

Prioritizing Marine Aquaculture Locations on the US West Coast

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Introduction

Marine aquaculture represents a critical opportunity for sustainable food production as global seafood demand continues to rise. Unlike land-based meat production, marine aquaculture can provide protein with lower environmental costs when strategically located in suitable environments (Hall et al., 2011). Recent research has demonstrated that global seafood demand could theoretically be met using less than 0.015% of the global ocean area (Gentry et al., 2017), highlighting the importance of identifying optimal locations for development.

This analysis evaluates the potential for marine aquaculture along the US West Coast by identifying Exclusive Economic Zones (EEZs) with suitable environmental conditions for oyster and black abalone cultivation. Using geospatial analysis of sea surface temperature and bathymetry data, we determine which regions offer the greatest area of suitable habitat for each species. The analysis addresses three key questions:

1. Which West Coast EEZs have the largest areas of suitable conditions for oyster aquaculture?
2. Which regions are most suitable for black abalone cultivation?
3. How do suitability patterns differ between these two species?

By developing a generalizable workflow, this analysis can be applied to assess aquaculture potential for additional species, supporting evidence-based decision-making for sustainable marine resource development.

Setup

```
# Load required packages
library(tidyverse)
library(sf)
```

```
library(terra)
library(here)
library(tmap)
library(kableExtra)
library(testthat)
```

Data Loading

Load Exclusive Economic Zones (EEZ)

```
# Load West Coast EEZ shapefile
wc_df <- st_read("data/wc_regions_clean.shp")
```

```
Reading layer `wc_regions_clean' from data source
  `C:\Users\ermil\Documents\MEDS\rellimylime.github.io\posts\eds223-final2\data\wc_regions_c
  using driver `ESRI Shapefile'
Simple feature collection with 5 features and 5 fields
Geometry type: MULTIPOLYGON
Dimension:      XY
Bounding box:   xmin: -129.1635 ymin: 30.542 xmax: -117.097 ymax: 49.00031
Geodetic CRS:   WGS 84
```

```
# Check CRS
st_crs(wc_df)$epsg
```

```
[1] 4326
```

Load Sea Surface Temperature (SST) Data

```
# Load SST rasters for 2008-2012

# Define a function to read rast
read_tif <- function(year) {
  rast(file.path("data", paste0("average_annual_sst_", year, ".tif")))
}
```

```
# Apply function
sst_list <- lapply(2008:2012, read_tif)

# Combine into raster stack
sst_stack <- rast(sst_list)

# Inspect properties
names(sst_stack)
```

```
[1] "average_annual_sst_2008" "average_annual_sst_2009"
[3] "average_annual_sst_2010" "average_annual_sst_2011"
[5] "average_annual_sst_2012"
```

```
st_crs(sst_stack["average_annual_sst_2008"])
```

Coordinate Reference System:

User input: WGS 84

wkt:

```
GEOGCRS["WGS 84",
  DATUM["unknown",
    ELLIPSOID["WGS84",6378137,298.257223563,
      LENGTHUNIT["metre",1,
        ID["EPSG",9001]]],
    PRIMEM["Greenwich",0,
      ANGLEUNIT["degree",0.0174532925199433,
        ID["EPSG",9122]]],
    CS[ellipsoidal,2],
      AXIS["latitude",north,
        ORDER[1],
        ANGLEUNIT["degree",0.0174532925199433,
          ID["EPSG",9122]]],
      AXIS["longitude",east,
        ORDER[2],
        ANGLEUNIT["degree",0.0174532925199433,
          ID["EPSG",9122]]]]
```

```
summary(values(sst_stack))
```

average_annual_sst_2008	average_annual_sst_2009	average_annual_sst_2010
Min. :278.7	Min. :278.1	Min. :279.9

1st Qu.:285.3	1st Qu.:285.7	1st Qu.:285.7
Median :287.1	Median :287.7	Median :287.4
Mean :287.1	Mean :287.7	Mean :287.5
3rd Qu.:289.1	3rd Qu.:289.7	3rd Qu.:289.4
Max. :301.4	Max. :301.5	Max. :301.0
NA's :82337	NA's :82481	NA's :82572
average_annual_sst_2011	average_annual_sst_2012	
Min. :278.9	Min. :278.1	
1st Qu.:285.5	1st Qu.:285.5	
Median :287.0	Median :287.0	
Mean :287.1	Mean :287.2	
3rd Qu.:288.8	3rd Qu.:289.0	
Max. :307.3	Max. :310.2	
NA's :82228	NA's :82130	

```
global(sst_stack, fun = "isNA")
```

```

isNA
average_annual_sst_2008 82337
average_annual_sst_2009 82481
average_annual_sst_2010 82572
average_annual_sst_2011 82228
average_annual_sst_2012 82130

```

Load Bathymetry (Depth) Data

```

# Load depth raster
depth <- rast(("data/depth.tif"))

# Check properties
st_crs(depth)

```

Coordinate Reference System:

User input: WGS 84

wkt:

```

GEOGCRS["WGS 84",
  ENSEMBLE["World Geodetic System 1984 ensemble",
    MEMBER["World Geodetic System 1984 (Transit)"],
    MEMBER["World Geodetic System 1984 (G730)"],
    MEMBER["World Geodetic System 1984 (G873)"],

```

```

    MEMBER["World Geodetic System 1984 (G1150)"],
    MEMBER["World Geodetic System 1984 (G1674)"],
    MEMBER["World Geodetic System 1984 (G1762)"],
    MEMBER["World Geodetic System 1984 (G2139)"],
    MEMBER["World Geodetic System 1984 (G2296)"],
    ELLIPSOID["WGS 84",6378137,298.257223563,
      LENGTHUNIT["metre",1]],
    ENSEMBLEACCURACY[2.0]],
  PRIMEM["Greenwich",0,
    ANGLEUNIT["degree",0.0174532925199433]],
  CS[ellipsoidal,2],
    AXIS["geodetic latitude (Lat)",north,
      ORDER[1],
      ANGLEUNIT["degree",0.0174532925199433]],
    AXIS["geodetic longitude (Lon)",east,
      ORDER[2],
      ANGLEUNIT["degree",0.0174532925199433]],
  USAGE[
    SCOPE["Horizontal component of 3D system."],
    AREA["World."],
    BBOX[-90,-180,90,180]],
  ID["EPSG",4326]]

```

Verify Coordinate Reference Systems

```

# Check sst crs
crs(sst_stack, describe = TRUE)$code

```

```
[1] NA
```

```

# Set epsg explicitly from metadata
crs(sst_stack) <- "EPSG:4326"

# Check again
crs(sst_stack, describe = TRUE)$code

```

```
[1] "4326"
```

```
# Check extent makes sense for this CRS  
ext(sst_stack)
```

SpatExtent : -131.98475233, -114.986717027556, 29.9930522526667, 49.988422964 (xmin, xmax, ymin, ymax)

```
# Verify sst and depth crs are the same  
crs(sst_stack) == crs(depth)
```

[1] TRUE

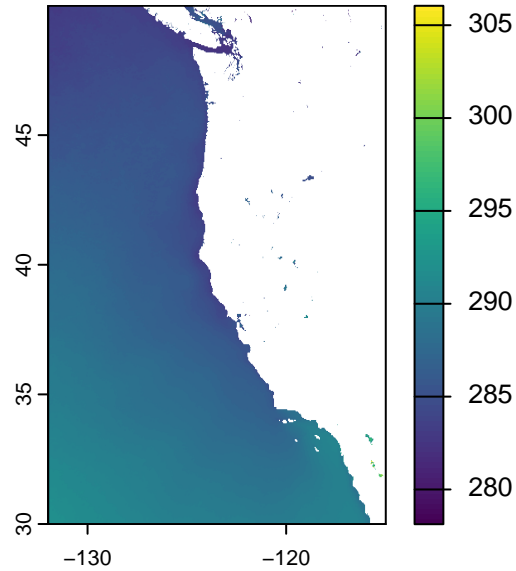
```
# Check if wc_df crs matches  
crs(sst_stack, describe = TRUE)$code == st_crs(wc_df)$epsg
```

[1] TRUE

Data Processing

Calculate Mean SST (2008-2012)

```
# Calculate mean SST across all years using app()  
sst_mean <- app(sst_stack, mean, na.rm = TRUE)  
  
# Visualize with plot()  
plot(sst_mean)
```



Convert SST from Kelvin to Celsius

```
# Convert from Kelvin to Celsius (subtract 273.15)
sst_mean_c <- sst_mean - 273.15

# Check temperature range with summary()
summary(sst_mean_c)
```

Warning: [summary] used a sample

```
      mean
Min.   : 7.02
1st Qu.:12.38
Median :14.07
Mean   :14.16
3rd Qu.:16.06
Max.   :32.90
NA's   :41907
```


Align Depth Data with SST

```
# Crop depth to SST extent using crop()
depth_crop <- crop(depth, sst_mean_c)

# Resample depth to match SST resolution using resample() with method = "near"
depth_resampled <- resample(depth_crop, sst_mean_c, method = "near")

# Verify alignment
res(depth_resampled) == res(sst_mean_c)
```

```
[1] TRUE TRUE
```

```
ext(depth_resampled) == ext(sst_mean_c)
```

```
[1] TRUE
```

```
crs(depth_resampled) == crs(sst_mean_c)
```

```
[1] TRUE
```

Species Suitability Analysis

Oyster Suitability

Define Oyster Requirements

Oysters require: - Sea surface temperature: 11-30°C - Depth: 0-70 meters below sea level

```
# Reclassify SST for oysters using classify()
# Create rcl matrix: <11°C = 0, 11-30°C = 1, >30°C = 0
sst_rcl_mat_oyster <- matrix(c(-Inf, 11, 0,
                                11, 30, 1,
                                30, Inf, 0),
                              ncol = 3,
                              byrow = TRUE)
```

```

sst_rcl_oyster <- classify(sst_mean_c, sst_rcl_mat_oyster)

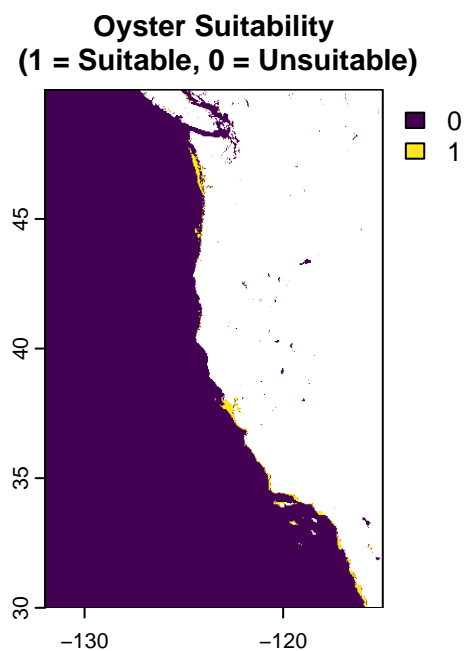
# Reclassify depth for oysters using classify()
# Create rcl matrix: <-70m = 0, -70m to 0m = 1, >0m = 0
depth_rcl_mat_oyster <- matrix(c(-Inf, -70, 0,
                                -70, 0, 1,
                                0, Inf, 0),
                                ncol = 3,
                                byrow = TRUE)

depth_rcl_oyster <- classify(depth_resampled, depth_rcl_mat_oyster)

# Combine suitability layers
suitable_zones_oyster <- sst_rcl_oyster * depth_rcl_oyster

# Visualize with plot()
plot(suitable_zones_oyster,
     main = "Oyster Suitability\n (1 = Suitable, 0 = Unsuitable)",
     cex.main = 0.9)

```



Black Abalone Suitability

Define Black Abalone Requirements

Black Abalone requires: - Sea surface temperature: 12.2-18.6°C - Depth: 0-6 meters below sea level - Source: SeaLifeBase

```
# Reclassify SST for black abalone using classify()
# Suitable: 12.2-18.6°C = 1, Unsuitable: <12.2°C or >18.6°C = 0
sst_rcl_mat_abalone <- matrix(c(-Inf, 12.2, 0,
                                12.2, 18.6, 1,
                                18.6, Inf, 0),
                              ncol = 3,
                              byrow = TRUE)

# Apply reclassification matrix to mean SST raster
sst_rcl_abalone <- classify(sst_mean_c, sst_rcl_mat_abalone)

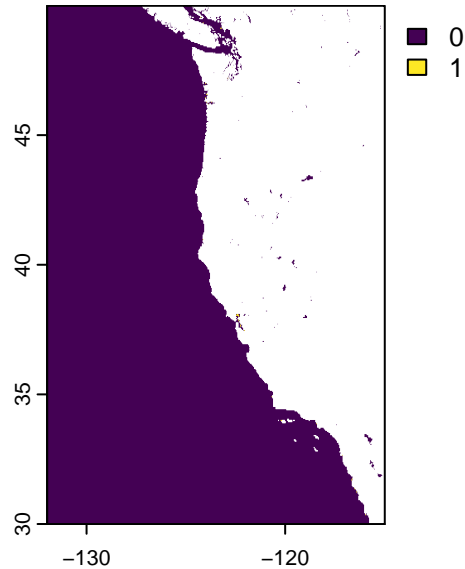
# Reclassify depth for black abalone using classify()
# Suitable: 0-6m below sea level = -6 to 0 in raster values
depth_rcl_mat_abalone <- matrix(c(-Inf, -6, 0,
                                  -6, 0, 1,
                                  0, Inf, 0),
                                ncol = 3,
                                byrow = TRUE)

# Apply reclassification matrix to resampled depth raster
depth_rcl_abalone <- classify(depth_resampled, depth_rcl_mat_abalone)

# Combine suitability layers
suitable_zones_abalone <- sst_rcl_abalone * depth_rcl_abalone

# Visualize with plot()
plot(suitable_zones_abalone, main = "Black Abalone Suitability\n (1 = Suitable, 0 = Unsuitable)",
     cex.main = 0.9)
```

Black Abalone Suitability (1 = Suitable, 0 = Unsuitable)



Calculate Suitable Area by EEZ

Prepare EEZ for Area Calculation

```
# Transform EEZ to match raster CRS
eez <- st_transform(wc_df, crs(sst_mean_c))

# Rasterize EEZ using rasterize() with rgn_id for field
eez_rast <- rasterize(eez, suitable_zones_oyster, field = "rgn_id")
```

Calculate Suitable Area for Oysters

```
# Mask suitable areas to EEZ using mask()
eez_suitable_oyster <- mask(suitable_zones_oyster, eez_rast)

# Calculate cell area in km2 using cellSize()
cell_area_km2 <- cellSize(eez_suitable_oyster, unit = "km")
```

```

# Multiply suitable locations by cell area to get suitable area per cell
suitable_area_oyster <- eez_suitable_oyster * cell_area_km2

# Sum area by EEZ region using zonal()
area_by_region_oyster <- zonal(suitable_area_oyster, eez_rast, fun = "sum", na.rm = TRUE)

# Rename area column for clarity
names(area_by_region_oyster)[2] <- "suitable_area_km2"

# Join results with EEZ sf object to get region names
area_by_region_oyster_joined <- eez %>%
  left_join(area_by_region_oyster, by = "rgn_id") %>%
  arrange(desc(suitable_area_km2))

oyster_results <- area_by_region_oyster_joined %>%
  st_drop_geometry() %>%
  select(rgn, suitable_area_km2) %>%
  head(10)

# Display top 10 regions in table with kable()
kable(oyster_results,
      caption = "Top 10 EEZ Regions for Oyster Aquaculture",
      col.names = c("Region", "Suitable Area (km²)"),
      digits = 2) %>%
  kable_styling(bootstrap_options = c("striped", "hover"))

```

Table 1: Top 10 EEZ Regions for Oyster Aquaculture

Region	Suitable Area (km ²)
Central California	4923.15
Southern California	4096.54
Washington	3224.74
Oregon	1533.09
Northern California	438.15

```

# Test that area calculations are reasonable
test_that("Oyster area calculations are valid", {
  # Check total area > 0
  expect_true(sum(area_by_region_oyster$suitable_area_km2, na.rm = TRUE) > 0)

  # Check all values >= 0

```

```
expect_true(all(area_by_region_oyster$suitable_area_km2 >= 0, na.rm = TRUE))

# Check that we have results for multiple regions
expect_true(nrow(area_by_region_oyster) > 0)
})
```

Test passed

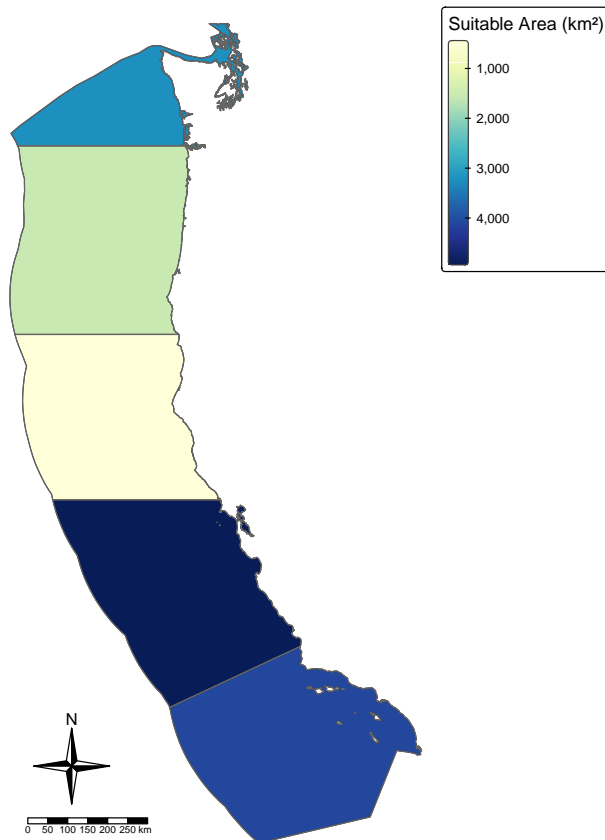
Map Oyster Suitability by EEZ

```
# Create map using tmap
tm_shape(area_by_region_oyster_joined) +
  tm_polygons(
    fill = "suitable_area_km2",
    fill.scale = tm_scale_continuous(values = "brewer.yl_gn_bu"),
    fill.legend = tm_legend(title = "Suitable Area (km²)")
  ) +
  tm_borders(col = "gray40", lwd = 0.5) +
  tm_title("Oyster Aquaculture Suitability by EEZ Region") +
  tm_compass(type = "4star", position = c("left", "bottom")) +
  tm_scalebar(position = c("left", "bottom")) +
  tm_layout(legend.outside = TRUE,
    legend.outside.position = "right",
    frame = FALSE)
```

[plot mode] fit legend/component: Some legend items or map components do not fit well, and are therefore rescaled.

i Set the tmap option `component.autoscale = FALSE` to disable rescaling.

Oyster Aquaculture Suitability by EEZ Region



Calculate Suitable Area for Black Abalone

```
# Mask suitable areas to EEZ using mask()
eez_suitable_abalone <- mask(suitable_zones_abalone, eez_rast)

# Calculate cell area in km² using cellSize()
cell_area_km2_abalone <- cellSize(eez_suitable_abalone, unit = "km")

# Multiply suitable locations by cell area to get suitable area per cell
suitable_area_abalone <- eez_suitable_abalone * cell_area_km2_abalone

# Sum area by EEZ region using zonal()
area_by_region_abalone <- zonal(suitable_area_abalone, eez_rast, fun = "sum", na.rm = TRUE)
```

```

# Rename area column for clarity
names(area_by_region_abalone)[2] <- "suitable_area_km2"

# Join results with EEZ sf object to get region names
area_by_region_abalone_joined <- eez %>%
  left_join(area_by_region_abalone, by = "rgn_id") %>%
  arrange(desc(suitable_area_km2))

abalone_results <- area_by_region_abalone_joined %>%
  st_drop_geometry() %>%
  select(rgn, suitable_area_km2) %>%
  head(10)

# Display top 10 regions in table with kable()
kable(abalone_results,
      caption = "Top 10 EEZ Regions for Black Abalone Aquaculture",
      col.names = c("Region", "Suitable Area (km²)"),
      digits = 2) %>%
  kable_styling(bootstrap_options = c("striped", "hover"))

```

Table 2: Top 10 EEZ Regions for Black Abalone Aquaculture

Region	Suitable Area (km ²)
Central California	186.24
Washington	103.47
Southern California	89.01
Northern California	16.22
Oregon	14.90

```

# Test that area calculations are reasonable
test_that("Black Abalone area calculations are valid", {
  # Check total area > 0
  expect_true(sum(area_by_region_abalone$suitable_area_km2, na.rm = TRUE) > 0)

  # Check all values >= 0
  expect_true(all(area_by_region_abalone$suitable_area_km2 >= 0, na.rm = TRUE))

  # Check that we have results for multiple regions
  expect_true(nrow(area_by_region_abalone) > 0)
})

```


Test passed

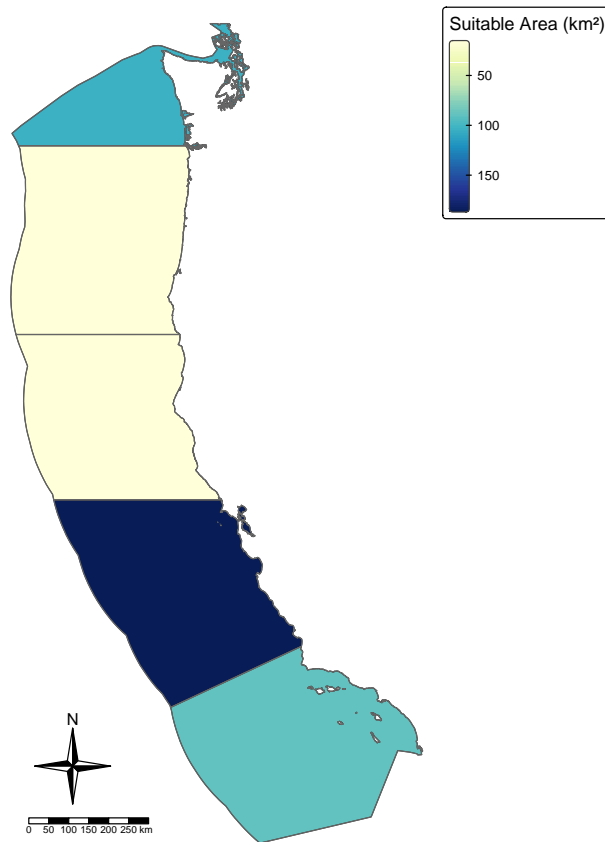
Map Black Abalone Suitability by EEZ

```
# Create map using tmap
tm_shape(area_by_region_abalone_joined) +
  tm_polygons(
    fill = "suitable_area_km2",
    fill.scale = tm_scale_continuous(values = "brewer.yl_gn_bu"),
    fill.legend = tm_legend(title = "Suitable Area (km²)")
  ) +
  tm_borders(col = "gray40", lwd = 0.5) +
  tm_title("Black Abalone Aquaculture Suitability by EEZ Region") +
  tm_compass(type = "4star",
    position = c("left", "bottom")) +
  tm_scalebar(position = c("left", "bottom")) +
  tm_layout(legend.outside = TRUE,
    legend.outside.position = "right",
    frame = FALSE)
```

[plot mode] fit legend/component: Some legend items or map components do not fit well, and are therefore rescaled.

i Set the tmap option `component.autoscale = FALSE` to disable rescaling.

Black Abalone Aquaculture Suitability by EEZ Region



Generalizable Function

Create Suitability Function

```
calculate_suitability <- function(min_temp, max_temp, min_depth, max_depth, species_name) {  
  
  # Validate inputs  
  if (min_temp >= max_temp) {  
    stop("min_temp must be less than max_temp")  
  }  
  
  if (min_depth >= max_depth) {  
    stop("min_depth must be less than max_depth")  
  }  
  
}
```

```

}

if (!is.character(species_name) || species_name == "") {
  stop("species_name must be a non-empty string")
}

# Create reclassification matrix for sea surface temperature
sst_rcl_matrix <- matrix(c(-Inf, min_temp, 0,
                           min_temp, max_temp, 1,
                           max_temp, Inf, 0),
                         ncol = 3,
                         byrow = TRUE)

# Apply reclassification matrix to mean SST raster
sst_rcl <- classify(sst_mean_c, sst_rcl_matrix)

# Create reclassification matrix for depth
depth_rcl_matrix <- matrix(c(-Inf, min_depth, 0,
                             min_depth, max_depth, 1,
                             max_depth, Inf, 0),
                           ncol = 3,
                           byrow = TRUE)

# Apply reclassification matrix to resampled depth raster
depth_rcl <- classify(depth_resampled, depth_rcl_matrix)

# Combine temperature and depth suitability using multiplication
suitable_zones <- sst_rcl * depth_rcl

# Mask suitable zones to only include areas within EEZ boundaries
eez_suitable <- mask(suitable_zones, eez_rast)

# Calculate the area of each raster cell in square kilometers
cell_area <- cellSize(eez_suitable, unit = "km")

# Multiply suitable locations (1s) by cell area (unsuitable cells will have 0 area)
suitable_area <- eez_suitable * cell_area

# Sum suitable area within each EEZ region
area_by_region <- zonal(suitable_area, eez_rast, fun = "sum", na.rm = TRUE)

# Rename the area column to be more descriptive
names(area_by_region)[2] <- "suitable_area_km2"

```

```

# Join area calculations with EEZ spatial data to add region names
results_joined <- eez %>%
  left_join(area_by_region, by = "rgn_id") %>%
  arrange(desc(suitable_area_km2))

# Create a summary table showing top 10 regions
results_table <- results_joined %>%
  st_drop_geometry() %>%
  select(rgn, suitable_area_km2) %>%
  head(10)

# Create choropleth map showing suitable area by EEZ region
map <- tm_shape(results_joined) +
  tm_polygons(
    fill = "suitable_area_km2",
    fill.scale = tm_scale_continuous(values = "brewer.yl_gn_bu"),
    fill.legend = tm_legend(title = "Suitable Area (km²)")
  ) +
  tm_borders(col = "gray40", lwd = 0.5) +
  tm_title(paste(species_name, "Aquaculture Suitability by EEZ Region")) +
  tm_compass(type = "4star", position = c("right", "top")) +
  tm_scalebar(stack = "vertical", position = c("left", "bottom")) +
  tm_layout(legend.outside = TRUE,
    legend.outside.position = "right",
    frame = FALSE)

return(list(
  results = results_table,
  full_data = results_joined,
  map = map
))
}

```

Apply Function to Oysters

```

# Call function with oyster parameters
oyster_output <- calculate_suitability(
  min_temp = 11,
  max_temp = 30,
  min_depth = -70,

```

```

max_depth = 0,
species_name = "Oyster"
)

# Display results table
kable(oyster_output$results,
      caption = "Top 10 EEZ Regions for Oyster Aquaculture (Function Output)",
      col.names = c("Region", "Suitable Area (km2)"),
      digits = 2) %>%
kable_styling(bootstrap_options = c("striped", "hover"))

```

Table 3: Top 10 EEZ Regions for Oyster Aquaculture (Function Output)

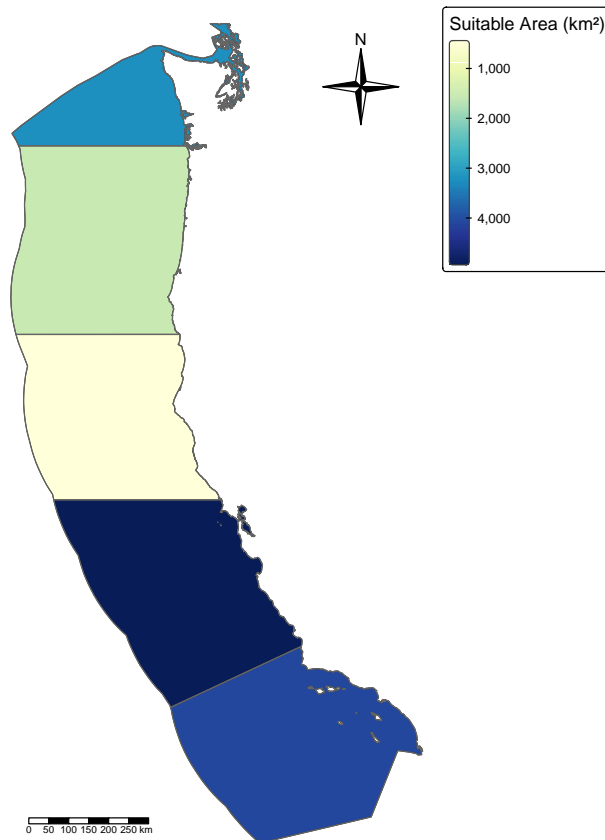
Region	Suitable Area (km ²)
Central California	4923.15
Southern California	4096.54
Washington	3224.74
Oregon	1533.09
Northern California	438.15

```

# Display map and table
print(oyster_output$map)

```

Oyster Aquaculture Suitability by EEZ Region



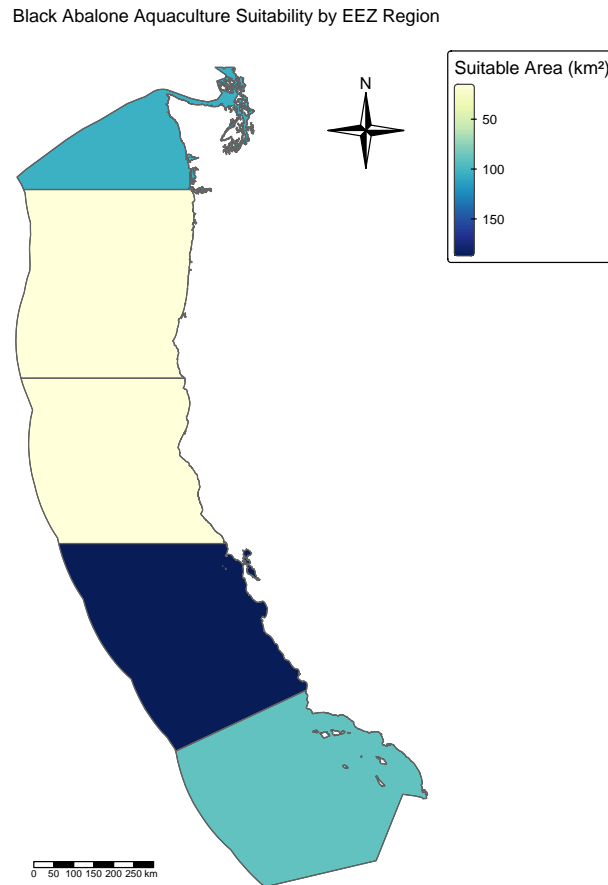
Apply Function to Black Abalone

```
# Call function with Black Abalone parameters
abalone_output <- calculate_suitability(
  min_temp = 12.2,
  max_temp = 18.6,
  min_depth = -6,
  max_depth = 0,
  species_name = "Black Abalone"
)

# Display map
abalone_output$map
```

[plot mode] fit legend/component: Some legend items or map components do not fit well, and are therefore rescaled.

i Set the tmap option `component.autoscale = FALSE` to disable rescaling.



```
# Display results table
kable(abalone_output$results,
      caption = "Top 10 EEZ Regions for Black Abalone Aquaculture (Function Output)",
      col.names = c("Region", "Suitable Area (km²)"),
      digits = 2) %>%
kable_styling(bootstrap_options = c("striped", "hover"))
```

Table 4: Top 10 EEZ Regions for Black Abalone Aquaculture (Function Output)

Region	Suitable Area (km ²)
Central California	186.24
Washington	103.47
Southern California	89.01
Northern California	16.22
Oregon	14.90

Discussion

Key Findings

Oyster Aquaculture: Central California has the most suitable area (4,069.88 km²), followed by Southern California (3,757.28 km²) and Washington (2,378.31 km²). These three regions contain over 90% of all suitable oyster habitat on the West Coast. Oregon has moderate potential (1,074.27 km²), while Northern California has minimal suitable area.

Black Abalone Aquaculture: Total suitable area is dramatically smaller—only 199.88 km² across the entire West Coast. Central California dominates with 152.37 km² (76% of suitable habitat). Washington and Southern California have limited suitable areas, while Oregon and Northern California have none.

Key Difference: Black abalone has 20 times less suitable habitat than oysters, potentially due to its narrow temperature tolerance (12.2-18.6°C vs. 11-30°C) and shallow depth requirement (0-6m vs. 0-70m). Both species show peak suitability in Central California, but oysters have viable habitat across all West Coast regions while black abalone is highly restricted.

Limitations

- **Temporal resolution:** Annual SST averages (2008-2012) may miss critical seasonal temperature extremes or variations important for specific life stages
- **Environmental factors excluded:** Does not account for ocean currents, dissolved oxygen, salinity, wave exposure, water quality, or harmful algal blooms
- **Socioeconomic constraints:** Ignores proximity to markets/infrastructure, conflicts with existing ocean uses, and regulatory/permitting requirements
- **Species requirements:** Based on optimal growth ranges from SeaLifeBase; may not reflect full tolerance ranges or life-stage-specific needs
- **Spatial resolution:** Bathymetry resolution (~500m) may be too coarse to accurately represent ideal depths, especially for shallow-water species like black abalone

Data Sources

Sea Surface Temperature: - Source: NOAA Coral Reef Watch. NOAA 5km Daily Global Satellite Sea Surface Temperature Anomaly v3.1 - URL: https://coralreefwatch.noaa.gov/product/5km/index_5km_ssta.php

Bathymetry: - Source: GEBCO Compilation Group (2022). GEBCO_2022 Grid - DOI: 10.5285/e0f0bb80-ab44-2739-e053-6c86abc0289c

Exclusive Economic Zones: - Source: Marine Regions - URL: <https://www.marineregions.org/>

Species Requirements: - Source: SeaLifeBase - URL: <https://www.sealifebase.org/>