

Quantifying the log reduction of pathogenic microorganisms by constructed wetlands: A literature review



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WWTPs

Efficient pathogen removal 🗸

Reliable

Environmental degradation

High energy consumption

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• Water can contain various microbial contaminants and can cause detrimental health effects if a specific dose is consumed







• Classification of human pathogens in five groups





- Classification of human pathogens in five groups
- Enumeration of human pathogens can be an expensive and time-consuming process
- Numerous methods have been developed that first quantify groups of indicator organisms that are easy and inexpensive to monitor, and then correlate them with their respective index pathogenic organisms



- Classification of human pathogens in five groups
- Enumeration of human pathogens can be an expensive and timeconsuming process
- Numerous methods have been developed that first quantify groups of indicator organisms that are easy and inexpensive to monitor, and then correlate them with their respective index pathogenic organisms
- The most common indicators are:
 - Coliforms (total and fecal)
 - o Escherichia Coli
 - Fecal Streptococcus



<u>Definition</u>: A man-made system designed to replicate the operation of a natural wetland (Nuttall et al., 1998).

- Constructed wetlands are alternative engineered systems that are based on the use of emerging plants for the purification of wastewater
- Series of physical, chemical and biological mechanisms for the removal of pathogens





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Systematic literature review

- Preferred Reporting Items for Systematic Reviews and Meta-Analyses (PRISMA) guidelines
- Preferred search engines "Scopus and "PubMed"
- After two series of screening, a total of 48 case studies qualified for both qualitative and quantitative analyses





- Hydraulic Loading Rate (HLR)
- Hydraulic Retention Time (HRT)
- Porosity (n) of the media grains
- A detailed description of CWs

- Type of influent wastewater
- Method of enumeration
- Physicochemical parameters like
 - Biological Oxygen Demand (BOD)
 - Chemical Oxygen Demand (COD)
 - Total Suspended Solids (TSS)

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Spatial distribution of case studies:

- Shows great variability
- Has different temperature/climatic zones
- Strengthens the validity of the results
- A relationship between temperature/climatic zones and log removal was examined



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Classification of pathogens:

- Three main categories
- Largest differences in removal between categories, and rather smaller within categories

Escherichia coli Total/fecal coliform Fecal streptococci/ enterococci Intestinal enterococciColiphages F-RNA specific phages Bacteriophages infecting GB124 MS2 bacteriophagesClostridia Clostridium perfringens sporesPseudomonas aeruginosa Campylobacter Salmonella AeromonasAdenovirus Aichi virus 1 BG/JC polyomavirus Enteric virus Norovirus GllGiardia/giardia lamblia Cryptosporidium		Gram-negative bacteria	Viruses	Protozoan parasites
Campylobacter Aichi virus 1 lamblia Salmonella BG/JC polyomavirus Cryptosporidium Aeromonas Norovirus Gll	Indicator	Total/fecal coliform Fecal streptococci/ enterococci	F-RNA specific phages Bacteriophages infecting GB124	Clostridium
	Pathogen	Campylobacter Salmonella	Aichi virus 1 BG/JC polyomavirus Enteric virus	lamblia



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Preliminary exploration between parameters:

The goal of this preliminary exploration was to identify potential patterns between parameters by plotting them to each other regarding either different types of wetland or different pathogen categories. Regarding parameters, the *Cin, Cout*, HRT, and HLR were assessed since those where the only parameters that were consistently reported in the literature review.



A sample of created dataset

Title	Type_of_CW	HLR H	HRT E	BOD5_remc	COD_removal	TSS_total	Medium	Dimensions_ol	 Vatertype	Group_Genus	Group	Type_of_	s Method_of_quar	n Conce	Concer Units	Concer	Concel	Inits	sign	Log_removal
Presence of		1.08	0.83	_	92.66247379					Escherichia			TBX medium	=	2.44 CHU/100 mL	=		:FU/100 mL	_	1.29
Presence of	VFCW	1.08	0.83	98.034996	92.66247379	0.32954	-			Faecal Coliforr	_		membrane filtratio	o =	2.86 CHU/100 mL	=	1.61 0	:FU/100 mL	. =	1.25
Presence of		1.08	0.83		92.66247379		-	L= 29 m,W= 40	SST effluent	Intestinal Enter	Gram_negat	indicator	SB agar	=	1.79 CHU/100 mL	=	0.48 0	FU/100 mL	. =	1.31
Presence of	VFCW	1.08	0.83	98.034996	92.66247379	0.32954	gravel	L= 29 m,W= 40	SST effluent	Somatic coliph	Viruses	indicator	triplicates by enu	ır =	4.61 CHU/100 mL	=	4.01 0	FU/100 mL	. =	0.6
Presence of	VFCW	1.08	0.83	98.034996	92.66247379	0.32954	gravel	L= 29 m,W= 40	SST effluent	F-RNA specific	Viruses	indicator	triplicates by enu	ır =	3.2 CHU/100 mL	=	2.2 0	:FU/100 mL	. =	1
Presence of	VFCW	1.08	0.83	98.034996	92.66247379	0.32954	gravel	L= 29 m,W= 40	SST effluent	phages infecti	r Viruses	indicator	triplicates by enu	ır =	2.57 CHU/100 mL	=	2 0	:FU/100 mL	. =	0.57
Comparative	VFCW	0.0028	4		91.82222222	0.01307	sand+ gravel	25-cm diamete	e raw wastewa	Escherichia	Gram_negat	indicator	membrane filtratio	o =	6.47 CFU/mL	=	2 0	FU/mL	=	4.47
Comparative	VFCW	0.0028	4		91.82222222	0.01307	sand+ gravel	25-om diamete	e raw wastewa	Total Coliform	Gram_negat	indicator	membrane filtratio	o =	7.41 CFU/mL	=	2.4 0	FU/mL	=	5.01
Comparative	VFCW	0.0028	4		91.82222222	0.01307	sand+ gravel	25-cm diamete	e raw wastewa	Faecal Coliforr	i Gram_negat	indicator	membrane filtratio	o =	6.08 CFU/mL	=	0 0	FU/mL	=	6.08
Comparative	VFCW	0.0028	4		89.22222222	0.01103	sand+ gravel	25-cm diamete	e raw wastewa	Escherichia	Gram_negat	indicator	membrane filtratio	o =	6.47 CFU/mL	=	3.04 0	FU/mL	=	3.43
Comparative	VFCW	0.0028	4		89.22222222	0.01103	sand+ gravel	25-cm diamete	e raw wastewa	Total Coliform	Gram_negat	indicator	membrane filtratio	o =	7.41 CFU/mL	=	3.6 0	FU/mL	=	3.81
Comparative	VFCW	0.0028	4		89.22222222	0.01103	sand+ gravel	25-om diamete	e raw wastewa	Faecal Coliforn	i Gram_negat	indicator	membrane filtratio	o =	6.08 CFU/mL	=	2.52 0	FU/mL	=	3.56
Comparative	VFCW	0.0028	4		91.8444444	0.01865	sand+marble o	25-om diamete	e raw wastewa	Escherichia	Gram_negat	indicator	membrane filtratio	o =	6.47 CFU/mL	=	2.54 0	FU/mL	=	3.93
Comparative	VFCW	0.0028	4		91.8444444	0.01865	sand+marble o	25-om diamete	e raw wastewa	Total Coliform	Gram_negat	indicator	membrane filtratio	o =	7.41 CFU/mL	=	2.59 0	FU/mL	=	4.82
Comparative	VFCW	0.0028	4		91.8444444	0.01865	sand+marble o	25-om diamete	e raw wastewa	Faecal Coliforn	Gram_negat	indicator	membrane filtratio	o =	6.08 CFU/mL	=	0 0	FU/mL	=	6.08
Comparative	VFCW	0.0028	4		92.59259259	0.01861	sand+marble o	25-cm diamete	e raw wastewa	Escherichia	Gram_negat	indicator	membrane filtratio	o =	6.47 CFU/mL	=	2.66 0	FU/mL	=	3.81
Comparative	VFCW	0.0028	4		92.59259259	0.01861	sand+marble o	25-om diamete	e raw wastewa	Total Coliform	Gram_negat	indicator	membrane filtratio	o =	7.41 CFU/mL	=	3.18 0	FU/mL	=	4.23
Comparative	VFCW	0.0028	4		92.59259259	0.01861	sand+marble o	25-om diamete	e raw wastewa	Faecal Coliforn	Gram_negat	indicator	membrane filtratio	o =	6.08 CFU/mL	=	1.26 0	FU/mL	=	4.82
Halophytes a		0.095	3.5		78.125		·	L= 1.20 m, V=0	1 6		Gram_negat	indicator	IDEXX Quanti-Tr	7	6.2 MPN/100mL	=		1PN/100mL	-	1.2
Halophytes a		0.095	3.5		78.125			L= 1.20 m, W=0			Gram_negat		IDEXX Quanti-Tr	a =	4.2 MPN/100mL	=	2.1 M	1PN/100mL	. =	2.1
Halophytes a	VFCW	0.095	3.5		79.01785714		2 layers of grav	L= 1.20 m, W=0), primary treat	Total Coliform	Gram_negat	indicator	IDEXX Quanti-Tr	a =	6.2 MPN/100mL	=	5.5 M	1PN/100mL	. =	0.7
Halophytes a	VFCW	0.095	3.5		79.01785714			L= 1.20 m, W=0), primary treat	Escherichia	Gram_negat	indicator	IDEXX Quanti-Tr	a =	4.2 MPN/100mL	=	2.4 M	1PN/100mL	. =	1.8
Emerging org	VFCW	0.044	6.34	91.2	82.94573643		1 layer of sand-		combined se	e Escherichia	Gram_negat	indicator	Chromogenic Me	er =	6.6 CFU/100 mL	=	5.95 0	CFU/100 mL	. =	0.65
Emerging org	FWS	0.06	2.3	94.4	88.75968992		siliceous grave		combined se	Escherichia	Gram_negat	indicator	Chromogenic Me	er =	5.95 CFU/100 mL	=	3.47 0	CFU/100 mL	. =	2.48
Emerging org	FWS	0.02	5.1	95.2	81.78294574	0.05263	siliceous grave	el	combined se	Escherichia	Gram_negat	indicator	Chromogenic Me	er =	3.47 CFU/100 mL	<	1.6 0	:FU/100 mL	. <	1.87
Integrated tre	VFCW	0.044	6.34	97.455471	91.46567718	0.02787	1 layer of sand-	D=0.8	combined se	Escherichia	Gram_negat	indicator	Chromogenic Me	er =	6.6 CFU/100 mL	=	5.18 0	:FU/100 mL	. =	1.424
Integrated tre	FWS	0.06	2.3	98.727735	94.43413729	0.02787	gravel		combined se	Escherichia	Gram_negat	indicator	Chromogenic Me	er =	5.176 CFU/100 mL	=	3.55 0	:FU/100 mL	. =	1.626
Integrated tre	FWS	0.02	5.1	98.21883	90.72356215	97.9094	gravel		combined se	Escherichia	Gram_negat	indicator	Chromogenic Me	er =	3.55 CFU/100 mL	=	4.5 0	:FU/100 mL	_ =	-0.95



Log removal and climatic zones

Although there seems to be a trend where higher removal values can be found in hot climatic zones, and rather low removal values can be found in cold and cool climatic zones, a clear correlation was not found since there is a significant variation within different climatic zones. Additionally, the log removal values are

averaged per country and in many cases there is only one case study per country







- Potential relationship between log removal and HLR and/or HRT
- Observed variation can be attributed to:
 - Different applications (full-scale vs pilot-scale) plotted together
 - Different technical features/experimental conditions of each study leads to different removal efficiency





Removal of pathogen categories

Comparison of pathogen categories per CW type:

- Gram-negative bacteria show great variability in their removal (0.01-6.08 log) compared to viruses and protozoan parasites categories where the ranges are between 0.02-3.62, and 0.18-3.63 log respectively.
- Average removal values ranging between 1 and 2 log for the three categories in all of the CW types except the viruses' category in VFCW where the average log removal is less than 1.







Removal of pathogen categories

Based on the CDF:

- 75 % of both protozoan parasites and viruses categories have a removal lower or equal to 2 log while their peak lies just before 4 log.
- Gram-negative bacteria category exhibits a smooth curve along the distribution where 60% of the observed data points have a removal lower or equal to 2 log while the rest 40% lies between 2 and 6 log.



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- Access to overall performance of CWs.
- Spatial variability of case studies.
- Benchmark for any new relevant research.

Pathogen categories

- Gram-negative Bacteria category has the highest log removal in all 3 types of CW.
- Difficult to draw conclusions due to small number of data points.

Constructed Wetlands

- Credible choice for WW polishing.
- The CDFs of different types of CWs and different pathogen categories simply provided an initial mapping of the situation (in terms of performance and removal capacity) according to the literature review and can be used as a reference point.
- Various patterns were observed between hydraulic characteristics and influent/effluent concentrations which gives room for further investigation.
- A potential meta-analysis of this database using statistical analysis can provide additional and insightful information on the significance of these parameters on pathogen removal.



Overall, the final outcome does provide an efficient approach to the scientific community by taking a step closer to a better understanding of these "black boxes" and pointing out where future research needs to focus, in order to fine-tune and quantify the factors that influence the performance of constructed wetlands.

