

Comparative estimates of groundfish biomass, abundance, and diversity using various optical sampling methods

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Abstract

TBD

Introduction

Many managed groundfish stocks on the U.S. west coast inhabit high-relief rocky areas that are inaccessible to traditional net-based survey methods (e.g., swept-area benthic trawls). Several species, such as cowcod (*Sebastes levis*) and yelloweye rockfish (*S. ruberrimus*), are strongly associated with these untrawlable habitats, have populations at various levels of depletion, and occupy habitats that have incurred substantial impacts from mobile fishing gear [Did I get this correct??] (Love and Yoklavich 2006; Yoklavich et al. 2007). Since much of the continental shelf and upper slope are closed to groundfishing, effective monitoring tools are required for species living in these untrawlable habitats.

Non-lethal and non-destructive survey methods, whether optical, acoustical, or some combination of both, are needed to assess these vulnerable species while minimizing impacts on the fishes and their habitat. To that end, this study compared the results from visual transect surveys conducted in untrawlable habitat using a human-occupied submersible (SUB), an autonomous underwater vehicle (AUV), and a remotely operated vehicle (ROV). Specifically, each survey method was used to (1) identify, count, and estimate the sizes of rockfishes (*Sebastes* spp.; both common and rare, large- and small-bodied, and semi-pelagic and highly demersal), lingcod (*Ophiodon elongatus*), thornyheads (*Sebastolobus* sp.), and Pacific hake (*Merluccius productus*); (2) estimate the density, abundance, and biomass (and associated precision) for these taxa; and (3) estimate diversity within a prescribed study area.

Methods

Survey area

Underwater surveys of demersal fishes and their seabed habitats were conducted on two rocky seamounts (Footprint Bank and Piggy Bank) off southern California (**Figure 1**). The study site is located inside the State and Federal Footprint Marine Reserves, offshore of Santa Cruz Island, inside the Channel Islands National Marine Sanctuary. The Piggy Bank is about 30 km² in area, ranging in depth from 275 to 900 m; the Footprint Bank is about 10 km² in area, ranging in depth from 80 to 500 m. High-resolution bathymetry and broad-scale seabed classification from multibeam echosounder backscatter data were available prior to our surveys (Dartnell et al. 2005). The survey area was stratified into 100-m depth strata down to 400 m depth. The total area of each stratum was estimated using GIS (ArcMap version 9.3; Esri, Inc.) and used to estimate total biomass and abundance from estimates of fish density. Large areas of soft sediment in the northeast section of the study site were excluded from our sampling frame.

Survey design

Visual surveys were conducted using a SUB (*Dual Deepworker*, Nuytco) deployed from the F/V *Velero IV* between 21 and 30 September 2011; using a remotely operated vehicle (ROV, *Phantom DS4*, Deep Ocean Exploration and Research) deployed from the F/V *Outer Limits* during four legs between 21 September and 8 December 2011 (Leg 1: 21-22 September; Leg 2: 4 October; Leg 3: 12-13 October; and Leg 4: 4-8 December; and using an autonomous underwater vehicle (AUV, *Lucille*, SeaBED) between **21 and 30 September 2011** aboard the NOAA Vessel *Shearwater*. The sampling unit for the SUB and ROV was a transect and the sampling unit for the AUV was a transect grid (see below).

Transects were planned to occur in only one depth stratum. Some transects spanned multiple depths strata and were split into multiple transects by 100-m-depth bins. To reduce potential variability arising from very short transects, only those transects with lengths greater than 200 m were included in the analysis. Transect lengths were calculated from the speed of the ROV, and transect area was estimated every 2-s from the camera height, tilt angle, and horizontal viewing angle (Stierhoff et al., 2016).

Each SUB dive included multiple, randomly located, 15-min-long transects within each stratum (**Figure X**). The number of transects per stratum was based on optimal sample variances from past visual surveys in the

study area (**Yoklavich, unpublished data**), the area of each depth stratum (**Table 1**), and the amount of time available for the entire survey. A bootstrap analysis of coefficients of variation in density of various fish species and species richness estimated from similar SUB transects conducted in 2005 on Footprint seamount at depths < 200 m. We concluded that 15 transects produced optimal sample variances for the area within the 100-200-m stratum at Footprint, and applied that ratio (15 transects/1.23 km²) to determine the number of transects to be conducted within the other depth strata in the study area. We increased the number of transects within the <100 meter stratum based on relatively high density of fishes and species richness at that depth in the earlier survey of this area. Using ArcMap 9.3, the appropriate number of spatially random points were generated within each depth stratum to locate transects.

For the AUV survey, a 250 x 250 m grid was superimposed on the survey area, then each cell was assigned to one of three focal areas: the top of Piggy Bank, the top of Footprint Bank and the deeper flank around Footprint Bank (**Figure 2**). Cells to be sampled were then selected randomly from each area. Within each cell, the AUV was programmed to survey pattern that covered the majority of the cell at a nominal height of **X m** above the seabed. The survey pattern consisted of five 200-m long transects connected by four 25-m long lines for a total transect distance of 1.1 km in each cell sampled. If randomly selected cells were close in proximity, multiple cells could be sampled during one deployment. One to four cells were sampled per day and deployment duration varied from 45 to 88 min (mean = 74 min). Digital still images were collected every 10 s throughout each transect (see Survey Vehicles below).

Survey vehicles

All visual transect surveys were conducted during daylight hours (~06:30 to 17:00 h PST) and spanned a variety of seabed types, from flat-sandy and mud seabeds to steeply sloping, high-relief rocky seabeds.

Remotely operated vehicle

The location of the ROV above the seabed was estimated using an ultra-short baseline (USBL) acoustic tracking system (TrackLink 5000HA, LinkQuest, Inc.) and differential global positioning system (dGPS, CSI Wireless dGPS MAX). The length of each transect was estimated from the ROV speed that was measured using a Doppler velocity log (DVL, Workhorse Navigator, Teledyne RD Instruments). Water-column and near-bottom water quality parameters [e.g., temperature, salinity, dissolved oxygen (DO) concentration and DO saturation (%)] were measured during each transect using a CTD (Citadel CTD-ES, Teledyne RD Instruments) and optode (Model 3930, Aanderaa, Inc.). All data were time-stamped and logged synchronously using WinFrog integrated navigation software (Fugro Pelagos, Inc.). Reference lasers (spaced 20 and 60 cm-apart) were used to estimate fish lengths and transect widths (see Effort analysis below). All video footage was recorded to digital-video tape (DVCAM) and later used for enumerating fishes and characterizing the seabed. To aid in the identification and measurement of fishes observed on the video tapes, and also for better characterizing seabed substrates, high-quality digital still images were collected haphazardly.

The location of transects during Leg 1 (**Figure X**, orange transects) were selected based on preliminary acoustic backscatter data collected by the SWFSC's Advanced Survey Technology (AST) group between 13 and 14 September. Subsequent transects were selected at random (Legs 2-4, **Figure X**).

Human-occupied submersible

A pilot operated the untethered SUB while an experienced scientist identified all fish species and estimated their total length (TL) to the nearest **centimeter**?. Each transect was documented with two external high-definition (HD) 3-color video cameras mounted at 45° on the starboard side of the SUB, one positioned in the same direction and field of view as the observer and the other camera located below the observer's field of view to record fishes in the area closest to the SUB (that may not have been seen by the observer). The videotape was time-stamped and annotated in real-time by the scientist inside the SUB.

The location of the SUB above the seabed was estimated using a USBL (Tracklink **1500**, LinkQuest) and dGPS. All data were time-stamped and logged synchronously every 3-s using integrated navigation software (WinFrog, Fugro Pelagos). The positioning system was linked to a GIS to track the SUB in real time and so that a navigator aboard the support vessel could guide its path; the pilot and observer inside the SUB did not influence the direction of travel. During a transect, we tried to maintain a constant distance within 2 m of the seafloor and a constant speed between 0.5 and 1.0 kns, depending on substratum type (i.e., generally slower speed in complex habitats). Those segments of a transect in which the seafloor was not clearly visible were excised and not considered as part of the 15-min sample. The scientist estimated fish sizes using paired lasers spaced 20-cm apart on either side of the main survey video camera as a guide. The length of each transect was determined using a DVL (**using speed, or dead reckoning between x/y/z estimates**) and ring-laser gyrocompass (**Make, Model**) attached to the outside of the submersible. An average[**and SD/SE??**] transect width of 2.5 m was estimated by the scientific observer with the aid of a hand-held sonar, the submersible sonar, and a crossing laser set **to intersect the parallel laser** at 3 m from the observer when the submersible was 1 m above the seabed. Temperature, salinity, depth, and dissolved oxygen were measured every **X** s using a CTD (SBE-19, Seabird) and oxygen sensor(**Make, Model**).

Autonomous underwater vehicle

Stereo images of the seabed were collected every 10 s using a **calibrated?**, downward-facing stereo camera pair (5-megapixel, 12-bit dynamic range machine vision cameras; GigE, Prosilica) synced to a strobe ([**model important?**]); a third GigE camera was angled forward at an angle of 30°strobe synced with the cameras. The location of the AUV above the seabed was estimated using a USBL acoustic tracking system (TrackLink **1500/5000HA**, LinkQuest, Inc.) and dGPS (**Model, Manufacturer??**), and the depth was determined using a depth sensor (**Model?**, Paroscientific). Scientists communicated with the AUV using an acoustic modem (256008 acoustic micromodem, WHOI) at depth and a radio modem (Model FGR-115 RCRF, FreeWave) at the surface. Physical oceanographic measurements were made using an onboard CTD (Model 49 FastCat, Sea-Bird).

Visual analysis methods

All target species observed in video recordings were identified, with the aid of high-resolution still images when available, to the lowest possible taxon and counted. When fishes could not be identified to species, they were identified to the genus (e.g., unidentified rockfish, *Sebastes* spp.; or unidentified thornyhead, *Sebastolobus* spp.) or subgenus level (e.g., rosy-group rockfish, *Sebastomus*).

ROV

For each observation, total length (*TL*; cm) was estimated to the nearest 10 cm (e.g., 0-10 cm, 10-20 cm, etc.) using the parallel lasers for reference. When fish were oriented normal to the camera lens and near the reference lasers, *TL* was measured more precisely (i.e., nearest 1 cm) using image analysis package (ImageJ, National Institute of Health).

SUB

Video transects and associated audio annotations made by the observer during each dive were reviewed following the survey. All observed fishes within 2 m of the seabed were identified to lowest possible taxon, counted, and measured (to nearest 5 cm).

Characterization of seabed type

[Not sure if we'll keep this section, but I've inserted text here in case.]

ROV

SUB

Seafloor substrate was classified using in order of decreasing particle size and vertical relief (as described in Greene et al. 1999): rock (R), boulder (B), cobble (C), and mud (M). A two-character code was used to quantify patches of uniform substratum type along each transect (as described in Yoklavich et al., 2000). The primary character in the code represented the substratum type that accounted for at least 50% of the patch, and the secondary character represented the substratum type accounting for at least 20% of the patch (e.g., CM represented a patch of at least 50% cobbles and at least 20% mud). The area of each substratum patch along a transect was estimated as the product of the transect width and the length of the patch.

AUV

Seafloor habitats in each photograph were categorized using a two-character code (Table 1) The first character signified the primary habitat type that covered greater than 50% of the field of view, while the second character defined the secondary habitat type covering between 20% and 50%. If the primary habitat coverage exceeded 80%, that letter was denoted twice (e.g., CC).

Estimation of abundance and biomass

ROV

[Methods may be same for ROV and SUB] The total abundance and biomass of each species was estimated within each stratum. Total abundance (N) in each transect was estimated using the strip transect method (Buckland et al. 2001) by multiplying the mean density of each species (D ; **get correct symbol**) within a stratum by the total area (A) within that stratum (**Table X**). For each transect, density was calculated as: $D = n/a$, where n is the number of individuals encountered and a is the transect area. The total area of each depth stratum was estimated using ArcGIS (**Table 2**). The biomass (B) for each species was estimated from TL as: $B = a * (TL)^b$, where a and b are taken from published relationships between TL and mass (**Appendix/Table X**). For fishes assigned to 10-cm length bins, the midpoint of each size class was used to estimate biomass (e.g., 5 cm for the 0-10 cm class). Many species for which few or no voucher specimens are available (e.g., dwarf-red rockfish, *S. rufinanus*; rosethorn rockfish, *S. simulator*; and pygmy rockfish, *S. wilsoni*), coefficients for closely related species (as described by Hyde & Vetter 2007) were used. The relationship for vermilion rockfish (*S. miniatus*) was substituted for the newly described sunset rockfish (*S. crocotulus*; Hyde et al. 2008). Since many of the unidentified rockfishes were of the semi-pelagic, aggregating variety, coefficients for squarespot rockfishes (*S. hopkinsi*, the most common species with similar behavior and vertical distribution) were used. For unidentified rosy-group fishes (*Sebastomus*), coefficients for swordspine rockfish (*S. ensifer*, the most common *Sebastomus* species observed) was used. Mean, CV of the mean, and 90%-quantile confidence intervals for abundance and biomass were estimated using a non-parametric bootstrap of 1,000 samples (Efron & Tibshirani 1993).

SUB

To estimate total biomass of species, we used the length-weight relationship

We substituted coefficients from closely related species for chameleon (*S. phillipsi*), dwarf-red (*S. rufinanus*), and pygmy (*S. wilsoni*) rockfishes and unidentified thornyheads and *Sebastomus* because coefficients were

unavailable for these taxa. Biomass density ($\text{kg } 100 \text{ m}^{-2}$) was then calculated for each taxon on each transect by summing the weights of individuals and dividing by the transect area. The mean and variance of biomass density were calculated from transects within each depth stratum, and expanded to total biomass and variance by multiplying by the area of the depth stratum. Total biomass and variance for each bank and the banks combined were estimated by summing biomass and variance for all depth strata. Coefficient of variation (CV) was calculated as $\text{CV} = \sqrt{\text{var}(\text{B})/\text{B}}$.

AUV

Digital still images were downloaded at the end of each mission and color-corrected. All non-overlapping images from the port, downward-facing camera were reviewed to identify and count all fish observed; images from the angled camera were used to aid in the identification of fishes observed. The area of each image was estimated using the measured altitude above the seabed and the specified camera field of view.

Biodiversity estimates

ROV

Species richness (S); species diversity (Shannon H'), and Simpson's lambda (λ) were computed for each transect and then summarized within each depth stratum and at each bank. The expected species richness (rarefaction) in a particular sample from that stratum was estimated and arefaction curves were generated for each bank using the. All biodiversity estimates were generated using the *vegan* package (Oksanen et al. 2012) in R (R Development Core Team 2011).

Results

Survey effort

[Assemble a table of n transects, distance, and area for each method, bank, and depth stratum.]

ROV

A total of 37 transects were surveyed from ~90 to 390 m. The majority of transects were conducted between the depths of 100 to 300 m on the Footprint, and from 200 to 300 m on Piggy Bank (Table 2, Fig. 1). The transect length ranged from ~300 to 1300 m (mean = X m, SD = SD), area ranged from (mean = X m^2 , SD = SD), and duration ranged from r20 to 140 min (mean = X min, SD = SD). The average transect width was 2.97 (SD = 1.34). Due to technical problems with the ROV, inclement weather, or both, the ROV portion of this comparative survey had to be completed in several "legs" spanning several months between September and December. **Calculate new estimates of area using method of Stierhoff et al. 2016.**

SUB

A total of 69 transects were conducted between 95-400 m during 17 dives on Footprint (n = 55 transects) and Piggy Bank (n = 14 transects) covering a total distance of 52000 m^2 (total area = 0.05 km^2) (Table X; Figure X). Mean transect length was 294 m (SE = 8 m). Dive duration ranged from 1.4 to 3.2 h (mean = 2.3 h, SE = 0.1 h).

AUV

Twenty-seven cells were sampled throughout the survey area covering a total of 70,006 m² and ranging in depth from 99 to 487 m: 13 on Footprint Bank, six on the flank of Footprint, and eight on Piggy Bank (**Table x, Figure 2**). Seventeen of the 27 cells was entirely within one 100-m depth stratum, while ten cells spanned two 100 m strata. The range of depths surveyed within a cell varied from 10 m to 83 m. Habitats encountered ranged from high-relief rock ridge to mud/sand bottom (**Figure x**). Cells surveyed on the top of Piggy Bank (200-300 m) contained rocky ridge substrate (30%), mud (33%) and similar amounts of boulder and cobble (17 and 19%, respectively). With increasing depth, the percentage of mud and cobble substrates increased. The top two depth strata on the Footprint were primarily cobble and boulder, with sand and rock ridge increasing with depth (**Figure X**). Mud was the primary substrate observed in all depth strata on the flank of the Footprint (86, 99 and 97%).

Length frequency composition

[GENERATE LENGTH FREQUENCY PLOTS FOR EACH SURVEY METHOD] Examine whether differences in length distributions contribute to any differences in biomass or abundance.

Abundance and biomass estimates

ROV

Over 37,700 individuals from thirty-three *Sebastes* species, *O. elongatus*, and *M. productus* were counted during this survey. The groundfish community throughout the survey area was numerically dominated by four small, semi-pelagic and aggregating species (squarespot rockfish, *S. hopkinsi*; swordspine rockfish, *S. ensifer*; halfbanded rockfish, *S. semicinctus*; and shortbelly rockfish, *S. jordani*) that comprised ~80% of all rockfishes (**Table 1**). Among non-aggregating species, bank (*S. rufus*, 5%) and swordspine rockfishes (*S. simulator*, 1.5%) were commonly observed. *Sebastes rufus* (n = 1,883) was the most abundant target species. In comparison, *S. paucispinis* (n = 232), *S. levis* (n = 62), and *S. crocotulus* (n = 16) were much less abundant (**Table X**).

Overall abundance and biomass estimates were calculated for 33 species of rockfish, three unidentified rockfish groups, *O. elongatus*, and *M. productus* (**Table 3**). Estimates of abundance and biomass of each of these species within each depth strata on each bank is provided in **Appendix 3**

Approximately 2.3 million rockfishes, *O. elongatus*, and *M. productus* were observed between 100-400 m on both banks (**Table 3**). *Sebastes ensifer* (619,114) was the most abundant species, which was slightly more abundant than *S. hopkinsi* (566,753). Among target species, *S. rufus* (177,981) were highly abundant. *Sebastes paucispinis* (12,624), *S. chlorostictus* (5,206), and *S. levis* (4,109) were much less abundant than those smaller, aggregating species (**Table 3**).

Total biomass of all species was approximately 287 metric tons (mt; 1 mt = 1,000 kg) in the same area (**Table 3**). Nearly 62% of the total fish biomass was comprised of the combination of *S. rufus* (60 mt) and small, aggregating species (*S. hopkinsi*, *S. ensifer*, *S. jordani*; 118 mt). The other target species, *S. paucispinis* (16.2 mt), *S. levis* (7.5 mt), and *S. chlorostictus* (2.5 mt) had relatively lower biomass.

The overall coefficient of variation (CV) for abundance (range = 0.12-1.00) and biomass estimates (range = 0.15-1.01) varied greatly among all species (**Table 3**). Species whose abundance and biomass estimates with high CVs (i.e., greater than ~0.50) were those that were either rarely encountered (e.g., *S. crameri*, *S. lentiginosus*, *S. rufinanus*, and *S. serranoides*) or whose densities varied greatly with depth (e.g., *S. hopkinsi*, *S. jordani*, *S. ovalis*, and *S. semicinctus*, which were densely aggregated in the shallower strata of the Footprint and almost entirely absent on Piggy Bank). Among target species, the CV values were relatively low and ranged from 0.28-0.46 and 0.26-0.44 for abundance and biomass, respectively. For *S. levis*, the overall CV values of abundance and biomass were quite low (0.28 and 0.30, respectively), and probably reflect the relatively even distribution of this species among transects on the Footprint and their total absence from any

transects on Piggy Bank (**Figure 2**). *Sebastes rufus* also had very low CV values (0.26 for both abundance and biomass), but in contrast to *S. levis*, was highly abundant in the deeper strata on both banks (**Fig. 2**). As expected, differences in the distribution of many species were apparent by depth and bank (**Fig. 2**). Small, more numerically abundant species (e.g., *S. ensifer*, *S. hopkinsi*, *S. semicinctus*, and *S. wilsoni*) were commonly encountered on the shallower (<200 m) portions of the Footprint. *Sebastes paucispinis* and *S. levis* were found almost exclusively on the Footprint. *Sebastes diploproa*, *S. rufus*, and *S. simulator* were found throughout the deeper strata on both banks, and were the only species commonly observed on Piggy Bank. n

SUB

A total of 25,085 individuals from 29 *Sebastes* species were observed, which as a group comprised ~80% of the total fishes encountered. Unidentified rockfishes (*Sebastes* spp.) comprised Only 0.5% of all rockfishes (excluding some young-of-year juveniles) to species or species group (i.e., Sebastomus). Dwarf rockfish species were most numerous and dominated by *S. hopkinsi* (20% of all rockfishes), *S. semicinctus* (17% of all rockfishes), *S. jordani* (12% of all rockfishes), *S. wilsoni* (pygmy 7%), and *S. ensifer* (7%). Of particular interest were the observations of 147 *S. paucispinis* (0.7%), 38 *S. levis* (0.2%), *S. miniatus* (0.1%), , *S. eos* (<0.1%), and 4*S. gilli* (<0.1%; a rare and elusive species). Among non-rockfish species, Pacific hake (mostly juveniles; 4%) were relatively abundant and lingcod were not (0.2%).

A total of 414,975 fishes and 60.3 mt of fish biomass on Piggy Bank (not including unidentified adult or juvenile rockfishes nor sharpchin rockfish (*S. zacentrus*) because meaningful length-weight relationships were not available) (**Table 5**). *Sebastes jordani* ($N = 163,022$), *S. rufus* ($N = 116,040$), and *S. diploproa* (64,489 individuals) were most abundant and comprised the greatest biomass (18%, 53%, and 12% of total biomass, respectively) on the Piggy Bank.

A total of 1,953,844 fishes representing 147.6 mt of fish biomass (excluding biomass estimates of unidentified rockfishes and sharpchin rockfish, as noted above) on The Footprint. Three dwarf species of rockfishes (*S. jordani*, $N = 339,855$); *S. semicinctus*, $N = 302,275$; and *Sebastomus* (likely *S. ensifer*), $N = 268,618$) were most abundant but together comprised only 27% of the total biomass. Among the large-bodied species [these are “large-bodied”?], *S. diploproa* ($N = 221,329$, $B = 14.5$ mt), juvenile *M. productus* ($N = 164,087$, $B = 19.5$ mt), and *S. rufus* ($N = 68,629$, $B = 14.9$ mt) were relatively abundant. *Sebastes levis* ($N = 4,325$, $B = 4.2$ mt) comprised only 0.2% of total abundance and 2.8% of total biomass on The Footprint. *Sebastes paucispinis* ($N = 13,342$, $B = 8.6$ mt) comprised 0.7% of total abundance and 5.8% of total biomass at this site.

Combining estimates from both banks, a total of 2,368,819 fishes with a biomass of 207.9 mt (**Table 7**). *Sebastes jordani*, *S. semicinctus*, and *S. diploproa* were most abundant overall. Juvenile *M. productus*, *S. rufus*, *S. ensifer*, and *S. hopkinsi* were also abundant. If *S. ensifer* and *Sebastomus* sp. (which were likely *S. ensifer*) were combined, the abundance would be second only to *S. jordani*. Four species (*S. rufus*, 23% of total biomass; *S. jordani*, 15%; *S. diploproa*, 10%; and juvenile *M. productus*, 10%) comprised over 58% of the total biomass in the entire study area.

The coefficient of variation (CV) ranged from 0.12 to 0.84 for estimates of total abundance and from 0.15 to 0.92 for estimates of total biomass over both banks combined (**Table 7**). The CV for total abundance was 0.30 for *S. levis*, 0.26 for *S. paucispinis*, and 0.32 for *S. rufus*; the CV for total biomass was 0.44 for *S. levis*, 0.27 for *S. paucispinis*, and 0.36 for *S. rufus*.

AUV

[WAITING FOR LIZ’S UPDATED RESULTS] A total of 21,228 fishes from 51 taxa were identified during the survey including 18 rockfish species (**Table X**). Rockfishes comprised 75% of the fishes encountered, including groups of unidentified rockfish (*Sebastes* spp., **X%**), rosy-group rockfishes (*Sebastomus*, **X%**), and thornyheads (*Sebastolobus* sp., **X%**), and other species that could be identified to species (**X%**). Notable were observations of 11 cowcod (*S. levis*), 29 bocaccio (*S. paucispinis*) and one bronzespotted rockfish (*S. gilli*). In addition, 53 lingcod (*Ophiodon elongatus*), 19 Pacific hake (*Merluccius productus*) and 10 sablefish

(*Anoplopoma fimbria*) were observed. Fish assemblages varied between depth strata and between the two banks. Twenty-nine taxa only occurred on the Footprint or along its Flank. Two species were observed on Piggy Bank only.

Biodiversity

ROV

Biodiversity also varied greatly by depth strata and bank (**Table 6**). The greatest number of species (richness = 15) was observed in the 0-100 m depth stratum on the Footprint, and the fewest were observed in the 300-400 m stratum at Piggy Bank. The rarefied species richness (the number of species expected per number of samples) was greatest in the 300-400 m stratum at the Footprint, with all other strata having rarefied species richness between 3.5 and 5.1. Neither of these richness parameters account for the abundance of different species so Shannon H' and Simpson λ , which do account for abundance differences, were also calculated. Again, the greatest diversity (both H' and λ , respectively) were observed in the 300-400 m stratum at the Footprint, but diversity was relatively uniform across all depth strata and banks. Biodiversity estimates for each transect is presented in **Appendix 4**.

The species accumulation (rarefaction) curves from each bank have much different shapes (**Figure 9**). The curve for the Footprint rises steeply in the first few samples and begins to plateau after ~ **10** transects. The rarefaction curve at Piggy Bank increased at a slower rate and never reached an asymptote. Although there were fewer transects at Piggy Bank, the other diversity estimates corroborate the finding that there is less biodiversity at Piggy Bank compared to the Footprint.

SUB

Diversity (**or richness??**; measured as cumulative number of species in transects within a depth stratum, excluding groups), was highest on The Footprint in the 100-200 m (23 species) and 200-300 m (26 species) strata (**Table X**). Species richness (S) was lowest on Piggy Bank at both depth strata (8 and 11 species in the 200-300 and 300-400 m strata, respectively).

AUV

[WAITING FOR LIZ'S UPDATED RESULTS] Species richness (total species, excluding groups) was the highest in the 100-200 m stratum on the Footprint ($N = 23$ species) and in the 300-400 m stratum on the Flank (18 species, **Table x**). Species richness was lowest in all three strata on Piggy Bank (six species at 200-300 m, eight species at 300-400 m, and seven species at 400-500 m).

Discussion

Survey effort

The AUV was able to sample the deeper 400-500 m stratum on the flank of Footprint, and to provide fish and habitat information that was not available from the ROV or SUB (**The ROV could have sampled deeper but .

Abundance and biomass estimates

AUV

The species encountered by the AUV, ROV, and SUB were comparable.

Species diversity was similar amongst all three methodologies showing greater species richness at the Footprint bank and lower richness at Piggy Bank. The ROV and the SUB included records of rockfish species not identified by the AUV. This could be due to the fact that the cameras on the AUV are positioned downward, rather than forward facing, allowing fish directly in front of the unit to escape detection. Another possible reason could be the difficulty in identifying some rockfish species from the dorsal view. The AUV detected comparable numbers of fishes, but the fish were identified to groupings (i.e. Unidentified rockfish, or Sebastomus) as opposed to species.

Tables

Table 1. Length-weight table [UPDATE]

Scientific name	Common name	a	b	Sex	Reference	Comment
<i>Merluccius productus</i>	Pacific hake	0.0347	2.5560	male	(Dark, 1975)	
<i>Ophiodon elongatus</i>	Lingcod	0.0113	2.9900	both	(RecFIN, 2009)	
<i>Sebastes aurora</i>	Aurora rockfish	0.0244	2.8320	both	(Wilkins et al., 1998)	
<i>Sebastes chlorostictus</i>	Greenspotted rockfish	0.0091	3.1632	both	(Love et al., 1990)	
<i>Sebastes constellatus</i>	Starry rockfish	0.0097	3.1598	both	(Love et al., 1990)	
<i>Sebastes crameri</i>	Darkbloched rockfish	0.0295	2.8240	both	(Wilkins et al., 1998)	
<i>Sebastes crocotulus</i>	Sunset rockfish	0.0216	2.9234	both	(Love et al., 1990)	borrowed from <i>S. miniatus</i>
<i>Sebastes diploproa</i>	Splitnose rockfish	0.0041	3.2440	both	(PSFMC, 1999)	
<i>Sebastes elongatus</i>	Greenstripe rockfish	0.0079	3.1275	both	(Love et al., 1990)	
<i>Sebastes ensifer</i>	Swordspine rockfish	0.0132	2.9702	both	(Love et al., 1990)	
<i>Sebastes entomelas</i>	Widow rockfish	0.0164	2.9426	both	(Love et al., 1990)	
<i>Sebastes eos</i>	Pink rockfish	0.0186	2.9573	both	(PSFMC, 1999)	
<i>Sebastes gilli</i>	Bronzespotted rockfish	0.0177	2.9807	both	(PSFMC, 1999)	
<i>Sebastes hopkinsi</i>	Squarespot rockfish	0.0146	2.9840	both	(Love et al., 1990)	
<i>Sebastes jordani</i>	Shortbelly rockfish	0.0056	3.1600	both	(PSFMC, 1999)	
<i>Sebastes lentiginosus</i>	Freckled rockfish	0.0067	3.3190	both	(Chen, 1971)	borrowed from <i>S. umbrosus</i>
<i>Sebastes levis</i>	Cowcod	0.0101	3.0933	both	(Love et al., 1990)	
<i>Sebastes macdonaldi</i>	Mexican rockfish	0.0445	2.6640	both	(PSFMC, 1999)	
<i>Sebastes melanostomus</i>	Blackgill rockfish	0.0123	3.0420	both	(Love et al., 1990)	
<i>Sebastes ovalis</i>	Speckled rockfish	0.0052	3.2170	female	(Love et al., 1990)	
<i>Sebastes paucispinis</i>	Bocaccio	0.0162	2.8810	female	(Love et al., 1990)	
<i>Sebastes phillipsi</i>	Chameleon rockfish	0.0244	2.8320	unsexed	(Wilkins et al., 1998)	borrowed from <i>S. aurora</i>
<i>Sebastes rosaceus</i>	Rosy rockfish	0.0052	3.3857	both	(Love et al., 1990)	
<i>Sebastes rosenblatti</i>	Greenblotched rockfish	0.0110	3.1057	both	(Love et al., 1990)	
<i>Sebastes ruberrimus</i>	Yelloweye rockfish	0.0074	3.2220	both	(Rosenthal et al., 1982)	
<i>Sebastes rubrivinctus</i>	Flag rockfish	0.0206	2.9431		(RecFIN, 2009)	
<i>Sebastes rufinanus</i>	Dwarf-red rockfish	0.0146	2.9840	both	(Love et al., 1990)	borrowed from <i>S. hopkinsi</i>
<i>Sebastes rufus</i>	Bank rockfish	0.0078	3.1469	both	(Love et al., 1990)	
<i>Sebastes saxicola</i>	Stripetail rockfish	0.0093	3.1201	both	(PSFMC, 1999)	
<i>Sebastes semicinctus</i>	Halfbanded rockfish	0.0127	3.0160	female	(Love et al., 1990)	
<i>Sebastes serranoides</i>	Olive rockfish	0.0111	3.0630	female	(Love and Westphal, 1981)	
<i>Sebastes simulator</i>	Pinkrose rockfish	0.0056	3.2785	both	(Love, unpublished data)	
<i>Sebastes sp.</i>	Rockfish-unidentified	0.0146	2.9840	both	(Love et al., 1990)	borrowed from <i>S. hopkinsi</i>
<i>Sebastes wilsoni</i>	Pygmy rockfish	0.0119	3.0230		(Moulton, 1977)	borrowed from <i>S. emphaeus</i>
<i>Sebastes zacentrus</i>	Sharpchin rockfish	0.0060	3.2800	both	(Wilkins et al., 1998)	
<i>Sebastolobus alascanus</i>	Shortspine thornyhead	0.0039	3.3570	both	(Wakefield, 1990)	
<i>Sebastomus sp.</i>	Rosy-group rockfish	0.0132	2.9700	both	(Love et al., 1990)	borrowed from <i>S. ensifer</i>

Figures

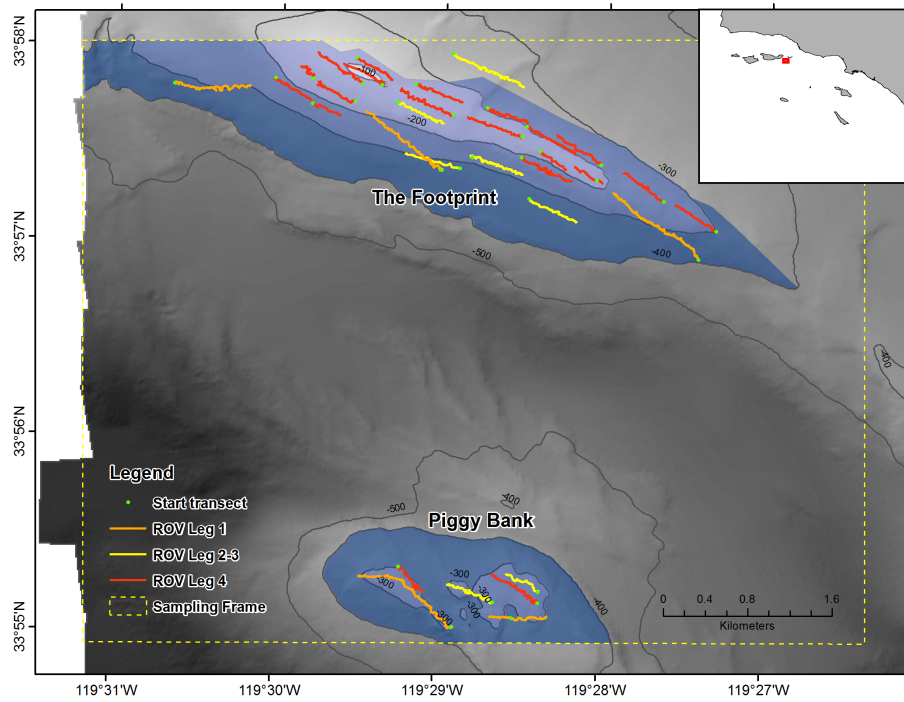


Figure 1. Map of the survey area indicating the extent of the sampling area, which includes The Footprint and Piggy Bank (dashed yellow box), depth strata (blue polygons), and the 250 x 250 m grid used to randomly select AUV sample locations. The area within each polygon is in **Table X**. [Replace with new figure without ROV transects, showing AUV 250x250 m grid and randomly selected cells].

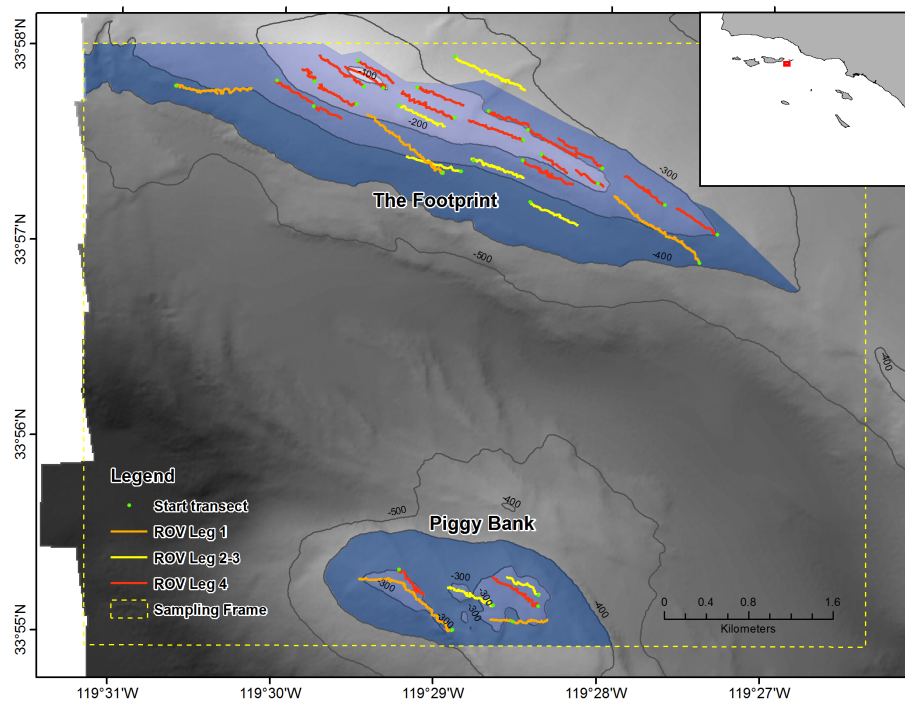


Figure 2. Locations of ROV, SUB, and AUV transects at the Footprint and Piggy Bank.

Literature cited