

# nwfscSurvey Vignette

2025-01-23

Set up: The first set of code is for the Bottom trawl survey The second set of code is for the triennial survey

Questions: is this a two area model as in WA/OR + CA? or WA + OR only? Is California a whole different assessment model? If we are seeing catches in CA, what is the strata doing again?

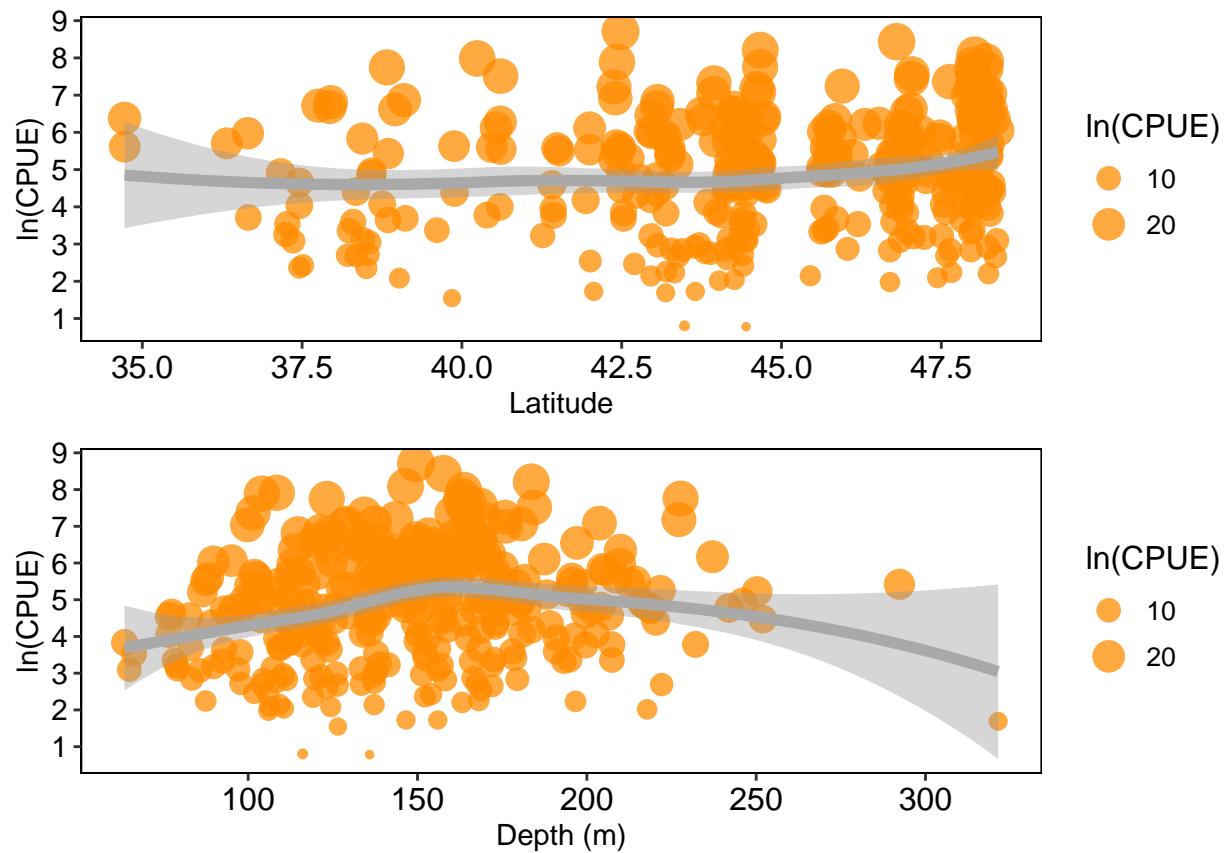
## NWFSC.Combo Data

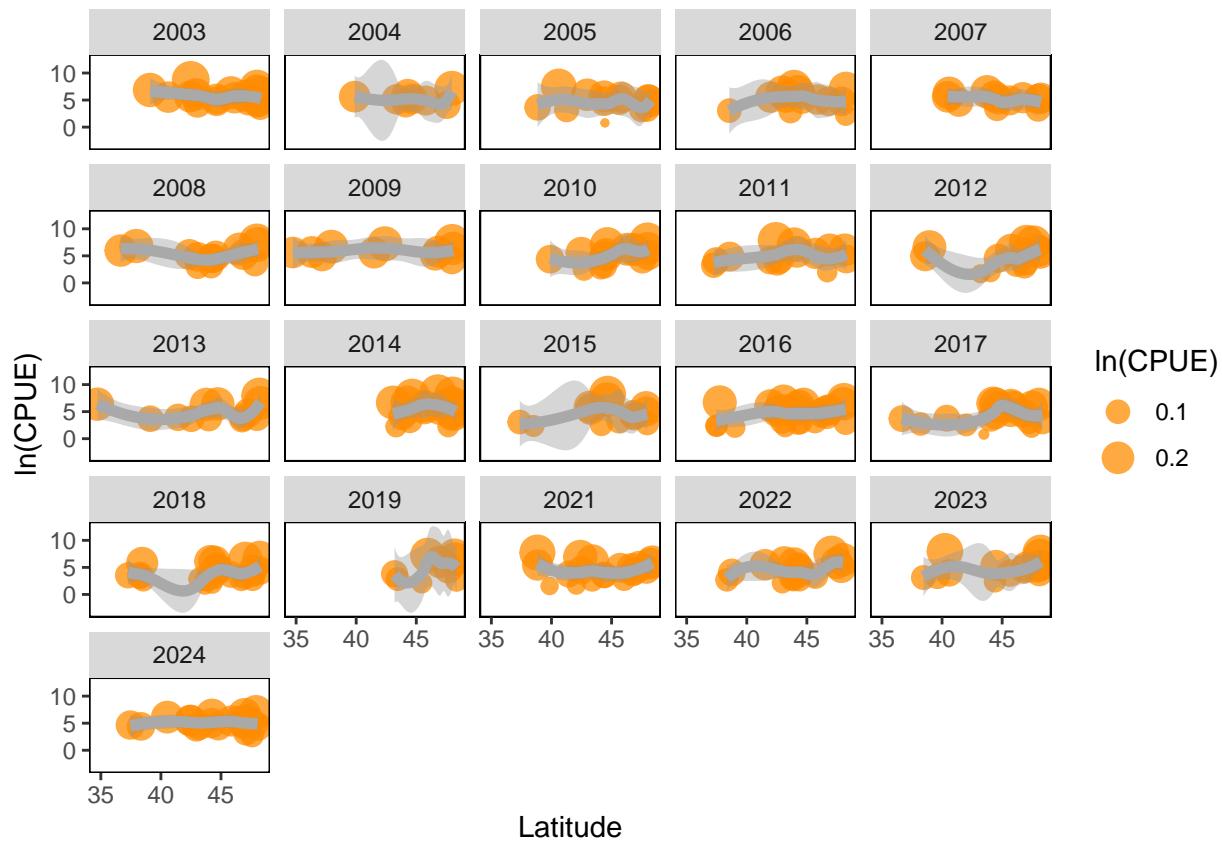
Initial pull and plotting of data: WCGBTS

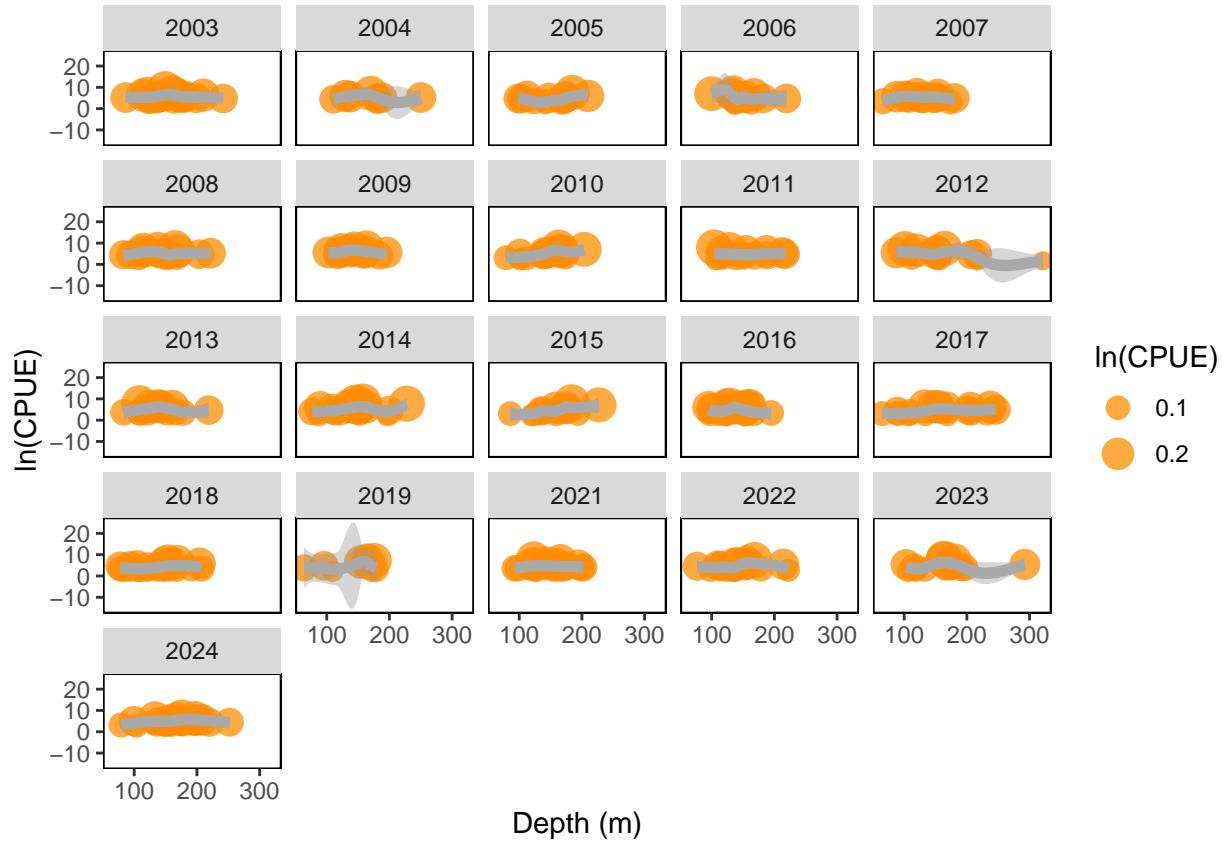
```
catch = pull_catch(
    common_name = "yelloweye rockfish",
    survey = "NWFSC.Combo")

bio = pull_bio(
    common_name = "yelloweye rockfish",
    survey = "NWFSC.Combo")

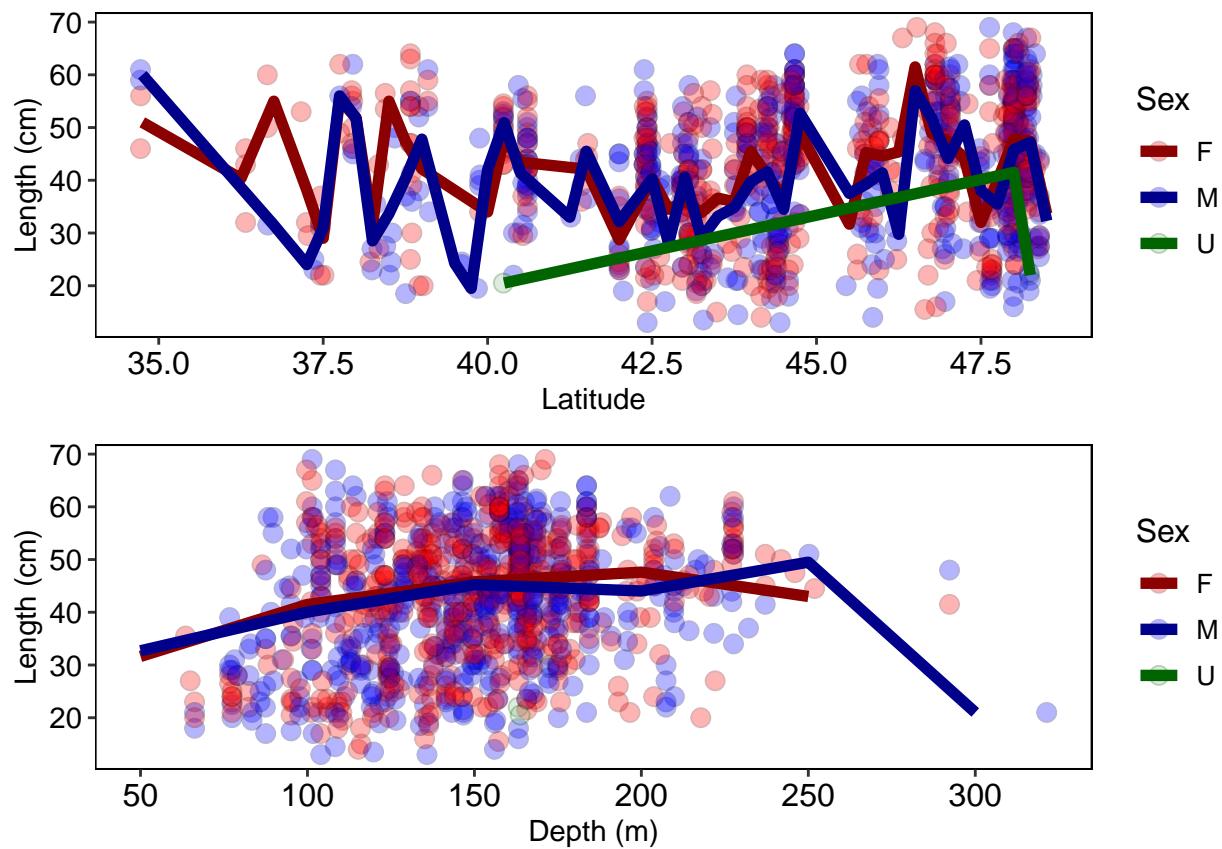
plot_cpue(
    catch = catch)
```

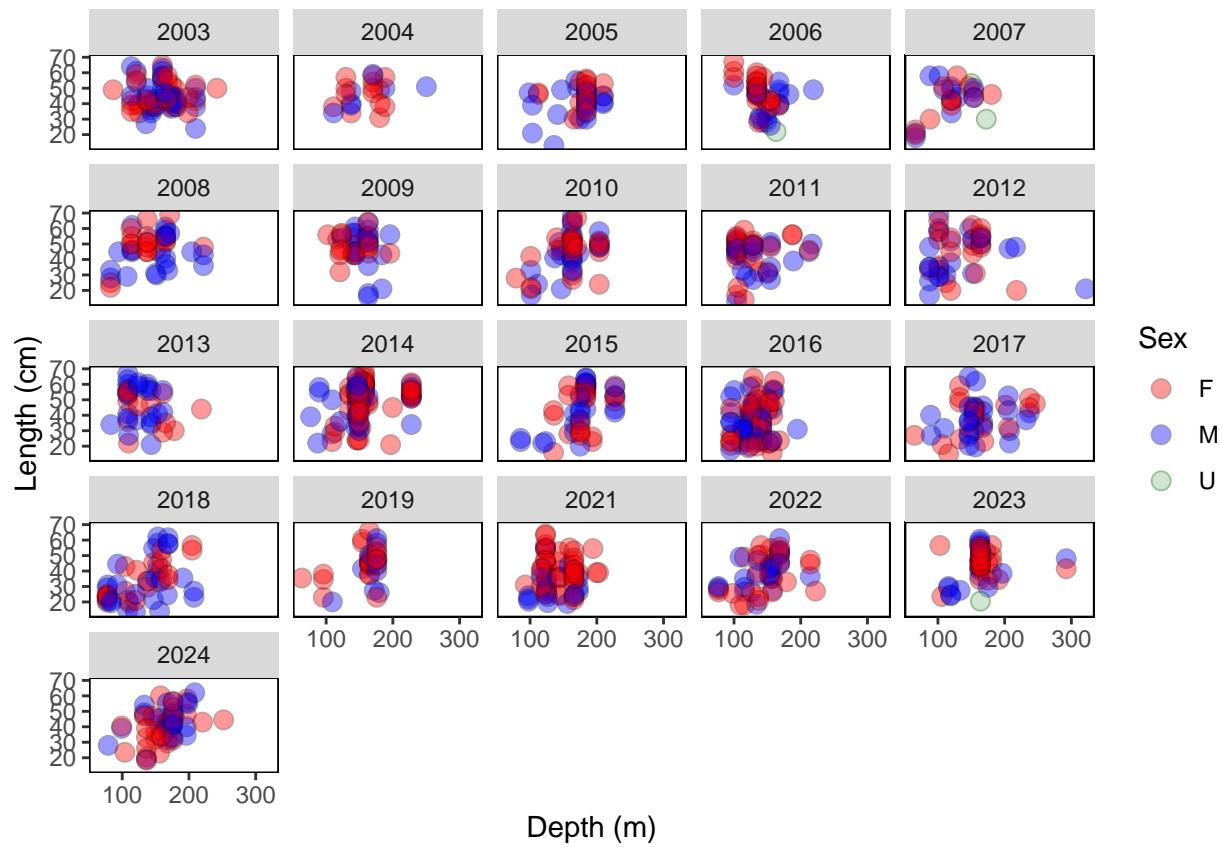


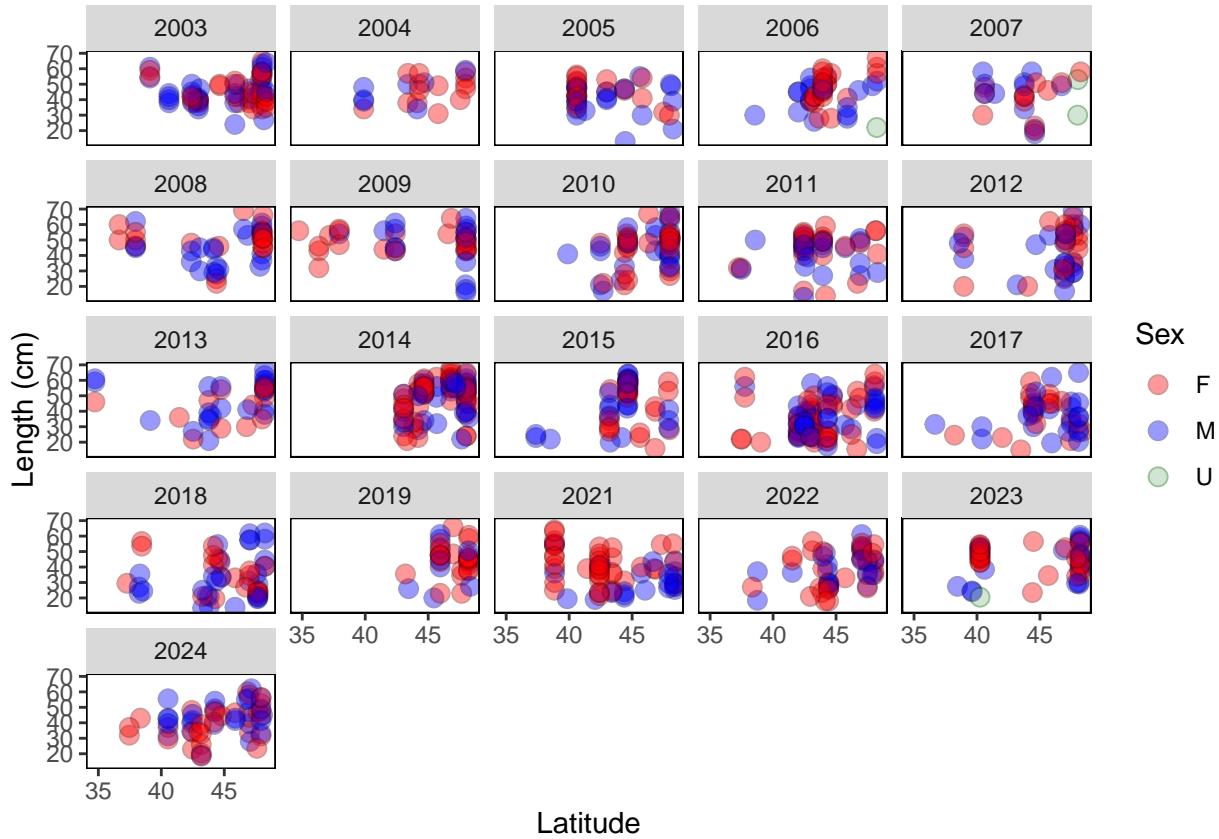




```
plot_bio_patterns(
  bio = bio,
  col_name = "Length_cm")
```







```
wh_plot_proportion(
  data_catch = catch,
  data_bio = bio
)

## [1] "C:/Users/elizabeth.gugliotti/Documents/github_repos/Sebastes_ruberimus_2025/Rcode/plots/presen"
## [2] "C:/Users/elizabeth.gugliotti/Documents/github_repos/Sebastes_ruberimus_2025/Rcode/plots/presen"
## [3] "C:/Users/elizabeth.gugliotti/Documents/github_repos/Sebastes_ruberimus_2025/Rcode/plots/sex_by"
## [4] "C:/Users/elizabeth.gugliotti/Documents/github_repos/Sebastes_ruberimus_2025/Rcode/plots/sex_by"
```

Define the strata

```
WCGBTS_strata <- CreateStrataDF.fn(
  names = c("shallow_OR", "deep_OR", "shallow_WA", "deep_WA"),
  depths.shallow = c(55, 183, 55, 183),
  depths.deep    = c(183, 350, 183, 350),
  lats.south     = c(42, 42, 46, 46),
  lats.north     = c(46, 46, 49, 49)
)
```

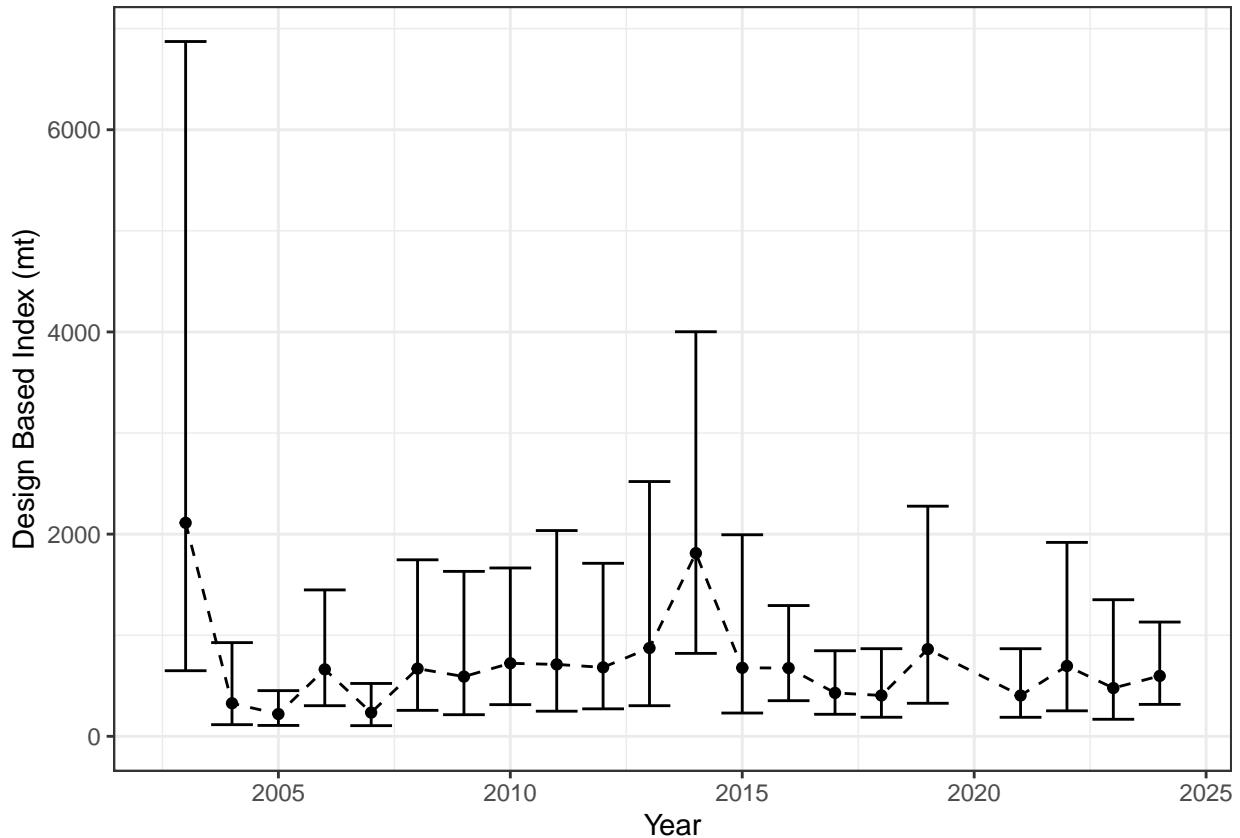
Calculate designed based index of abundance

```
biomass = get_design_based(
  data = catch,
  strata = WCGBTS_strata,
  dir = file.path(dirname(getwd()), "Data", "NWFSC.Combo_and_Tri_design_based_indices"),
  printfolder = "")
```

```
## [1] TRUE
```

Plot coastwide

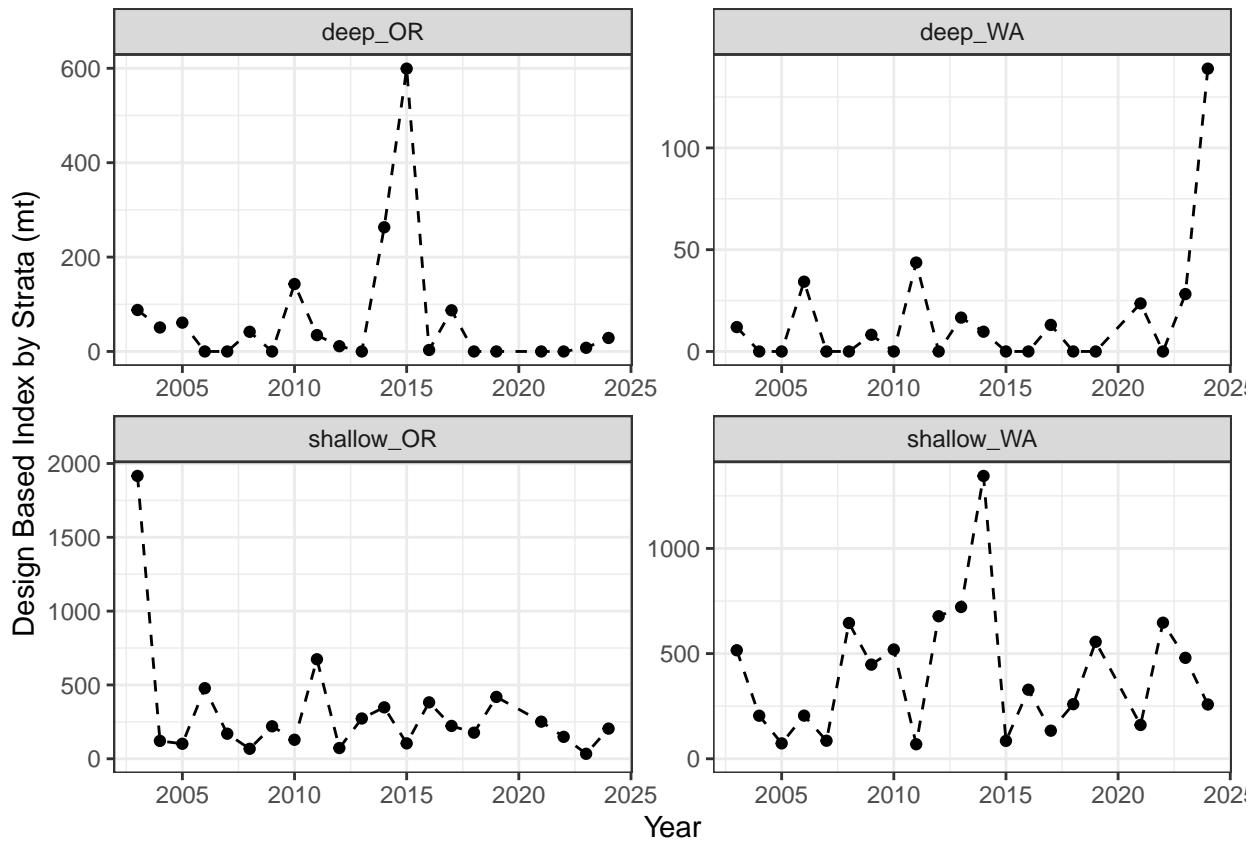
```
plot_index(  
  data = biomass,  
  plot = 1)
```



# that's only about a maximum of 200,500 fish.... very few...

Plot index for each strata

```
plot_index(  
  data = biomass,  
  plot = 2)
```



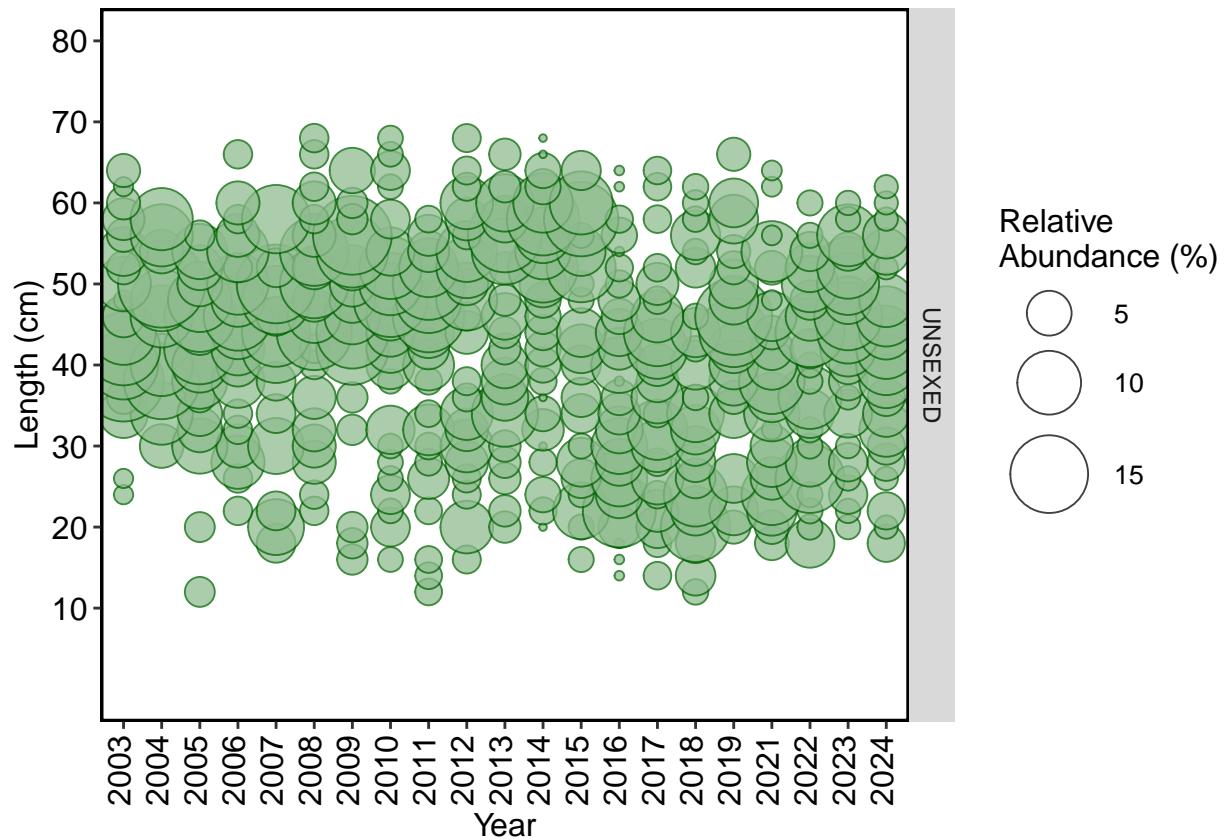
Length composition data

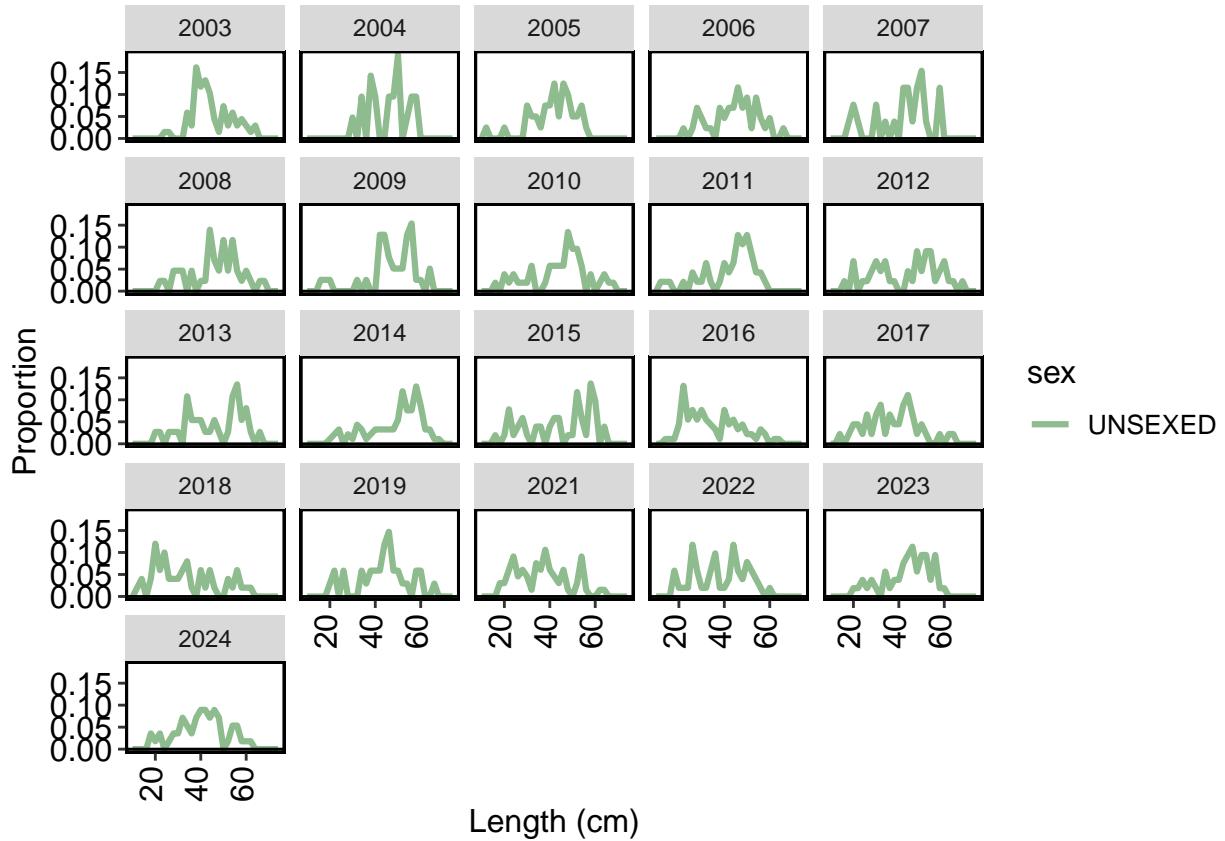
```
# Expanded length comps are not used for the yelloweye assessment
# length_comps <- get_expanded_comps(
#   bio_data = bio,
#   catch_data = catch,
#   comp_bins = seq(10, 74, 2),
#   strata = WCBTS_strata,
#   comp_column_name = "length_cm",
#   output = "full_expansion_ss3_format",
#   two_sex_comps = FALSE, #single sex model
#   input_n_method = "stewart_hamel")
#
# plot_comps(
#   data = length_comps)

# Use raw length comps, you can tell this is what was used in the previous assessment because whole num
raw_length_comps <- get_raw_comps(
  data = bio,
  comp_bins = seq(10, 74, 2),
  comp_column_name = "length_cm",
  dir = getwd(),
  printfolder = "",
  two_sex_comps = FALSE) #single sex model

plot_comps(
  data = raw_length_comps
```

)





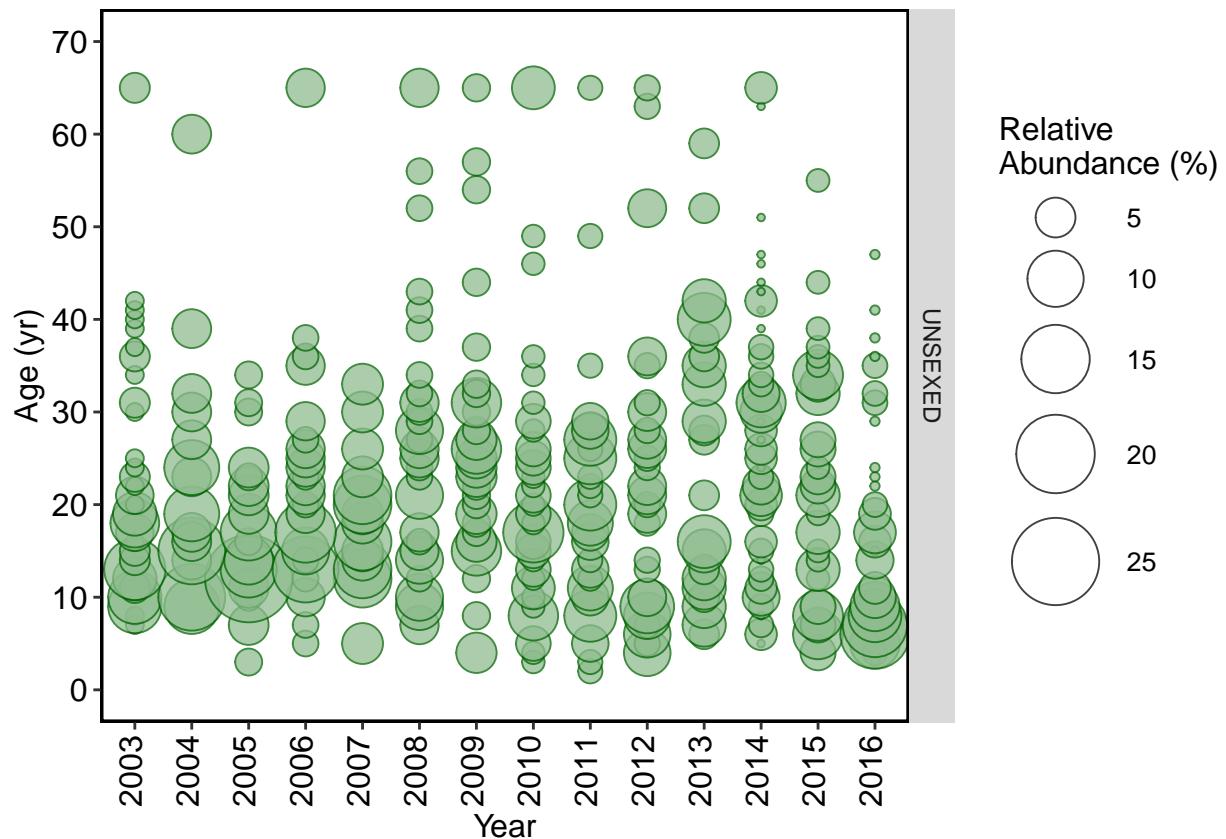
```
## [1] TRUE
## [1] TRUE
## [1] TRUE
## [1] TRUE
```

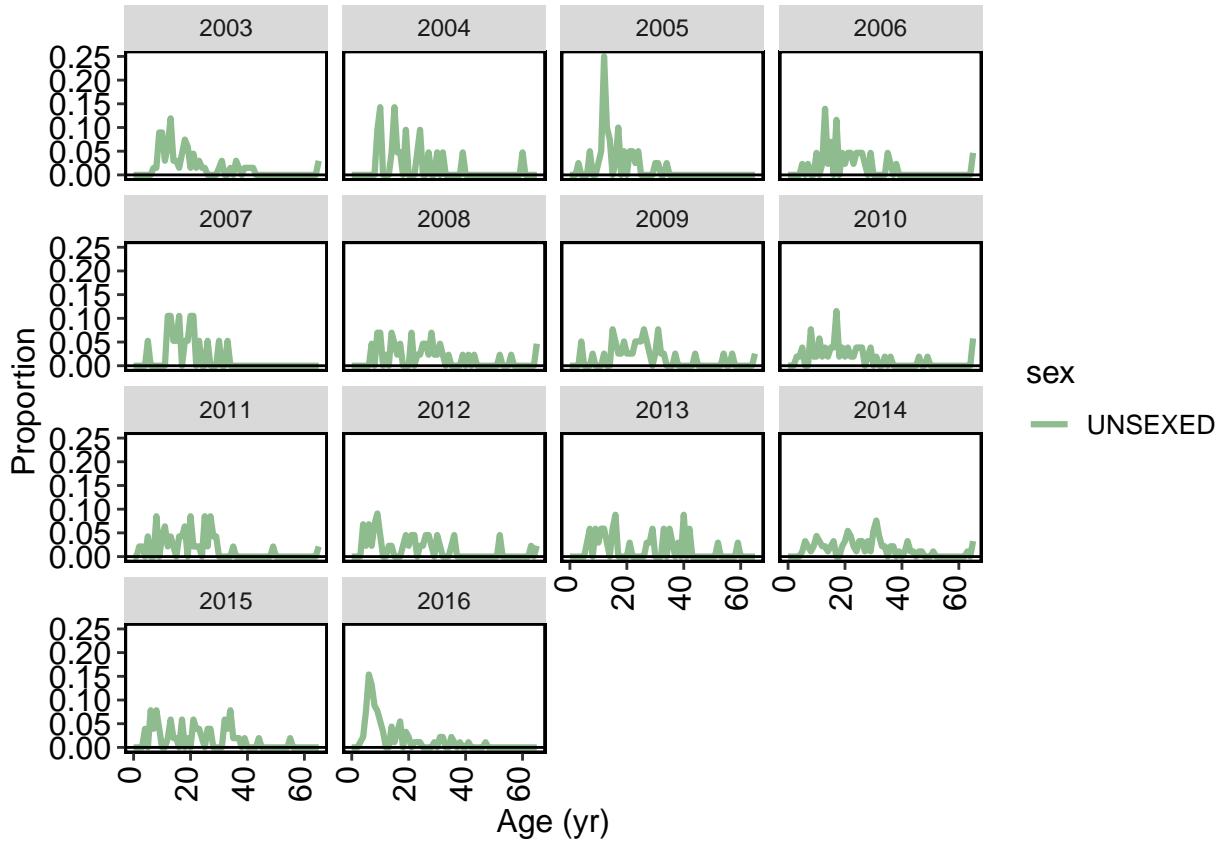
Marginal age composition data

```
# Expanded age comps are not used in the yelloweye assessment
# age_comps <- get_expanded_comps(
#   bio_data = bio,
#   catch_data = catch,
#   comp_bins = 0:65,
#   strata = WCGBTS_strata,
#   comp_column_name = "age",
#   output = "full_expansion_ss3_format",
#   two_sex_comps = FALSE, #single sex model
#   input_n_method = "stewart_hamel")
#
# plot_comps(
#   data = age_comps)

raw_age_comps <- get_raw_comps(
  data = bio,
  comp_bins = 0:65,
  comp_column_name = "age",
  dir = getwd(),
```

```
printfolder = "",  
two_sex_comps = FALSE) #single sex model  
  
plot_comps(  
  data = raw_age_comps)
```





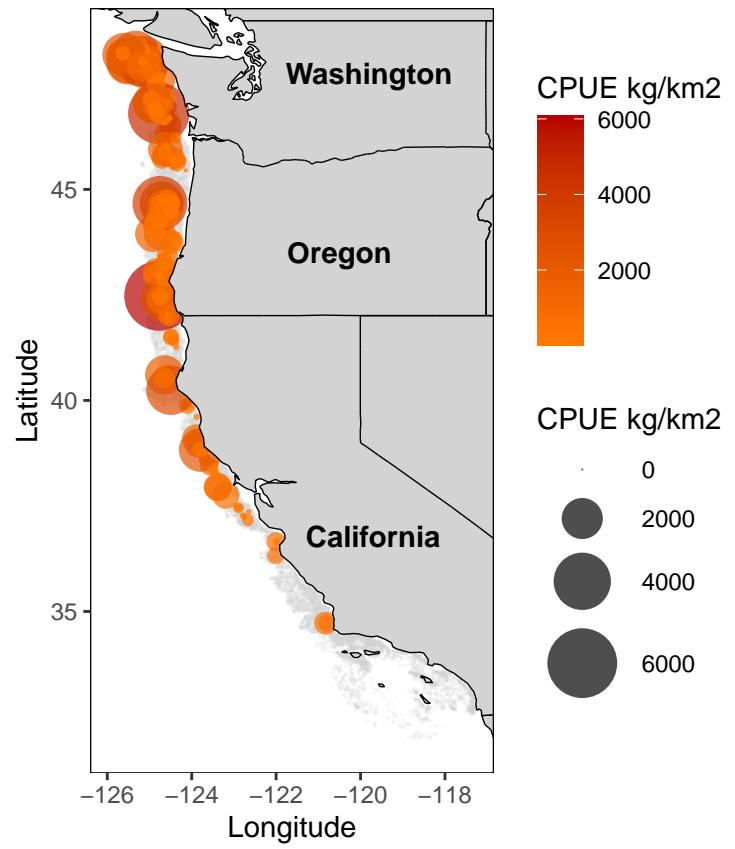
```

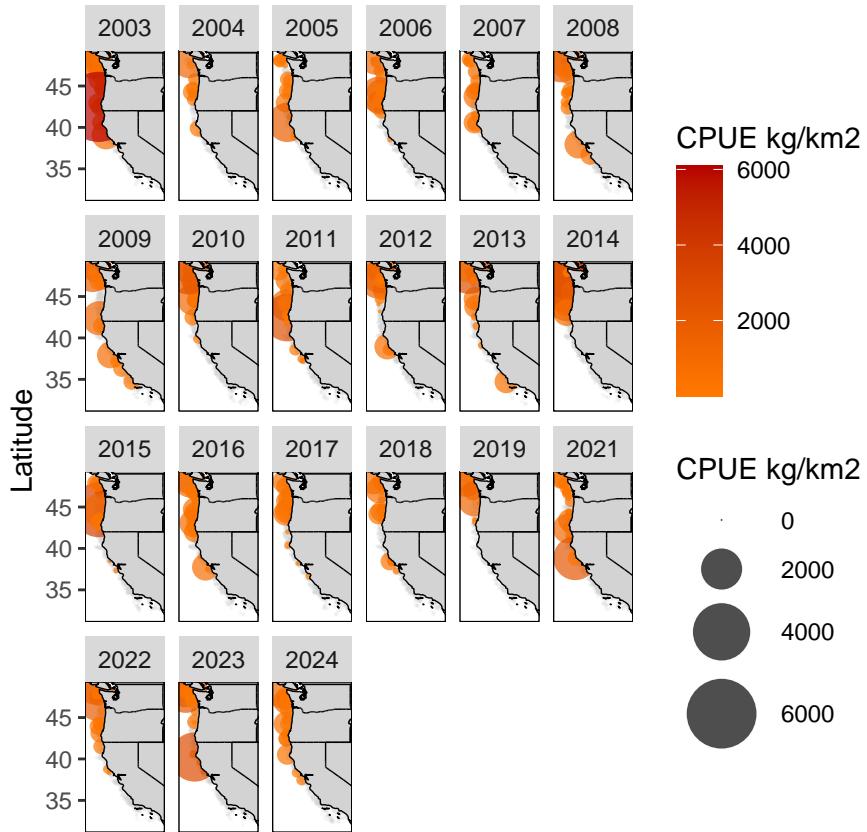
## [1] TRUE
## [1] TRUE
## [1] TRUE
## [1] TRUE

Conditional age-at-length data
caal <- SurveyAgeAtLen.fn(
  datAL = bio,
  datTows = catch,
  strat.df = WGBTStrata,
  lgthBins = seq(10, 74, 2),
  ageBins = 0:65,
  dir = file.path(dirname(getwd()), "Data", "NWFSC.Combo_CAA"),
  printfolder = "")
```

```

PlotMap.fn(
  dat = catch)
```





## Triennial Data

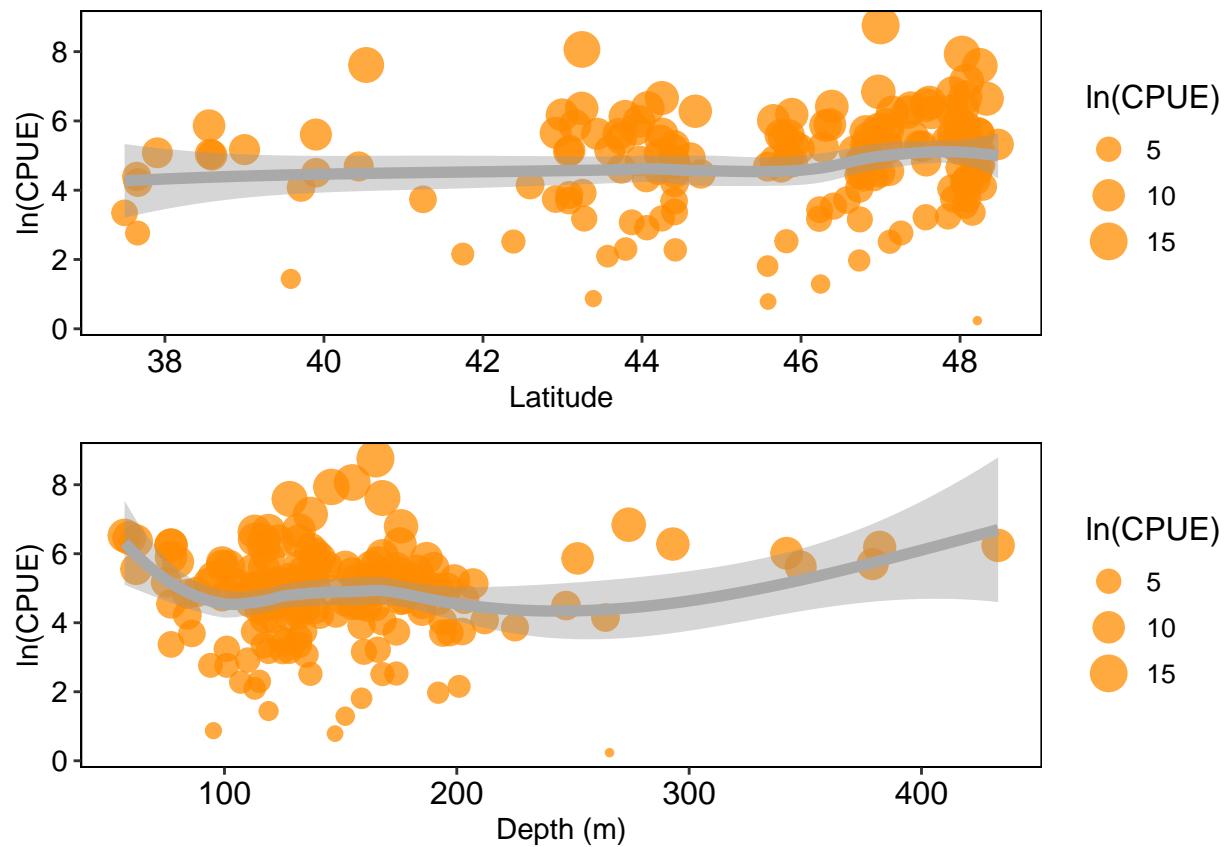
Initial pull and plotting of data: Triennial

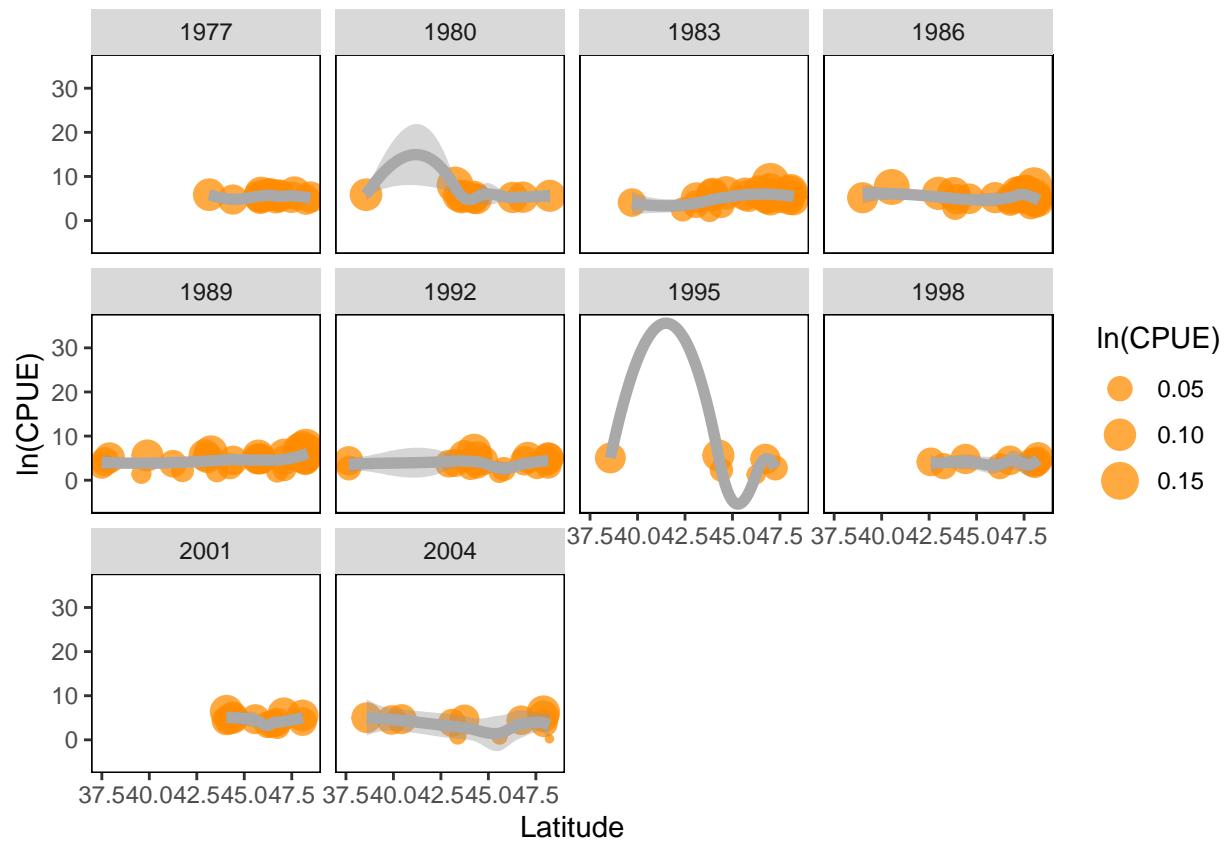
```
catchT = pull_catch(
  common_name = "yelloweye rockfish",
  survey = "Triennial")

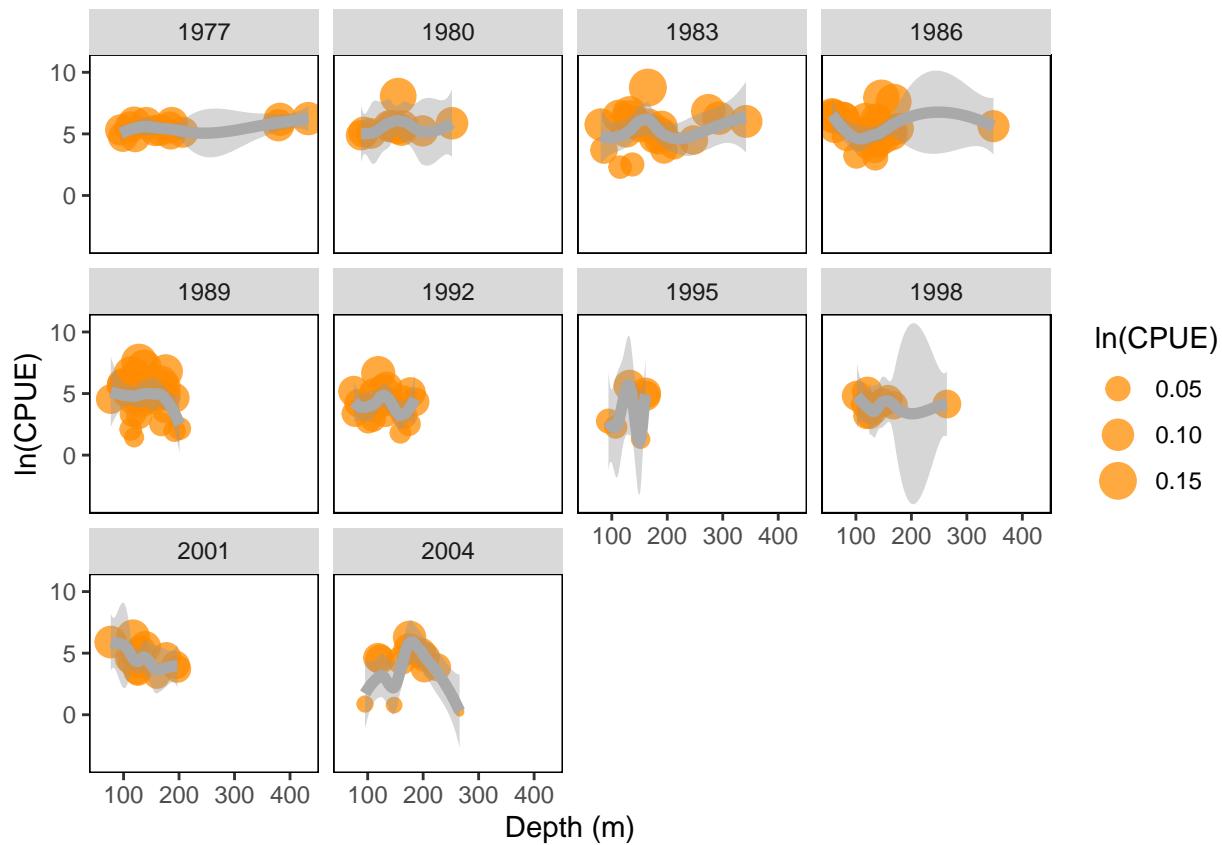
#this is creating a list, just use $age_data
bioT = pull_bio(
  common_name = "yelloweye rockfish",
  survey = "Triennial")

bioT <- bioT$length_data # Ian said to use length data

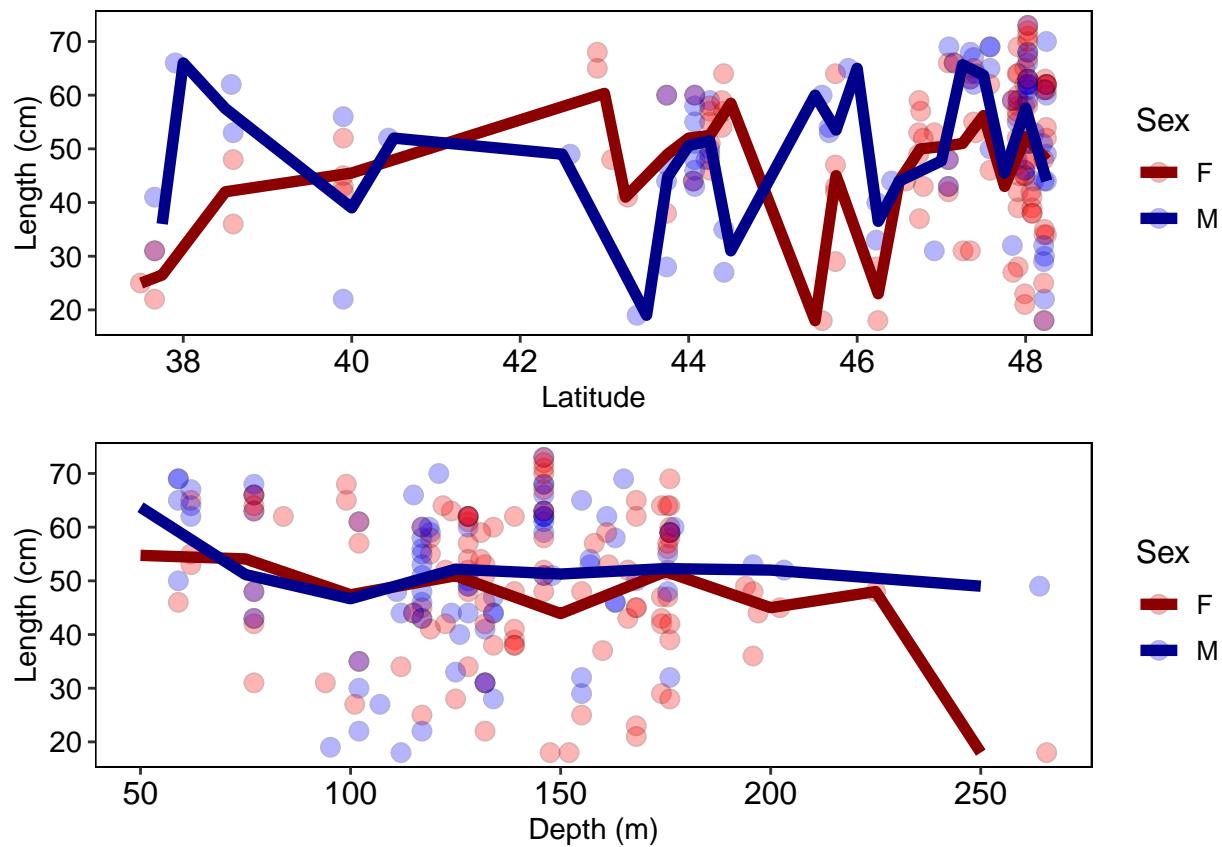
plot_cpue(
  catch = catchT)
```

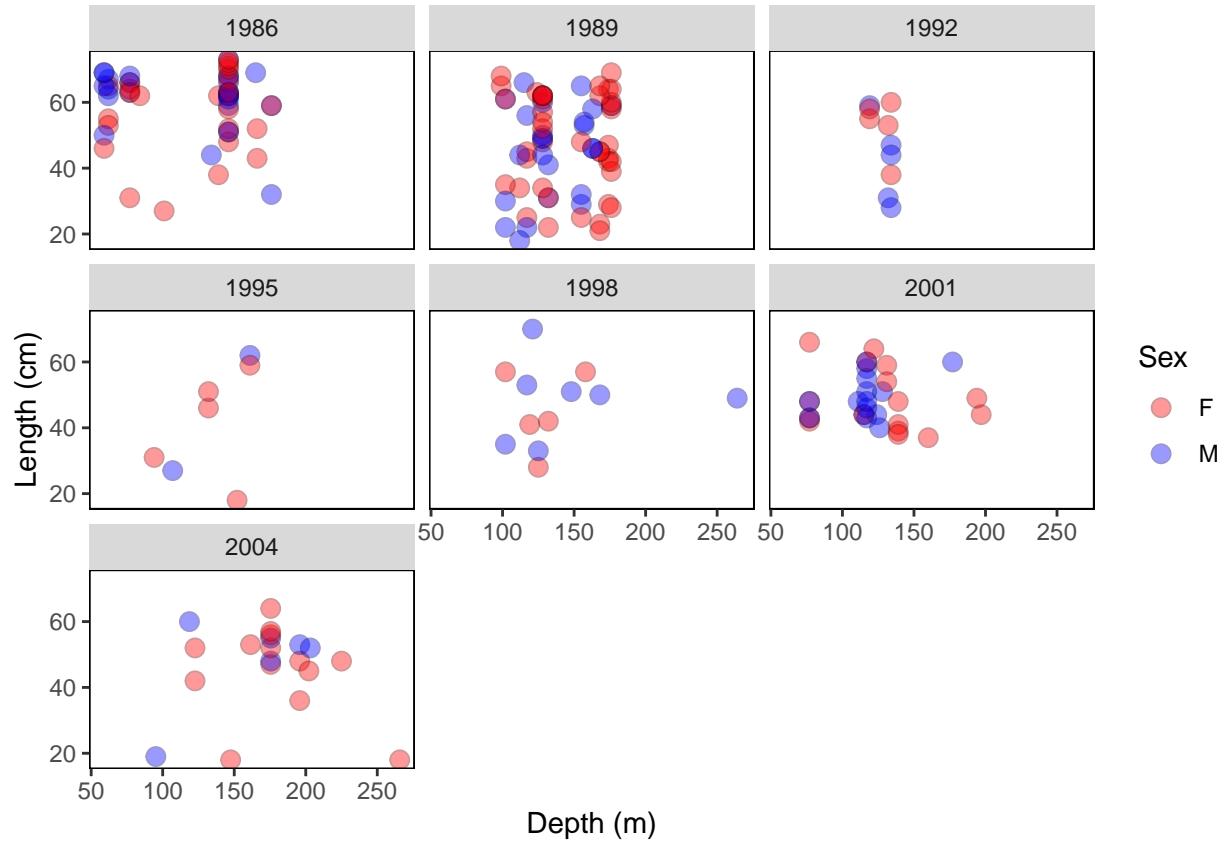


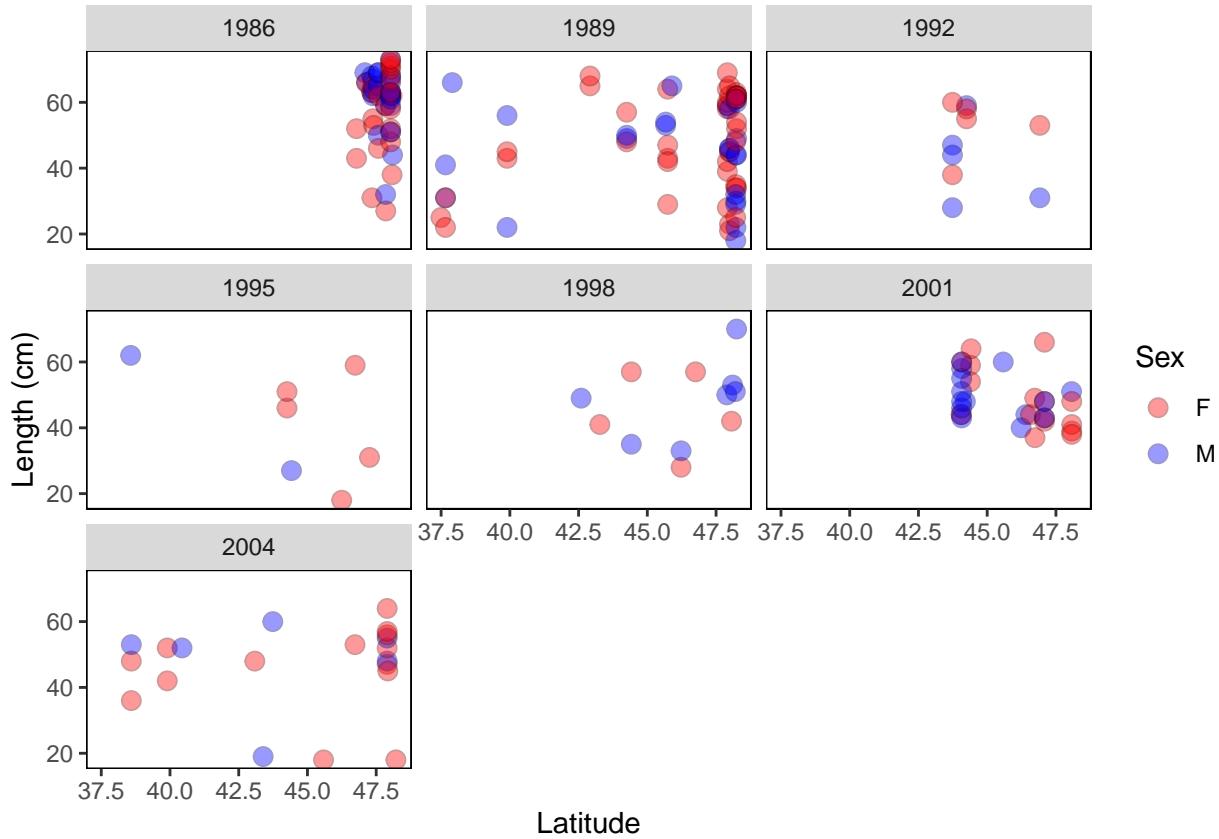




```
plot_bio_patterns(
  bio = bioT,
  col_name = "Length_cm")
```







```

wh_plot_proportion(
  data_catch = catchT,
  data_bio = bioT
)
## [1] "C:/Users/elizabeth.gugliotti/Documents/github_repos/Sebastes_ruberrimus_2025/Rcode/plots/presen"
## [2] "C:/Users/elizabeth.gugliotti/Documents/github_repos/Sebastes_ruberrimus_2025/Rcode/plots/presen"
## [3] "C:/Users/elizabeth.gugliotti/Documents/github_repos/Sebastes_ruberrimus_2025/Rcode/plots/sex_by"
## [4] "C:/Users/elizabeth.gugliotti/Documents/github_repos/Sebastes_ruberrimus_2025/Rcode/plots/sex_by"

```

Define the strata

```

Triennial_strata = CreateStrataDF.fn(names=c("OR", "WA"),
                                       depths.shallow = c(55, 55),
                                       depths.deep    = c(350, 350),
                                       lats.south     = c(42, 46),
                                       lats.north     = c(46, 49))

```

Calculate designed based index of abundance

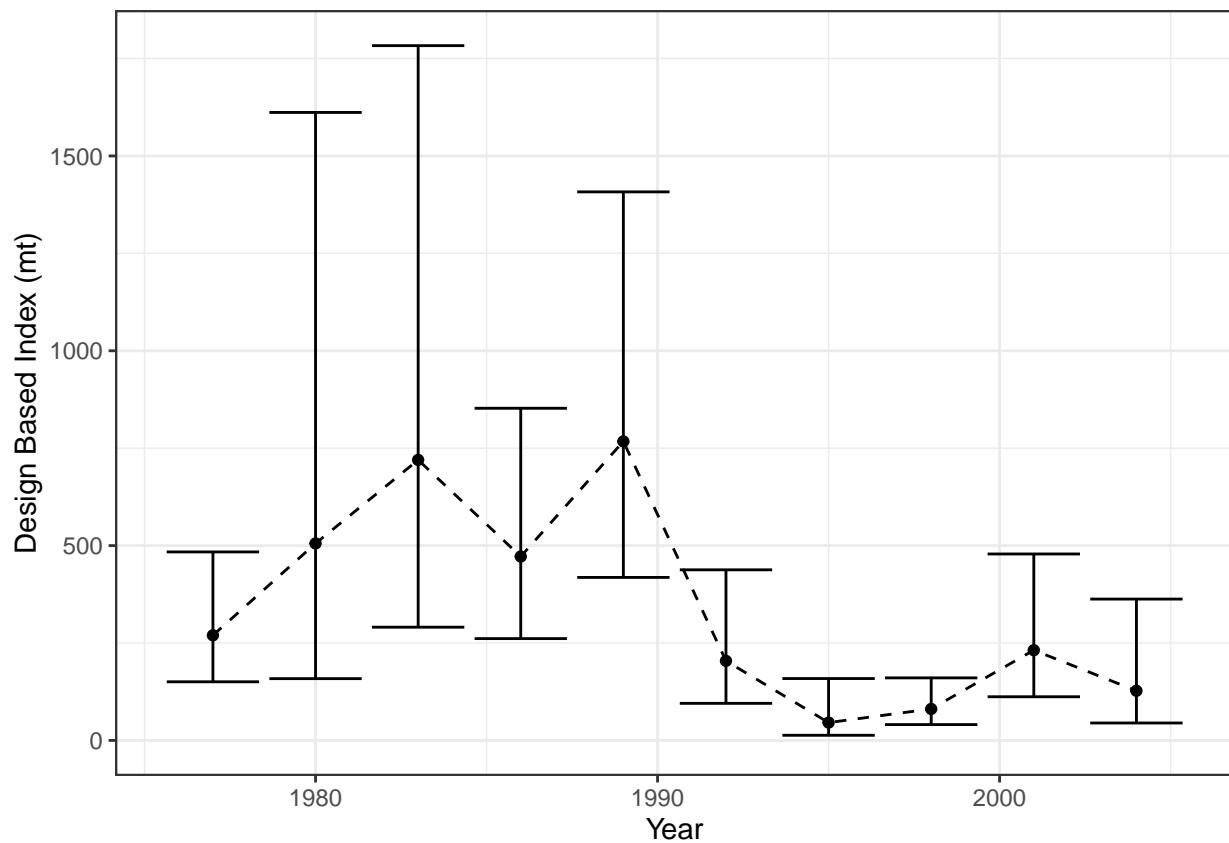
```

biomassT = get_design_based(
  data = catchT,
  strata = Triennial_strata,
  dir = file.path(dirname(getwd()), "Data", "NWFSC.Combo_and_Tri_design_based_indices"),
  printfolder = "")
## [1] TRUE

```

Plot coastwide

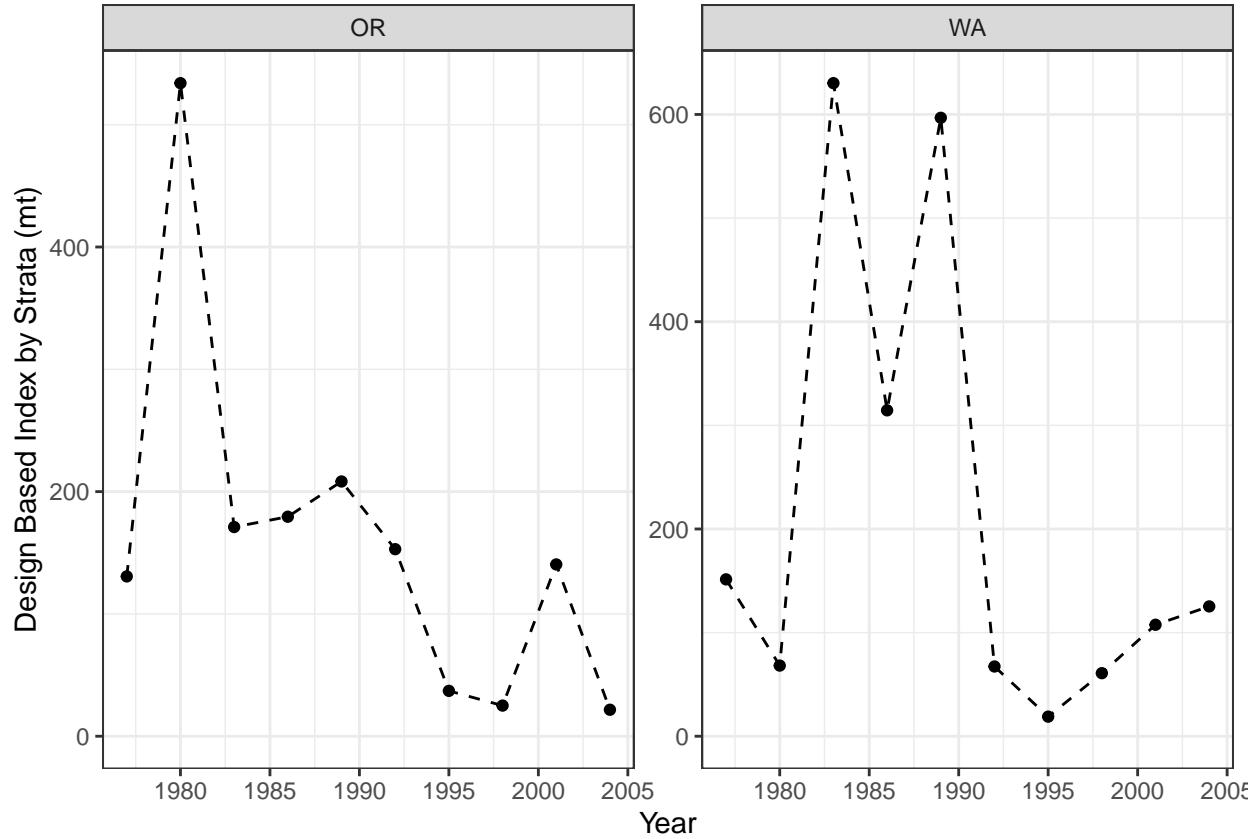
```
plot_index(  
  data = biomassT,  
  plot = 1)
```



```
# not a lot of fish...
```

Plot index for each strata

```
plot_index(  
  data = biomassT,  
  plot = 2)
```

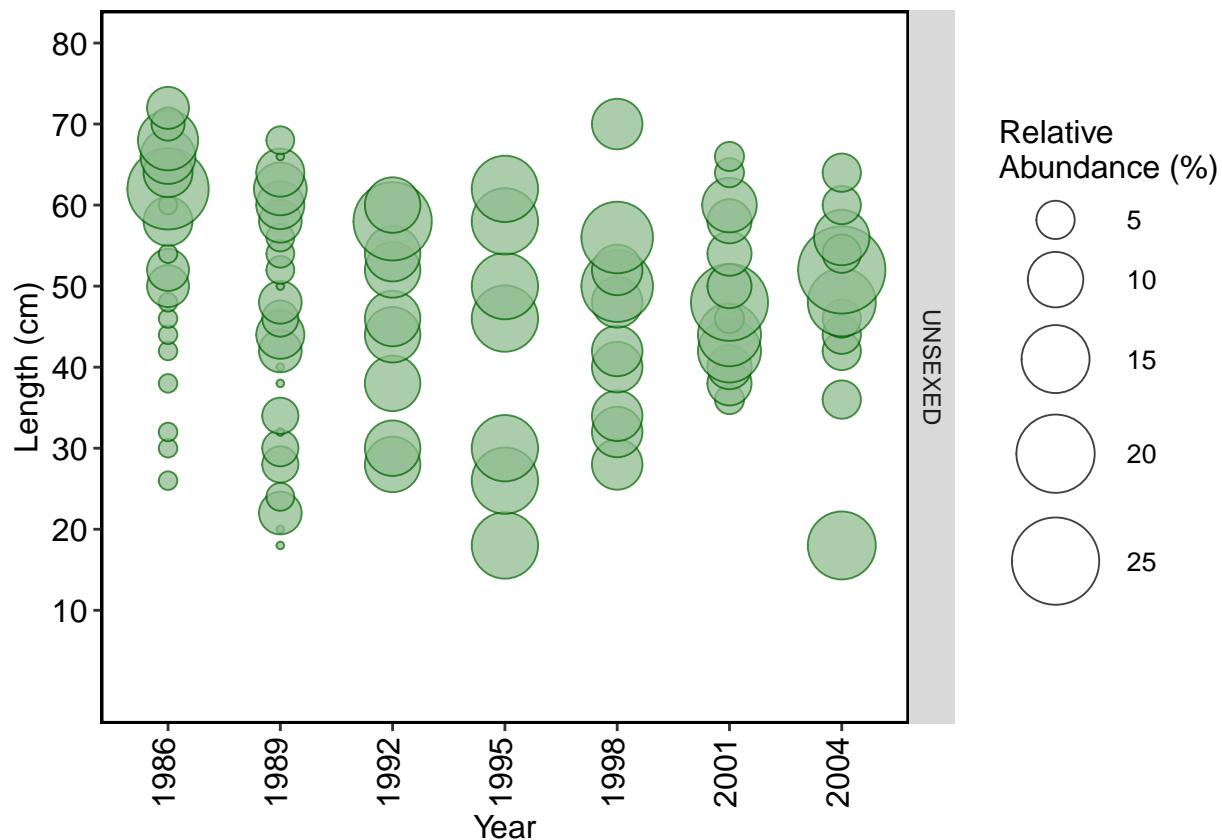


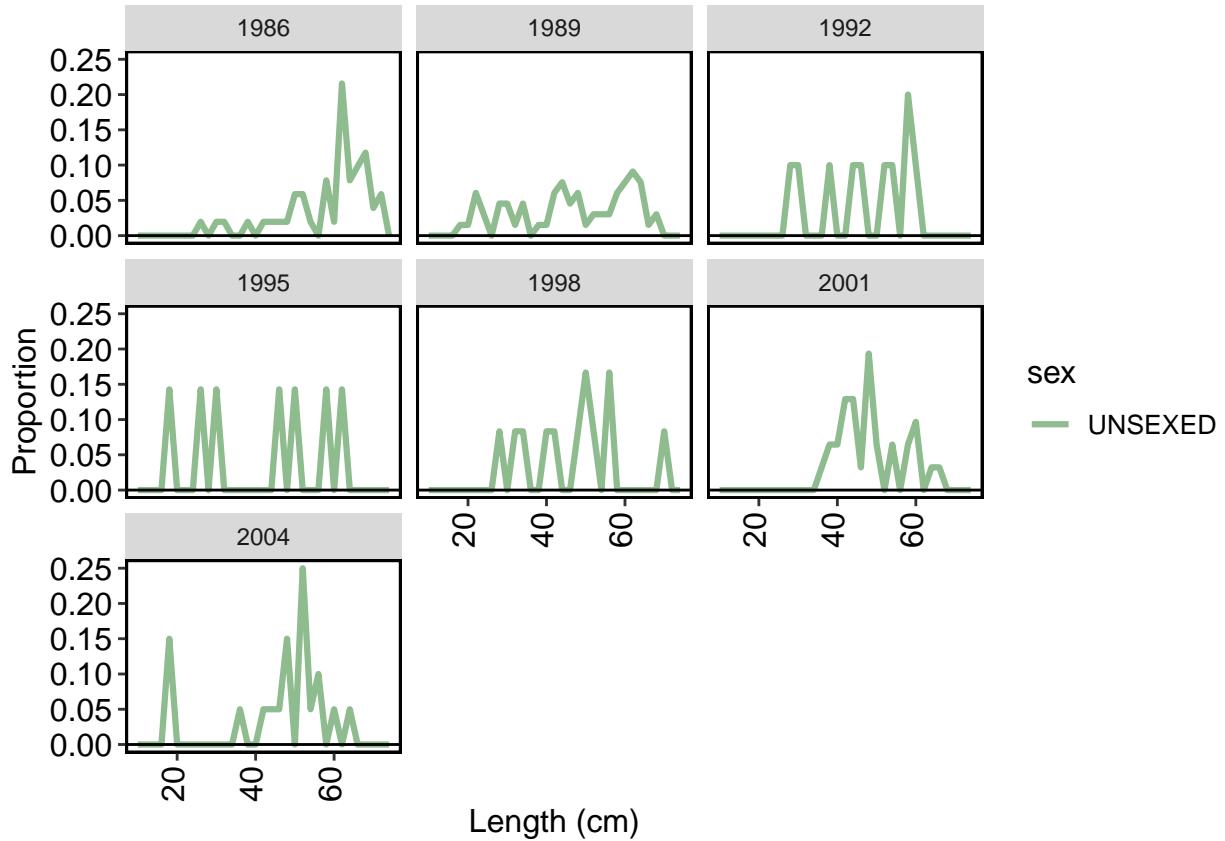
Length composition data

```
# Expanded length comps are not used for the yelloweye assessment
# length_compsT <- get_expanded_comps(
#   bio_data = bioT,
#   catch_data = catchT,
#   comp_bins = seq(10, 74, 2),
#   strata = Triennial_strata,
#   comp_column_name = "length_cm",
#   output = "full_expansion_ss3_format",
#   two_sex_comps = FALSE, #single sex model
#   input_n_method = "stewart_hamel")
#
# plot_comps(
#   data = length_compsT)

raw_length_compsT <- get_raw_comps(
  data = bioT,
  comp_bins = seq(10, 74, 2),
  comp_column_name = "length_cm",
  dir = getwd(),
  printfolder = "",
  two_sex_comps = FALSE) #single sex model

plot_comps(
  data = raw_length_compsT)
```





```
## [1] TRUE
## [1] TRUE
## [1] TRUE
## [1] TRUE
```

No marginal age composition data for triennial survey

*#no age data from Triennial survey?*

```
#age_compsT <- get_expanded_comps(
#  bio_data = bioT,
#  catch_data = catchT,
#  comp_bins = 0:65,
#  strata = Triennial_strata,
#  comp_column_name = "age",
#  output = "full_expansion_ss3_format",
#  two_sex_comps = FALSE, #single sex model
#  input_n_method = "stewart_hamel")

#plot_comps(
#  data = age_compsT)

#raw_age_compsT <- get_raw_comps(
#  data = bioT,
#  comp_bins = 0:65,
#  comp_column_name = "age",
```

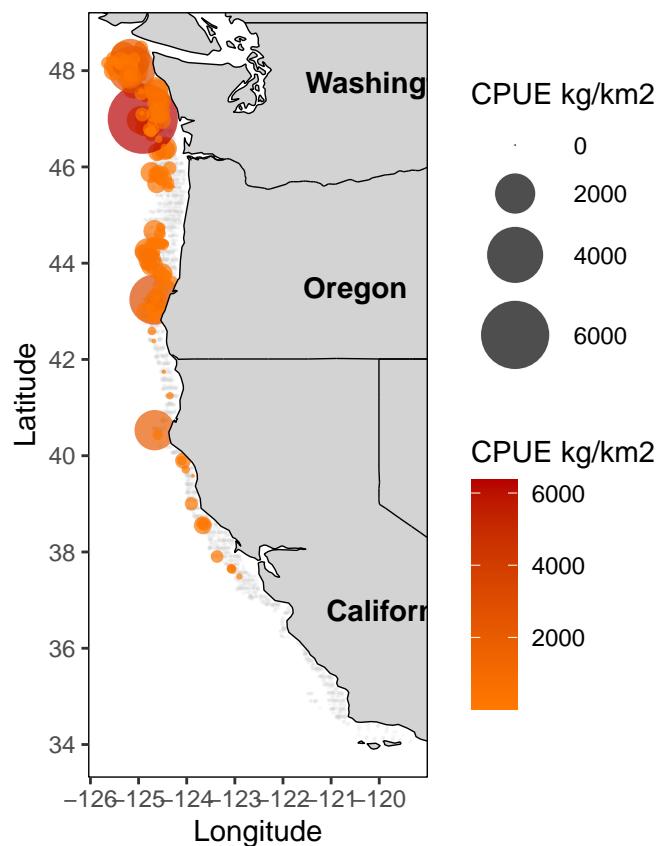
```
#      two_sex_comps = FALSE) #single sex model
```

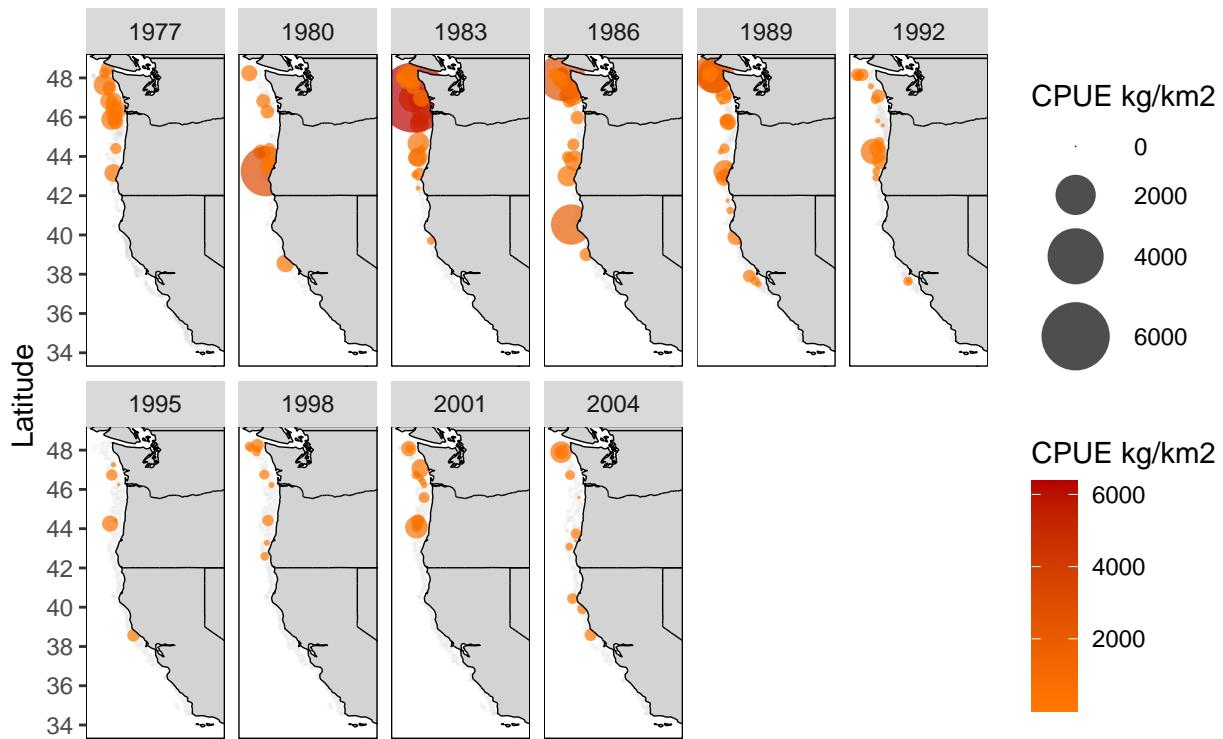
No conditional age-at-length data for triennial survey

```
#no age data from Triennial survey?
```

```
#caalT <- SurveyAgeAtLen.fn(  
#  datAL = bioT,  
#  datTows = catchT,  
#  strat.df = Triennial_strata,  
#  lgthBins = seq(10, 74, 2),  
#  ageBins = 0:65)
```

```
PlotMap.fn(  
  dat = catchT)
```





## Weight-Length Relationship

```
# Ian said we can just use the nwfsc.combo data and don't need to filter by the same latitudes as the i
wt_len_est <- estimate_weight_length(
  bio,
  col_length = "Length_cm",
  col_weight = "Weight_kg",
  verbose = FALSE
)

wt_len_est

##      sex median_intercept          SD          A          B
## 1 female    7.027898e-06 0.11126214 7.071533e-06 3.254269
## 2   male    7.328479e-06 0.09047821 7.358537e-06 3.233642
## 3    all    7.094611e-06 0.10339147 7.132633e-06 3.247382

plot_weight_length(
  bio,
  dir = getwd(),
  estimates = wt_len_est,
  col_length = "length_cm",
  col_weight = "weight_kg",
  two_sex = FALSE,
  add_save_name = NULL,
  height = 7,
```

```

width = 7,
dpi = 300
)

```

## WCGOP Fisheries Discards

```

gemm_data <- pull_gemm("yelloweye rockfish",
                        dir = file.path(dirname(getwd()), "Data"),
                        verbose = TRUE)

head(gemm_data, n = 20)

##           cv      grouping          sector        species
## 29       NA Yelloweye rockfish At-Sea Hake CP Yelloweye Rockfish
## 166      NA Yelloweye rockfish At-Sea Hake MSCV Yelloweye Rockfish
## 239      NA Yelloweye rockfish California Recreational Yelloweye Rockfish
## 371      NA Yelloweye rockfish CS - Bottom Trawl Yelloweye Rockfish
## 639      NA Yelloweye rockfish CS EM - Bottom Trawl Yelloweye Rockfish
## 804      NA Yelloweye rockfish CS EM - Pot Yelloweye Rockfish
## 933      NA Yelloweye rockfish Incidental Yelloweye Rockfish
## 1226     0.2010 Yelloweye rockfish LE Sablefish - Hook & Line Yelloweye Rockfish
## 1313     0.7071 Yelloweye rockfish LE Sablefish - Pot Yelloweye Rockfish
## 1393      NA Yelloweye rockfish Midwater Hake Yelloweye Rockfish
## 1451      NA Yelloweye rockfish Midwater Rockfish Yelloweye Rockfish
## 1648     0.1962 Yelloweye rockfish Nearshore Yelloweye Rockfish
## 1953     0.4029 Yelloweye rockfish OA Fixed Gear - Hook & Line Yelloweye Rockfish
## 2064      NA Yelloweye rockfish Oregon Recreational Yelloweye Rockfish
## 2297      NA Yelloweye rockfish Research Yelloweye Rockfish
## 2411      NA Yelloweye rockfish Tribal Shoreside Yelloweye Rockfish
## 2449      NA Yelloweye rockfish Washington Recreational Yelloweye Rockfish
## 2734      NA Yelloweye rockfish California Recreational Yelloweye Rockfish
## 2828      NA Yelloweye rockfish CS - Bottom Trawl Yelloweye Rockfish
## 3133      NA Yelloweye rockfish CS EM - Bottom Trawl Yelloweye Rockfish
##           total_discard_and_landings_mt total_discard_mt
## 29             0.008090189   0.008090189
## 166            0.005692350   0.001974691
## 239            23.110335868  22.226033249
## 371            0.331968730   0.032594694
## 639            0.079833076   0.000000000
## 804            0.000907194   0.000000000
## 933            0.056699628   0.000000000
## 1226           4.959751182   4.957029600
## 1313           0.013064238   0.005353089
## 1393           0.009554114   0.000028577
## 1451           0.013154508   0.000000195
## 1648           3.823393417   3.823393417
## 1953           4.541011780   4.536022212
## 2064           15.252869152  15.002164320
## 2297           2.547652171   0.000000000
## 2411           8.510387372   0.000000000
## 2449           6.134381586   5.936130949
## 2734           11.444385043  10.392699488
## 2828           0.736515570   0.013035885

```

```

## 3133          0.027669418      0.000000000
##      total_discard_with_mort_rates_applied_and_landings_mt
## 29                  0.008090189
## 166                 0.005692350
## 239                 9.588233406
## 371                 0.331968730
## 639                 0.079833076
## 804                 0.000907194
## 933                 0.056699628
## 1226                4.959751182
## 1313                0.013064238
## 1393                0.009554114
## 1451                0.013154508
## 1648                1.785947257
## 1953                4.541011780
## 2064                3.903488643
## 2297                2.547652171
## 2411                8.510387372
## 2449                0.198250637
## 2734                3.801367438
## 2828                0.736515570
## 3133                0.027669418

##      total_discard_with_mort_rates_applied_mt total_landings_mt      type year
## 29                  0.008090189      0.000000000 groundfish 2023
## 166                 0.001974691      0.003717658 groundfish 2023
## 239                 8.703930788      0.884302619 groundfish 2023
## 371                 0.032594694      0.299374036 groundfish 2023
## 639                 0.000000000      0.079833076 groundfish 2023
## 804                 0.000000000      0.000907194 groundfish 2023
## 933                 0.000000000      0.056699628 groundfish 2023
## 1226                4.957029600      0.002721582 groundfish 2023
## 1313                0.005353089      0.007711149 groundfish 2023
## 1393                0.000028577      0.009525538 groundfish 2023
## 1451                0.000000195      0.013154314 groundfish 2023
## 1648                1.785947257      0.000000000 groundfish 2023
## 1953                4.536022212      0.004989567 groundfish 2023
## 2064                3.652783811      0.250704832 groundfish 2023
## 2297                0.000000000      0.000000000 groundfish 2023
## 2411                0.000000000      8.510387372 groundfish 2023
## 2449                0.000000000      0.198250637 groundfish 2023
## 2734                2.749681884      1.051685555 groundfish 2022
## 2828                0.013035885      0.723479685 groundfish 2022
## 3133                0.000000000      0.027669418 groundfish 2022

```