2 Optimization of the hemolytic assay*

The hemolytic assay was standardized considering different experimental parameters reported 232 by previous published studies. In the first series of experiments, the hemolytic assay was 233 carried out following the most recent published method, using human erythrocytes (Malagoli, 234 2007). Following Malagoli's protocol, the sensitivity of purified human erythrocytes to 235 PLTX-induced hemolysis was compared to that of the whole human blood. The hemolytic 236 activity of PLTX (7.8x10⁻¹⁰-5.0x10⁻⁸ M) towards purified human erythrocytes was 237 significantly higher than that displayed by the whole human blood: 65% and 7% hemolysis 238 were recorded at 5.0×10^{-8} M PLTX, respectively (Fig. 1A). 239

Previous studies demonstrated that borates, as H_3BO_3 or $Na_2B_4O_7$ (> 5.0x10⁻⁶ M), and calcium ions (> 2.0x10⁻⁵ M) in the buffer solution increases PLTX-induced hemolysis, probably by promoting the interaction between the toxin and its molecular target, the Na^+/K^+ ATPase (Ahnert-Hilger et al., 1982; Habermann, 1983). Thus, the influence of these factors on PLTXinduced lysis of purified erythrocytes was evaluated. As shown in Fig. 1B, no significant differences in PLTX-induced hemolysis were observed between buffer solution containing 1 mM H₃BO₃ and that containing 1 mM Na₂B₄O₇, in agreement with literature data, where H₃BO₃ or Na₂B₄O₇ are interchangeably used. Regarding Ca²⁺ ions, no significant differences were recorded between PLTX-induced hemolysis in D-PBS with or without CaCl₂ (Fig. 1C). However, the Ca²⁺ concentration in D-PBS solution (137 mM) is higher than that (>20 μ M) reported to promote the interaction between the toxin and its target (Ahnert-Hilger et al., 1982). Thus, additional Ca²⁺ ions at millimolar concentrations in the D-PBS buffer solution containing 1mM H₃BO₃ are not necessary to increase PLTX hemolytic activity.

Based on the mechanism of action of the toxin (i.e. interaction with the Na^+/K^+ ATPase), the 253 influence of Na⁺ and K⁺ ions on PLTX-induced hemolysis was also evaluated. The presence 254 of Na⁺ ions (1.8x10⁻⁴M or 3.6x10⁻⁴M NaCl) in the buffer solution was associated to a 255 significant reduction of PLTX-induced hemolysis, probably due to the medium hypertonicity 256 causing erythrocytes shrinking and volume reduction (Kregenow, 1971) (Fig. 1D). In 257 contrast, while the K^+ -free buffer containing H_3BO_3 did not significantly influence the 258 hemolytic activity of PLTX, the latter was significantly increased using K^+ -free buffer 259 without H₃BO₃ (Fig. 1E). 260

Finally, the temperature influence on the hemolytic activity was evaluated. After erythrocytes incubation with PLTX for 5 h at 41 °C, hemolysis was significantly higher than that recorded at 37°C, with EC₅₀ values of 6.2×10^{-9} M (95% confidence intervals, CI = $5.3 - 7.2 \times 10^{-9}$ M) and 4.9×10^{-8} M (95% CI = $4.1 - 5.9 \times 10^{-8}$ M), respectively (Fig. 1F). This result is in agreement to that reported by Habermann et al. (1981), suggesting that PLTX-induced hemolysis is temperature-dependent.

267 On the whole, these results allowed to optimize a protocol for the hemolytic assay, using 268 purified human erythrocytes exposed to PLTX in a K⁺-free D-PBS buffer at 41 °C for 5 h.

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270 *3.3 Characterization of the hemolytic assay*

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Using the standardized hemolytic assay, the calibration curve for PLTX represented in Fig. 2A was obtained. The working range for PLTX detection was 3.9×10^{-10} - 2.5×10^{-8} M, with a 273 limit of detection (LOD) and quantitation (LOQ) of 1.4x10⁻¹⁰ M and 3.4x10⁻¹⁰ M, respectively. Analyzing the working range by linear regression, plotting the theoretical toxin concentrations against the PLTX concentrations measured by the hemolytic assay, a good 276 correlation coefficient was found ($r^2 = 0.9979$; n = 10) (Fig. 2B). A mean *Bias* value (%) of -277

0.8% (range: -2.0% to 2.4%) highlights the accuracy of the measures (Table 1). 278

Intra-assay and inter-assay repeatability were then evaluated. A good correlation was 279 observed, with r^2 values of 0.9736 for intra-assay (Fig. 3A) and 0.9977 for inter-assay 280 repeatability (Fig. 3B). These data were confirmed by the intra-day and inter-day repeatability 281 coefficients (relative standard deviation of repeatability, RSDr) of 15% (n=10) and 6% (six 282 months period, n=10), respectively (Table 1). 283

The haemolytic effect of PLTX in these conditions (described in section 3.2; 94% hemolysis 284 at 2.5x10⁻⁸ M) is much higher than that reported by Malagoli (2007) and Taniyama (2001): 285 they recorded a maximum hemolysis lower than 50% at the highest tested PLTX 286 concentrations (i.e. 10³ and 10² ng PLTX/ml, corresponding to 3.7x10⁻⁶ M and 3.7x10⁻⁷ M 287 288 PLTX, respectively). These concentrations were 1 to 2 orders of magnitude higher than the highest concentration of the assay working range presented in this study (i.e. 2.5x10⁻⁸ M, 289 giving 94% of hemolysis). Thus, an improved PLTX-induced hemolysis was achieved by 290 lowering the osmolarity of the working buffer (K⁺ ions withdrawal) and increasing the assay 291 temperature (41°C) in the optimized hemolytic assay protocol. 292

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3.3.2 Cross-reactivity with PLTX analogues 294

The hemolytic assay was then evaluated for its ability to detect some PLTX analogues and N-295 biotinyl-PLTX within the working range set up for PLTX $(3.9 \times 10^{-10} - 2.5 \times 10^{-8} \text{ M})$. Although 296 the hemolytic activity of nanomolar concentrations of 42S-OH-50S-PLTX (from P. toxica) 297 was slightly lower than that of the same PLTX concentrations, its hemolytic potency (EC_{50} = 298 5.8×10^{-9} M, 95% CI = 4.1 - 8.2 × 10⁻⁹ M) was comparable to that of PLTX (EC₅₀ = 6.2 × 10⁻⁹ M, 299 95% Cl = 5.3×10^{-9} - 7.2×10^{-9} M) (Fig. 4). The latter result is in agreement with a previous 300 study demonstrating a similar hemolytic activity of the two compounds, using mouse 301 erythrocytes (Tubaro et al., 2011a). On the other hand, hemolysis induced by 42S-OH-50R-302 PLTX (stereoisomer from *P. tuberculosa*) was significantly lower than that of PLTX (Fig. 4), 303 in agreement with its lower cytotoxic effects, previously observed on HaCaT cells (Ciminiello 304 et al., 2014b). Also the hemolytic activity of pure OVTX-a and of a mixture of OVTXs 305 (OVTX-a, -d and -e) was significantly lower than that of PLTX, similarly to their cytotoxicity 306 307 toward skin keratinocytes, as recently demonstrated by Pelin et al. (2016). In contrast, the latter result is not in agreement with other studies hypothesizing that PLTX and OVTXs 308 possess similar hemolytic activity (Brissard et al., 2014; Pezzolesi et al., 2012). However, 309 since these studies were carried out using sea urchins and Ostreopsis ovata extracts 310 containing OVTX analogues other than OVTX-a, -d and -e as well as isobaric PLTX 311 (Brissard et al., 2014; Pezzolesi et al., 2012), we can speculate that those mixtures of toxins 312 could display a hemolytic activity different than that of OVTX-a or of the other single 313 OVTXs. Moreover, since complete studies on this matrices were not carried out, a significant 314 interference of the extract's matrix cannot be excluded. 315

In addition, the hemolytic potency of a semisynthetic PLTX derivative, biotinylated PLTX, was significantly lower than that of PLTX, inducing only 5% hemolysis at the highest concentration (2.5×10^{-8} M). Intriguingly, a biotin linked to the terminal PLTX amino group reduces PLTX-induced hemolysis, indicating the importance of the primary amine for the

hemolytic effect via Na^+/K^+ -ATPase interaction. This finding is in agreement with previous 320 evidences for N-acetyl-PLTX which biological activity via Na⁺/K⁺-ATPase were over 100 321 times weaker than those of the parent compound PLTX, tentatively due to a change in the 322 global toxin conformation that prevents its dimerization (Kudo and Shibata, 1980; Ohizumi 323 and Shibata, 1980; Inuzuka et al., 2008). Moreover, this result suggests that the functional 324 hemolytic assay detects only biologically active PLTX analogues. In contrast, other non-325 functional analytical methods can detect also biologically inactive PLTX-like compounds 326 (Boscolo et al., 2013), which might not contribute the whole toxic potential of PLTXs-327 contaminated seafood samples. 328

In conclusion, these results demonstrated that the hemolytic assay has a good sensitivity for PLTX and 42*S*-OH-50*S*-PLTX from *P. toxica*. On the contrary, the stereoisomer 42*S*-OH-50*R*-PLTX from *P. tuberculosa*, as well as OVTX-a and OVTXs mixture have lower hemolytic activity. Anyway, all the tested natural PLTX analogues exert hemolytic effects, which suggest a common mechanism of action.

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335 *3.3.3 Hemolysis neutralization*

It is known that the cardiac glycoside ouabain, which binds to the same molecular target of 336 PLTX, inhibits the in vitro effects of PLTX (Habermann and Chhatwal, 1982; Pelin et al., 337 2013). Thus, to confirm the specific delayed hemolysis by PLTXs, excluding a possible 338 hemolysis by other constituents of seafood samples, the standardized assay was carried out 339 pre-incubating the red blood cells with 100 µM ouabain for 30 min at 37 °C (Aligizaki et al., 340 2008; Biré et al., 2013; Brissard et al., 2014; Gleibs et al., 1995; Malagoli, 2007; Onuma et 341 al., 1999; Pezzolesi et al., 2012; Riobo et al., 2006; Riobo et al., 2008; Taniyama et al., 2001; 342 Taniyama et al., 2003; Volpe et al., 2014). Similarly, the mouse monoclonal anti-PLTX 343 antibody (mAb-PLTX, 50 µg/ml) was used to neutralize the hemolysis induced by PLTX. The 344

348 *3.4 Matrix effect*

To assess the interference of mussel matrix on PLTX-induced hemolysis, different extracts of 349 Mytilus galloprovincialis were prepared. Firstly, 80%, 50% and 20% aqueous methanol and 350 ethanol extracts of toxin-free mussels were tested after 1:1, 1:10, 1:50, 1:100 and 1:1000 (v/v) 351 dilutions, to verify whether the matrix (mussel extracts) could produce false positive results. 352 All the aqueous ethanol extracts required dilutions higher than 1:100 to avoid matrix effects in 353 354 the hemolytic assay (Fig. S2). On the other hand, no significant interferences were recorded for the 80% aqueous methanol extract at 1:50 dilution or 50% and 20% aqueous methanol 355 extracts at 1:10 dilution (Fig. S3). Thus, the subsequent studies were carried out using the 356 357 aqueous methanol extracts at these dilutions. Specifically, 80% aqueous methanol extract (1:50), as well as 50% and 20% aqueous methanol extracts (1:10) were spiked with known 358 amounts of PLTX (final concentrations ranging from 3.9x10⁻¹⁰ M to 2.5x10⁻⁸ M). Then, the 359 spiked extracts were analyzed by the hemolytic assay in comparison to the same PLTX 360 concentrations without matrices. The extract which did not interfere with the hemolytic assay 361 362 was the 80% aqueous methanol extract diluted 1:50 (Fig. S4). Notably, this extraction method is the most suitable for PLTX extraction from mussels, giving 90-100% toxin recovery, by 363 LC-HRMS and a sandwich ELISA (Ciminiello et al., 2011; Boscolo et al., 2013). Thus, 80% 364 aqueous methanol at 1:50 dilution was subsequently used for the matrix effect study. 365

Fig. 6 shows the results of the linear regression analysis carried out comparing the theoretical PLTX concentrations in the spiked extract to those measured by the hemolytic assay. The LOQ for PLTX in the mussels extract was 1.3 ng/ml, corresponding to 13 μ g/kg meat. Considering the 1:50 dilution, the LOQ for PLTX in mussels corresponds to 640 μ g/kg meat,

a value about 20 times higher than the maximum limit suggested by EFSA (30 µg PLTXs/kg 370 meat). Moreover, the linear regression analysis yielded a correlation coefficient ($r^2 = 0.9259$) 371 and a *Bias* range from -27.0 to 33.6% (mean *Bias* = 3.7%, Table 2) index of high dispersion 372 of the data, suggesting low accuracy and precision of the measurement. Thus, the 373 standardized and characterized hemolytic assay suffers from a significant mussel matrix 374 effect, which does not allow PLTX quantitation in *Mytilus galloprovincialis* at concentrations 375 lower than that proposed by EFSA. Interestingly, a significant interference by mussels matrix 376 was recently observed also for a biosensor exploiting lactate dehydrogenase (LDH) release 377 from sheep erythrocytes as hemolysis parameter to quantify PLTX (Volpe et al., 2014): 378 379 similarly to our study, due to the significant matrix effect, a 1:50 dilution of the mussels extract was required to quantify PLTX in mussels, not meeting the EFSA requirements 380 (Volpe et al., 2014). Intriguingly, the observed interference of the mussels extract is not due 381 382 to the solvent used for extraction (i.e. 80% aqueous methanol) since the relevant mussels-free solvent did not induce hemolysis (data not shown). On the contrary, low dilutions of mussels 383 methanol extract (i.e. 1:1 or 1:10, Fig. S3) displayed an increased hemolytic activity as 384 compared to that of higher extract dilutions, sometimes comparable to that of the positive 385 control (0.1% Tween 20). Hence, the significant matrix effect could be due to hemolytic 386 compounds different from PLTX, that could be at least partially extracted from mussels by 387 aqueous methanol. Indeed, a series of potentially hemolytic compounds, such as 388 glycoglycerolipids, lysophospholipids and unsaturated fatty acids, are known constituents of 389 mussel edible parts. The concentration of these compounds in mussel meat could be also 390 influenced by season, living site of mussels, dietary composition of phytoplankton as well as 391 temperature and enzymes action during transportation and storage of seafood (Colles and 392 Chisolm, 2000; Facchini et al., 2016; Ginsburg et al., 1989; Parrish et al., 1998; Pleissner et 393 al., 2012). 394

All together, these results suggest that PLTX-induced erythrocytes lysis is not a suitable endpoint for the toxin quantitation in mussels. In addition, since the hemolytic assay has been worldwide used to detect PLTXs in different field marine samples (also concomitantly to potential human poisonings ascribable to PLTXs), those data should be carefully considered because of possible matrices effect that could have affected the analytical outcomes.

400

401 **4.** Conclusions

In conclusion, this hemolytic assay employing purified human erythrocytes is characterized 402 by a good sensitivity, accuracy, specificity and repeatability. However, it does not allow any 403 PLTX quantitation in mussels (Mytilus galloprovincialis) at concentrations below the 404 maximum limit suggested by EFSA (30 µg/kg; EFSA, 2009), due to the high matrix 405 interference. In fact, hemolytic substances different from PLTX could be extracted from 406 mussels together with PLTX (Colles and Chisolm, 2000; Facchini et al., 2016; Ginsburg et 407 al., 1989; Parrish et al., 1998; Pleissner et al., 2012), causing false positive results. Thus, the 408 hemolytic assay for PLTX quantitation in seafood could be used only after a careful 409 evaluation of the specific matrix effects in each natural sample. 410

411

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416

417 **Conflict of interest**

418 The authors declare that there are no conflicts of interest.

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Fig. 1. Optimization of the hemolytic assay. A: PLTX-induced hemolysis using whole human 608 blood or purified human erythrocytes. B: PLTX-induced hemolysis on human erythrocytes in 609 presence of 1 mM boric acid or 1 mM sodium tetraborate. C: PLTX-induced hemolysis on 610 human erythrocytes with and without 1 mM CaCl₂. D: PLTX-induced hemolysis on human 611 erythrocytes with and without 1.8x10⁻⁴M or 3.6x10⁻⁴M NaCl, 2x or 4x. E: PLTX-induced 612 hemolysis on human erythrocytes with and without K⁺ ions (2.7 mM KCl, 8.1 mM KH₂PO₄). 613 F: PLTX-induced hemolysis on human erythrocytes at 37° C and 41° C. Each point represents 614 the mean \pm SE of 3 different experiments.*p < 0.05; **p < 0.01; ***p < 0.001 (for D and E as 615 compared to PBS + H₃BO₃ + NaCl 1x; two-way ANOVA and Bonferroni post test). 616

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Fig. 2. A: Calibration curve of the hemolytic assay for PLTX. Each point represents the mean \pm SE of ten different experiments. **B**: Linear regression analysis performed within the working range of the hemolytic assay (3.9x10⁻¹⁰ M - 2.5x10⁻⁸ M) by plotting the theoretical PLTX concentrations against the toxin concentrations measured by the hemolytic assay (n = 10).

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Fig. 3. Repeatability of the hemolytic assay. Linear regression analysis performed within the working range of the hemolytic assay $(3.9 \times 10^{-10} \text{ M} - 2.5 \times 10^{-8} \text{ M})$ by plotting the theoretical PLTX concentrations against toxin concentrations measured by the hemolytic assay. A: Intraday repeatability (n = 10). **B**: Inter-day repeatability (six months period, n = 10).

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Fig. 4. Hemolytic activity of N-biotinyl-PLTX, PLTX and its natural analogues within the working range for PLTX ($3.9 \times 10^{-10} \text{ M} - 2.5 \times 10^{-8} \text{ M}$). Each point represents the mean \pm SE of 3 different experiments. *p < 0.05; **p < 0.01; ***p < 0.001 as compared to PLTX (two-way ANOVA and Bonferroni post test).

Fig. 5. Neutralization of PLTX-induced hemolysis within the working range of the hemolytic assay $(3.9 \times 10^{-10} \text{ M} - 2.5 \times 10^{-8} \text{ M})$ by ouabain and anti-PLTX monoclonal antibody, mAb. Each point represents the mean \pm SE of 3 different experiments.*p < 0.05; **p < 0.01; ***p <0.001 as compared to PLTX (two-way ANOVA and Bonferroni post test).

Fig. 6. Matrix effect. Linear regression analysis within the working range of the hemolytic assay $(3.9 \times 10^{-10} \text{ M} - 2.5 \times 10^{-8} \text{ M})$ performed on 80% aqueous methanol mussels extracts diluted 1:50. Linear regression analysis was performed within the working range of the hemolytic assay $(3.9 \times 10^{-10} \text{ M} - 2.5 \times 10^{-8} \text{ M})$ by plotting the theoretical PLTX concentrations against toxin concentrations measured by the hemolytic assay.

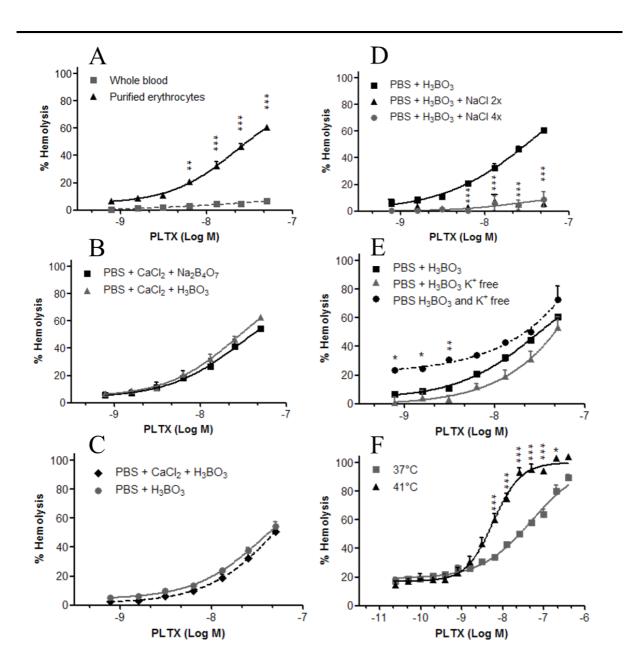
Table 1. *Bias* values (%) for PLTX analysis and intra-day (n=10, 1 day) and inter-day (n=10, 6 months) relative repeatability (RSDr %).

| Intra-day repeatability Inter-day repeatability PLTX (M) Bias (%) | | | | | | |
|---|-------------|------------------------|--------|------------------------|--------|--|
| | , 21005 (70 | Mean | RSDr % | Mean | RSDr % | |
| 3.91x10 ⁻¹⁰ | -1.2 | 4.55x10 ⁻¹⁰ | 17 | 3,86x10 ⁻¹⁰ | 10 | |
| 7.81x10 ⁻¹⁰ | -1.7 | 7.60x10 ⁻¹⁰ | 12 | 7,68x10 ⁻¹⁰ | 3 | |
| 1.56x10 ⁻⁹ | -2.0 | 1.63x10 ⁻⁹ | 20 | 1,53x10 ⁻⁹ | 8 | |
| 3.13x10 ⁻⁹ | 2.4 | 2.47x10 ⁻⁹ | 24 | 3,24x10 ⁻⁹ | 7 | |
| 6.25x10 ⁻⁹ | 0.3 | 6.51x10 ⁻⁹ | 12 | 6,27x10 ⁻⁹ | 3 | |
| 1.25x10 ⁻⁸ | -1.4 | 1.28x10 ⁻⁸ | 8 | 1,24x10 ⁻⁸ | 7 | |
| 2.50x10 ⁻⁸ | -1.7 | 3.18x10 ⁻⁸ | 14 | 2,50x10 ⁻⁸ | 7 | |
| Mean | -0.8 | | 15 | | 6 | |

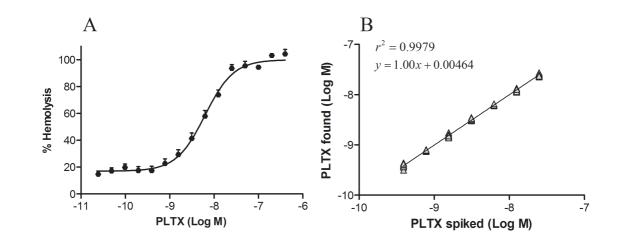
Table 2. Bias values (%) for PLTX detected in 80% aqueous methanol mussels extracts 649 spiked with the toxin after 1:50 dilution in comparison to the theoretical PLTX concentrations 650 (n=6). 651 652 653 PLTX (M) Bias (%) 654 3.91x10⁻¹⁰ -13.8 655 7.81×10^{-10} -12.8 656 1.56x10⁻⁹ 12.9 3.13x10⁻⁹ 31.7 657 6.25×10^{-9} -27.0 658 1.25x10⁻⁸ 33.6 659 2.50x10⁻⁸ 0.9 Mean 3.7 660

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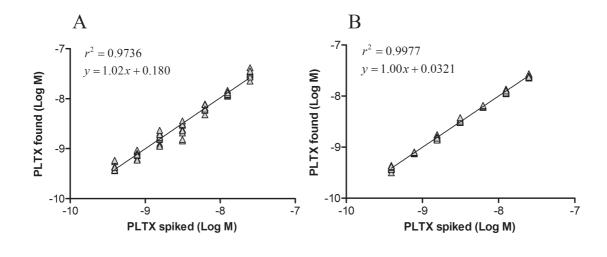


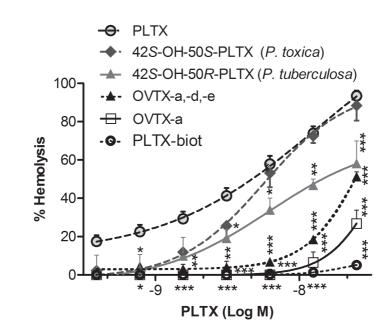




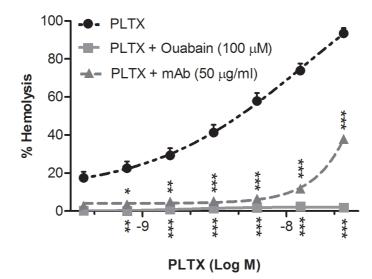


669 Figure 3





676 Figure 5



680 Figure 6

