

# Seminoff EPac green turtle Stable Isotope Data Analysis

Lisa Komoroske

2016-12-22

## Metadata

**Site** - an ordinal code for each site

**Site code** - 3 letter code for each site

**Ordered\_SITE** - combined site code with ordered # roughly North to South for graphing ordering **Location\_Label** - shortened locatin names for graphing labeling purposes **Location** - location of turtle capture

**LAB ID** - self explanatory

**Collection Date** - self explanatory

**Run Date** - self explanatory

**%N** - elemental concentration of N. that is, how much each sample is made up of nitrogen. this is used as a diagnostic to know sample quality (anything outside of ~9-17% N raises a red flag)

**%C** - elemental concentration of C. that is, how much each sample is made up of carbon. this is used as a diagnostic to know sample quality (anything outside of ~40-60% C raises a red flag)

**d15N** - stable isotope value for N

**d13C** - stable isotope value for C

**Color** - rarely filled in. This is largely for the Galapagos and Colombia, where black turtles (eastern Pacific stock) and yellow turtles (west Pacific origins) co-exist. Safe to say that anything that is not filled in here would be a 'black' morph.

**SCL** - straight carapace length

**CCL\_calc\_fromSCL** - used formula from Seminoff et al. 2003 to interpolate CCLs from SCLs **CCL\_empirical** - curved carapace length-these are only the empirically collected values

**CCL\_combined** - curved carapace length-I pasted over all the empirical values, and then for ones that were missing empirical CCL but had **CCL\_calc\_fromSCL**, I added these in; so this is the combined variable that we'll use for size relationships

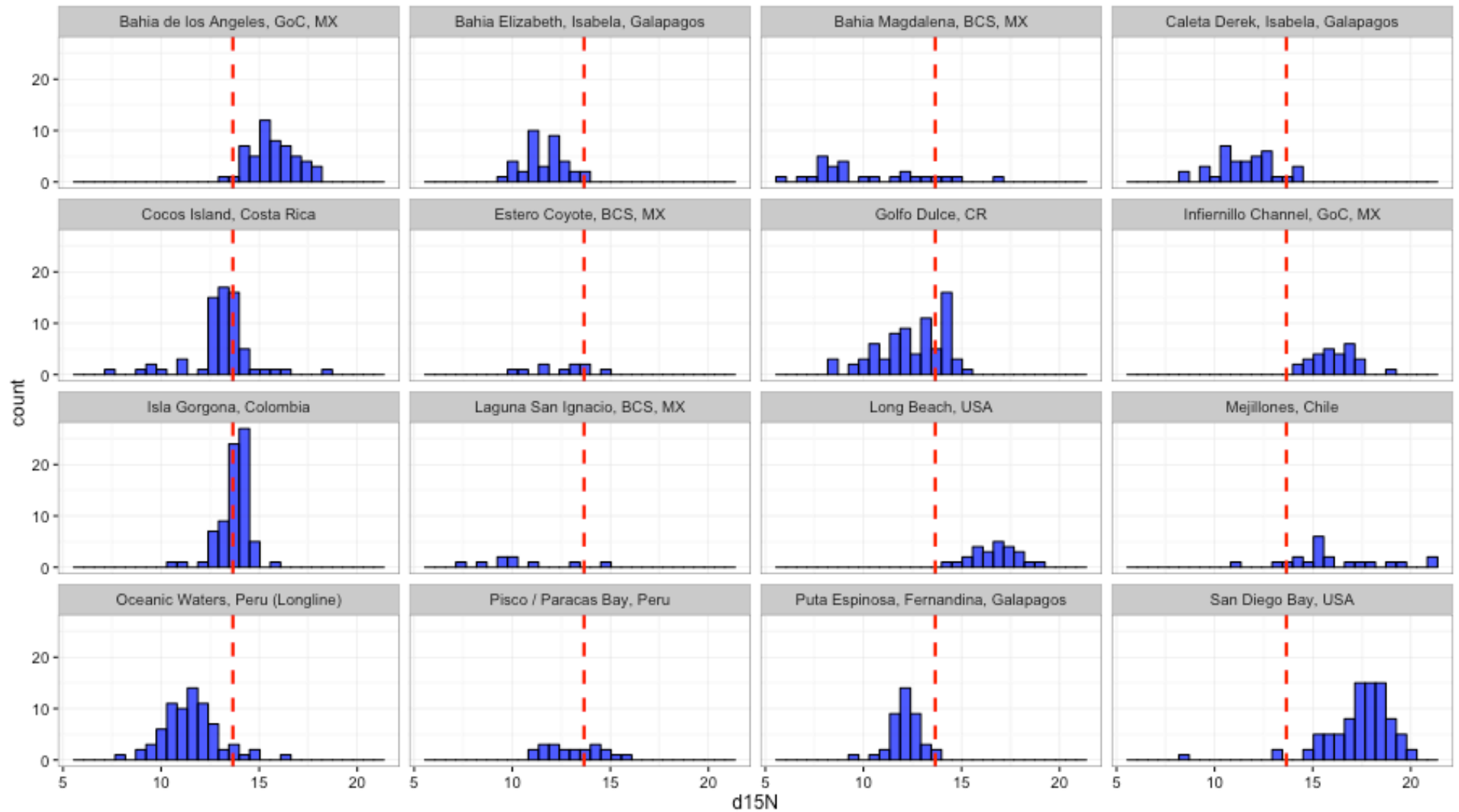
## Summary (Sample Sizes)

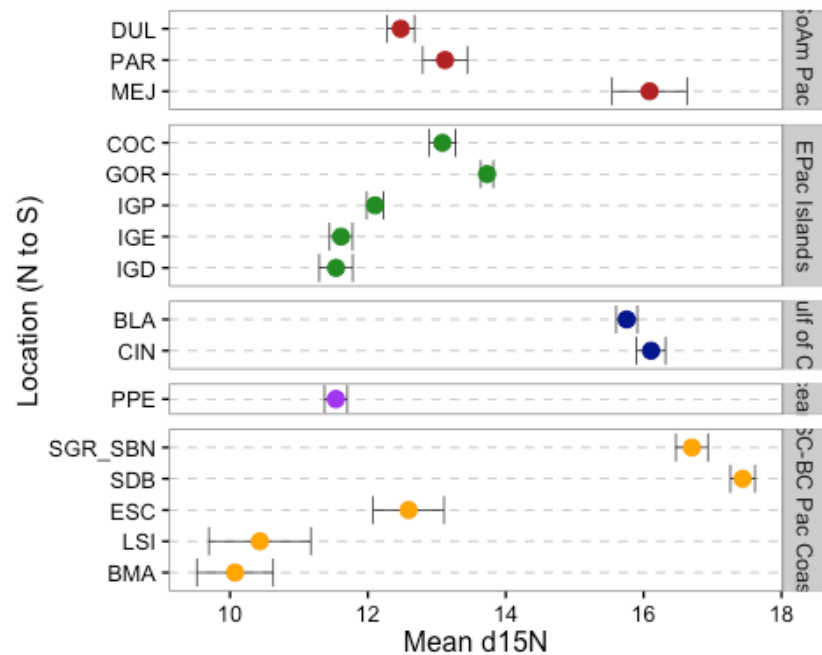
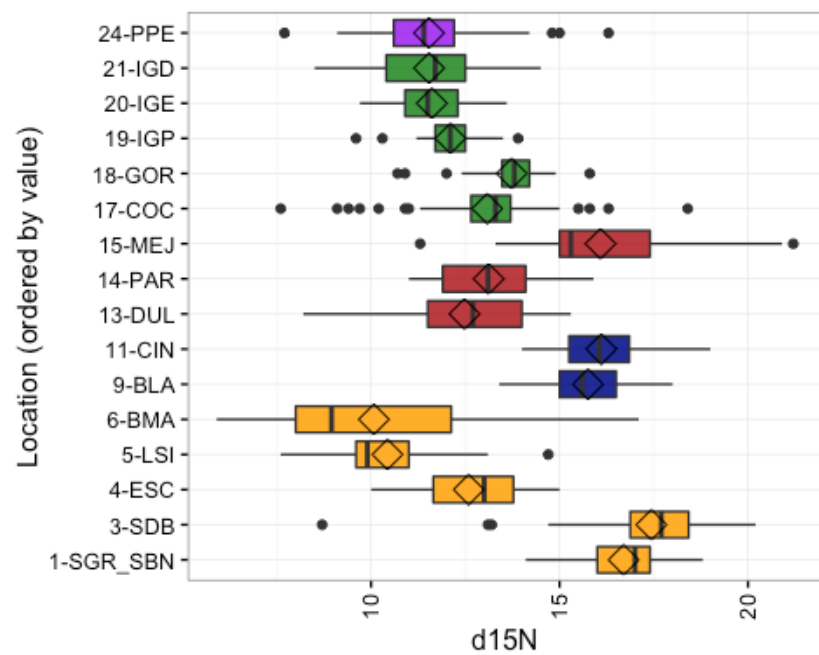
##

##	1-SGR_SBN	3-SDB	4-ESC	5-LSI	6-BMA	9-BLA	11-CIN
##	25	88	10	9	26	53	28
##	13-DUL	14-PAR	15-MEJ	17-COC	18-GOR	19-IGP	20-IGE
##	74	21	21	67	76	41	37

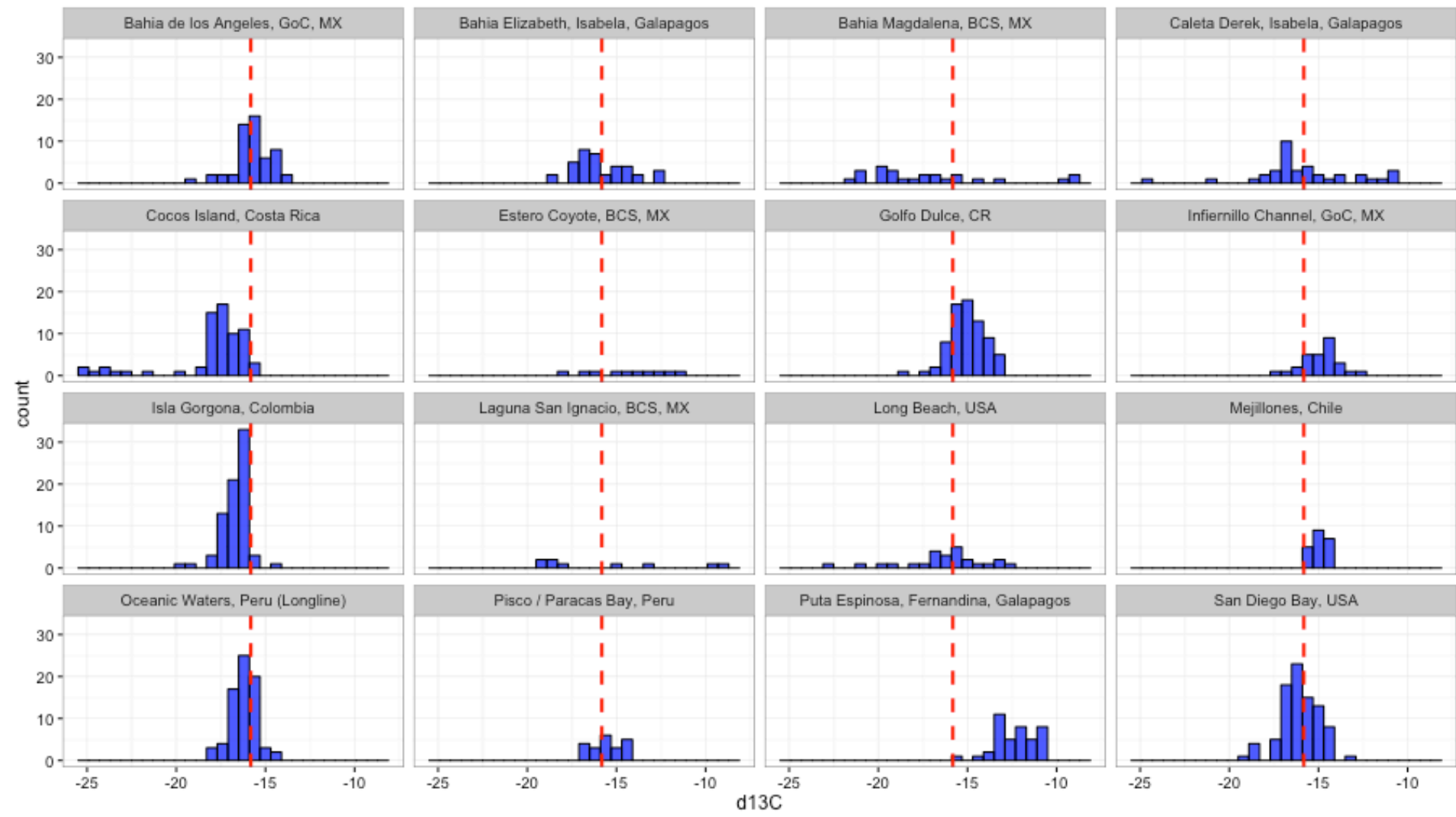
##	21-IGD	24-PPE
##	37	74

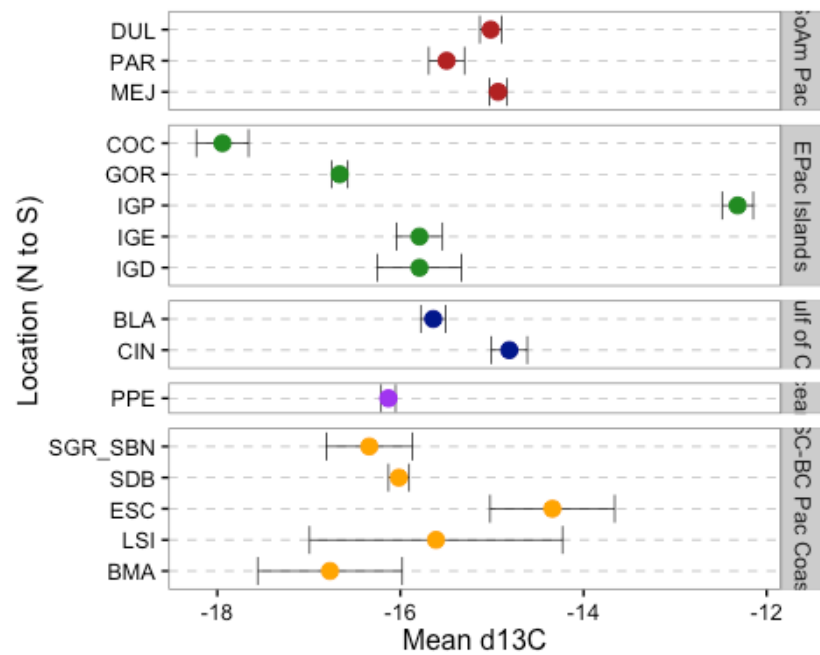
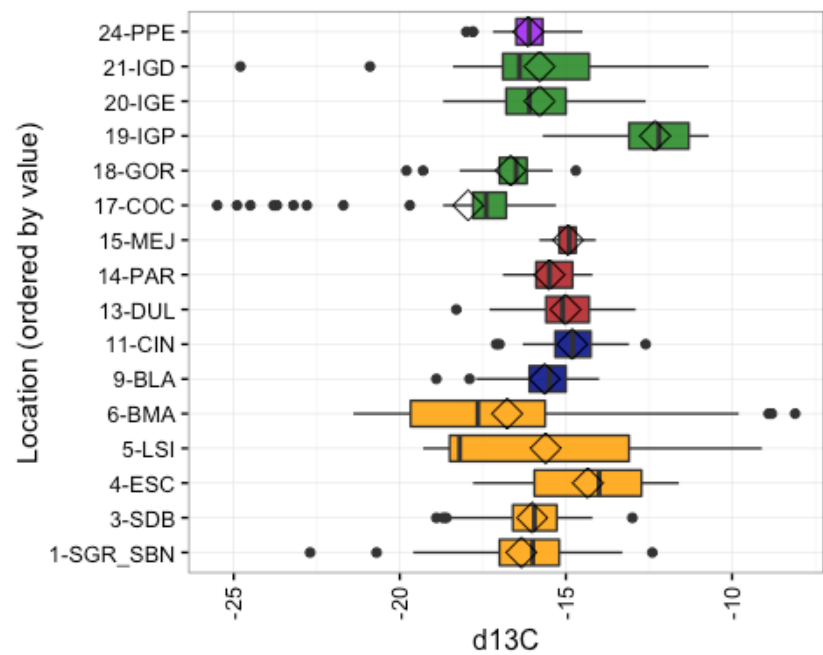
Laguna San Ignacio now only one with lower sample #s-confirm with Jeff keeping/seperate with Estero Coyote



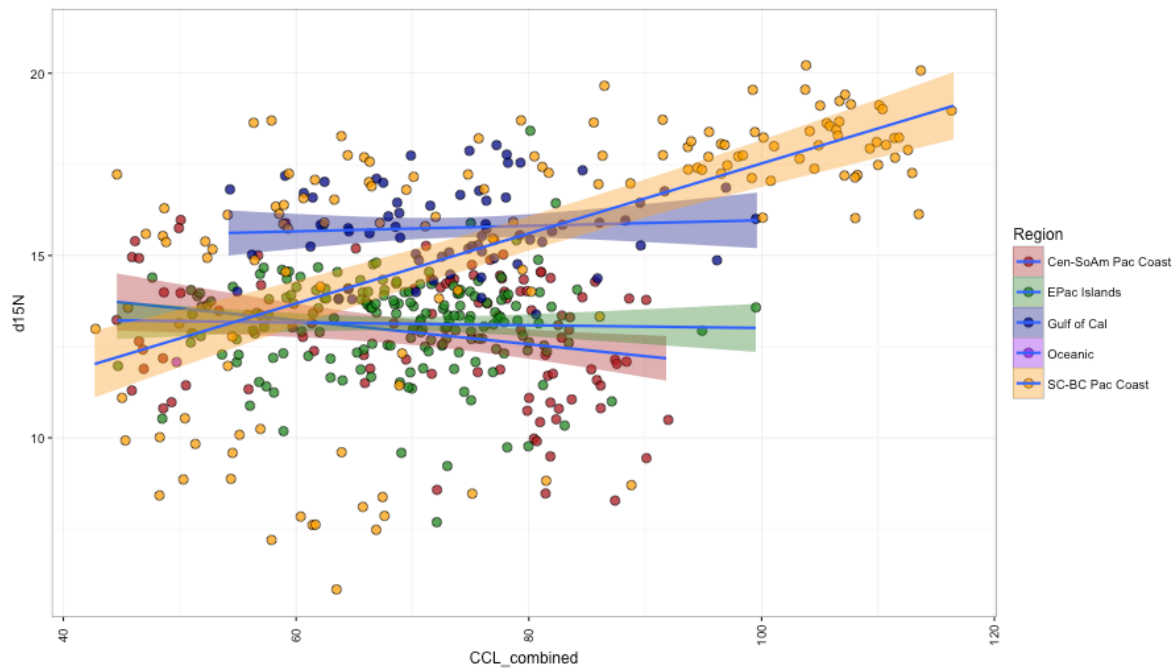
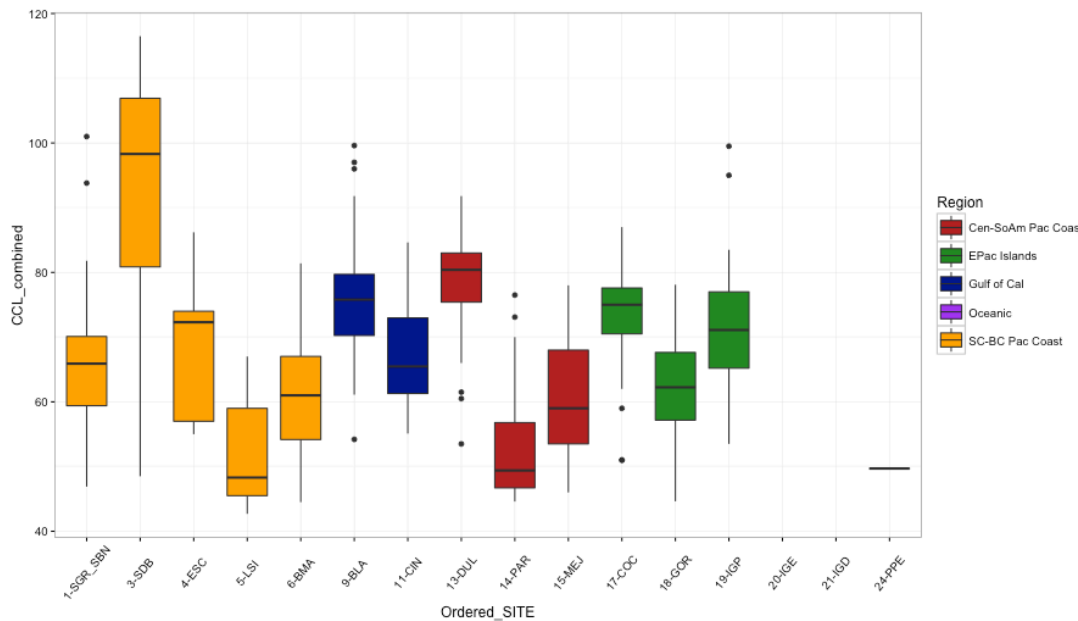


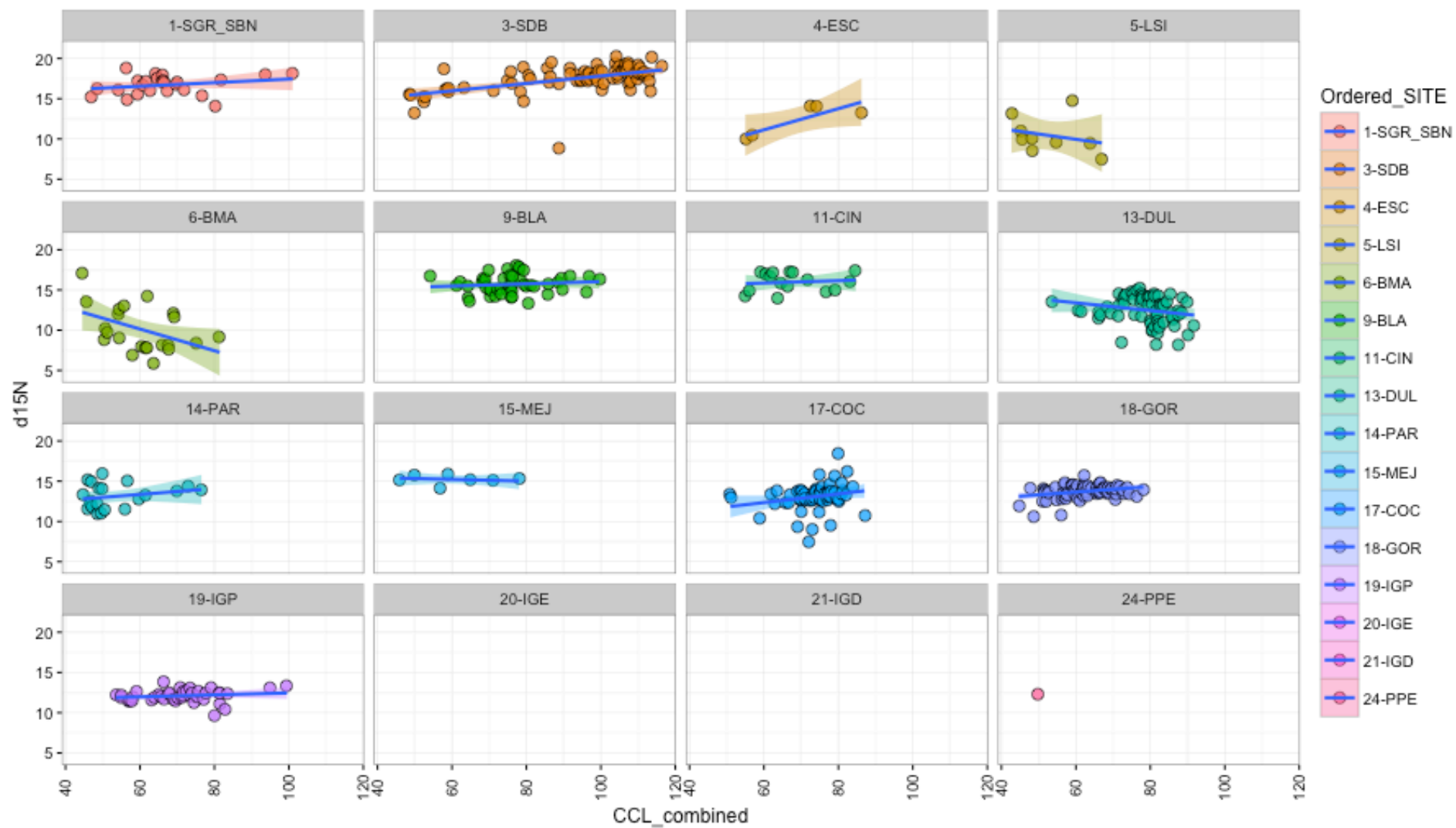
Carbon

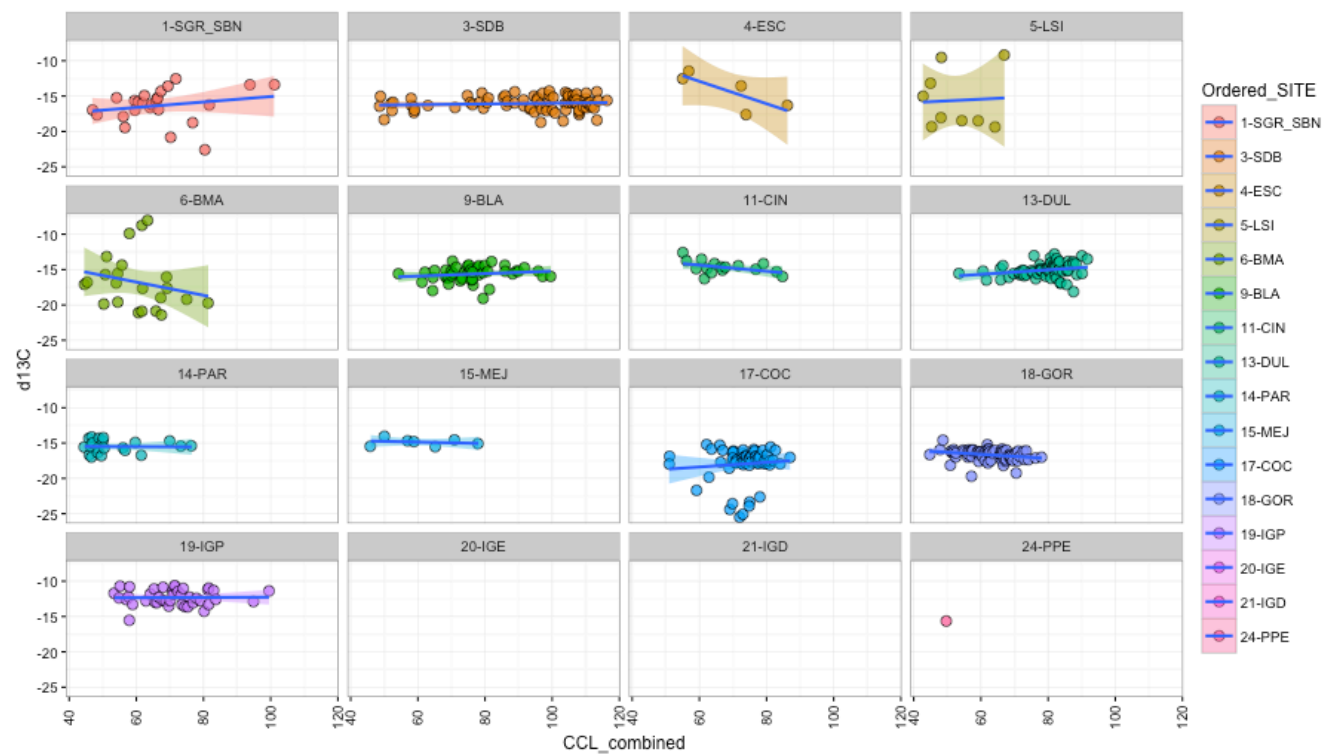
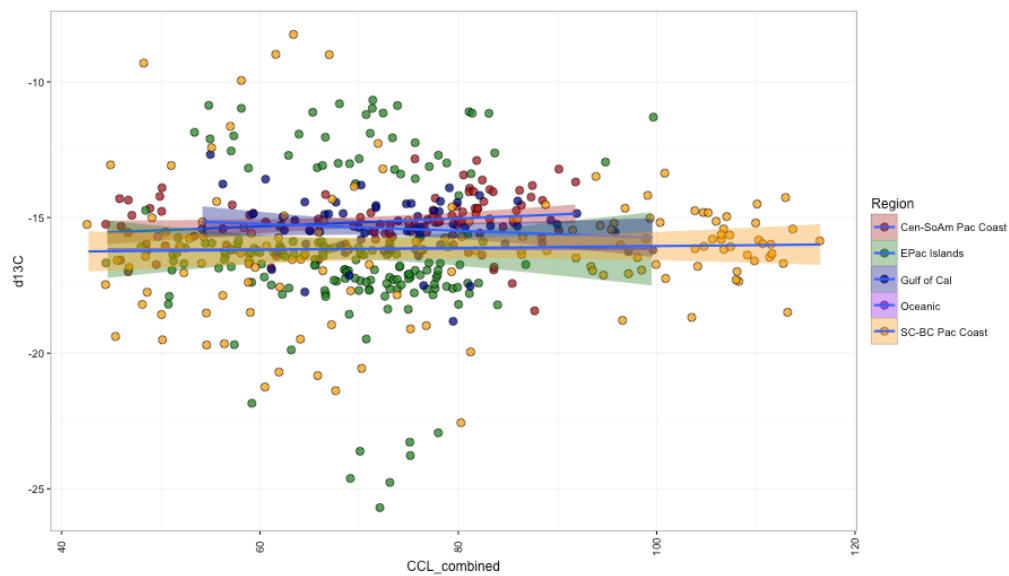




Turtle Size or Color









```
m1<-lm(data=data, d15N~CCL_combined*Ordered_SITE)
```

```
## CCL_combined:Ordered_SITE6-BMA -0.156964 0.039628 -3.961 8.64e-05 ***
```

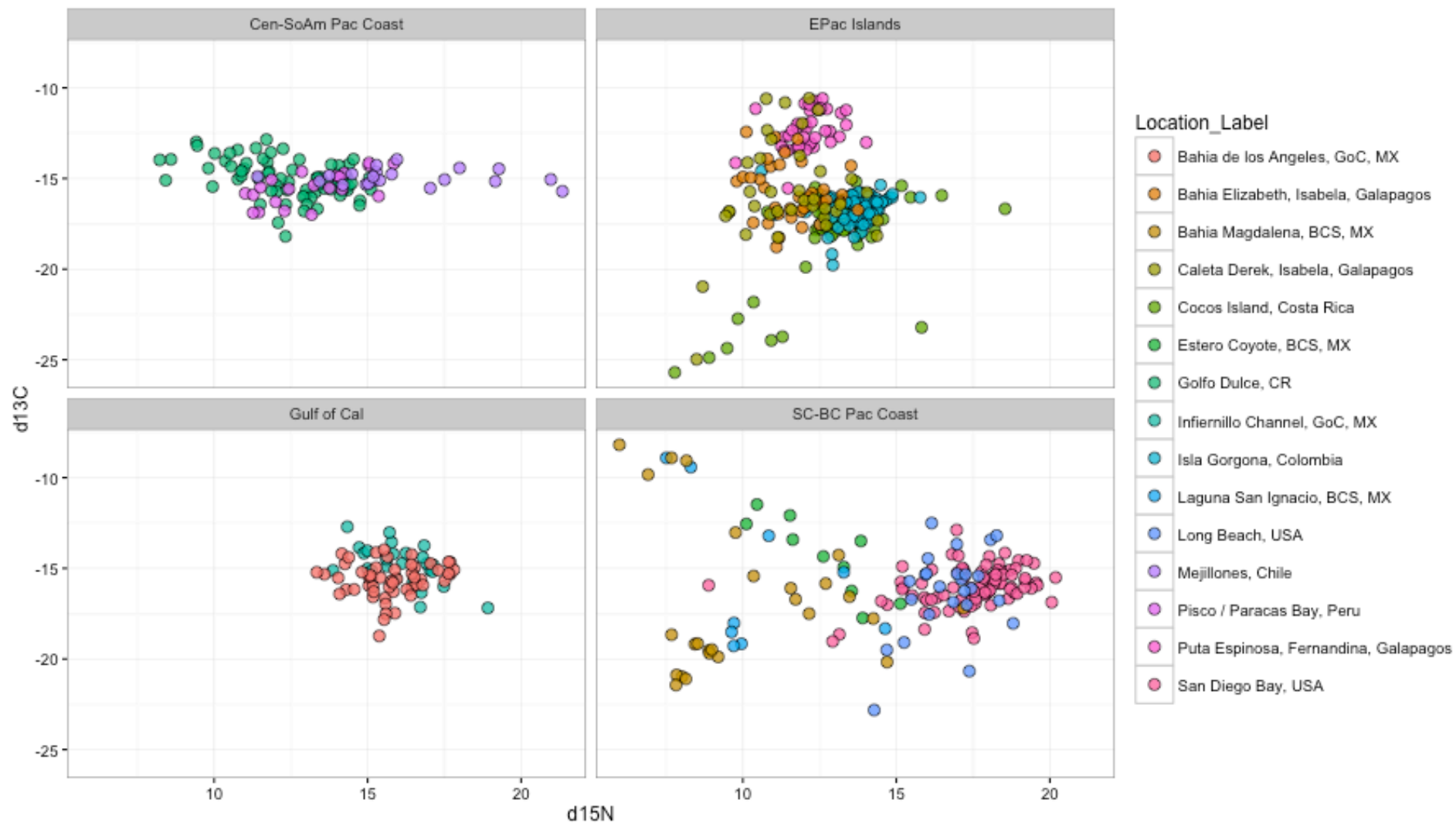
```
## CCL_combined:Ordered_SITE13-DUL -0.070234 0.032281 -2.176 0.03008 *
```

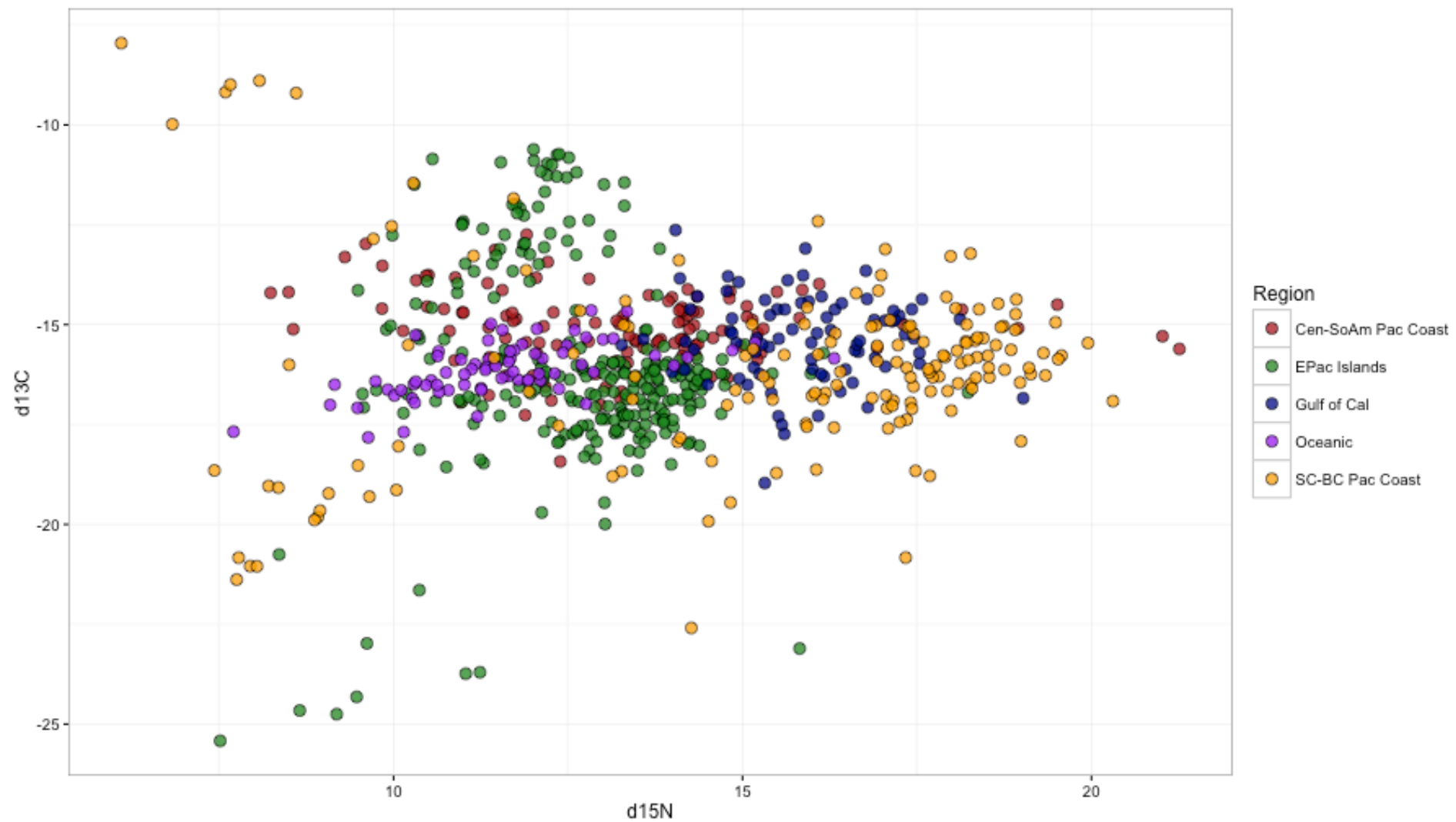
```
## Anova Table (Type II tests)
```

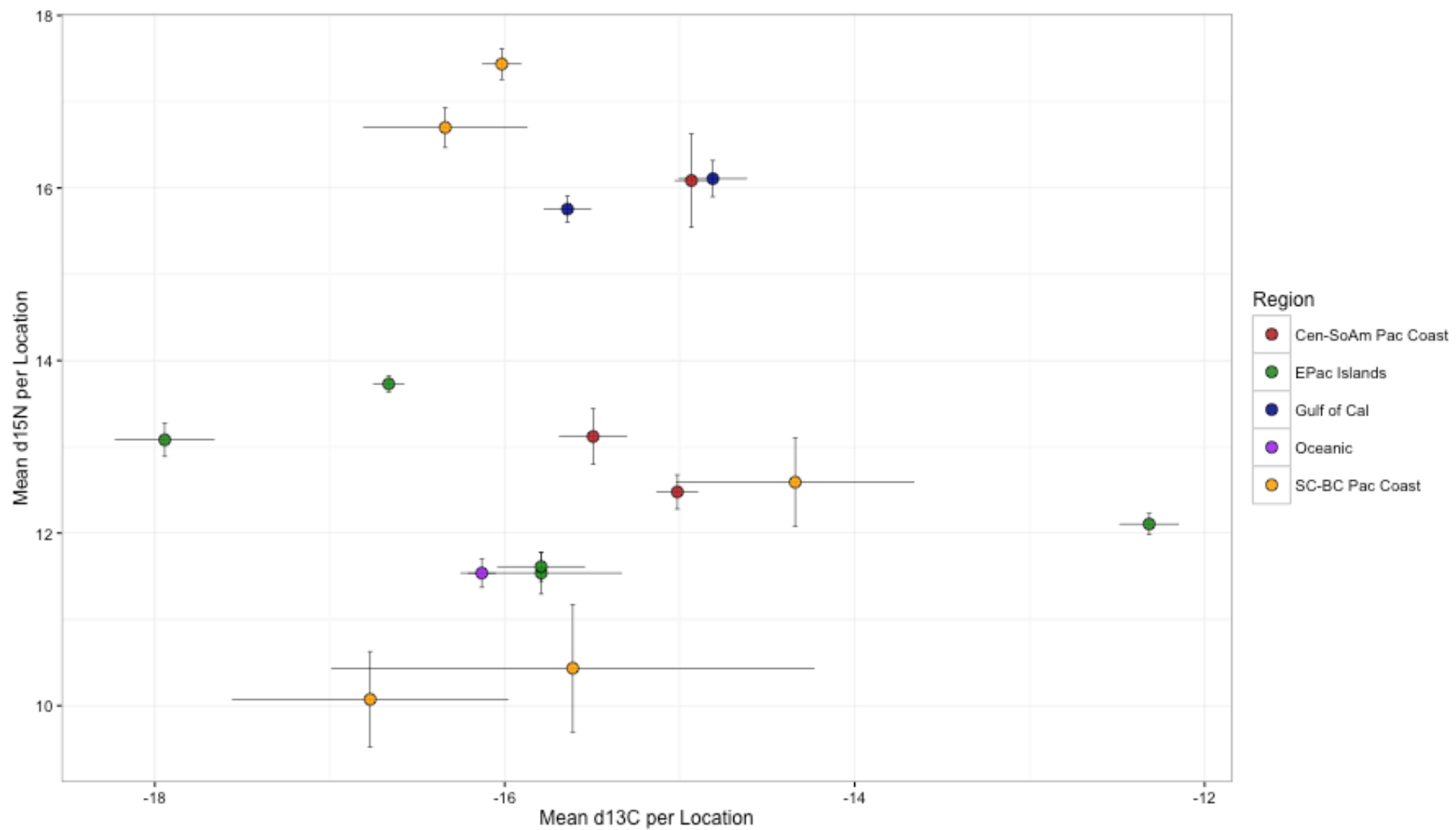
```
## Response: d15N
```

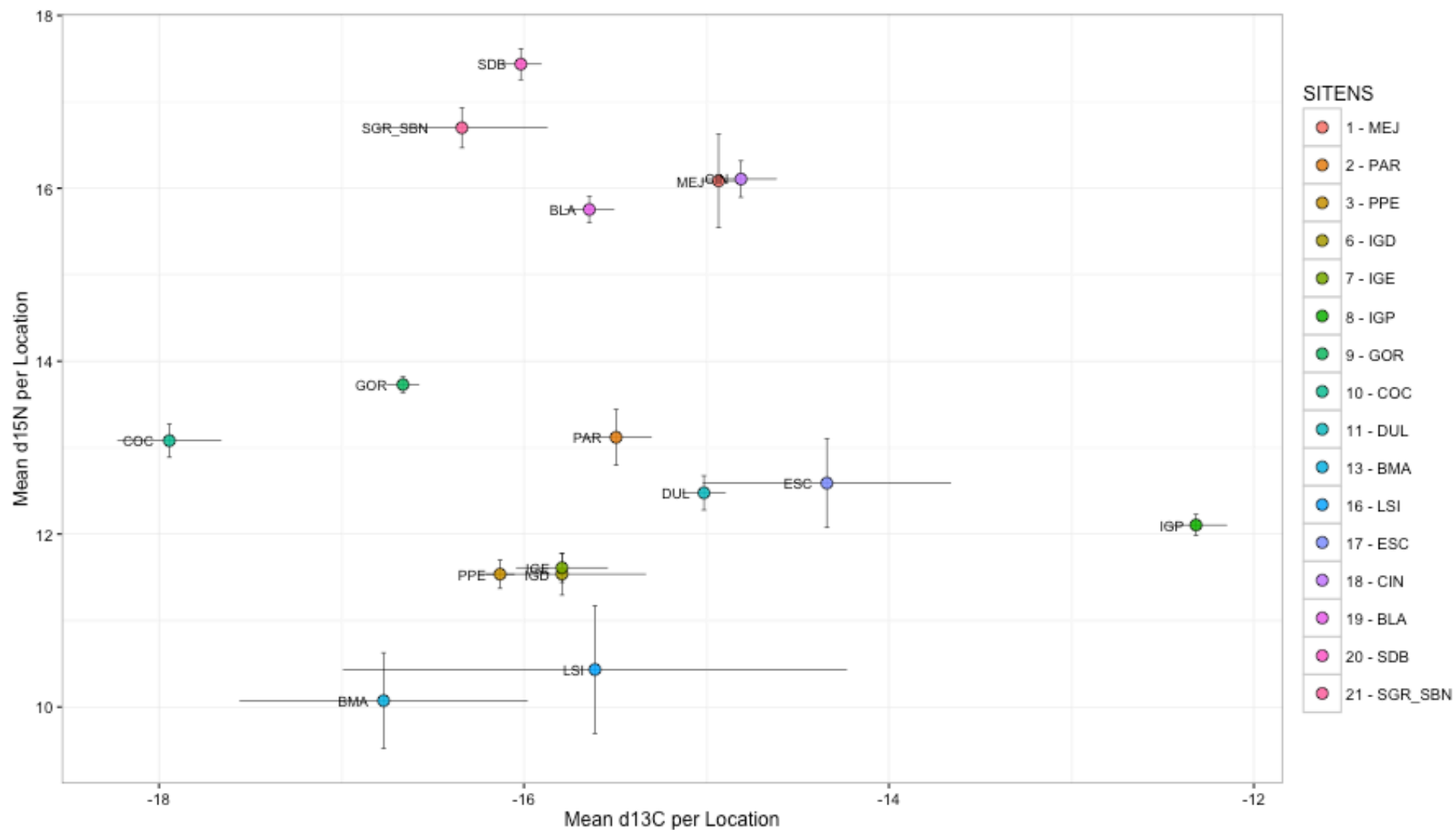
##	Sum Sq	Df	F value	Pr(>F)	
## CCL_combined	38.70	1	19.9084	1.021e-05	***
## Ordered_SITE	1647.81	13	65.2055	< 2.2e-16	***
## CCL_combined:Ordered_SITE	97.68	12	4.1874	2.868e-06	***
## Residuals	900.04	463			

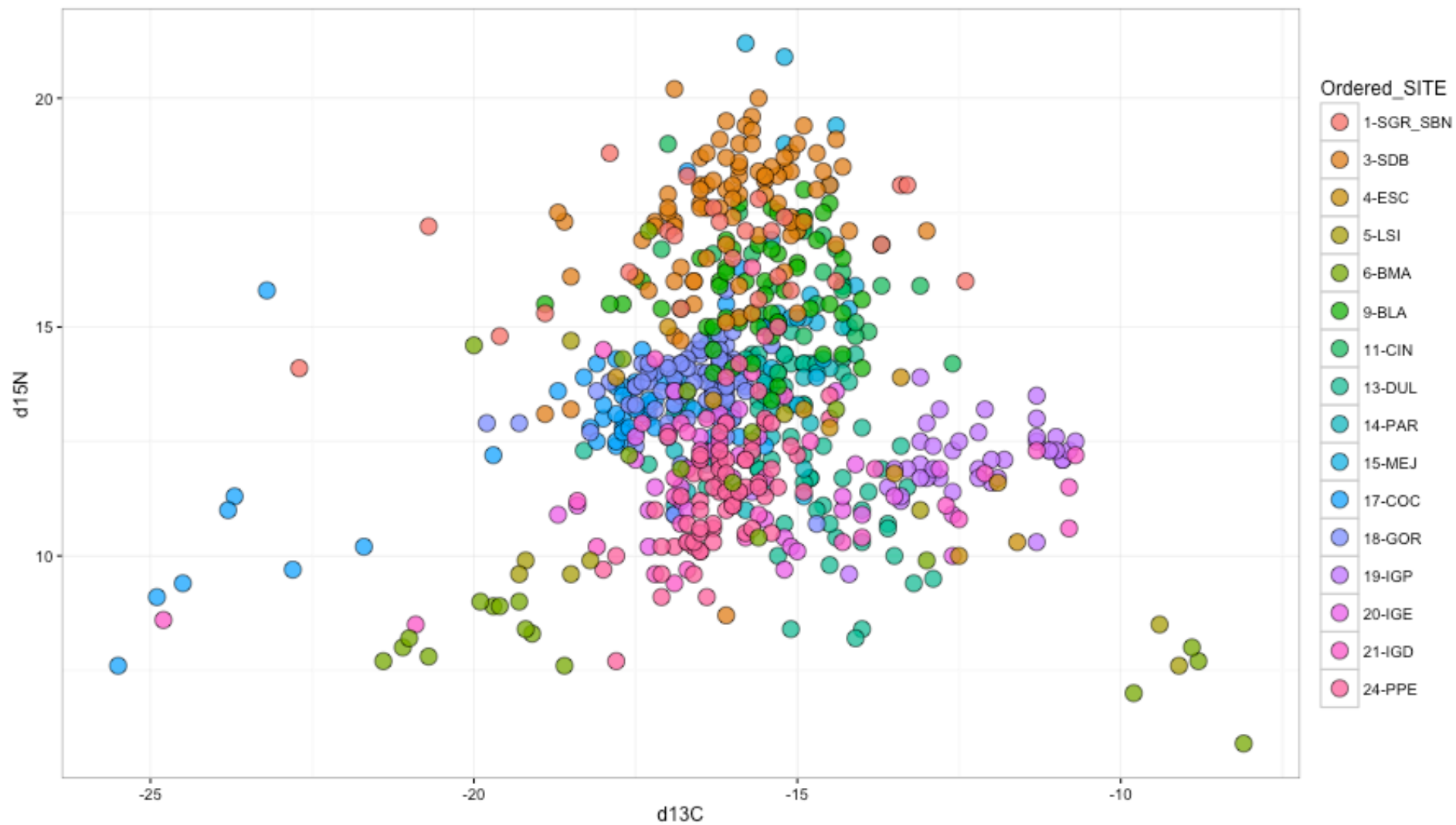
```
#remember to come back and remove point to see if makes difference in weight of sign. at BMA
```

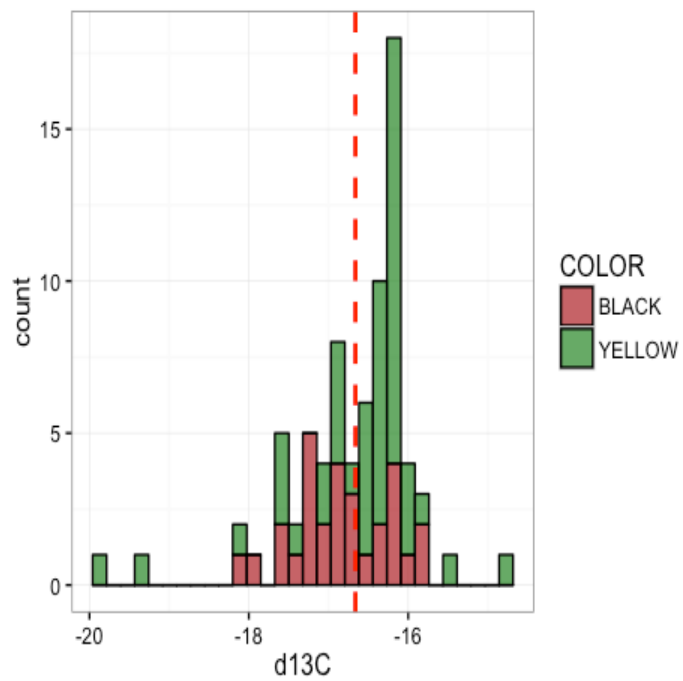
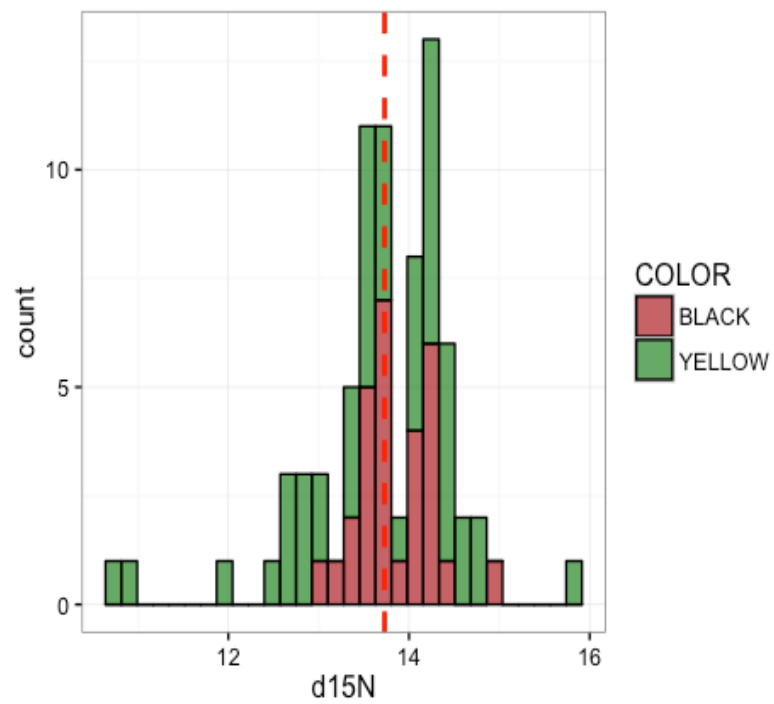
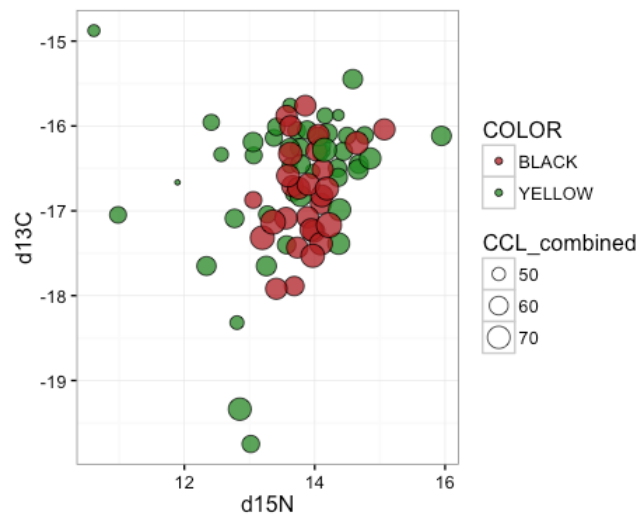








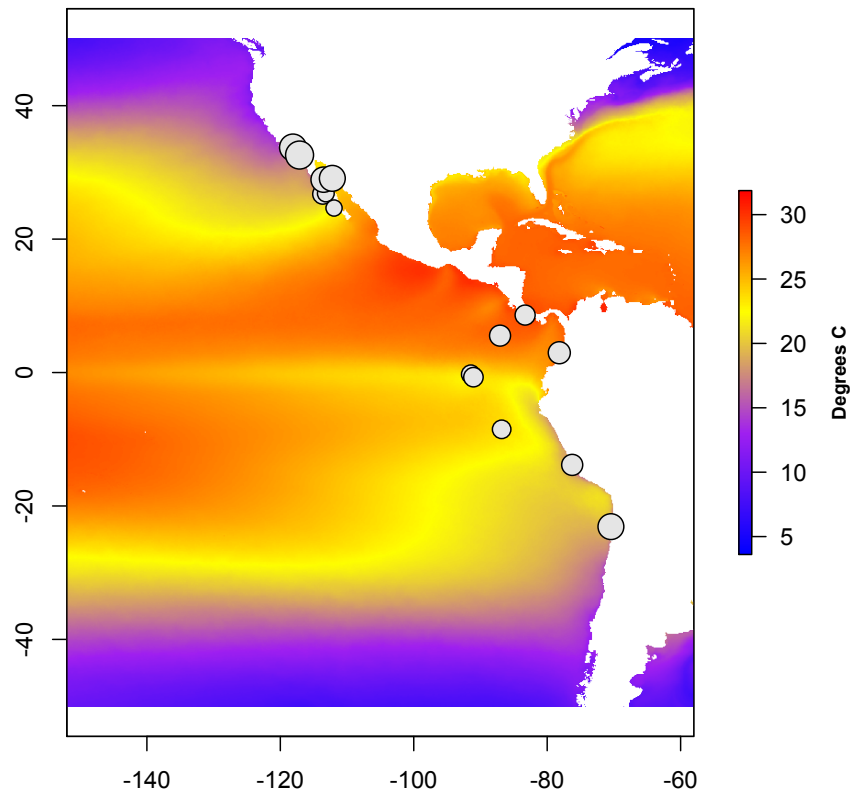








### East Pacific



### East Pacific

