Host-Specific Viruses for the Detection of Faecal Pollution in **Coastal Waters**

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Aims

□ Investigate the host-specificity and -sensitivity of human-specific adenoviruses (HS-AVs), bovine-specific adenoviruses (BS-AVs), and human-specific polyomaviruses (HS-PVs) for microbial source tracking in coastal waters

□ Application of viral markers to identify the sources of faecal pollution in a coastal river affected by faecal pollution in Southeast Queensland (SEQ), Australia.

Results

Viral markers	Host-specificity	Host-sensitivity
HS-AVs	1.0	0.78
BS-AVs	1.0	0.73
HS-PVs	1.0	0.99

Materials and Methods

• Culture based methods were used for the enumeration of faecal indicator bacteria (FIB).

□ Host-specificity and -sensitivity of the markers were assessed by screening 182 wastewater and animal faecal samples.

□ PCR and qPCR assays were used for the detection of MST markers and quantification of zoonotic pathogens in environmental water samples (n = 20).



Table 1: Host-specificity and -sensitivity of viral markers



Figure 3: Percentage of samples exceeded ANZECC recreational water quality guideline value for primary contact

□ The numbers of FIB in water samples collected ranged from 48 300 (for *E. coli*) and from 60 20 to 1586 11 to 2906 180 (for

Figure 1: Host-groups screened for the host-specificity and -sensitivity assays



enterococci).



Figure 4: Percentage of samples positive for the viral markers and zoonotic pathogens

□ The numbers of *Salmonella* spp. ranged from 350 40 to 430 70 genomic copies per 500 mL of water.

 \Box Weak correlation was found between *E. coli* with HS-AVs (p =0.02). However, significant correlations were observed between *E*. *coli* with BS-AVs (p = 0.007) and *Salmonella* spp. (p = 0.007)

 \Box The numbers of enterococci correlated with BS-AVs (p =0.006), *C. jejuni* (p = 0.01) and *Salmonella* spp. (p = 0.006). No significant correlations were found between E. coli and enterococci with HS-PVs.

□ The results suggest that human and bovine specific viral markers detection using PCR could be a useful tool for the identification of human and bovine faecal pollution in coastal waters.

Further information

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