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2 **Exploring the use of tertiary reclaimed water in dairy cattle production**

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26 **Abstract**

27 The objective of this study was to explore through both *in vitro* and *in vivo* experiments
28 the use of reclaimed urban wastewater in dairy cattle production systems with the aim
29 of improving water efficiency and sustainability. Firstly, the use of different tertiary
30 treatments (ultrafiltration (UF), ultraviolet disinfection (UV), chlorination process, and
31 their combination) to improve the quality of an urban secondary effluent was studied in
32 intestinal primary cell cultures evaluating the expression of genes related to apoptosis,
33 cell damage, and inflammation. The results revealed that secondary treated wastewater
34 and waters that were treated with a chlorination process (even tap water) caused an
35 increase in apoptosis, intestinal primary cell damage, and inflammation. The *in vivo*
36 experiment evaluated the short-term effects on health and performance of using UF- and
37 UV-treated secondary effluent compared with the use of tap water for drinking and
38 preparing milk replacer in young calves from 5 to 47 days of age. Calves previously fed
39 with UF+UV treated secondary effluent clearly preferred tap water when they were
40 exposed to a double water choice at the end of the study. This reduction of the
41 palatability and acceptability was probably due to a greater level of water salinity of the
42 treated reclaimed water (570 vs $1,437 \pm 76.5$ $\mu\text{S}/\text{cm}$ of conductivity for tap water and
43 UF-UV treated secondary effluent, respectively), which potentially entailed a reduction
44 of calf concentrate intake (466 vs 351 ± 32.2 g/d for calves fed with tap water and UF-
45 UV treated water, respectively). The use of reclaimed water did not pose an acute risk to
46 animal health. It is concluded that improvements on the tertiary treatment to reduce
47 water salinity should be considered when using reclaimed water for drinking purposes
48 in livestock production systems. This study is a first approach to a more sustainable and
49 efficient use of water in animal husbandry for countries with water scarcity. However,

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50 more studies are required before its implementation to further study long-term effects
51 and the presence of new-contaminants not defined in the current legislation.

52 Keywords: livestock drinking water; reclaimed water; ultrafiltration; water reuse

53 **Abbreviation list**

54 ACTB: β -actin

55 BNIP3: Adenovirus E1B 19 kDa protein-interacting protein 3

56 CASP3: Caspase 3

57 DM: dry matter

58 HSPA1A: Heat shock 70 kDa protein 1A

59 HSPB1: Heat shock protein family B member 1

60 IL-1 β : Interleukin 1 beta

61 IL-10: Interleukin 10

62 MR: Milk replacer

63 TDS: total dissolved solids

64 TIC: total inorganic carbon

65 TNF- α : Tumor necrosis factor alpha

66 TOC: total organic carbon

67 TSS: total suspended solids

68 UF: ultrafiltration

69 UV: ultraviolet

70 WW: wastewater

71 **1. Introduction**

72 The agricultural sector accounts for around 70% of water use worldwide, and it remains
73 one of the major sources of water pollution with fertilizer run-off, pesticide, and
74 livestock effluents (FAO, 2015). Furthermore, the prediction of rising world human

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75 population will increase water demand in this sector (Gulbenkian Think Tank, 2014),
76 and future policies must look for more sustainable food production systems to avoid
77 serious food and water crisis in the upcoming years. Water reduction and sustainable
78 systems in crop production, such as more efficient irrigation systems (Kusakabe et al.,
79 2016; Singh et al., 2016) or crops more adapted to drought areas (Daryanto et al., 2016;
80 Vurukonda et al., 2016) merit current attention. Within the primary sector, it is
81 estimated that livestock production (including irrigation of grains, forages, and pastures
82 plus water usage for animal husbandry) uses 29% of the total agriculture water demand
83 (Mekonnen and Hoekstra, 2012). However, much less attention has been paid to the
84 livestock production system regarding efficiency of water utilisation (Ran et al., 2016).
85 To face the increasing water scarcity and water pollution, some initiatives involving
86 water reuse, mainly from urban sources, have been implemented in some countries
87 (Kihila et al., 2014) as an economically-feasible method of increasing existing water
88 supply, especially when compared with expensive alternatives such as desalination or
89 development of new water sources involving dams and reservoirs (Shannon et al.,
90 2008). The most common water reuse application in the agricultural sector involves
91 irrigation of food crops, pastures, and industrial non-food crops (Maestre-Valero et al.,
92 2016; Jiang et al., 2016), and to a lesser extent aquaculture (Feldlite et al., 2008), and
93 silviculture (House et al., 1999). In the livestock sector either groundwater or surface
94 water is used to supply water to animals depending on their locations. However, only
95 the Environment Protection Authority of Victoria in Australia has regulated the
96 implementation of reclaimed water usage in animal husbandry. The two main water
97 uses in intensive dairy cattle production systems are for drinking (82%) and cleaning
98 farm facilities (18%) (Drastig et al., 2010), and these needs are fairly constant
99 throughout the year. In contrast, for crops irrigation water demand is seasonally, which

100 makes the implementation of more reclaimed water systems difficult. In general, quality
101 requirements for livestock drinking water and their impact on livestock health and
102 performance are poorly investigated. There are no specific legal requirements
103 concerning quality of drinking water for dairy cattle (in most of the legislation is
104 mentioned suitable and healthy water), with most documents being mere guidelines
105 from governmental and academic institutions (South Africa Department of Water
106 Affaires and Forestry, 1996; Olkowski, 2009; Schlink, 2010; Department of
107 Environment, Food and Rural Affairs of United Kingdom, 2012; Department of
108 Primary Industries and Regional Development of Australia, 2017).

109 The motivations of the present study were the need for new strategies to improve water
110 efficiency and sustainability in livestock production coupled with the availability of
111 economical reasonable water treatment technologies that apparently result in sufficient
112 water quality that could have no negative effects on health and productivity of dairy
113 cattle. Therefore, the two main objectives of the study were: 1) to evaluate, in an *in vitro*
114 system, the most suitable tertiary wastewater (WW) treatment process to obtain
115 reclaimed water of sufficient quality for dairy cattle drinking purposes, and 2) to
116 evaluate in an *in vivo* study the short-term effects on health and performance of offering
117 reclaimed water to dairy calves.

118 **2. Materials and methods**

119 *2.1. In vitro study*

120 This study was performed in the facilities of IRTA in Torre Marimon (Caldes de
121 Montbui, Spain), and WW was obtained from the urban WW treatment plant in Caldes
122 de Montbui (Barcelona, Spain.), which received mostly municipal discharges. This
123 plant was situated at approximately 1 km from IRTA facilities. Water treatment in this

124 urban WW treatment plant includes a physicochemical primary treatment followed by
125 biological and settling secondary treatments.

126 *2.1.1. Tertiary wastewater treatment selection*

127 The WW (composition depicted in Table 1) intended for tertiary reclamation was the
128 effluent from the secondary treatment of the WW treatment plant. The water tertiary
129 treatment to obtain water of suitable quality for calves drinking should pursue a
130 reduction in the microbiological load and a reduction of water turbidity, following the
131 Australian guidelines for the use of reclaimed water (Class B, pH 6-9, < 100 cfu of
132 *Escherichia coli* /100 mL, < 20 mg/L Biological Oxygen Demand, < 30 mg/L
133 suspended solids, and a reduction of helminth eggs). In the present study, several
134 tertiary treatments were initially tested in intestinal primary cell cultures to select the
135 most suitable for conducting a subsequent *in vivo* study with dairy calves. The tertiary
136 treatments consisted of a combination of several technologies: ultrafiltration (UF) with a
137 30 nm pore membrane (66.03 I8 Berhof Membrane Technology GmbH, Eningen,
138 Spain); ultraviolet (UV) disinfection (STERILUX MINI-1000 CEASA, Castellví de
139 Rosanes, Barcelona, Spain) with a targeted UV dose of 80 mJ/cm² (UV); and
140 chlorination with addition of sodium hypochlorite to achieve 1 mg/L of free chlorine
141 (ClO⁻).

142 *2.1.2. Intestinal primary cell cultures*

143 Jejunum tissue was obtained at a slaughterhouse from an 11-mo old bull and
144 immediately transported in chilled phosphate-buffered saline with 100 µg/mL
145 streptomycin, 100 U/mL penicillin, and 2.5 µg/mL amphotericin B to the laboratory. In
146 the laboratory, tissue was cut into small pieces and washed in phosphate-buffered saline
147 with 0.1 mM ethylenediaminetetraacetic acid and 0.1 mM dithiothreitol for 10 min at
148 37°C in 5% CO₂ at 150 rpm. Then, supernatant was removed and Roswell Park

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149 Memorial Institute (RPMI) 1640 media with 0.25% collagenase was added and
150 incubated for 15 min at 37°C in 5% CO₂ at 150 rpm. The supernatant containing
151 isolated epithelial cells was added to a same volume of RPMI 1640 media with 0.02
152 mg/mL DNase. This step was repeated 3 times. Then, supernatants (containing the cells)
153 were centrifuged at 300 g for 5 min and the cell pellets resuspended in Dulbecco's
154 Modified Eagle Medium/Nutrient Mixture F-12 (DMEM/F12) media with 8 µg/mL
155 bovine insulin, 10 µg/mL gentamycin, 50 µg/mL hydrocortisone, 10% fetal bovine
156 serum, 100 µg/mL streptomycin, 100 U/mL penicillin, and 2.5 µg/mL amphotericin.
157 Cells were quantified by microscopy and incubated at 80,000 cells/cm² in 175 cm²
158 flasks for 24 h at 37°C in 5% CO₂ at 150 rpm. Epithelial cell phenotype was confirmed
159 by immunofluorescence staining against anti-cytokeratin antibodies (Sigma-Aldrich,
160 Saint Louis, US).

161 *2.1.3. Primary cell culture*

162 Jejunum cells were cultured in 24-well plates at 37°C under a 5% CO₂ atmosphere, at a
163 cell density of 50,000 cells/well during 16 h. Then, cells were treated during 2 h with
164 eight different types of water: 1) cell media culture (MC) as a negative control to
165 evaluate the effects of incubating the intestinal cells in the plates, 2) tap water (TW) to
166 evaluate a potable water source, 3) water from a drinker from a dairy farm filled with
167 chlorinated groundwater (ThW) to evaluate the water that animals are consuming in the
168 field conditions, 4) secondary effluent of the WW treatment plant (SW) to evaluate the
169 improvements of the tertiary treatments proposed in the study, 5) secondary effluent
170 with an UF treatment (UF), 6) secondary effluent with an UV disinfection (UV), 7)
171 secondary effluent with UF and UV treatments (UF+UV), and 8) secondary effluent
172 with UF and chlorination (UF+ClO⁻). After the incubation, cells were washed and lysed
173 with TriZol (Invitrogen, Paisley, UK) to extract RNA and quantify, by qPCR,

174 expression of apoptotic (*BNIP3* and *CASP3*), cell damage (*HSPA1A* and *HSPB1*), and
175 inflammation (*TNF α* , *IL-1 β* , and *IL-10*) genes.

176 *2.1.4. Sampling and analyses*

177 Five-litres water samples were obtained in a plastic container in the wastewater
178 treatment plant, and within the same day they were sent refrigerated (5-7 °C) to the
179 laboratory. Samples were kept refrigerated in the laboratory until the different
180 treatments to produce the studied waters were applied. Samples of the different
181 treatments were obtained to analyse them for pH, conductivity, turbidity, chemical
182 oxygen demand (COD), ammonium (NH_4^+), chloride (Cl^-), phosphate (PO_4^{3-}), nitrate
183 (NO_3^-), sulphate (SO_4^{2-}), aerobic bacteria counts, and *E. coli* counts following analytical
184 standard methods for water quality. COD was determined by the method 5220 defined
185 in the Standard Methods for the Examination of Water and Wastewater (1998) and
186 anions and cations were analysed by ionic chromatography (Dionex ICS-2100).
187 Microbiological characterisation was performed following the standard methods UNE-
188 EN ISO 6222 and UNE-EN ISO 9308 - 1 for aerobic counts and *E. coli*, respectively.

189 Total RNA from the cells was extracted using Trizol (Invitrogen, Paisley, UK). One
190 microgram of RNA was retrotranscribed to cDNA using IScript cDNA synthesis kit
191 (Bio-Rad, Hercules, CA) following the manufacturer's instructions. Real-time PCR was
192 performed using specific primers described in Table 2. A total reaction volume of 20 μL
193 containing 100 ng of cDNA, 10 μL SYBR Green (Bio-Rad Laboratories) was used at
194 the optimized primer concentration for each gene (Table 2). The relative expression of
195 selected genes was calculated using the delta cycle threshold (ΔCt) method with β -actin
196 (*ACTB*) as the reference gene, and a randomly-chosen sample of the media culture
197 treatment as a calibrator following Pfaf (2014).

198 *2.1.5. Statistical analysis*

199 An analysis of variance with the type of water as the main effect was performed for all
200 data. Outcome variables that did not follow a normal distribution were log-transformed.
201 Least square means and the standard error of the mean (SEM) presented herein
202 correspond to non-transformed data, and the *P*-values correspond to the results with the
203 log-transformed model. Significance was declared at $P < 0.05$ and tendencies at ≤ 0.10 ,
204 using the Fisher's protected LSD test to assess differences among treatments.

205 *2.2. In vivo study*

206 *2.2.1. Tertiary reclaimed water production*

207 The secondary effluent from the wastewater treatment plant was transported by a tanker
208 truck to a 27 m³ closed tank in the facilities of IRTA. The onsite system for the tertiary
209 treatment consisted of a UF and a UV disinfection process. The UF consisted of a
210 HyperFlux tubular module (model 66.03 I8, Berghof Membrane Technology GmbH,
211 Eningen, Germany) and was operated in crossflow mode to have around 300L/d of
212 permeate. All piping was made of plastic black tubing to avoid algae growth. The
213 system worked intermittently and started/stopped automatically following a set program
214 (running from 3:00 to 6:00 a.m., from 7:00 to 10:00 a.m., and from 10:00 a.m. to 1:00
215 p.m.) to avoid overheating of the circulation pump. The UF system consisted of some
216 programmable logic controllers with programs and security sensors (i.e., water level).
217 To check the working pressure and pressure drop during filtration manometers were
218 installed at the intake and outtake of the UF module. Permeate was diverted to a black
219 storage tank of 1,000 L, which fed the UV module. The UV operated daily at a flow rate
220 of 300 L/h (UV dose of 80 mJ/cm²) to obtain the amount of reclaimed water required
221 for the preparation of milk replacer (MR) and water drinking for 10 calves during the
222 study.

223 *2.2.2. Animals and treatments*

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224 Eighteen Holstein dairy calves of 5 ± 3.2 d of age and 40 ± 6.3 kg of body weight were
225 gathered from several farms, and raised at the facilities of IRTA according to the
226 recommendations of the animal Care Committee of IRTA. Calves were housed
227 individually and bedded with sawdust. Two different types of MR feeding programs
228 were tested in order to achieve different amounts of reclaimed water consumption by
229 the animals (as calves would consume much larger quantities of MR than water alone).
230 The experiment followed a 2x2 factorial design, two different MR feeding programs (4
231 vs 8 L/d of MR at 12.5% dry matter (DM) throughout the study), and two different
232 water sources (TW vs UF+UV) that calves consumed through both MR feeding and
233 drinking water. Concentrate and barley straw were offered *ad libitum* from the
234 beginning of the study until calves reached 47 d of age (study end).

235 2.2.3. Sampling and analysis

236 Calves were weighed at the beginning of the study and once weekly thereafter. Milk
237 replacer, concentrate, straw and water intakes were measured daily, and veterinary
238 treatments were recorded. Calf faecal consistency was evaluated daily using a 3-point
239 scale (1: normal, 2: loose, 3: watery). Blood samples were obtained at the beginning and
240 at 35 d of study to determine glucose, insulin, urea, creatinine, hepatic enzymes (AST
241 and GGT), non-esterified fatty acids (NEFA), triglycerides (TG), and thyroid hormone
242 (T_3) serum concentrations, and conduct a full haematological profile. Faecal samples
243 were also obtained at the beginning and at 35 d of study to assess the presence of
244 helminthic eggs, *Cryptosporidium* cysts, and coccidia oocysts.

245 The last day of study, a preference test was performed to evaluate the capacity of
246 animals to distinguish between TW and UF+UV waters and to determine whether
247 calves had a preference for any of them. During that day, all animals were offered the
248 two types of water (TW and UF+UV) in two separate buckets to drink. At 9:00 a.m. all

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249 calves were offered 5,500 mL of each type of water, then at 1:30 p.m. water
250 consumption up to that point was recorded and additional water was offered if the
251 buckets contained less than 4,000 mL (consumption >1,500 mL). At 4:30 p.m. water
252 consumption was determined again (by weighing the buckets) and the water leftover
253 was completely replaced with new water until 9:00 a.m. of the following day, when all
254 buckets were weighed again to assess water consumption.

255 During the study, water samples of secondary water and the effluents after each
256 treatment step (UF effluent, storage tank before UV, UV effluent, and water from both
257 TW and UF+UV calf buckets) were collected fortnightly to determine water pH,
258 conductivity, turbidity, total organic carbon (TOC), total suspended solids (TSS), and
259 counts of aerobic bacteria, *Clostridium perfringens*, total coliforms, and *E. coli*. In
260 addition, total inorganic carbon (TIC), total dissolved solids (TDS), and concentration
261 of anions, cations, total Mn, and Fe, and contents of several toxic organic and inorganic
262 trace constituents including pesticides, halogenated toxic compounds, polycyclic
263 aromatic hydrocarbons, nonylphenols, and heavy metals were determined in reclaimed
264 water (UF+UV) and TW at least once throughout the duration of the study, following
265 standard analytical methods. Water samples were collected in different containers
266 depending on the parameter to be analysed: for the determination of organic
267 contaminants samples were kept in amber glass bottles, for the microbiological
268 parameters in sterile plastic containers and for the remaining parameters in plastic.
269 Samples were sent refrigerated to the laboratories and kept refrigerated until analysis.

270 Briefly, turbidity, TSS, and TDS were determined following methods 2130D, 2540G,
271 and 2540C from the Standard Methods for the Examination of Water and Wastewater
272 (1998), respectively. Water was cultivated at 37°C during 24 h in plate count agar
273 media, and in chromogenic media Compact Dry EC (Hardy Diagnostics, Santa Maria,

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274 CA) to determine counts of aerobic, coliforms, and *E. coli*, respectively. TOC and TIC
275 were analysed using a total carbon analyser (Analytik-Jena 3100 N/C). Metal
276 concentrations in water were determined by inductively coupled plasma mass
277 spectrometry (7500 CX, Agilent), and analyses of organic constituents were determined
278 by high-resolution gas chromatography coupled to mass spectrometry.

279 Reclaimed water samples were analysed three times during the study for potential
280 microbiological risks. Reference indicators for microbial hazards determined in the
281 present study were: bovine polyomaviruses (as indicator of bovine faecal
282 contamination), human adenoviruses (a human fecal viral indicator), somatic
283 coliphages, *Clostridium perfringens*, *Cryptosporidium spp.*, and helminth and *Taenia*
284 *spp.* Eggs. For virus detection, 10 L of reclaimed water samples were concentrated by
285 skimmed milk flocculation (Calgua et al., 2013) while wastewater samples were
286 concentrated by ultracentrifugation (Pina et al., 1998) and viral nucleic acids were
287 extracted using the QIAmp Viral RNA kit (QIAGEN, Inc.) to further quantify human
288 adenoviruses and bovine polyomaviruses by qPCR (Hernroth et al., 2002; Hundesa et
289 al., 2010). For *Cryptosporidium*, one litre of water was filtered using cellulose acetate
290 filters of 0.2 µm pore diameter (Whatman, GE Healthcare, Germany), and DNA was
291 extracted using the DNeasy blood and tissue kit (QIAGEN, Hilden, Germany) to further
292 quantify by qPCR *Cryptosporidium* following Guy et al. (2003). To detect *Clostridium*
293 *perfringens*, water samples were cultured in the selective media tryptone sulphite
294 neomycin agar during 24 h at 37°C in an anaerobiosis jar with Anaerocult A (Merck) to
295 consume the oxygen. Somatic coliphages were determined by incubating 100 µL of a
296 dilution of the tested water in BBL™ media and *E.coli* BL21 on Luria-Bertani media,
297 and monitoring plaque formation after 8-12 h of incubation at 37°C. Lastly, 10 L of

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298 water were used to detect by optical microscopy the presence of eggs from intestinal
299 nematodes and *Taenia spp.*

300 2.2.4. Statistical analysis

301 Data pertaining to growth performance and feed intake were analyzed with a mixed-
302 effects model with repeated measures, including the fixed effects of milk-feeding
303 program, type of water, week of study and their 2- and 3-way interactions, plus the
304 random effect of calf. Initial body weight, initial age, and farm of origin were used as
305 covariates in the model, and week entered the model as a repeated measure using an
306 autoregressive covariance matrix.

307 The incidence of scours was analyzed with a mixed-effects logistic regression being the
308 proportions of observation of score 2 the independent variable. The model considered
309 milk-feeding program, type of water, and their interaction as fixed effects.

310 Hematological and blood biochemical profiles were analyzed with an analysis of
311 variance including the effects of milk-feeding program, type of water, and their 2-way
312 interactions plus initial age and initial values (day 0 of study) as covariates. Parameters
313 that did not follow a normal distribution were log-transformed. Least square means and
314 SEM presented herein correspond to non-transformed data, and *P*-values correspond to
315 the results with the log-transformed model. Significance was declared at $P < 0.05$ and
316 tendencies at $P \leq 0.10$.

317 Water preferences were determined as a preference ratio for TW as follows:

$$318 \text{ Preference ratio} = (\text{TW intake}) / (\text{TW intake} + (\text{UF+UV}) \text{ intake})$$

319 Values for preference ratio greater than 0.5 indicate a preference for TW, values equal
320 to 0.5 denote no preference for any type of water, and preference ratio values lower than
321 0.5 indicate a preference for UF+UV water.

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322 Preference ratios for all calves (independently of their previous water experience) were
323 analyzed for a difference from 0.5 (lack of preference) using a t-test. Then, to check the
324 effect of the previous exposure to different water sources, preference ratio for TW
325 throughout the 24-h were also analyzed using a mixed-effects model considering type of
326 milk feeding program, type of water, and their 2-way interaction as fixed effects.

327 **3. Results and discussion**

328 *3.1. In vitro study*

329 The present *in vitro* experiment explored the effects that different tertiary water
330 treatments applied on a secondary effluent from a municipal wastewater treatment plant
331 had when they came into contact with bovine intestinal cells. Chemical and
332 microbiological quality of the different water types and treatments are shown in Table
333 1. Intestinal primary cells cultured with TW, ThW, SW, UV, and UF+ClO⁻ had an
334 increased ($P < 0.05$) expression of *BNIP3*, *HSPA1A*, *TNF- α* , and *IL-10* genes
335 compared with those cultured with MC, UF, and UF+UV (Figure 1), denoting an
336 increase of cellular apoptosis, cell damage, and inflammation. Although the
337 experimental design does not allow determining which chemical or microbial
338 parameters were responsible of the differences in gene expression, some hypothesis can
339 be made to explain the results herein. Interestingly, TW, ThW, and UF+ClO⁻ had in
340 common a chlorination process and this may be, in part, the cause of the negative
341 impacts on intestinal primary cells. It has been previously described that chlorine
342 disinfection by-products can cause cellular oxidative stress, and they may have
343 carcinogenic and mutagenic properties (Yuan, et al., 2006; Richardson, et al., 2007). On
344 the other hand, waters that achieved similar results than those obtained with MC
345 (considered the optimum media for intestinal cells) and had the lowest impact on
346 intestinal primary cells, were the treatments including a UF process. The UF treatment

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347 was intended to eliminate suspended particles and colloids and partly reduce the
348 microbiological load. It was not possible to associate the low gene expression elicited
349 by the UF treatment with an elimination of the microbial load because UV disinfection
350 alone did not generate similar results to MC. Thus, perhaps the good performance of UF
351 could be associated to the elimination of toxic constituents attached to suspended
352 particles (e.g., polycyclic aromatic hydrocarbons, heavy metals), which should have
353 been removed through the UF process (Smol et al., 2012). To our knowledge, this is the
354 first study that clearly indicates that the least harmful water for intestinal cells was the
355 reclaimed water treated by UF and UV. Therefore, UF treatment followed by UV
356 disinfection was the tertiary treatment selected to offer to calves in the *in vivo* study.

24 357 3.2. *In vivo* study

26 358 The *in vivo* study evaluated the short-term effects on performance and health of feeding
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28 359 dairy calves with a tertiary treated effluent from a wastewater treatment plant or tap
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30 360 water. The tertiary treatment consisted of an UF needed to reduce part of the microbial
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32 361 load such as parasite eggs and spores that are not removed with an UV disinfection
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34 362 needed to eliminate other microbial hazards such as bacteria and viruses. The
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36 363 combination of both techniques allowed having a multi-barrier treatment. Reclaimed
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38 364 water quality achieved by the proposed UF+UV reuse scheme fulfilled the water quality
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40 365 objectives proposed by the Australian guidelines (Environment Protection Authorities
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42 366 of Victoria, 2003) for the use of reclaimed water for livestock drinking in ruminants.
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44 367 Physicochemical characterization of reclaimed water in comparison to TW is shown in
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46 368 Table 3. Levels of toxic inorganic and organic constituents in reclaimed water were low
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48 369 and fulfilled water guidelines for both human and livestock drinking water (Schlink,
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51 370 2010; Olkowski, 2009; Spanish RD 140/2003). However, the concentration of certain
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371 ions, such as K^+ , Na^+ , PO_4^{3-} , and Cl^- , slightly exceeded the upper levels indicated in
372 Schlink et al. (2010).

373 There was no interaction between MR feeding program and type of water on
374 performance and feed intake parameters (Table 4). Calves fed 8L of MR had a greater
375 growth performance ($P < 0.05$), reduced concentrate intake ($P < 0.001$), and improved
376 feed efficiency ($P < 0.001$) in comparison with those fed 4L, independently of the type
377 of water offered. This effect was expected because there is a negative relationship
378 between the amount of MR offered to calves and concentrate intake (Terré et al., 2009),
379 and as MR is more digestible than concentrate, feed efficiency of calves improves when
380 more MR is offered. Calves that were offered UF+UV consumed less concentrate than
381 TW-fed calves ($P < 0.05$). There exists a close relationship between water and
382 concentrate intake (Kertz et al., 1984), and low availability of water is usually related to
383 a decrease in calf starter concentrate intake. However, water intake was similar in both
384 treatments. Constituents associated with salinity may affect water acceptability and
385 palatability and livestock performance. Generally, increasing salt content in water for
386 dairy cows decreases water and feed intake, and milk yield (Challis et al., 1987;
387 Solomon et al., 1995).

388 Regarding the microbiological quality, obtained UF+UV water did not present any
389 potential microbial risk (Table 3), and all reference microbiological indicators were
390 below threshold limits. Sewage samples obtained from the wastewater treatment plant
391 used for wastewater collection were evaluated for the potential presence of bovine fecal
392 contamination by analysing bovine adenoviruses. Since, bovine adenoviruses were not
393 detected; the probability of bovine fecal contamination of the wastewater used to
394 produce reclaimed water for calves drinking purposes was low. These results were in
395 concordance with the lack of bovine exploitations or slaughterhouses in the area

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396 surrounding the wastewater treatment plant evaluated. For this reason, human
397 adenoviruses that are present in all urban sewage samples, being excreted persistently
398 by human population, were used as indicators of the efficiency of the UF+UV treatment
399 applied. Since they are extremely stable to UV inactivation, they represent ideal
400 indicators of UV viral inactivation. It is well known that the absence of bacteria
401 indicators do not correlate with the absence of viruses so it is relevant to test reclaimed
402 water used for drinking purposes for the presence of viruses. Since viruses are host-
403 specific microorganisms, bovine adenoviruses may be used as indicators of bovine fecal
404 contamination while other animal viruses have been described to indicate animal fecal
405 contamination from other sources (Bofill-Mas et al., 2013).

406 Total aerobic counts increased 3-log cfu/mL in the storage UF tank after the UF
407 treatment, but after the UV treatment a 5-log cfu/mL reduction was observed in the
408 reclaimed water ($P < 0.05$). Loss of water quality occurred in some steps of the
409 treatment scheme because the treatment did not work continuously and some aerobic
410 bacteria re-growth occurred during storage time in the tanks. However, the UV
411 treatment was able to reduce bacterial counts again before water was used to prepare
412 MR and also offered to calves as drinking water (Figure 2). The regrowth of bacteria in
413 reclaimed water after storage has been described elsewhere (Jjemba et al., 2010; Li et
414 al., 2013). Bacteria regrowth during the storage step in the present study could be
415 envisaged because UF+UV water contained organic carbon and phosphorous, which are
416 the two main limiting nutrients needed for microbial growth (Table 3).

417 Faecal score was measured in calves fed with UF+UV and TW as indicator of faeces
418 consistency and assess incidence of diarrhoea. There were no differences in the
419 probability of faecal score of 2 in any of the treatments, and the number of veterinary

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420 treatments against diarrhoea and respiratory problems was similar between UF+UV and
421 TW calves.
422 Haematological and biochemical parameters were measured in all animals to detect
423 possible variations in general health and metabolic status. Haematological blood
424 profiles were similar between animals fed TW and UF+UV either at 4 or 8 L/d, with the
425 exception of blood eosinophil and platelet counts. Blood eosinophil counts tended ($P =$
426 0.08) to be greater in calves reared with UF+UV than those offered TW (0.61 vs $0.24 \pm$
427 0.163 , $10^3/\mu\text{L}$, respectively), and platelet counts tended ($P = 0.05$) to be lower in
428 UF+UV than in TW calves (605 vs 766 ± 49.8 , $10^3/\mu\text{L}$, respectively). Most of the
429 haematological parameters were in the range reported elsewhere (Knowles et al., 2000;
430 Brun-Hansen et al. 2006). However, blood eosinophil counts in the present study were
431 within physiological ranges of adult cattle (Roland et al., 2014), but in greater
432 concentration than in those observed in calves (Knowles et al., 2000; Brun-Hansen et al.
433 2006). Conditions commonly associated with eosinophilia included hypersensitivity
434 reactions and parasitic infections. Additional causes are neoplasia, infections, and drug
435 reactions (Roland et al., 2014). In the present study, faeces of calves were checked for
436 coccidia oocysts, nematodes eggs, and the presence of *Cryptosporidium* cysts. Only
437 *Cryptosporidium* cysts were found at the beginning of the study in both treatments ($P =$
438 0.34 ; 44.4 vs 55.6 % of positive calves were distributed in the UF+UV and TW,
439 respectively), but no helminth eggs or cysts were detected at 35 d of study in any of the
440 treatments. Therefore, either parasitic infections other than those caused by coccidia,
441 nematodes or cryptosporidia occurred in UF+UV calves to explain the deviations on
442 blood eosinophil and platelets counts or, the presence of toxic constituents in the
443 secondary effluent, which may not have been eliminated by the UF treatment, could be
444 the 2 hypothetical explanations for the observed haematological variations. In any of the

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445 cases, the problem should be considered as mild, since a greater infection or problem
446 would have changed more parameters of the haematological profile.
447 The biochemical blood profile presented some differences among treatments: serum
448 GGT concentrations were lower ($P < 0.05$) in 8-L than in 4-L fed calves, serum NEFA
449 concentration lower ($P < 0.05$) in UF+UV than in TW fed calves, serum TG
450 concentrations were lower ($P < 0.05$) in 4L-TW and 8L-UF+UV than in 8L-TW and
451 4L-UF+UV, and serum thyroid hormone concentrations were greater ($P < 0.01$) in 8-L
452 than in 4-L calves. Blood glucose and insulin are related to carbohydrate metabolism,
453 and they were within the range of calves at this age (Knowles et al., 2000). Similarly,
454 serum urea and creatinine are indicative of protein catabolism, and kidney damage,
455 respectively, and their values were similar in all treatments. Hepatic enzyme AST is an
456 indicator of soft tissue damage and no differences were found between groups of calves.
457 Although differences appeared in the hepatic enzyme GGT, normal values of GGT in
458 calves are around 20 U/L (Klinkon and Jezek, 2014), and values under this level are not
459 considered a health problem. Similarly, an increase in serum NEFA concentrations is an
460 indicator of body fat reserve mobilization; however, serum NEFA concentrations in the
461 present study were too low to be indicative of a negative energy balance in TW calves
462 (which had greater values than UF+UV-fed calves). Lastly, an increase of serum thyroid
463 hormone in 8L-fed calves was expected since it is a hormone related to growth and
464 basal metabolic rate, and 8L-fed calves grew more compared with 4L-fed calves.
465 To determine the preference and acceptability of UF+UV water, a preference water test
466 was performed. When data were analysed with all calves together with a t-test without
467 considering their previous experience, calves clearly showed a preference for TW (0.76
468 ± 0.297). However, the confidence limits for no preference were very wide (between
469 0.91 ± 0.30 and 0.22 ± 0.45), indicating a high variability among animals. When

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470 analysing the effects of a previous experience on water exposure, calves that were
471 consuming TW throughout the study, afterwards did not show any preference for any of
472 the waters during the preference test (preference ratio of 0.53 ± 0.08); in contrast, calves
473 previously exposed to UF+UV had a clear preference for TW during the preference test
474 (preference ratio of 0.91 ± 0.07). This outcome demonstrated firstly, the ability of
475 calves to distinguish between the two types of water, and secondly that the UF+UV
476 water obtained in the present study, in spite of not posing any important risk for the
477 animal health, has a lower acceptability and palatability than the TW water.

478 **4. Conclusions**

479 The experiment involving intestinal primary cell cultures pointed the ultrafiltration
480 treatment of reclaimed water as necessary to prevent intestinal cell damage, apoptosis,
481 and inflammation. Wastewater treated with an ultrafiltration and ultraviolet treatment
482 seems a plausible potential option for livestock drinking. However, some
483 recommendations can be drawn from the present work when considering the use of
484 reclaimed water for livestock drinking:

485 • Multi-barrier technologies for water reclamation, such as ultrafiltration and
486 ultraviolet disinfection, achieve a desirable water physicochemical and microbiological
487 quality for livestock drinking. However, high contents of soluble salts, found in the
488 present study at concentrations around the threshold limits set up for human and
489 livestock drinking water, may reduce water palatability and acceptability and
490 consequently impair water and concentrate intakes and ultimately negatively affect
491 animal performance, but without causing noticeable health afflictions.

492 • The livestock sector is a promising candidate for the reuse of urban wastewater,
493 especially during severe drought periods. However, equilibrium between reclaimed
494 water availability, water demand, technology cost, environment sustainability, and

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495 animal health and performance should be evaluated when proposing a water reuse
496 scheme for livestock.

497 • Testing for the presence of human and/or animal viral fecal indicators in the
498 wastewater used for producing reclaimed water may be relevant since it could serve for
499 tracing the origin of fecal contamination that may pose a risk for animal or/and human
500 health.

501 • Using reclaimed water in the livestock sector is challenging because it requires
502 relatively important economic investments and further studies are needed to evaluate
503 long-term effects on animal health and performance are required for its potential impact
504 on food safety.

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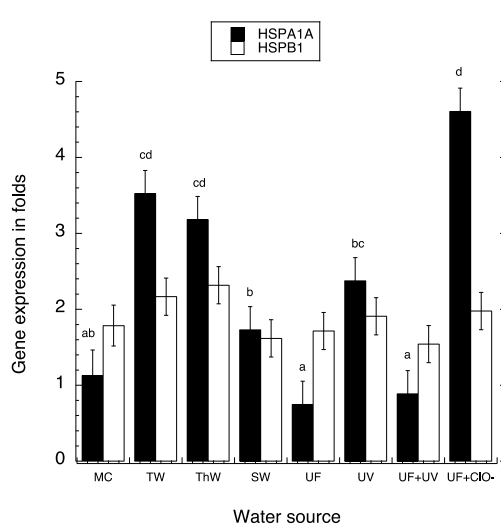
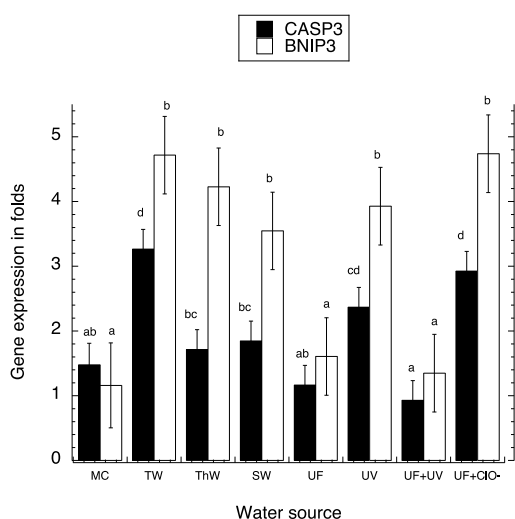
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1 **Figure 1.** Expression of apoptotic (a), cell damage (b) and inflammation (c) genes,
 2 expressed as relative media culture sample folds, of *in vitro* intestinal cells cultured with
 3 different types of water: cells media culture (MC), tap water (TW), dairy cow trough
 4 water that was filled with chlorinated ground water (ThW), secondary effluent from
 5 Caldes WWTP (SW), secondary effluent with an ultrafiltration (UF), secondary effluent
 6 with an UV disinfection (UV), secondary effluent with UF and UV treatments
 7 (UF+UV), and secondary effluent with UF and chlorination (UF+ClO⁻). Columns with
 8 different letters indicate differences within gene among water treatments.

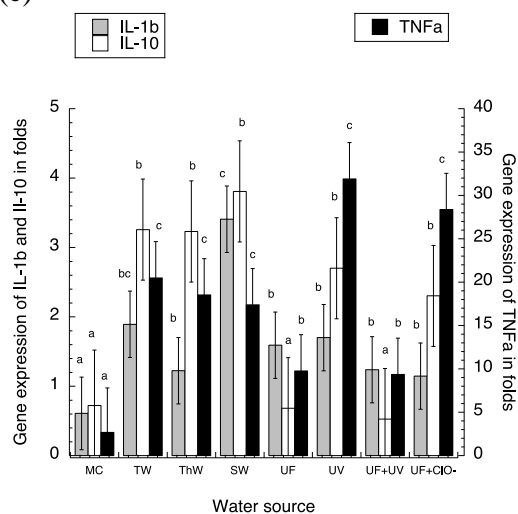
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(b)



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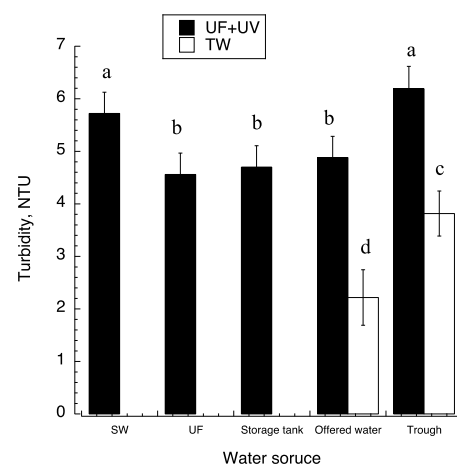
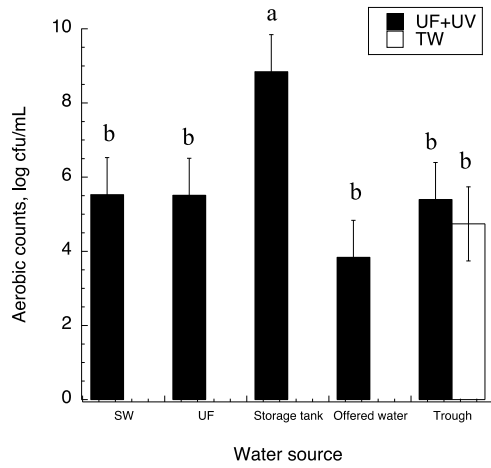


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1 **Figure 2.** Evolution of aerobic counts (a) and turbidity (b) of UF+UV reclaimed water
2 during the production process in the discontinuous system, and the final quality in
3 animal troughs.

4 (a)

(b)



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Table 1. Water chemical and microbiological quality parameters of the different waters used in the *in vitro* Study 1.

	Source of water ¹						
	TW	ThW	SW	UF	UV	UF+UV	UF+ClO ⁻
pH	7.8	7.8	7.5	7.3	7.5	7.5	7.5
Conductivity, $\mu\text{S}/\text{cm}$	570	601	980	943	985	971	951
Turbidity, NTU	0.6	3.7	2.5	0.6	2.3	1.1	1.3
COD, mgO_2/L	< 50	< 50	< 50	< 50	< 50	< 50	< 50
NH ₄ ⁺ , mg/L	< 0.1	< 0.5	< 0.1	< 0.1	< 0.1	< 0.1	< 0.1
Chloride, mg/L	36	41.5	173	165	173	174	167
Phosphate, mg/L	< 0.2	< 0.5	1.5	1.6	1.2	1.2	1.2
Nitrate, mg/L	9.9	< 0.5	19	18	17	17	17
Sulphate, mg/L	45	ND	45	42	43	43	42
Aerobic counts, cfu/mL	0	1,540,000	205,000	< 150	4	1	0
<i>E. coli</i> , cfu/mL	< 5	58	7,600	< 5	< 5	< 5	< 5

¹ TW= tap water; ThW= water from an animal trough; SW= secondary effluent of a wastewater treatment plant; UF= secondary effluent with an ultrafiltration treatment; UV= secondary effluent with an ultraviolet disinfection; UF+UV= secondary effluent with an ultrafiltration treatment and an ultraviolet disinfection; UF+ClO⁻= secondary effluent with an ultrafiltration treatment and a chlorination

ND = not determined

Table 2. Gene names, primer sequences, annealing temperature, primer concentration, and efficiency of the used genes in Study 1.

Gene name	Primer sequence (5' to 3')	Tm	μ M	Efficiency	Reference	
β -actin (<i>ACTB</i>)	Fw	CTGGACTTCGAGCAGGAGAT	57°C	0.125	1.82	Bach et al., 2018
	Rv	CCCGTCAGGAAGCTCGTAG				
Tumor necrosis factor alpha (TNF- α)	Fw	AACAGCCCTCTGGTTCAAAC	60°C	0.5	1.89	Riollet et al., 2001
	Rv	TCTTGATGGCAGACAGGATG				
Interleukin 1 beta (IL-1 β)	Fw	TGGGAGATGGAAACATCCAG	50°C	0.3125	1.82	Riollet et al., 2001
	Rv	TTTATTGACTGCACGGGTGC				
Interleukin 10 (IL-10)	Fw	ACTTTAAGGGTTACCTGGGTTG	57°C	0.5	1.90	Bruno et al., 2010
	Rv	GAAAGCGATGACAGCGCCGC				
Adenovirus E1B 19 kDa protein-interacting protein 3 (BNIP3)	Fw	GAAGGAATGCCGACACTAGG	55°C	0.5	1.85	Nishimura et al., 2008
	Rv	CAAAGCCAGCAGACACTCAG				
Caspase 3 (CASP3)	Fw	AAGCCATGGTGAAGAAGGAA	55°C	0.5	1.88	Nishimura et al., 2008
	Rv	GGCAGGCCTGAATAATGAAA				
Heat shock 70 kDa protein 1A (HSPA1A)	Fw	GGCACCAGAGCTTCACGATGT	60°C	0.5	1.91	Bach et al., 2018
	Rv	CCTACGCAGGAGTAGGTGGT				
Heat shock protein family B member 1 (HSPB1)	Fw	CCTGAAACACCGCCTGCTAA	60°C	0.5	1.92	Bach et al., 2018
	Rv	CGGAGAAGCGAGAGAAGTGG				

Table 3. Physicochemical and microbiological analysis of tap water (TW) and ultrafiltered and ultraviolet treated wastewater (UF+UV) used in Study 2 to prepare milk replacer and as drinking water for young calves.

	Type of water		SEM	P-value	Upper limits
	TW	UF+UV			(Schlink et al., 2010)
pH	7.8	8.4	0.07	0.001	-
Conductivity, $\mu\text{S}/\text{cm}$	570	1,437	38.8	<0.001	-
Total suspended solids, mg/L	1.33	0.92	0.236	0.29	-
Turbidity, NTU	0.6	1.8	1.23	0.53	-
Total organic carbon, mg/L	2.0	4.9	0.43	<0.01	-
Total inorganic carbon, mg/L	40.0	46.9	-	-	-
Total dissolved solids, mg/L	384	756	-	-	-
Chloride, mg/L	36	269	5.03	<0.01	100
Sulphate, mg/L	45.3	56.7	4.11	0.19	50
Bromide, mg/L	0.55	0.45	0.403	0.88	-
Nitrate, mg/L	9.9	16.4	1.05	0.05	89
Nitrite, mg/L	< 0.2	< 0.2	-	-	-
Phosphate, mg/L	< 0.2	4.5	0.57	0.03	2.15
Ca ²⁺ , mg/L	57.8	74.8	4.75	0.13	100
Mg ²⁺ , mg/L	18.5	19.5	2.68	0.82	50
Na ⁺ , mg/L	20.3	188.1	1.34	<0.001	50
K ⁺ , mg/L	2.8	20.2	0.11	<0.001	20
NH ₄ ⁺ , mg/L	< 0.2	< 0.2	0.03	0.67	-
Total Mn, $\mu\text{g}/\text{L}$	2.3	11	-	-	50

Total Fe, µg/L	5.7	15	-	-	200
Aerobic counts, cfu/mL	ND ¹	1.4E+05	2.04E+05	-	-
Total coliforms, cfu/mL	0	1.6	-	-	-
<i>Escherichia coli</i> , cfu/mL	0	0.4	0.9	-	-
<i>Clostridium perfringens</i> , cfu/mL	0	0	-	-	-
<i>Cryptosporidium</i> , spp, oocyst/L	ND ¹	< 8	-	-	-
Bovine polyomaviruses, GC/L	ND ¹	< 105	-	-	-
Human adenoviruses, GC/L	ND ¹	< 105	-	-	-
Somatic coliphages, pfu/mL	ND ¹	5	5.8	-	-
Helminth eggs, egg/10L	ND ¹	< 1	-	-	-
<i>Taenia</i> spp eggs, egg/10 L	ND ¹	< 1	-	-	-

¹ Not determined

1 **Table 4.** Performance and dry matter (DM) intake of calves fed two different milk
 2 feeding programs (4 L/d vs 8 L/d of milk replacer), with two different sources of water:
 3 tap water (TW) vs ultrafiltered and ultraviolet (UF+UV) treated wastewater.

Water source	Treatments				SEM ²	P-values ¹		
	TW		UF+UV			W	M	WxM
Milk program	4L	8L	4L	8L				
Number calves	4	4	5	5	-	-	-	-
Initial age, d	5.3	3.5	4.8	5.4	1.63	0.66	0.73	0.48
Initial body weight, kg	40.6	37.8	40.0	41.0	3.21	0.69	0.79	0.57
Final body weight, kg	66.0	71.9	63.3	69.3	1.57	0.35	0.001	0.60
Average daily gain, g/d	616	778	556	692	49.1	0.14	0.003	0.79
Dry matter intake, g/d								
Milk replacer	487	861	487	872	0.02	0.72	<0.001	0.73
Concentrate	650	282	532	170	45.6	0.01	<0.001	0.94
Straw	17	15	14	15	7.6	0.81	0.94	0.85
Water intake, L/d	3.4	3.2	2.9	2.8	0.41	0.19	0.64	0.86
Feed efficiency ³	0.53	0.68	0.53	0.69	0.04	0.89	<0.001	0.94

4 ¹ W: effect of the type of water used to prepare the MR and for drinking calves; M:
 5 effect of the volume of MR offered to calves; WxM: effect of the interaction of source
 6 of water and amount of milk replacer

7 ² standard error of the mean

8 ³ Expressed as ratio between daily gain and daily feed consumption

9