2021 Outfall Benthic Monitoring Results

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2021 Outfall Benthic Monitoring Results

Submitted to

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EXECUTIVE SUMMARY

Background. The Massachusetts Water Resources Authority (MWRA) operates the Deer Island Wastewater Treatment Plant, where sewage from more than 40 communities in Greater Boston undergoes primary and secondary treatment. Following treatment, the effluent is discharged to Massachusetts Bay more than 5 miles (8 km) from the nearest shoreline. In 2000, MWRA moved the Deer Island discharge from Boston Harbor to Massachusetts Bay as part of the Boston Harbor Project to alleviate harbor pollution. MWRA has monitored in the bay since 1992 to gather data used to evaluate the potential effects of the discharge.

MWRA is required to monitor the benthos (seafloor community) in Massachusetts Bay. MWRA's discharge permit requires benthic monitoring to detect any effects of the effluent on the ocean environment. The monitoring focuses on three main concerns: (1) eutrophication (excess organic material and nutrients) and related low levels of dissolved oxygen; (2) deposits of toxic contaminants; and (3) smothering of animals by sewage effluent solids, or other changes to benthic communities. MWRA measures levels of total organic carbon to assess whether effluent discharge has resulted in organic enrichment; traces areas of possible contamination based on levels of a sewage indicator bacterium, *Clostridium perfringens*; and reports on benthic animal abundance and diversity. Since animal communities vary naturally across different benthic habitats, monitoring is designed to detect changes in habitat that can be associated with sediment grain size. Within the context of natural variation, potential outfall impacts can then be assessed.

Benthic habitat quality has remained high near the outfall. The discharge is located in an area dominated by hydrodynamic and physical factors, including tidal and storm currents, turbulence, and sediment transport (Butman et al. 2008). These physical factors, combined with the high quality of the effluent discharged into the Bay (Taylor 2010, Werme et al. 2021), are the principal reasons that benthic habitat quality has remained high near the outfall.

MWRA's discharge permit includes a Contingency Plan which sets out caution and warning threshold levels of key indicators including the diversity of benthic animals. There were no exceedances of these thresholds for any infaunal (soft-sediment) community in 2021. Data on these macrobenthic (larger than 0.3 mm) animals continue to suggest that the communities near the outfall remain healthy. In addition, years of monitoring have shown that solids from wastewater discharge do not reach levels that disturb or smother animals near the outfall.

2021 results show that the effluent discharge has had little influence on sediment conditions. This is consistent with prior results (Rutecki et al. 2022, Nestler et al. 2020, Maciolek et al. 2008). Concentrations of the sewage tracer, *Clostridium perfringens* during 2021 were relatively low, with the highest values at sites closest to the discharge. This spatial pattern has remained consistent since outfall relocation to the bay in 2000 (e.g., Rutecki et al. 2022, Maciolek et al. 2007, 2008). These results provide evidence that effluent solids have settled at sites within about 2 km (1.25 miles) of the outfall, but neither sediment grain size nor levels of total organic carbon have been affected.

1 INTRODUCTION

Under its Ambient Monitoring Plan (MWRA 1991, 1997, 2001, 2004, 2010, 2021) the MWRA has collected extensive information over a nine-year baseline period (1992–2000) and a twenty-one-year period (2001–2021) after the wastewater discharge was moved to Massachusetts Bay. These studies included surveys of sediments and soft-bottom communities using traditional grab sampling and sediment profile imaging (SPI; 1992–2019) as well as surveys of hard-bottom communities using a remotely operated vehicle (ROV). Data collected by this program allow for a more complete understanding of the bay system and provide a basis to explain any changes in benthic conditions and the question of whether MWRA's discharge has contributed to any such changes.

Benthic monitoring during 2021 was conducted following the current Ambient Monitoring Plan (MWRA 2021) which is required under MWRA's effluent discharge permit for the Deer Island Treatment Plant. Under this current plan, annual monitoring includes soft-bottom sampling for sediment conditions and infauna at 14 nearfield and farfield stations. Every third year, hard-bottom surveys are conducted at 23 nearfield stations. The most recent hard-bottom survey was conducted in 2020. Monitoring results for 2020 indicated that hard-bottom benthic communities near the outfall have not changed substantially during the post-diversion period as compared to the baseline period (Rutecki et al. 2022). Some modest changes in hard-bottom communities (e.g., coralline algae, upright algae cover, and sponge abundance) have been observed; nonetheless, factors driving these changes are unclear. Since declines in upright algae started in the late 1990s, it is unlikely that the decrease was attributable to diversion of the outfall (Rutecki et al. 2022). Under the current monitoring plan (which went into effect in July 2020), the SPI study in Massachusetts Bay and the sediment contaminant evaluations have been discontinued; these studies had answered their monitoring questions fully.

This report summarizes key findings from the 2021 benthic surveys, with a focus on the most noteworthy observations relevant to understanding the potential effects of the discharge on the offshore benthic environment. Results of 2021 benthic monitoring were presented at MWRA's Annual Technical Workshop on April 11, 2022. This report builds on the presentations and discussions at that meeting.

2 METHODS

Methods used to collect, analyze, and evaluate all sample types remain largely consistent with those reported for previous monitoring years (Rutecki et al. 2022, Maciolek et al. 2008). Detailed descriptions of the methods are contained in the Quality Assurance Project Plan (QAPP) for Benthic Monitoring 2020–2023 (Rutecki et al. 2020). A brief overview of methods, focused on information that is not included in the QAPP, is provided in Sections 2.1 to 2.3.

2.1 FIELD METHODS

Sediment and infauna sampling was conducted at 14 stations on August 4, 2021 (Figure 2-1). To aid in analyses of potential spatial patterns reported herein, these stations are grouped, based on distance from the discharge, into four "monitoring areas" within Massachusetts Bay[1]:

- Nearfield stations NF13, NF14, NF17, and NF24, located in close proximity (less than 2 km) to the offshore outfall
- Nearfield stations NF04, NF10, NF12, NF20, NF21, and NF22, located in Massachusetts Bay but farther than 2 km (and less than 5 km) from the offshore outfall
- Transition area station FF12, located between Boston Harbor and the offshore outfall (just under 8 km from the offshore outfall)
- Farfield reference stations FF01A, FF04, and FF09, located in Massachusetts Bay but farther than 13 km from the offshore outfall

Sampling effort at these stations has varied somewhat during the monitoring program. In particular, from 2004-2010 some stations were sampled only during even years (NF22, FF04 and FF09), Stations NF17 and NF12 were sampled each year, and the remaining stations were sampled only during odd years.

Sampling at Station FF04 within the Stellwagen Bank National Marine Sanctuary was conducted in accordance with Research Permit SBNMS-2016-003-A1.

Soft-bottom stations were sampled for grain size composition, total organic carbon (TOC), and the sewage tracer *Clostridium perfringens*. Infauna samples were also collected using a 0.04-m^2 Ted Young-modified van Veen grab; infauna samples were rinsed with filtered seawater through a 300-µm-mesh sieve.

[1] The current monitoring areas form a subset of stations that were sampled before 2011. For example, the transition area formerly included station FF12 and two others that are no longer sampled.

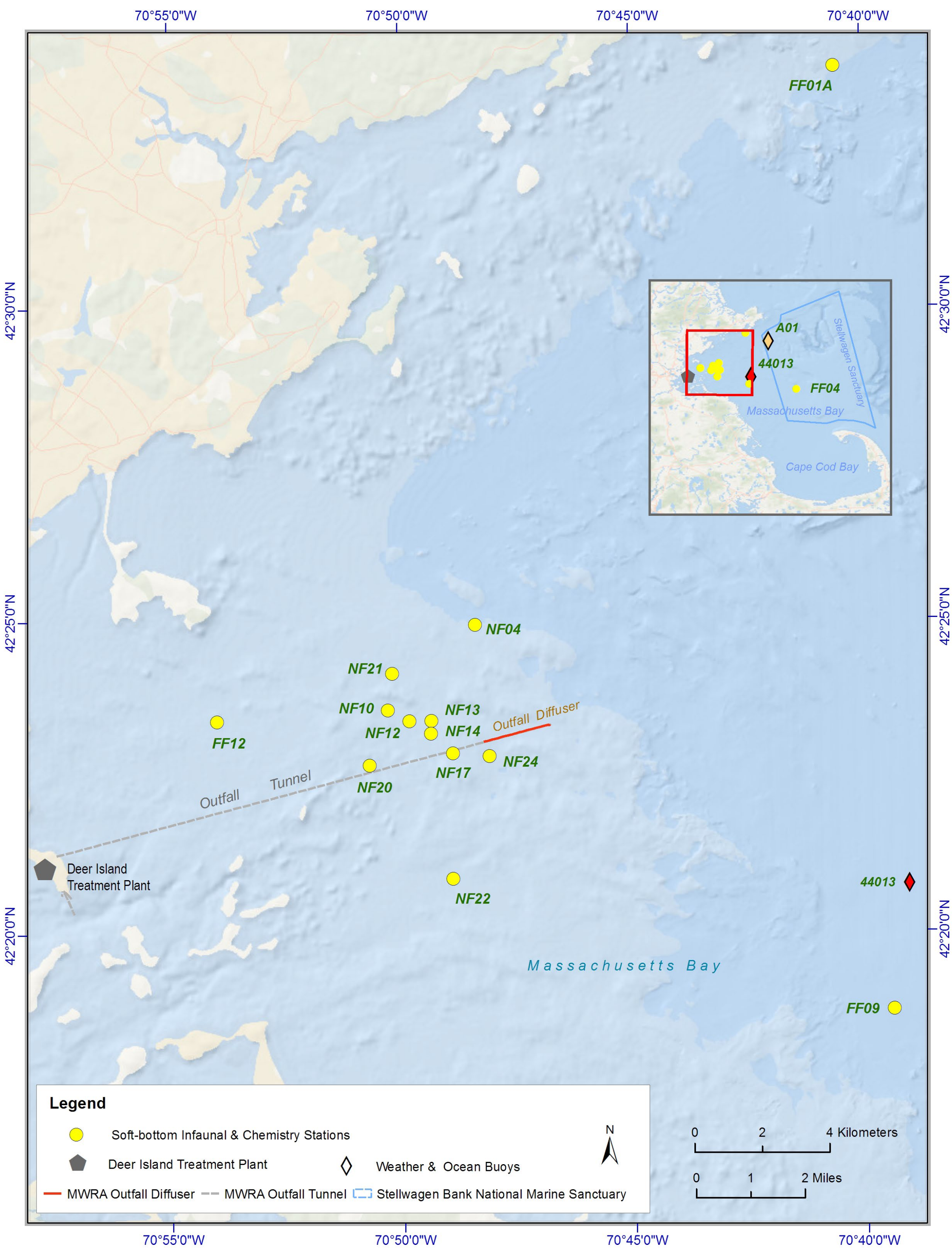

Figure 2-1. Locations of soft-bottom sampling stations for 2021. Inset map shows farfield station FF04.

2.2 LABORATORY METHODS

All bacteriological, physical and chemical analyses were conducted by MWRA's DLS Central Laboratory or its contractors following the procedures described in Constantino et al. (2014). All sample processing, including sorting, identification, and enumeration of infaunal organisms, was done following methods consistent with the QAPP (Rutecki et al. 2020).

2.3 DATA HANDLING, REDUCTION, AND ANALYSIS

All benthic data were extracted directly from the MWRA's database and imported into Excel. Data handling, reduction, graphical presentations and statistical analyses were performed as described in the QAPP (Rutecki et al. 2020) or by Maciolek et al. (2008).

Additional multivariate techniques were used to evaluate infaunal communities. Multivariate analyses were performed using PRIMER v7 (Plymouth Routines in Multivariate Ecological Research) software to examine spatial patterns in the overall similarity of benthic assemblages in the survey area (Clarke 1993, Warwick 1993, Clarke and Green 1988). These analyses included classification (cluster analysis) by hierarchical agglomerative clustering with group average linking and ordination by non-metric multidimensional scaling (MDS). Bray-Curtis similarity was used as the basis for both classification and ordination. Prior to analyses, infaunal abundance data were fourth-root transformed to ensure that all taxa, not just the numerical dominants, would contribute to similarity measures.

Cluster analysis produces a dendrogram that represents discrete groupings of samples along a scale of similarity. This representation is most useful when delineating among sites with distinct community structure. MDS ordination produces a plot or "map" in which the distance between samples represents their rank ordered similarities, with closer proximity in the plot representing higher similarity. Ordination provides a more useful representation of patterns in community structure when assemblages vary along a steady gradation of differences among sites. Stress provides a measure of adequacy of the representation of similarities in the MDS ordination plot (Clarke 1993). Stress levels less than 0.05 indicate an excellent representation of relative similarities among samples with no prospect of misinterpretation. Stress less than 0.1 corresponds to a good ordination with no real prospect of a misleading interpretation. Stress less than 0.2 still provides a potentially useful two-dimensional picture, while stress greater than 0.3 indicates that points on the plot are close to being arbitrarily placed. Together, cluster analysis and MDS ordination provide a highly informative representation of patterns of community-level similarity among samples. The "similarity profile test" (SIMPROF) was used to provide statistical support for the identification of faunal assemblages (i.e., selection of cluster groups). SIMPROF is a permutation test of the null hypothesis that the groups identified by cluster analysis (samples included under each node in the dendrogram) do not differ from each other in multivariate structure.

To help with assessment of spatial patterns, stations have been grouped into regions according to distance from the outfall. The monitoring areas include nearfield stations <2 km from the outfall, nearfield stations >2 km from the outfall, a transition station, and farfield stations (see Section 2.1).

3 RESULTS AND DISCUSSION

3.1 SEDIMENT CONDITIONS

3.1.1 *Clostridium perfringens*, Grain Size, and Total Organic Carbon

Sediment conditions were characterized by three parameters measured during 2021 at each of the 14 sampling stations: (1) *Clostridium perfringens*, (2) grain size (categories from coarse to fine: gravel, sand, silt, and clay), and (3) total organic carbon (Table 3-1).

Table 3-1. Monitoring results for sediment condition parameters in 2021.

Monitoring Area	Station	*Clostridium perfringens* (cfu/g dry/%fines)	Total Organic Carbon (%)	Gravel (%)	Sand (%)	Silt (%)	Clay (%)	Percent Fines (Silt + Clay)
Nearfield (<2 km from outfall)	NF13	84.4	0.16	0.0	96.7	1.5	1.8	3.3
	NF14	25.6	0.34	18.0	70.0	7.4	4.7	12.1
	NF17	151.9	0.08	0.0	97.9	0.5	1.6	2.1
	NF24	79.5	1.46	0.0	29.7	54.2	16.1	70.3
Nearfield (>2 km from outfall)	NF04	6.6	0.18	2.5	58.5	29.7	9.4	39.0
	NF10	25.8	0.64	3.2	59.9	29.4	7.5	36.9
	NF12	1.0	1.64	0.0	25.7	52.7	21.6	74.3
	NF20	39.2	0.67	9.5	71.1	12.0	7.4	19.4
	NF21	5.6	1.07	0.0	39.1	50.5	10.4	60.9
	NF22	44.2	0.87	3.1	57.1	30.4	9.5	39.8
Transition Area (~ 8 km from outfall)	FF12	48.9	0.57	8.8	85.3	2.5	3.5	6.0
Farfield (>13 km from outfall)	FF01A	3.7	0.31	0.4	85.9	11.1	2.6	13.7
	FF04	4.1	1.69	0.0	11.3	58.1	30.6	88.7
	FF09	10.8	0.59	0.6	78.1	12.1	9.3	21.3

Spores of the anaerobic bacterium *Clostridium perfringens* provide a sensitive tracer of effluent solids. *C. perfringens* abundances were reported as colony forming units per gram dry weight, normalized to percent fines (silt and clay). Abundances were normalized because the distribution of C. perfringens varies with the proportion of fine-grained material in the sediments, and normalization provides a more conservative means of evaluating the data for trends (Parmenter and Bothner, 1993). A sharp increase in *C. perfringens* concentrations at sites within two kilometers from the diffuser occurred coincident with diversion of effluent to the offshore outfall (Figure 3-1). *C. perfringens* concentrations have declined or remained comparable to the baseline at all other monitoring locations during the post-diversion period. Statistical analyses reported in Maciolek et al. (2007, 2008) confirmed that concentrations of *C. perfringens* were significantly higher at stations close to the outfall in 2006 and 2007 compared to pre-diversion concentrations and consistent with an impact of the outfall discharge. *C. perfringens* counts in samples collected during 2021 were relatively low at most locations. Counts were similar to the previous year at nearfield locations within two kilometers from the discharge and decreased at the nearfield locations greater than two kilometers (Figure 3-1). *C. perfringens* counts were also similar to the previous year in the transition area of the Bay. The farfield locations had slightly lower *C. perfringens* counts in 2021 compared to 2020. As in past years during the post-diversion period, *C. perfringens* concentrations during 2021 continued to indicate a footprint of the effluent plume at sites closest to the discharge. Normalized *C. perfringens* spore counts in samples collected in 2021 were highest at NF17, NF13, and NF24; three stations located within two kilometers from the discharge (Table 3-1, Figure 3-2).

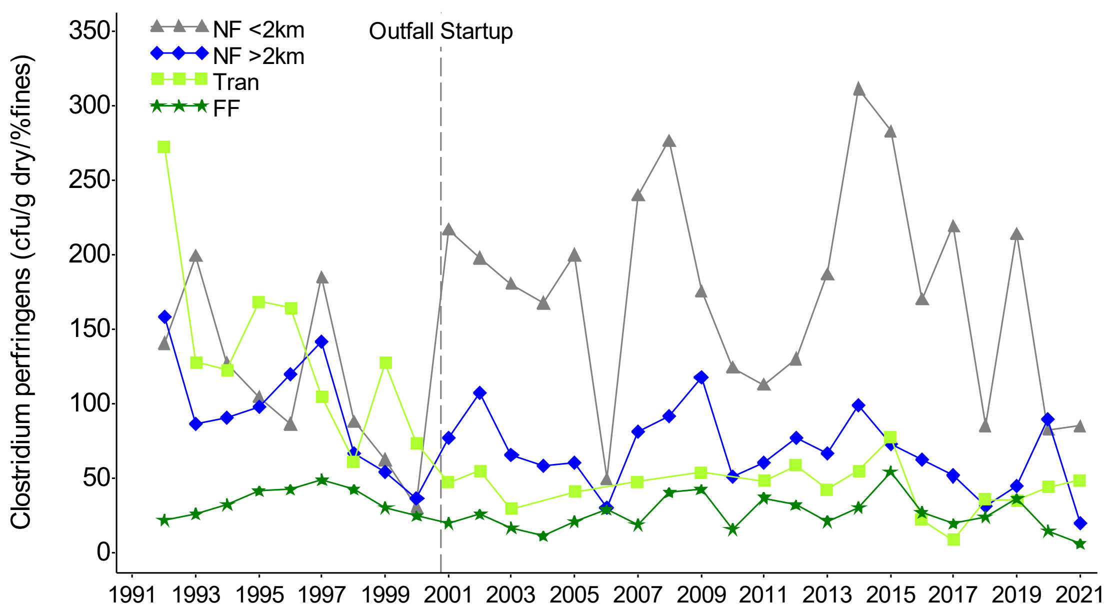

Figure 3-1. Mean concentrations of *Clostridium perfringens* in four areas of Massachusetts Bay, 1992 to 2021. Tran=Transition area; NF<2km=nearfield, less than two kilometers from the outfall; NF>2km=nearfield, more than two kilometers from the outfall; FF=farfield.

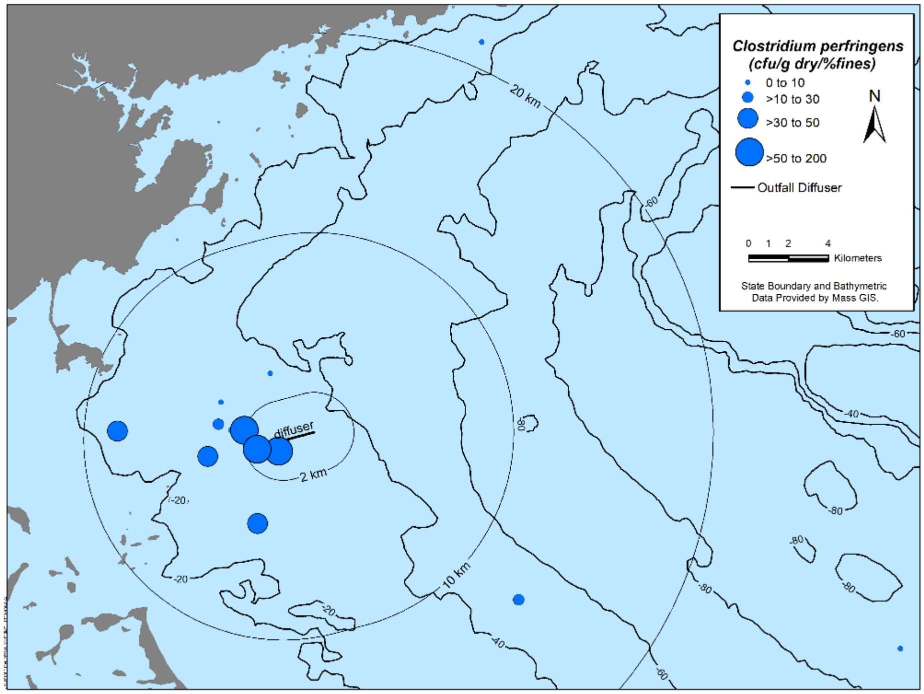

Figure 3-2. Monitoring results for *Clostridium perfringens* in 2021.

Sediment texture in 2021 varied considerably among the 14 stations, ranging from almost entirely sand (e.g., NF17 and NF13) to predominantly silt and clay (e.g., FF04), with most stations having mixed sediments (Figure 3-3). Sediment texture has remained generally consistent over time, with relatively small year-to-year changes in the percent fine sediments at most stations (Figure 3-4). Annual variability in sediment texture at the Massachusetts Bay stations has typically been associated with strong storms. Sediment transport at water depths less than 50 meters near the outfall site in Massachusetts Bay occurs largely as a result of wave-driven currents during strong northeast storms (Bothner et al. 2002).

Concentrations of TOC in 2021 were lower than previous years at FF04 and within the range of historical variability at FF01A, NF17 and NF12 (Figure 3-5). Despite year-to-year variability, TOC values have generally remained consistent with historically reported values at different locations within the Bay (Figure 3-5). Higher TOC values are generally associated with higher percent fines (compare Figures 3-4 and 3-5). To further assess spatial patterns in TOC concentrations while accounting for the association between TOC and percent fine sediments, TOC values were normalized to percent fines (Figure 3-6).

C. perfringens counts continue to provide evidence of effluent solids depositing near the outfall. There is no indication, however, that the wastewater discharge has resulted in changes to the sediment grain size composition at the Massachusetts Bay sampling stations, and there is no indication of organic enrichment. Overall, TOC concentrations remain comparable to, or lower than, values reported during the baseline period, even at sites closest to the outfall (Figures 3-7 and 3-8).

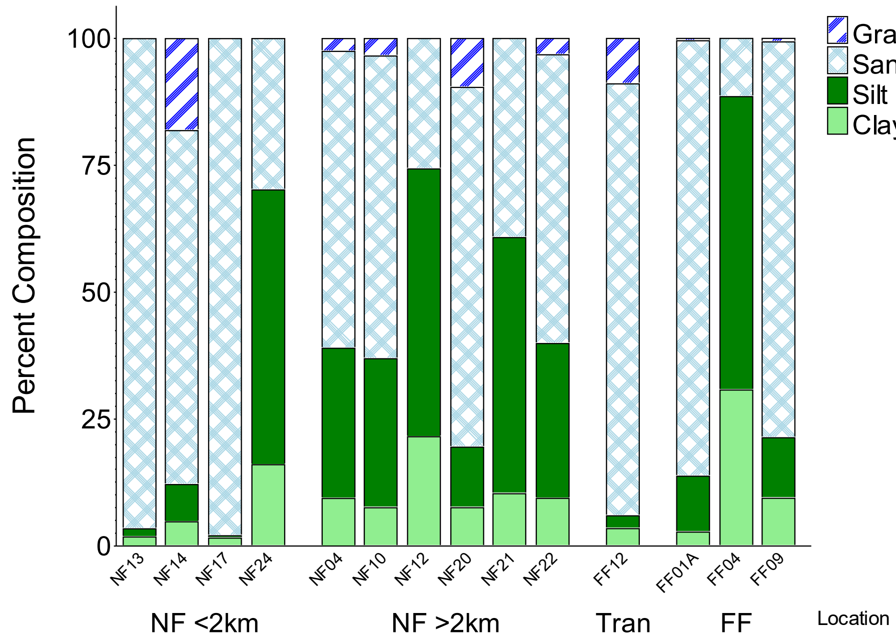

Figure 3-3. Monitoring results for sediment grain size in 2021.

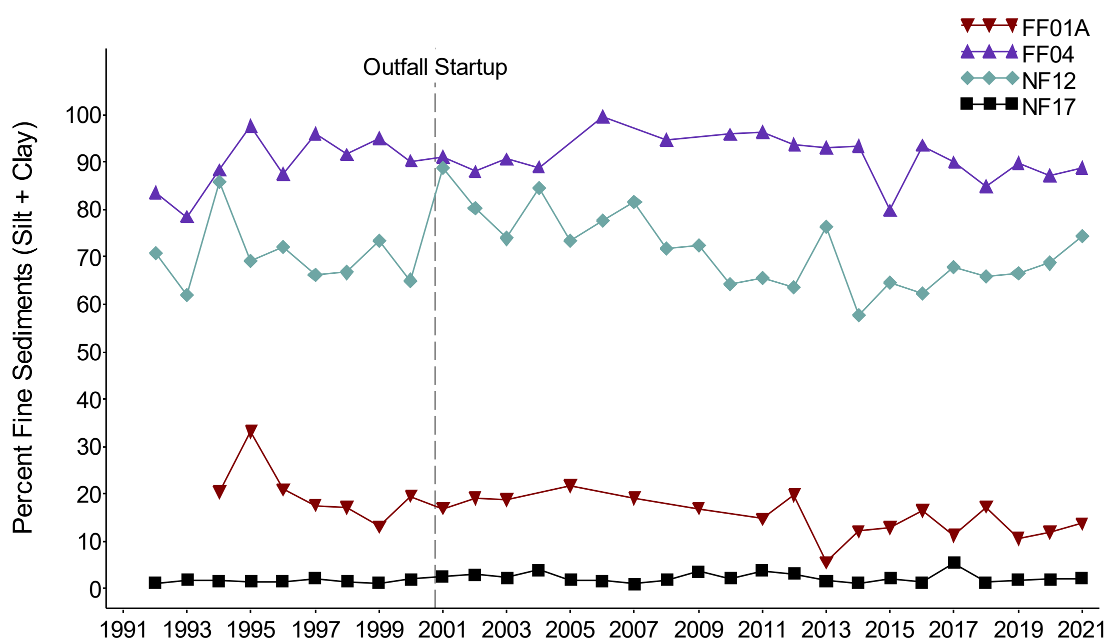

Figure 3-4. Mean percent fine sediments at FF01A, FF04, NF12 and NF17; 1992 to 2021.

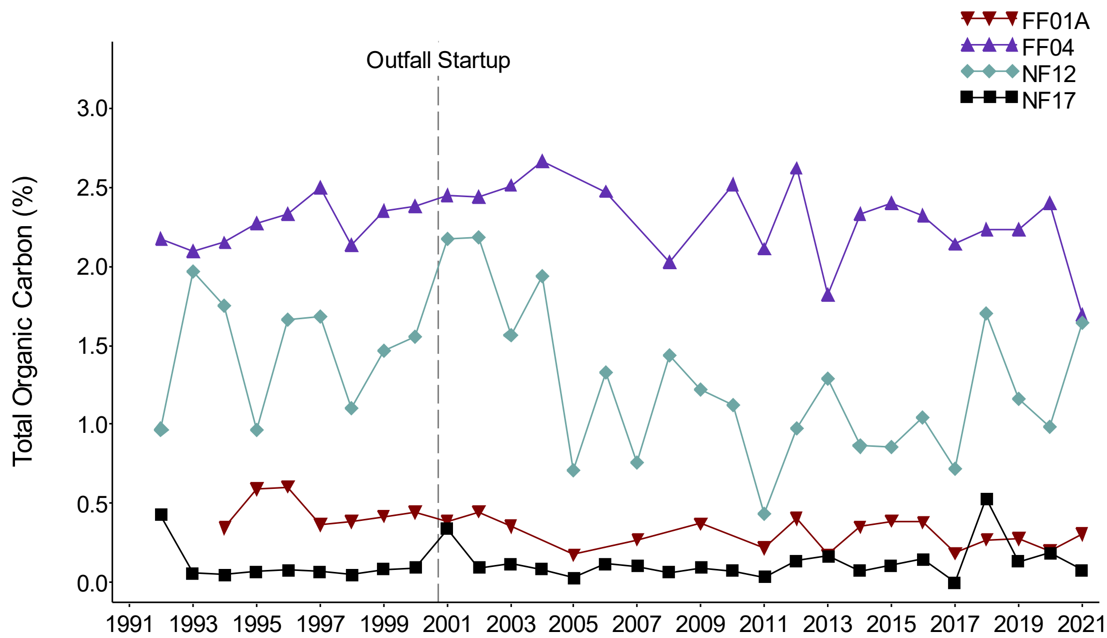

Figure 3-5. Mean concentrations of TOC at four stations in Massachusetts Bay, 1992 to 2021.

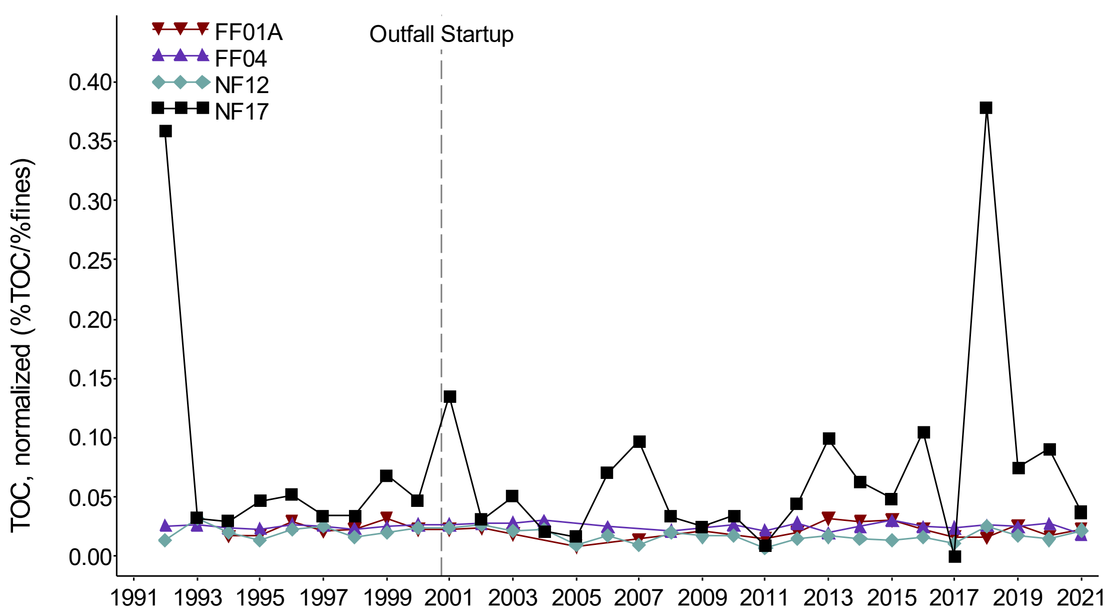

Figure 3-6. Normalized mean concentrations of TOC at four stations in Massachusetts Bay, 1992 to 2021.

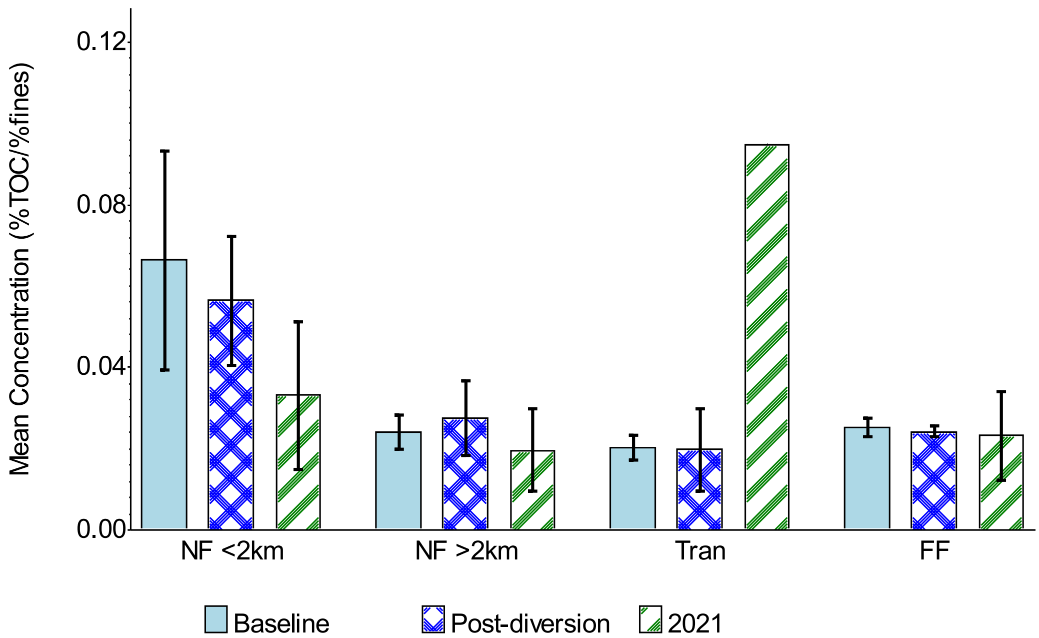

Figure 3-7. Normalized mean (with 95% confidence intervals) concentrations of TOC at four areas in Massachusetts Bay during the baseline (1992 to 2000) and post-diversion (2001 to 2020) periods compared to 2021. Tran=Transition area; NF<2km=nearfield, less than two kilometers from the outfall; NF>2km=nearfield, more than two kilometers from the outfall; FF=farfield.

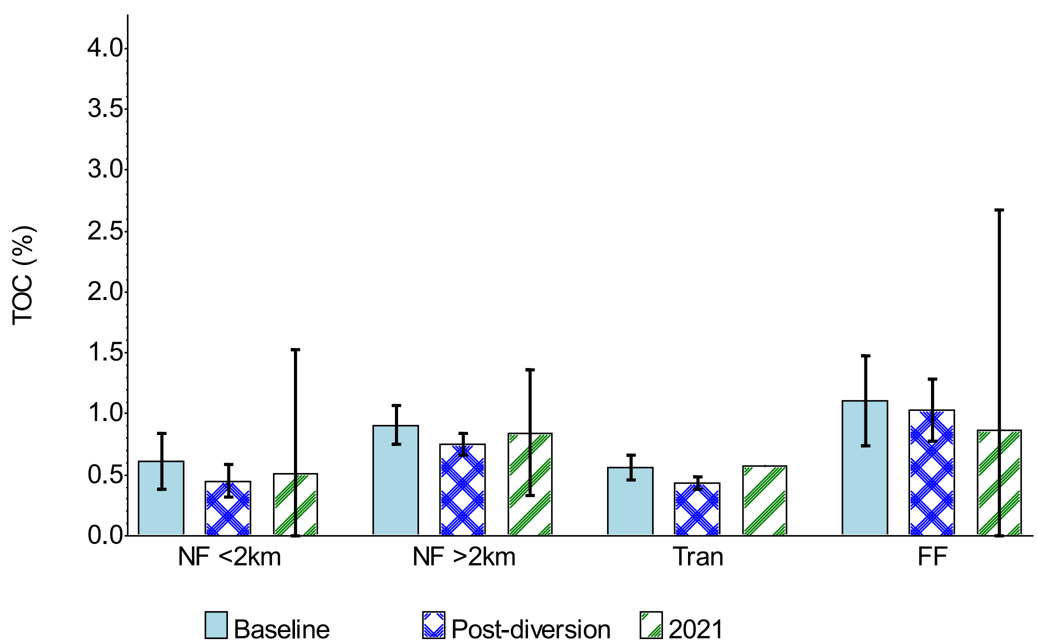

Figure 3-8. Mean (with 95% confidence intervals) concentrations of TOC at four areas in Massachusetts Bay during the baseline (1992-2000) and post-diversion (2001 to 2020) compared to 2021.

3.2 BENTHIC INFAUNA

3.2.1 Community Parameters

A total of 25,086 infaunal organisms were counted from the 14 samples in 2021. Organisms were classified into 208 discrete taxa; 182 of those taxa were species-level identifications. The abundance values reported herein reflect the total counts from both species and higher taxonomic groups, while all diversity measures and multivariate analyses are based on the species-level identifications only (Table 3-2).

Table 3-2. Monitoring results for infaunal community parameters in 2021.

Monitoring Area	Station	Total Abundance (per grab)	Number of Species (per grab)	Log-series alpha	Shannon-Wiener Diversity (H′)	Pielou's Evenness (J')
Nearfield (<2 km from outfall)	NF13	1518	73	16.00	4.17	0.67
	NF14	1929	71	14.49	4.11	0.67
	NF17	907	50	11.39	3.88	0.69
	NF24	2474	68	12.93	3.76	0.62
Nearfield (>2 km from outfall)	NF04	1736	79	17.05	4.40	0.70
	NF10	2096	69	13.70	4.37	0.72
	NF12	1219	57	12.40	4.06	0.70
	NF20	2950	66	11.98	4.02	0.66
	NF21	1568	64	13.42	4.37	0.73
	NF22	1777	56	11.00	4.10	0.71
Transition Area (~ 8 km from outfall)	FF12	2797	61	11.01	3.97	0.67
Farfield (>13 km from outfall)	FF01A	1036	63	14.77	4.07	0.68
	FF04	799	43	9.73	4.05	0.75
	FF09	1592	93	21.55	4.83	0.74

Total abundance values in 2021 were higher than the 2020 values at all areas in Massachusetts Bay except the farfield stations located farther than 13 km from the discharge (Figure 3-9). The numbers of species per sample in 2021 were also higher than in 2020 at most locations; higher species richness occurred at the "Nearfield < 2 km", the "Transition" and the "Farfield" locations, species richness was lower at the "Nearfield >2 km" locations (Figure 3-10). Shannon-Wiener Diversity (H') and Pielou's Evenness (J') values were higher in 2021 compared to the previous year at the nearfield stations, and within the range of variability reported historically (Figures 3-11 and 3-12). The spionid polychaete *Prionospio steenstrupi* remained relatively abundant in 2021, though slightly less abundant than in 2019 and 2020 (Figure 3-13).

This species was the numerical dominant at the nearfield stations during 2021, while other dominant species included the polychaetes *Aricidea catherinae* and *Mediomastus californiensis* (Figure 3-13).

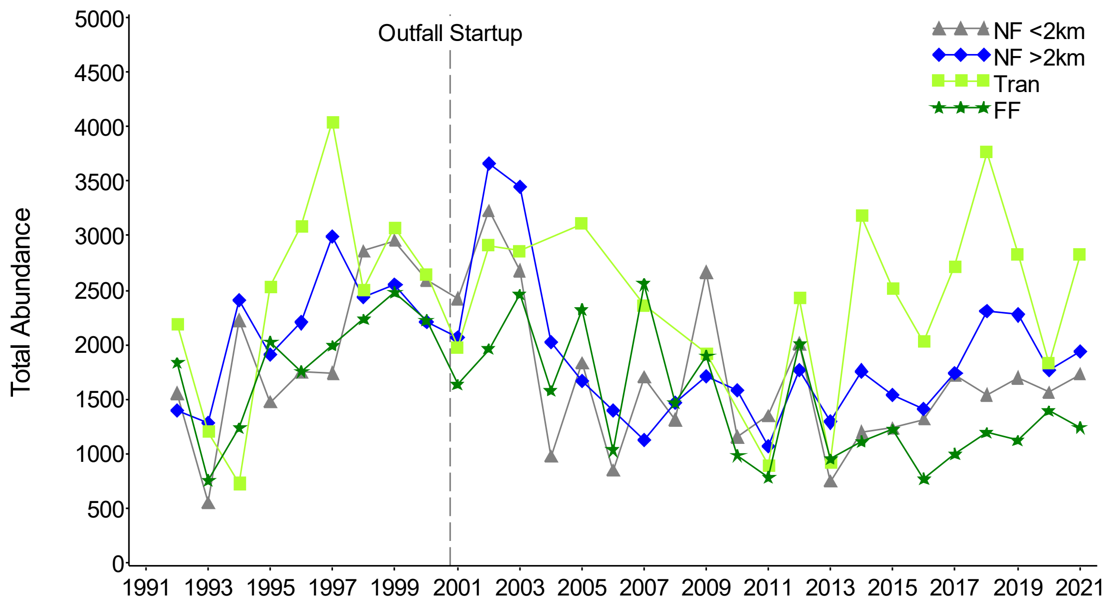

Figure 3-9. Mean infaunal abundance per sample at four areas of Massachusetts Bay, 1992 to 2021. Tran=Transition area; NF<2km=nearfield, less than two kilometers from the outfall; NF>2km=nearfield, more than two kilometers from the outfall; FF=farfield.

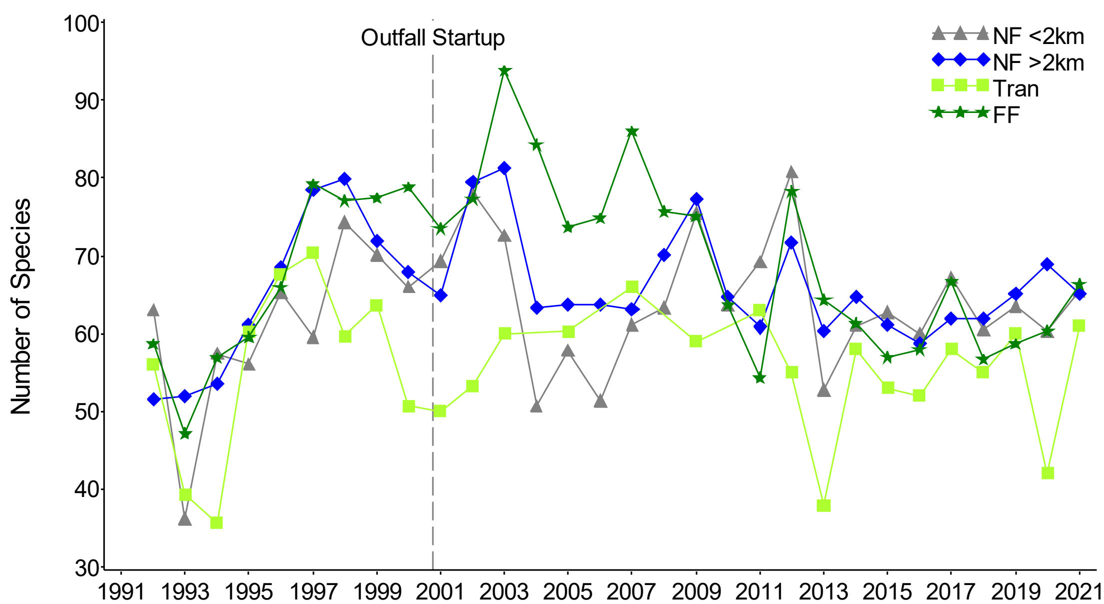

Figure 3-10. Mean number of species per sample at four areas of Massachusetts Bay, 1992 to 2021. Tran=Transition area; NF<2km=nearfield, less than two kilometers from the outfall; NF>2km=nearfield, more than two kilometers from the outfall; FF=farfield.

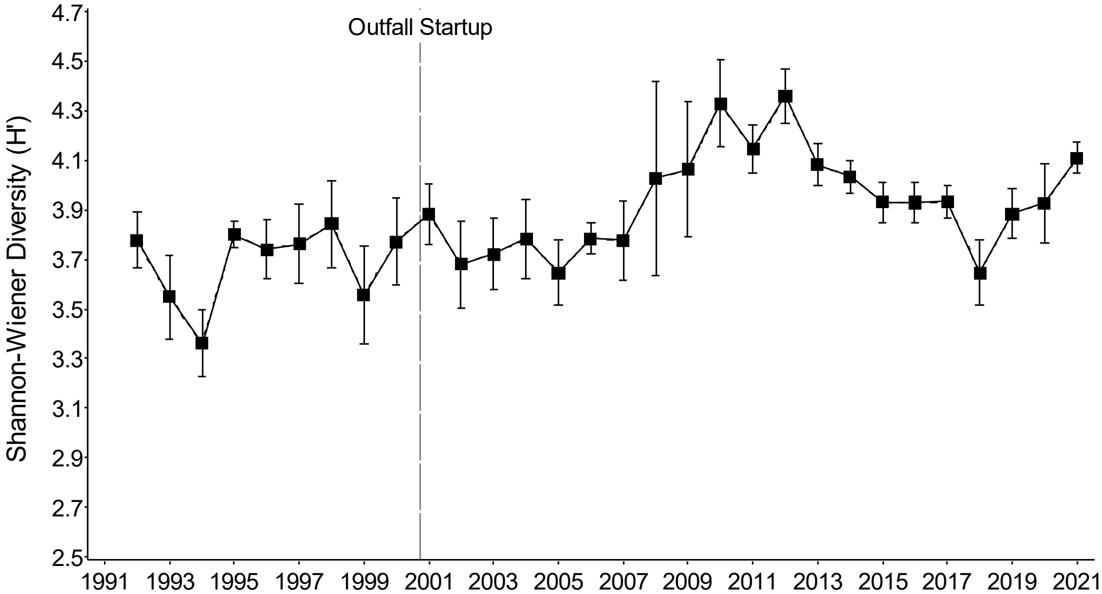

Figure 3-11. Mean (and 95% confidence intervals) Shannon-Wiener Diversity (H′) at nearfield stations in comparison to threshold limit, 1992 to 2021. The nearfield annual means and associated threshold limit are both based on the list of stations sampled following the 2010 revision to the Ambient Monitoring Plan (MWRA 2010).

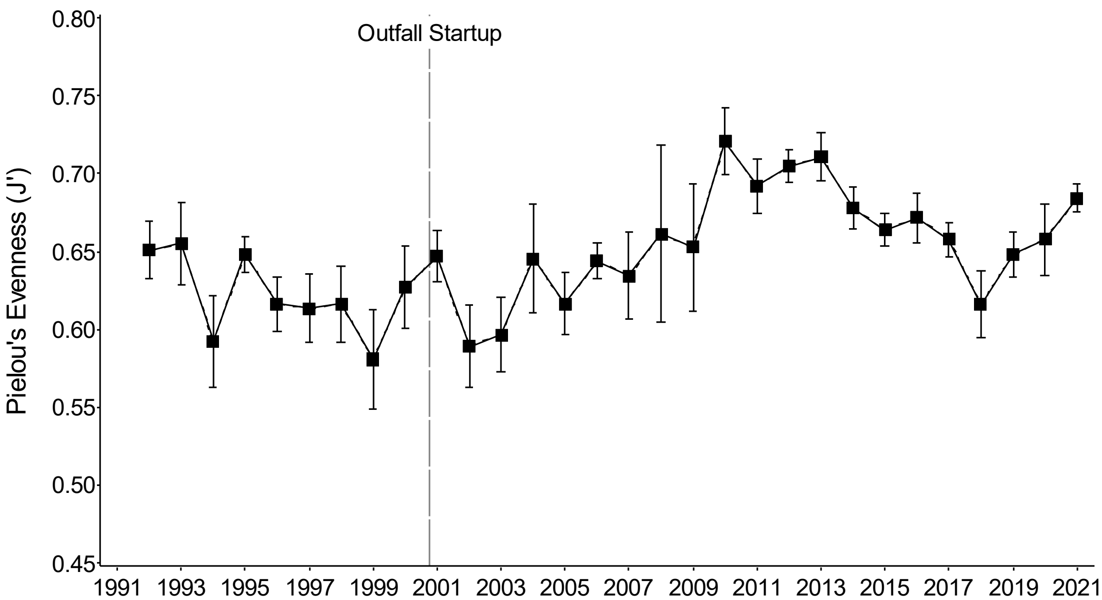

Figure 3-12. Mean (and 95% confidence intervals) Pielou's Evenness (J′) at nearfield stations in comparison to threshold limit, 1992 to 2021. The nearfield annual means and associated threshold limit are both based on the list of stations sampled following the 2010 revision to the Ambient Monitoring Plan (MWRA 2010).

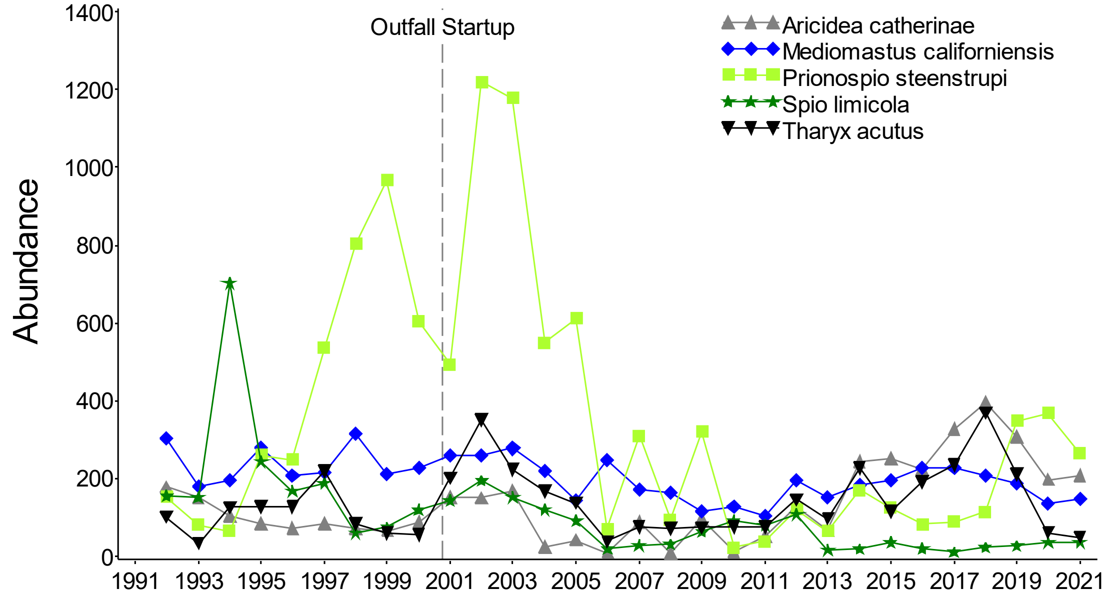

Figure 3-13. Mean abundance from 1992 to 2021, of the five numerically dominant taxa at nearfield stations in Massachusetts Bay.

There were no Contingency Plan threshold exceedances for any infaunal diversity measures in 2021 (Table 3-3).

Table 3-3. Infaunal monitoring Contingency Plan threshold results, August 2021 samples.

Parameter	Thresholds*		Result	Exceedance?
	Value	Limit		
Total species	42.99	Low	64.90	No
Log-series Alpha	9.42	Low	13.20	No
Shannon-Weiner H′	3.37	Low	4.11	No
Pielou's J'	0.57	Low	0.68	No
Percent opportunists	10% (Caution)	High	0.53	No
Percent opportunists	25% (Warning)	High	0.53	No

*Threshold exceedances occur when current year results are below threshold values for a "low" limit or above the values for a "high" limit for a given parameter.

Spatial and temporal patterns of abundance, species richness, species diversity and evenness generally support the conclusion that there is no evidence of negative impacts caused by operation of the offshore outfall.

3.2.2 Infaunal Assemblages

Multivariate analyses based on Bray-Curtis Similarity were used to assess spatial patterns in the faunal assemblages at the Massachusetts Bay sampling stations. Two main assemblages (Groups I and II) and an outlier assemblage (Group III) were identified in a cluster analysis of the 14 samples from 2021 (Figure 3-14). The groups were distinguished based on species composition and the relative abundances of each taxon in the samples. Clear differences in the mean abundances of dominant taxa were identified. Abundances at the stations included in Groups I and III were generally lower than Group II (compare Figure 3-14 with Table 3-2). The three assemblages were dominated by polychaetes, although arthropods were a dominant in Group I and molluscs were dominants in Group IIA (Table 3-4). Several species were dominant only in Group I (e.g., *Capitella capitata* complex, *Crassicorophium crassiorne, Polygordius jouinae, Parexogone hebes,* and *Tanaissus psammophilus*), while others were more prevalent in Group II (e.g., *Solamen glandula, Mediomastus californiensis, Prionospio steenstrupi,* and *Spio limicola*) or in Group III (e.g., *Chaetozone anasimus* and *Cossura longocirrata*). *Capitella capitata* complex is a taxon often associated with organic enrichment (Grassle and Grassle 1976). This taxon is one of the "opportunistic species" that are tracked by MWRA as a Contingency Plan threshold for infaunal communities at the nearfield stations. *C. capitata* complex has not typically been a dominant taxon at the Massachusetts Bay sampling stations. Nonetheless, during 2021, *C. capitata* complex was found in relatively high numbers within the Group IB assemblage, which occurred only at farfield station FF01A (Table 3-4). Since these higher numbers of *C. capitata* complex were found only at a farfield station, this finding is not consistent with a potential impact from the offshore outfall.

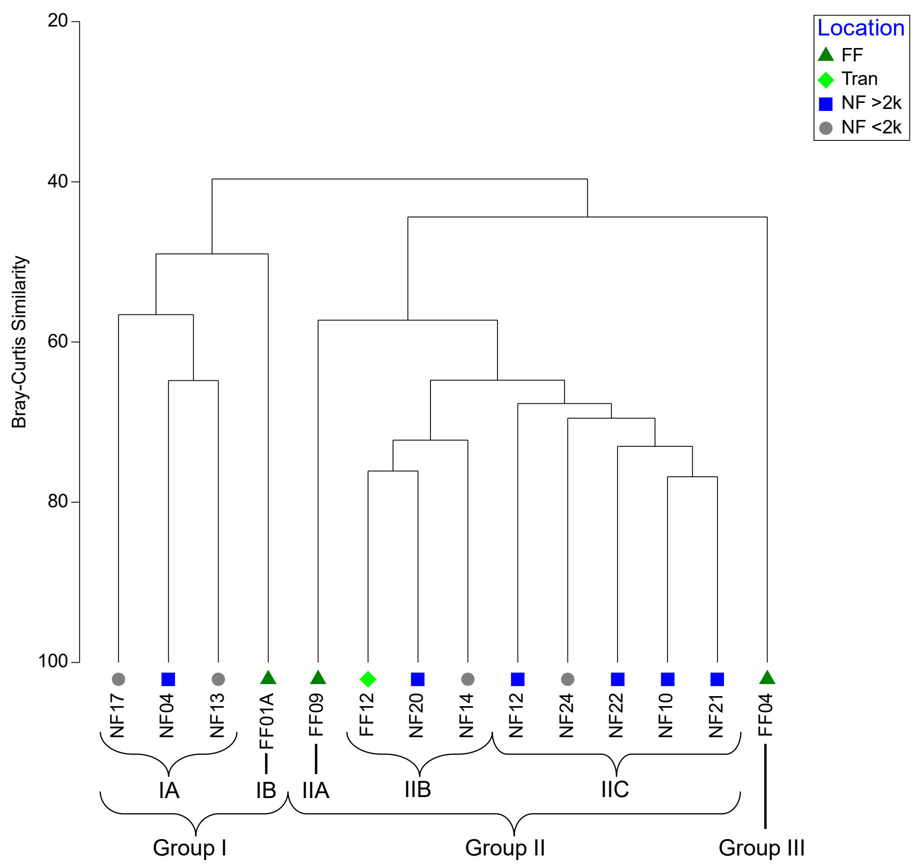

Figure 3-14. Results of cluster analysis of the 2021 infauna samples.

Table 3-4. Abundance (mean # per grab) of numerically dominant taxa (10 most abundant per group) composing infaunal assemblages identified by cluster analysis of the 2021 samples.

Family	Species	Group I		Group II			Group
		IA	IB	IIA	IIB	IIC	FF04
Mollusca (Bivalvia)							
Mytilidae	*Solamen glandula*	1.7	-	153.0	6.3	0.8	-
Nuculidae	*Ennucula*	12.3	-	98.0	52.0	10.6	5.0
Thyasiridae	*Thyasira gouldii*	-	-	43.0	-	3.4	20.0
Annelida (Polychaeta)							
Paraonidae	*Aricidea (acmira)*	90.0	15.0	8.0	483.7	111.4	-
	Aricidea (strelzovia)	-	-	43.0	1.7	33.6	54.0
Ampharetidae	*Ampharete lindstroemi*	19.3	32.0	8.0	2.7	0.6	-
	Anobothrus gracilis	1.7	-	101.0	1.3	1.2	105.0
Capitellidae	*Capitella capitata*	-	277.0	-	-	1.4	-
	Mediomastus	4.7	38.0	48.0	204.0	201.6	10.0
Apistobranchidae	*Apistobranchus typicus*	-	1.0	1.0	0.3	11.2	72.0
Cirratulidae	*Chaetozone anasimus*	13.3	8.0	3.0	-	-	93.0
	Kirkegaardia	7.3	40.0	-	98.7	96.4	-
	Kirkegaardia	1.3	11.0	3.0	102.0	61.6	-
	Tharyx acutus	23.7	35.0	23.0	68.0	45.2	-
Cossuridae	*Cossura longocirrata*	-	3.0	3.0	-	8.4	65.0
Lumbrineridae	*Ninoe nigripes*	0.7	-	54.0	79.0	82.0	16.0
Nephtyidae	*Aglaophamus circinata*	35.3	14.0	3.0	3.3	-	-
Orbiniidae	*Leitoscoloplos acutus*	0.3	-	12.0	23.0	45.8	19.0
Paraonidae	*Levinsenia gracilis*	1.3	16.0	175.0	142.0	238.8	114.0
Polygordiidae	*Polygordius jouinae*	191.0	220.0	11.0	17.0	1.6	-
Sabellidae	*Euchone incolor*	9.3	34.0	92.0	95.3	157.6	89.0
ScalibregmatidaeSpionidae	*Scalibregma inflatum*	45.7	12.0	1.0	45.7	29.8	1.0
Spionidae	*Prionospio steenstrupi*	10.7	49.0	149.0	484.7	287.4	29.0
	Spio limicola	0.7	-	162.0	4.0	73.6	11.0
	Spiophanes bombyx	91.3	19.0	-	192.3	50.6	-
Syllidae	*Erinaceusyllis*	-	24.0	-	-	-	-
	Parexogone hebes	193.7	25.0	10.0	71.0	13.6	-
Arthropoda							
Corophiidae	*Crassicorophium*	227.7	9.0	-	1.7	-	-
Idoteidae	*Edotia montosa*	26.3	-	6.0	12.7	5.6	-
Nototanaidae	*Tanaissus*	58.7	-	-	-	0.4	-

Group I consisted of two subgroups and was composed of three nearfield stations (Group IA: Stations NF17, NF04, and NF13; Group IB: Station FF01A). The Group II assemblage included three subgroups (Group IIA: Station FF09; Group IIB: Stations FF12, NF20, and NF14; and Group IIC: Stations NF12, NF24, NF22, NF10, and NF21) that could be differentiated by species composition and total abundance. The relatively deep Station FF04 was characterized by low abundances and species richness. The outlier assemblage that was found at this station was labeled as Group III. Dominant species at Station FF04, including *Levinsenia gracilis, Cossura longocirrata,* and *Chaetozone anasimus*, are characteristic of the soft sediment community observed throughout Stellwagen Basin (e.g., Maciolek et al. 2008).

Both main assemblages (Groups I and II) occurred at one or more of the four stations within two kilometers of the discharge as well as at stations more than two kilometers from the discharge (Figure 3-15). Thus, stations closest to the discharge were not characterized by a unique faunal assemblage reflecting effluent impacts. Comparisons of faunal distribution to habitat conditions indicated that patterns in the distribution of faunal assemblages follow differences in habitat types at the sampling stations and are associated with the sediment types at the sampling stations (Figure 3-16) and with station depth (not shown).

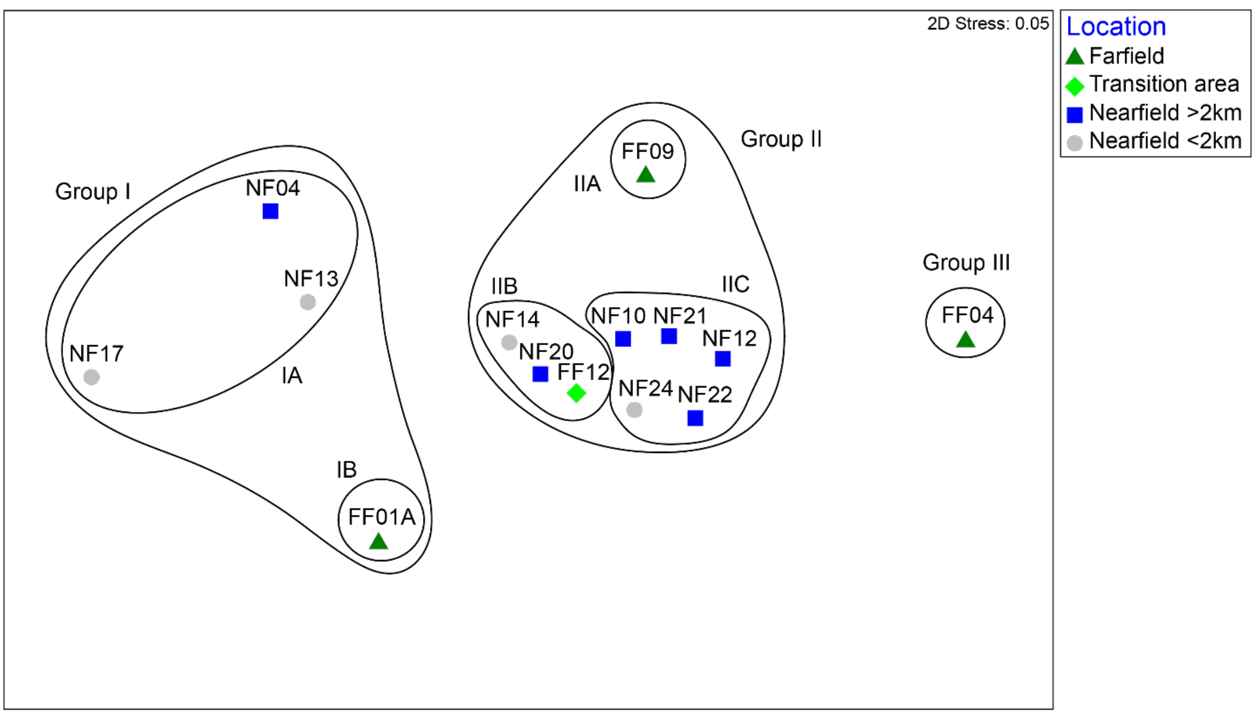

Figure 3-15. Results of a MDS ordination of the 2021 infauna samples from Massachusetts Bay showing distance from the outfall.

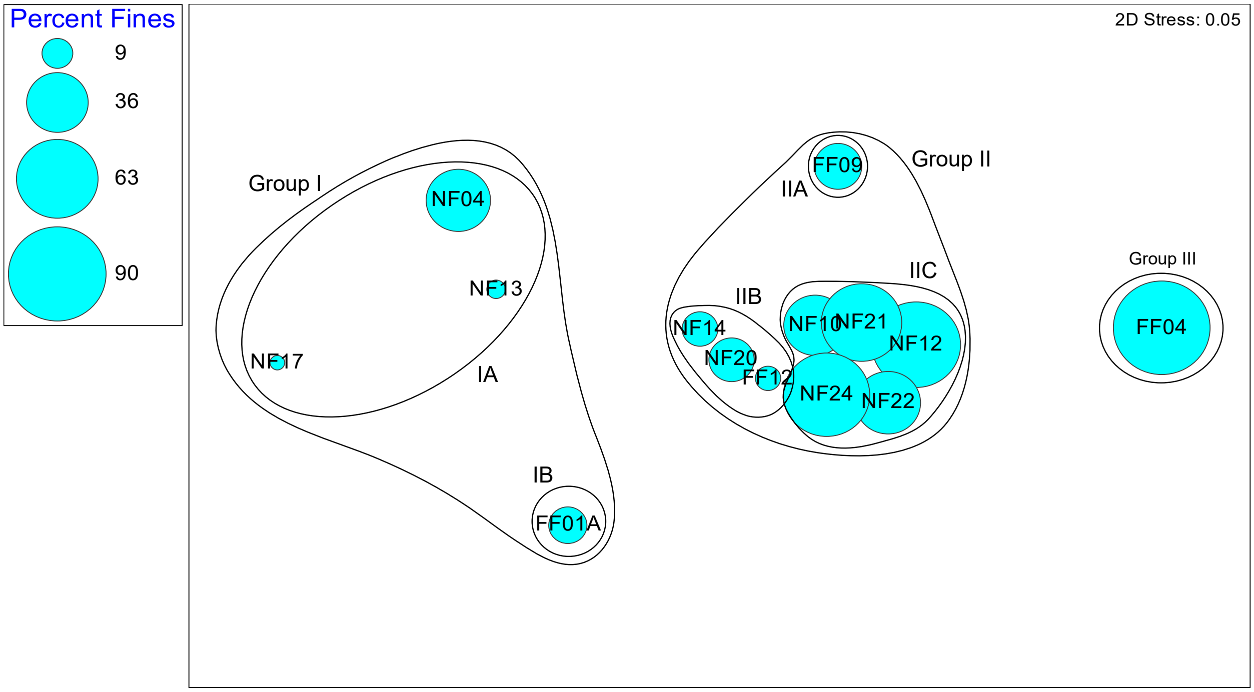

Figure 3-16. Percent fine sediments superimposed on the MDS ordination plot of the 2021 infauna samples. Each point on the plot represents one of the 14 samples; similarity of species composition is indicated by proximity of points on the plot. Faunal assemblages (Groups I-II, and sub-groups) identified by cluster analysis are circled on the plot. The ordination and cluster analysis are both based on Bray-Curtis Similarity.

Patterns identified in these analyses were highly consistent with previous years. No evidence of impacts from the offshore outfall on infaunal communities in Massachusetts Bay was found.

The outfall is located in an area dominated by hydrodynamic and physical factors, including tidal and storm currents, turbulence, and sediment transport (Butman et al. 2008). These physical factors, combined with the high quality of the effluent discharged into the Bay (Taylor 2010, Werme et al. 2021), are the principal reasons that benthic habitat quality has remained high in the nearfield area. Previous assessments have indicated that changes in the benthic habitat quality and infaunal communities in the nearfield are related to physical processes associated with increased storminess (Nestler et al. 2020).

4 SUMMARY OF MONITORING RESULTS

Benthic monitoring for MWRA's offshore ocean outfall focused on addressing three primary concerns regarding potential impacts to the benthos from the wastewater discharge: (1) eutrophication and related low levels of dissolved oxygen; (2) accumulation of toxic contaminants in depositional areas; and (3) smothering of animals by particulate matter.

Findings from previous assessments found no indication that the wastewater discharge has resulted in low levels of dissolved oxygen or the accumulation of toxic contaminants in nearfield sediments (Nestler et al. 2018, 2020; Maciolek et al. 2008). As result, SPI surveys in Massachusetts Bay and the sediment contaminant evaluation every third year at the nearfield and farfield stations were discontinued beginning in 2020. Hard-bottom benthic community monitoring in 2020 also found no evidence that particulate matter from the wastewater discharge has smothered benthic organisms (Rutecki et al. 2022). Although some modest changes in this community (e.g., decreased coralline algae, temporal variability in upright algae cover, and decreased sponge abundance) have been observed, comparisons between the post-diversion and baseline periods indicate that these changes are not substantial (Rutecki et al. 2022).

Surveys of soft-bottom benthic communities presented in this report continue to suggest that animals near the outfall have not been smothered by particulate matter from the wastewater discharge or experienced stress resulting from increased deposition of organic matter. The percentage of fine grain sediments has not increased in the nearfield stations since the diversion indicating no pattern of settlement of particulate matter from the discharge. There were no Contingency Plan threshold exceedances for any infaunal diversity measures in 2021.

The spatial extent of particulate deposition from the wastewater discharge is measurable in the *Clostridium perfringens* concentrations in nearfield sediments. *C. perfringens* concentrations provide evidence of the discharge footprint at stations close to the outfall. Within this footprint, no other changes to sediment composition and infaunal communities have been detected. Nonetheless, subtle variations in the species composition of infaunal assemblages clearly delineate natural spatial variation in the benthic community based on habitat (e.g., associated with different sediment grain sizes) and bottom energy (e.g., turbulence and sediment transport associated with storm events). Changes over time have also been detected including region-wide shifts in diversity, with peaks from 2010 to 2012, in the Massachusetts Bay infaunal assemblages. Detection of these spatial and temporal patterns in the benthos suggests that any ecologically significant adverse impacts from the outfall would be readily detected by the monitoring program if those impacts had occurred.

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