SEACAR SAV Analysis

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Important Notes

The purpose of this script is to provide a report summary of SAV analysis. The script used for analysis is SEACAR SAV BB script website.R.

All scripts and outputs can be found on the SEACAR GitHub repository:

 $https://github.com/FloridaSEACAR/SEACAR_Trend_Analyses$

This script is based off of code originally written by Stephen Durham with comments by Marcus W. Beck.

Summary of SEACAR_SAV_BB_script_website.R

- Objective: Import and format SAV data, create summary plots and maps of changes over time, model changes over time using Bayesian and mixed-effects models. Results are separate for each managed area, species, and parameter (e.g., Braun Blanquet, percent cover, etc.).
- Packages: bayesplot, brms, broom.mixed, data.table, grid, gridExtra, gtable, nlme, scales, sf, tictoc, tidybayes, tidyverse
- File inputs: Combined_SAV_column_All-2021-Sep-20.csv, seacar_dbo_SampleLocation_Point.shp, seacar_dbo_SampleLocation_Line.shp, ORCP_Managed_Areas.shp, Counties_-_Detailed_Shoreline.shp, MApolygons_corners.csv
- Steps by line number:
 - 1 30: load libraries and import SAV file
 - 31 143: format SAV data including renaming columns, removing NA values, and reformatting abundance/cover values. For the latter, this included removal of NA values and those out of range, and ensuring appropriate values for Braun Blanquet, modified Braun Blaunqet, percent occurrence, and percent cover.
 - 146 316: function for plotting model predictions
 - 327 426: if loop for rotating a spatial object, used for plotting to create a presentation of stacked
 2-d layers.
 - 399 426: import spatial data objects, project all to WGS 1984 datum. Spatial data objects include sample locations and relevant boundaries (counties, management area boundaries)
 - 428 663: setup parameter list and objects for looping through parameters to create models and summary output, parameters include Braun Blanquet, median percent cover, visual percent cover, percent occurrence, frequency of occurrence

- 670 1848: loop through parameters to create models and summary output, the following is an outline of steps in this loop.
 - * 683 1848: loop through managed areas using parameter from outside loop
 - * 703 717: create and save plot of parameter score for managed area over time by species
 - * 719 732: create and save plot of parameter score for managed area over time by program ID
 - * 734 747: create and save plot of parameter score for managed area over time grouped by species, by program ID
 - * 749 762: create and save plot of quadrat sizes for managed area over time by species
 - * 764 777: create and save plot of quadrat sizes for managed area over time by program ID
 - * 779 792: create and save plot of method for managed area over time by species
 - * 794 807: create and save plot of method for managed area over time by program ID
 - * 809 823: create and save plot of method for managed area by quadrat size and species
 - * 825 839: create and save plot of method for managed area by quadrat size and program ID
 - * 842 871: create and save plots of grid values over time by species and program ID if data available
 - * 873 903: create and save plots of depth values over time by species and program ID if data available
 - * 907 923: create and save a plot legend of species
 - * 925 949: loop through species to create and save a plot of parameter score over time
 - * 953 972: create and save a plot of totals for the species of parameter score over time
 - * 974 991: create and save a plot legend of species
 - * 993 1017: loop through species to create and save a plot of parameter score as boxplots over time
 - * 1019 1040: create and save a plot of boxplots for the species of parameter score over time
 - * 1042 1368: create and save stacked maps of parameter values for the managed area by year
 - * 1370 1401: setup empty objects for model results
 - * 1405 1533: Loop through species to fit models, with separate exception statements for different parameters. The modeling workflow is similar for each parameter, with minor exceptions. The general goal of each is to assess trends in a parameter over time for a particular species and managed area. Each workflow includes error handling if models did not converge, produces summary tables of model fit, and summary plots showing model result. The models vary in the Gaussian distribution family for the response variable depending on parameter. Random effects (e.g., for LocationID) are used for all models.
 - * 1539 1592: reads and sets modeled and umodeled data for plots
 - * 1595 1761: create base plot of seagrass percent cover data over time for managed area. Add model fits if applicable, and saves objects
 - * 1781 1841: create and save barplots of parameter results for managed area over time by species, only for Braun Blanquet and percent cover, save model results from prior loops
 - $\ast\,$ 1843 1946: print statements for the console
- 1876 1880: get models, plots, and files. save list of failed models and R session info
- 1890 1943: Save trendplot figures as png files
- 1948 1961: Save barplot figures as png files
- − 1964 1994: Crop geographic scope figure images & add metadata stamp
- File outputs: Multiple binary RDS files.