

1 Toxicity Equivalency Factors for Tetrodotoxin Analogues

2 Determined with Automated Patch Clamp on Voltage-Gated

3 Sodium Channels in Neuro-2a Cells

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15 Abstract

16 Tetrodotoxin (TTX) is a potent marine neurotoxin, responsible for numerous poisoning incidents and some
17 human fatalities. The pufferfish *Lagocephalus sceleratus* is one of the main carriers of this toxin in nature,
18 which coexists with a variety of TTX analogues. To date, more than thirty TTX analogues have been
19 identified, but their individual toxicities and roles in poisoning remain largely unknown. In this work, the
20 toxicity equivalency factors (TEFs) of five TTX analogues were determined by assessing the blockade of
21 voltage-gated sodium channels (VGSCs) in Neuro-2a cells using automated patch clamp (APC). Among the
22 TTX analogues tested, 11-norTTX-6(S)-ol exhibited the highest toxicity, followed by 11-deoxyTTX, 6,11-
23 dideoxyTTX, 5,11-dideoxyTTX and 5,6,11-trideoxyTTX. All analogues were less toxic than TTX. The derived
24 TEFs were applied to the individual TTX analogues concentrations measured in fifteen pufferfish samples
25 from Greece, using liquid chromatography coupled to tandem mass spectrometry (LC-MS/MS). A
26 comparison of these results with those obtained from APC analysis demonstrated that TEFs can be
27 effectively used to translate LC-MS/MS analytical data into meaningful toxicological information. This is
28 the first study to utilise APC device for the toxicological assessment of TTX analogues, highlighting its
29 potential as a bioanalytical tool for seafood safety management and human health protection.

30 Keywords

31 Tetrodotoxin; tetrodotoxin analogue; toxicology; single-cell analysis; pufferfish; food safety.

33 Tetrodotoxins (TTXs) are a family of highly toxic neurotoxins, which received this name because they were
34 first isolated from *Tetraodontidae* fish (Katikou et al., 2022). In natural samples, TTX coexists with several
35 TTX analogues, sharing a similar overall structure but differing at specific molecular positions. To date,
36 over thirty TTX analogues have been described, yet their individual toxicities and role in poisoning
37 incidents remain largely undefined (Katikou et al., 2022). The pufferfish *Lagocephalus sceleratus* is one of
38 the main carriers of TTXs, although these toxins are also found in a variety of marine and terrestrial
39 organisms (Bane et al., 2014). The presence of TTXs in organisms from different phyla suggests an
40 exogenous origin, which has been linked to various bacteria from the *Proteobacteria* phylum (Magarlamov
41 et al., 2017). However, the metabolic pathways for TTXs biosynthesis and/or biotransformation in hosts
42 are still unclear (Kono et al., 2008; Ueyama et al., 2018). The biological target of TTXs is the voltage-gated
43 sodium channel (VGSC, also known as Na_v), an important membrane protein involved in the generation
44 and propagation of action potentials in neuronal and other excitable tissues (Catterall, 2012). The toxicity
45 mechanism of TTXs relays on the blockade of the VGSCs, inhibiting Na^+ influx into the cell and therefore
46 preventing cell membrane depolarisation (Tukker et al., 2023). Depending on the dose, TTX can be useful
47 for some medical applications (Nieto et al., 2012; Bucciarelli et al., 2021; González-Cano et al., 2021) or
48 can cause poisoning (Guardone et al., 2019). In this case, symptoms ranging from mild neurological or
49 gastrointestinal disorders (e.g., oral tingling or vomiting) to severe systemic failures have been described
50 that, in severe cases, can lead to death (Bentur et al., 2008; Katikou et al., 2022). Currently, there is no
51 specific treatment for TTX poisoning other than palliative care to relieve the symptoms. For this reason,
52 the prevention of TTX poisoning highly depends on the early detection of TTX and TTX analogues in
53 seafood.

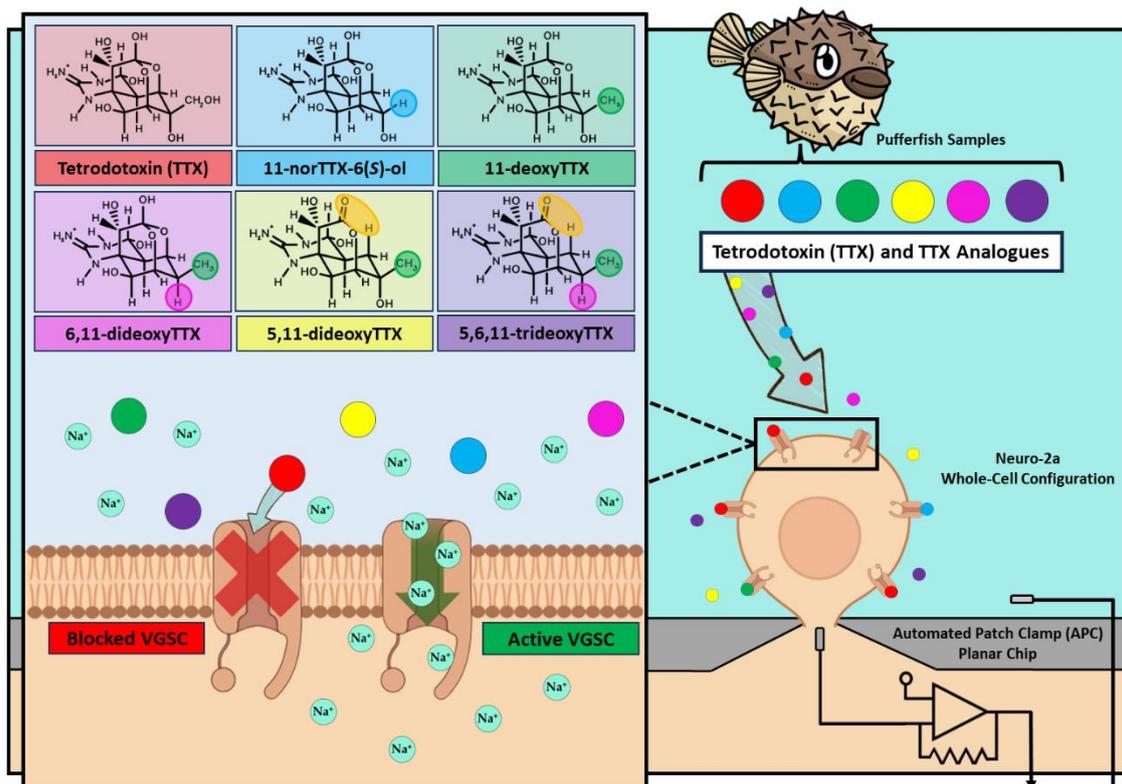
54 Tetrodotoxin poisoning events are predominantly reported in Asian countries where pufferfish is eaten
55 as a delicacy (Bane et al., 2014). However, poisoning cases have also been reported in non-endemic
56 regions, highlighting a potential new food safety concern if appropriate control measures are not
57 implemented (Reverté et al., 2023). In Europe, poisonous fish of the *Tetraodontidae*, *Molidae*,
58 *Diodontidae* and *Canthigasteridae* families must not be placed on the market due to the threat they may
59 pose to human health (Regulation (EC) No 854/2004). However, with *L. sceleratus* establishing itself as a
60 new persistent invasive species in the Mediterranean (Akyol et al., 2005; Katikou et al., 2009; Rambla-
61 Alegre et al., 2017; Coro et al., 2018; Guardone et al., 2018; Akbora et al., 2020; Christidis et al., 2021;
62 Anastasiou et al., 2023; Kosker et al., 2023), concerns arise regarding the potential risks associated with
63 the accidental consumption of toxic pufferfish by local populations as a consequence of recreational
64 fishing (Anastasiou et al., 2023; Ulman et al., 2024) and the possible intermingling of juveniles with other
65 small pelagic species (e.g., anchovy, sardines, picarel and bogue) (Leonardo et al., 2019). Moreover, the
66 risk of exposure to TTX through the consumption of common edible seafood species (originally non-toxic)
67 that have been cross-contaminated as a result of accidental pufferfish predation (Malloggi et al., 2023;
68 Tinacci et al., 2023) cannot be ignored. In addition, over the last few decades, shellfish has been identified

69 as another important TTX carrier in Europe (Rodriguez et al., 2008; Silva et al., 2012; Turner et al., 2015;
70 Vlamis et al., 2015; Gerssen et al. 2018; Leão et al., 2018; Reverté et al., 2018; Blanco et al., 2019;
71 Dell'Aversano et al., 2019; Campàs et al., 2020; Hort et al., 2020; Bacchiocchi et al., 2021; Bordin et al.,
72 2021; Réveillon et al., 2021; Alkassar et al., 2024). Although TTX concentrations in shellfish are generally
73 lower than those in pufferfish, they still pose a poisoning risk. Indeed, the first TTX poisoning case
74 originated in Europe was linked to the consumption of contaminated gastropods (Rodriguez et al., 2008;
75 Fernández-Ortega et al., 2010).

76 In general, instrumental analysis techniques, such as liquid chromatography coupled to tandem mass
77 spectrometry (LC-MS/MS), are the gold standard for monitoring regulated toxins in Europe, including
78 lipophilic marine toxins (Regulation (EC) No 15/2011). The main advantage of these techniques is the
79 ability to detect and quantify not only the target toxin but also several of its individual analogues, thus
80 providing a toxin profile for a sample (Jang et al., 2010; Rambla-Alegre et al., 2017; Rambla-Alegre et al.,
81 2018; Alkassar et al., 2023; Hong et al., 2023; Park et al., 2024). However, these analyses do not provide
82 direct toxicological insights (Reverté et al., 2023), requiring the use of toxicity equivalency factors (TEFs)
83 for the different analogues to transform analytical data into practical toxicological information. The scant
84 knowledge about the individual toxicities of emerging marine toxins, including the TTX analogues,
85 emphasises the increasing need for their thorough toxicological evaluation (Botana et al., 2017).

86 Historically, animal bioassays, such as the mouse bioassay (MBA), were the primary methods for the
87 toxicological characterisation of marine toxins (Reverté et al., 2023). However, MBA is now of limited use
88 in many countries due to its low specificity, high variability and the ethical issues concerning the
89 experimentation with live animals (Campàs et al., 2021). Cell-based assays (CBAs) have been proposed as
90 an alternative approach to assess the toxicity of a sample without involving live animals. The detection of
91 toxins with CBA relies on assessing changes in the morphology or viability of cells after their exposure to
92 them (Kogure et al., 1988; Manger et al., 1993; Manger et al., 1995). However, cells used for CBA
93 sometimes require pretreatments with auxiliary drugs to ensure optimal assay performance, necessitating
94 cautious interpretation of the toxicological data that they generate. Recently, a new approach that
95 leverages the electrophysiological activity of Neuro-2a cells has emerged as a promising high throughput
96 bioanalytical method for detecting TTXs in pufferfish samples, with potential applications in food safety
97 and broader toxicological studies (Campàs et al., 2024).

98 In this work, we characterised the toxicity of five TTX analogues using an automated patch clamp (APC)
99 device (Figure 1). By assessing the toxicological response of Neuro-2a cells to 11-norTTX-6(S)-ol, 11-
100 deoxyTTX, 6,11-dideoxyTTX, 5,11-dideoxyTTX and 5,6,11-trideoxyTTX, the respective TEF values were
101 determined. Then, we applied these TEFs as correction factors to the concentrations of individual TTX
102 analogues measured in various pufferfish samples from Greece using LC-MS/MS. The overall toxicity levels
103 calculated in TTX equivalents (equiv.) through this indirect approach closely matched those directly
104 measured in the samples using APC, suggesting a consistent agreement between the two techniques. This
105 study represents the first application of APC in evaluating the toxicological potency of TTX analogues.



106 **Figure 1.** The diagram on the right outlines the strategy for assessing the activity of voltage-gated sodium channels
 107 (VGSCs) in Neuro-2a cells exposed to TTX or the TTX analogues studied using the APC method. The close-up view on
 108 the left illustrates the VGSC blockade by the toxins and the inhibition of Na⁺ influx into the cells. Structural differences
 109 between TTX and its analogues are highlighted by different colours (upper left).

110 [Colour should be used for Figure 1 in print]

111 2. Materials and Methods

112 2.1. Reagents

113 TTX (purity ≥98% by HPLC) was purchased from Tocris Bioscience (Bristol, UK) and a standard solution was
 114 prepared at 1 mg/mL in 1% (v/v) acetic acid. In the APC method, the external solution (140 mM NaCl, 4
 115 mM KCl, 1 mM MgCl₂, 2 mM CaCl₂, 5 mM D-glucose monohydrate, 10 mM HEPES/NaOH, pH 7.4), the
 116 internal solution (50 mM CsCl, 10 mM NaCl, 60 mM CsF, 20 mM EGTA, 10 mM HEPES/CsOH, pH 7.2) and
 117 the seal enhancer solution (10 mM HEPES, 130 mM NaCl, 5 mM glucose, 4 mM KCl, 10 mM CaCl₂, 10 mM
 118 MgCl₂, pH 7.4, osmolarity: 302 mOsmol) were obtained from Nanion Technologies GmbH (Munich,
 119 Germany). Murine neuroblastoma (Neuro-2a) cells were purchased from ATCC LGC Standards (Manassas,
 120 VA, USA). Foetal bovine serum (FBS), penicillin/streptomycin solution, Roswell Park Memorial Institute
 121 (RPMI-1640) medium, sodium pyruvate and trypsin-EDTA enzyme were purchased from Merk KGaA
 122 (Darmstadt, Germany).

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125 *2.2. Pufferfish samples and tetrodotoxin extraction*

126 Three pufferfish specimens, two males (PF1, PF2) and one female (PF3), identified as *L. sceleratus* by
127 morphological evaluation, were collected from the Libyan Sea near Crete, Greece in May (PF1, PF2) and
128 March (PF3) of 2019. The specimens were dissected and their tissues, including skeletal muscle, skin, liver,
129 intestinal tract and gonads, were separated. The extraction of TTXs was performed following the protocol
130 described by Reverté and co-workers (Reverté et al., 2015). Briefly, 10 g of each tissue was homogenised
131 using an Ultraturrax blender at full speed. To each homogenised sample, 25 mL of 0.1% (v/v) acetic acid
132 was added and the mixture was vortexed for 2 min at 2500 rpm. Subsequently, the tubes were placed in
133 a boiling water bath for 10 min with occasional stirring. The homogenates were then cooled down and
134 centrifuged at 2500 rpm for 5 min (4 °C). The supernatants were collected and the pellets were subjected
135 to a second extraction with additional 20 mL of 0.1% (v/v) acetic acid. The two supernatants were pooled,
136 and their final volume was adjusted to 50 mL with 0.1% (v/v) acetic acid. For liver samples, a liquid-liquid
137 partitioning of the crude extract with hexane (1:1) was necessary to remove fats. For APC analysis, the
138 extracts were passed through 0.45-µm nylon syringe filters. For LC-MS/MS analysis, filtration was
139 conducted using 3000-Da molecular sieve filters followed by 0.2-µm nylon filters. The final extracts, at 200
140 mg pufferfish tissue per mL, were stored at -20 °C until analysed.

141 *2.3. Isolation and purification of tetrodotoxin analogues*

142 TTX analogues were sourced from PF2 liver tissue, as detailed in Alkassar et al., 2023. In brief, a 30 g liver
143 tissue extract was reduced to 5 mL using a rotary evaporator system, mixed with 5 mL of acetonitrile and
144 fractionated by hydrophilic interaction liquid chromatography (HILIC) at room temperature utilising a
145 prep-LC column Luna HILIC AXIA (250 mm × 21.2 mm, 5 µm particle size; Phenomenex, Torrance, CA, USA).
146 A binary gradient elution at 10 mL/min was applied with Milli-Q water (mobile phase A) and
147 acetonitrile/water (90/10 v/v, mobile phase B), both containing 30 mM ammonium acetate at pH 5.8. The
148 gradient program started at 100% B, and it was kept isocratic for 5 min; then, phase B was reduced to 95%
149 B at 35 min, further decreased to 82.5% B at 80 min, held isocratic for 5 min and returned 100% B at 90
150 min. A total of ten chromatographic runs were performed on 1 mL aliquots and the TTXs fractions
151 collected (10-mL per run) were pooled together (100 mL in total) and stored at -20 °C until used. The TTXs
152 present in the pooled fractions were analysed by LC-MS/MS, using a triple quadrupole mass spectrometer
153 (Xevo TQ-XS, Waters Corporation, Milford, MA, USA) coupled to a UPLC binary pump system (Acquity
154 UPLC I-plus-Class, Waters Corporation, Milford, MA, USA). Further experimental details are provided in a
155 previous work (Alkassar et al., 2023). The fractions were dried under a N₂ stream at room temperature
156 and reconstituted in the external buffer before being analysed with APC.

157 *2.4. Neuro-2a maintenance and automated patch clamp recording*

158 Neuro-2a cells were cultured and maintained in RPMI-1640 medium supplemented with 10% (v/v) FBS,
159 1% (v/v) penicillin-streptomycin and 1% (v/v) sodium pyruvate at 37 °C in a 5% CO₂ humid atmosphere.
160 All Neuro-2a cells used in this work were between passages 245 and 255.

161 The electrophysiological changes of Neuro-2a cells in response to the TTX analogues were evaluated using
162 a Patchliner (Nanion Technologies GmbH, Munich, Germany), an automated planar patch clamp device
163 (Campàs et al., 2024). Briefly, eight wells of a medium resistance NPC-16 borosilicate planar chip (Nanion
164 Technologies GmbH, Munich, Germany) were filled up with the internal and external solutions. Then, a
165 suspension of Neuro-2a cells, prepared at 100,000 cells/mL in a 1:1 mixture of RPMI and external solution,
166 was introduced into the chip. A single cell was immobilised on the hole located at the bottom of each well
167 with a holding potential of -30 mV. Then, after adding the enhancer solution, the potential was changed
168 to -100 mV. Successful patching was confirmed by achieving a stable Giga-sealing with resistances (higher
169 than 1 G Ω).

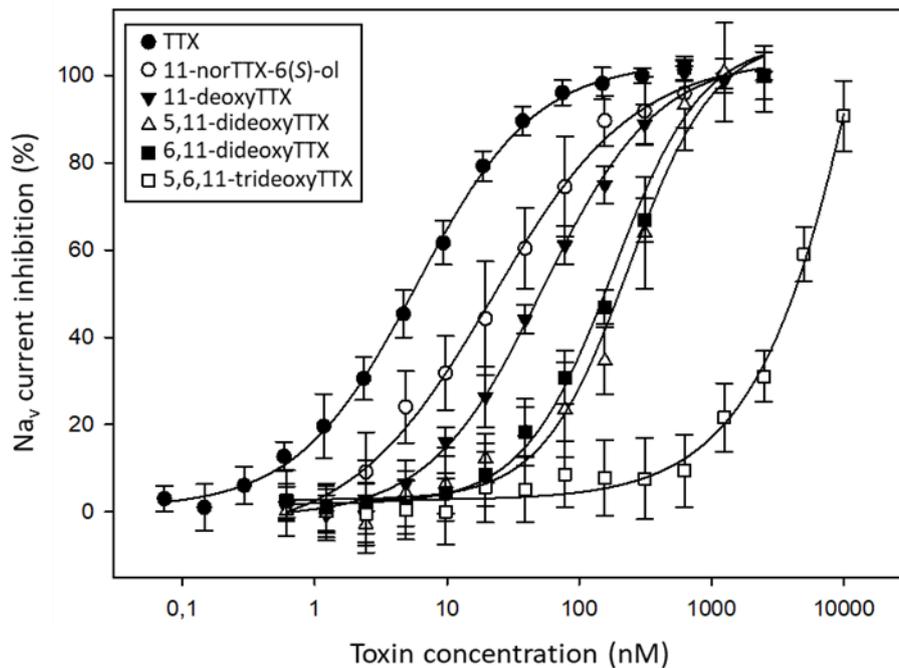
170 The I_{NaV} currents were measured by applying the potential in increments of 10 mV from -80 mV to 40 mV
171 using two EPC Quatro USB amplifier units (8 probes) (HEKA Elektronik, Stuttgart, Germany) controlled and
172 digitalised in real time with the Patchmaster software (Nanion Technologies GmbH, Munich, Germany).
173 For preparing the dose-response curves of TTX and TTX analogues, as well as for analysing the pufferfish
174 extracts, 15 μ L of sample was injected into each well of the chip at a flow rate of 30 μ L/s. Dose-response
175 concentrations ranged as follows: for TTX from 0.07 nM to 300 nM; for 11-norTTX-6(S)-ol, 11-deoxyTTX,
176 6,11-dideoxyTTX and 5,11-dideoxyTTX analogues from 0.7 nM to 2500 nM; for 5,6,11-trideoxyTTX from
177 2.44 nM to $10,000$ nM. The concentrations of pufferfish tissue equivalents for their analysis ranged from
178 0.002 to 10 mg/mL. All measurements were performed, at least, in quadruplicate.

179 **3. Results and Discussion**

180 *3.1. Toxicological characterisation of tetrodotoxin analogues*

181 The detailed profile of TTXs in the concentrated PF2 liver extract, which served as a source material for
182 the isolation of TTX analogues, was previously characterised by LC-MS/MS (Alkassar et al., 2023) and it
183 was shown to contain TTX and thirteen of its analogues. However, after the fractionation process, only
184 five of these analogues (i.e., 11-norTTX-6(S)-ol, 11-deoxyTTX, 6,11-dideoxyTTX, 5,11-dideoxyTTX and
185 5,6,11-trideoxyTTX), which were the most abundant, were isolated in sufficient quantity and grade of
186 purity (90% to 95%) for the subsequent toxicological characterisation with APC. The grade of purity of the
187 fractions corresponding to these TTX analogues was calculated as the percentage of the major TTX
188 analogue with respect to the total amount of all TTX analogues found in those fractions. The composition
189 of the fractions used in this work is detailed in the "Supplementary material" section (Figure S1).

190 To characterise the toxic potential of the isolated toxins, dose-response curves describing the I_{NaV} current
191 inhibition in Neuro-2a cells exposed to varying concentrations of each TTX analogue were constructed by
192 assessing cells' electrophysiological activity with APC and compared with the dose-response curve of TTX.
193 All data were normalised with respect to the I_{NaV} currents measured for each individual cell after patching
194 stabilisation and before toxin exposure and fitted to a sigmoidal logistic four-parameter equation (Figure
195 2).



196 **Figure 2.** Dose-response curves describing the sodium current inhibition in Neuro-2a cells exposed to tetrodotoxin
 197 (TTX) and TTX analogues through electrophysiological activity assessment with automated patch clamp (APC). Each
 198 point represents the average \pm standard deviation (at least, $n = 4$).

199 [Colour should not be used for Figure 2 in print]

200 The half-maximal inhibitory concentration (IC_{50}) of TTX and its analogues were derived from the dose-
 201 response curve equations and served as a measure of toxin potency. These values were then used to
 202 calculate the toxicity equivalency factors (TEFs) for the different TTX analogues (Table 1), defined as the
 203 ratio of the IC_{50} value of TTX to that of each analogue. All TTX analogues presented lower toxicity than TTX
 204 ($TEF < 1$). Among the analogues tested, 11-norTTX-6(S)-ol was the most toxic followed by 11-deoxyTTX,
 205 6,11-dideoxyTTX, 5,11-dideoxyTTX and 5,6,11-trideoxyTTX. The differences in toxicity can be attributed to
 206 the distinct structures among TTX and its analogues, which influence how these molecules interact with
 207 the target sites on VGSCs (Yotsu-Yamashita et al., 1999; Bane et al., 2016). Additionally, changes in the
 208 amino acid sequences of the VGSC have been shown to result in different TTX affinities (Tsukamoto et al.,
 209 2017). The five TTX analogues can be categorised into three groups based on the number of structural
 210 modifications compared to TTX: 11-norTTX-6(S)-ol and 11-deoxyTTX with a single modification, 6,11-
 211 dideoxyTTX and 5,11-dideoxyTTX with a double modification, and 5,6,11-trideoxyTTX with a triple
 212 modification (Figure 1).

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216 **Table 1.** Half-maximal inhibitory concentration (IC₅₀) and toxicity equivalency factors (TEFs) for tetrodotoxin (TTX) and
217 TTX analogues determined by electrophysiological activity assessment of Neuro-2a cells with automated patch clamp
218 (APC).

Toxin	IC ₅₀ (nM)	TEF
TTX	5.43	1.000
11-norTTX-6(S)-ol	22.9	0.238
11-deoxyTTX	50.8	0.107
6,11-dideoxyTTX	157	0.035
5,11-dideoxyTTX	200	0.027
5,6,11-trideoxyTTX	4196	0.001

219 The main difference between the two TTX analogues with a single modification is that the hydroxymethyl
220 group at position C-6 of TTX is fully substituted by a hydrogen in 11-norTTX-6(S)-ol, whereas it is replaced
221 by a methyl group in 11-deoxyTTX. Interestingly, the APC analysis revealed that the toxicity of 11-norTTX-
222 6(S)-ol and 11-deoxyTTX was 4- and 9-fold lower compared to TTX, respectively. Considering that both
223 analogues were lacking the hydroxyl-group at C-11 position, it is reasonable to suggest that this group is
224 crucial for the efficient interaction of TTXs with VGSCs and its absence leads to significantly lower toxicity.
225 These findings align well with those of previous studies, where it was hypothesised that the specific
226 hydroxyl group was responsible for the formation of a hydrogen bond between TTX and VGSCs (Yotsu-
227 Yamashita et al., 1999; Choudhary et al., 2003). Nevertheless, it is important to note that although none
228 of these two TTX analogues have the hydroxyl at C-11 position, 11-norTTX-6(S)-ol is twice as toxic as 11-
229 deoxyTTX, suggesting that the exact type of modification at this site has also an important effect on
230 toxicity. It appears that the hydrogen substituent in 11-norTTX-6(S)-ol has a less adverse impact on other
231 non-covalent interactions with VGSCs as compared to the bulkier methyl substituent in 11-deoxyTTX,
232 implying that some steric effects also play a role in the toxicity of these analogues.

233 Regarding the two TTX analogues with double modification, both share the same alteration at C-11
234 position as seen in 11-deoxyTTX. Additionally, 6,11-dideoxyTTX has a hydroxyl group at C-6 position
235 replaced by a hydrogen, whereas the ether group at C-5 position of 5,11-dideoxyTTX, which is part of the
236 2,4-dioxaadamantane skeleton, is also replaced by a hydrogen. APC analysis revealed that the toxicity of
237 6,11-dideoxyTTX and 5,11-dideoxyTTX was 29- and 37-fold lower compared to TTX, respectively. Based
238 on these results, the hydroxyl group at C-6 position seems to have a lower impact on toxicity compared
239 to the hydroxyl group at C-11 position, as evidenced by the greater toxicity reduction between TTX and
240 11-deoxyTTX than between 11-deoxyTTX and 6,11-dideoxyTTX. These results are aligned with the findings
241 of other studies (Kudo et al., 2014). As for the C-5 modification in the 2,4-dioxaadamantane structure, its
242 impact on toxicity appears to be greater than that of the hydroxyl group modification at C-6 position (i.e.,
243 the toxicity of 5,11-dideoxyTTX is 22% lower than the toxicity of 6,11-dideoxyTTX). A notable reduction in

244 toxicity in 5-deoxyTTX (that has the same modification at C-5 position than 5,11-dideoxyTTX) was also
245 observed in other works (Satake et al., 2014).

246 Finally, 5,6,11-trideoxyTTX combines the three modifications described for 6,11-dideoxyTTX and 5,11-
247 dideoxyTTX. Its toxicity was 773-fold lower than that of TTX, which is aligned with results from previous
248 papers (Yotsu-Yamashita et al., 1995). This huge decrease in toxicity indicates that the combination of the
249 different modifications exerts a synergistic effect. It is clear that C-11, C-6 and C-5 positions are important
250 for the overall toxicity of TTXs to a greater or lesser extent.

251 This is the first time that APC technology has been used to characterise the toxicity of TTX analogues and
252 elucidate their TEFs. Literature on the relative toxicity of TTX analogues is rather limited and fragmented,
253 complicating the comparison of our results with those of previous studies. Nevertheless, all TTX analogues
254 tested by APC exhibited lower toxicity than TTX, aligning with the findings from MBA or CBA methods
255 (Knutsen et al., 2017). The TEFs obtained with APC for 11-norTTX-6(S)-ol, 11-deoxyTTX, 6,11-dideoxyTTX
256 and 5,6,11-trideoxyTTX are aligned with some previously obtained with CBA and MBA (Table 2). The only
257 analogue which diverged from the toxicity trend of TTXs was 5,11-dideoxyTTX, with a TEF of 0.027 via APC
258 compared to 0.750 via CBA (Alkassar et al., 2023). The discrepancy in TEFs could stem from differences in
259 the grade of purity of the TTX analogues used, since the presence of other compounds may either increase
260 or decrease the toxicological response. However, this does not seem to be the case as the purities of the
261 5,11-dideoxyTTX fractions used for the determination of the TEF with CBA (Alkassar et al., 2023) and APC
262 (this work) were similarly high (i.e. 98 and 95%, respectively). Since the same cell model has been used
263 for CBA and APC (i.e., Neuro-2a cells), this variability on TEFs is more likely due to the distinct detection
264 principles of the two methods. The detection principle of APC is based on the assessment of the changes
265 in the electrophysiological activity of cells after their exposure to toxins. In contrast, CBA evaluates the
266 changes in cell viability caused by the toxins in cell cultures pretreated with two auxiliary drugs: ouabain,
267 a Na^+/K^+ ATPase pump blocker, and veratridine, a VGSCs inactivation inhibitor. The combination of these
268 two drugs enhances the accumulation of Na^+ in the cytosol, inducing cell mortality. The blocking of VGSCs
269 by TTX or its analogues counteracts the cytotoxic effect of these two drugs. Therefore, the toxicological
270 response obtained with CBA after exposing the cells to these toxins is a consequence of an induced cellular
271 state that is not physiologically natural and that is influenced by the activity of ouabain and veratridine
272 (any change in ouabain/veratridine concentrations may affect the response of the cells to toxins). As a
273 result, toxicological data obtained with CBA must be interpreted with caution, whereas APC results may
274 more accurately reflect the actual effects encountered in the human body during poisoning events.

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278 **Table 2.** Toxicity equivalency factors (TEFs) for the tetrodotoxin (TTX) analogues evaluated with automated patch
 279 clamp (APC), cell-based assay (CBA) and mouse bioassay (MBA).

TTX analogue	TEF	Method	Reference
	0.238	APC	(This work)
11-norTTX-6(S)-ol	0.404	CBA	(Alkassar et al., 2023)
	0.19	MBA	(Yotsu et al., 1992)
	0.107	APC	(This work)
11-deoxyTTX	0.139	CBA	(Alkassar et al., 2023)
	0.14	MBA	(Yasumoto et al., 1988)
	0.035	APC	(This work)
6,11-dideoxyTTX	0.005	CBA	(Kudo et al., 2014)
	0.02	MBA	(Jang et al., 2007)
	0.027	APC	(This work)
5,11-dideoxyTTX	0.750	CBA	(Alkassar et al., 2023)
	0.001	APC	(This work)
5,6,11-trideoxyTTX	0.011	CBA	(Alkassar et al., 2023)
	0.01	MBA	(Yotsu-Yamashita et al., 1995)

280 *3.2. Pufferfish toxicity analysis with automated patch clamp*

281 To understand the impact that multi-TTX mixtures from real biological extracts may have on APC response,
 282 fifteen pufferfish samples (five different tissues from three *L. sceleratus* specimens) were analysed with
 283 APC and the results were compared with those from CBA and LC-MS/MS analyses performed on the same
 284 pufferfish extracts in a previous work (Alkassar et al., 2023) (Table 3).

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293 **Table 3.** Concentration of tetrodotoxins (TTXs) in five tissues (gonads (G), skeletal muscle (M), liver (L), digestive tract
 294 (D) and skin (S)) from three *Lagocephalus sceleratus* pufferfish (PF1, PF2 and PF3) determined by automated patch
 295 clamp (APC), cell-based assay (CBA) and liquid chromatography coupled to mass spectrometry (LC-MS/MS). For APC
 296 and CBA, toxin concentrations are expressed as μg TTX equiv./kg of pufferfish tissue. For LC-MS/MS, the concentration
 297 of only TTX or all TTXs concentrations are expressed as μg TTX or TTX+TTX analogues/kg of pufferfish tissue.

Sample	APC	CBA*	LC-MS/MS*		
			[TTX]	Σ [TTXs]	
PF1	G	5150 \pm 575	3657 \pm 472	1490 \pm 101	2129
	M	5738 \pm 1160	5559 \pm 1027	4488 \pm 1161	7640
	L	24,982 \pm 1505	21,454 \pm 2598	18,616 \pm 668	38,917
	D	25,050 \pm 276	19,584 \pm 4105	17,907 \pm 2299	34,646
	S	6324 \pm 312	8032 \pm 454	5036 \pm 356	14,251
PF2	G	13,588 \pm 1959	6365 \pm 257	5193 \pm 119	10,541
	M	22,484 \pm 3105	14,091 \pm 1028	14,313 \pm 2572	36,486
	L	74,215 \pm 4259	51,351 \pm 2318	48,745 \pm 7611	188,240
	D	86,994 \pm 8655	92,425 \pm 16,021	66,731 \pm 10,052	205,770
	S	14,147 \pm 3096	16,117 \pm 1012	12,550 \pm 27	63,178
PF3	G	222,320 \pm 48,038	228,881 \pm 12,229	161,798 \pm 3420	1,324,439
	M	6505 \pm 855	8479 \pm 943	5902 \pm 260	11,563
	L	17,456 \pm 2850	26,760 \pm 2443	15,763 \pm 624	57,179
	D	87,619 \pm 5896	113,128 \pm 9840	71,809 \pm 4773	210,873
	S	16,031 \pm 3239	14,659 \pm 1031	8334 \pm 836	34,746

298 *Data from Alkassar and co-workers (Alkassar et al., 2023).

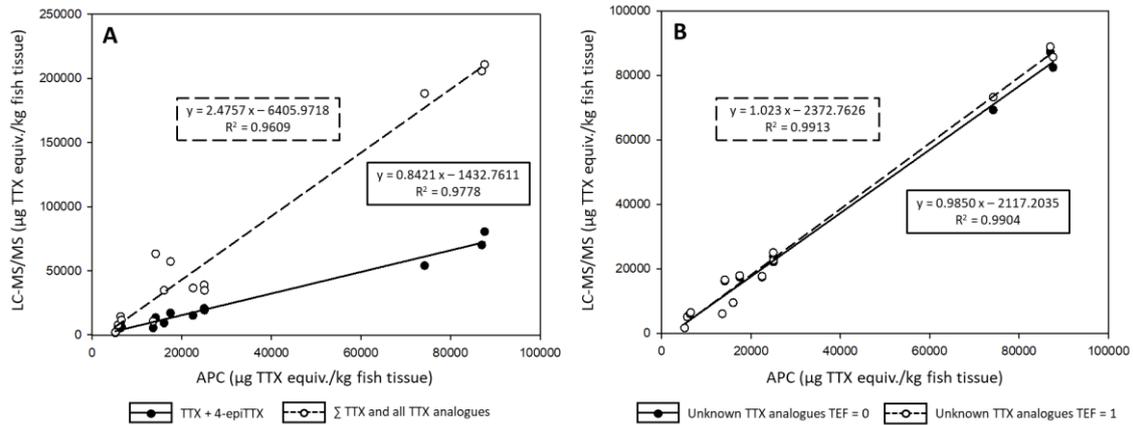
299 All samples analysed with APC showed TTXs concentration exceeding the Japanese official regulatory limit
 300 of 2000 μg TTX equiv./kg. Considering the total TTXs concentration, PF3, which was a female, was the
 301 most toxic pufferfish followed by PF2 and PF1, which were males. Although limited in number, these
 302 results are consistent with those reported in other studies suggesting gender-based toxicity differences,
 303 where females typically exhibit higher toxicity levels (Ikeda et al., 2010; Itoi et al., 2012; Christidis et al.,
 304 2021; Anastasiou et al., 2023). The reason why females are generally more toxic than males is not fully
 305 clear. Originally, it was thought that TTXs served exclusively as a defence/protection mechanism against
 306 predators (Williams et al., 2010). However, several studies suggested that TTXs may also play a role in
 307 sexual development and reproduction. For example, females have been proposed to use TTX as a male-
 308 attracting pheromone (Okita et al., 2013). In this work, the gonads from PF3 (ovaries) was the tissue with
 309 the highest concentration of TTXs. The accumulation of TTXs in female gonads has been observed in
 310 previous works, and it was hypothesised that the toxin-transferring mechanism during spawning may be
 311 useful for conferring protection to eggs (Wang et al., 2011; Wood et al., 2012). Regarding tissue-specific
 312 distribution, liver and digestive tract generally contained higher TTXs levels than skeletal muscle and skin,

313 which is consistent with the findings of other studies (Rodriguez et al., 2012; Kosker et al., 2016; Reverté
314 et al., 2017; Akbora et al., 2020; Christidis et al., 2021; Anastasiou et al., 2023).

315 Overall, the results from APC, CBA and LC-MS/MS analysis show consistent trends in the distribution of
316 TTXs across the different specimens and tissues. However, the TTXs concentrations estimated as TTX
317 equiv. with APC and CBA are on average 30% and 26% higher, respectively, than those measured for TTX
318 alone by LC-MS/MS. On the contrary, the concentrations derived by LC-MS/MS for the total TTXs
319 concentration (TTX and its analogues) are nearly two times higher than those obtained by APC and CBA.
320 These results may indicate that the toxin profile of a sample is influencing the TTX quantifications due to
321 the non-equally detection of TTX analogues by the three analytical techniques. As mentioned above, APC
322 and CBA are toxicological approaches, and their detection principle is based on the toxicological response
323 of cells to the sample under investigation (composite response) and depends on the toxic potencies of
324 the individual compounds. On the contrary, LC-MS/MS employs a structural detection approach that
325 identifies TTX and its analogues based on their distinct physicochemical properties and provides the molar
326 concentrations of each separate analyte independently of their toxicity. A notable example of this fact is
327 the analysis of the gonads from the PF3 sample. The TTXs concentration determined by APC for this tissue
328 was $\approx 220,000 \mu\text{g TTX equiv./kg}$, whereas by LC-MS/MS the concentration of TTX was $\approx 160,000 \mu\text{g TTX/kg}$
329 and that of TTX analogues was $\approx 1,160,000 \mu\text{g TTX analogues/kg}$. It is evident that, despite the high
330 abundance of 5,6,11-trideoxyTTX (Supplementary material, Figure S2, PF3, G, purple bar), its low toxicity
331 ($\text{TEF} = 0.001$) makes its relative contribution to the toxicological response by APC very low. Despite this
332 fact, the toxic effects from TTX analogues are still considerable and should not be overlooked, as they can
333 play an important role in poisoning especially when pufferfish tissues contain even higher concentrations
334 of these compounds.

335 *3.3. Application of toxicity equivalency factors to LC-MS/MS data*

336 The feasibility of estimating the overall toxicity of a sample by applying the TEF values determined by APC
337 to the individual TTX analogues concentrations measured by LC-MS/MS was evaluated for the fifteen
338 pufferfish samples discussed above. Initially, the APC data were plotted against the LC-MS/MS-measured
339 concentration of TTX (Figure 3A, solid line) or the sum of TTX and its analogues (Figure 3A, dotted line)
340 without applying any TEF value. In the former case, the concentration of TTX was combined with that of
341 4-*epi*TTX, since they are in chemical equilibrium in nature (epimers). Subsequently, TEFs were applied to
342 the LC-MS/MS measurements of the individual TTX analogues, and their sum was again plotted against
343 the APC-derived results (Figure 3B). Given the inability to determine TEFs for all TTX analogues present in
344 the samples, two extreme scenarios were considered: the unknown TTX analogues were assumed to be
345 non-toxic ($\text{TEF} = 0$, Figure 3B solid line) or the unknown TTX analogues were assumed to be as toxic as TTX
346 ($\text{TEF} = 1$, Figure 3B dotted line). TTXs concentrations after application of TEFs to LC-MS/MS measurements
347 of the individual TTX analogues and corresponding to correlations of Figure 3B are in Supplementary
348 material, Table S1. The gonads from PF3 were excluded from all correlations since the TTXs values
349 determined by both APC and LC-MS/MS were outliers (z-scores of 3.55 and 3.87, respectively).



350 **Figure 3.** Correlations between the tetrodotoxins (TTXs) contents determined with automated patch clamp (APC) and
 351 liquid chromatography coupled to mass spectrophotometry (LC-MS/MS) before (A) and after (B) the toxicity
 352 equivalency factors (TEFs) application. All correlations were fitted to a linear regression model and the obtained
 353 equations are shown.

354 [Colour should not be used for Figure 3 in print]

355 As illustrated in Figure 3A, LC-MS/MS tends to underestimate the TTXs concentration compared to APC
 356 when only the TTX and 4-*epi*TTX are considered (slope of 0.84), suggesting that not all the toxicological
 357 response obtained with APC can be explained by the amount of TTX/4-*epi*TTX present in sample. On the
 358 contrary, LC-MS/MS leads to an overestimation of TTXs concentration compared to APC when, in addition
 359 to TTX/4-*epi*TTX, the other TTX analogues are also taken into account (slope of 2.47). This result confirms
 360 that the different TTX analogues do not contribute equally to the toxicological response measured
 361 through APC. As shown in Figure 3B, this overestimation is corrected when TEFs are applied. Regardless
 362 of whether a TEF of 0 or 1 is applied to the unknown TTX analogues, the slope obtained is close to 1 with
 363 a coefficient of determination of 0.99. These findings demonstrate that TEFs derived from APC technique
 364 can be effectively applied as correction factors to LC-MS/MS quantifications and enable the translation of
 365 those data into practical toxicological information.

366 4. Conclusions

367 In this study, the toxicity of five TTX analogues was characterised through the assessment of the changes
 368 in the electrophysiological activity of Neuro-2a cells using APC, following their exposure to various doses
 369 of these VGSC blockers. All tested TTX analogues exhibited lower toxicity compared to TTX, with TEFs
 370 ranging from 0.238 to 0.001. The most toxic analogue was 11-norTTX-6(S)-ol, followed by 11-deoxyTTX,
 371 6,11-dideoxyTTX, 5,11-dideoxyTTX and 5,6,11-trideoxyTTX. Naturally contaminated pufferfish samples
 372 displaying diverse multi-TTX profiles were analysed using APC and the results were compared with the
 373 TTXs concentrations measured by LC-MS/MS. The differences in the TTXs concentrations determined by
 374 the two analytical techniques were greatly influenced by the presence of TTX analogues, which were not
 375 equally detected by the different techniques. These discrepancies were corrected after applying TEFs to
 376 the individual TTX analogues concentrations determined by LC-MS/MS, indicating that the TEFs derived

377 in this study can effectively serve as correction factors to convert instrumental analysis data into
378 meaningful toxicological information. The robust performance of the APC device across the different
379 specimens analysed underscores its potential as a valuable, new bioanalytical tool for assessing the toxic
380 potency of marine toxins and enhancing consumers protection against seafood-borne TTX poisonings.

381 **Acknowledgements**

382 This work would not have been possible without the intervention of Marisa Vinyals from Almirall, S.A. The
383 authors want to thank Almirall, S.A. for the kind transfer of use of the automated patch clamp equipment.
384 The authors would also like to express their gratitude to Alison Obergrussberger and her team from
385 Nanion Technologies GmbH for their technical assistance.

386 **Funding sources**

387 This research was supported by the Ministerio de Ciencia e Innovación (MICIN) and the Agencia Estatal de
388 Investigación (AEI) (Spain) through the CELLECTRA (PID2020-112976RB-C21 and PID2020-112976RB-C22)
389 project. J.R. acknowledges IRTA for his PhD grant (CPI0422). Andres Sanchez-Henao acknowledges the
390 Canary Islands Research Agency (ACIISI) for his research staff training aid through the Catalina Ruiz
391 Program (APCR2022010011). The authors also acknowledge the support from the CERCA
392 Program/Generalitat de Catalunya.

393 **CRedit author statement**

394 Conceptualisation, M.C.; methodology, J.R., M.R.-A., F.X.S. and M.C.; investigation, J.R., A.S.-H., M.M. and
395 P.P.; writing—original draft preparation, J.R. and M.C.; writing—review and editing, all authors;
396 supervision, M.C.; project administration, M.C.; funding acquisition, F.X.S and M.C. All authors have read
397 and agreed to the published version of the manuscript.

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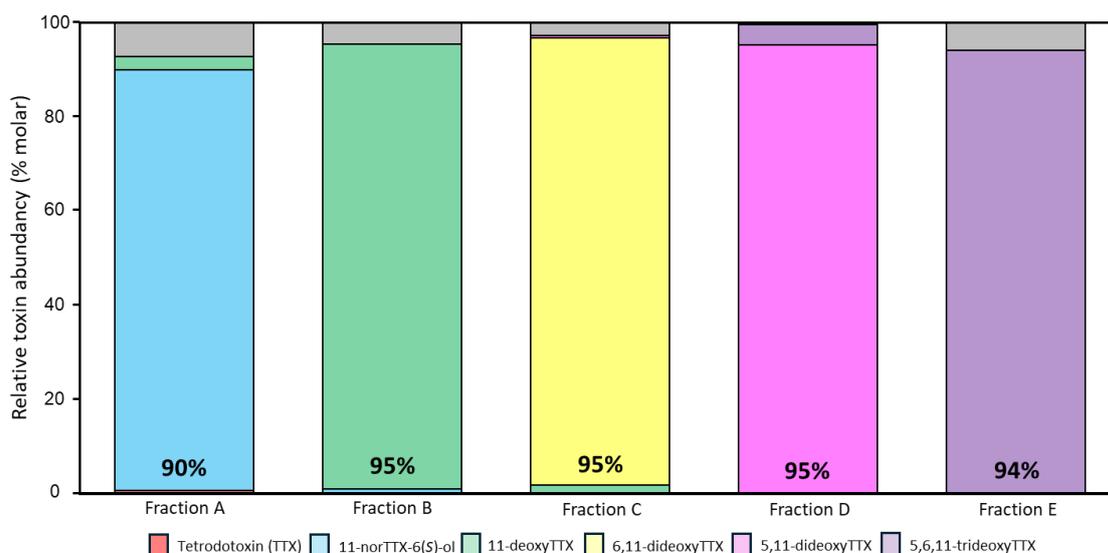
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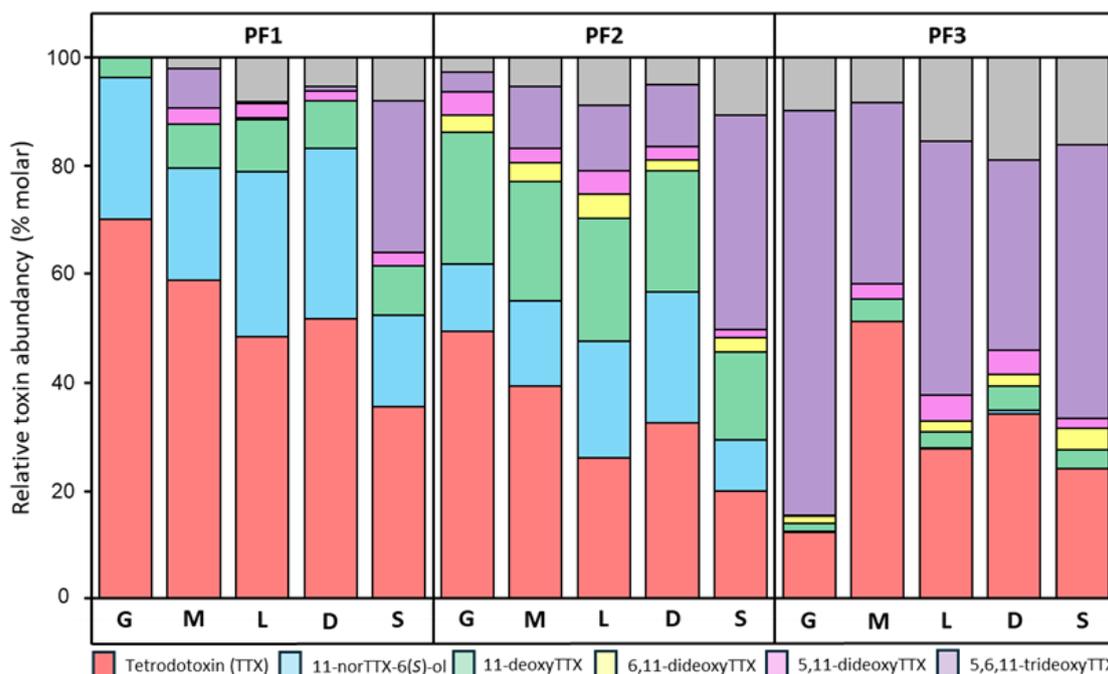
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614 **Supplementary material**



615 **Figure S1.** Relative abundance of TTX and TTX analogues in the different fractions used in this work (A-E), which have
 616 been obtained from the fractionation of a liver of a pufferfish (PF2). The TTX analogues other than TTX, 11-norTTX-
 617 6(S)-ol, 11-deoxyTTX, 6,11-dideoxyTTX, 5,11-dideoxyTTX or 5,6,11-trideoxyTTX have not been individually detailed
 618 but they have been expressed as a set in the toxin profile of the fractions (grey bars).

619 [Colour should be used for Figure S1 in print]



620 **Figure S2.** Toxin profile of the gonads (G), skeletal muscle (M), liver (L), digestive tract (D) and skin (S) of three
 621 pufferfishes from Greece (PF1, PF2 and PF3). The analogues with unknown TEFs have not been individually detailed
 622 but they have been expressed as a set in the toxin profile (grey bars).
 623

624 [Colour should be used for Figure S2 in print]

625 **Table S1.** Concentration of tetrodotoxins (TTXs) in five tissues (gonads (G), skeletal muscle (M), liver (L), digestive
626 tract (D) and skin (S)) from three *Lagocephalus sceleratus* pufferfish (PF1, PF2 and PF3) after application of TEFs to LC-
627 MS/MS measurements of the individual TTX analogues. The concentrations are expressed as µg TTX equiv./kg of
628 pufferfish tissue.

Sample	LC-MS/MS		
	Σ [TTXs]	Σ [TTXs]	
	TEF _{unknown} = 0	TEF _{unknown} = 1	
PF1	G	1631	1631
	M	5063	5063
	L	23,893	25,018
	D	22,127	22,622
	S	6018	6021
PF2	G	6022	6025
	M	17,386	17,682
	L	69,254	73,297
	D	87,300	88,921
	S	16,179	16,529
PF3	G	176,353	180,008
	M	6422	6422
	L	17,380	17,884
	D	82,469	85,700
	S	9408	9485

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Declaration of interests

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

The authors declare the following financial interests/personal relationships which may be considered as potential competing interests.