

## Erratum

### Journal of Water and Health 2(4), 249–260: Tracking the origin of faecal pollution in surface water. An ongoing project within the European Union research programme, A. Blanch *et al.*

Unfortunately due to an oversight during production, Table 1 of the above paper was published with an error in the Table caption. The correct caption is published below, with the relevant table.

**Table 1** | Data from municipal or hospital sewage (human) and slaughterhouse wastewater (animal). Results are the average of log CFU/100 mL or log PFU/100 mL for faecal coliforms (FC), enterococci (FE), clostridia (CL), somatic coliphages (SOMCPH), total F bacteriophages (FTOTAL), F-RNA bacteriophages (FRNAPH), *Bact. fragilis* bacteriophages (RYC2056). Faecal sterols are in  $\mu\text{g/g}$ : coprostanol (COP), stigmastanol or 24-ethylcoprostanol (ETHYLCOP), epicoprostanol (EPICOP) and cholestanol (CHOL). The percentage of the rest of the values is also the average of samples: diversity of enterococci (DiE), *Ent. faecium* plus *Ent. faecalis* (FM-FS), *Ent. hirae* (Hir), *E. coli* Phene System patterns (ECP), *E. coli* confirmed (ECT). *N*: number of samples

	Human			Animals		
	Average	SD	<i>N</i>	Average	SD	<i>N</i>
FC	7.05	0.58	89	7.60	0.94	95
FE	6.00	0.81	89	6.52	0.83	95
CL	5.20	0.69	89	5.29	1.26	95
SOMCPH	6.07	0.93	89	6.77	1.10	94
FTOTAL	5.48	0.76	87	5.43	1.52	85
FRNAPH	5.23	1.01	87	5.31	1.58	81
Percentage of genotypes of F-RNA phages						
I	7	–	83	34	–	71
II	40	–	83	12	–	71
III	48	–	83	28	–	71
IV	5	–	83	26	–	71
RYC2056	3.82	1.01	85	3.76	1.23	85
COP	1042.73	3436.66	70	252.92	1147.86	69
ETHYLCOP	657.96	3151.15	70	2840.69	10210.25	69
EPICOP	643.79	3316.43	70	2033.00	10347.60	69
CHOL	1361.60	6351.94	70	441.26	1351.80	69
DiE	0.85	0.19	84	0.76	0.25	95
FM-FS	66.10	20.69	84	49.03	30.63	94
Hir	17.15	18.48	84	28.69	31.12	94
DiC	0.89	0.16	88	0.88	0.09	95
ECP	79.10	22.65	77	97.11	5.37	84
ECT	72.13	24.29	77	93.30	9.10	84