

Associations of demersal fish with sponge grounds on the continental slopes of the northwest Atlantic

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ABSTRACT: Association of demersal fish taxa with *Geodia*-dominated temperate sponge grounds was examined in 104 research vessel survey trawl sets from 500 to 1500 m depth along the continental slopes of the Grand Banks and Flemish Cap, northwest Atlantic. The total number of fish taxa and their total biomass were negatively correlated with sponge biomass and catch depth, and sponge biomass increased with depth. ANOSIM identified significantly different fish communities (biomass and abundance composition) associated with 3 sponge catch weight classes, and multi-dimensional scaling plots showed separation of trawl sets labelled by this factor to be strongest for abundance. SIMPER analysis identified the fish assemblage associated with the sponge grounds (≥ 250 kg km⁻¹): roughhead grenadier *Macrourus berglax*, roundnose grenadier *Coryphaenoides rupestris*, blue hake *Antimora rostrata*, longnose eel *Synaphobranchus kaupii*, turbot *Reinhardtius hippoglossoides*, black dogfish *Centroscyllium fabricii* and deep-sea cat shark *Apristurus profundorum*. These contributed to 92 and 81 % of the similarity (~50 %) among trawl set catches using biomass and abundance data, respectively. Four additional taxa contributed to 92 % similarity in abundance: common grenadier *Nezumia bairdii*, shortnose snipe eel *Serrivomer beanii*, lanternfish (Myctophidae) and goitre blacksmelts *Bathylagus euryops*. SIMPER dissimilarity identified 6 fish taxa with larger biomass in catches with high sponge catch weight: shortnose snipe eel, deep-sea cat shark, eelpout *Lycodes* spp., spinytail skate *Bathyraja spinicauda*, white skate *Dipturus linteus* and deepwater chimaeras *Hydrolagus affinis*. The first 3 species were also more abundant in those catches and are considered to be strongly associated with the sponge grounds. Future research focussing on these species identified as being associated with the sponge grounds may provide greater insight into the strength of the associations and the ecological linkages between these benthic and pelagic communities.

KEY WORDS: Demersal fish assemblages · Sponge grounds · Trawl catch

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INTRODUCTION

The importance of individual sponges as microhabitat for invertebrate species has been widely demonstrated and includes a wide range of ecological interactions including both facultative and obligate commensalisms (see recent reviews by Wulff 2006 and Bell 2008, and articles specific to the North Atlantic by Bett & Rice 1992, Klitgaard 1995, Klit-

gaard & Tendal 2004, ICES 2009, Hogg et al. 2010, Barrio Froján et al. 2012). At these small spatial scales, sponge architecture is an important determinant of the type and strength of such interactions, as sponge morphology includes a wide diversity of forms.

At larger spatial scales, the use of temperate sponge grounds as habitat for fish assemblages has been less well documented, especially in terms of

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obligative, facultative, or opportunistic relationships (Hixon et al. 1991, ICES 2009, Hogg et al. 2010). Sponges can form dense aggregations in some temperate waters, including biogenic reefs (e.g. Hectate Glass Sponge Reefs in British Columbia, Canada; Chu & Leys 2010, Hogg et al. 2010) and sponge grounds (e.g. continental slope off Labrador, Canada; Kenchington et al. 2010). The latter are characterized by the presence of 1 or more dominant taxa which have a larger individual- and population-level biomass than smaller associated sponge species. Such structure-forming species include members of the genera *Geodia*, *Stryphnus*, *Asconema*, *Vazella* and *Stelletta*, amongst others (ICES 2009). Fish often use the structural habitat that sponge grounds provide for shelter, reproduction and to forage for food (Bell 2008). The 3-dimensional spatial complexity of sponge grounds also provides important nursery grounds for juvenile fish in their early stages of growth (Auster et al. 2003, Freese & Wing 2003) and shelter for all life stages (Brodeur 2001). Rockfish (or 'redfish') of the genus *Sebastes* are particularly prevalent in sponge grounds in some areas, living both inside and between the sponges (Richards 1986, Freese & Wing 2003, Marliave et al. 2009, Du Preez & Tunnicliffe 2011). Marliave et al. (2009) describe habitat partitioning by *Sebastes maliger* in which adults are associated with the reef structures (bioherms) and juveniles are associated with single sponges or lower density 'sponge gardens'. *S. maliger* feeds on benthic crustaceans, and the authors hypothesize that increased species richness in the food resource on the sponge gardens drives this distribution pattern. However, avoidance of certain sponges by some species may also occur. Bell (2008) cites a number of examples in which the chemical compounds of the sponges act as deterrents to other organisms. In temperate waters, Marliave et al. (2009) show regional patterns in British Columbian waters (Northeast Pacific) in the association of adult *S. maliger* with the sponge reef structures, with the fish absent from some sponge reefs entirely but present in nearby areas. There is also some evidence that removal of temperate sponge grounds by bottom trawling changes the composition of the fish fauna, providing weak evidence for the association of some fish with such habitats (cf. Klitgaard & Tendal 2004). Thus, it seems that temperate sponge grounds may be an important refuge and habitat for fish as well as invertebrates, although little ecological work has been carried out to understand the exact nature of this habitat use in the deep sea, and most studies to date are limited to tropical waters (e.g. McCormick 1994, Cleary & de Voogd 2007).

Given the limited evidence in the literature regarding temperate sponge and fish interactions, we identified a need to look at large-scale species associations. Fish species that are consistently associated with temperate sponge grounds, when alternative habitats are available, are likely to be actively selecting the sponge grounds for some purpose. Here we present novel data on the presence/absence of fish assemblages compared with sponge biomass in the temperate waters of the northwest Atlantic. We describe the relationship between the fish fauna associated with the sponge grounds on the continental slopes of the Grand Banks and Flemish Cap, off Newfoundland, Canada. We give a particular focus to the Northwest Atlantic Fisheries Organization (NAFO) Regulatory Area (NRA, seaward of the Canadian 200 mile limit), where these sponge grounds are located and have been verified with *in situ* photography (NAFO 2010). The sponges in this area are dominated by *Geodia* species, which are massive ball sponges found throughout the North Atlantic (ICES 2009). However, at least 27 different species have been identified in the study area (Murillo et al. 2012). *Geodia barretti*, *G. phlegraei*, *G. macandrewi*, *Stryphnus ponderosus* and *Stelletta normani* are the main structural sponges and constitute >99% of the total invertebrate biomass (Murillo et al. 2012). These species all have similar morphologies, being massive spherical/lobed sponges reaching diameters in the range of 20 to 55 cm (Best et al. 2010). Although pelagic fish may use sponge grounds as habitat, we expect that if there are any large-scale associations between fish assemblages and sponge grounds, they will be strongest in demersal fish assemblages which are more closely associated with the sea floor. Our hypothesis is that demersal fish assemblages caught in research vessel trawl catches will differ in areas of low, medium and/or high sponge biomass.

MATERIALS AND METHODS

Data sources and selection

Data used for these analyses come from the Fisheries and Oceans Canada (DFO) Newfoundland Region fall multispecies surveys of the Grand Banks inside and adjacent to Canada's exclusive economic zone. These surveys followed a depth-stratified random design to 1500 m and employed a Campelen 1800 shrimp trawl. The Campelen carries out 15 min tows using a towing speed of 3.0 knots. The catches

were standardized to a 1 km distance towed. The average wingspread was 16.84 m. Because catchability of the standard survey trawl was unknown for most species, the indices were considered to be relative estimates only. A trawl-mounted CTD was used to collect temperature data. The trawl sets were almost evenly split between day and night times. The catch was sorted at sea, and the number and weight (kg) of each taxon were recorded using a standard set of species codes. Fish were generally identified to species level, whereas sponges were only recorded as Porifera and weighed collectively for each set (not counted). Only records from 2001 to 2007 were used in order to avoid confounding the results by temporal trends due to environmental factors (cf. Colbourne 2004) and to ensure consistency of reporting. These records were further reduced to include: (1) only those sets deeper than 500 m, to minimize confounding of the results by including both shelf and slope taxa; (2) only those sets from south of 50°N latitude, in order to reduce confounding the results by introducing biogeographic differences in community composition through disproportionate representation of arctic and boreal faunas; and (3) only records in which the sponge catch was verified at sea through species keys or identified via representative samples post-

survey (to avoid false identification of bryozoans and other taxa as sponges). For (3), zero sponge catch records were not included, as we could not be certain that sponges were not caught, as they may have just not been recorded. These criteria produced 104 trawl records for analysis, with an average depth of 1096 m (range: 578 to 1446 m) (Fig. 1).

The 104 selected trawls contained non-zero records for 200 taxa of fish. The weights and abundance of each of the taxa were standardized to a 1 km trawl, using the start and end positions of the trawl to calculate distance on bottom. This involved only a minor adjustment to the data, as the average trawl length was 0.8 ± 0.07 km (range: 0.6 to 1.0).

The 200 taxa were reduced by: (1) combining some species to higher level groupings and (2) eliminating all rarities after combining the data to include only taxa $>0.1\%$ of total biomass/abundance. The first of these steps was done to avoid introducing errors due to taxonomic imprecision among trips and also sets within trips. The second was to eliminate taxa that may not be reliably caught in the trawl, or whose rarity may escape detection in the routine sorting and sub-sampling processes. To determine the low end cut off, decisions were made on the size of the taxon relative to the biomass record before removing it from the list.

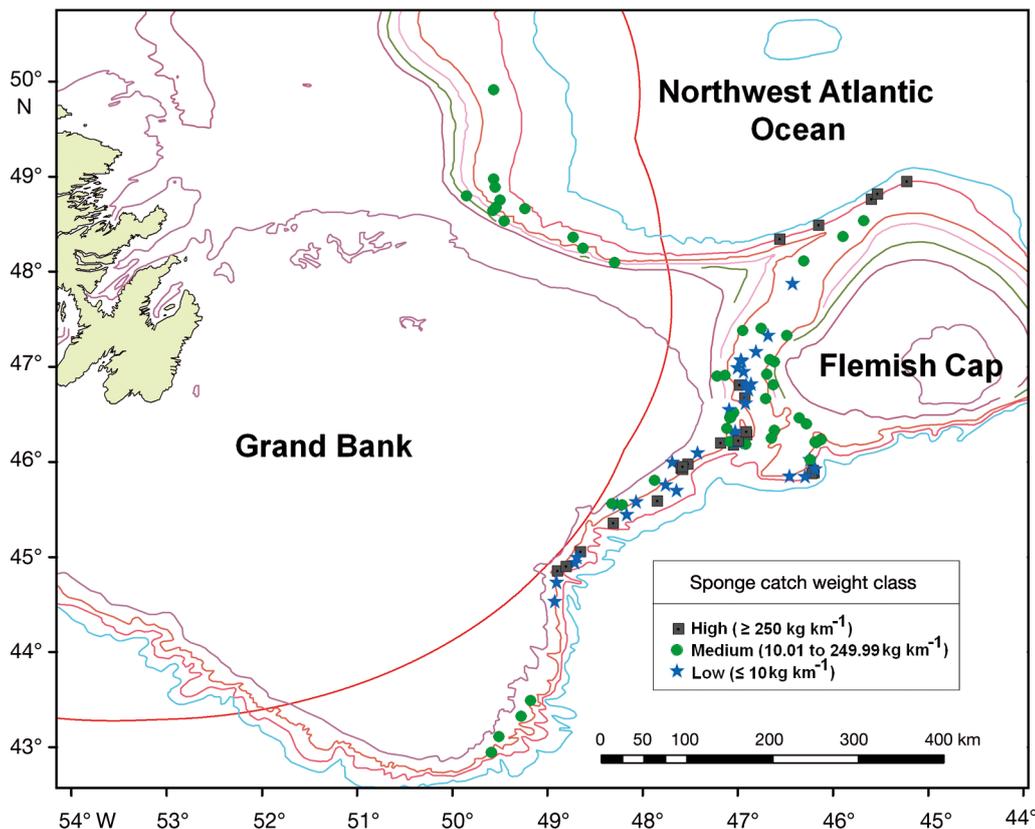


Fig. 1. Locations off Newfoundland, Canada, of the research vessel trawls used in our analyses, with corresponding sponge catch weight classes. Thick red line is the Canadian Exclusive Economic Zone (EEZ). Depth contours are shown for 200, 400, 600, 800, 1000, 1500 and 2000 m

Data analyses

The analytical variables used to describe the fish communities were: total number of fish taxa (number km^{-1} of the 34 species for which biomass was analyzed; see Table 1), total fish biomass (kg km^{-1}), total fish abundance (number km^{-1}), fish taxon biomass (weight in kg of each taxon, usually species, km^{-1}) and fish taxon abundance (number km^{-1}). Total sponge biomass (kg km^{-1}), average trawl depth (m) and bottom temperature ($^{\circ}\text{C}$) were also recorded for each trawl set.

The cumulative distribution of total sponge biomass is highly right-skewed, that is, there are large numbers of very small catches and very few large ones (Kenchington et al. 2009). This distribution was used to categorize the trawl sets according to 1 of 3 arbitrary sponge catch weight classes which were selected to represent sponge grounds, transition areas and non-sponge ground, respectively (sponge catch weight class; see Supplement 1 at www.int-res.com/articles/suppl/m477p217_supp.pdf): high ($\geq 250 \text{ kg km}^{-1}$, which is the upper quartile of the distribution; $N = 24$), medium (10.01 to $249.99 \text{ kg km}^{-1}$; $N = 30$) and low ($\leq 10 \text{ kg km}^{-1}$, which is approximately the median of the distribution; $N = 50$). The high class equates to sponge grounds. The medium class may arise through trawl sets encountering the fringes of sponge grounds (research vessel trawl catches $> 75 \text{ kg}$ of sponge per trawl set for this survey and area; Kenchington et al. 2009), or through the accumulation of smaller sponges that are not associated with sponge grounds (Kenchington et al. 2011). The low class is outside of the sponge grounds. The small weight threshold for this class could be achieved through collection of a single sponge.

The variance in average trawl depth (Supplement 1) was tested for equality among sponge catch weight classes using Levene's test and was found to be unequal ($p = 0.04$). This variable was analyzed in its untransformed state, and Welch's ANOVA was used to test for an effect of sponge catch weight class on average trawl depth. Tukey's HSD test was used for post hoc comparisons of the mean values to identify differences between factor levels. These and similar analyses below were performed with JMP v.6.0.3 software (SAS Institute). The locations of the trawl sets used in these analyses, identified by their sponge weight class, are illustrated in Fig. 1.

The variance of the total number of fish taxa (Supplement 1) was tested for equality among sponge catch weight classes using Levene's test and was found to be unequal; attempts to transform it did not

improve its variance structure. This variable was analyzed in its untransformed state, and Welch's ANOVA was used to test for an effect of sponge catch weight class on total number of fish taxa. Tukey's HSD test was used for post hoc comparisons of the mean values to identify differences between factor levels. The total number of fish taxa was regressed against average trawl depth and log₁₀-transformed total sponge biomass using a rank correlation.

Total fish biomass and total fish abundance (Supplement 1) were log₁₀ transformed and their variances were found to be homogeneous among sponge catch weight classes using Levene's test. ANOVAs were used to assess whether mean total fish biomass and mean total fish abundance were equal among sponge catch weight classes. Tukey's HSD test was used for post hoc comparisons of the mean values to identify differences between factor levels. The variables were independently regressed against average trawl depth and log₁₀-transformed total sponge biomass.

Differences in community composition among sponge weight classes were examined with ANOSIM (analysis of similarities; Clarke 1993). The test statistic, R , is based on comparing distances between groups with distances within groups using permutation. The degree of sensitivity of R to heterogeneity may be influenced by our unbalanced design, depending on whether the group with the larger numbers of samples (low sponge catch weight class) had greater or lesser dispersion than the other groups. In our data set, variance in fish taxon biomass was not statistically different among the sponge catch weight classes. Bray-Curtis similarity matrices of trawl set pairs were constructed based on the transformed fish taxon biomass (Table 1) and fish taxon abundance (Table 2) of the 34 fish taxa identified for each variable. A similarity of percentages (SIMPER) analysis was performed to quantify the level of similarity within and between groups and to identify the taxa contributing most to the dissimilarities. A multi-dimensional scaling (MDS) ordination of the Bray-Curtis similarity matrix was used to visualize the relationships among the trawl set catches. To select the environmental variables (total sponge biomass, average trawl depth and bottom temperature) that best explain the similarity of the demersal fish communities, the BIOENV procedure (BIOta ENVironment matching, Euclidean distance matrix for normalized environmental data, Spearman rank correlation method, 10 restarts, 99 permutations) was used. Unfortunately, the effect of sampling year could not be robustly tested due to small sample sizes and

Table 1. List of the 34 fish taxa analyzed for fish biomass association with sponge grounds, their common names, total fish taxon biomass (total B) and percent of total biomass (%) in 104 trawl sets standardized to 1 km distance

Fish taxon	Common name	Total B (kg km ⁻¹)	%
<i>Somniosus microcephalus</i>	Greenland shark	3555.56	26.56
<i>Macrourus berglax</i>	Roughhead grenadier	2477.42	18.51
<i>Antimora rostrata</i>	Blue hake	1738.51	12.99
<i>Centroscyllium fabricii</i>	Black dogfish	1091.65	8.16
<i>Reinhardtius hippoglossoides</i>	Turbot	846.83	6.33
<i>Sebastes mentella</i>	Deepwater redfish	574.43	4.29
<i>Hippoglossoides platessoides</i>	American plaice	546.04	4.08
<i>Synphobranchus kaupii</i>	Longnose eel	543.07	4.06
<i>Coryphaenoides rupestris</i>	Roundnose grenadier	374.73	2.80
<i>Apristurus profundorum</i>	Deep-sea cat shark	185.44	1.39
<i>Bathyraxa spinicauda</i>	Spinytail skate	171.28	1.28
<i>Nezumia bairdii</i>	Common grenadier	156.29	1.17
<i>Anarhichas denticulatus</i>	Broadhead wolffish	134.40	1.00
<i>Notacanthus chemnitzii</i>	Largescaled tapirfish	112.52	0.84
<i>Amblyraja radiata</i>	Thorny skate	109.71	0.82
<i>Glyptocephalus cynoglossus</i>	Witch flounder	96.09	0.72
<i>Gaidropsarus</i> spp.	Threebeard rockling	89.31	0.67
<i>Hydrolagus affinis</i>	Deepwater chimaera	87.46	0.65
<i>Amblyraja jenseni</i>	Jensen's skate	81.82	0.61
<i>Dipturus linteus</i>	White skate	62.07	0.46
Myctophidae	Lanternfishes	51.96	0.39
Notacanthidae	Spiny eels	38.78	0.29
<i>Lycodes</i> spp.	Eelpout	37.96	0.28
<i>Lycodes vahlii</i>	Vahl's eelpout	37.37	0.28
<i>Serrivomer beanii</i>	Shortnose snipe eel	31.25	0.23
<i>Harriotta raleighana</i>	Longnose chimaera	25.57	0.19
<i>Phycis chesteri</i>	Longfin hake	19.33	0.14
<i>Bathylagus euryops</i>	Goitre blacksmelts	19.22	0.14
<i>Anarhichas minor</i>	Spotted wolffish	16.41	0.12
<i>Bathytroctes</i> spp.	Black herring	16.39	0.12
<i>Amblyraja hyperborea</i>	Arctic skate	16.30	0.12
<i>Chauliodus sloani</i>	Viperfish	16.19	0.12
<i>Rajella bathyphilia</i>	Abyssal skate	12.63	0.09
<i>Stomias boa ferox</i>	Boa dragonfish	11.31	0.08

Table 2. List of the 34 fish taxa analyzed for fish abundance association with sponge grounds, their common names, total fish taxon abundance (total A) and percent of total abundance (%) in 104 trawl sets standardized to 1 km distance

Fish taxon	Common name	Total A (km ⁻¹)	%
Myctophidae	Lanternfishes	10214.69	25.25
<i>Antimora rostrata</i>	Blue hake	7006.51	17.32
<i>Macrourus berglax</i>	Roughhead grenadier	4875.68	12.05
<i>Synphobranchus kaupii</i>	Longnose eel	4038.52	9.98
<i>Coryphaenoides rupestris</i>	Roundnose grenadier	3435.24	8.49
<i>Nezumia bairdii</i>	Common grenadier	1645.77	4.07
<i>Sebastes mentella</i>	Deepwater redfish	1570.75	3.88
<i>Centroscyllium fabricii</i>	Black dogfish	961.89	2.38
<i>Reinhardtius hippoglossoides</i>	Turbot	961.67	2.38
<i>Hippoglossoides platessoides</i>	American plaice	897.63	2.22
<i>Stomias boa ferox</i>	Boa dragonfish	867.42	2.14
<i>Bathylagus euryops</i>	Goitre blacksmelts	673.56	1.67
<i>Chauliodus sloani</i>	Viperfish	485.04	1.20
Notacanthidae	Spiny eels	333.03	0.82
Paralepididae	Barracudinas	322.96	0.80
<i>Serrivomer beanii</i>	Shortnose snipe eel	289.72	0.72
<i>Gaidropsarus</i> spp.	Threebeard rockling	283.74	0.70
<i>Glyptocephalus cynoglossus</i>	Witch flounder	209.09	0.52
<i>Notacanthus chemnitzii</i>	Largescale tapirfish	160.21	0.40
<i>Lycodes vahlii</i>	Vahl's eelpout	156.67	0.39
Polymixiidae	Beardfishes	114.87	0.28
<i>Lycodes</i> spp.	Eelpout	111.25	0.28
<i>Phycis chesteri</i>	Longfin hake	110.62	0.27
<i>Apristurus profundorum</i>	Deep-sea cat shark	99.65	0.25
<i>Nemichthys scolopaceus</i>	Atlantic snipe eel	97.60	0.24
<i>Malacosteus niger</i>	Loosejaw	89.99	0.22
<i>Chiasmodon niger</i>	Black swallower	83.95	0.21
<i>Bathytroctes</i> spp.	Black herring	61.06	0.15
<i>Polyacanthonotus rissoanus</i>	Shortspine tapirfish	58.04	0.14
Alepocephalidae	Smoothheads	52.30	0.13
<i>Amblyraja radiata</i>	Thorny skate	49.92	0.12
<i>Cottunculus microps</i>	Polar deep-sea sculpin	48.14	0.12
Stomiinae	Scaled dragonfishes	44.28	0.11
<i>Anarhichas denticulatus</i>	Broadhead wolffish	42.17	0.10

empty cells (Supplement 1). This procedure maximizes the rank correlation between the respective similarity matrices, and all permutations of the variables are assessed. The analyses and community analyses described above were performed with Primer v.6.1.5 software (2006 Primer-E).

RESULTS

There was a significant difference in depth among the sponge catch weight classes ($p < 0.0001$), and post hoc analyses showed that this difference was due to a shallower average trawl depth in the low sponge catch weight class than in the medium and high classes, which were not significantly different

from each other. The average depths \pm SD (range) for each class were: low: 981 ± 247 m (578 to 1420 m), medium: 1165 ± 205 m (589 to 1385 m) and high: 1237 ± 168 m (827 to 1446 m).

Thirty-four fish taxa had biomass values $>0.1\%$ of the total biomass for the trawl sets (Table 1). Total fish biomass (kg km⁻¹) summed across the 104 trawl sets for the 34 selected taxa (Supplement 1) was 13514.92 kg. Greenland shark *Somniosus microcephalus* accounted for 26% of the total biomass in the data, with roughhead grenadier *Macrourus berglax* accounting for 19%. Eleven taxa accounted for 90% of the total biomass in these sets (Table 1).

A different set of 34 fish taxa showed abundance values $>0.1\%$ of the total abundance for the trawl sets (Table 2). The total number of fish caught (km⁻¹)

summed across the 104 trawl sets for the 34 selected taxa was 40 453.63. Eleven taxa accounted for 90 % of the total abundance, with lanternfish (Myctophidae) being the largest taxon and accounting for 25 % of total abundance (Table 2). Blue hake *Antimora rostrata* was the second most abundant taxon, and this species also ranked high in total biomass (Table 1).

The total number of fish taxa and the total fish biomass were significantly different among the sponge catch weight classes ($p < 0.001$). Post hoc analyses showed that fish biomass was not significantly different between the medium and high classes, but that the low class had significantly higher fish biomass. The total number of fish taxa was significantly lower in the high sponge catch weight class and did not differ between medium and low classes. Total fish abundance did not differ significantly among sponge catch weight classes ($p = 0.079$).

Linear regressions between fish biomass and sponge biomass by depth were statistically significant ($p < 0.0001$). Sponge biomass increased signifi-

cantly with depth ($R^2 = 0.197$), while fish biomass significantly decreased ($R^2 = 0.182$). In contrast, the total number of fish taxa and total fish abundance were not significantly correlated with depth ($p = 0.249$ and $p = 0.797$, respectively). Total fish biomass and the total number of fish taxa were both significantly negatively correlated with total sponge biomass ($p < 0.0001$, $R^2 = 0.133$ and $R^2 = 0.209$, respectively), while there was no significant relationship with fish abundance. These p-values remained significant after correcting for multiple tests.

Community analyses: biomass

ANOSIM of the 104 trawl sets found a significant difference in community composition of the biomass between sponge catch weight classes (Global $R = 0.129$, $p = 0.001$). There was no significant difference between low and medium classes ($R = 0.056$, $p = 0.087$); however, there was a significant difference

Table 3. Taxa contributing to >90 % of the cumulative percent similarity in the biomass (kg km^{-1}) of research vessel trawl set catch composition (2001 to 2007) within each of the low, medium and high sponge catch weight classes

Species	Common name	Average similarity	% contribution	Cumulative %
Low sponge catch weight class—average similarity overall: 54.85				
<i>Macrourus berglax</i>	Roughhead grenadier	12.52	22.83	22.83
<i>Antimora rostrata</i>	Blue hake	10.79	19.68	42.50
<i>Reinhardtius hippoglossoides</i>	Turbot	8.06	14.69	57.20
<i>Synaphobranchus kaupii</i>	Longnose eel	4.76	8.68	65.88
<i>Centroscyllium fabricii</i>	Black dogfish	3.56	6.50	72.38
<i>Coryphaenoides rupestris</i>	Roundnose grenadier	3.48	6.35	78.73
<i>Nezumia bairdii</i>	Common grenadier	2.91	5.31	84.04
Myctophidae	Lanternfishes	1.46	2.66	86.70
<i>Notacanthus chemnitzii</i>	Largescaled tapirfish	1.28	2.34	89.04
<i>Gaidropsarus</i> spp.	Threebeard rockling	1.15	2.10	91.14
Medium sponge catch weight class—average similarity overall: 57.59				
<i>Antimora rostrata</i>	Blue hake	14.10	24.48	24.48
<i>Macrourus berglax</i>	Roughhead grenadier	13.34	23.16	47.63
<i>Synaphobranchus kaupii</i>	Longnose eel	6.50	11.29	58.93
<i>Reinhardtius hippoglossoides</i>	Turbot	5.51	9.56	68.49
<i>Centroscyllium fabricii</i>	Black dogfish	3.94	6.84	75.33
<i>Coryphaenoides rupestris</i>	Roundnose grenadier	3.82	6.64	81.97
<i>Apristurus profundorum</i>	Deep-sea cat shark	1.69	2.93	84.90
<i>Nezumia bairdii</i>	Common grenadier	1.59	2.76	87.67
<i>Gaidropsarus</i> spp.	Threebeard rockling	1.52	2.64	90.31
High sponge catch weight class—average similarity overall: 49.73				
<i>Macrourus berglax</i>	Roughhead grenadier	17.77	35.74	35.74
<i>Reinhardtius hippoglossoides</i>	Turbot	9.04	18.18	53.91
<i>Antimora rostrata</i>	Blue hake	8.66	17.41	71.32
<i>Centroscyllium fabricii</i>	Black dogfish	2.90	5.83	77.15
<i>Synaphobranchus kaupii</i>	Longnose eel	2.88	5.79	82.93
<i>Coryphaenoides rupestris</i>	Roundnose grenadier	2.87	5.77	88.70
<i>Apristurus profundorum</i>	Deep-sea cat shark	1.81	3.64	92.34

between the low and high classes ($R = 0.232$, $p = 0.001$) and between the medium and high classes ($R = 0.122$, $p = 0.001$) even after adjusting the p-value for multiple tests. SIMPER indicated an overall similarity of the fish assemblages within groups of 54%. Similarities within the low (55%), medium (58%) and high (50%) classes were not widely variant and included many of the same taxa in differing proportions. The taxa which accounted for 90% of the similarity in biomass within each class are listed in Table 3. Average dissimilarity in the fish communities between the low and medium sponge weight classes was 46%, between the low and high classes 52% and between the medium and high classes 49%. The taxa contributing to the dissimilarity between the low and high classes are shown in Table 4. Of the 22 taxa accounting for 90% of the dissimilarity in biomass between these 2 classes, black dogfish *Centroscyllium fabricii*, blue hake and longnose eel *Synaphobranchus kaupii* accounted for a quarter of the dissimilarity (Table 4). All had higher biomass in the low sponge catch weight class, although they were also found in the high class. Only 6 taxa showed increased biomass in the high class, namely the deep-sea cat shark, spinytail skate *Bathyrāja spinicauda*, white skate *Dipturus linteus*, shortnose snipe eel *Serrivomer beanii*, eelpout *Lycodes* spp. and deepwater chimaera *Hydrolagus affi-*

nis (Table 4). Another 4 taxa were never found in association with high sponge catches, namely deepwater redfish *Sebastes mentella*, American plaice *Hippoglossoides platessoides*, witch flounder *Glyptocephalus cynoglossus* and thorny skate (*Amblyraja radiata*) (Table 4). Most other taxa showed increased biomass in the low class to varying degrees. These relationships are visualized in the MDS plot (Fig. 2), with trawl sets with an average similarity of 54% circled. Owing to the high stress of the 2-dimensional presentation (0.20), the 3-dimensional presentation is also shown (stress = 0.15). Although some sets with high sponge catch cluster together, most of the trawl sets share a similarity of taxa that is not explained by the sponge catch weight class (Fig. 2). The relative proportions of biomass for selected pairs of taxa are indicated in the MDS ordination in Fig. 3. Each graph highlights a taxon which has increased biomass in the high or low sponge catch weight class and shows the discreteness of the groups. The BIOENV analysis identified 2 variables, total sponge biomass and average trawl depth, as having the largest rank correlation with the fish community patterns of similarity ($\rho = 0.490$, $p = 0.001$). All 3 variables together produced a lower correlation ($\rho = 0.450$), while average trawl depth had the highest rank correlation of the variables tested singly ($\rho = 0.446$).

Table 4. Taxa contributing to >90% of the dissimilarity in the biomass (kg km^{-1}) of research vessel trawl set catch composition (2001 to 2007) between the low and high sponge catch weight classes. Arrowheads indicate the direction of change in biomass from low to high sponge catch weight class. ▼: decrease in biomass; ▼▼: decrease to zero biomass; ▲: increase in biomass

Taxon	Common name	Direction of change	Ave. \log_{10} biomass		% contr. to diss.	Cum. contr. to diss.
			Low	High		
<i>Centroscyllium fabricii</i>	Black dogfish	▼	1.53	1.22	9.23	9.23
<i>Antimora rostrata</i>	Blue hake	▼	2.58	2.13	7.83	17.05
<i>Synaphobranchus kaupii</i>	Longnose eel	▼	1.50	0.96	6.67	23.73
<i>Reinhardtius hippoglossoides</i>	Turbot	▼	2.10	1.76	6.04	29.77
<i>Macrourus berglax</i>	Roughhead grenadier	▼	3.00	2.79	5.87	35.64
<i>Coryphaenoides rupestris</i>	Roundnose grenadier	▼	1.24	0.74	5.74	41.38
<i>Apristurus profundorum</i>	Deep-sea cat shark	▲	0.29	0.93	5.08	46.46
<i>Nezumia bairdii</i>	Common grenadier	▼	0.95	0.37	4.54	51.00
<i>Sebastes mentella</i>	Deepwater redfish	▼▼	0.76	0.00	4.17	55.17
<i>Hippoglossoides platessoides</i>	American plaice	▼▼	0.79	0.00	4.12	59.29
<i>Anarhichas denticulatus</i>	Broadhead wolffish	▼	0.64	0.15	3.94	63.23
<i>Bathyrāja spinicauda</i>	Spinytail skate	▲	0.34	0.42	3.73	66.96
Notacanthidae	Spiny eels	▼	0.59	0.31	3.65	70.61
<i>Gaidropsarus</i> spp.	Threebeard rockling	▼	0.55	0.16	3.20	73.80
Myctophidae	Lanternfishes	▼	0.47	0.11	2.53	76.34
<i>Glyptocephalus cynoglossus</i>	Witch flounder	▼▼	0.44	0.00	2.34	78.68
<i>Dipturus linteus</i>	White skate	▲	0.03	0.41	2.25	80.93
<i>Amblyraja radiata</i>	Thorny skate	▼▼	0.40	0.00	2.16	83.10
<i>Serrivomer beanii</i>	Shortnose snipe eel	▲	0.18	0.32	1.96	85.05
<i>Lycodes</i> spp.	Eelpout	▲	0.09	0.33	1.94	86.99
<i>Hydrolagus affinis</i>	Deepwater chimaera	▲	0.01	0.33	1.61	88.60
<i>Amblyraja jenseni</i>	Jensen's skate	▼	0.19	0.10	1.61	90.21

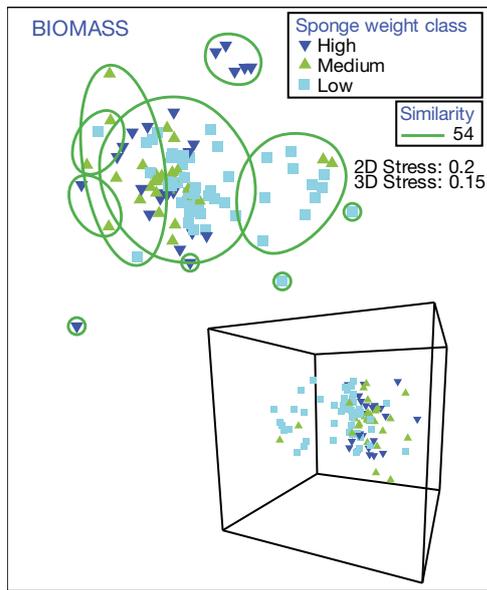


Fig. 2. Multidimensional scaling (MDS) configuration of the trawl catches (2001 to 2007) in 2D and 3D based on the Bray-Curtis similarity matrix calculated from \log_{10} -transformed biomass (kg km^{-1}) data for each of 34 fish taxa. Trawl set catches are labelled according to sponge catch weight class. In the 2D representation, trawl sets with 54% similarity to each other are indicated. This is the average similarity level within each of the 3 classes (see 'Results: Community analyses: biomass')

Community analyses: abundance

ANOSIM of the 104 trawl sets found a significant difference in community composition in abundance of taxa between sponge catch weight classes (Global $R = 0.245$, $p = 0.001$), and between all weight class pairs. However, there was a low R (0.083) produced between the low and medium classes, and the associated probability was 0.034, a value which would not be considered significant if corrected for the total number of tests performed. SIMPER indicated an overall similarity of the fish communities within groups of 59%; however, there was greater similarity within the low (63%) and medium (63%) classes than within the high class (51%). The taxa which account for 90% of the similarity in abundance within each class are listed in Table 5. Average dissimilarity in the fish communities between the low and medium classes was 39%, between the low and high classes was 52% and between the medium and high classes was 48%. The taxa contributing to the dissimilarity between the low and high classes are provided in Table 6. Of the 25 taxa accounting for 90% of the dissimilarity in abundance between these 2 classes, lanternfish, common grenadier *Nezamia bairdii* and roundnose grenadier

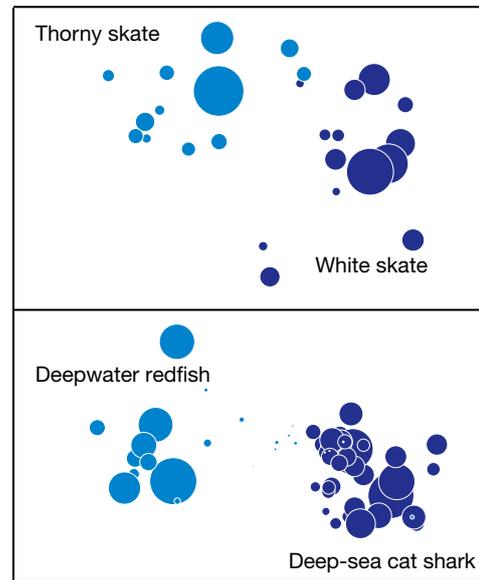


Fig. 3. MDS configuration of the trawl catches (2001 to 2007) based on the Bray-Curtis similarity matrix calculated from \log_{10} -transformed biomass data for each of 34 taxa. The proportional biomass (kg km^{-1}) of pairs of taxa which favour the sponge grounds (high sponge catch weight class; dark blue) or the low sponge areas (light blue) are illustrated. Note that proportions are relative to each taxon, and the values have not been added to allow comparisons as the purpose was largely to illustrate the non-overlap of the selected taxa

Coryphaenoides rupestris accounted for a quarter of the dissimilarity (Table 6). These all have higher abundance in the low sponge catch weight class than in the high class. The only taxa which had increased abundance in the high class were the deep-sea cat shark, eelpout and the shortnose snipe eel. All other taxa ($N = 22$) had decreased abundance in the high class, and 4 taxa were not found in that class: deep-water redfish, American plaice, witch flounder and Vahl's eelpout *Lycodes vahlii* (Table 6). The relationships between the trawl sets labelled by sponge catch weight class are visualized in the MDS plot in Fig. 4, with trawl sets with an average similarity of 59% circled. It can be seen that, although some sets with high sponge catch cluster together, most of the trawl sets share a similarity of taxa that is not explained by the sponge catch weight classes (Fig. 4). Total sponge biomass and average trawl depth, which had the highest correlation in the BIOENV analyses with the pattern of similarity in fish biomass, also had the largest rank correlation with the fish community pattern calculated from abundance ($\rho = 0.558$, $p = 0.001$). However, in contrast to the analyses of biomass, total sponge biomass had the highest rank correlation of the variables tested singly ($\rho = 0.383$).

Table 5. Taxa contributing to >90 % of the cumulative % similarity in the abundance (km^{-1}) of research vessel trawl set catch composition (2001 to 2007) within each of the low, medium and high sponge catch weight classes

Species	Common name	Average similarity	% contribution	Cumulative %
Low sponge catch weight class—average similarity overall: 62.58				
Myctophidae	Lanternfish	9.45	15.10	15.10
<i>Antimora rostrata</i>	Blue hake	9.27	14.81	29.91
<i>Macrourus berglax</i>	Roughhead grenadier	7.75	12.38	42.29
<i>Synaphobranchus kaupii</i>	Longnose eel	6.01	9.60	51.89
<i>Coryphaenoides rupestris</i>	Roundnose grenadier	4.73	7.55	59.45
<i>Nezumia bairdii</i>	Common grenadier	4.47	7.14	66.59
<i>Reinhardtius hippoglossoides</i>	Turbot	4.09	6.54	73.13
<i>Stomias boa ferox</i>	Boa dragonfish	3.36	5.37	78.50
<i>Chauliodus sloani</i>	Viperfish	1.67	2.66	81.16
<i>Gaidropsarus</i> spp.	Threebeard rockling	1.61	2.57	83.73
Paralepididae	Barracudinas	1.58	2.52	86.25
<i>Bathylagus euryops</i>	Goitre blacksmelts	1.55	2.48	88.73
<i>Centroscyllium fabricii</i>	Black dogfish	1.55	2.48	91.21
Medium sponge catch weight class—average similarity overall: 63.13				
<i>Antimora rostrata</i>	Blue hake	10.51	16.64	16.64
<i>Macrourus berglax</i>	Roughhead grenadier	8.86	14.04	30.69
<i>Synaphobranchus kaupii</i>	Longnose eel	7.47	11.84	42.52
Myctophidae	Lanternfishes	6.74	10.68	53.21
<i>Coryphaenoides rupestris</i>	Roundnose grenadier	5.49	8.69	61.90
<i>Chauliodus sloani</i>	Viperfish	3.46	5.49	67.39
<i>Nezumia bairdii</i>	Common grenadier	2.76	4.37	71.75
<i>Reinhardtius hippoglossoides</i>	Turbot	2.73	4.33	76.08
<i>Stomias boa ferox</i>	Boa dragonfish	2.51	3.97	80.05
<i>Gaidropsarus</i> spp.	Threebeard rockling	1.87	2.97	83.02
<i>Centroscyllium fabricii</i>	Black dogfish	1.74	2.75	85.76
<i>Lycodes</i> spp.	Eelpout	1.14	1.80	87.56
<i>Serrivomer beanii</i>	Shortnose snipe eel	1.10	1.74	89.31
Polymixiidae	Beardfishes	1.02	1.61	90.92
High sponge catch weight class—average similarity overall: 51.13				
<i>Macrourus berglax</i>	Roughhead grenadier	14.55	28.46	28.46
<i>Antimora rostrata</i>	Blue hake	8.34	16.30	44.76
<i>Coryphaenoides rupestris</i>	Roundnose grenadier	5.56	10.88	55.64
<i>Synaphobranchus kaupii</i>	Longnose eel	5.07	9.92	65.56
<i>Reinhardtius hippoglossoides</i>	Turbot	4.96	9.70	75.26
<i>Centroscyllium fabricii</i>	Black dogfish	2.01	3.93	79.19
<i>Nezumia bairdii</i>	Common grenadier	1.89	3.69	82.88
<i>Serrivomer beanii</i>	Shortnose snipe eel	1.60	3.14	86.02
<i>Bathylagus euryops</i>	Goitre blacksmelts	0.98	1.92	87.94
Myctophidae	Lanternfishes	0.86	1.67	89.61
<i>Apristurus profundorum</i>	Deep-sea cat shark	0.85	1.66	91.27

DISCUSSION

Temperate sponge grounds are considered vulnerable marine ecosystems (sensu FAO 2009) in the northwest Atlantic (Fuller et al. 2008), and in 2009 NAFO implemented Sponge Protection Zones in the international waters of the Grand Banks (NAFO 2009). Their upright nature and poor recovery potential if dislodged from the seafloor render them vulnerable to bottom-contact fishing gear. Large structure-forming sponges are considered to be

ecosystem engineers through their modification of the seafloor (through altering bottom currents and depositing spicules), and, by adding structural complexity, provide habitat for a wide variety of benthic invertebrates (Klitgaard 1995). However, the association of demersal fish communities with these temperate sponge grounds has not been well documented. Our results show that the high biomass sponge grounds on the continental slopes off Newfoundland, Canada, have lower biomass and numbers of taxa of demersal fish relative to areas with low sponge bio-

Table 6. Taxa contributing to >90% of the dissimilarity in abundance (km^{-1}) of research vessel catch composition (2001 to 2007) between the low and high sponge catch weight classes. Arrowheads indicate the direction of change in fishtaxon abundance in going from the low to the high sponge catch weight class. ▼: decrease in abundance; ▼▼: decrease to zero abundance; ▲: increase in abundance

Taxon	Common name	Direction of change	Ave. \log_{10} abundance		% contr. to diss.	Cum. contr. to diss.
			Low	High		
Myctophidae	Lanternfishes	▼	4.46	0.93	11.61	11.61
<i>Nezumia bairdii</i>	Common grenadier	▼	2.58	1.11	5.95	17.55
<i>Coryphaenoides rupestris</i>	Roundnose grenadier	▼	2.90	2.01	5.81	23.36
<i>Synaphobranchus kaupii</i>	Longnose eel	▼	3.21	2.23	5.80	29.16
<i>Stomias boa ferox</i>	Boa dragonfish	▼	2.02	0.55	5.28	34.44
<i>Antimora rostrata</i>	Blue hake	▼	4.13	3.06	5.22	39.66
<i>Centroscyllium fabricii</i>	Black dogfish	▼	1.41	1.24	4.68	44.34
<i>Bathylagus euryops</i>	Goitre blacksmelts	▼	1.44	0.74	4.35	48.69
<i>Sebastes mentella</i>	Deepwater redfish	▼▼	1.17	0.00	3.66	52.35
Paralepididae	Barracudinas	▼	1.24	0.28	3.55	55.90
<i>Reinhardtius hippoglossoides</i>	Turbot	▼	2.20	1.61	3.48	59.38
<i>Chauliodus sloani</i>	Viperfish	▼	1.28	0.60	3.45	62.84
<i>Serrivomer beanii</i>	Shortnose snipe eel	▲	0.79	1.04	3.20	66.04
<i>Macrourus berglax</i>	Roughhead grenadier	▼	3.66	3.48	3.08	69.12
<i>Gaidropsarus</i> spp.	Threebeard rockling	▼	1.11	0.45	2.99	72.11
<i>Hippoglossoides platessoides</i>	American plaice	▼▼	0.89	0.00	2.70	74.81
<i>Notacanthus chemnitzii</i>	Largescale tapirfish	▼	0.78	0.28	2.32	77.13
<i>Apristurus profundorum</i>	Deep-sea cat shark	▲	0.19	0.69	2.04	79.17
<i>Lycodes</i> spp.	Eelpout	▲	0.25	0.54	1.85	81.02
<i>Glyptocephalus cynoglossus</i>	Witch flounder	▼▼	0.59	0.00	1.82	82.84
<i>Malacosteus niger</i>	Loosejaw	▼	0.44	0.37	1.75	84.59
<i>Chiasmodon niger</i>	Black swallower	▼	0.41	0.31	1.57	86.16
<i>Nemichthys scolopaceus</i>	Atlantic snipe eel	▼	0.46	0.15	1.57	87.73
<i>Phycis chesteri</i>	Longfin hake	▼	0.51	0.03	1.55	89.27
<i>Lycodes vahlii</i>	Vahl's eelpout	▼▼	0.47	0.00	1.41	90.69

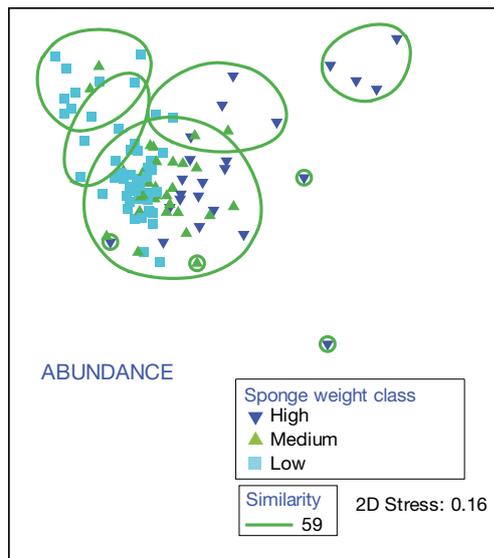


Fig. 4. MDS configuration of the trawl catches (2001 to 2007) in 2 dimensions (2D) based on the Bray-Curtis similarity matrix calculated from \log_{10} -transformed abundance data for each of 34 taxa. Trawl catches are labelled according to sponge catch weight class. Trawl sets with 59% similarity to each other are indicated. This is the average similarity level within each of the 3 classes (see 'Results: Community analyses: abundance')

mass outside of the sponge grounds, a correlation not reflected in the abundance of fish. This scenario is created by small numbers of the larger bodied fish living where sponge biomass is significantly lower and where water depth is also shallower.

Although we attempted to narrow the depth range of our samples *a priori* by selecting trawl sets from the continental slope (578 to 1446 m) for analysis, depth together with sponge biomass explained the highest percentage of the variation in the fish communities, whether measured by biomass or abundance. Bottom temperature was not an important determinant either on its own or in combination with depth and sponge biomass. A significantly distinct fish assemblage based on both biomass and abundance metrics was associated with the sponge grounds (trawl sets with high sponge biomass $\geq 250 \text{ kg km}^{-1}$), despite the statistically similar depth of the trawl sets with medium sponge biomass (10 to 250 kg km^{-1} exclusive). The coherence amongst trawl set fish composition is relatively high for both biomass and abundance at about 50% similarity over the sponge grounds. Species which form the fish assemblage associated with the sponge grounds are

the roughhead grenadier, roundnose grenadier, blue hake, longnose eel, turbot *Reinhardtius hippoglossoides*, black dogfish and deep-sea cat shark. SIMPER showed these fish to characterize sponge grounds both in terms of biomass and abundance, contributing to 92 and 81 %, respectively, of the similarity among trawl set fish composition. An additional 4 taxa, common grenadier, shortnose snipe eel, lanternfish and goitre blacksmelts *Bathylagus euryops*, were identified through characterization of abundance only, contributing an additional 10% of the similarity among trawl sets. The association of these demersal fish with the sponge grounds further strengthens the role that sponge grounds play in marine ecosystems.

Only 6 taxa had higher abundance and/or biomass in trawl sets with high sponge catches compared to sets with low sponge catches ($\leq 10 \text{ kg km}^{-1}$) (sponge grounds vs. non-sponge grounds). None are targeted in commercial fisheries in this area. These were deep-sea cat shark, spinytail skate, white skate, shortnose snipe eel, eelpout and deepwater chimaeras (Table 7), although none were exclusive to those sets. Only 2 (deep-sea cat shark and shortnose snipe eel) contributed to the SIMPER description of similarity amongst fish caught in the high sponge bycatch sets. This is because the remainder were also found in sets with medium sponge catch. All of these are deep-living fish (Rose 2005), and their associations with the sponges may be a coincidental one with depth, particularly for the 4 taxa that also occurred with medium sponge catches, as there was no significant difference in the depth of trawl sets between the medium and high sponge catch weight classes. However, all 6 taxa are also active predators, which may indicate a foraging relationship with the sponge grounds.

Previous studies have shown that the invertebrate fauna associated with sponge grounds has a higher species diversity compared to surrounding bottoms. This associated fauna is largely comprised of other smaller sponges (ICES 2009), hydroids, zoanthids, bryozoans and ascidians (Klitgaard 1995, Klitgaard & Tendal 2004). Klitgaard (1995) identified 242 taxa of epifauna and infauna associated with the sponge grounds, dominated by *Geodia* spp. and *Strychnus* spp. near the Faroe Islands, and considered that almost half of these formed obligate associations with the sponges. The spicule mats associated with the sponge communities' support increased biomass of macrofaunal species (Bett & Rice 1992). Preliminary evidence suggests that the sponge grounds in the northwest Atlantic also host a high diversity of ben-

thic invertebrates (NAFO 2010, Barrio Froján et al. 2012), and foraging may be the mechanism which has produced the associated fish assemblage observed in our study. Eelpout are known to eat sponge remains (Table 7; Coad & Reist 2004) and so may feed directly on the sponges in the sponge grounds. The diet of shortnose snipe eel consists of crustaceans, cephalopods and teleosts (Geidner 2008). *Harriotta raleighana*, a chimaera related to the deep-water chimera, feeds on endo- and epibenthic organisms on the Grand Bank and at Flemish Cap (González et al. 2007). They found that *H. raleighana* in particular consumed a high proportion of benthic polychaetes and associated sediment. Little information is available on the diets of white or spinytail skate, but, in the Barents Sea, the white skate feeds on benthic invertebrates, including brittlestars, gammarids and shrimp, while the spinytail is largely piscivorous as an adult (Dolgov 2005), but feeds mainly on benthos as a juvenile (Dolgov 2002). Similarly, the diet of the deep-sea cat shark is not well documented, although congeneric species feed heavily on crustaceans (Cortés 1999), especially euphausiids and penaeid shrimp. Therefore, all of the 6 taxa identified as having a high biomass/abundance in association with the sponge grounds (Table 7) have dietary links to that habitat. The other fish taxa identified by SIMPER as descriptive of the high-density sponge grounds are also benthic feeders, that is, the roughhead grenadier (Eliassen & Jobling 1985), black dogfish (González et al. 2007), blue hake (Mauchline & Gordon 1984), turbot (Rodríguez-Marín et al. 1995) and juvenile longnose eel (Marques 1998), with only the lanternfish and goitre blacksmelts being predominantly pelagic feeders. However, most demersal fish are benthivores; thus, more direct evidence is needed before conclusions can be drawn on how these fish may use the habitat offered by the sponge grounds.

There is no strong evidence for fish avoidance of the sponge grounds. The lower biomass of black dogfish, blue hake and longnose eel and the lower abundance of lanternfish, common grenadier and roundnose grenadier drove the distinction in the fish assemblages between trawl sets with high sponge biomass in the catch and those with low sponge catch, accounting for a quarter of the dissimilarity in both instances. These taxa only differ between the sponge weight classes in terms of relative biomass as all are found in both types of trawl sets. Only 5 species common in the low sponge biomass trawl sets were never reported in the catches with high sponge biomass and had only a small proportion of their bio-

Table 7. Summary of known habitat and diet characteristics of those fish taxa showing a positive association with sponge grounds in biomass (B) and/or abundance (A) and those never captured with high sponge catches in the study (Tables 4 & 6). This table was produced using information recorded on FishBase (www.fishbase.org) and from Coad & Reist (2004) and Troyanovsky (1992)

Common name (taxon)	Metric response	Zone	Ecology	Diet	Depth information
Fish taxa showing a positive association with sponge grounds in biomass and/or abundance					
Shortnose snipe eel (<i>Serrivomer beanii</i>)	Increase B, increase A	Epibenthic-mesopelagic	Exhibits vertical migrations to feed during the night	Shrimps, other crustaceans, small fish	To 5998 m, Canadian Atlantic: 850–925 m. Mostly at depths of 550–1000 m; present study (N = 57 sets): range = 768–1446 m median = 1211 m
Deep-sea cat shark (<i>Apristurus profundorum</i>)	Increase B, increase A	Bathydemersal, inhabits the continental slope	Relatively small, sluggish sharks; egg cases may be attached to biogenic structures as they are laid	Euphausiids, penaeid shrimp, crustaceans	1100–1750 m; present study (N = 37 sets): range = 871–1446 m median = 1303 m
Eelpout (<i>Lycodes</i> spp.)	Increase B, increase A	Demersal	Muddy bottoms. It seems to get the bulk of its food by burrowing in the sediment.	Small bivalves, polychaetes, small crustaceans. <i>L. terraenovae</i> eats sponge	19–1750 m; present study (N = 40 sets): range = 589–1410 m median = 1164 m
Spinytail skate (<i>Bathyraja spinicauda</i>)	Increase B	Benthic, bathydemersal	Boreal and arctic species found in water temperatures <8°C	Invertebrates as juveniles, fish as adults	140–1463 m, usually 65–255 m; present study (N = 17 sets): range = 601–1384 m, median = 1202 m
White skate (<i>Dipturus linteus</i>)	Increase B	Benthic, bathydemersal	Little known species is widespread but distributed deeply	Worms, crustaceans, fishes	150–1455 m, usually 250 m; present study (N = 14 sets): range = 1153–1446 m, median = 1357 m
Deepwater chimaera (<i>Hydrolagus affinis</i>)	Increase B	Epibenthic, bathydemersal	Deepwater slope, seamount and sea plain dwelling fish	Fish, invertebrates	Deep waters to 2400 m; 300–2400 m; present Study (N = 9 sets): range = 1058–1370 m, median = 1243 m
Fish taxa never captured with high sponge catches in the present study					
Deepwater redfish (<i>Sebastes mentella</i>)	Decrease B, decrease A	Epibenthic–pelagic, bathypelagic	Long-lived species (>65 yr)	Crustaceans, fish	300–1441 m; 500–700 m in NAFO Div.2J,3K; present study (N = 27 sets): range = 578–1243 m, median = 812 m
American plaice (<i>Hippoglossoides platessoides</i>)	Decrease B, decrease A	Benthic, demersal	Soft bottoms	Worms, molluscs, echinoderms, crustaceans, fish	10–3000 m, usually 9–250 m; present study (N = 18 sets): range = 589–1161 m, median = 739 m
Witch flounder (<i>Glyptocephalus cynoglossus</i>)	Decrease B, decrease A	Benthic, demersal	Soft mud bottoms in fairly deep water	Worms, crustaceans, molluscs, brittle stars, fish	8–1570 m, usually 45–366 m; present study (N = 14 sets): range = 578–1384 m, median = 622 m
Vahl's eelpout (<i>Lycodes vahlii</i>)	Decrease A	Benthic, bathydemersal	Lives at temperatures of 0–5°C	Worms, crustaceans, molluscs	65–1200 m; present study (N = 12 sets): range = 578–985 m, median = 630 m
Thorny skate (<i>Amblyraja radiata</i>)	Decrease B	Benthic, demersal	Sandy and muddy bottoms	Worms, crustaceans, fish	20–1000 m; present study (N = 12 sets): range = 578–1190 m, median = 655 m

mass/abundance in areas with medium sponge biomass: deepwater redfish, American plaice, witch flounder, Vahl's eelpout and thorny skate. These fish could be avoiding the sponges, as has been noted elsewhere (Bell 2008). American plaice, thorny skate and witch flounder are associated with mud or sandy bottoms and may actively avoid the sponge grounds which can have dense spicule mats forming biogenic substratum (Bett & Rice 1992) and creating a heterogeneous surface. Further, some of these fish tend to prefer shallower waters (Table 7), and the significantly shallower average depth of the low sponge catch weight class may contribute to our findings.

Collectively our data suggest that the *Geodia*-dominated sponge grounds of the continental slopes off Newfoundland, northwest Atlantic, host distinctive fish assemblages, although the active or passive nature of this association is not known. We speculate that these fish use the sponge grounds to forage, although they may only be taking advantage of the presence of the physical structure created by the sponges. A more detailed analysis of these data using less coarse taxonomic categories may have revealed greater differences in community composition. However, trawl survey catch can only give a generalized picture of the species associations for those species that are caught by the gear. Other sampling tools are required to provide details for the smaller invertebrate and fish species and life-history stages that have been reported elsewhere as being associated with sponge grounds. Future research focussing on the species identified to be associated with the sponge grounds may provide greater insight into the strength and temporal stability of the associations and the ecological linkages between these benthic and pelagic communities.

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