Groundfish multivariate analyses - Compare Years by Area

Colette Ward May 5, 2016

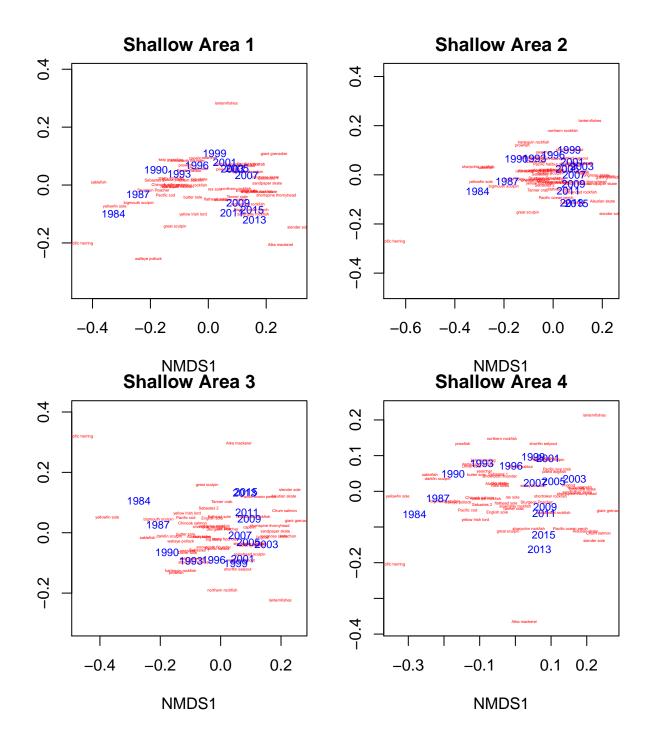
1. nMDS

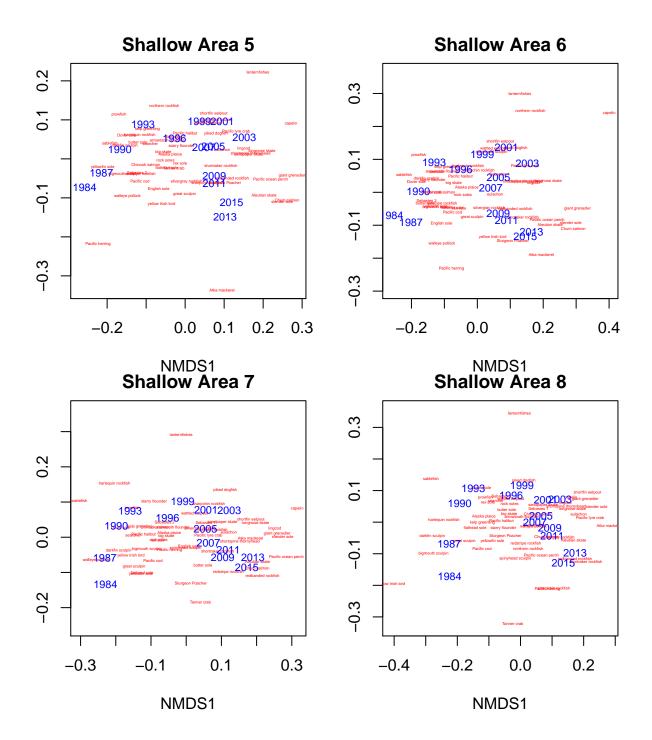
Transformation is Wisconsin-style double transformation (normalizes taxa to % abundance, then normalizes abundances to the maximum for each species) of sqrt-transformed data.

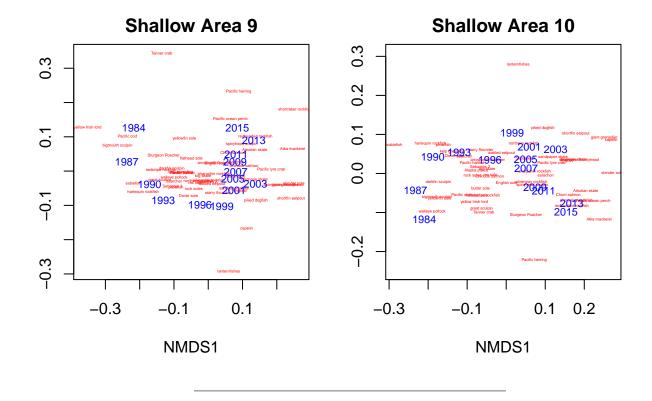
Sorry the text is so small. It's necessary for legibility because there are so many species. Enlarge the output pdf to see species names.

Shallow areas

Area 10 is Total

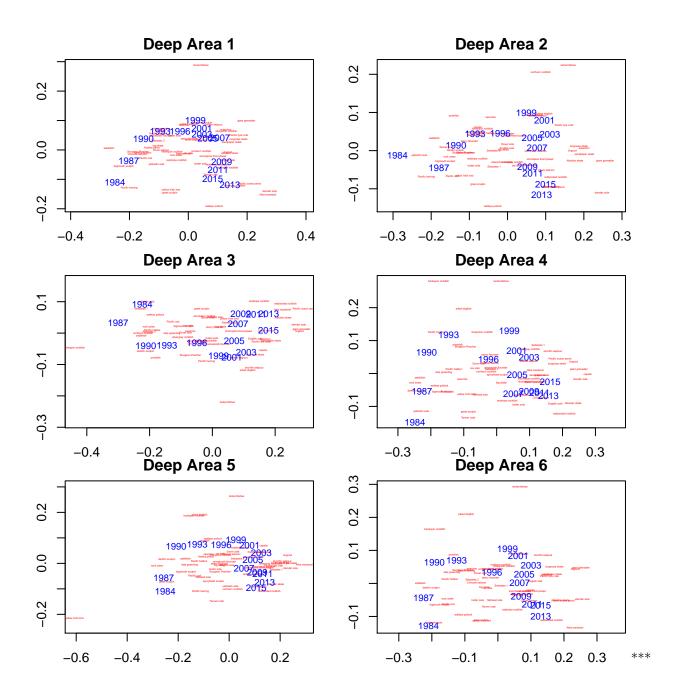






Deep areas

Area 6 is Total



2. PCA

Methods: Used vegan package in R, function rda(). Raw CPUEs vary by many orders of magnitude $(10^-7 \text{ to } 10^2 \text{ in shallow areas}, 10^-5 \text{ to } 10^2 \text{ in deep areas})$, therefore, to prevent the most abundant species from driving the result and allow all species to contribute equally to the result, cpues were standardized prior to analysis (Legendre & Legendre 1998) - means that PCs were extracted from correlation matrices, not dispersion matrices.

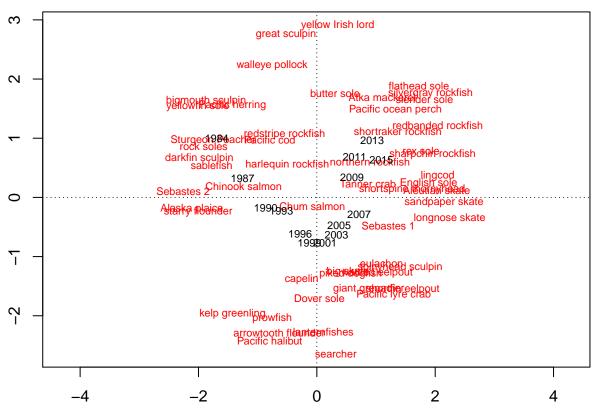
Shallow areas

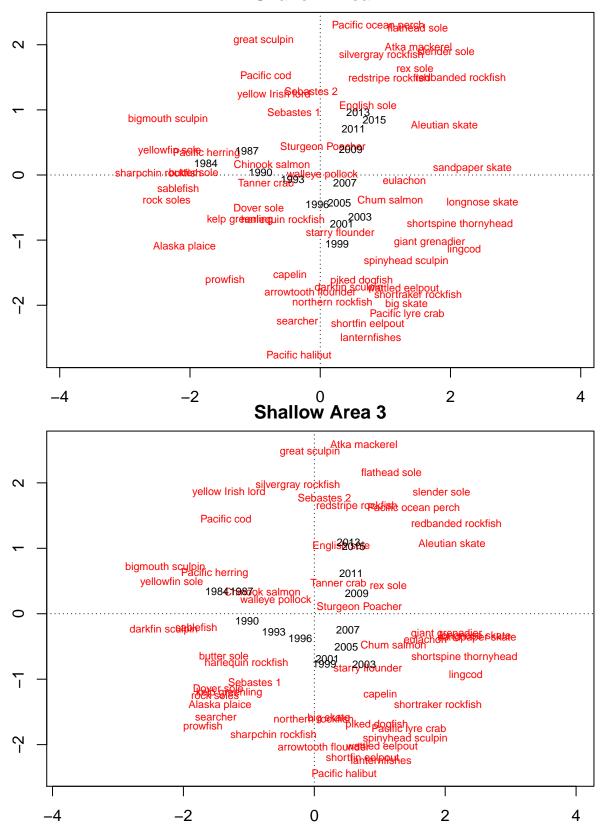
Area 10 is Total

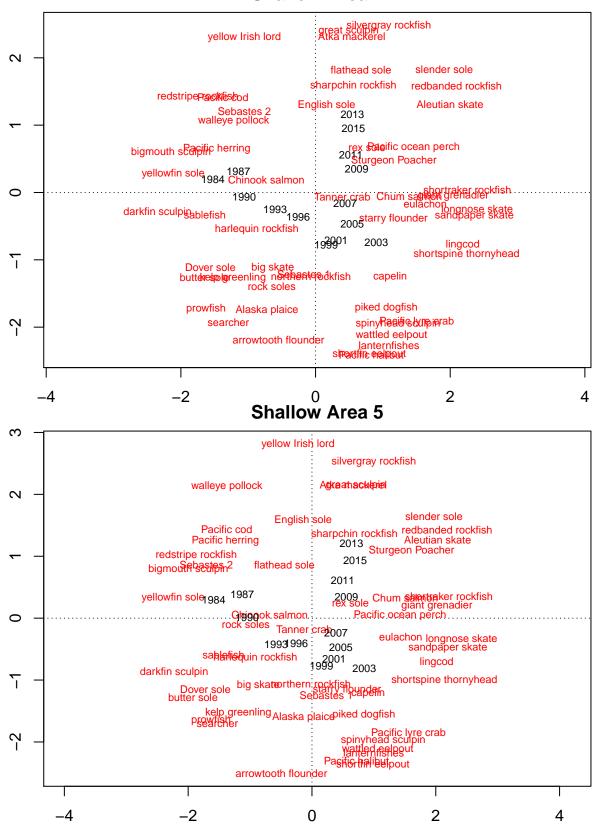
% of total variance explained by PC1 & PC2 (Area 10 = Total):

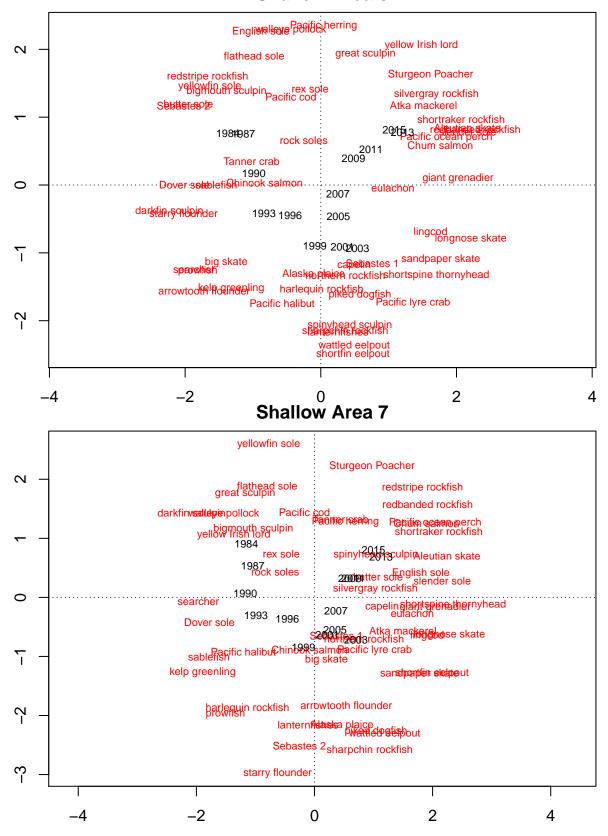
##		Area	PC1	PC2
##	1	1	0.3519725	0.2048984
##	2	2	0.2915416	0.1887890
##	3	3	0.2834110	0.1964664
##	4	4	0.2998676	0.1878989
##	5	5	0.3035660	0.2050751
##	6	6	0.3427620	0.2280674
##	7	7	0.3380583	0.1769048
##	8	8	0.3498791	0.2463860
##	9	9	0.3640909	0.1794389
##	10	10	0.3373172	0.2248949

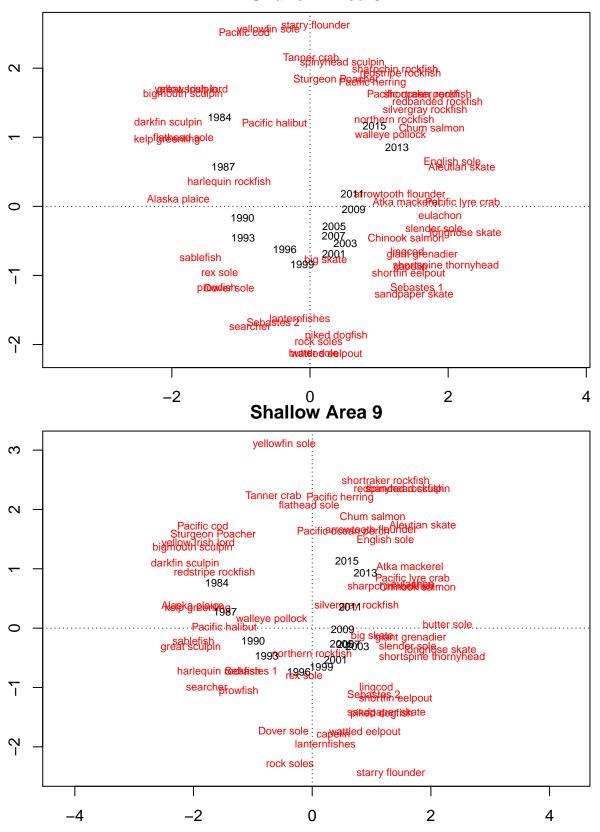
Plots for shallow areas. Note that I've removed arrows for species scores and did some wonky scaling of species scores to make them legible in the plots (divided each species score by species StDev - although their relative positions are preserved in the plots, this is technically incorrect because I've already standardized data prior to analysis). This will need to be changed if used in a publication - see .Rmd file for correct code.

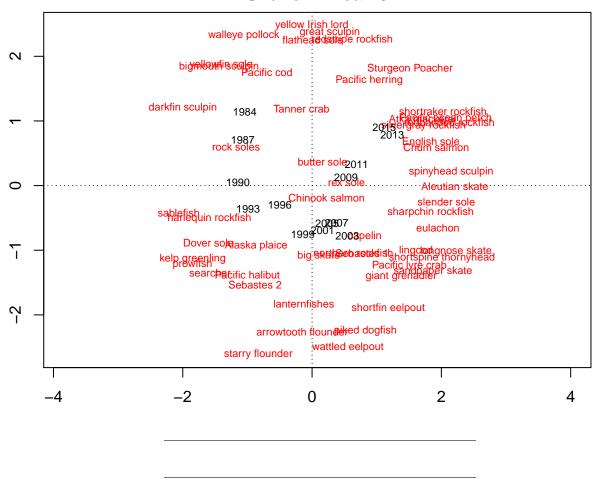












Deep areas

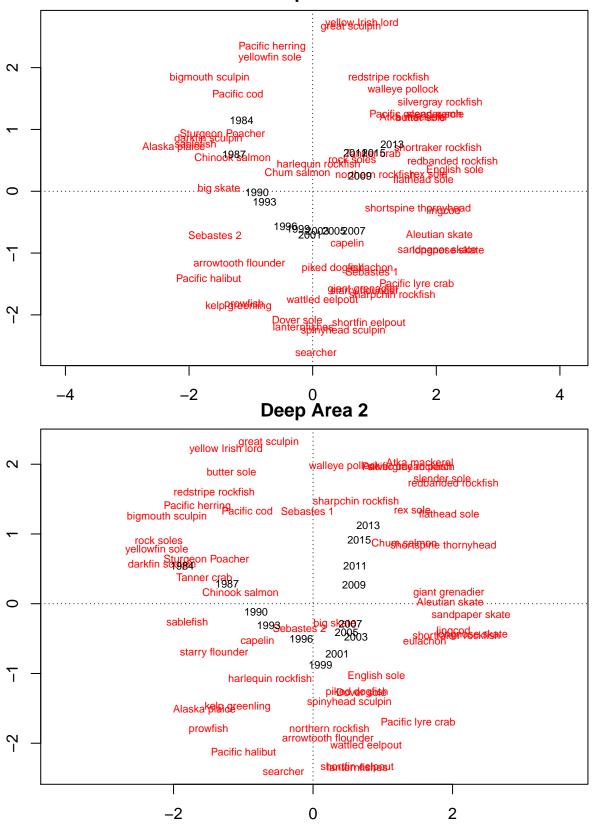
Area 6 is Total

% of total variance explained by PC1 & PC2 (Area 6 = Total):

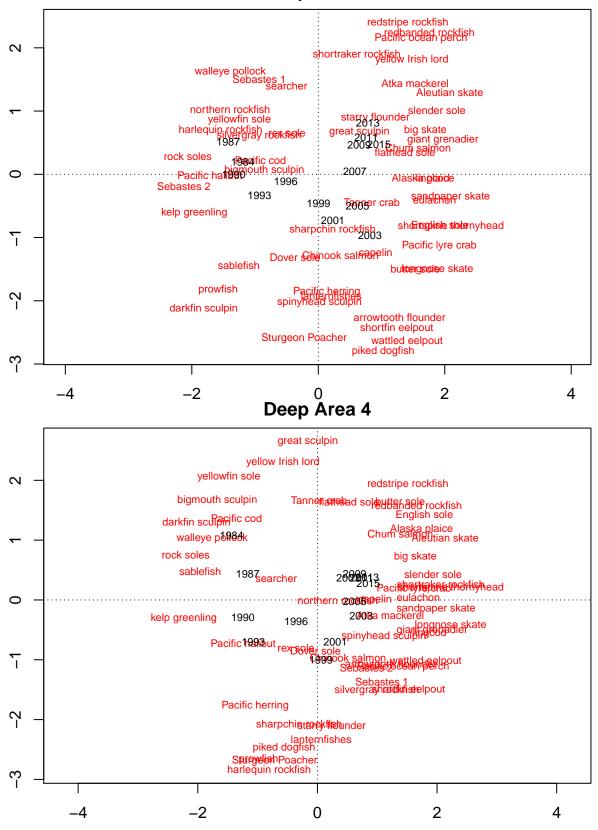
```
## Area PC1 PC2
## 1 1 0.3378644 0.2105648
## 2 2 0.3405090 0.1898041
## 3 3 0.3935400 0.1444878
## 4 4 0.3759332 0.1649310
## 5 5 0.3554989 0.1661971
## 6 6 0.3765467 0.1725329
```

Plots for deep areas. See note above about scaling of species scores.

Deep Area 1



Deep Area 3



Deep Area 5

