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## Research papers

# Transport and fate of viruses in sediment and stormwater from a Managed Aquifer Recharge site



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## ABSTRACT

Enteric viruses are one of the major concerns in water reclamation and reuse at Managed Aquifer Recharge (MAR) sites. In this study, the transport and fate of bacteriophages MS2, PRD1, and ΦX174 were studied in sediment and stormwater (SW) collected from a MAR site in Parafield, Australia. Column experiments were conducted using SW, stormwater in equilibrium with the aquifer sediment (EQ-SW), and two pore-water velocities (1 and 5 m day<sup>-1</sup>) to encompass expected behavior at the MAR site. The aquifer sediment removed >92.3% of these viruses under all of the considered MAR conditions. However, much greater virus removal (4.6 logs) occurred at the lower pore-water velocity and in EQ-SW that had a higher ionic strength and Ca<sup>2+</sup> concentration. Virus removal was greatest for MS2, followed by PRD1, and then ΦX174 for a given physicochemical condition. The vast majority of the attached viruses were irreversibly attached or inactivated on the solid phase, and injection of Milli-Q water or beef extract at pH = 10 only mobilized a small fraction of attached viruses (<0.64%). Virus breakthrough curves (BTCs) were successfully simulated using an advective–dispersive model that accounted for rates of attachment ( $k_{att}$ ), detachment ( $k_{det}$ ), irreversible attachment or solid phase inactivation ( $\mu_s$ ), and blocking. Existing MAR guidelines only consider the removal of viruses via liquid phase inactivation ( $\mu_l$ ). However, our results indicated that  $k_{att} > \mu_s > k_{det} > \mu_l$ , and  $k_{att}$  was several orders of magnitude greater than  $\mu_l$ . Therefore, current microbial risk assessment methods in the MAR guideline may be overly conservative in some instances. Interestingly, virus BTCs exhibited blocking behavior and the calculated solid surface area that contributed to the attachment was very small. Additional research is therefore warranted to study the potential influence of blocking on virus transport and potential implications for MAR guidelines.

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## 1. Introduction

The availability of high-quality drinking water has decreased due to climate variability and population growth, while demand has increased to meet agricultural, industrial, environmental, and municipal water needs (Levantesi et al., 2010; Shannon et al., 2008; Yates et al., 1987). The United Nations has estimated that about 1.8 billion people around the globe will face severe water stress and scarcity by 2025 (UN, 2013). Consequently, there is an urgent need to find alternative sources of freshwater, such as by developing economic and effective water reclamation, recycling, and preservation techniques (Levantesi et al., 2010; Shannon

et al., 2008; Yates et al., 1987). Water recycling can be achieved by various engineering techniques and natural or passive treatment. Major water recycling techniques employ multiple barrier approaches, including secondary treatment, riverbank filtration, reverse osmosis, UV disinfection, chlorination, and ultrafiltration (Page et al., 2010a). However, many countries across the world do not have access to cheap and efficient wastewater treatment plants (Vega et al., 2003). Managed Aquifer Recharge (MAR) is a collection of natural treatment techniques, including Aquifer Storage, Transfer, and Recovery (ASTR) or Aquifer Storage and Recovery (ASR), that has gained a lot of attention recently (Ayuso-Gabella et al., 2011; Bekele et al., 2014, 2013, 2011; Stevens, 2014; Toze and Bekele, 2009; Ward and Dillon, 2009). Water recycling is facilitated during MAR by purposefully recharging lower quality water into aquifers for natural treatment prior to recovery (Dillon et al., 2009; Page et al., 2010a). Currently, MAR is considered as an effec-

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tive and economical option to store water in the treatment train; e.g., a sequence of multiple stormwater treatments, which are designed to meet the needs of a particular environment, in order to maximize results (MelbourneWater, 2017). MAR provides a natural buffer, increases public perception, provides a residence time that can facilitate removal of biodegradable organic matter and pathogens, improves the quality of the treated wastewater or stormwater, and reduces the cost of seasonal peak demands (Dillon et al., 2009; Levantesi et al., 2010; Page et al., 2010a; Page et al., 2010c). However, current MAR guidelines require expensive and energy-intensive pre-treatment of injected water and post-treatment of recovered water depending on the end use, with drinking water supply requiring the highest level of treatment (Dillon et al., 2009).

One of the major concerns with potable water reuse is the microbiological quality of recovered water and the possibility of transmitting infectious diseases from pathogenic microorganisms (virus, bacteria, and protozoa) that are not eliminated by conventional wastewater treatment (Costán-Longares et al., 2008; Levantesi et al., 2010; Shannon et al., 2008). Enteric viruses pose the greatest public health concern because they can travel long distances (Schijven and Hassanizadeh, 2000) and are infectious at very low doses (Ward et al., 1986). MAR systems can improve the microbiological quality of water by natural attenuation processes during soil filtration and/or aquifer transport (Asano et al., 2007; Dillon et al., 2008; Levantesi et al., 2010; Mayotte et al., 2017). For example, inactivation occurs when viruses lose their ability to infect host cells and replicate because of the disruption of proteins and the degradation of nucleic acid (Gerba, 1984; Schijven et al., 2003). The most important factors affecting virus inactivation rate include temperature, groundwater microbial activity, pH, salt species and concentration, some forms of organic matter, and virus type (McCarthy and McKay, 2004; Schijven and Hassanizadeh, 2000). However, decay rates for human enteric viruses, determined using diffusion chambers in monitoring wells at MAR sites, have been found to be slow and nonlinear (Sidhu et al., 2015; Sidhu and Toze, 2012). Quantitative microbial risk assessment calculations for MAR systems have been developed from a detailed hydrogeological assessment of the aquifer and in-situ decay studies (Donald et al., 2011; Page et al., 2010b, 2015a; Toze et al., 2010). These risk assessment studies considered that liquid phase virus inactivation was the only reliable mechanism for virus removal in the aquifer, and neglected the processes of virus attachment, detachment, and solid phase inactivation (Abu-Ashour et al. (1994); Dillon et al., 2008).

In addition to liquid phase inactivation, virus removal from groundwater may occur by attachment from the bulk solution to the solid phase (Shen et al., 2012a,b), whereas detachment refers to the reverse process of virus release from the solid phase to the bulk solution (Bergendahl and Grasso, 2000). Virus attachment to aquifer materials is a strong function of many physicochemical variables, including pore-water velocity (Hijnen et al., 2005); solution ionic strength (IS) (Da Silva et al., 2011; Knappett et al., 2008; Xu et al., 2005); solution ionic composition (Bales et al., 1991; Lipson and Stotzky, 1983; Sadeghi et al., 2013; Sasidharan et al., 2014; Walshe et al., 2010); and the presence of humic materials (Zhuang and Jin, 2003), and metal oxides (Foppen et al., 2006). Consequently, an accurate assessment of virus attachment at MAR sites must consider realistic solution chemistries and aquifer mineralogy. In contrast, most virus transport studies have been conducted under highly idealized conditions using clean sand or glass beads, and simple electrolyte solutions.

Attachment during MAR can only serve as an effective, long-term treatment when viruses are irreversibly retained or inactivated on the solid phase. It is difficult to separately quantify the processes of irreversible attachment and solid phase inactivation

because they both decrease the number of infective viruses that can detach from the solid phase (Ryan et al., 2002). Consequently, much less is known about solid than liquid phase inactivation (Murray and Laband, 1979; Ryan et al., 2002). Solid phase inactivation has been reported to increase with the strength of the adhesive interaction and the temperature (Loveland et al., 1996; Murray and Laband, 1979; Ryan et al., 2002). Yates et al. (1985) found that the inactivation rate of MS2 in eleven groundwater samples increased with the  $\text{Ca}^{2+}$  concentration. However, low levels of virus detachment have been commonly observed under steady-state physicochemical conditions (Bales et al., 1993). Furthermore, changes in solution chemistry (e.g., a decrease in IS or an increase in pH) have been observed to produce large pulses of released colloids (Bales et al., 1993). Consequently, there is a concern that attached viruses can be remobilized with a decrease in IS or divalent cation concentrations during rainfall events (Gerba, 1983; Yates et al., 1988).

The main objective of this work was to systematically examine virus removal processes in urban stormwater from a wetland (Urrbrae, South Australia) when in contact with aquifer sediment collected from an ASTR site (Parafield Gardens, South Australia). Three bacteriophages (PRD1, MS2, and  $\Phi$ X174) were used as surrogate viruses for enteric human viruses. Laboratory scale virus transport experiments and inactivation studies were conducted. Additional studies investigated the release of attached viruses by sequentially injecting step pulses of solutions with alternating solution chemistry. Virus breakthrough curves (BTCs) were simulated using the numerical solution of the advection–dispersion equation with terms for attachment, detachment, Langmuirian blocking, and a sink term that accounted for irreversible attachment and solid phase inactivation. Model parameters were obtained by inverse fitting to the observed BTCs. Results from this study provide valuable insight on the relative importance of natural attenuation processes for viruses at MAR sites and indicate that microbial risk assessments that only consider liquid phase inactivation may be overly conservative in some instances.

## 2. Materials and methods

### 2.1. Stormwater

Stormwater (SW) samples were collected from the Urrbrae Wetland located in Mitcham, South Australia (Fig. S1). The wetlands were constructed to collect urban stormwater for flood protection in the nearby area and for potential future use in MAR operations (Lin et al., 2006). The stormwater chemistry (major and minor elements) was analyzed using an Inductively Coupled Plasma Mass Spectrometry (ICP-MS) and the carbon content was measured using Varo TOC Cube (Analytical Chemistry, CSIRO, Adelaide). The pH and electrical conductivity (EC) were measured using a Eutech PC 700 (Eutech Instruments).

Dissolved calcium is a weathering product of almost all rocks and is, consequently, abundant in most groundwater sources. Water from limestone aquifers may contain 30–100  $\text{mg L}^{-1}$  of calcium, gypsiferous shale aquifers may contain several hundred milligrams per liter (Sadeghi et al., 2013), and dolomite produces water with high levels of calcium and magnesium (Wade, 1992). The calcium concentrations range from 0.03 to 36.5  $\text{mg L}^{-1}$  in Australian groundwaters (Radke et al., 1998). When water percolates through soils, the concentration of divalent cations in soil solutions often increases with depth along the vertical weathering-leaching gradient (Sadeghi et al., 2013; Sverdrup and Warfvinge, 1993). It is, therefore, logical to anticipate that the solution IS and divalent cation concentration will increase with a travel distance and residence time during ASTR, due to mineral dissolution, along with

mixing in aquifers containing groundwater of higher salinity. In a limestone aquifer, increases in calcium are prevalent due to the dissolution of calcium carbonate. This increase in solution IS and  $\text{Ca}^{2+}$  concentration may increase virus attachment. Virus transport and survival experiments were therefore also conducted using stormwater after it was equilibrated with the aquifer sediment (50 g of sediment with 200 mL of stormwater), denoted as EQ-SW. Table S1 provides a summary of selected chemical properties for EQ-SW.

## 2.2. Porous media

Aquifer materials were collected from an ASTR site situated in Parafield Gardens, Adelaide, Australia (Fig. S1) established for stormwater storage and treatment (Page et al., 2015b). The target aquifer for ASTR is the lower Tertiary marine sediments of the Port Willunga Formation (T2 aquifer), a well-cemented sandy limestone aquifer which is intersected between  $-149$  and  $-214$  m Australian Height Datum (AHD) (160–220 m below ground surface). At this location, the target aquifer is approximately 60 m thick and overlain by 7 m thick clay aquitard of Munno Para Clay which prevents the migration of injected stormwater to the overlying aquifers. A schematic cross-section of the aquifer is given in Fig. S2. Karstic features were not identified during construction of the site (Vanderzalm et al., 2010). The mineralogy in the storage zone is dominated by calcite ( $65 \pm 23\%$ ), quartz ( $30 \pm 22\%$ ), and a trace amount of ankerite, goethite, hematite, pyrite, albite, and microcline (Vanderzalm et al., 2010). Goethite and hematite largely account for the 2.1%  $\text{Fe}_2\text{O}_3$  quantified by X-ray Fluorescence (XRF). The transmissivity of the aquifer ranges from 100 to  $>200 \text{ m}^2 \text{ d}^{-1}$ , depending on the silt content and degree of weathering (Miotliński et al., 2014). Considerable variability exists in hydraulic conductivity within the aquifer and, therefore, the ASTR was constructed with partially penetrating wells to avoid the zone of high hydraulic conductivity in the lower part of the aquifer and to exclude excessive mixing between native groundwater and injection water (Miotliński et al., 2014). The aquifer is low in organic carbon ( $<0.5\%$ ) and has an average cation exchange capacity of 1.5 meq/100 g (Page et al., 2010b). The aquifer has a temperature of 25 °C at ambient conditions. The design, operation, hydraulic, chemical, and physical properties of this aquifer has been extensively studied and reported in the literature (Adkinson et al., 2008; Dillon et al., 2008; Kremer et al., 2008; Page et al., 2010b; Pavelic et al., 2004; Rinck-Pfeiffer et al., 2005; Vanderzalm et al., 2010). The aquifer material used in this study was obtained from intact core samples taken at a depth of 171.30 m below the ground surface. The collected aquifer sediments were directly used in the experiments without any further treatment. Selected physical and chemical properties of the sediment are provided in Table S2. The bulk mineralogy of the sediment sample was determined by X-ray Diffraction (XRD). The major elemental composition was determined by XRF and Inductively Coupled Plasma Optical Emission Spectrometry (ICP-OES) following reverse aqua regia acid digest. Sieve analysis was used to determine the particle size distribution.

Additional transport experiments were conducted using ultra-pure clean quartz sand (Charles B. Chrystal CO., Inc., NY, USA) with size ranging from 125 to 300  $\mu\text{m}$ . This sand was cleaned using an acid wash and boiling procedure described by Sasidharan et al. (2014). Transport experiments with the ultra-pure quartz sand represent a worst-case scenario for virus transport because of minimal chemical heterogeneity.

## 2.3. Viruses

Bacteriophage MS2, PRD1, and  $\Phi\text{X174}$  were used as model viruses in this study. These microbes were chosen because of their

structural resemblance to many human enteric viruses and they have been used in numerous investigations as surrogates for human enteric viruses (Chu et al., 2003; Schijven and Hassanizadeh, 2000). It should be mentioned that recent studies have demonstrated that the transport and fate of human enteric viruses may not always be well correlated with that of bacteriophages (Bellou et al., 2015). Additional research is, therefore, warranted to identify the best surrogate for the transport and fate of pathogenic viruses, but this issue is beyond the scope of the present study. The bacteriophages were analyzed using their respective *Escherichia coli* host. The production of bacteriophages and their analysis using double layer agar (DLA) methods were detailed in Sasidharan et al. (2016). Stock solutions of viruses were diluted in stormwater to obtain an initial concentration ( $C_0$ ) of  $\sim 5 \times 10^6$  plaque forming unit  $\text{mL}^{-1}$ . The survival of viruses over a 7 h interval was determined in both SW and EQ-SW solutions.

## 2.4. Zeta potential and size measurements

The electrophoretic mobility (EM) of viruses, crushed aquifer sediment, and quartz ( $<2 \mu\text{m}$ ) was measured in both SW and EQ-SW using a Zetasizer (Malvern, Zetasizer Nano Series, Nano-ZS). The EM measurements were repeated five times with more than twenty runs per measurement. The Smoluchowski equation (Elimelech et al., 1994) was used to convert the measured EM values to zeta potentials. The size distribution of viruses in both SW and EQ-SW was measured using a dynamic light scattering (DLS) (Malvern Instruments Ltd, 2004; Sikora et al., 2016) process (Malvern, Zetasizer Nano Series, Nano-ZS).

## 2.5. Transport experiments

The column experiments were set up in a constant temperature laboratory (20 °C). Sterilized polycarbonate columns (1.9 cm inside diameter and 11 cm height) were wet packed using aquifer material or ultra-pure quartz sand while the column was being vibrated. After packing, the column was preconditioned with  $>10$  pore volumes (PV) of stormwater water using a syringe pump (Model 22, Harvard Apparatus) at a flow rate of 0.394  $\text{mL min}^{-1}$ . A suspension with known  $C_0$  of viruses (PRD1, MS2, and  $\Phi\text{X174}$ ) and solution chemistry (SW or EQ-SW) was introduced into the column using a syringe pump at a constant pore water velocity (1 or 5  $\text{m day}^{-1}$ ) for 13 PV (Phase I), followed by injection of  $\sim 7$  PV of virus-free solution at the same solution chemistry and velocity (Phase II). To study the reversibility of attached viruses, the experiments were continued by flushing the columns with  $\sim 11$  PV of Milli-Q water (Phase III), followed by injection of 3% Beef Extract with pH = 10 (Phase IV). Virus release is enhanced in Milli-Q water by a reduction in solution IS which expands the double layer thickness, and increases the magnitude of the sediment and virus zeta potentials (Chen et al., 2014; Sasidharan et al., 2017b). Beef extract is a mixture of peptides, amino acids, nucleotide fractions, organic acids, minerals, and some vitamins, with a pH of 10 and its injection further enhances the virus release by masking positively charged sites with organic matter, and reversing the charge of some pH dependent sites (Landry et al., 1978; McMinn, 2013). Effluent samples were collected using a fraction collector (CF-2, Spectrum, USA). The effluent concentrations ( $C$ ) of viruses were enumerated using the methods described above. Separate column experiments were run for each porous medium (aquifer sediment or ultra-pure quartz sand), solution chemistry (SW or EQ-SW), and pore-water velocity (1 or 5  $\text{m day}^{-1}$ ) combination.

Breakthrough curves (BTCs) were plotted as a dimensionless concentration ( $C/C_0$ ) of viruses as a function of PVs. A mass balance was conducted for the viruses in the column experiments using information on injected and recovered viruses during Phases I–



IV. The percentage of virus mass retained on the solid phase ( $M_s$ ) was determined as the difference in the mass of injected virus and mass of virus recovered in the effluent BTC ( $MBTC = M_I + M_{II}$ ) during Phases I and II. The percentage of injected viruses that was recovered during Phases III and IV were denoted as  $M_{III}$  and  $M_{IV}$ , respectively. The percentage of injected viruses that were irreversibly retained ( $M_{irr}$ ) was determined as  $100 - MBTC - M_{III} - M_{IV}$ . The log removal of the viruses in the column effluent experiments was determined as  $-\log_{10}(M_{BTC})$ .

### 2.6. Simulation of virus BTCs

Experimental BTCs for viruses were simulated using the HYDRUS-1D model (Šimůnek et al., 2016). The HYDRUS-1D program numerically solves the Richards' equation for variably saturated water flow and Fickian-based advection–dispersion equations for heat and solute transport. The governing continuum-scale flow and transport equations are solved numerically using Galerkin-type linear finite element schemes (Šimůnek et al., 2016). The following aqueous and solid phase mass balance equations were considered in this model for each virus.

$$\frac{\partial C}{\partial t} = \lambda v \frac{\partial^2 C}{\partial z^2} - v \frac{\partial C}{\partial z} - k_{att}\psi C - k_{det} \frac{\rho_b}{\theta} S - \mu_l C \quad (1)$$

$$\frac{\rho_b}{\theta} \frac{\partial S}{\partial t} = k_{att}\psi C - k_{det} \frac{\rho_b}{\theta} S - \mu_s \frac{\rho_b}{\theta} S \quad (2)$$

where  $t$  (T; T denotes unit of time) is time,  $z$  (L; L denotes units of length) is the direction of mean water flow,  $C$  ( $NL^{-3}$ ; N denotes the virus number) is the aqueous phase virus concentration,  $\lambda$  (L) is the dispersivity,  $v$  ( $LT^{-1}$ ) is the average pore water velocity,  $\rho_b$  ( $ML^{-3}$ ; M denotes the unit of mass) is the bulk density,  $\theta$  is the water content,  $S$  ( $NM^{-1}$ ) is the solid phase concentrations of virus,  $k_{att}$  ( $T^{-1}$ ) is the virus attachment rate coefficient,  $k_{det}$  ( $T^{-1}$ ) is the virus detachment rate coefficient, and  $\mu_s$  ( $T^{-1}$ ) is a sink term which accounts for irreversible attachment and inactivation of viruses attached on the solid phase, and  $\mu_l$  ( $T^{-1}$ ) is the inactivation rate coefficient for viruses in the liquid phase. The parameter  $\psi$  is a dimensionless Langmuirian blocking function that is given as (Adamczyk et al., 1994)

$$\psi = \left(1 - \frac{S}{S_{max}}\right) \quad (3)$$

where  $S_{max}$  ( $NM^{-1}$ ) is the maximum solid phase concentrations of retained virus.

The fraction of the solid surface area that is available for retention ( $S_f$ ) was calculated from  $S_{max}$  as (Kim et al., 2009; Sasidharan et al., 2014):

$$S_f = \frac{A_c \rho_b S_{max}}{(1 - \gamma) A_s} \quad (4)$$

where  $A_c$  ( $L^2N^{-1}$ ) is the cross sectional area of a virus,  $A_s$  ( $L^{-1}$ ) is the solid surface geometric area per unit volume, and  $\gamma$  is the porosity of a monolayer packing of viruses on the solid surface that was taken from the literature to be 0.5 (Johnson and Elimelech, 1995).

The value of the sticking efficiency ( $\alpha$ ) was determined from the fitted  $k_{att}$  value and the filtration theory as (Schijven and Hassanizadeh, 2000; Yao et al., 1971):

$$\alpha = \frac{2d_c k_{att}}{3(1 - n)v\eta} \quad (5)$$

where  $n$  is the porosity (0.4) and  $d_c$  (L) is the collector (median grain) diameter. The value of the single collector-efficiency,  $\eta$ , was

calculated using the correlation equation presented by (Messina et al., 2015).

## 3. Results and discussion

### 3.1. Characterization of solution chemistry

SW had a pH of 7 and EC of  $260 \mu S cm^{-1}$  and after equilibration with the aquifer sediment the pH and EC of the EQ-SW increased to 7.3 and  $2230 \mu S cm^{-1}$ , respectively (Table S1). In addition, the concentration of all major ions also increased in EQ-SW, notably, with  $\sim 7$  times increase in Ca and K, and  $\sim 11$  times increase in Na concentration. Therefore, the ionic strength of the EQ-SW (0.014 mM) increased 10 times compared to SW (0.002 mM). This increase in pH and EC of EQ-SW compared to SW was attributed to the dissolution of limestone in the aquifer sediment (Table S2) that leaches  $CaCO_3$  and increases the solution pH (Earle, 2013; Panthi, 2003). The measured concentration value of  $Ca^{2+}$  in EQ-SW ( $151 mg L^{-1}$ , Table S1) was very close to  $Ca^{2+}$  ( $135 \pm 5 mg L^{-1}$ ) in ambient groundwater from the T2 aquifer (Page et al., 2010b) and, therefore, it confirms that the solution chemistry of the EQ-SW was representative of the target aquifer.

### 3.2. Characterization of aquifer sediment

The average particle size of the sediment was  $\sim 110 \mu m$ , with 93.5% within the 20–200  $\mu m$  size fraction (Table S2) and only 6.5% in smaller or larger size fractions. The quantitative bulk mineralogy of the sediment in Table S2 showed that the aquifer material was made of quartz (58.1%), calcite (35.0%), and goethite (2.2%). Major element analysis (Table S2) indicates the presence of a large fraction of  $SiO_2$  (53.4%),  $Al_2O_3$  (1.2%),  $Fe_2O_3$  (3.5%), and CaO (19.6%). Acid digest data (Table S2) revealed the presence of other major ions such as K, Mg, and Na in the sediment. These results demonstrated that the sediment was rich in major metal oxides such as CaO, MgO,  $Fe_2O_3$ , and  $Al_2O_3$ , as well as minerals such as quartz and clays. The metal oxide surfaces are positively charged and the mineral surfaces are negatively charged at the experimental pH ( $\sim 7.3$ ) due to surface chemical reactions with  $H^+/OH^-$  ions, respectively (Tombácz, 2009). Consequently, the aquifer sediment surface is expected to be chemically heterogeneous and possess a distribution of negative and positive surface charges. It has been shown that minor degrees of positive charge heterogeneity on the collector surface result in attachment rates that are an order of magnitude larger than similar surfaces having no charge heterogeneity (Schijven and Hassanizadeh, 2000).

Table 1 shows that the crushed aquifer sediment and quartz were negatively charged in the presence of SW and EQ-SW. We assume that some of the positively charged surfaces were masked

**Table 1**

The measured zeta potential values of viruses and sediment and the measured values of the size of viruses in stormwater (SW) and stormwater equilibrated with aquifer sediment (EQ-SW).

Virus	Solution	$\zeta$ + [mV]	Size [nm]
MS2	SW	$-22.4 \pm 1.2$	$27.6 \pm 3.4$
	EQ-SW	$-12.6 \pm 1.1$	$28.3 \pm 3.3$
$\Phi X174$	SW	$-23.8 \pm 1.2$	$29.6 \pm 2.6$
	EQ-SW	$-13.7 \pm 0.9$	$30.0 \pm 2.9$
PRD1	SW	$-23.3 \pm 1.4$	$69.0 \pm 0.82$
	EQ-SW	$-14.8 \pm 1.3$	$68.3 \pm 2.5$
Sediment	SW	$-19.8 \pm 0.8$	–
	EQ-SW	$-16.2 \pm 1.5$	–
Quartz	SW	$-31.5 \pm 2.8$	–
	EQ-SW	$-27.2 \pm 1.3$	–

by negatively charged P, Dissolved Organic Carbon (DOC), and Dissolved Inorganic Carbon (DIC) present in the SW (Table S1), which will lead to a reduction in the overall surface potential and a net negative zeta potential value for the sediment ( $\zeta_{sand}^-$ ) (Karageorgiou et al., 2007). Furthermore, the presence of  $\text{SiO}_2$  and clays will significantly decrease the value of  $\zeta_{sand}^-$  (Chen et al., 2014). Consequently, the zeta potential for quartz was lower than the aquifer sediment (Table 1). Table 1 also indicates that the zeta potential for the aquifer sediment and quartz was more negatively charged in the presence of SW than EQ-SW. This could be explained by an increase in EC, divalent cation ( $\text{Ca}^{2+}$  and  $\text{Mg}^{2+}$ ) concentration, and IS by more than 8 times in EQ-SW compared to SW, and the compression of the electrostatic double layer (Elimelech, 1994). Increased  $\text{Ca}^{2+}$  concentration can also lead to charge reversal (Lipson and Stotzky, 1983; Moore et al., 1981; Redman et al., 1999; Roy and Dzombak, 1996) and charge neutralization (Bales et al., 1991).

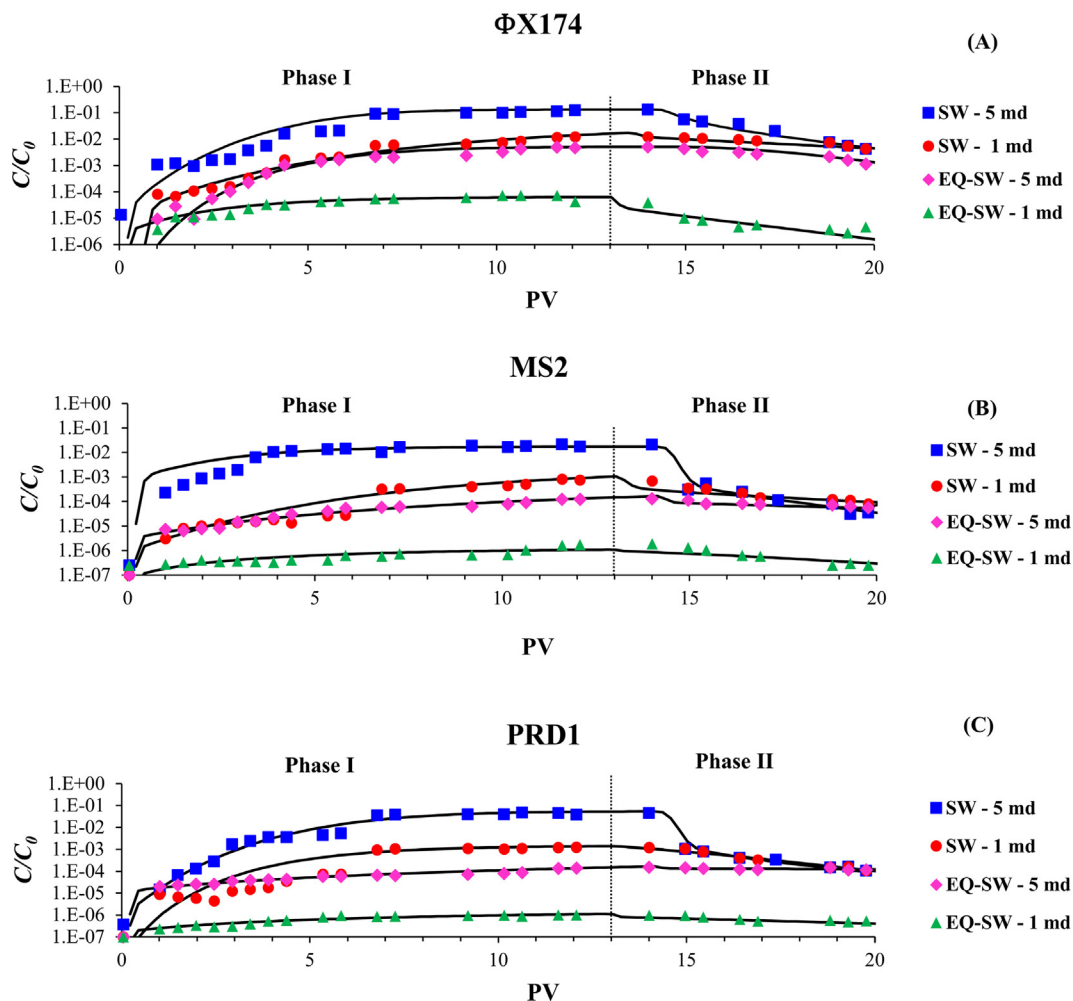
### 3.3. Characterization of viruses

The average size of MS2,  $\Phi\text{X174}$ , and PRD1 was measured to be 27.6–28.3 nm, 29.6–30.0 nm, and 68.0–69.0 nm, respectively. These values are within 3–5 nm of previously reported values (Chrysikopoulos and Aravantinou, 2012; Sasidharan et al., 2016; Thomson, 2005), and this indicates that aggregation of these

viruses was negligible under our experimental solution chemistry conditions. Table 1 shows the measured zeta potential values of viruses in SW and EQ-SW. The absolute value of the zeta potential for these viruses was always negative, but smaller in magnitude in the presence of EQ-SW compared to SW. Similar to the sediment, this can be explained by an increase in IS, compression of the double layer, and higher  $\text{Ca}^{2+}$  concentration for EQ-SW. In addition,  $\text{Ca}^{2+}$  is expected to bind to the carboxyl functional groups on the viral protein capsid which reduces the negative surface charge density (Harvey and Ryan, 2004).

### 3.4. Virus retention under various physicochemical conditions

Fig. 1 shows observed and fitted BTCs for the viruses under various solution chemistry (SW and EQ-SW) and pore-water velocity (1 and 5  $\text{m day}^{-1}$ ) conditions. The normalized effluent concentrations  $C/C_0$  (where  $C_0$  is the influent and  $C$  is the effluent virus concentration) were plotted as a function of pore volumes. Table 2 presents experimental mass balance ( $M_{BTC}$ ,  $M_s$ ,  $M_{III}$ ,  $M_{IV}$ , and  $M_{irr}$ ) information. This data is summarized in Fig. 2 which presents bar plots of the log removal (e.g.,  $-\log_{10}(M_{BTC})$ ) of viruses under different physicochemical conditions. The value of  $M_s = 26.2\%$  is in the worst-case transport scenario using clean quartz sand and SW at 5  $\text{m day}^{-1}$  (Fig. S3). In contrast, values of  $M_s$  in the aquifer sediment were always much higher (>92.3%) than this control



**Fig. 1.** The observed breakthrough concentrations of viruses (A)  $\Phi\text{X174}$ , (B) MS2, and (C) PRD1 at Phase I (injection of virus in SW or EQ-SW) and the observed effluent concentration in Phase II (injection of virus-free SW or EQ-SW). The Phases I and II were conducted at pore water velocity of 1 or 5  $\text{m day}^{-1}$  using either stormwater (SW) or stormwater equilibrated in the aquifer sediment (EQ-SW). Here, the markers are observed data and the solid black line is the fitted model.

**Table 2**

Experimental conditions and mass balance information from the column experiments. Here,  $M_{BTC}$ ,  $M_s = 100 - M_{BTC}$ ,  $M_{irr} = (100 - M_{BTC} - M_{III} - M_{IV})$ ,  $M_{III}$ , and  $M_{IV}$  denote the percentage of the injection MS2, PRD1, and  $\Phi$ X174 viruses that was recovered in the breakthrough curve, retained on the solid phase, irreversibly retained on the solid phase following the completion of Phases I–IV, and recovered with the injection of Milli-Q water in Phase III and Beef extract with pH 10 in Phase IV, respectively.

Bacteriophage	Velocity [m d <sup>-1</sup> ]	Solution Chemistry	Retention			Release	
			$M_{BTC}$	$M_s$	$M_{irr}$	$M_{III}$	$M_{IV}$
			[%]	[%]	[%]	[%]	[%]
$\Phi$ X174	1	SW	0.98	99.02	98.37	0.14	0.50
		SW EQ	0.00	100.00	99.94	0.01	0.04
	5	SW	7.70	92.30	91.86	0.06	0.38
		SW EQ	0.35	99.65	99.40	0.01	0.24
MS2	1	SW	0.04	99.96	99.94	0.01	0.01
		SW EQ	0.00	100.00	100.00	0.0001	0.001
	5	SW	1.32	98.68	98.67	0.002	0.006
		SW EQ	0.01	99.99	99.98	0.002	0.009
PRD1	1	SW	0.09	99.91	99.87	0.02	0.02
		SW EQ	0.00	100.00	100.00	0.0002	0.0004
	5	SW	2.51	97.49	97.46	0.01	0.02
		SW EQ	0.01	99.99	99.97	0.001	0.01

experiment. The maximum retention for all three viruses was observed in the experiment conducted under EQ-SW at 1 m day<sup>-1</sup> (>4.6 logs), whereas the least retention was observed under SW at 5 m day<sup>-1</sup> ( $M_s = 92.3\%$ ). A detailed discussion of the dependence of virus retention on water velocity, solution chemistry, and virus type is given below.

Fig. 2 indicates that all three viruses always had a higher removal in EQ-SW than SW at a given velocity. Table S1 indicates that the IS and concentration of Ca<sup>2+</sup> were much higher for EQ-SW than SW. This increase in IS and concentration of Ca<sup>2+</sup> lowered the magnitude of the zeta potential of the sediment and viruses (Table 1) in EQ-SW and thereby increased the adhesive interaction between the sediment and viruses. Consequently, one key consideration in the determination of virus removal is the effect of ionic strength and the presence of multivalent cations (Harvey and Ryan, 2004). The effects of multivalent cations on virus attachment can be attributed to a number of factors, including the larger ionic radius of Ca<sup>2+</sup> (1.61 Å) compared to Na<sup>+</sup> (1.02 Å) (Gutierrez et al., 2010), change in electrostatic interactions between virus and mineral surfaces (Carlson et al. (1968)), cation bridging (Bales et al., 1991; Chu et al., 2003; Pham et al., 2009), charge neutralization (Bales et al., 1991; Lukasik et al., 2000), screening of repulsive surface interaction energies between virus and grain surfaces (McCarthy and McKay, 2004), reduction of the net charge within the electrokinetic shear plane (Simoni et al., 2000), compression of the double-layer (Huysman and Verstraete, 1993), inner sphere complexation of the cations at the virus surfaces (Sadeghi et al., 2013), and the calcium binding to the carboxyl functional groups on the viral protein capsid (Harvey and Ryan, 2004).

Greater removal of viruses occurred at a lower (1 m day<sup>-1</sup>) than higher (5 m day<sup>-1</sup>) pore-water velocity under given solution chemistry conditions (Fig. 2). Previous studies have similarly demonstrated that the retention of colloids such as viruses, nanoparticles, and bacteria in porous media is velocity dependent (Hendry et al., 1999; Kim and Lee, 2014; Sasidharan et al., 2017a, 2014; Toloni et al., 2014; Torkzaban et al., 2007). Greater retention of viruses at low flow velocity can be explained by: (i) an increase in the virus residence time (Meinders et al., 1994; Xu et al., 2005); (ii) an increase in the virus adhesive interaction (Xu and Logan, 2006); and (iii) a decrease in the applied hydrodynamic torque that acts on the virus (Bradford et al., 2011; Sasidharan et al., 2017a). It should be mentioned that an increase in residence time increases

the time for virus removal. It also increases the probability that viruses can diffuse over shallow energy barriers on physically and chemically heterogeneous surfaces. Consequently, virus removal at a MAR site will be influenced by the flow field, with less removal near the injection well. Careful consideration of the flow field and separation distance between the injection and recovery wells is therefore necessary to achieve the maximum virus removal.

PRD1 is generally considered as the most conservative model for enteric viruses in subsurface viral transport studies (Harvey and Ryan, 2004; Schijven and Hassanizadeh, 2000; Stevenson et al., 2015). In contrast, Fig. 2 indicates that MS2 had the highest removal followed by PRD1 and then by  $\Phi$ X174 for a given physicochemical condition. Consequently,  $\Phi$ X174 was the most conservative model virus (had the least removal) in our sediment and stormwater. It is commonly believed that the removal and interactions of viruses with a solid surface are the results of their electrical charge and hydrophobicity (Shields and Farrah, 1987). Table 1 shows that the measured zeta potential values for all three viruses in each solution chemistry were in the same range, so differences in the net zeta potential of the viruses cannot explain these variations in retention. Both PRD1 and MS2 are known to be partially hydrophobic and  $\Phi$ X174 is hydrophilic in nature (Sasidharan et al., 2016; Schijven and Hassanizadeh, 2000). The presence of some forms of hydrophobic organic matter in the solution (Table S1) and the sediment surface may enhance the hydrophobic interaction between the partially hydrophobic viruses (PRD1 and MS2), which lead to their higher retention. In addition, nanoscale chemical and especially physical heterogeneity on the surfaces of colloids are known to strongly influence their adhesive interaction (Attinti et al., 2010; Bradford et al., 2017). The viral protein coat may contain weakly acidic and basic amino acid groups which act as localized positive and negative charges (Gerba, 1984) and it has a span of hydrophobic amino acids which will determine the hydrophobicity of viruses (Bendersky and Davis, 2011; Bradford and Torkzaban, 2012; Shen, et al., 2012d). The virus surface also contains nanoscale roughness features such as spikes (Huiskonen et al., 2007). Additional research is needed to fully characterize nanoscale variations in chemical heterogeneity and roughness features on the surfaces of our viruses and to assess the influence of these factors on virus retention. Considerable experimental and theoretical research would be needed to address this issue, and it is beyond the scope of this applied study.

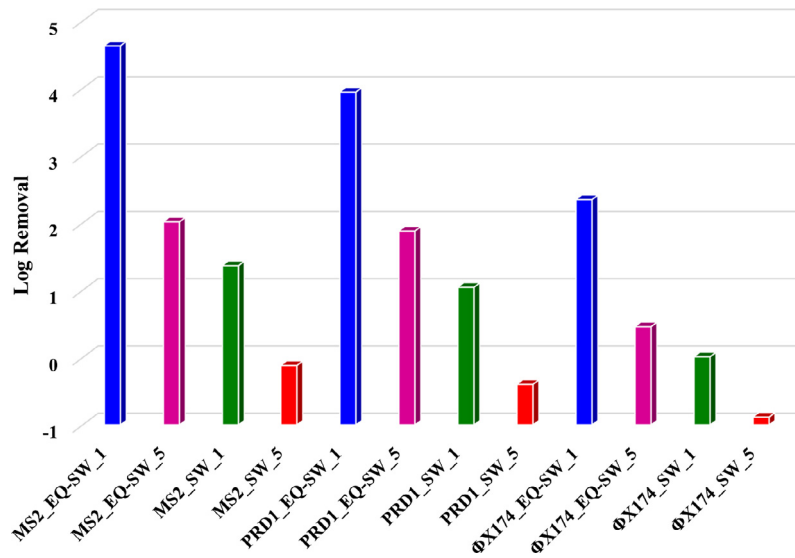


Fig. 2. Bar plots of the log removal (e.g.,  $-\log_{10}(M_{BTC})$ ) of viruses under the different physicochemical conditions shown Fig. 1.

### 3.5. Mathematical modeling of virus retention

Table 3 shows the liquid phase inactivation rate ( $\mu_l$ ) for viruses over the course of the transport and release experiments. The value of  $\mu_l$  was negligible ( $\sim 10^{-7} \text{ s}^{-1}$ ) in both SW or EQ-SW, and was consequently neglected in the mathematical model. A number of different model formulations were employed to describe the virus BTCs. However, low values of the Pearson correlation coefficient ( $R^2$ ) and nonunique parameter estimates were obtained in many instances. The best model description ( $R^2$  ranged from 67 to 98%) was obtained when considering attachment, detachment, blocking, and a solid phase sink term that accounted for irreversible attachment and inactivation. Straining, clogging, and wedging were not considered as important mechanisms for virus retention in this study, as the ratio of virus to a sediment grain diameter is far below the suggested threshold of 0.003 (Bradford and Bettahar, 2006). Table 4 presents fitted model ( $k_{att}$ ,  $k_{det}$ ,  $S_{max}/C_0$ , and  $\mu_s$ ) or calculated ( $\alpha$ ,  $\eta$ , and  $S_f$ ) parameters, and the  $R^2$  for the goodness of model fit.

Blocking decreases the attachment rate coefficient as available retention sites become filled, and has typically been neglected in most previous virus transport studies (Schijven and Hassanizadeh, 2000). However, Fig. 1 (BTC is plotted in log scale) and Fig. S4 (BTC is plotted in normal scale) indicate that blocking occurred for our viruses and sediment. In particular, BTCs were initially delayed (arriving after 1 PV), next they rapidly increased, and then slowly approached the influent virus concentration. Consistent with our observations, Xu et al. (2017) observed blocking

behavior for  $\Phi X174$  on a goethite-coated sand. Many others have reported on similar blocking behavior for nanoparticles (Li et al., 2008; Sasidharan et al., 2014; Virkutyte et al., 2014).

Fitted values of  $S_{max}$  in the blocking model were subsequently used to determine  $S_f$  (Table 4). Calculated values of  $S_f$  for the viruses were very small ( $1.30 \times 10^{-8}$ – $1.45 \times 10^{-4}$ ). Similarly, Xu et al. (2017) reported small values of  $S_f$  ( $0.3 \times 10^{-6}$ – $0.5 \times 10^{-6}$ ) for  $\Phi X174$  on a goethite-coated sand. In contrast, values of  $S_f$  for 50 and 100 nm latex nanoparticles were much higher on clean quartz sand (0.009–0.39) (Sasidharan et al., 2017b, 2014). Natural solid surfaces like sand grains always contain a wide distribution of physical (e.g., roughness) or chemical (e.g., metal oxides) heterogeneities (Bhattacharjee et al., 1998; Shen et al., 2012c). Previous studies have demonstrated that roughness height and fraction, and positive zeta potential and fraction, at a specific location on the collector (sand) surface can significantly reduce the magnitude of the energy barrier to attachment and the depth of the primary minimum for viruses (Bradford et al., 2017; Bradford and Torkzaban, 2013, 2015; Sasidharan et al., 2017b; Torkzaban and Bradford, 2016). In contrast to smooth latex nanoparticles, the virus exhibits chemical (e.g., lipid membrane and protein coat) (Meder et al., 2013) and physical heterogeneity (e.g., spikes and tail) (Huiskonen et al., 2007; Kazumori, 1981) on their surface. The combination of physical and chemical heterogeneities on both virus and collector surfaces apparently created a shallow primary minimum with negligible energy barrier to attachment and detachment for viruses, and thus, only a very small fraction of the collector surface contributed to  $S_f$ .

Previous research has shown that the relative effluent concentration increased and the relative retention decreased with increasing input concentration of colloids as a result of blocking (Leij et al., 2015; Wang et al., 2012). All three viruses were run in the same experiment and had a total input concentration of  $\sim 4.9 \times 10^8$  viruses in this study. Conversely, urban stormwater has a much lower enterovirus concentration of 6–170 virus/10 L (Strassler et al., 1999). Consequently, blocking and the subsequent exhaustion of retention sites may not be apparent in some natural systems with low input concentrations. However, recharging stormwater with a high concentration of some forms of organic matter or negatively charged phosphate ions (Table S1) can mask and reverse the charge of positive sites that are favorable for virus retention (Schijven and Hassanizadeh, 2000). This competition for

Table 3

The inactivation rate coefficients ( $\mu_l$ ) of MS2, PRD1, and  $\Phi X174$  in the stormwater (SW) and stormwater equilibrated with aquifer sediment (EQ-SW) for the experiment duration (4500 min = 75 h) and the Pearson correlation coefficient ( $R^2$ ) value calculated using linear regression analysis.

	Solution	$\mu_l$ [ $\text{s}^{-1}$ ]	$R^2$ [%]
$\Phi X174$	SW	$8.33 \times 10^{-7}$	99.63
	EQ-SW	$6.67 \times 10^{-7}$	90.98
MS2	SW	$5.00 \times 10^{-7}$	95.37
	EQ-SW	$3.33 \times 10^{-7}$	100
PRD1	SW	$1.17 \times 10^{-7}$	99.78
	EQ-SW	$1.00 \times 10^{-7}$	100



**Table 4**  
 A summary of experimental conditions (solution chemistry and water velocity), model parameters that were fitted to the virus BTCs (the attachment coefficient,  $k_{att}$ ; the detachment coefficient,  $k_{det}$ ; the maximum solid phase virus concentration,  $S_{max}$ ; and a sink term that accounts for irreversible attachment and solid phase inactivation,  $\mu_s$ ) and parameters  $S_f$ ,  $\alpha$ , and  $\eta$  which were calculated from  $S_{max}$ ,  $k_{att}$ , and the correlation equation of Messina et al. (2015), respectively. The table also includes the Pearson correlation coefficient ( $R^2$ ) for the goodness of model fit, and the standard error values (S.E.Coeff) on fitted parameters.

Virus	Solution	Velocity (m day <sup>-1</sup> )	R <sup>2</sup> (%)	$k_{att}$ (s <sup>-1</sup> )	S.E.Coeff – $k_{att}$	$k_{det}$ (s <sup>-1</sup> )	S.E.Coeff – $k_{det}$	$\mu_s$ (s <sup>-1</sup> )	S.E.Coeff – $\mu_s$	$S_{max}/C_0$ (cm <sup>3</sup> gr <sup>-1</sup> )	S.E.Coeff – $S_{max}/C_0$	$S_f$	$\alpha$	$\eta$
MS2	SW	1	94.28	$3.25 \times 10^{-3}$	$9.6 \times 10^{-3}$	$2.83 \times 10^{-6}$	$7.8 \times 10^{-5}$	$1.94 \times 10^{-5}$	$2.7 \times 10^{-4}$	2.11	0.35	$1.72 \times 10^{-6}$	0.162	0.211
	EQ-SW	1	67.37	$4.26 \times 10^{-3}$	$1.4 \times 10^{-3}$	$1.08 \times 10^{-5}$	$7.6 \times 10^{-4}$	$3.99 \times 10^{-5}$	$1.2 \times 10^{-3}$	$2.49 \times 10^1$	0.19	$2.01 \times 10^{-5}$	0.213	0.211
	SW	5	96.81	$5.80 \times 10^{-3}$	$6.1 \times 10^{-2}$	$2.65 \times 10^{-6}$	$1.5 \times 10^{-5}$	$2.32 \times 10^{-4}$	$3.2 \times 10^{-3}$	1.12	0.45	$9.12 \times 10^{-7}$	0.145	0.085
	EQ-SW	5	97.21	$1.43 \times 10^{-2}$	$1.7 \times 10^{-2}$	$1.63 \times 10^{-5}$	$1.6 \times 10^{-4}$	$7.14 \times 10^{-5}$	$7.6 \times 10^{-4}$	4.94	0.58	$3.99 \times 10^{-6}$	0.356	0.085
PRD1	SW	1	79.63	$4.86 \times 10^{-3}$	$5.8 \times 10^{-2}$	$4.81 \times 10^{-5}$	$3.2 \times 10^{-3}$	$5.93 \times 10^{-5}$	$8.9 \times 10^{-4}$	$7.77 \times 10^{-1}$	0.13	$2.74 \times 10^{-6}$	0.007	0.127
	EQ-SW	1	93.53	$4.00 \times 10^{-3}$	$2.4 \times 10^{-3}$	$3.93 \times 10^{-6}$	$6.2 \times 10^{-5}$	$2.11 \times 10^{-5}$	$1.9 \times 10^{-4}$	2.31	0.75	$1.45 \times 10^{-4}$	0.006	0.127
	SW	5	98.60	$1.52 \times 10^{-2}$	$6.3 \times 10^{-2}$	$3.95 \times 10^{-6}$	$8.4 \times 10^{-5}$	$1.87 \times 10^{-4}$	$1.7 \times 10^{-3}$	$7.32 \times 10^{-1}$	0.007	$2.58 \times 10^{-6}$	0.011	0.048
	EQ-SW	5	96.51	$1.21 \times 10^{-2}$	$1.2 \times 10^{-2}$	$2.43 \times 10^{-5}$	$3.2 \times 10^{-4}$	$4.89 \times 10^{-5}$	$8.9 \times 10^{-4}$	$1.53 \times 10^1$	0.007	$9.62 \times 10^{-5}$	0.01	0.048
ΦX174	SW	1	96.17	$1.18 \times 10^{-3}$	$2.4 \times 10^{-3}$	$8.09 \times 10^{-6}$	$2.2 \times 10^{-4}$	$1.92 \times 10^{-5}$	$2.5 \times 10^{-4}$	2.24	0.22	$4.01 \times 10^{-8}$	0.001	0.203
	EQ-SW	1	96.84	$2.86 \times 10^{-3}$	$6.1 \times 10^{-3}$	$4.14 \times 10^{-6}$	$8.9 \times 10^{-5}$	$4.85 \times 10^{-5}$	$3.4 \times 10^{-4}$	2.17	0.31	$5.54 \times 10^{-8}$	0.002	0.203
	SW	5	93.55	$1.08 \times 10^{-2}$	$5.8 \times 10^{-3}$	$1.25 \times 10^{-4}$	$2.3 \times 10^{-3}$	$1.86 \times 10^{-4}$	$1.4 \times 10^{-4}$	$7.27 \times 10^{-1}$	0.003	$1.30 \times 10^{-8}$	0.005	0.081
	EQ-SW	5	97.11	$3.06 \times 10^{-2}$	$1.9 \times 10^{-1}$	$1.08 \times 10^{-3}$	$1.7 \times 10^{-2}$	$3.11 \times 10^{-4}$	$8.6 \times 10^{-4}$	1.06	0.12	$2.71 \times 10^{-8}$	0.013	0.081

the same retention sites and  $S_{max}$  could potentially cause blocking to be enhanced.

Colloid filtration theory (CFT) has been developed to predict the value of  $k_{att}$  under different physicochemical conditions (Yao et al., 1971). CFT considers that  $k_{att}$  is proportional to the product of  $\alpha$  and  $\eta$  that account for adhesion and mass transfer, respectively. Consistent with CFT predictions, the value of  $k_{att}$  (Table 4) increased with increasing fluid velocity at a given solution chemistry. Similar to Fig. 2, values of  $k_{att}$  for MS2 and ΦX174 were always higher for EQ-SW than SW at a given water velocity. However, values of  $k_{att}$  for PRD1 were nearly the same for EQ-SW and SW. The observed dependence of  $k_{att}$  on the solution chemistry and the virus type was likely complicated by blocking which reduced the apparent value of  $k_{att}$ . Both  $\alpha$  and  $S_f$  are strong functions of the interaction energy between the virus and sediment (Shen et al., 2010; Tufenkji and Elimelech, 2004), so it may not be possible to separately quantify these effects especially when  $S_f$  is very low.

The virus BTCs shown in Fig. 1 exhibited low levels of concentration tailing when the columns were eluted with virus-free SW or EQ-SW. The model did not provide an accurate description of this tailing region when  $k_{det}$  was considered and  $\mu_s$  was neglected. Conversely, a good fit was obtained to both the retention and the tailing portion of the BTCs when  $\mu_s$  and  $k_{det}$  (Table 4) were included in the model. Little is known about solid phase inactivation of viruses other than the need for a strong binding force which may disintegrate the protein structure and thereby inhibit the ability of viruses to infect their host (Harvey and Ryan, 2004). Only a few approaches have been developed to separately quantify the rates of virus attachment and solid phase inactivation (Grant et al., 1993; Harvey and Ryan, 2004) such as radiolabeling polio virus capsid and RNA to track the fate of these components (Murray and Parks, 1980). These approaches are difficult, time consuming, and/or require specialized equipment. In this research, these technical challenges were partially overcome through numerical modeling that allows the determination of attachment ( $k_{att}$ ) and detachment ( $k_{det}$ ) rates, and a sink term to account for the combined rate of irreversible attachment and solid phase inactivation ( $\mu_s$ ).

Guidelines for MAR have only considered inactivation of viruses in the liquid phase, but have neglected irreversible attachment and solid phase inactivation of viruses (Abu-Ashour et al. (1994); Dillon et al., 2008). Tables 3 and 4 indicate that  $k_{att} > \mu_s > k_{det} > \mu_i$ , and  $k_{att}$  is 3–4 orders of magnitude higher than  $\mu_i$ . Consequently, guidelines for MAR that only consider liquid phase inactivation will be overly conservative for the considered experimental conditions. Recognizing the removal of viruses via irreversible attachment and/or solid phase inactivation during aquifer storage would help to eliminate some of the expensive post-treatment for recovered MAR water. However, we acknowledge that the virus removal via irreversible attachment and/or solid phase inactivation is highly site specific. Therefore, conducting site specific microbial risk assessment and subsequent development of appropriate MAR design that considers adequate residence time and travel distance based on the horizontal and vertical flow field, soil heterogeneity, and chemical characterization is necessary to achieve the maximum virus removal.

### 3.6. Release of virus

Fig. 3 shows plots of ΦX174, MS2, and PRD1 concentrations in the column effluent during Phases III and IV when Milli-Q water and 3% beef extract with pH = 10 were injected into the column, respectively. Small pulses of released viruses were observed during both phases, with a small peak followed by long concentration tailing. Table 2 summarizes mass balance information from Phases III

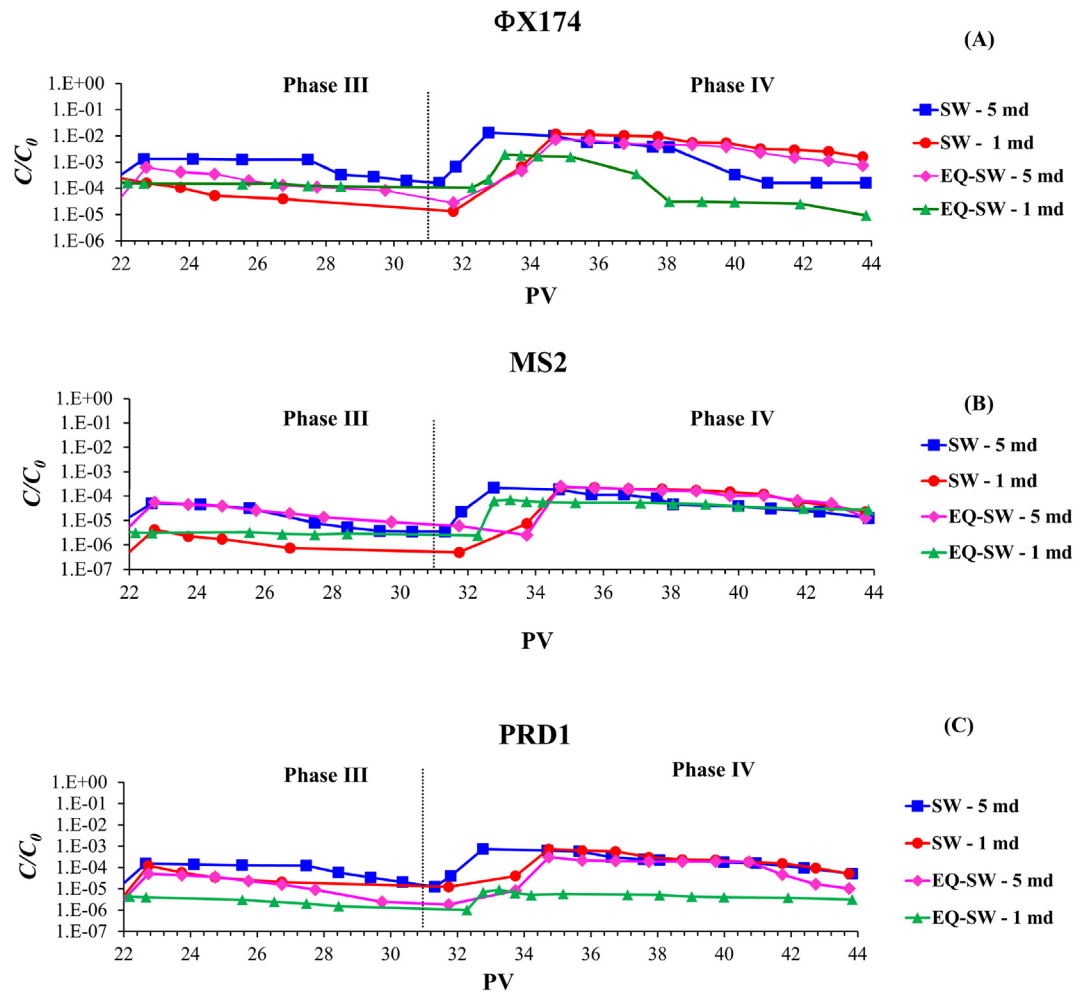


Fig. 3. Observed effluent concentrations of viruses (A)  $\Phi$ X174, (B) MS2, and (C) PRD1 in Phase III (Milli-Q water) and IV (3% beef extract with pH 10) at a pore-water velocity of 1 and 5 m day<sup>-1</sup>.

and IV. Only a very small percentage (<0.5%) of the initially attached viruses were released even under the worst-case scenario of beef extract with pH = 10.

Table 2 shows the mass of viruses on the sediment surface that was not recovered after the release Phases III and IV ( $M_{irr} = 100 - M_{BTC} - M_{III} - M_{IV}$ ). The negligible release of viruses and the presence of large  $M_{irr} > 91\%$  indicate that the viruses were irreversibly attached or inactivated on the solid surface. Values of  $M_{irr}$  followed similar trends to  $M_s$  with physicochemical conditions. In particular,  $M_{irr}$  was higher for EQ-SW than SW for a given velocity,  $M_{irr}$  was higher at lower (1 m day<sup>-1</sup>) than higher (5 m day<sup>-1</sup>) pore-water velocity at a given chemistry, and  $M_{irr}$  was highest for MS2 followed by PRD1 and then  $\Phi$ X174. The presence of a large  $M_s$  and  $M_{irr}$  value for the EQ-SW experiment implies that viruses will irreversibly attach and/or inactivate on the solid before the water reaches the recovery well during an ASTR operation, provided that a sufficient residence time is achieved. An injection event will only release a very tiny fraction of the viruses which may mobilize and re-attach to the surface as the water equilibrates with the sandy limestone aquifer sediment and the IS and the divalent cation concentration increases. Therefore, during ASR operations the chance of releasing the attached viruses during the reverse flow is negligible. However, we assume that the ASTR operation has added advantage of long residence time and travel distance compared to ASR and, therefore, ASTR would be the best option when removal of viruses via attachment/solid phase inactivation is considered.

#### 4. Conclusions and future directions

This research was conducted to better understand and quantify the relative importance of various virus removal processes during MAR in a sandy limestone aquifer. The aquifer sediment was found to remove >92.3% of bacteriophage MS2, PRD1, and  $\Phi$ X174 from SW and EQ-SW when the pore-water velocity was 1 or 5 m day<sup>-1</sup>. However, much greater virus removal (4.6 logs) occurred at the lower pore-water velocity and in EQ-SW because of an increase in residence time, IS, and Ca<sup>2+</sup> concentration. Bacteriophage  $\Phi$ X174 showed less removal than either PRD1 or especially MS2, and was, therefore, the most conservative model virus for this site. Negligible virus detachment (<0.64%) occurred when columns were flushed with Milli-Q water or beef extract at pH = 10, and this indicates that viruses were irreversibly attached or quickly inactivated on sediment surfaces.

Virus BTCs were successfully simulated using the advection-dispersion equation with terms for attachment, detachment, Langmuirian blocking, and solid phase removal by irreversible attachment and/or surface inactivation. Values of  $k_{att} > \mu_s > k_{det} > \mu_i$ , and  $k_{att}$  was 3–4 orders of magnitude greater than  $\mu_i$ . The value of  $k_{det}$  was always small under steady-state physicochemical conditions. The remaining viruses on the solid phase were either irreversibly retained or inactivated. This process was modeled using the  $\mu_s$  term. Blocking has commonly been neglected in previous virus transport studies. In contrast, our results clearly demonstrated blocking behavior. The relative importance of blocking is expected

to decrease in a natural setting with lower input concentrations and/or higher  $S_{max}$  (Leij et al., 2015). Furthermore, the value of  $S_f$  was found to be very small for viruses. This result was attributed to the presence of a shallow primary minimum due to roughness on both the virus and sediment surfaces. Additional research is needed to further assess the potential influence of blocking on virus transport at MAR sites.

MAR has only been considered as a storage option in a water recycling train. Consequently, all previous field scale studies for MAR risk assessments have only considered virus removal by liquid phase inactivation. Current MAR guidelines do not acknowledge the removal of viruses by attachment and solid phase inactivation. This research clearly demonstrated that viruses were irreversibly attached or inactivated on the sediment surface when given enough residence time and that  $k_{att} \gg \mu_1$ . Natural treatment of injected water during infiltration through the unsaturated zone is widely recognized. However, the mechanisms for natural treatment of virus within the saturated zone is not well understood and the selection of a sub-section of aquifer material with lower carbonate content/higher quartz content represents an environment that is less favorable for virus removal (worst-case scenario). Furthermore, greater virus removal is expected as water moves away from an injection well due to a decrease in water velocity and an increase in IS and  $Ca^{2+}$  ions. However, field-scale experiments under actual artificial recharge conditions, using the same bacteriophages employed in this study, showed that the safe setback distance depended on site-specific physicochemical conditions. For example, the setback distance was estimated to be just a few meters (~10 m) in a sandy aquifer in Los Angeles County (Anders and Chrysikopoulos, 2005), whereas the reported setback distance was  $8000 \pm 4800$  m in a fractured aquifer in Southern Italy (Masciopinto et al., 2008). Future MAR guidelines and microbial risk assessment may therefore need to consider site specific removal of viruses by irreversible attachment and solid phase inactivation during the storage period. This could help to eliminate some of the expensive post-treatment to achieve desired water quality.

## Notes

The authors declare no competing financial interest.

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## Appendix A. Supplementary data

Supplementary data associated with this article can be found, in the online version, at <https://doi.org/10.1016/j.jhydrol.2017.10.062>.

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