

BIOMARKERS AS A TOOL FOR ASSESSING DIFFUSE CONTAMINATION OF COASTAL WETLAND

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ABSTRACT

Playa Penino is a natural reserve located on the south coast of Uruguay. It hosts 244 species of birds (more than 50% of the bird biodiversity recorded for Uruguay). The area is included in the International *BirdLife* Program for biodiversity preservation. Urbanization and pollution generally have affected the quality of the water and biodiversity in wetlands. The analysis of the major wetland of Playa Penino revealed high levels of organic compounds. The biomonitoring was done using the caviomorph rodent *Ctenomys pearsoni* (commonly known as tuco tuco) as an indicator species because they inhabit burrows around the wetland. Genetic effects were determined using the comet assay and micronucleus test. The significant correlations between chemical and microbiological parameters and genetic damage might indicate that macronutrients from sewage could be one of the causes of the genetic damage. There is an urgent need to conserve the biodiversity of this natural area by introducing sewage treatment, cesspool control and by controlling settlement in the area etc.

Keywords: biomarkers; comet assay; *Ctenomys pearsoni*; micronucleus test; Playa Penino; wetland

Introduction

The net cumulative effect of humanity has been to reduce biodiversity and most societies have decreased it. The direct and indirect effects of humans on the environment are ubiquitous and almost all biodiversity research and conservation must take the possibilities of human influence into consideration (Kopnina 2016). Communities that live in the proximity of protected areas have exceeded the carrying capacity of their natural environment and are unintentionally depleting resources (Sponsel 2014).

Many wetlands harbour highly diverse biological communities and provide extensive ecosystem services. However, these important ecological features are being altered, degraded and destroyed around the world (Kopnina 2016; Sievers et al. 2017). Our current lack of knowledge of individual-level responses may therefore limit our capacity to manage wetland ecosystems effectively (Sponsel 2014; Kopnina 2016).

In South America, many wetlands are remote and far from agricultural and urban pressures. However, coastal areas have been subjected to severe anthropogenic effects (Kopnina 2016; Sievers et al. 2017). Notwithstanding their critical importance, many freshwater ecosystems are considered useless. Widespread ignorance about the important benefits that wetlands provide human societies has contributed to this notion and has promoted their destruction and degradation (Kopnina 2016; Sievers et al. 2017).

Playa Penino is a natural reserve located in the south of Uruguay. It hosts 244 species of birds (more than 50% of bird biodiversity recorded in this country); 69% are

residents and 31% are migratory species, some are vulnerable or close to threat (Arballo and Bresso 2007). It is part of the International *BirdLife* Program for biodiversity preservation (CFPP 2013).

Although, Aldabe et al. (2009) mention the effect of urbanization and pollution in this area, currently the landscape has been completely altered by anthropogenic activities such as; overpopulation, irregular settlements, deforestation, introduction of livestock, invasion of exotic species, sand mining and habitat fragmentation (Rocha 1999; CFPP 2013; MVOTMA 2018). Part of the livelihood of the residents is dependent on the natural resources of the Playa Penino wetlands, such as juncus, totora, shells, sand and fish (Arballo and Bresso 2007). Although local authorities declared the area a nature reserve, protected under the category of “ecological beach” and is part of the National Protected Areas System (Arballo and Bresso 2007; MVOTMA 2018), there is no effective control over the area. Under this area is the Raigón aquifer, whose depth is less than 40 meters and is the sole source of water for more than 15,000 people, including towns, industry and rural areas (Martino et al. 2008). The shallow aquifer in sandy subsoil is severely affected by wastewater discharges and the absence of a sanitary network (MVOTMA 2018). Domestic effluents empty into filtration chambers or discharged directly into ditches next to houses.

GeoUruguay report (Martino et al. 2008) indicates that the major ecological problems are erosion, mining sand for construction and loss of biodiversity due to urban expansion and pollution of natural areas. Despite its ecological importance, there are no field studies on the levels of contamination.



Fig. 1 Aerial photograph of the main wetland of Playa Penino with the sites, where the indicator species were captured, are indicated by the white circles. The number of circles does not represent the number of individuals captured. Modified version of the photograph in Google Earth.

A biomarker is a molecular or cellular indicator of the effects of a particular environment on the cells of an indicator species (Mudry and Carballo 2006). This makes it possible to assess the environmental conditions affect an organism. The indicator species is an organism that inhabits the ecosystem studied and has some important features. For example, it is widely distributed in the area studied and its biology is well-known, and could have a key role in the comprehension of the consequences of exposure to a xenobiotic. Identifying incipient deterioration in the indicator species indicates potential dangerous situations at sub-lethal levels, either for a point or diffuse source of a xenobiotic (Ribeiro et al. 2003; Mudry and Carballo 2006).

In environmental biomonitoring, genotoxic biomarkers are used to assess the risks to living organisms that inhabit a particular environment that is thought to be exposed to mutagenic substances (Clemente et al. 2004; Machella et al. 2006; Emmanouil et al. 2007; Yin et al. 2009; Kumar et al. 2010; Capriglione et al. 2011; Mohanty et al. 2011; Parolini et al. 2011).

The aim of this study was to determine the degree of contamination of wetland by measuring certain chemical and microbiological parameters and whether they negatively affect the site (Playa Penino) and its fauna. For the latter, we used an indicator organism, which could be affected at the molecular level by different levels of contamination: *Ctenomys pearsoni* (Lessa and Langguth 1983), commonly known as tuco tuco, which was chosen

because they inhabit burrows around the wetland. They imbibe water in the form of raw sap from the roots of wetlands plants and rarely leave their burrows, so we do not have to consider the effects of air pollution. Genetic biomarkers of genetic damage were identified by using the comet assay and micronucleus test.

Methods

Playa Penino is a natural reserve, located in the Department of San Jose (34°47'S 56°24'W; south of Uruguay). It extends for 7 km along the coast. The area studied is located in the wetland of Playa Penino (route 1 South, kilometer 30) (Fig. 1). Locations where the indicator animals were captured are indicated by white circles. Animals were captured using live traps. Blood was obtained without sedation by cutting the base of a rear claw. To avoid stress, each animal was placed in an opaque tube leaving only its rear end exposed. Adult individuals were selected and sexed. Following disinfection of the cut, the animals were returned to their burrows. Blood samples were collected by dripping and placed in tubes containing heparin (da Silva et al. 2000). Individuals were collected in four months of each year (February, May, August and October), which includes the widest range in seasonal variability (i.e. wet and dry months) a total of 25 females / 25 males in 2013, 23 females / 20 males in 2015 and 22 females/21 males in 2017.

Control

Thirteen individuals (6 males and 7 female) of *Ctenomys pearsoni* from Playa Penino were also maintained in captivity in temperature-controlled conditions, in tanks containing sand and vegetables sufficient for a month, as negative controls. After a month, they were released back into the burrows that they previously inhabited, which were marked by flags. Blood was obtained from them during the last two days of their captivity and kept in liquid nitrogen until analysed.

Genetic biomarkers

The comet assay, or single cell gel electrophoresis (SCGE), is a standard method for determining Genotoxicity *in vivo* and *in vitro*, which records the genetic damage caused by clastogenic agents by measuring breaks in the DNA chains in specific cells (Ribeiro et al. 2003; Mudry and Carballo 2006). This characteristic and its sensitivity, makes it especially well-suited for Eco toxicological studies, both in terrestrial and aquatic environments (Ribeiro et al. 2003; Mudry and Carballo 2006; De Lapuente et al. 2015). Micronuclei are a portion or whole chromosomes that are not in the nucleus during anaphase. Their presence implies the loss of a part or an entire chromosome in daughter cells (Mudry and Carballo 2006; Luzhna et al. 2013). More than one micronucleus can be produced during cell division depending on the extent of the genetic damage (Luzhna et al. 2013).

Comet assay

The genotoxic evaluation was done using the comet assay. The standard procedure described by Hartmann et al. (2003) was used. Blood samples were centrifuged, washed in PBS and re-suspended in low melting agarose. Samples were incubated for 30 min at 37 °C and slides were prepared as outlined by Hartmann et al. (2003). After incubation, 100 µl of each sample was added to each slide, covered with normal agarose and subjected to lysis for 1 h in a lysis solution (NaCl 2.5 M, EDTA 100 mM, Tris 10 mM, Triton X100 1% and DMSO 10%, pH 10). The DNA was denaturalized in electrophoresis buffer for 20 min and after that subjected to electrophoresis for 30 min (20 V, 250 mA). Then the slides were neutralized in Tris 0.4 M, pH 7.5 for 15 minutes and stained with propidium iodide (PI) (10 µg/ml). The comet visualization and measurements were done using Comet Score™ software (Tritek) and 100 cells for each individual were analysed under a fluorescence microscope (Shiner XY-FL™). To assess cytotoxicity, one slide was removed from lysis after 1 hour and neutralized without undergoing electrophoresis. Air dried slides were stained with PI and 100 cells per slide were scored visually. Cells were classified as normal, if their DNA was mostly condensed (normal cells) and apoptotic cells were visualized as diffused or condensed DNA without the round normal shape. Samples with a percentage of apoptosis lower than

15% were used for this study as recommended in the protocols (Hartmann et al. 2003; Mudry and Carballo 2006).

Micronucleus test

Following the procedure described by Fenech (2007) with modifications, smears were used to determine the number of micro nucleated polychromatic erythrocytes (MN-PCE) and total percentage of micronuclei (% MN) in more than 2,000 cells. A micronucleus was 1/16 to 1/3 of the size of the main nucleus and not attached to it (Garriott et al. 2002; Norppa and Falck 2003; Zalacain et al. 2005). Frequencies were analysed using parametric and non-parametric statistics (the latter for cases, in which the conditions of normality and homoscedasticity were not met using Centurion Statgraphics XVI.I software). The observations were made under a fluorescence microscope and slides were stained with propidium iodide (PI) (10 µg/ml).

Chemical analysis

Water samples (one per season) were analysed in the field using the kit colour (Merckmillipore™) for ammonium (NH₄₊) and test strips (Hach™) for nitrates (NO₃-N) and phosphates (reactive orthophosphate PO₄) following the procedures indicated by the providers. Samples were from the surface along the shoreline of the wetland. Superficial water was collected in 500 ml sterilized plastic bottles at five locations in the wetland. Then, they were mixed in sterilized plastic container for the analyses. Samples were obtained when the indicator animals were captured.

Faecal coliform bacteria

3M™ Petrifilm™ Enterobacteriaceae plates were used following the procedures indicated by the provider and the samples described in the chemical analysis section. Water samples were obtained along with the indicator animals using the same procedure as in the previous section.

Statistical analyses

The normality and homoscedasticity of data obtained using biomarkers of genetic damage (comet assay and MN test) were tested by methods based on Kolmogorov-Smirnov for normality and Levene and Tukey for homoscedasticity (Zar 2010). To compare the data on these variables, analysis of variance (ANOVA) for parametric data and Kruskal-Wallis test for nonparametric data were used. Results of all the analyses are listed in tables along with the relationships between variables analysed using multiple correlation analysis (Statgraphic Centurion XVII software). The variables that were not normally distributed were transformed prior to correlation.

Agrochemical analyses

Samples of water (obtained as already described), soil (from the burrows of the captured animals) and the

Table 1 Values of % damaged DNA and OTM per month/year. Control values were: % damaged DNA 5.1; OTM 0.24.

Month ¹ /Year	% DNA damaged	OTM
February 2013	31.40	9.08
May 2013	17.11	2.06
August 2013	44.97	7.45
October 2013	51.91	11.99
Annual average	36.35	7.65
February 2015	34.80	8.78
May 2015	23.10	1.66
August 2015	46.00	7.15
October 2015	59.50	11.19
Annual average	40.85	7.20
February 2017	33.20	10.21
May 2017	18.90	2.34
August 2017	50.14	8.05
October 2017	59.30	12.14
Annual average	40.39	8.19

¹ February corresponds to summer, May to autumn, August to winter and October to spring. This is valid for all tables.

plants they consumed were tested for the presence of agrochemicals by LATU (Laboratorio Tecnológico del Uruguay) in every season of the year 2013. Gas exchange chromatography was used for the analyses, which is accredited by the United Kingdom Accreditation Agency, UKAS. The detection limit was 1 ng/l.

The presence or absence of the following pesticides was recorded: aldrin (chlorinated insecticide), carbaryl (insecticide, carbamate), cypermethrin (insecticide, pyrethroid), chlorothalonil (fungicide and chlorinated broad-spectrum insecticide), chlorpyrifos (organophosphate insecticide), dieldrin (chlorinated insecticide, prohibited use), α , β and λ endosulfan (organochlorine insecticide and acaricide, prohibited), malathion (organophosphate insecticide), methylchlorpyrifos (insecticide, organophosphate), methyl parathion (organophosphate, prohibited) (insecticide, pyrethroid), procymidone (chlorinated fungicide).

Results

Average values for the percentage of DNA damaged (% DNA damaged) and Olive tail moment (OTM) in the lymphocytes of *Ctenomys* did not differ between males and females (Table 1), Mann-Whitney U test ($p = 0.08$).

Data provided is for all the individuals sampled. Both variables are normally distributed according to Kolmogorov-Smirnov and homoscedastic according to Levene and Tukey tests. ANOVAs show highly significant differences ($p < 0.01$) between samples in all months in each year in both variables and the control samples, and no

Table 2 Values of PCE-MN and ‰ MN per month/year.

*Significantly different from the other months. Control values were: MN-PCEs 0.03; ‰ MN 0.04.

Month/Year	MN-PCEs	‰ MN
February 2013	2.4	2.8
May 2013	1.5	1.7
August 2013	2.0	2.6
October 2013	5.6*	5.9*
Annual average	2.9	3.3
February 2015	2.3	3.2
May 2015	1.6	1.8
August 2015	2.1	2.7
October 2015	5.7*	6.2*
Annual average	3.0	3.5
February 2017	2.2	2.6
May 2017	1.6	1.8
August 2017	2.1	2.5
October 2017	5.3*	5.5*
Annual average	2.8	3.1

significant differences between the annual values for the three years ($p < 0.05$).

In Table 2, values for polychromatic erythrocytes (PCE) with one or more micronuclei (MN-PCEs) and ‰ MN are shown. There are no significant differences between males and females according to Mann-Whitney U test ($p = 0.075$). Data provided is for all the total individuals sampled. The MN-PCEs counted were 2000 (2 smears per individual) and values were averaged. Only polychromatic erythrocytes were counted as the spleen of *Ctenomys* removes abnormal erythrocytes from its blood, therefore normochromatic erythrocytes (NCE) were not detected. According the Kruskal-Wallis tests, there are significant differences ($p < 0.05$) between the October samples for each year and the other months analysed for both variables (MN-PCEs and ‰ MN). All samples showed significant differences relative to the controls ($p < 0.05$). There were no significant differences between the annual values of the three years ($p < 0.05$).

Phosphate reached its highest values in May and February in each year and nitrates increased in May only in 2013. Values for ammonium, on the other hand, were lowest in May and high in the February and October samples, which is coincident with the increase in the ammonium values. Faecal coliform bacteria (FC) also showed an increase in the months of February and October in each year (Table 3).

FC concentration was positively and significantly correlated with OTM. The concentration of ammonia was positively correlated ($p < 0.05$) with the biomarkers of genotoxicity (% damaged DNA and OTM). The percentage FC correlates positively and significantly with the concentration of ammonia in water (0.985, data not

Table 3 Values of the chemical and microbiological parameters of the compound samples from the wetland for each month/year.

Month/Year	Phosphates (mg/l)	Nitrates (mg/l)	Ammonium (mg/l)	Faecal coliform bacteria (CFU/100 ml)
February 2013	1.0	0	1.0	1300
May 2013	1.0	0.15	0	300
August 2013	0	0	0	800
October 2013	0	0	1.0	1500
Annual average	0.5	0.004	0.5	975
February 2015	1.0	0.15	1.0	1300
May 2015	0.5	0.15	0.5	300
August 2015	0	0	0	800
October 2015	0	0	1.0	1700
Annual average	0.4	0.08	0.6	1025
February 2017	1.0	0.15	1.0	1200
May 2017	0.5	0.3	0	300
August 2017	0	0	0.5	700
October 2017	0	0	1.0	1600
Annual average	0.4	0.11	0.6	950

shown in the Table). Correlation coefficient between phosphates and % MN was moderate (Table 4).

There were no agrochemicals present in the water, soil and plants in any season of the year when they were recorded.

Discussion

Genetic damage

The biomarkers used revealed higher levels of genetic damage compared to other studies using the same genus (*Ctenomys*). The percentages of MN-PCEs and % MN reveal the existence of serious genetic damage in the indicator species. The MN-PCEs values exceed those reported by Heuser et al. (2002) in *Ctenomys minutus* exposed to elevated levels of hydrocarbons and heavy metals (MN-PCEs value = 0.76) at locations along route 30 (coast of Rio Grande do Sul, Brazil). They are also higher than the maximum values of MN-PCEs (value = 2.8) obtained by da Silva et al. (2000) in *Ctenomys torquatus* for individuals exposed to polycyclic aromatic hydrocarbons (PAHs)

Table 4 Multiple correlation test showing significant coefficients at $p < 0.05$. NS = not significant.

Correlation coefficients $p < 0.05$				
	Phosphates	Nitrates	Ammonium	Faecal coliforms
% damaged DNA	NS	NS	0.980	NS
OMT	NS	NS	0.969	0.975
PCE-MN	NS	NS	NS	NS
% MN	NS	0.531	NS	NS

in a coal mining area located at Candiota in southern Brazil. The values of MN-PCEs also are higher than the maximum value obtained by Ayla et al. (2005) for rats exposed to lead acetate (a solvent used in the production of cosmetics and furniture) (value = 2.41).

The average % of DNA damaged ranged from 36 to 41%. Kopjar et al. (2008) report values of 0.85–7.37 for % DNA damaged and 0.11–0.86 for OTM, in peripheral blood of fish from rivers that were contaminated with urban effluents (nitrates, nitrites, and ammonium). Festa et al. (2003) report maximum values for % of DNA damaged of 32.8 and Tail Moment of 38.28 in *Mus spretus* blood cells during a toxic spill at the Aznalcollar mine in Doñana National Park, which dumped acidic water, mud full of toxic metals and arsenic into the Guadiamar River.

The highest values recorded in this study indicate the presence of the genotoxic agents lead, hydrocarbons, heavy metals and arsenic in the Playa Penino wetland. It is likely that these xenobiotics are entering the blood system of the indicator species via the plants it consumes.

Effects of the chemicals and bacteria

There were no registered agrochemicals in the samples of water, soil and plants analysed. Although antibiotics were not tested for in the water, it is important to point out that Rodríguez (2010) reports values for these substances of less than 1 $\mu\text{g/l}$ in surface waters in areas with larger populations than at Playa Penino. This value is the minimum that results in adverse effects on microorganisms *in vitro* (Rodríguez 2010). Moreover, the concentration of microorganisms should be affected by the presence of antibiotics in the water, so that the high concentrations of enterobacteria recorded in this study is not consistent with the presence of toxic levels of antibiotics (Rodríguez 2010).

Even though the presence of detergents was not recorded, many of them include phosphate, which would generate a very big increase in the phosphate values of water samples (Romero 2006).

According to ordinance 253/79 governing the quality of water in Uruguay (MVOTMA 1979), the Playa Penino wetland corresponds to class 2 b, which includes “waters intended for recreation by direct contact with the human body for many residents and visitors of the area” or class 3 “waters destined to the preservation of fish in general and other members of the water flora and fauna, or also waters destined to the irrigation of crops whose product is not consumed in a natural way or in those cases that being consumed in natural way irrigation systems are applied that do not cause the product to wet”. For either class, the value for ammonia exceeded the allowable limits in all samples (allowable limit 0.02 mg/l). In contrast, nitrates are below the limits set by this ordinance (10 mg/l).

Regarding the nitrogen-related contributions, the average annual values ranged from 0.5 to 0.6 mg/l for ammonium. For nitrates, the average annual values ranged from 0.004 to 0.11 mg/l. The average annual nitrate val-

ues and their maximum values (0.11 mg NO₃/l) were below the national and international standards (Arocena et al. 2008). The maximum average ammonium values (0.6 mg NH₄/l) exceeded not only these ranges, but the international values for the preservation of aquatic life and drinking water (Arocena et al. 2008). Ammonia is not toxic, but its presence even in low concentrations may mean the presence of faecal bacteria and pathogenic microorganisms in the water. Ammonia is produced during the first inorganic stage of the bacterial breakdown of urea and proteins, and it is a good indirect chemical indicator of the faecal contamination of water (Laws 1993; Rodríguez et al. 2006).

The average values for phosphate (PO₄) ranged from 0.4 to 0.5/l, which is greater than the internationally recommended level (Delgadillo et al. 2010; Putz 2010). Phosphates are directly related to the eutrophication of rivers, and especially lakes and reservoirs, considering that only one gram of phosphate (PO₄) causes the growth of up to 100 g of algae (Delgadillo et al. 2010; Putz 2010). Directive UE 91/271/EEC indicates that the critical concentrations for incipient eutrophication are between 0.1–0.2 mg/l PO₄ in running water and between 0.005–0.01 mg/l PO₄ in still water (De Azevedo and Da Matta 2004; Márquez et al. 2007; Putz 2010).

Phosphorus, unlike nitrogen, is not obtained from the atmosphere; it comes from the excreta of living organisms and is mainly contained in fertilizers, pesticides, detergents and sewage (Márquez et al. 2007). Rainfall is higher from January to May than in other months (110 mm/month) in the San José Department (climate-data.org), which may account for the higher levels of phosphate in water being a product of runoff and filtration from rudimentary septic chambers.

Nitrogen and phosphorus are essential for all living organisms, as they are a basic part of molecules such as proteins and nucleic acids (Spiro and Stigliani 2004), and play a key role in their growth. Under normal conditions, these substances come mainly from external natural contributions and the decomposition of dead organic matter (recycling). When the environmental conditions are altered, however, the additional contributions come mostly from the discharge of urban waste and certain industrial processes, as well as from the increasing use of fertilisers and pesticides in agriculture (De Azevedo and Da Matta 2004; Spiro and Stigliani 2004; Cicerone et al. 2007).

In urban or domestic wastewater effluents, nitrogen is present primarily as organic nitrogen or ammonium and, to a lesser extent, as nitrites and nitrates (De Azevedo and Da Matta 2004; Spiro and Stigliani 2004; Cicerone et al. 2007; Delgadillo et al. 2010). The reduced nitrogen is converted into nitrite (nitrification) and subsequently into nitrate (De Azevedo and Da Matta 2004; Cicerone et al. 2007; Newman and Clements 2008; Delgadillo et al. 2010).

Ammonium particularly inhibits the symbiotic fixation of nitrogen by microorganisms associated with

plants, which are in close contact with water in the wetland environment (Newman and Clements 2008; Delgadillo et al. 2010). This process is indirectly affected by the increase in the cytoplasmic concentration of ammonium in the roots exceeding their assimilative capacity (Newman and Clements 2008; Delgadillo et al. 2010). The chemical compounds detected in the wetland at Playa Penino might be affecting the genome of the indicator species, which lives in burrows, in the immediate proximity of water, consumes the roots of the vegetation associated with the wetland, absorbing the sap and pollutants present especially in those species with a high water content (*Equisetum sp.*, *Cynodon dactylon* and *Paspalum nicorae*).

In an aquatic environment, the presence of nitrogen in the form of ammonia is regulated by a chemical equilibrium that determines the coexistence of two forms: ammonia (NH₃) and ammonium ions (NH₄⁺). The relative proportion of each is determined by pH, temperature and ionic strength (Delgadillo et al. 2010). The concentration of ammonia increases with increase in pH and temperature and decreases with increase in ionic strength; the ammonium ion predominates at pH values lower than 7 (Pepe and Lombardi 2003). The toxicity of ammonia for organisms is related to the non-ionized form (ammonia), mainly because it is highly soluble in lipids, which facilitates its passage through biological membranes, causing damage to respiratory surfaces. In fish, for example, the distal parts of the fins lose their original colour and turn white or transparent (Pepe and Lombardi 2003; Spiro and Stigliani 2004). Ammonium salts induce skin secretions and destroy gills (Pepe and Lombardi 2003; Spiro and Stigliani 2004). Relevant to this, the mass death of fish recorded in 2013 and 2015 in Playa Penino could have resulted from the process described above.

In the digestive system of mammals, nitrogen compounds (ammonium and nitrates) are transformed into nitrites by microbes. An excess of nitrite induces the formation of methaemoglobin (Khademikia et al. 2013), which blocks haemoglobin transporting oxygen. In addition, these nitrogen compounds (nitrates and nitrites) in water can be transformed into nitrosamines, whose teratogenic and carcinogenic potential is recognized (Camargo and Alonso 2006). As mentioned before, tucu tucu obtains water from the roots of the plants (raw sap, mostly from grass in this area), so they may be accumulating these compounds, as they are in the water and their metabolic transformation could occur later, in other parts of the plants. However, this needs to be tested under controlled conditions in the laboratory.

The ability of faecal bacteria to reproduce outside the intestine of mammals is favoured by suitable environmental conditions (Eaton and Franson 2005). Their presence is interpreted as an indication that pathogenic organisms may be present and their absence that the water or food is free of disease-producing organisms (Eaton and Franson 2005). Uruguayan ordinance 253/79

(MVOTMA 1979) establishes a maximum limit of 1000 cfu/100 ml for faecal coliform bacteria (FC) for class 3 water and a maximum limit of 500–1000 cfu/100 ml for class 2 water. The microbiological contamination by FC recorded in this study exceeded the guide levels for these classes of water, as well as for drinking water (according to national and international regulations) and are a risk even for the recreational use of water (Delgadillo et al. 2010). The values of FC recorded in this study were similar (May) or higher (October and February) to those recorded in sections of the Traiguén IX Region river in Chile (540 cfu/ml; 1100 cfu/ml; 920 cfu/ml) (Rivera et al. 2009). Pérez et al. (2004) report FC values for the water in Lake Izabal and Río Dulce, in Guatemala, which are lower than those recorded in this study (Lake Izabal 454 cfu/100 ml and Río Dulce 62 cfu/100 ml). These authors mention that these levels of contamination pose a health hazard for the population inhabiting the margins of these environments. Something similar occurs in the Playa Penino wetland, which is also used for irrigation, fishing and recreational uses by the surrounding population.

Pollution by both bacteria and chemicals pose a high risk to human health, affect the crops and reduce the availability and quality of fish, which are a source of food for communities where the incidence of poverty is high.

The multiple correlation analysis (MCA) revealed significant positive correlations between the concentration of ammonia and the presence of enterobacteria, and the genetic damage recorded in the blood cells of *Ctenomys*. Although positive these relationships do not prove there is causal connection between the genetic and environmental variables. It might indicate that the discharge of sewage has a negative effect on the wetland. This requires additional *in vivo* and *in vitro* studies.

Squatter settlements around Playa Penino along with the lack of an urban sanitation system has resulted in an increase in nutrients to above the maximum allowed by current national and international regulations, which has disturbed natural nutrient cycling.

Conservation implications

The results of this study might indicate the decrease in biodiversity at Playa Penino is due to an acute and chronic exposure of the genetic material of the organisms there to pollutants that in the long-term might reduce their fitness and survival.

An indicator species was used to detect a potentially dangerous situation associated with the presence and action of pollutants, which to some extent can be extrapolated to humans (Alonso and Tabor 2004; Den Besten and Munawar 2005; Bischoff et al. 2010; Fernández-Tajes et al. 2011). The extent of the genetic damage to individual cells measured using OTM and percentage of DNA damaged revealed acute and recent exposure to xenobiotics, which may be associated with the discharge of domestic wastewater. The MN test also revealed the chronic

exposure of the animals showing genetic damage to clastogenic and/or aneugenic events (Mudry and Carballo 2006).

It is essential to maintain the ecosystem integrity of Playa Penino and ensure the protection of the health of humans by a strict surveillance and control of septic tanks and design of an adequate sanitation system. The regularization of the squatter settlements through a housing plan that contemplates the socio-economic possibilities of the inhabitants, would reduce the pollutant load from sewage.

The restoration of the physical-chemical balances between the soil, water and sediment will also be beneficial, as is described in the GeoUruguay report (Martino et al. 2008).

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