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Recovery of Normal Cytogenetic Records in Mussels After Cessation of Pollutant Effluents in Puerto

Madryn (Patagonia, Argentina)

Machado-Schiaffino · Bala · Garcia-Vazquez

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41	Abstract	The capacity of coast recovering after interruption of pollution sources has been explored employing mussels as biomarkers. In an area polluted by sewage sludge in Puerto Madryn (Argentina), abnormally high cytogenetic records (micronuclei) had been detected in the mussel <i>Mytilus edulis</i> , even higher than those obtained in this and other mussel species (<i>Brachydontes rodriguezii</i> , <i>Aulacomya atra atra</i> , <i>Perumytilus purpuratus</i>) sampled from heavily polluted industrial areas, and much higher than those recorded in samples from unpolluted areas of the same region. Normal cytogenetic patterns were recovered in Puerto Madryn less than 1 year after cessation of sewage sludge discharges, without additional treatment of the affected area. This discovery opens the possibility of considering restored coastal areas for aquaculture purposes instead of endangering natural populations in virgin areas.
42	Keywords separated by ' - '	Ecosystem recovery - Micronucleus test - Mollusks - Monitoring - <i>Mytilidae</i>
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3 NOTE

4 **Recovery of Normal Cytogenetic Records in Mussels**
 5 **After Cessation of Pollutant Effluents in Puerto Madryn**
 6 **(Patagonia, Argentina)**

7 **Gonzalo Machado-Schiaffino · Luis O. Bala ·**
 8 **Eva Garcia-Vazquez**

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 27

28 **Keywords** Ecosystem recovery · Micronucleus test ·
 29 Mollusks · Monitoring · *Mytilidae*

Introduction 30 Q2

31 Scientific analyses are critical to the ongoing effort to
 32 understand events and to improve guidelines for pollution
 33 control (Woodfield et al. 2003; Moss et al. 2005). Although
 34 recovery of marine ecosystems from pollution has tended to
 35 receive less attention than the study of new or continuing
 36 impacts, such studies are important to deal with chronic
 37 contamination (Hawkins et al. 2002). As pollution load
 38 decreases, responses would be expected at the molecular
 39 and cellular levels. These should be detectable by the
 40 growing battery of sensitive biomarkers (Depledge 1999).
 41 Among them, the micronuclei (MN) test has been applied
 42 in surveys for detection of mutagens in water ecosystems
 43 employing mussels as target species (Scarpato et al. 1990;
 44 Mersch and Beauvais 1997; Izquierdo et al. 2003).

45 Disposal of sewage sludge in the marine environment
 46 has been practiced globally and affects the seafloor and its
 47 biota even in apparently clean areas (Costello and Read
 48 1994; Studholme et al. 1995). Puerto Madryn city, located
 49 in the protected area of Peninsula Valdes, is one of the most
 50 relevant touristic points in Patagonia, principally for its
 51 marine fauna. Evidence of contamination by urban effluents
 52 had been detected in the beach in the year 2000 employing
 53 mussels as bioindicators (Izquierdo et al. 2003). Those
 54 effluents were interrupted later due to the construction of a
 55 system for urban wastes disposal.

56 The aim of this study was to evaluate the capacity of
 57 environmental recovery after cessation of urban discharges
 58 in Puerto Madryn, for exploring the possibility of re-
 59 utilizing formerly polluted sites for purposes of mussel
 60 culture. To discard the general effects of diffuse pollution in
 61 the region as a source of variation in the results obtained,
 62 two additional sites were studied as controls: a positive
 63 control subjected to intense pollution by discharge of

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64 industrial effluents and a negative control sampled in a
 65 pristine area without human activity. The micronucleus test
 66 in all mussel species present in the selected areas was
 67 chosen as a tool to monitor environmental health. Other
 68 authors used jointly MN tests and evaluation of DNA
 69 strand breaks, allowing the detection of a recent exposure
 70 (Bolognesi et al. 2004). We have chosen the MN test based
 71 on its simplicity, ease, and low cost (Sanchez-Galan et al.
 72 1998), for encouraging its future use in long-term monitor-
 73 ing in the area, where systematic pollution monitoring does
 74 not exist at the present.

Q2 75 **Materials and Methods**

Q2 76 **Sampling Locations and Characterization**

- 77 Site 1. Puerto Madryn beach. In the beach of Puerto
 78 Madryn (Patagonia, Argentina, 43° S 65° W),
 79 urban effluents (sewage outfall) had been dis-
 80 charged without treatment until September 2001,
 81 when they were interrupted. Further cleaning of
 82 the affected area was not carried out.
- 83 Site 2. Positive control. A neighboring beach with
 84 industrial effluents (fish-processing enterprise
 85 discharges) was chosen as a positive control site
 86 with conspicuous pollution.
- 87 Site 3. Negative control. Punta Este beach, 14 km east
 88 from Puerto Madryn, was chosen as an unpolluted
 89 negative control. This point can be considered
 90 completely clean because it is a virgin beach
 91 without urban, industrial, or other wastes.

92 The three locations considered are shown in a map
 93 (Fig. 1). Although there is no public information available

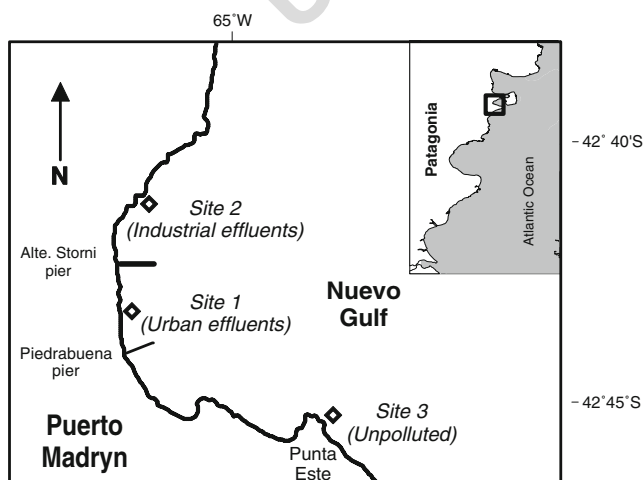


Fig. 1 Map showing the study area and the three sampling locations (Patagonia, Argentina)

about water quality in the two control sites 2 and 3, heavy
 pollution (site 2) and high water quality, respectively, are
 evident. Data on some physical–chemical parameters
 monitored in site 1 before and after interruption of sludge
 discharge are shown in Table 1. They were provided by the
 Municipality of Puerto Madryn and corresponded to official
 records of water quality obtained five times during the
 15 months previous to the interruption of discharge and one
 time 15 months after.

Target Species and Sampling Schedule 103 Q2

Mussel individuals (sample size, 12 individuals per species
 and sampling point) of similar sizes (major axis 3.4±
 1.2 cm) were sampled from each site. Gamete production
 was not observed in any case. *Mytilus edulis* was the only
 species found in site 1. In site 2 and site 3, four *Mytilidae*
 species were present: *Brachydontes rodriguezii*, *Aulacomya*
atra atra, *Perumytilus purpuratus*, and *M. edulis*. All
 mussels sampled were immediately transported to the
 laboratory alive in marine water, to avoid desiccation and
 anoxia conditions, then processed for slide preparations.

In site 1, samples were obtained just before the
 interruption of the sewage discharge (time 0, September
 2001) and 6 (time 1) and 12 (time 2) months after the
 interruption.

Tissue Sampling and Slide Preparation 118 Q2

We analyzed MN in the subpopulation of cells prevailing in
 gill tissue, following Scarpato et al. (1990). A portion of
 gill was removed with tweezers and dragged along a slide
 in a single layer of well-spread cells, then allowed to dry for
 a few minutes. Two slides per animal were prepared.

Staining procedure followed Ayllon and Garcia-Vazquez
 (2000). Briefly, slides were sequentially stained with May–
 Grünwald for 2 min; May–Grünwald/distilled water 1:1 for
 3 min; and Giemsa/distilled water 1:6 for 10 min; then
 rinsed with distilled water, allowed to dry, and mounted
 with Eukitt.

For each animal, 1,000 main gill cells (500 per slide
 whenever possible) were scored under ×1,000 magnifica-
 tion to determine the frequency of micronucleated cells.
 Coded slides were randomly sorted and scored by a single
 observer.

Statistical Analysis 135 Q2

Micronuclei frequencies were expressed per 1,000 cells (per
 mill). ANOVA tests were employed to compare MN
 frequencies between sampling sites and species. Statistical
 analyses were carried out with the SPSS 8.0 program (SPSS
 Inc.) for PC computers.

t1.1 **Table 1** Physico-chemical characteristics of sea water from site 1 (before and after, 16 months prior, and consecutive to the cessation of sewage discharge, respectively)

t1.2	Parameter	Site 1 (before)	Site 1 (after)
t1.3	pH	6.75 (1.25)	8.8
t1.4	Conductivity ($\mu\Omega\text{ cm}^{-1}$)	1,406 (433)	958
t1.5	Dissolved oxygen (mg l^{-1})	4.8 (4.7)	9
t1.6	BOD (mg l^{-1})	1,12.5 (27.5)	78
t1.7	DRP (mg l^{-1})	7.7 (0.0)	4.43

Average (standard deviation) of five and one records before and after discharge interruption, respectively

BOD biochemical oxygen demand, DRP dissolved reactive phosphorus

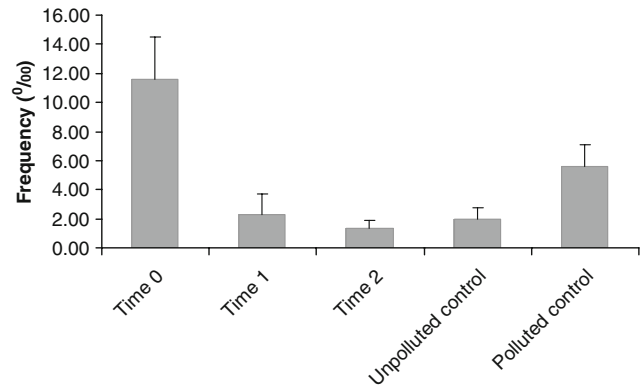


Fig. 3 *Mytilus edulis*. Mean micronuclei frequency (per mill) and standard deviation before (time 0) and after 6 (time 1) and 12 (time 2) months of the urban effluent interruption (September 2001), and in the unpolluted and polluted control areas (sites 3 and 2, respectively)

Q2 141 Results

142 Micronuclei frequencies for the different species in the
 143 polluted and unpolluted controls (sites 2 and 3, respective-
 144 ly) are shown in Fig. 2. Micronuclei averages ranged from
 145 1.5 (*P. purpuratus*, site 3) to 7.3 (*A. atra atra*, site 2).
 146 Micronuclei frequencies were significantly higher for
 147 individuals exposed to industrial effluents (site 2) than for
 148 unpolluted controls (site 3) for all four species considered
 149 ($P < 0.001$ in all cases). The four Mytilidae species were
 150 similarly sensitive to in situ pollution. Between species,
 151 significant differences were not found for MN frequency in
 152 the polluted (site 2) and in the unpolluted (site 3) stations
 153 ($P = 0.065$ and 0.3974 , respectively). The micronucleus test
 154 was sensitive enough for detecting pollution.

155 With respect to the urban sewage sludge, it was
 156 associated with noticeable nuclear damage in mussels.
 157 MN frequencies found for *M. edulis* sampled in site 1
 158 before and after the interruption of urban effluents
 159 discharges (times 0 to 2) are shown in Fig. 3, together
 160 with the average MN frequencies found in the control
 161 polluted and unpolluted sites (sites 2 and 3, respectively).
 162 The mean frequency of micronuclei scored in site 1 when

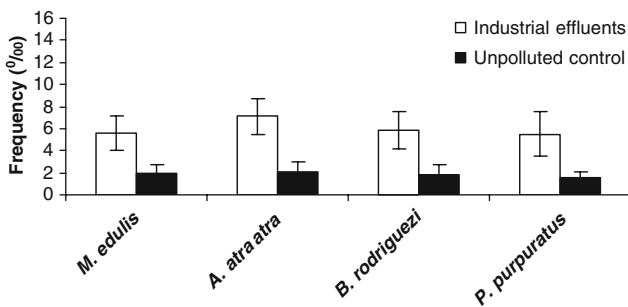


Fig. 2 Mean micronuclei frequency (per mill) and standard deviation for the four *Mytilidae* species sampled near the industrial effluents (site 2, white bars) and in the control unpolluted beach Punta Este (site 3, black bars)

the area was still subjected to pollutant discharges was
 163 11.58‰, as previously reported by Izquierdo et al. (2003)
 164 in the same area. This value was even higher than the
 165 average record obtained for this species near the heavily
 166 polluted industrial effluent in site 2. MN frequency
 167 significantly declined in the urban area to 2.29‰ and
 168 1.38‰ at 6 and 12 months, respectively, after discharge
 169 interruption ($P < 0.01$). The MN average obtained for the *M.*
 170 *edulis* samples after 12 months from the cessation of the
 171 discharges was significantly lower ($P < 0.001$) than that
 172 obtained for the polluted control (site 2), and not signifi-
 173 cantly different in MN frequencies of *M. edulis* sampled
 174 from the unpolluted site 3 ($P = 0.085$). Although it is not
 175 possible to evaluate the *M. edulis* MN base level in the area
 176 due to the lack of records previous to the beginning of the
 177 effluents, as MN frequency was similar in time 1, time 2,
 178 and unpolluted control, we could roughly take the average
 179 value (1.9‰) as the base level.
 180

181 Decreased MN frequencies after effluent cessation were
 182 consistent with improved values of physical-chemical
 183 parameters in seawater after the interruption of the urban
 184 discharge (Table 1), as revealed by increased dissolved
 185 oxygen and decreased biological oxygen demand and
 186 dissolved reactive phosphorus (signals of eutrophication).

Discussion

187 **Q2**
 188 The most remarkable result in this study was a rapid
 189 recovery of normal cytogenetic records in a mussel species
 190 (*M. edulis*) after interruption of urban discharges, without
 191 any additional management strategies such as cleaning the
 192 affected area or others. In controlled conditions, Majone et
 193 al. (1987) found persistent increased micronuclei frequen-
 194 cies in mussels for more than 1 month after interrupting

195 treatment with mitomycin C. Our results showed that less
196 than 6 months was enough to allow mussels to recover
197 baseline micronucleus frequencies after cessation of urban
198 effluents. This result is encouraging because it demonstrates
199 that genotoxic damage, detected by micronuclei tests, can
200 revert spontaneously in the wild by simple interruption of
201 the pollutant source.

202 We are aware that the main shortcoming of the paper is
203 the lack of details on the pollution status of the sampling
204 sites due to the fact that there are no pollution surveys in
205 this coast. Only some physiochemical data, such as pH and
206 biochemical oxygen demand, have been measured one time
207 after the recovery intervention. Previous evaluation of water
208 quality, where only bacterial records were made, reported
209 total and fecal coliforms in the area (Izquierdo et al. 2003),
210 those values exceeding the allowed sanitary levels in 91%
211 water samples taken along the year before effluent
212 cessation.

213 Genotoxicity biomarkers must be an integral part of the
214 suite of biomarkers considered as exposure to genotoxic
215 agents may exert a damage beyond that of the individual
216 and may be active through several generations (Magni et al.
217 2006). This study provides some additional information
218 useful for pollution-biomonitoring purposes. First, as the
219 effect of pollution on cytogenetic abnormalities can revert
220 in less than 6 months, periodic surveys should be
221 considered in shorter periods (may be monthly) if MN
222 tests are intended to be employed for routine monitoring of
223 coastal ecosystems (Smolders et al. 2002; Izquierdo et al.
224 2003); otherwise, sporadic pollution events would remain
225 undetected. Moreover, high levels of inter-individual
226 variability of the responses of the aquatic organisms should
227 be considered before applying these biomarkers for routine
228 monitoring (Burgeot et al. 1996; Bolognesi et al. 2004).
229 Seasonality is one of the factors accounting for biomarker
230 variation (Solé et al. 1995; Bolognesi et al. 2004) and
231 should also be considered, for example, in implementing
232 year-round biomonitorization protocols. Finally, in the
233 present study, micronuclei frequencies were very similar
234 for the four Mytilidae species considered in each site (with
235 higher and lower abnormality records in polluted and
236 unpolluted sites, respectively). Therefore, although *M.*
237 *edulis* is more popular for cytogenetic studies, any of them
238 could be a potential bioindicator. Although species-specific
239 characteristics have to be generally considered when
240 monitoring the health status and possible toxic effects of
241 the contaminant load in marine animals (Nyman et al.
242 2003), the cosmopolite group of mussel species could be
243 considered collectively as a suitable universal indicator for
244 in situ biomonitoring of coastal pollution. Local species
245 could be employed to monitor coastal areas, avoiding risks
246 of genetic introgression associated to specimens transfer to
247 the monitoring sites (García-Vázquez et al. 2007), as would

be required if only one target species is considered (Mersch 248
and Beauvais 1997). 249

250 Bivalve aquaculture initiatives are promoted by the
251 Argentine government, with interest of the private sector
252 for the culture of the flat oyster *Ostrea puelchana* and the
253 mussel *M. edulis*. Aquaculture of mussels has been
254 proposed as an alternative or complement of artisanal
255 shellfish harvest (Narvarte et al. 2007). The location of
256 new hatcheries is one of the first points under consider-
257 ation. The Patagonian coastline is almost totally unpolluted
258 except near urban settlements (Gil et al. 1999; Izquierdo et
259 al. 2003), and aquaculture may endanger virgin locations,
260 some of them declared natural reserves and protected areas
261 like the nearby Peninsula Valdes. Alternative usage of sites
262 already anthropized could be envisaged if environmental
263 conditions in those sites are safe and pollution does not
264 compromise hatchery productivity and consumer health.
265 The results found in this study may open the possibility of
266 envisaging the use of anthropized areas for mussel culture
267 after interrupting pollutants. However, many other analyses
268 are necessary before starting aquaculture in formerly
269 polluted areas, even if they are apparently clean. Persistent
270 chemicals such as heavy metals, PCB, and specific
271 pesticides produce an unpredictable long-term hazard in
272 the marine environment. Evidence of long-term adverse
273 effects of pollution in marine animals, for example heavy
274 metals, has been demonstrated (Domingo 1994). Although
275 from our results it seems evident that cytogenetic damage
276 revealed with micronucleus test was a reversible process,
277 additional complementary tests to evaluate long-term
278 chronic ecotoxicological effects, largely unknown for
279 aquatic biota until now (Fent 2004), should also be
280 envisaged.

281 Another issue can be the effect of mussel aquaculture on
282 the recuperated environment. Aquaculture has enormous
283 economic potential but its main drawback is the phosphorus
284 pollution it generates. In Table 1, we can observe that
285 dissolved reactive phosphorus has been reduced by almost
286 one half, from a level even higher than that existing in
287 closed systems like marine aquariums down to a level
288 which ranges within normal values (Trépanier et al. 2002).
289 If aquaculture facilities were installed in the area, phospho-
290 rus load would increase again with subsequent eutrophica-
291 tion risks (Read and Fernandes 2003). Other impacts of
292 similar nature can also be expected derived from the
293 metabolic activity of cultured mussels. On the other hand,
294 aquaculture encompasses a risk of disturbance of the
295 existing wild community. The perceived risks are often
296 associated with interactions between cultured and native
297 stocks, and the adverse effects to ecosystems; public health
298 issues are also a matter of concern (Svåsand et al. 2007).
299 The siting of aquaculture facilities plays a major role in
300 determining the impact of farmed stocks on wild popula-

301 tions (Triantafyllidis et al. 2007). Escapes or introgression
 302 of domestic individuals in wild populations is not the only
 303 risk associated with shellfish cultivation: human-induced
 304 disturbance from farming operations may also contribute to
 305 the biological patterns around mollusk cultures (Forrest and
 306 Creese 2006). A balance between socioeconomic benefits
 307 and environmental issues should be reached for sustainable
 308 shellfish culture.

309 In conclusion, this monitoring study demonstrated that
 310 the cessation of the marine disposal of sewage sludge was
 311 enough to allow recovery of normal cytogenetic records
 312 (micronucleus test) in mussels in 1 year, without additional
 313 treatment of the affected area. Although a combination of
 314 different indicators should be employed before considering
 315 the area totally recovered, this result is encouraging for
 316 opening the possibility of usage of recovered coastal
 317 systems as mollusk aquaculture sites.

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 326

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