

Report Comments			
Collection Date	Location	Test	Comment
6/1/2026	RAW	BOD_C	Sample volume <100% and <2 mg/L depletion.
6/2/2026	RAW	BOD_C	
6/5/2026	OUTFALL30	BOD_C	Sample dilutions show more than 30% between highest and lowest value
6/8/2026	OUTFALL30	BOD_C	Residual Dissolved Oxygen <1.5 mg/L for all dilutions. (over-depleted)
6/10/2026	OUTFALL30	BOD	Sample dilutions show more than 30% between highest and lowest value
6/11/2026	OUTFALL30	BOD_C	Sample volume <100% and <2 mg/L depletion.
6/11/2026	OUTFALL30	BOD	Sample volume <100% and <2 mg/L depletion.
6/11/2026	OUTFALL10	BOD	Sample dilutions show more than 30% between highest and lowest value
6/11/2026	OUTFALL30		Result is confirmed by the Lab that BOD results is <100.
6/15/2026	OUTFALL10	BOD	Sample dilutions show more than 30% between highest and lowest value
6/18/2026	OUTFALL10	BOD	Sample dilutions show more than 30% between highest and lowest value
6/20/2026	OUTFALL10	BOD	Sample dilutions show more than 30% between highest and lowest value
6/20/2026	OUTFALL10	BOD_C	Sample dilutions show more than 30% between highest and lowest value
6/20/2026	OUTFALL20	BOD_C	BOD/CBOD outside of the method uncertainty.
6/20/2026	OUTFALL20	BOD	BOD/CBOD outside of the method uncertainty.
6/23/2026	RAW	BOD_C	Sample dilutions show more than 30% between highest and lowest value
6/23/2026	OUTFALL30	BOD_C	Result calculated with seed correction factor based on highest depletion, as no seed control achieved 2.0 mg/L depletion.
6/23/2026	RAW	BOD	Result calculated with seed correction factor based on highest depletion, as no seed control achieved 2.0 mg/L depletion.
6/23/2026	RAW	BOD_C	Result calculated with seed correction factor based on highest depletion, as no seed control achieved 2.0 mg/L depletion.
6/23/2026	OUTFALL10	BOD	Result calculated with seed correction factor based on highest depletion, as no seed control achieved 2.0 mg/L depletion.
6/23/2026	OUTFALL30	BOD	Result calculated with seed correction factor based on highest depletion, as no seed control achieved 2.0 mg/L depletion.
6/23/2026	OUTFALL10	BOD_C	Result calculated with seed correction factor based on highest depletion, as no seed control achieved 2.0 mg/L depletion.
6/23/2026	RAW	BOD	Sample dilutions show more than 30% between highest and lowest value
6/24/2026	RAW	BOD_C	Results calculated with seed correction factor based on the highest depletion as no seed control achieved 2.0mg/L depletion.
6/24/2026	OUTFALL10	BOD_C	Results calculated with seed correction factor based on the highest depletion as no seed control achieved 2.0mg/L depletion.
6/24/2026	OUTFALL30	BOD_C	Results calculated with seed correction factor based on the highest depletion as no seed control achieved 2.0mg/L depletion.
6/24/2026	OUTFALL30	BOD	Results calculated with seed correction factor based on the highest depletion as no seed control achieved 2.0mg/L depletion.
6/24/2026	OUTFALL10	BOD	Results calculated with seed correction factor based on the highest depletion as no seed control achieved 2.0mg/L depletion.
6/24/2026	RAW	BOD	Results calculated with seed correction factor based on the highest depletion as no seed control achieved 2.0mg/L depletion.
6/24/2026	RAW	BOD_C	Sample dilutions show more than 30% between highest and lowest value
6/25/2026	OUTFALL30	BOD_C	Result calculated with seed correction factor based on highest depletion, as no seed control achieved 2.0 mg/L depletion.
6/25/2026	RAW		Insufficient sample volume to perform analysis of COD, OP04, D, CL, CL.
6/25/2026	OUTFALL30	BOD	Result calculated with seed correction factor based on highest depletion, as no seed control achieved 2.0 mg/L depletion.

Averages and geometric means may be calculated using only the numeric component of censored values

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Paul Jacquier **MBA, BMSc.**
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Report finalized by Ayesha Prasad Jul 28, 2026 09:34