

**Benthic invertebrate samples and data from the
Ocean Survey 20/20 voyages to the
Chatham Rise and Challenger Plateau, 2007**

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EXECUTIVE SUMMARY

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This report describes data sets derived from benthic invertebrate samples collected during the Ocean Survey 20/20 Chatham-Challenger Hydrographic, Biodiversity and Seabed Habitats Project in 2007. To characterise assemblages across a range of organism sizes and spatial scales, samples were collected from the Chatham Rise and the Challenger Plateau using several gear types: coarse mesh 'seamounts' epibenthic sled and beam trawl (sampling mega-epifauna); multicorer (meiofauna); fine mesh 'Brenke' epibenthic sled (macro-epifauna and hyperbenthic fauna); the still image camera of NIWA's Deep Towed Imaging System (DTIS) (mega- and macro-epifauna, bioturbation, and substrate types), and the video camera of DTIS (mega-epifauna and substrate types). The report: (1) details the samples that were available for analysis, (2) describes the methods used to process and extract data from these samples, and (3) presents summaries of the final data sets.

Large numbers of physical specimens and photographic samples were analysed successfully for taxonomic identities and abundances of fauna, bioturbation, trawl marks, and substrate types. Comprehensive data sets were compiled from each of the gear types, but spatial coverage and the level of taxonomic resolution varied with gear type and the size fraction of the biota sampled, with consequent differences in the utility of each data set for addressing the overarching objectives of the OS 20/20 project. The greatest spatial coverage, in terms of both number of sites and area of seabed sampled, was achieved using the towed underwater video system, DTIS, whereas the greatest numbers of taxa and the finest taxonomic resolution were identified from the seamount sled and beam trawl samples. These two gear types (DTIS and sled/beam trawl) were deployed successfully at more than 90% of the OS 20/20 sampling sites and thus provide consistent quantitative data across the entire study area. The least representative data are those for meiofauna and macro-hyperbenthos, where fewer samples, extended sample processing times, and difficult taxonomies resulted in data sets that, in their present form, are of less value for biodiversity studies at the scale of the Chatham-Challenger project.

The samples from these OS 20/20 voyages have generated one of the most comprehensive, spatially extensive, and internally consistent data sets available describing benthic invertebrate distributions in New Zealand waters to date. These data will be of considerable value for future assessments of the ecology of the region.

INTRODUCTION

1.1 Ocean Survey 20/20 Chatham-Challenger project

In November 2004, the Government of New Zealand established Ocean Survey 20/20 (OS 20/20) as a 15 year programme with the aim to:

“Complete by 2020 an ocean survey that will provide New Zealand with the knowledge of its ocean territory to:

- *demonstrate our stewardship and exercise our sovereign rights;*
- *conserve, protect, manage and sustainably utilise our ocean resources; and*
- *facilitate safe navigation and enjoyment of the oceans around New Zealand.”*

The Chatham-Challenger Hydrographic Biodiversity and Seabed Habitats Project was identified by Cabinet as the priority OS 20/20 project for 2006–07 and the Ministry of Fisheries (MFish), Land Information New Zealand (LINZ), Department of Conservation (DoC), and NIWA collaborated in its development.

The overall goal of the Chatham-Challenger OS 20/20 Project was to map and compare the distribution of seabed habitats and their associated biological diversity across the Chatham Rise and the Challenger Plateau. These areas are in contrasting oceanographic environments and are subject to different levels of anthropogenic disturbance from fisheries. The Chatham Rise lies under waters with high biological productivity which support substantial populations of commercially important fish species, whereas the Challenger Plateau has generally lower productivity and is of less importance for fisheries.

The principal objectives of the project were: to determine the distribution of bottom habitats and biodiversity on the seabed in depths from 200 to 1200 m on the Chatham Rise and the Challenger Plateau; to assess the utility of the Marine Environment Classification (Snelder et al. 2006) as a proxy for habitat types and biodiversity distribution and further develop habitat mapping techniques; and to assess the influence of bottom trawling as a broad-scale driver of sea-bed biodiversity.

Three voyages were undertaken in 2006 and 2007 to complete the field sampling phase of the project. The first voyage conducted multi-beam echo-sounder swath mapping of pre-selected transects across the Chatham Rise and the Challenger Plateau (TAN0610, Mitchell & Nodder 2006). These transects were then targeted for biodiversity and habitat studies on two subsequent voyages (TAN0705 to Chatham Rise and TAN0707 to Challenger Plateau; Nodder 2007a/2007b).

1.2 Sampling design

The initial sampling design for the Chatham-Challenger project (Nodder et al. 2007) was based on a multivariate statistical classification of environmental data to determine the optimal number of environmental classes. In this classification, available oceanographic environmental data layers and parameters from the multi-beam transects sampled during voyage TAN0610 were combined. Classifications were run separately for the two locations, identifying eight environmental classes on the Chatham Rise and nine classes on the Challenger Plateau (Figure 1). However, because it was anticipated that time at sea would be insufficient to sample adequately all nine classes on the Challenger Plateau, a secondary classification level was used in which the region was divided into five classes. This is important in relation to the distribution of sampling effort described below. These classes were then used as the strata for the benthic sampling programme, with sampling effort allocated evenly across strata in each location.

To characterise substrate types and biological assemblages across a faunal size range from bacteria to megafauna, samples were collected using several gear types, the principal gears together with their target fauna being: coarse mesh ‘seamounts’ epibenthic sled and beam trawl (mega-epifauna); multicorer (meiofauna); fine mesh ‘Brenke’ epibenthic sled (macro-epifauna and hyperbenthic fauna); the still image camera of NIWA’s Deep Towed Imaging System (DTIS) (mega- and macro-epifauna, bioturbation, and substrate types), and the video camera of DTIS (mega-epifauna and substrate types). Time constraints dictated that not all sampling gear types could be deployed at all sites. Thus, a hierarchy of site types was defined: *A* sites, at which all gear types were deployed with replicate deployments of each gear; *B* sites at which the DTIS camera platform and the seamounts epibenthic sled were deployed once each; and *C* sites, at which only the DTIS was deployed. In practice, it was possible to deploy both the DTIS and the epibenthic sled at all *C* sites, and thus, for the most part, they are the same as *B* sites for analyses. However, the decision to adopt this approach at sea was not made until the first 10 *C* sites had been sampled and thus these sites are represented only by a DTIS transect (Appendix - Table A1).

A further modification to the sampling programme, made possible by favourable conditions at sea, was the addition of a number of extra sites which were not part of the original sampling design. These were designated as *D* sites and were placed in areas where more detailed coverage was anticipated to be of use to subsequent analyses. On the northeast Chatham Rise, 11 *D* sites were targeted in the orange roughly spawning “box” (Dunn 2007), as two sets of sampling sites arranged as north-south transects: a western transect in an area of high fishing intensity (sites D013–D018, Figure 1), and an eastern transect in an area of relatively low fishing intensity (sites D020–D024, Figure 1). These sites were sampled with DTIS, multicorer, and beam trawl.

For analysis, a minor change was made to the original nomenclature of the sites on the Challenger Plateau. In initial site designations, some site codes were duplicated between areas. For instance, the site code D006 existed on both the Chatham Rise and on the Challenger Plateau. To ensure unique codes for all sites across the entire study, all Challenger Plateau sites were given a double letter prefix; e.g., D006 became DD006.

One hundred sites were sampled on the Chatham Rise and 49 sites were sampled on the Challenger Plateau (Figure 1). These totals include 29 and 9 extra *D* sites in each area respectively (Table 1).

Table 1: Total number of sites sampled on Chatham Rise and Challenger Plateau during TAN0705 and TAN0707, respectively. Categories indicate the range of sampling gears deployed at each site; see text for details.

Site category	Chatham Rise	Challenger Plateau
A	11	5
B	37	14
C	23	21
D	29	9
Total	100	49

1.3 Post-voyage analysis of samples

This report describes the methods used for initial processing of Chatham-Challenger OS 20/20 benthic faunal samples under the post-voyage analysis project (MFish code ZBD2007-01) and presents the data sets derived from them. The specific objectives of this research were:

- 1) To count, measure, and identify to species level (where possible) all macro invertebrates (larger than 2 mm) collected by the seamounts sled and beam trawl.

- 2) To count, measure, and identify to the finest practicable taxonomic resolution all meiofauna (from 32 μm to 500 μm) from multicorer samples.
- 3) To count, measure, and identify to species level (where possible) all fauna collected by hyperbenthic sled.
- 4) To count, measure, and identify to species level (where possible) all macrofauna observed in DTIS still images together with estimates of the number of biogenic features (burrows, mounds etc.) and habitat complexity.
- 5) To count, measure, and identify to species level (where possible) all macrofauna observed in DTIS video footage.

The following sections detail the numbers and locations of samples that were available for analysis for each gear type, describe the methods used to process and extract data from the samples, and present summaries of the final data sets. Detailed analyses and ecological interpretation of the data sets are outside the scope of this report and will be reported separately.

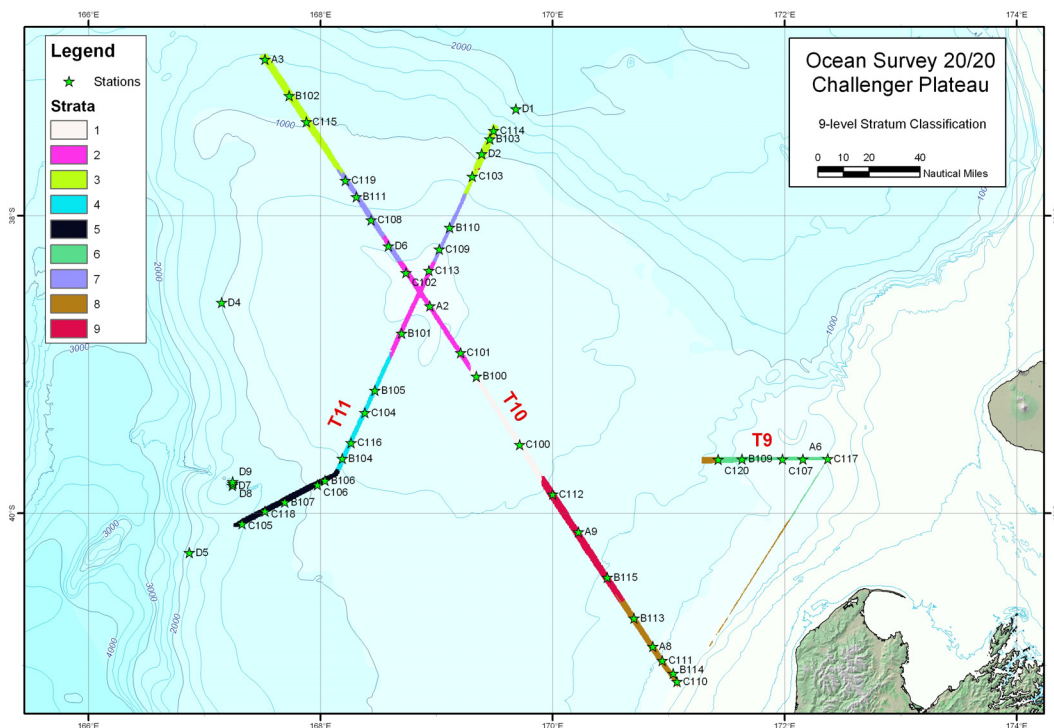
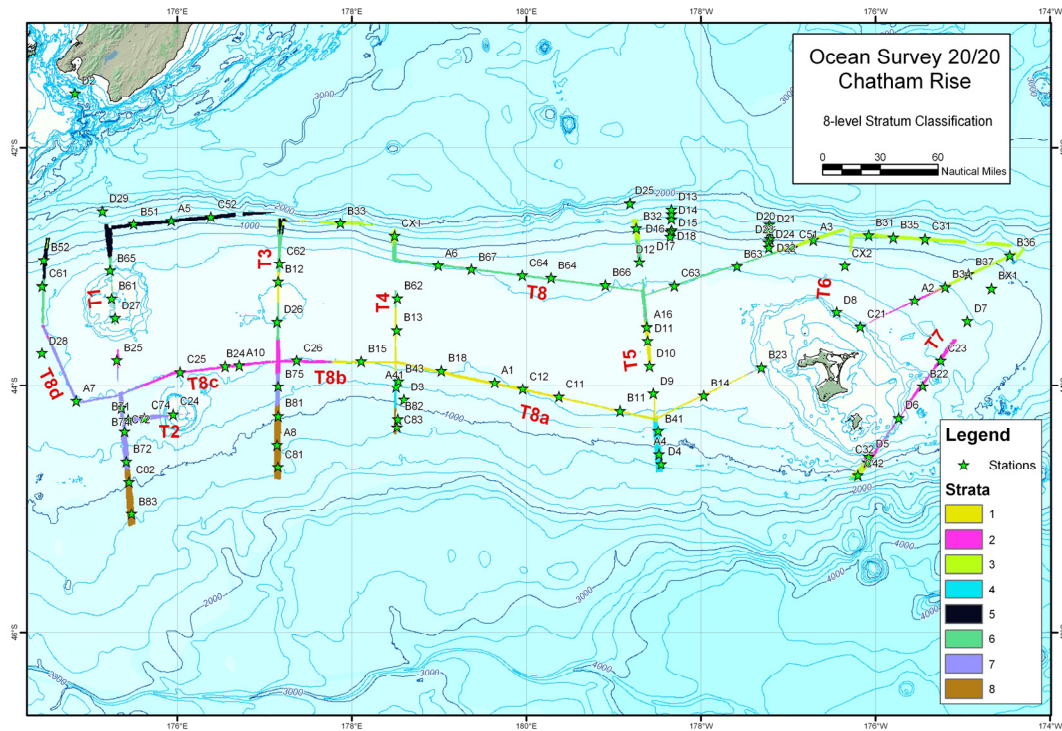


Figure 1: Sites sampled during Ocean Survey 20/20 voyages to Chatham Rise (TAN0705, top) and Challenger Plateau (TAN0707, bottom). Sites are shown as labelled stars (see Section 1.2 for site numbering protocols) superimposed on the survey strata determined *a priori* from acoustic and environmental data. Red ‘T’ prefix numbers identify the multibeam sonar transects collected during TAN0610.

SAMPLES AND PROCESSING METHODS

1.4 Seamounts sled and beam trawl samples (mega-epifauna)

A seamount sled and a beam trawl were deployed to sample organisms larger than ~10 mm (beam trawl) or 25 mm (seamount sled) (*megafauna*) living on the surface of the seabed (*epifauna*) or in surficial sediments where these were soft enough for the lower edge of the gear to dig in.

Depending on substrate suitability, either the seamount sled (SEL) or the beam trawl (TB) was deployed at each site across the two locations. The seamount sled is designed for use on hard, irregular substrata, but sweeps a relatively small seabed area (1 m mouth width), whereas the beam trawl can be used only on smooth substrata but sweeps a larger area (3 m mouth width). More importantly, the SEL has a codend mesh size of 25 mm, but the TB, as used in these surveys, had a 10 mm mesh liner. Both gears were towed for approximately 15 minutes at 2–3 knots. Across all deployments in both geographical regions, the average (± 1 SD) seabed area swept by the SEL was 929 ± 137 m² and for the TB was 2065 ± 479 m².

During the voyages (TAN0705 and TAN0707), initial plans to use SEL as the standard sampling method for collection of mega-epifauna at all sites were modified. The paucity of fauna and small catch sizes recovered with this gear at some sites led to the TB being used instead of the SEL at 4 sites on the Challenger Plateau and 17 sites on the Chatham Rise (Figure 2 and Figure 3). Because of the differences in swept area and mesh size, this change in gear types during the survey has consequences for analysis of the data. The greater seabed area sampled by the TB and its smaller mesh size (10 mm versus 25 mm as configured for these surveys) mean it is likely to have captured a greater number and diversity of organisms than the SEL. Thus, data from these two sets of samples are not directly comparable. However, the sites at which only the TB was deployed were all *D* class sites, additional to the original sampling plan, and most are either outside the planned survey area (sites DD001, DD003, DD004, DD005) or are part of the additional site transects conducted in the orange roughly spawning box (sites D013–D025 see Sampling design above and Figure 1). Thus, the coverage of sites sampled with the SEL was extensive across both survey regions and only a relatively small proportion of the data would be lost from the core survey if the TB-only sites were to be excluded from analyses (Figure 2, Table 1).

All fauna were sorted to higher level taxonomic groups at sea (class, phylum), preserved appropriately (in ethanol, or formalin, or frozen, depending on taxon), and catalogued in the *Specify* database of the NIWA Invertebrate Collection (NIC). Preserved specimens were then distributed to specialist taxonomists at NIWA and abroad for identification. For most megafaunal groups, identifications were made to species or genus level.

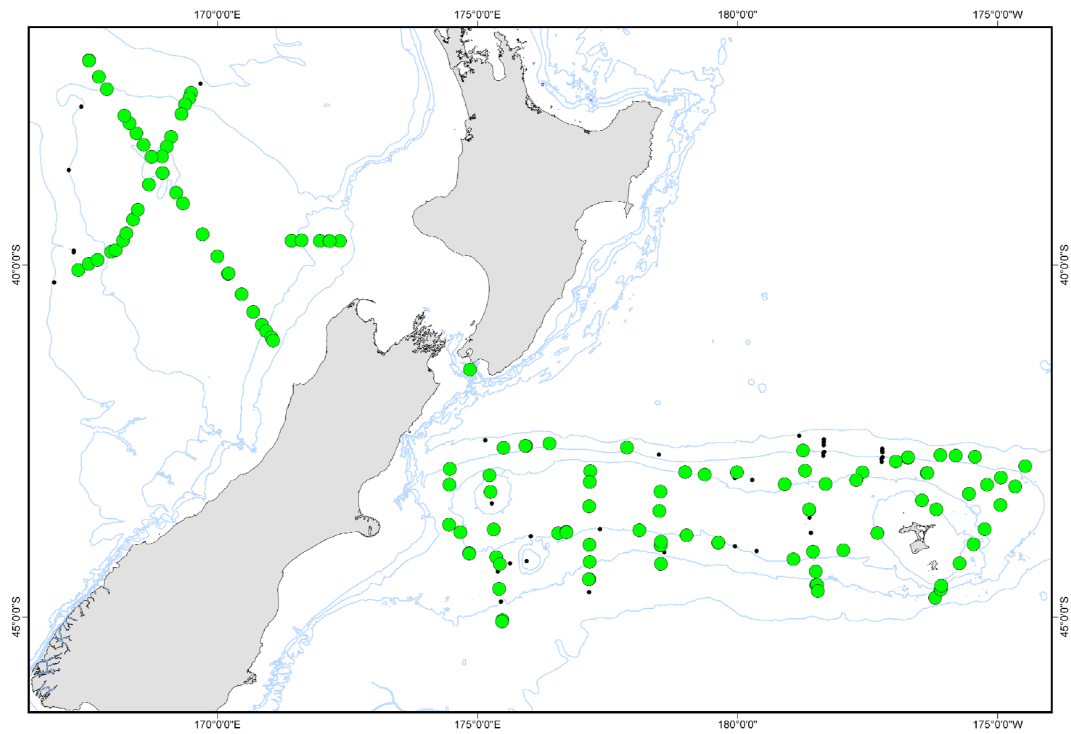


Figure 2: Seamount sled (SEL): sites sampled during the 2007 Ocean Survey 20/20 voyages (filled circles). Black points show sample sites at which SEL was not deployed.

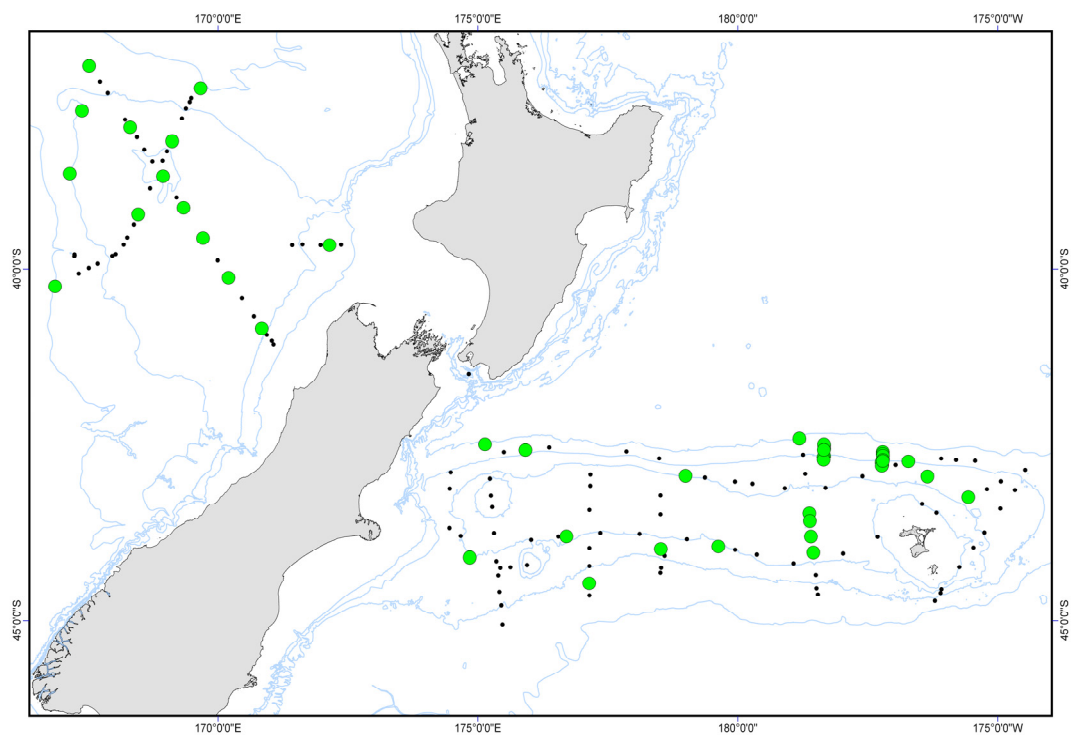


Figure 3: Beam trawl (TB): sites sampled during the 2007 Ocean Survey 20/20 voyages (filled circles). Black points show sample sites at which TB was not deployed.

1.5 Multicorer samples (meiofauna)

The multicorer recovers undisturbed sediment samples of up to 50 cm sediment depth complete with the overlying seawater. This gear was used to sample sediment characteristics, meiofauna (metazoan organisms under 0.5 mm in size), and surficial bacteria. Sediments and bacteria will be reported elsewhere.

The multicorer was deployed at all *A* sites where soft sediments were present: 10 sites on the Chatham Rise and 5 sites on the Challenger Plateau. Multicorer samples were also obtained from 7 *D* sites on Chatham Rise and 1 *C* site on Challenger Plateau (Figure 4, Table 1).

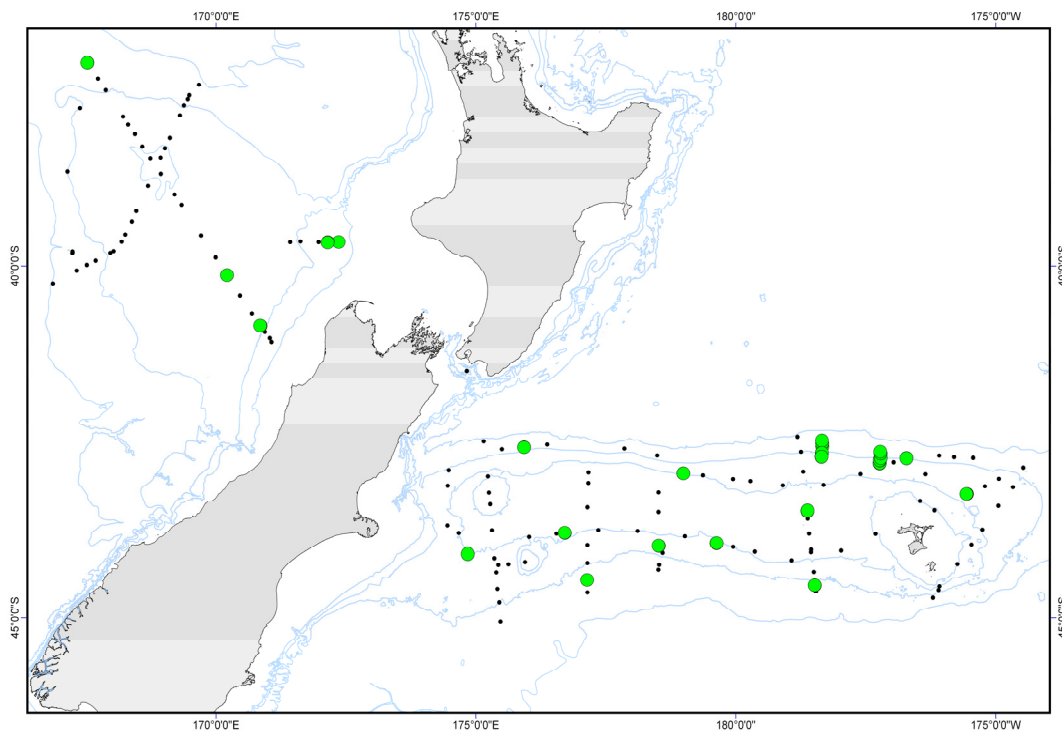


Figure 4: Multicorer (CM): sites sampled during the 2007 Ocean Survey 20/20 voyages (filled circles). Black points show sample sites at which CM was not deployed.

For assessment of meiofauna diversity and abundance, the top 5 cm of sediment from a single core at each site was subsampled using a 5.3 cm² syringe and separated into 0–1 cm and 1–5 cm layers. Samples were preserved at sea in formalin with Rose Bengal dye (to stain body tissues of live organisms). For analysis, samples were sieved to isolate meiofauna in the size range 32 µm to 500 µm and size fractionated on 64, 45, and 32 µm sieves. All specimens were then identified to higher taxonomic groups (phylum, class) and counted.

1.6 Hyperbenthic ('Brenke') sled samples (macro-hyperbenthos)

The Brenke sled (SEH) has two fine mesh nets (0.5 mm mesh) with rigid plastic codend containers (Brenke 2005). The nets are positioned one above the other, so that they sample motile macrofauna from the water layers immediately above (lower net) and about 1 m above (upper net) the seabed ('macro-hyperbenthos'). The mouth width of the sled is 1 m and a lever mechanism ensures that both nets remain closed unless the sled is in contact with the seabed, thus preventing contamination of the sample by planktonic fauna during ascent and descent through the water column.

The SEH was deployed at all *A* sites where flat substrates were present: 11 sites on the Chatham Rise and 5 on the Challenger Plateau. SEH samples were also obtained from 4 other sites on the Chatham Rise (Figure 5, Table 1).

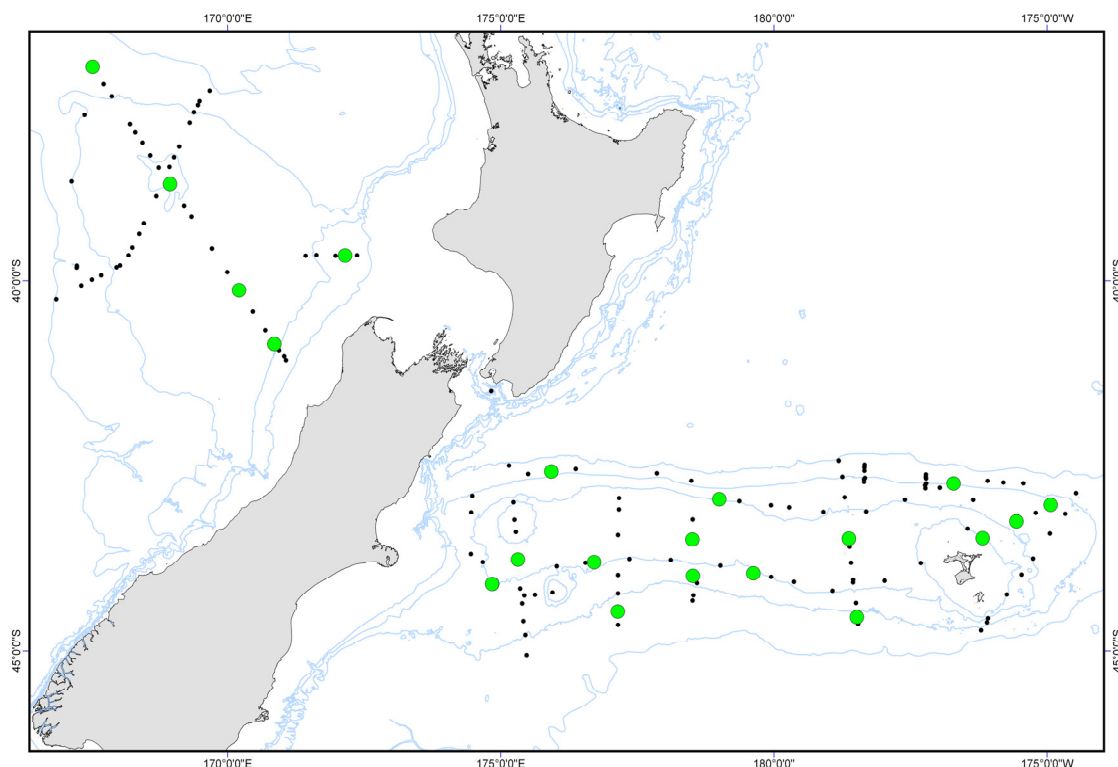


Figure 5: Brenke sled (SEH): sites sampled during the 2007 Ocean Survey 20/20 voyages (filled circles). Black points show sample sites at which SEH was not deployed.

Samples from both nets were preserved, but only those from the upper net were analysed. This is because sample processing is very labour-intensive due to the large numbers of small organisms from groups which are often problematic to identify, and because upper net samples are generally smaller and contain less sediment than those from the lower net. Fauna from these samples were identified to higher taxonomic groups (phylum, class, sub-class) and counted. Data were then standardised to abundances per 1000 m² of seabed.

1.7 DTIS still photographs (macro and mega-epifauna, bioturbation, substrate type)

NIWA's Deep Towed Imaging System (DTIS) take continuous high-definition, digital video (Sony HD1080 format) of the seabed, with high resolution digital still photographs (8 megapixel JPEG format) taken automatically at 15 s intervals along each station transect. For the Chatham Rise and Challenger Plateau voyages, the focal axes of both cameras were oriented vertically downwards. A pair of parallel red lasers spaced 20 cm apart projected on to the seabed enable scaling of the images and video. DTIS was towed at a target altitude of 2–3 m above the seabed at speeds of 0.25–0.5 m s⁻¹ and for a standard deployment time of 1 hour.

DTIS was deployed at all 149 sampling sites (Figure 6). In total, about 35 000 usable still images were collected across the study locations. At target altitude, the seabed area sampled per image is about 2 m². At one image every 15 s, this gives the total seabed area sampled by still photographs per transect of about 480 m².

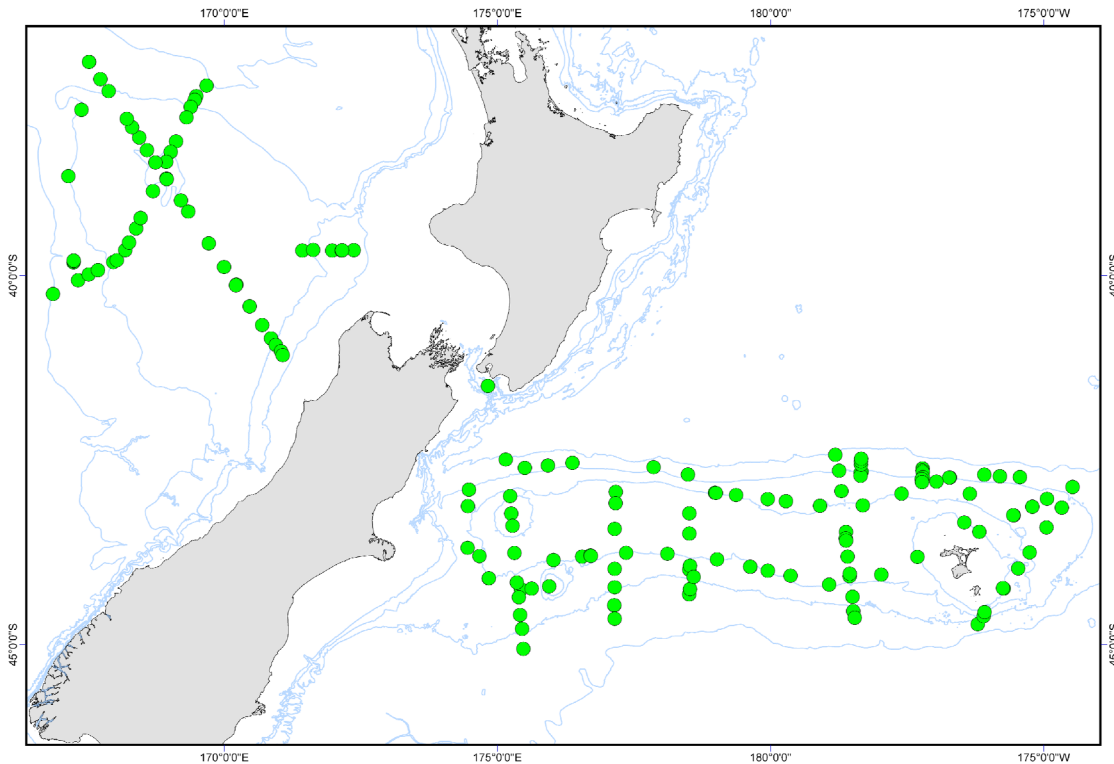


Figure 6: Deep Towed Imaging System (DTIS): sites sampled during the 2007 Ocean Survey 20/20 voyages (filled circles).

Analysis of high resolution still images can yield exceptionally detailed data at small spatial scales (less than 1 m). Such analysis is time-consuming, however, and it was necessary to subsample from the full set of DTIS still images, both in terms of the number of images analysed per site, and the number of sites analysed per location.

1.7.1 Sample selection for DTIS still image analysis

To assess the number of images that could be analysed, a pilot study was undertaken in which 20–40 images from selected sites were analysed in detail for fauna and substrate. The time taken per image was recorded and taxon accumulation curves were plotted to assess the rate of addition of new taxa with increasing number of images analysed (Figure 7).

The rate of accumulation of taxa within a single transect is affected by the level of habitat heterogeneity. Because the probability of encountering new habitats and new taxa increases with increasing area sampled, particularly along a linear transect, it is also unlikely that DTIS transects will reach a true asymptote. Despite this, the pilot study indicated that analysing about 20 images per transect would capture much of the total megafaunal diversity on homogeneous muddy sediment substrata, but to assess the diversity on more heterogeneous substrata higher levels of sampling would be required (Figure 7). As the original observation logs from DTIS deployments showed that more than 90% of sites were on uniform muddy seabed, the number of images analysed was standardised at 24 images per transect. At this level of sampling, it was possible to analyse 55 transects in total.

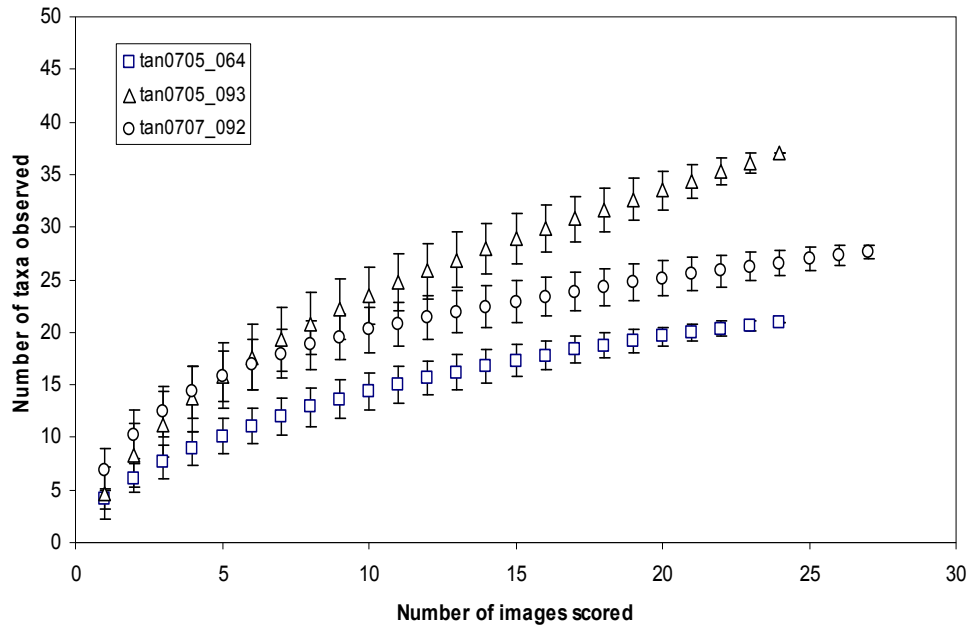


Figure 7: Taxon accumulation curves for still images scored along 3 example DTIS seabed transects. Transect TAN07050_064 is from site A004 on Chatham Rise, substrate >99% muddy sediments. TAN0707_092 is from site CC100 on Challenger Plateau, substrate 100% muddy sediments. TAN0705_093 is from site A041 on Chatham Rise, substrate 64% sand, 14% muddy sediments, 9% gravel, 8% boulders, 2% shell hash, 1% cobbles. The plotted number of new taxa per image is the mean calculated from 999 random permutations of sample (image) order. Error bars show ± 1 SD.

Within each selected transect, every tenth image was analysed provided that: (1) it was correctly exposed, (2) both scaling lasers were visible, and (3) no part of the image was obscured (e.g., by suspended sediment). In cases where the tenth image in sequence was rejected, the next usable image along the transect was selected.

1.7.2 Taxonomic identifications from seabed images

In order to make reliable and consistent identifications from DTIS photographs and video, it was necessary first to generate a reference library of identification images. All DTIS photographs from all transects were first searched for clearly recognisable images of benthic taxa, bioturbation marks, and substrate types present and identifiable at the image resolution (effectively larger than 20 mm). Example images of each taxon were then cropped, colour-corrected, scaled by reference to laser points in the images, and grouped by higher taxa (phylum, class). Specialist taxonomists for each group then assigned identifications for each taxon at the lowest practicable taxonomic resolutions. Because these identifications were across a range of taxonomic levels, including, in some instances, just numbered putative species (e.g., “Corallimorpharia 1”), they are referred to here as operational taxonomic units (OTUs). The result was a reference library of digital images in which all organisms seen in the DTIS transects were represented. This library ensured consistency of identifications between different analysts and between transects.

1.7.3 Image analysis protocols

Data extraction from DTIS images was conducted using the image analysis application *ImageJ* (<http://rsbweb.nih.gov/ij/>). Images were first corrected for colour balance. The seabed area was then calculated by reference to the two parallel laser points (20 cm separation) projected onto the seabed from the DTIS. The proportions of different substrate types visible in the image were measured by either: (1) drawing polygons where distinctions between substrates were clear, or (2) where substrates

were mixed, estimation with reference to an overlay of polygon templates of known areas. Substrates were classified by reference to a table of 39 descriptors, ranging from bedrock to muddy sediments (Table A4) and recorded as percentage of the total image area. All benthic megafauna and bioturbation marks (Table A5) were counted using the *Point selection tool* in *ImageJ*. To avoid over estimation of abundances through edge effects (i.e., if individual organisms intersect the frame edge, counting ‘half organisms’ at different edges effectively includes more individuals than are present in the measured image area), organisms intersecting the edge of the image frame were counted only at the top or left edge of the image.

This resulted in four classes of observation:

- (1) Mobile megafauna (larger than 2 cm), as individuals per m² of seabed
- (2) Sessile megafauna (larger than 2 cm), as individuals per m² of seabed
- (3) Bioturbation marks (tracks, mounds, burrows, etc.) as numbers per m² of seabed
- (4) Substrate type, as percent area of the image for substrate classes

The distinction between motile and sessile epifauna was driven by the observation that many sessile organisms are clonal and that counts of colonies may not be ecologically equivalent to counts of unitary organisms. However, because both data sets are in the same units, they may be combined in later analyses.

1.8 DTIS video transects (mega-epifauna, bioturbation, substrates)

DTIS video transects were captured at all 149 sampling sites across the Chatham Rise and the Challenger Plateau. Each transect was of 1 h duration and across all deployments the average (± 1 SD) swept area of seabed per transect was 2359 ± 644 m².

Because the continuous video samples from the DTIS camera cover a considerably greater area of seabed than the still images and can be analysed for megafauna more rapidly, they were given higher priority in analyses. There are obviously trade-offs associated with this decision. Working from video enables analysis of the full length of each transect and thus captures data on rarer mega-epibenthic species that are likely to be missed when sampling discontinuous still frames. The principal drawback, however, is that the moving video image is at lower resolution than the still images and, in consequence, smaller benthic fauna (under 5 cm) are under-represented in the data, particularly when they are present in abundances that are too high to allow counting of all individuals in the transect.

1.8.1 Video analysis protocols

Post-voyage analyses of video transects were run using the software Ocean Floor Observation Protocol (OFOP; <http://ofop.texel.com>). This software enables analysts to record spatially referenced data on the occurrence of objects or events seen in video by clicking on pre-defined observation types in a graphical user interface. Raw position data for the DTIS (from the ultra-short baseline acoustic tracking system on *RV Tangaroa*) were first smoothed using an adaptive running median and splined along with all associated metadata (e.g., time, depth, heading) to yield corrected seabed tracks with position coordinates and metadata values at 1 s intervals. The digital video files (in *.avi format) were then synchronised with the corrected position files to enable re-running of the transects in the laboratory, with full video playback control (playback speed, reverse, freeze-frame, etc.) and precise spatial and temporal logging of events. The identities and abundances of mega-epifauna (larger than 5 cm) together with substrate descriptors were logged over the full length of each transect.

The output from OFOP is in the form of a list of spatially referenced point observations for each transect. For fauna, this makes immediate sense, because each individual is represented as a single event and total abundance is simply the sum of all individual observations of that taxon. Substrata, however, are continuous rather than point data. To generate continuous substrate records, all video

data were loaded into a PostgreSQL-postGIS database incorporating scripts to separate substrate and fauna observations into different database fields, and then fill down substrate observations at 1 s intervals throughout each transect. The fill-down script was of the form; “if $t_{2\text{-original}}$ is null, make $t_{2\text{-fill}} = t_1$, and if $t_{2\text{-original}}$ is not null, make $t_{2\text{-fill}} = t_2$ ” (where t_x is a value entered in the database relating to time x). Because analysts recorded each substratum type as it occurred along the transects, even for highly heterogeneous transects, this yielded a detailed, continuous record of the main substrate types for all transects.

The resulting substrate data were extracted from the database as percentages for each substrate type in each transect. Abundances of fauna and bioturbation features were recorded as counts of individuals or colonies per transect and then standardised to number per 1000 m² of transect. Thus, although data were recorded in a form that would allow analysis of spatial pattern within transects, for first analyses they were summarised to give an integrated, point-sample measurement of taxon richness and substrate composition per transect.

RESULTS AND DISCUSSION

1.9 Mega-epibenthos

In total, 4200 individual specimens were identified from the SEL and TB samples, representing 867 taxa, most at species level, from 15 phyla (Table A1). The most abundant and widespread phylum caught by both gears was Echinodermata. Crustacea, Mollusca, Annelida, and Cnidaria were also common, but the proportions of these taxa caught varied between gear types. In particular, Annelida were more abundant in SEL samples, with Echinodermata, Crustacea, Mollusca, and Foraminifera being more abundant in TB samples (Figure 8). Given the differences between gear types in the total number of stations (SEL, 130 stations; TB, 45 stations) and sites (SEL, 111 sites; TB, 42 sites) at which each gear was deployed, it is perhaps surprising that overall the TB collected more specimens of these taxa than the SEL. This is likely to be a function of the greater seabed area swept by the TB per deployment, which resulted in broadly similar total swept areas per gear type (SEL 120 000 m²; TB 90 000 m²), combined with its smaller mesh size.

For both Echinodermata and Crustacea, across all samples, more taxa were identified from TB samples than from SEL samples (Figure 9 and 10). The opposite was true for Annelida (all of which are Polychaeta, Table A2) and Porifera, of which substantially more taxa were captured by the SEL than the TB. There are several potential explanations for these differences. For instance, the greater incidence of Porifera in SEL samples is likely to be because this gear was deployed on hard and heterogeneous substrata, where sponges are more common, whereas the TB was used only on level, homogeneous substrates. Similarly, the SEL, being heavier than the TB, tends to dig in to the sediment more and thus might sample infaunal polychaetes more effectively. However, with the data available here – and particularly bearing in mind the lack of matched SEL and TB samples from the same sites – all that we can reliably conclude is that direct comparisons between data from these two gears are problematic.

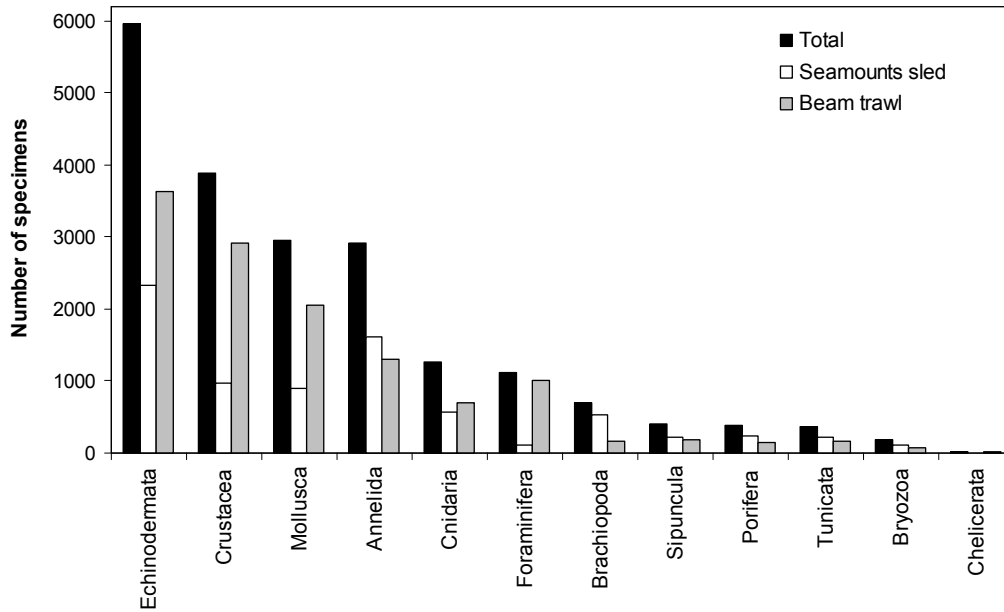


Figure 8: Epibenthic megafauna collected and identified from the seamounts sled (SEL) and beam trawl (TB): total number of specimens per higher taxonomic group (phylum, superclass) collected by each gear type and in total.

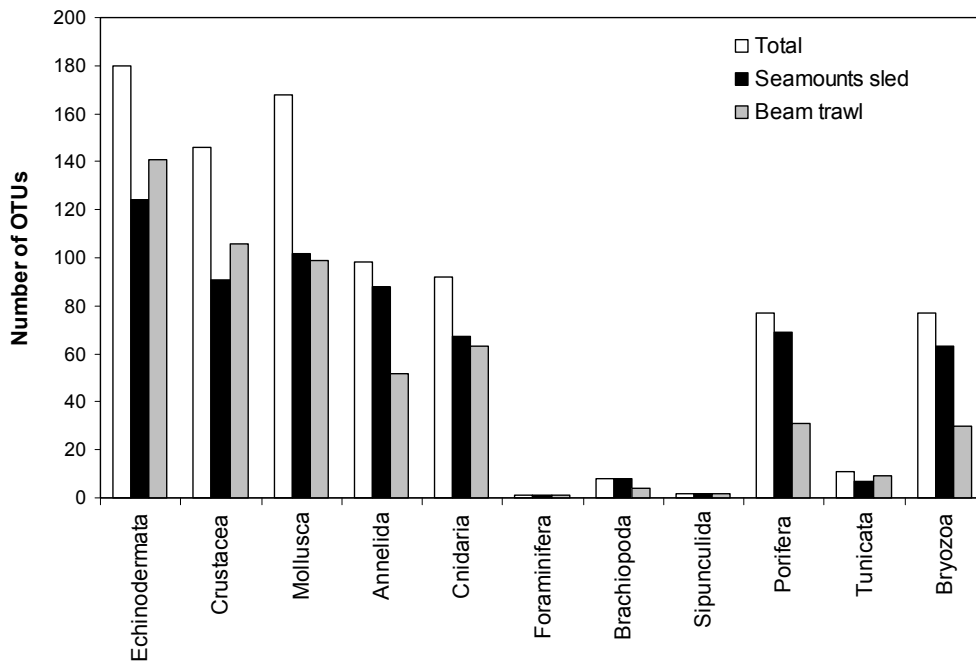


Figure 9: Epibenthic megafauna collected and identified from the Seamounts sled and beam trawl: number of taxa (operational taxonomic units – OTUs – mostly species level) per higher taxonomic group (phylum, superclass – showing the 11 most abundant taxa only).

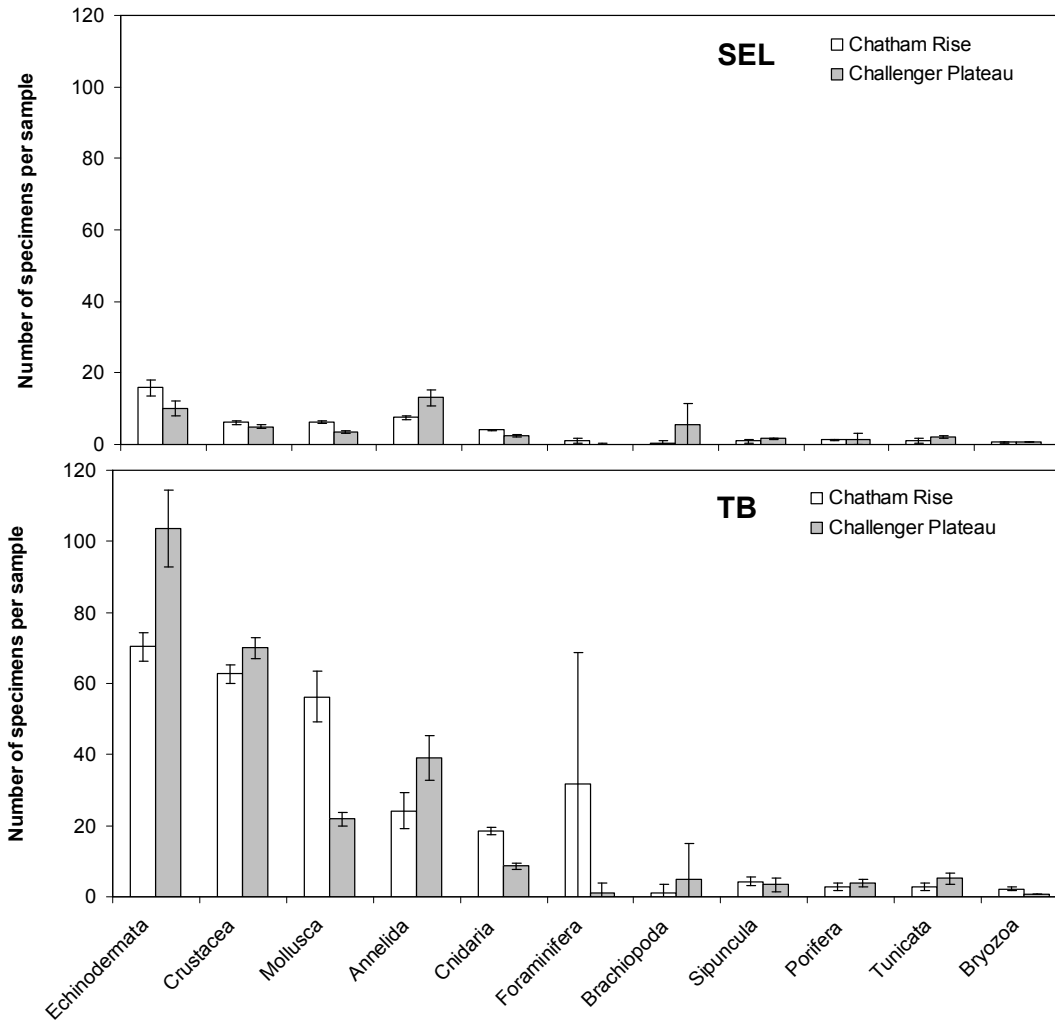


Figure 10: Number of specimens per sample for the seamounts sled (SEL, upper panel) and beam trawl (TB, lower panel) across Chatham Rise and Challenger Plateau. Values are means of n=109 (Chatham Rise) and n=31 (Challenger Plateau) stations. Error bars ± 1 se.

1.9.1 Biomass estimation

In the initial goals of the project, it was planned to derive biomass estimates for epifauna using the wet weights and measurements of identified specimens combined with measurements and abundance counts made from the DTIS video. In practice, however, both aspects of this work proved to be problematic. This is primarily because there are inherent inaccuracies in relating measurements taken from live organisms in the images to those taken from preserved specimens.

First, when they are preserved, soft-bodied taxa, such as holothurians and anemones, contract and, as a consequence, measurements cannot be related directly to length or width measurements taken from live animals in the seabed images. Second, because of varying orientations and degrees of cover by sediment, rocks, or shadows of animals in seabed images, it is often not possible to consistently take the same measurements *in situ* as are taken from the preserved specimens. Third, for infaunal taxa, such as heart-urchins, burrowing anemones, sabellid polychaetes, and bivalve molluscs, there is often no measurement from the image that can be compared directly with the preserved specimen.

To illustrate the first of these points, Figure 11 shows weight-length relationships for preserved specimens of two commonly caught megafauna species: the commensal anthozoan *Epizoanthus* sp. and the holothurian *Molpadia musculus* which have different responses to preservation. *Epizoanthus*

sp. cannot contract significantly when preserved because it grows commensally on the shells of pagurid crabs and consequently the length-weight relationship is strong ($R^2 = 0.89$). By contrast, *M. musculus* contracts when preserved and the length-weight relationship is poor ($R^2 = 0.5$). For these reasons, biomass estimates derived from image measurement methods are likely to incorporate unquantifiable errors.

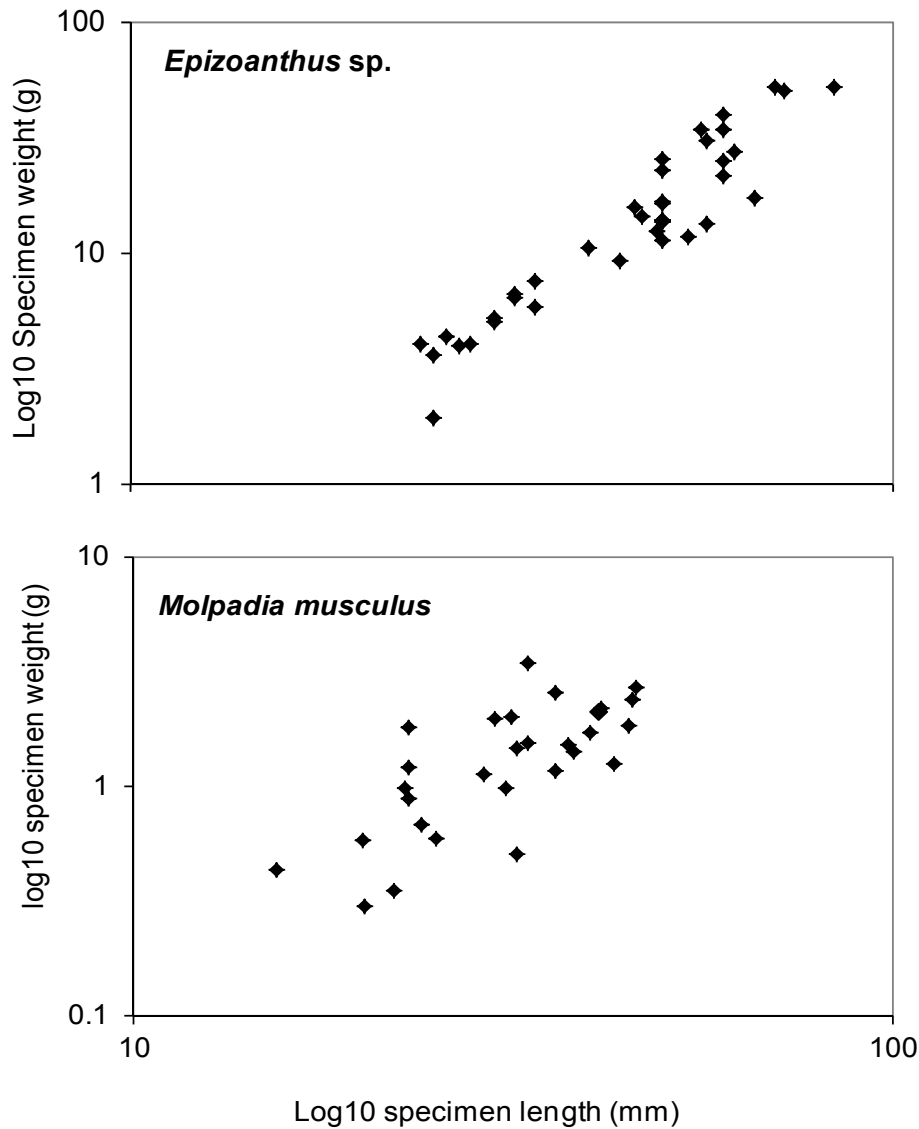


Figure 11: Example length-weight relationships for preserved megafaunal specimens. Top, the anthozoan *Epizoanthus sp.* cannot contract significantly when preserved because it grows commensally on the shells of pagurid crabs and consequently the length-weight relationship is clear ($R^2 = 0.89$). Bottom, the holothurian *Molpadia musculus* contracts when preserved and the length-weight relationship is poor ($R^2 = 0.5$).

Some interesting patterns emerged from the biomass data, however, which merit further investigation. For example, the common ophiuroid *Ophiomusium lymani* shows a strong relationship between disc diameter and body weight (Figure 12), but size frequency distributions differ markedly between the two study locations, with unimodal distribution on the Chatham Rise and bimodal distribution and a higher proportion of small individuals on the Challenger Plateau (Figure 13). While the generally smaller body size of this species on the Challenger Plateau could result from lower food availability than on the Chatham Rise, the bimodal distribution on Challenger Plateau also suggests the presence of two distinct cohorts there. If this is the case, it is likely that any overall biomass differences between

the two locations could change considerably with time and thus that measured patterns may not represent consistent differences in benthic productivity between locations.

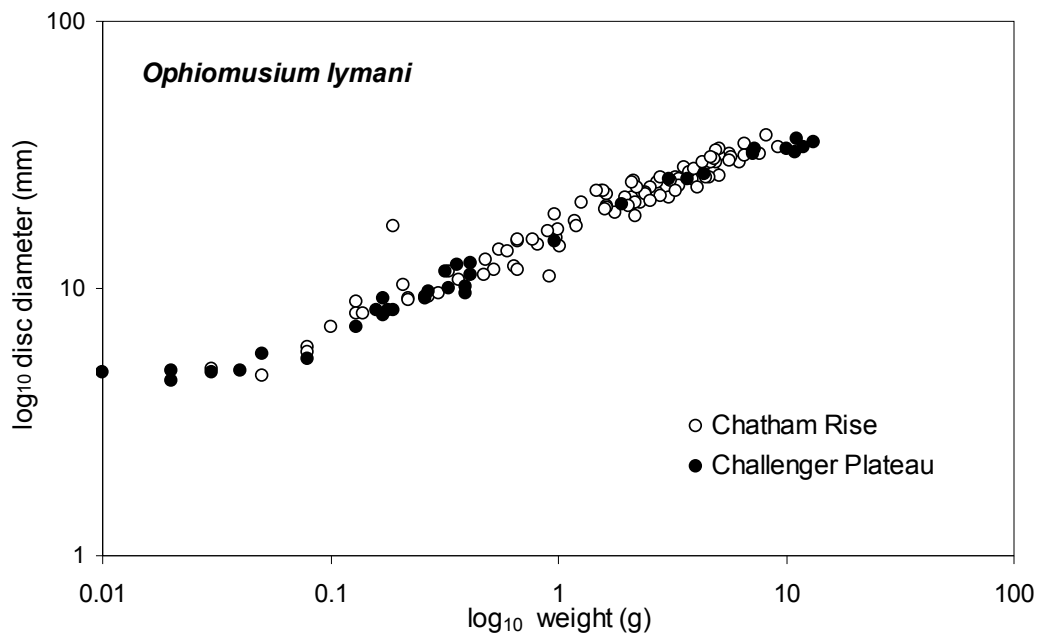


Figure 12: Size to weight relationship for the ophiuroid *Ophiomusium lymani* on Challenger Plateau and Chatham Rise.

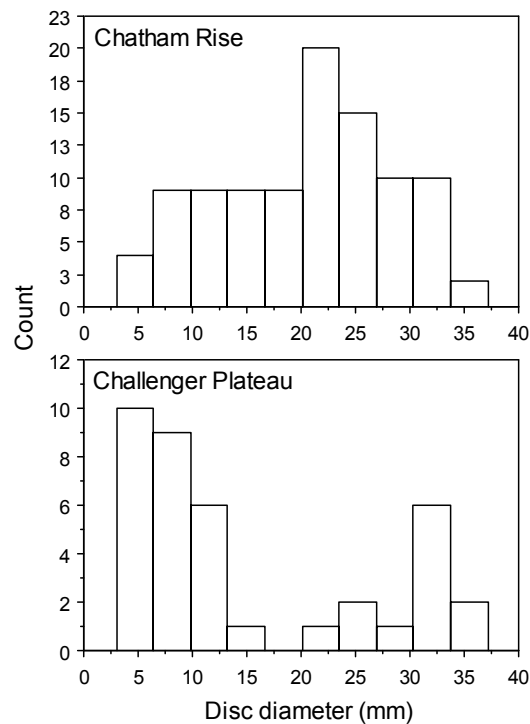


Figure 13: Size frequency histograms for the ophiuroid *Ophiomusium lymani* on Chatham Rise and Challenger Plateau (note different y-axis scales).

1.10 Meiofauna

Multicore samples were processed from 10 *A sites* and 7 *D sites* on the Chatham Rise and 5 *A sites* and 1 *C site* on the Challenger Plateau. More than 11 000 individual organisms representing 18 taxonomic groups (phylum, class, sub-class) were counted and identified. Nematode worms were an order of magnitude more abundant than any other taxon in all samples (Figure 14). Pyriform copepods and nauplii larvae of crustaceans, primarily copepods, were also common, but all other taxa occurred in very low numbers.

The coarse level of taxonomic resolution achieved for meiofauna samples is particularly striking. Although 90% of all specimens were nematode worms, this group was not resolved here beyond the level of phylum. At this level of resolution, therefore, the meiofauna dataset provides limited scope for analyses of spatial patterns of diversity. The abundance data (which are also expressed in the data set as biomass) are, however, potentially important for assessing spatial variations in benthic secondary productivity and will link with data on total sediment oxygen demand collected by the University of Waikato during the OS 2/20 Chatham-Challenger voyages.

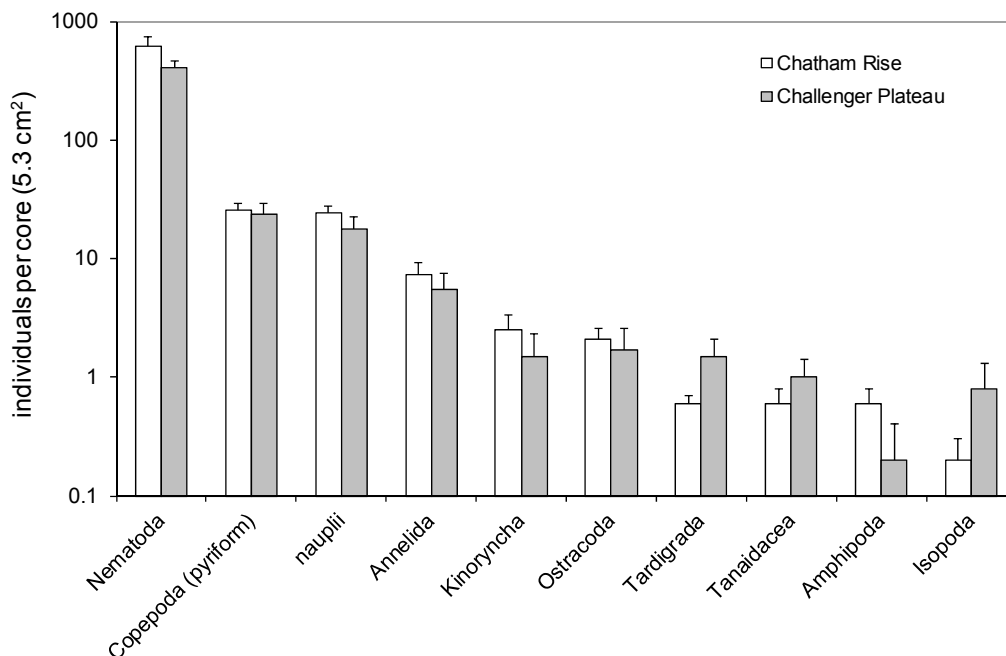


Figure 14: Meiofauna abundances on Chatham Rise (n = 17 sites) and Challenger Plateau (n = 6 sites). Note log scale for abundance and the predominance of Nematoda at both locations. Means +1se.

1.11 Macro-hyperbenthos

More than 79 500 individual invertebrate specimens were identified from the 20 Brenke sled samples (Figure 15, Table A3). Copepod and peracarid crustaceans were the most abundant taxa but at this stage, and at the relatively coarse taxonomic level of the identifications, the only obvious difference between Chatham Rise and Challenger Plateau samples is the greater overall abundance of Tanaidacea on the Challenger Plateau.

The hyperbenthos of the continental shelf and slope had not been sampled directly in New Zealand before this study and the high abundance of crustaceans recorded here suggests that this component of the seabed fauna may play a significant role in benthic ecosystem processes. There is considerable potential for future research using these samples, including assessment of species-level diversity and elucidation of their trophic relationships, particularly with respect to their role in the processing of

organic detrital material at the seabed and their potential significance as prey items for demersal fish species. As with the meiofauna, however, the coarse level of identification achieved to date, combined with the limited spatial coverage of sampling (Table A1, Figure 5) render these data of less use for analyses of biodiversity in the OS 20/20 project than the more extensive data from other gear types

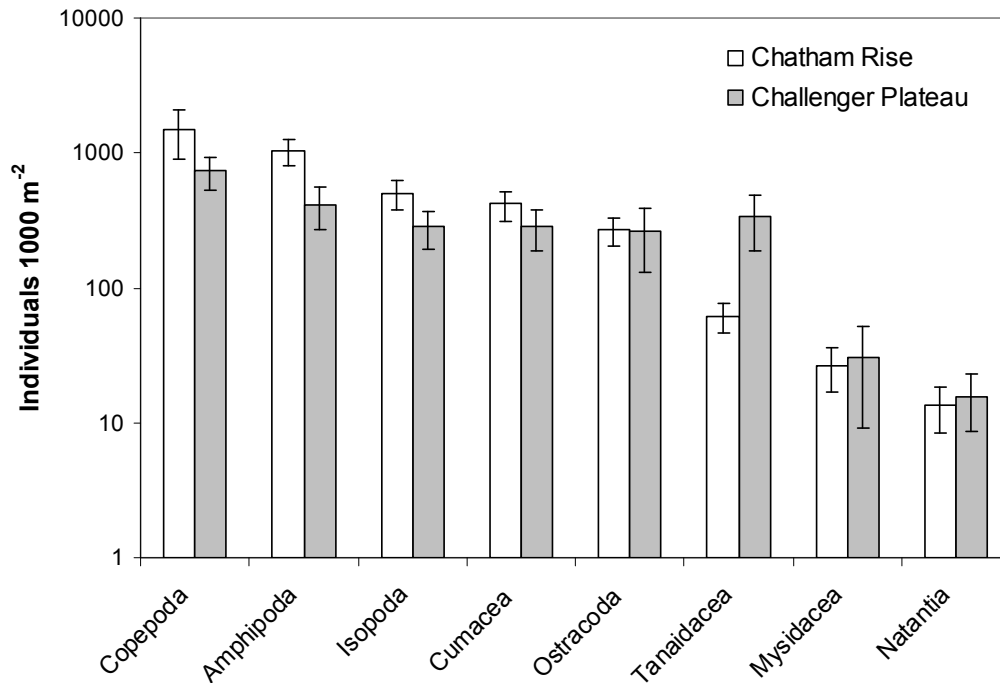


Figure 15: Abundance of macrofaunal hyperbenthic crustacean taxa in Brenke sled upper net samples from Chatham Rise (n = 16 sites) and Challenger Plateau (n = 5 sites). Means \pm 1se. Note log scale on y axis.

1.12 Macro and mega-epifauna, bioturbation, and substrates from DTIS still photographs

A total of 1275 still images were analysed from 55 DTIS transects representing 45 sites: 32 on the Chatham Rise and 13 on the Challenger Plateau. Data were extracted from 10 to 25 (mean 22.9, SE 0.35) still images per transect. At least one transect from each site (16 sites) was selected as first priority for analysis and those from a further 29 sites were selected to give broad spatial coverage across both Chatham Rise and Challenger Plateau (Figure 16).

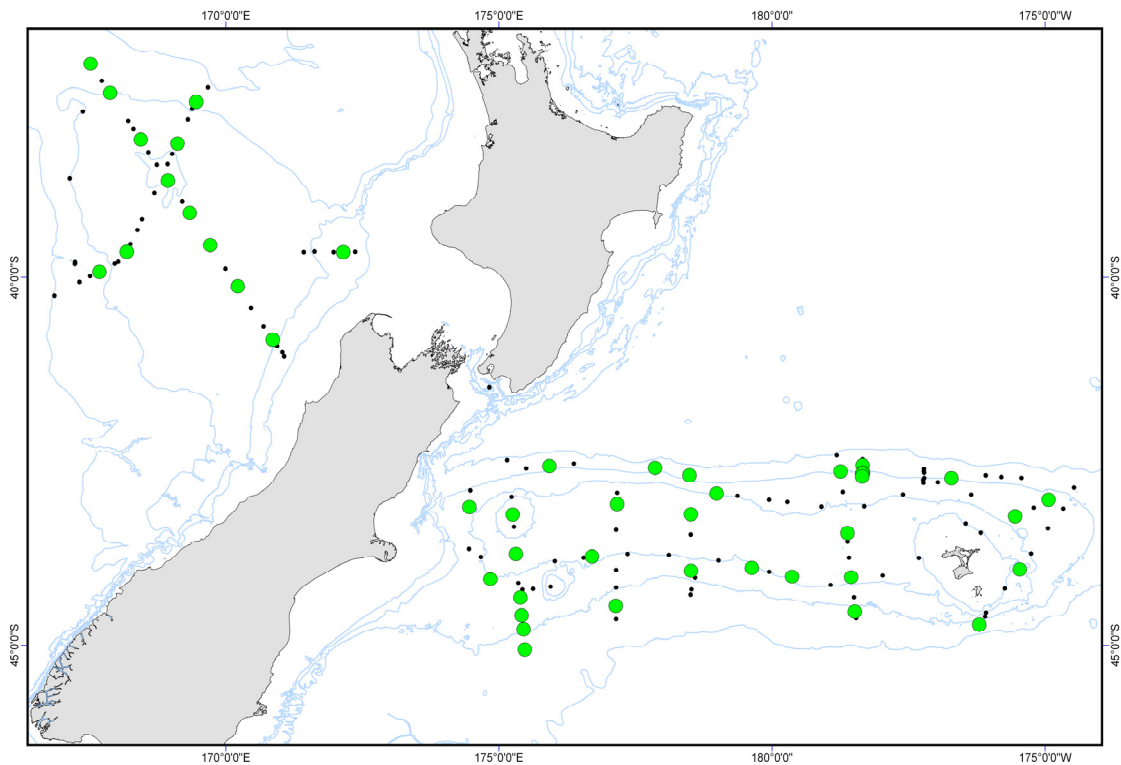


Figure 16: Deep Towed Imaging System (DTIS) still images: filled circles show sites from which still images were analysed for benthic fauna, substrates, and bioturbation marks. Black dots show all sites sampled during the 2007 Ocean Survey 20/20 voyages.

A total of 9428 records of individual organisms, representing 93 motile and 84 sessile taxa (Table A6), were identified from the images. More than 5000 records of bioturbation marks were also compiled, together with substrate classifications for all images (see Table A4 and Table A5 for substrate and bioturbation categories). The most taxonomically rich groups were Cnidaria (48 OTUs) and Echinodermata (43) and for all major groups more taxa were recorded on the Chatham Rise than on the Challenger Plateau (Figure 17). Differences in abundances between locations were less pronounced, with only Mollusca and Porifera being in appreciably higher abundance on the Chatham Rise than the Challenger Plateau. The mean abundance of Foraminifera (Xenophyophora) was highest on the Challenger Plateau, but with very high variance resulting from locally abundant populations at some sites (Figure 18).

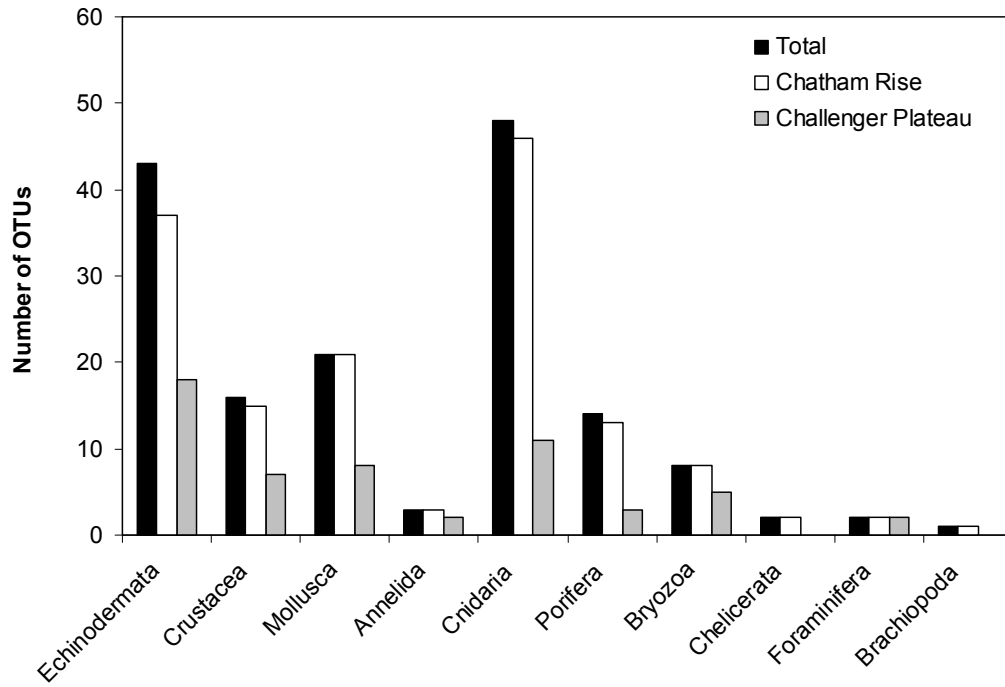


Figure 17: DTIS still images: numbers of taxa (operational taxonomic units – OTUs) recorded across the whole survey, on Chatham Rise, and on Challenger Plateau.

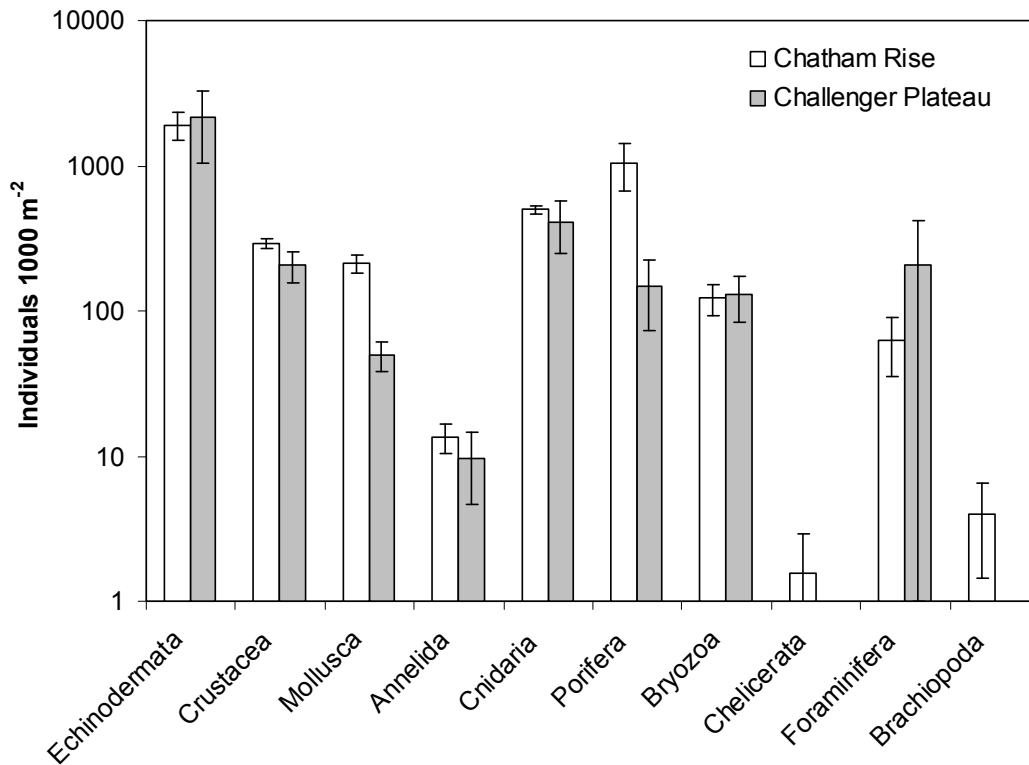


Figure 18: DTIS still images: number of individuals per 1000 m² of seabed. Values are means of n=42 (Chatham Rise) and n=13 (Challenger Plateau) transects. Error bars ±1 SE. (Note log scale for abundance).

1.13 Mega-epifauna, bioturbation, and substrates from DTIS video transects

The full length of at least one DTIS video transect was analysed from each of the planned sampling sites across the Chatham Rise and the Challenger Plateau (Figure 19). At all *A* sites across both locations, the two replicate transects were both analysed. Transects from another seven sites, all of which were additions to the sampling plan, were not analysed: six *D* sites on the Challenger Plateau, five of which were either outside the New Zealand EEZ or not on the original multibeam transects, and one extra *B* site on the eastern end of the Chatham Rise (Figure 19). This is the most comprehensive coverage of the study area of any of the seabed sampling gears, both in terms of the number of sites and the area of seabed sampled.

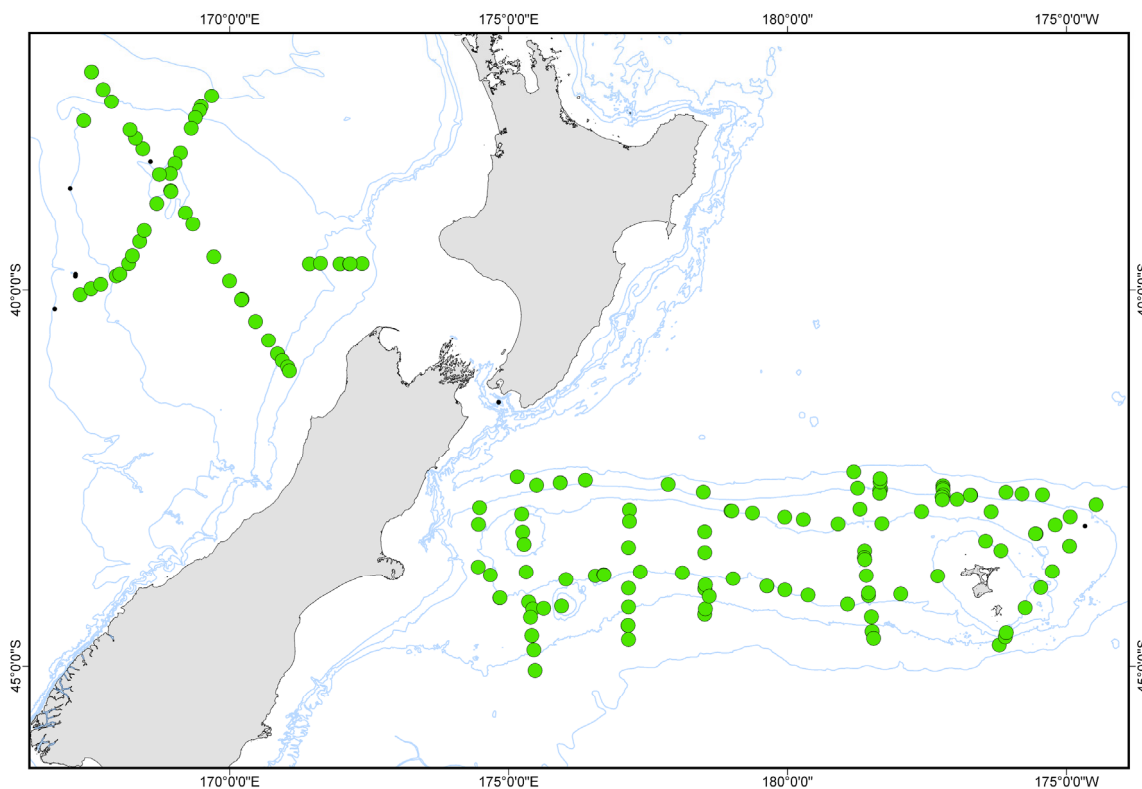


Figure 19: Deep Towed Imaging System (DTIS) video: filled circles show sites from which video transects were analysed for benthic fauna, substrates, and bioturbation marks. Black dots show sites from which video was not analysed.

A total of 55 066 records of individual organisms, representing 318 OTUs, were made from the DTIS video data, together with 8225 records of bioturbation features. Substrate type was recorded throughout each transect (see Methods). The most taxonomically rich group in video transects was Echinodermata (75 OTUs), followed by Cnidaria (38) and Arthropoda (34) (Figure 20). For all phyla, more taxa were recorded from the Chatham Rise than from the Challenger Plateau. This is likely to be a consequence of the greater seabed area sampled on the Chatham Rise, and the greater range of habitats encountered there. Echinodermata was also the most abundant phylum across all sites and both locations. At the phylum level, mean abundances of mega-epifauna recorded from video were not significantly different between locations except for Polychaeta and Bryozoa, which were in higher abundance on the Chatham Rise than the Challenger Plateau (ANOVA on \log_{10} transformed data, $P < 0.05$) (Figure 21).

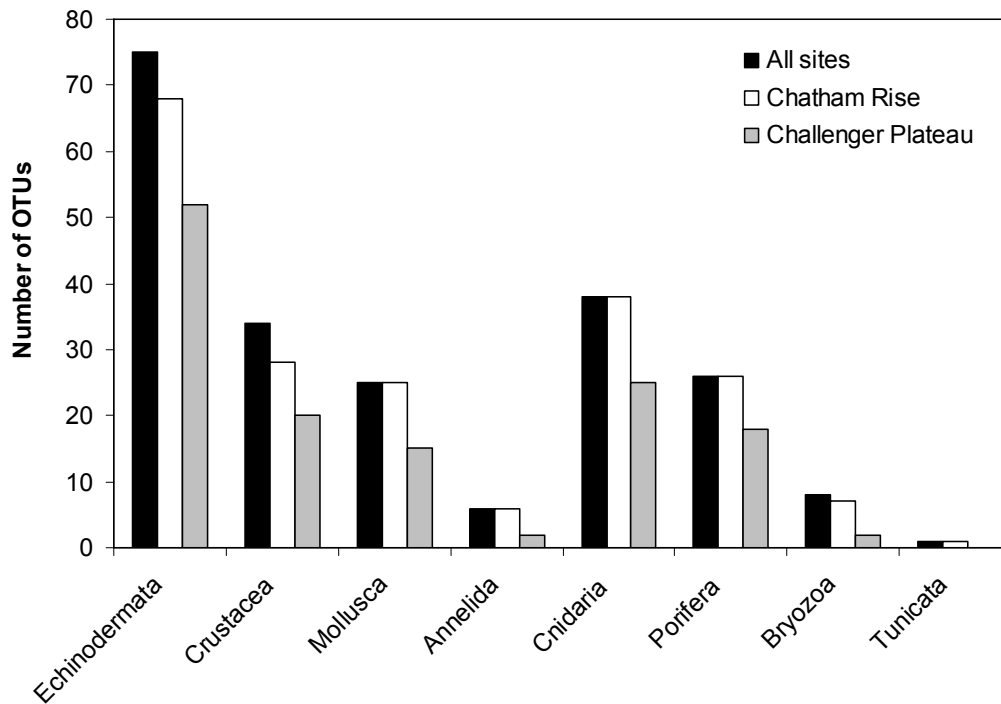


Figure 20: DTIS video transects: number of taxa (operational taxonomic units – OTUs) recorded across the whole survey (All sites) and in each of the study locations (Chatham Rise and Challenger Plateau).

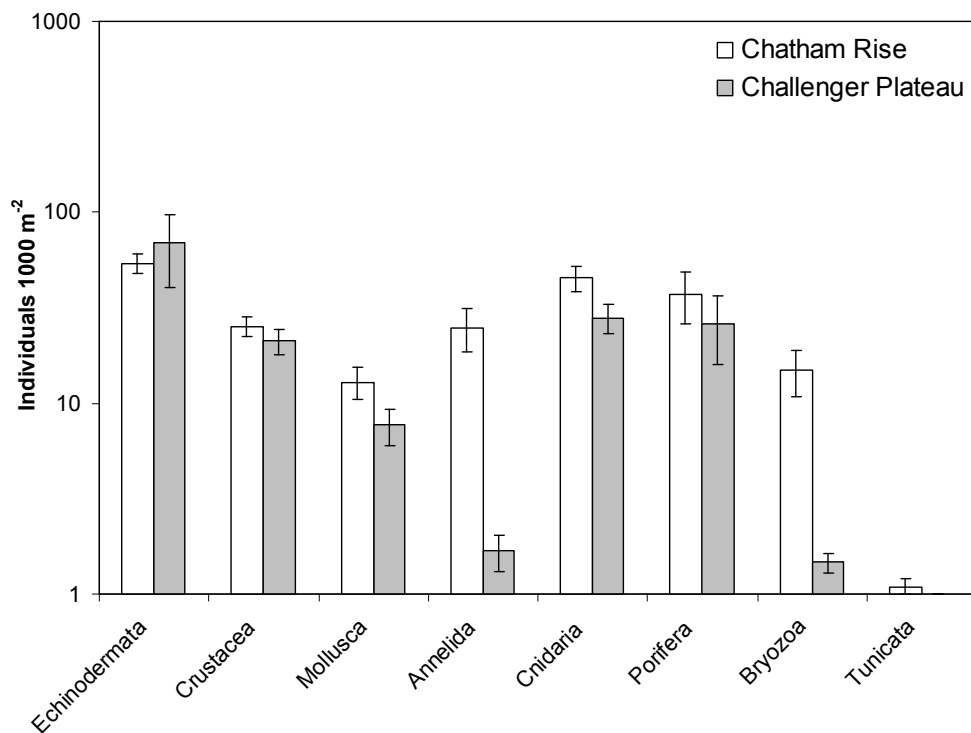


Figure 21: DTIS video transects: mean number of individuals per 1000 m² of seabed for the principal phyla recorded. Values are means of n=108 (Chatham Rise) and n=46 (Challenger Plateau) transects. Error bars 1 SE. Note log scale for abundances.

As noted in the Methods, the principal drawbacks to the video data are that identifications are generally to a coarser taxonomic level than for other megafaunal sampling gears (SEL, TB, DTIS still images) and that smaller organisms (under ca. 50 mm) are not counted. This latter point was important in relation to small-bodied, highly abundant taxa, such as the brittle star *Ophiomusium lymani* and the

quill worm *Hyalinoecia* spp., which were impracticable to count from the video (e.g., mean abundances for *O. lymani* were over 10 individuals m⁻² in some transects). To generate complete data sets for these taxa in video transects, counts were made from still images. Thus, if a taxon was observed to be present in high abundance in sections of a video transect, this was noted in the video analysis file and representative still images were analysed to calculate a mean abundance per square metre. This value was then used to generate an estimate of the total abundance of the taxon along the transect and this was then appended to the original video analysis output. This estimation of abundant, small, taxa was necessary in only 11 transects, and in all cases 8–10 images were sufficient to generate acceptably precise estimates of population density.

For medium- to large-scale habitat mapping, the video transect data have been amalgamated to derive values per transect (abundances summarised as numbers of individuals per unit area (1000 m²) and substrate types summarised as percentages of the overall transect, e.g., 75% muddy sediments, 25% cobbles). However, because each observation is referenced to a position along the transect, the data contain a great deal more spatial information which affords considerable potential for finer scale ecological analyses in future. Thus, the spatial extent of substrate patches and the distribution of organisms in relation to them are inherent in the data (Figure 22 and Figure 23) and could be used, for instance, to examine fine-scale habitat associations, commensal interactions, and spatial aggregation of populations.

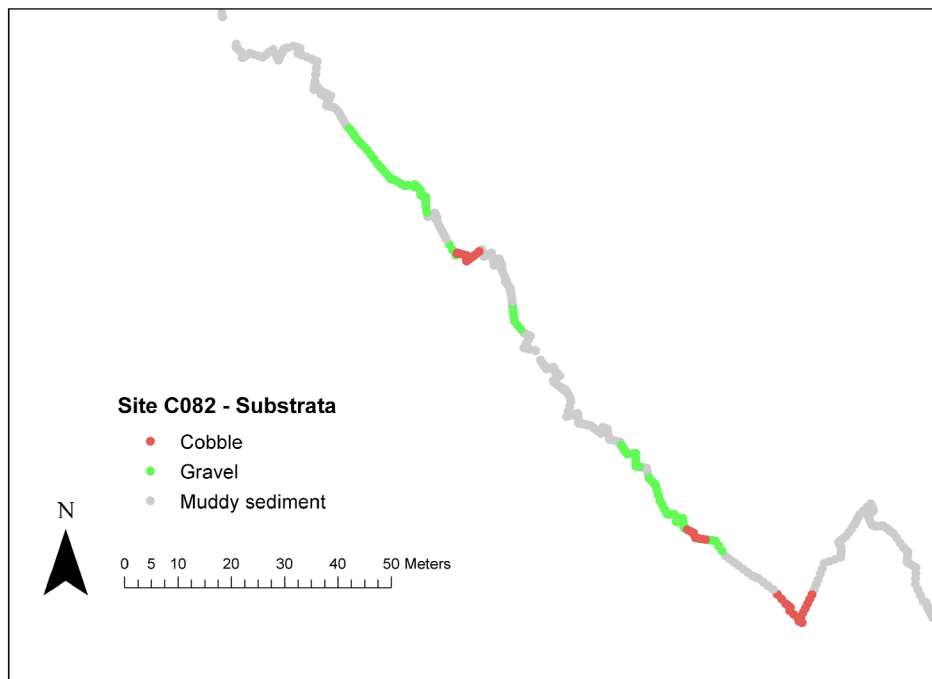


Figure 22: Within-transect DTIS video detail: example of continuous substrate description throughout transect at Site C082 on Chatham Rise.

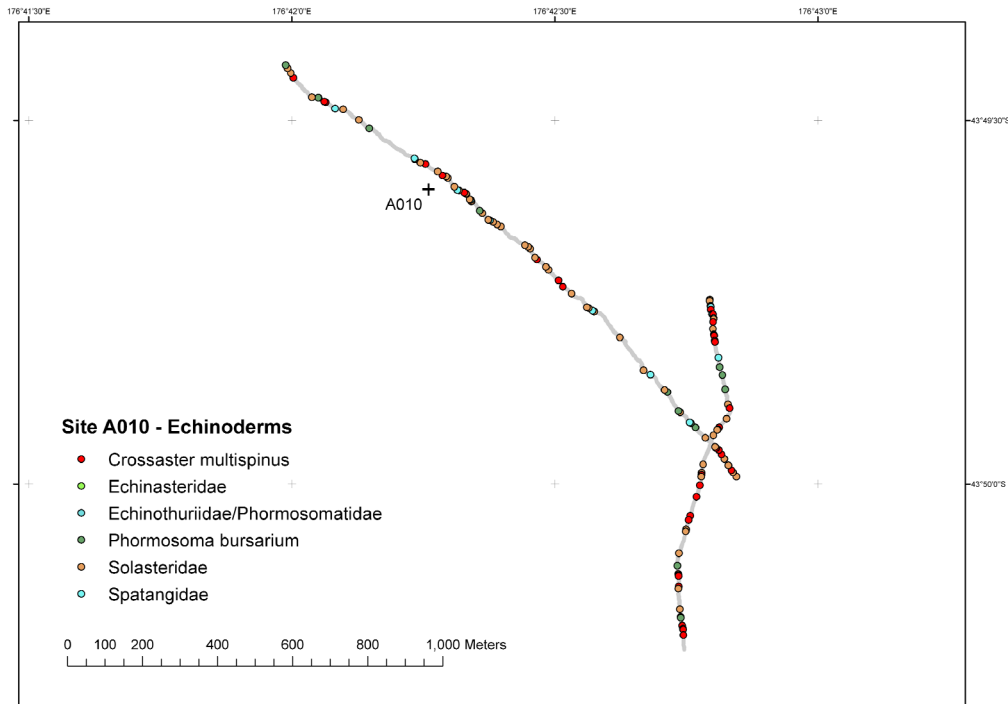


Figure 23: Within-transect DTIS video data detail: example at site A010 (cross marks the nominal site location) on Chatham Rise showing all echinoderm observations against substrate type (grey = muddy sediment).

SUMMARY AND CONCLUSIONS

The Chatham Rise and Challenger Plateau project was the first in the Ocean Survey 20/20 programme and the sampling voyages in 2006 and 2007 were ambitious in terms of the area of seabed covered, the number of gear types deployed, and the range of measurements that were planned from them. A considerable volume of novel, high quality data has been generated from the processing of these samples under the present project and these data will provide the basis for a much-improved understanding of the distribution of benthic biodiversity and habitats across the study area through subsequent analyses. These analyses will centre on validation and testing of the original Marine Environments Classification (MEC) (Snelder et al. 2005, 2006) and its later developments (e.g., Leathwick et al. 2006) but will also include assessment of the utility of multibeam sonar data for predicting seabed habitats and biota at large scales, and the influence of fisheries on benthic habitats.

The sampling strategy and range of gears deployed for the Chatham-Challenger OS 20/20 surveys were decided using the best available information about the survey locations and the types of analyses that were envisaged at the time. Most importantly, the sampling strategy had to balance the conflicting requirements for, on the one hand, broad spatial coverage to adequately characterise variability across the study areas, and on the other, intensive local sampling to provide detail across a broad spectrum of faunal groups and habitats. As with all sampling programmes there were trade-offs involved in these decisions and from this first examination of the samples it is clear that some of the resulting data sets are more useful for addressing the higher-level objectives of the OS 20/20 project than others. These differences are discussed below and result from two principal factors: the number of sites at which each gear type was deployed, and the taxonomic level to which different sample types are routinely identified.

In the present project, most effort was directed to the processing and identification of epibenthic megafauna from the sled and trawl samples. This is the component of the benthos that is most easily

sampled over large spatial scales, and for which taxonomies are most complete and New Zealand's taxonomic expertise is strongest. These megafauna samples also represent the largest proportion of sampling effort at sea and the greatest bulk of samples collected. The samples have yielded taxonomically detailed and spatially extensive data sets that are unparalleled in New Zealand waters and will be central to assessments of biodiversity and subsequent analyses of distributional and functional ecology. Of the two data sets derived here from these gears, the seamounts sled samples afford the most complete spatial coverage across the study area and therefore are of most interest for spatial analyses of biodiversity. The seamounts sled is a versatile tool and was selected as the standard gear for these surveys because it allows sampling on any substrate type. However, because it sweeps a smaller area than the beam trawl and because, in this instance, the beam trawl was fitted with a finer mesh net, it catches fewer animals in total and a more restricted range of taxa. This is particularly apparent on the predominantly soft-sediment seabeds that characterise much of the Chatham Rise and the Challenger Plateau (DTIS video data show only 5 sites out of 149 across both locations where bedrock or boulder substrates constituted more than 10% of the transect). Mega- and macro-epifauna are generally scarcer on soft sediments and thus the greater swept area of the beam trawl is more likely to capture representative samples in these environments. For these reasons, it would be more effective to use the beam trawl as the standard gear for sampling mega-epifauna in continental shelf and deep sea habitats and use the seamounts sled only as necessary on the rarer rocky sites.

The DTIS video and still images provide quantitative data across a range of scales, and analysis of the video in particular was very effective for enumeration of mega-epifaunal taxa and their spatial relationships to substrate type throughout the study area. Analysis of the DTIS video and stills samples was dependent on the provision of identifications from the sled and trawl samples and there is a very effective synergy between these two sampling methods; the physical samples from the sled and trawl providing taxonomic validation for the photographic samples. The resulting data sets afford the potential for detailed analyses of distributions across scales ranging from less than 10 m² within transects to 10–100 km² between sites and locations.

The two data sets that have the least potential for use in broad-scale ecological analyses in their present form are the meiofauna and the hyperbenthic fauna from the multicorer and Brenke sled, respectively. This is because the level of taxonomic resolution that was achieved is too coarse to enable reliable discrimination between samples and because these gears were deployed only at a relatively small subset of sites. The lack of taxonomic resolution is a consequence of long sample sorting times for the very large numbers of small organisms collected, current lack of taxonomic expertise in New Zealand for many of the faunal groups (e.g., nematodes, isopods), and lack of established taxonomies and identification keys. Both these components of the benthos are highly diverse, are of considerable interest taxonomically and, most importantly, are likely to play significant roles in ecosystem functioning (e.g., Hinz et al. 2008, Ingels et al. 2009). Thus, there are strong arguments for expanding research in these fields as long as the resources required are weighed against the objectives of individual projects and available funding.

The initial decision to deploy five different gear types during sampling was a consequence of wanting to sample a very broad spectrum of faunal sizes. With the resources available both during and after the sample collection phase, this decision has had two important effects on the scope and detail of the data sets described here. First, by committing to deploying many gear types, it was impractical to collect replicate samples other than at a small subset of sites (*A* sites) or to deploy all gear types at all sites. Thus the potential for quantification of within-site variability is limited, and for the multicorer, beam trawl, and Brenke sled, spatial coverage of the samples is restricted. This was known at the time of planning and the strategy adopted was a pragmatic and well informed approach designed to ensure broad coverage while enabling some assessment of variability at a small subset of sites. Second, by collecting samples of many different types, from sediments and bacteria to mobile macro- and megafauna, each of which requires specialist skills and taxonomic knowledge to process, the number of samples that could be analysed and the level of taxonomic detail that could be achieved for some faunal components has, to date, been restricted.

From this preliminary assessment of data sets from the first OS 20/20 voyages, and given that the focus is on epifaunal, rather than infaunal, habitats, the sampling gears that provide the most appropriate data at relevant scales for regional biodiversity research and for which taxonomic expertise currently exists in New Zealand are the DTIS camera system, and either the beam trawl or the epibenthic sled. When used together, these gears provide a practical and effective means for collecting broad-scale quantitative information across large areas of the seabed. This combination of gears was central to the Chatham-Challenger OS 20/20 surveys and has generated an extensive, detailed, and consistent data set, which is of direct relevance to the objectives of the OS 20/20 project and will be of considerable value for future ecological analyses of the region.

Sediment characteristics are also fundamental to ecological or geological studies of the seabed and the multicorer samples from the Chatham-Challenger surveys have generated a wide range of data relating to physical sediment characteristics, bacteria, meiofauna, macro-infauna, and biological oxygen demand. The multicorer is a versatile and practical tool but the value of the resulting data for analyses at spatial scales relevant to the OS 20/20 objectives could be considerably enhanced in future by increasing the spatial coverage of sampling: ideally cores would be taken routinely at all sites. Although the fundamental trade-off between the number of gear types deployed and the number of sites sampled will always remain, as will the limitations of processing time, increasing the spatial coverage of the multicorer sampling would provide a wealth of data for future studies and would resolve one of the main short-comings identified with some of the data described here: the lack of spatial coverage.

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Constructive reviews by Scott Nodder, Judi Hewitt, and Mary Livingston improved earlier versions of this report.

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APPENDICES

Table A1: Challenger Plateau (TAN0707) and Chatham Rise (TAN0705); gear deployments at each sampling site. CB; box corer, CM; multicorer, DTIS; camera, SEH; Brenke sled, SEL; seamount sled, TB; beam trawl.

Region	Stratum	Site	longitude	latitude	CB	CM	DTIS	SEH	SEL	TB
Challenger	1	BB100	169.34008	-39.08889			1		1	1
		CC100	169.71519	-39.54406			1		1	1
	2	AA002	168.94528	-38.62290	2	2	2	1	2	1
		BB101	168.69013	-38.80054			1		1	
		CC101	169.20700	-38.92946			1		1	
		CC102	168.73717	-38.38679			1		1	
		CC113	168.93763	-38.37821			1		1	
	3	AA003	167.52723	-36.91874	2	2	2	1	2	1
		BB102	167.72800	-37.17063			1		1	
		BB103	169.46000	-37.48575			1		1	
		CC103	169.30967	-37.73488			1		1	
		CC114	169.49296	-37.41583			1		1	
		CC115	167.87933	-37.35233			1		1	
		DD001	169.67096	-37.27188			1			1
		DD002	169.38154	-37.58433			1		1	
		DD003	167.38596	-37.61967			1			1
		DD004	167.14767	-38.58250			1			1
	4	BB104	168.18763	-39.63946			1		1	
		BB105	168.46872	-39.18319			1		1	1
		CC104	168.38325	-39.32929			1		1	
		CC116	168.25350	-39.53129			1		1	
	5	BB106	168.03967	-39.78179			1		1	
		BB107	167.69313	-39.92596			1		1	
		CC105	167.32363	-40.07071			1		1	
		CC106	167.96425	-39.80763			1		1	
		CC118	167.52217	-39.98383			1		1	
		DD005	166.86713	-40.25804			1			1
		DD007	167.23992	-39.81367			1			
		DD008	167.24342	-39.80033			1			
	6	AA006	172.15342	-39.64543		2	2	1	2	1
BB109		171.62325	-39.63646			1		1		
CC107		171.97413	-39.64463			1		1		
CC117		172.36213	-39.64273	1	1	1		1		
CC120		171.42913	-39.64454			1		1		
DD006		168.58471	-38.20604			1		1		
7	BB110	169.11589	-38.08239			1		1	1	
	BB111	168.31133	-37.87439			1		1	1	
	CC108	168.44367	-38.02508			1		1		
	CC109	169.02450	-38.23183			1		1		
	CC119	168.21504	-37.75725			1		1		
	DD006	168.58471	-38.20604			1		1		
8	AA008	170.85563	-40.87888	2	2	2	1	2	1	
	BB113	170.69142	-40.69229			1		1		
	BB114	171.03929	-41.05963			1		1		
	CC110	171.07163	-41.10971			1		1		
	CC111	170.93854	-40.96983			1		1		
9	AA009	170.21211	-40.12983	2	2	2	1	2	1	

Region	Stratum	Site	longitude	latitude	CB	CM	DTIS	SEH	SEL	TB	
Challenger		BB115	170.46613	-40.43146				1	1		
		CC112	169.99933	-39.87563				1	1		
Chatham	1	A001	179.63038	-43.97924		3	2	1	2	1	
		A016	181.37951	-43.52803		4	2	1	2	1	
		B011	181.07596	-44.21025			1		1		
		B012	177.16375	-43.13079			1		1		
		B013	178.50833	-43.53569			1	1	1		
		B014	182.03142	-44.08208			1		1		
		B015	178.11479	-43.80529			1		1		
		B017	181.45171	-44.10304			1		1		
		B018	179.02263	-43.87883			1		1		
		B043	178.52858	-43.96438						1	
		C011	180.36933	-44.09625				1			
	C012	179.95283	-44.02925				1				
	2	A002	184.44616	-43.29189			3	2	1	2	1
		A010	176.70989	-43.83266	1	2	2	1	2	1	
		B022	184.53883	-44.00042			1		1		
		B023	182.69133	-43.84571			1		1		
		B024	176.55454	-43.84492			1		1		
		B025	175.31467	-43.79714			1	1	1		
		B0X1	185.33738	-43.19238			1		1		
		C021	183.82358	-43.51539			1	1	1		
		C023	184.74879	-43.79008			1		1		
		C024	175.94983	-44.23617			1				
		C025	176.02900	-43.89017			1				
	C026	177.35825	-43.79333			1					
	3	A003	183.28313	-42.78437			3	2	1	3	1
		B031	183.90983	-42.74904				1		1	
		B032	181.25942	-42.68983				1		1	
		B033	177.86788	-42.64479				1		1	
		B034	184.79746	-43.17471				1		1	
		B035	184.19954	-42.76417				1		1	
		B036	185.53308	-42.91050				1		1	
		B037	185.06603	-43.07003				1	1	1	
C031		184.56938	-42.77692				1		1		
C032		183.90496	-44.62663				1		1		
CX1		178.49042	-42.74542				1				
CX2	183.64860	-43.00392				1		2	1		
4	A004	181.51960	-44.56104			2	2	1	2		
	A041	178.52010	-44.01517			3	2	1	2	1	
	B041	181.50600	-44.37550				1		1		
	B043	178.52858	-43.96438				1				
C042	183.79813	-44.73550				1		1			
5	A005	175.92653	-42.62298			2	2	1	2	1	
	B051	175.50325	-42.65221				1		1		
	B052	174.47563	-42.94742				1		1		
	C051	183.04154	-42.84079				1		1		
C052	176.38096	-42.58733				1		1			
6	A006	179.11933	-42.99245			2	2	1	2	1	
	B061	175.25204	-43.26933				1		1		
	B062	178.51746	-43.26588				1		1		
	B063	182.40267	-42.99879				1		1		
	B064	181.28454	-43.10438				1		1		

Region	Stratum	Site	longitude	latitude	CB	CM	DTIS	SEH	SEL	TB
Chatham		B065	175.23413	-43.03663			1		1	
		B066	180.90654	-43.16425			1		1	
		B067	179.37163	-43.02333			1		1	
		C061	174.46142	-43.16983			1		1	
		C062	177.16917	-42.97781	1		1		1	
		C063	181.69354	-43.15971			1		1	
		C064	179.95092	-43.07442			1			
	7	A007	174.83039	-44.10292	2	2	2	2	2	1
		B071	175.43263	-44.27517			1		1	
		B072	175.41725	-44.61392			1		1	
		B073	174.67375	-43.83417			1		1	
		B074	175.35996	-44.18129			1		1	
		B075	177.15146	-44.00596			1		1	
		C072	175.39525	-44.37900			1			
		C074	175.62958	-44.26550			1			
	8	A008	177.14280	-44.48548		2	2	1	2	1
		B081	177.15358	-44.24654			1		1	
		B082	178.52804	-44.27229			1		1	
		B083	175.47667	-45.05339			1		2	
		C081	177.14900	-44.66208			1			
		C082	175.45425	-44.79358			1			
		C083	178.51417	-44.34250			1			
	9	D002	174.82725	-41.53683			1		1	
		D003	178.59625	-44.11125			1			
		D004	181.54350	-44.64679			1		1	
		D005	183.91896	-44.57396			1		1	
		D006	184.26558	-44.26275			2		1	
		D007	185.05508	-43.45479			1		1	
		D008	183.54917	-43.38954			1		1	
	D009	181.45442	-44.06775			1			1	
	D010	181.41129	-43.84246			1			1	
	D011	181.38796	-43.63008			1			1	
	D012	181.29917	-42.97175			1		1		
	D013	181.66133	-42.53433		1	1			1	
	D014	181.66235	-42.56604		2	1			1	
	D015	181.60201	-42.67243		2	1			2	
	D016	181.66310	-42.70640		2	1			1	
	D017	181.65192	-42.72621			1			1	
	D018	181.65135	-42.76179		2	1			1	
	D019	182.78713	-42.65629			1			1	
	D020	182.78511	-42.68694		1	1			1	
	D021	182.78854	-42.71196		2	1			1	
	D022	182.79010	-42.77640		2	1			2	
	D023	182.77331	-42.80679		2	1			1	
	D024	182.77275	-42.84829		3	1			2	
	D025	181.18738	-42.46975			1			1	
	D026	177.14858	-43.47392			1		1		
	D027	175.27833	-43.43275			1				
	D028	174.45463	-43.72767			1		1		
	D029	175.14692	-42.54171			1			1	

Table A2: Epibenthic megafaunal taxa identified from seamount sled (SEL) and Beam Trawl (TB) samples collected during voyages TAN0705 and TAN0707 to Chatham Rise and Challenger Plateau.

Phylum	Class	Order	Number of taxa (species or higher level)	
Foraminifera	Xenophyophorea		1	
Porifera	Demospongiae	Astrophorida	8	
		Dendroceratida	1	
		Dictyoceratida	3	
		Hadromerida	11	
		Halichondrida	6	
		Haplosclerida	8	
		Poecilosclerida	21	
		Spirophorida	6	
		Hexactinellida	Amphidiscosida	5
	Hexactinosida	2		
	Lyssacosida	6		
	Cnidaria	Anthozoa	Actiniaria	15
			Alcyonacea	7
Antipatharia			2	
Corallimorpharia			2	
Gorgonacea			10	
Pennatulacea			14	
Scleractinia			12	
Telestacea			2	
Zoanthidea			4	
Hydrozoa			Anthoathecata	8
Hydroida		1		
Leptothecata		11		
Scyphozoa			1	
Mollusca		Aplacophora	Aplacophora	1
		Bivalvia	Arcoida	4
	Dimyidae		1	
	Limoida		1	
	Myoida		9	
	Mytiloida		1	
	Nuculoidea		13	
	Ostreoida		4	
	Pholadomyoida		2	
	Pterioida		2	
	Veneroida		5	
	Cephalopoda		Octopoda	2
	Sepiida		1	
	Sepiolida		1	
	Spirulida	1		
	Teuthida	2		
	Gastropoda	Kapala	1	
		Speoides	1	
	Gastropoda Opisthobranchia	Cephalaspidea	4	
		Nudibranchia	1	

Phylum	Class	Order	Number of taxa (species or higher level)	
Mollusca	Gastropoda Prosobranchia	Archaeogastropoda	10	
		Cocculiniformia	1	
		Heterostropha	1	
		Mesogastropoda	18	
		Neogastropoda	31	
		Neotaenioglossa	6	
		Stenoglossa	29	
		Vetigastropoda	2	
		Polyplacophora	Neoloricata	1
		Polyplacophora Neoloricata	Ischnochitonida	2
Scaphopoda	Dentaliida	3		
Solenogastera		1		
Nemertea	Nemertea	Nemertea	1	
Echiura			1	
Priapulida	Priapulida	Priapulida	1	
Sipuncula	Sipunculidea		1	
Annelida	Polychaeta	Amphinomida	3	
		Eunicida	22	
		Phyllodocida	11	
		Phyllodocida Aphroditiformia	13	
		Phyllodocida Nereidiformia	3	
		Sabellida	13	
		Scolecida	19	
		Spionida	5	
		Terebellida Cirratuliformia	2	
		Terebellida Terebelliformia	6	
Brachiopoda	Articulata	Terebratulida	6	
	Rhynchonellida		1	
Bryozoa	Gymnolaemata	Cheilostomata	70	
		Ctenostomata	1	
		Stenolaemata	5	
Arthropoda Chelicerata	Pycnogonida	Pycnogonida	2	
Arthropoda Crustacea	Malacostraca	Amphipoda	20	
		Decapoda	87	
		Euphausiacea	1	
		Isopoda	18	
		Mysidacea	2	
		Stomatopoda	1	
		Tanaidacea	1	
		Maxillopoda	Pedunculata	11
			Sessilia	2
		Ostracoda		1

Phylum	Class	Order	Number of taxa (species or higher level)
Echinodermata	Asteroidea	Brisingida	6
		Forcipulatida	5
		Notomyotida	9
		Paxillosida	18
		Spinulosida	4
		Valvatida	16
		Velatida	9
		Articulata	4
	Crinoidea	Bourgueticrinida [aka Millericrinida]	3
		Cyrtocrinida	1
		Cidaroida	5
	Echinoidea	Echinoida	3
		Echinothurioida	5
		Pedinoida	1
		Spatangoida	9
		Aspidochirotida	6
	Holothuroidea	Dactylochirotida	1
		Dendrochirotida	9
		Elasipodida	6
		Holothuroidea	1
Molpadiida		8	
Ophiuroidea		Euryalinida	3
		Ophiurida	48
	Chordata	Ascidiacea [Tunicates]	Enterogona Aplousobranchia
Pleurogona Stolidobranchia			7
Thaliacea [Salps]			1

Table A3: Taxa and total number of individuals identified from upper net samples of the Brenke epibenthic sled (SEH). Totals are summed across all 20 sites at which the SEH was deployed.

Phylum	Class	Superorder	Order	Taxon	No. individuals	
Cnidaria	Hydrozoa or Scyphozoa			Medusa	26	
Mollusca	Bivalvia			Bivalvia	1888	
	Gastropoda			Gastropoda	1845	
	Scaphopoda			Scaphopoda	676	
	Solenogastres			Solenogastres	705	
Nematoda			Nematoda	1567		
Nemertea			Nemerteans	3		
Sipuncula	Sipunculida			Sipunculida	8	
Annelida	Polychaeta indeterminate worms			Polychaeta	5012	
				worms	1572	
Brachiopoda				Brachiopoda	4850	
Chaetognatha				chaetognaths	44	
Arthropoda	Malacostraca	Eucarida	Decapoda	Anomura	67	
				Brachyura	58	
				Natantia	249	
				Euphausiacea	20	
		Peracarida	Amphipoda	14493		
			Cumacea	6294		
			Isopoda	7740		
			Mysida	483		
	Tanaidacea		2214			
	Leptostraca		178			
	Maxillopoda		Cirripedia	5		
			Copepoda	21851		
	Ostracoda			Ostracoda	4264	
	Pycnogonida			Pycnogonida	5	
	Echinodermata	Asteroidea			Asteroidea	432
		Crinoidea			Crinoidea	90
Echinoidea				Echinoidea	929	
Holothuroidea				Holothuroidea	441	
Ophiuroidea				Ophiuroidea	1218	
Chordata	Ascidiacea			Ascidiacea	5	
	Thaliacea		Salpida	salps	345	

Table A4: Substrate descriptors available during analysis of DTIS still images. Note that this is the full substrate table available in the NICAMS database and only a subset of these descriptors was actually present in the Chatham-Challenger images that were analysed. Substrate types were recorded as percentages of the full image area.

Category	Substratum	qualifier	Label	definition
Hard substrata	Bedrock		Hard substrata Bedrock	
		outcrop	Bedrock outcrop	
		smooth	Bedrock smooth	
		irregular	Bedrock irregular	
		eroded	Bedrock eroded	
	Lava	scoured	Bedrock scoured	
			Larva	
		glassy	Larva glassy	
	Breccia	pillow	Larva pillow	
		lobate	Larva lobate	
	sheet	Larva sheet		
	Breccia	Breccia	rock composed of angular fragments of rocks or minerals in a matrix of cementing material.	
	Talus	Talus	rock that occurs in fragments or particles lying on, above, or adjacent to the place in which it was originally formed or deposited	
Soft sediments	Boulders		Boulders	>25.6 cm
	Boulders	smooth	Boulders smooth	
	Boulders	irregular	Boulders irregular	
	Cobbles		Cobbles	6.5 - 25 cm
	Pebbles		Pebbles	0.4 - 6.4 cm
			Soft sediment	
	Gravel		Gravel	0.2 - 0.4 cm
		waves	Gravel waves	wavelength > ca 50 cm
		ripples	Gravel ripples	wavelength < ca 30 cm
		flat	Gravel flat	
	overlay	Gravel overlay	gravel overlying hard substrata	
	Sand	Sand	0.063 - 0.2 cm	
	waves	Sand waves	wavelength > ca 50 cm	
	ripples	Sand ripples	wavelength < ca 30 cm	
	flat	Sand flat		
	overlay	Sand overlay	Sand overlying hard substrata	
	Mud	Mud	<0.063 cm	
	overlay	Mud overlay	Mud overlying hard substrata	
Biogenic			Biogenic substrata	
	Coral rubble		Coral rubble	broken fragments of stony corals
	Shell hash		Shell hash	non seep or vent related taxa
	Coral (intact)		Coral (intact)	structurally intact stands of reef-forming corals: may be live, dead, or mixture

Category	Substratum	qualifier	Label	definition
Biogenic	Epifauna high		Epifauna high	mixed complex assemblage of erect epifauna
	Epifauna low		Epifauna low	mixed complex assemblage, epifaunal turf
	Pteropod shell		Pteropod shell	dead pteropod shells
	Salps		Salps	dead salps on the seafloor
Anthropogenic	Trawl marks		Trawl marks	e.g., sled tracks, grab pits
	Fishing gear		Fishing gear	
	Rubbish		Rubbish	
	Sampling marks		Sampling marks	
Vents & seeps	Carbonate rock		Carbonate rock	weathered carbonate rock, few or no live seep fauna associated
	Chemoherm		Chemoherm	'soft' appearance, high relief, associated with live chemosynthetic fauna and active seepage
	Sulphidic sediment		Sulphidic sediment	dark sediment, often with characteristic 'raindrop' texture
	Bacterial mat		Bacterial mat	white patches, usually on soft sediments
	Vesicomylid shell		Vesicomylid shell	dead vesicomylid clam shells
	Bathymodiolin shell		Bathymodiolin shell	dead Bathymodiolin mussel shells
	<i>Acharax</i> sp. Shell		<i>Acharax</i> sp. Shell	dead <i>Acharax</i> sp. shells

Table A5: Bioturbation mark descriptors available during analysis of DTIS still images. Marks were recorded as numbers per m².

Category	Substratum	qualifier	Label	definition
Bioturbation			Bioturbation	
	Burrow		Burrow	
	Track		Track	
	Mound		Mound	
	Pit		Pit	
	Faecal coil		Faecal coil	e.g. holothurian casts
	Ring of burrows		Ring of burrows	raised mound surrounded at base by burrow entrances
	Spiral		Spiral	hemichordate worm feeding trace
	Resting trace	asteroid	Resting trace asteroid	impression of body outline
	Resting trace	ophiuroid	Resting trace ophiuroid	impression of body outline
Paleodictyon		Paleodictyon	hexagonal dot matrix pattern	

Table A6: DTIS still images: operational taxonomic units (OTU) for mega-epibenthic fauna identified from seabed still images.

Phylum	Class	Order	OTU
Annelida	Echiura	Aciculata Canalipalpata	Echiuran
	Polychaeta		Hyalinoecia
			Sabellid
			Errant polychaete
			Polychaeta
			Worm indet.
Arthropoda	Malacostraca	Decapoda	<i>Agononida nielbrucei</i>
			Atelecyclidae
			Brachyura
			<i>Campylonotus rathbunae</i>
			Chirostylidae
			Galatheidae
			Galatheoidea
			<i>Gastroptychus novaezelandiae</i>
			Glyphocrangon
			Goneplacidae
			<i>Haliporoides sibogae</i>
			<i>Ibacus alticrenatus</i>
			Inachidae
			<i>Leptomithrax longipes</i>
			<i>Lithodes cf. longispinus</i>
			<i>Lithodes murrayi</i>
			Lithodidae
			Majidae
			<i>Metanephrops challengerii</i>
			<i>Munida gracilis</i>
			<i>Natant decapod</i>
			<i>Nematocarcinus</i> sp.
			<i>Neolithodes brodiei</i>
			<i>Neommatocarcinus huttoni</i>
			Nephropidae
			Paguridae
			<i>Paralomis zealandica</i>
			<i>Platymaia maoria</i>
			<i>Plesionika</i> sp.
			<i>Polycheles</i> spp.
			Polychelidae
			<i>Pycnoplax victoriensis</i>
			Scyllaridae
			<i>Teratomaia richardsoni</i>
			<i>Trichopeltarion fantasticum</i>
			<i>Vitjazmaia latidactyla</i>
		Isopods	
		Serolidae	
		Peracida	
	Pycnogonida	<i>Collossendeis</i> sp.	
		<i>Pycnogonid</i> indet	
		Pycnogonids	

Phylum	Class	Order	OTU		
Brachiopoda	Gymnolaemata	Cheilostomata	Brachiopoda		
Bryozoa (Ectoprocta)			Bryozoan		
			Bryozoan - antler form		
			Bryozoan - bushy form		
			Bryozoan - erect cheilostome		
			Bryozoan - feather form		
			Bryozoan - branched form		
			Bryozoan - branched white		
			Bryozoan - encrusting cheilostome		
			Bryozoan - lace form		
Chordata			Ascidiacea	Actiniaria	Ascidian
Cnidaria			Anthozoa		Anemone (hermit)
					Anemone (large columnar)
		Anemone (mauve)			
		Anemone (small red)			
		Anemone indet.			
		Anenome 10			
		Anenome 11			
		Anenome 12			
		Anenome 13			
		Anenome 14			
		Anenome 15			
		Anenome 16			
		Anenome 17			
		Anenome 18			
		Anenome 19			
		Anenome 4			
		Anenome 5			
		Anenome 6			
		Anenome 7			
		Anenome 8			
		Anenome 9			
		Anenome 1			
		Anenome 2			
		Anenome 3			
		Anenome indet.			
		Alcyonacea			
		Alcyoniidae			
		Anthomastus 3			
		<i>Anthomastus</i> sp.			
		<i>Clavularia</i> sp.			
		Taiaroa tauhou			
		<i>Telesto</i> sp.			
		Antipatharia			
		Antipathes			
		Bathypathes			
		Dendrobathypathes			
		Leiopathes			
		Parantipathes			
		Trissopathes			
		Ceriantheria			
		<i>Ceriantharia</i> spp			

Phylum	Class	Order	OTU
Cnidaria	Anthozoa	Corallimorpharia	Corallimorpharia Corallimorpharia 1 Corallimorpharia 2 Corallimorpharia 3
		Gorgonacea	<i>Callogorgia</i> sp. Chrysogorgiidae Coralliidae <i>Corallium</i> sp. Gorgonacea Isididae <i>Lepidisis</i> sp. <i>Paragorgia</i> sp. Paragorgiidae <i>Primnoella</i> sp. Primnoidae Primnoidea/Callogorgia <i>Radicipes</i> sp. <i>Sibogorgia</i> spp. <i>Thourella</i> sp.
		Pennatulacea	<i>Acanthoptilum</i> sp. <i>Anthoptilum grandiflorum</i> <i>Anthoptilum</i> sp. <i>Distichoptilum gracile</i> <i>Funiculina quadriangularis</i> <i>Gyrophyllum sibogae</i> <i>Halipteris</i> sp. <i>Kophobelemnon</i> sp. <i>Kophobelemnon stelliferum</i> <i>Pennatula aculeata</i> <i>Pennatula inflata</i> <i>Pennatula</i> sp. Pennatulacea Pennatulacea 1 Pennatulacea 2 Pennatulacea 3 Pennatulacea 4 Pennatulacea 5 <i>Stylatula</i> sp.
		Scleractinia	Cup coral <i>Enallopsammia</i> spp. Flabellum Flabellum 1 Flabellum 3 <i>Flabellum knoxi</i> <i>Flabellum loure kexei</i> <i>Flabellum rubrum</i> <i>Goniocorella dumosa</i> <i>Madrepora oculata</i> <i>Madrepora</i> sp. Scleractinia

Phylum	Class	Order	OTU		
Cnidaria	Anthozoa	Scleractinia	<i>Solenosmillia variabilis</i> Solitary coral		
		Stolonifera	<i>Rhodelina</i> sp.		
		Zoanthidea	Epizoanthidea Coral indet.		
	Hydrozoa	Anthoathecatae		Athecate hydroid 1 <i>Errina</i> sp. Stylasteridae	
			Leptothecatae	Hydroid Hydroid 2 branches Hydroid A Hydroid orange	
		Asterioidea	Brisingida		<i>Brisinga chathamica</i> <i>Brisinga tasmani</i> Brisingid 1 Brisingid 2 Brisingid 3 Brisingid 4 Brisingidae Hymenodiscididae
				Forcipulatida	Asteriidae <i>Cosmasterias dyscrita</i> <i>Pseudechinaster rubens</i> <i>Zoroaster</i> sp. Zoroasteridae Zoroasteridae/Asteriidae
				Notomyotida	<i>Benthopecten</i> sp. Benthopectinidae
				Paxillosida	Astropectinidae <i>Dipsacaster magnificus</i> <i>Dipsacaster</i> sp. Paxillosida? <i>Radiaster</i> sp. Radiasteridae
				Spinulosida	<i>Crossaster multispinus</i> <i>Crossaster</i> sp. Echinasteridae <i>Henricia</i> sp. <i>Hymenaster</i> sp. Pterasteridae <i>Solaster torulatus</i> Solasteridae
Echinodermata	Asterioidea	Valvatida	<i>Ceramaster</i> sp. Goniasteridae <i>Hippasteria</i> sp. <i>Lithosoma novazealandiae</i> Lithosoma/Pseudarchaster <i>Mediaster</i> sp. <i>Pillsburiaster</i> sp. Valvatida		
			Velatida	Myxaster? Asteroids	

Phylum	Class	Order	OTU		
Echinodermata	Crinoidea	Bourgueticrinida	Crinoidea (stalked)		
		Comatulida	Crinoidea (motile)		
	Echinoidea	Cidaroida	Cidaridae	Crinoids	
			Cidaroida	Cidaridae	
			Goniocidarinae	Cidaroida	
			<i>Goniocidaris parasol</i>	Goniocidarinae	
			<i>Goniocidaris</i> sp.	<i>Goniocidaris parasol</i>	
			Histocidaridae	<i>Goniocidaris</i> sp.	
			<i>Ogmocidaris benhami</i>	Histocidaridae	
			Stereocidaridae	<i>Ogmocidaris benhami</i>	
			<i>Peronella hinemoae</i>	Stereocidaridae	
			<i>Dermechinus horridus</i>	<i>Peronella hinemoae</i>	
		Clypeasteroida	Echinoida	Echinidae	<i>Dermechinus horridus</i>
				Echinoida	Echinidae
		Echinothurioida	Echinothurioida	Echinothuriidae	Echinoida
				Echinothuriidae/Phormosomatidae	<i>Gracilechinus multidentatus</i>
				<i>Phormosoma bursarium</i>	Echinothuriidae
				Phormosomatidae	Echinothuriidae/Phormosomatidae
		Pedinoida	Pedinoida	<i>Sperosoma</i> sp.	<i>Phormosoma bursarium</i>
				Caenopedina	Phormosomatidae
<i>Caenopedina</i> spp.	<i>Sperosoma</i> sp.				
Spatangoida	Spatangoida	Pedinidae	Caenopedina		
		<i>Paramaretia peloria</i>	<i>Caenopedina</i> spp.		
		Spatangidae	Pedinidae		
Temnopleuroidea	Temnopleuroidea	<i>Spatangus</i> sp.	<i>Paramaretia peloria</i>		
		<i>Pseudechinus flemingi</i>	Spatangidae		
		Temnopleuridae	<i>Spatangus</i> sp.		
Holothuroidea	Aspidochirotida	Echinoids	<i>Pseudechinus flemingi</i>		
		<i>Bathyplores moseleyi</i>	Temnopleuridae		
		<i>Bathyplores</i> sp.	Echinoids		
		<i>Bathyplores sulcatus</i>	<i>Bathyplores moseleyi</i>		
		<i>Benthodytes incerta</i>	<i>Bathyplores</i> sp.		
		<i>Pseudostichopus mollis</i>	<i>Bathyplores sulcatus</i>		
	Elasipodida	Elasipodida	<i>Pseudostichopus mollis</i>	<i>Benthodytes incerta</i>	
			<i>Pseudostichopus peripatus</i>	<i>Pseudostichopus mollis</i>	
			Stichopodidae	<i>Pseudostichopus peripatus</i>	
			<i>Stichopus mollis</i>	<i>Pseudostichopus</i> sp.	
			Synallactidae	Stichopodidae	
			Elasipoda	<i>Stichopus mollis</i>	
Elasipoda 1	Synallactidae				
Elasipoda 2	Elasipoda				
<i>Enypniastes eximia</i>	Elasipoda 1				
<i>holothurian indet < 25 mm (Pale)</i>	Elasipoda 2				
<i>Laetmogone violacea</i>	<i>Enypniastes eximia</i>				
<i>Pannychia</i> sp.	<i>holothurian indet < 25 mm (Pale)</i>				
Pelagothuridae	<i>Laetmogone violacea</i>				
Psychropotidae	<i>Pannychia</i> sp.				
<i>holothurian indet < 25 mm (Pale)</i>	Pelagothuridae				
<i>holothurian indet.</i>	Psychropotidae				

Phylum	Class	Order	OTU	
Echinodermata	Holothuroidea		holothurian uni 1	
			holothurian uni 2	
			holothurian uni 3	
			holothurian uni 4	
	Ophiuroidea	Euryalinida		Holothurians
				Euryalinida
		Ophiurida		Gorgonocephalidae
				Amphiuridae
				Ophiacanthidae
				<i>Ophiomusium lymani</i>
				<i>Ophiomyxa brevirima</i>
				Ophiurida (apricot small)
				Ophiurida (Mauve small)
			Ophiurida unspecified	
	Ophiuridae			
	Unknown ophiurida 0			
	Ophiuroids			
Echiura			Echiuran	
Foraminifera (Protozoa)	Foraminifera	Foraminiferida	Foram (giant)	
			Foraminifera	
Mollusca	Granuloreticulosea	Foraminiferida	Bathysiphon	
			Bivalvia	Ostreoida
	Cephalopoda	Pholadomyoida		<i>Delectopecten fosterianus</i>
			<i>Euciroa galathea</i>	
			Bivalvia	
			Bathypolypodinae	
			<i>Benthoctopus</i> sp.	
			Cirroteuthididae/Luteuthididae	
			<i>Enteroctopus zealandicus</i>	
			<i>Graneledone</i> sp.	
Graneledoninae				
Octopodinae				
Opisthoteuthididae				
<i>Opisthoteuthis</i> sp.				
<i>Pinnoctopus cordiformis</i>				
Cephalopoda				
Gastropoda	Archaeogastropoda		<i>Calliostoma alertae</i>	
			Callostomatidae	
			Aeneator recens	
			<i>Amalda</i> sp.	
			<i>Austrofusus glans</i>	
			Buccinidae	
			<i>Coluzea</i> sp.	
			<i>Comitas onokeana vivens</i>	
			Muricidae	
			Olividae	
			<i>Pagodula</i> sp.	
			<i>Penion</i> sp.	
			Turbinellidae	
			Turridae	
			Volutidae	
			<i>Volutomitira banksi</i>	
			Volutomitridae	

Phylum	Class	Order	OTU	
Porifera		Neotaenioglossa	Cassidae <i>Fusitriton magellanicus</i> Naticidae Ranellidae	
		Nudibranchia	Opisthobranchia Gastropoda Opisthobranchia Scaphopoda	
		Scaphopoda	Scaphopoda	
		Calcarea	Leucosolenida	Leucosolenia Calcarea
		Demospongiae	Astrophorida	Astrophorid <i>Geodia regina</i> <i>Geodinella vestigifera</i> Pachastrellidae <i>Tethyopsis n. sp.</i> <i>Thenea sp.</i>
			Hadromerida	Hadromerid <i>Suberites affinis</i>
			Halichondrida	Axinella or Pararaphoxya <i>Axinella spp.</i> Halichondrid
			Haplosclerida	Haplosclerid Petrosia
			Lithistida	<i>Awhiowhio sepulchrum</i> <i>Costifer wilsoni</i> Lithistid <i>Neoaulaxinia persicum</i>
			Poecilosclerida	Cladhorizidae <i>Cladhorizidae sp. nov.</i> <i>Latrunculia spp.</i> Poecilosclerid
	Spirophorida		Spirophorida <i>Tetilla leptoderma</i> Demospongiae Encrusting sponges Demospongiae	
	Demspungiae Hexactinellida	Amphidiscosida	<i>Hyalonema sp.</i> <i>Pheronema sp.</i>	
		Hexactinosida	Farreidae Hexactinosida	
		Lyssacinosida	<i>Euplectella regalis</i> <i>Hyalascus n. sp.</i> Hexactinellida Demospongiae Sponge (mauve)	
		Amphipoda Euphausiacea Mysida	Amphipoda Euphausiacea Mysida Buccinidae/Ranellidae Galatheidae (white)	