

Install necessary libraries

Load Libraries

Read in data

Process and Join data

Visualize linear and nonlinear regressions: Richness and Volume

NOT INCLUDING VSEEP

Community vs SGD Regression

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Install necessary libraries

```
if("tidyverse" %in% rownames(installed.packages()) == FALSE){install.packages("tidyverse")}
if("here" %in% rownames(installed.packages()) == FALSE){install.packages("here")}
if("ggrepel" %in% rownames(installed.packages()) == FALSE){install.packages("ggrepel")}
if("PNWColors" %in% rownames(installed.packages()) == FALSE){install.packages("PNWColors")}
if("vegan" %in% rownames(installed.packages()) == FALSE){install.packages("vegan")}
if("pairwiseAdonis" %in% rownames(installed.packages()) == FALSE){ devtools::install_github("pmartinezarbizu/pairwiseAdonis/pairwiseAdonis")}
if("patchwork" %in% rownames(installed.packages()) == FALSE){install.packages("patchwork")}
if("ggmap" %in% rownames(installed.packages()) == FALSE){install.packages("ggmap")}
```

Load Libraries

```
library(tidyverse)
```

```
## — Attaching packages ————— tidyverse 1.3.2 —
## ✓ ggplot2 3.3.6      ✓ purrr   0.3.5
## ✓ tibble  3.1.8      ✓ dplyr   1.0.10
## ✓ tidyr   1.2.1      ✓ stringr 1.4.1
## ✓ readr   2.1.3      ✓ forcats 0.5.2
## — Conflicts ————— tidyverse_conflicts() —
## ✗ dplyr::filter() masks stats::filter()
## ✗ dplyr::lag()     masks stats::lag()
```

```
library(here)
```

```
## here() starts at C:/Users/Danielle Barnas/Documents/Repositories/Community_Functional_Diversity
```

```
library(ggrepel)  
library(PNWColors)  
library(vegan)
```

```
## Loading required package: permute  
## Loading required package: lattice  
## This is vegan 2.6-2
```

```
library(pairwiseAdonis)
```

```
## Loading required package: cluster
```

```
library(patchwork)  
library(ggmap)
```

```
## Google's Terms of Service: https://cloud.google.com/maps-platform/terms/.  
## Please cite ggmap if you use it! See citation("ggmap") for details.
```

```
library(wakefield)
```

```
##  
## Attaching package: 'wakefield'  
##  
## The following object is masked from 'package:patchwork':  
##  
##     area  
##  
## The following object is masked from 'package:dplyr':  
##  
##     id
```

Read in data

```
meta <- read_csv(here("Data", "Full_Metadata.csv"))

#comp <- read_csv(here("Data", "Surveys", "Species_Composition_2022.csv"))
taxa <- read_csv(here("Data", "Surveys", "Distinct_Taxa.csv"))
chem <- read_csv(here("Data", "Biogeochem", "Nutrients_Processed_All.csv"))
turb <- read_csv(here("Data", "Biogeochem", "July2022", "Turb_NC.csv"))
# richness, % richness of community pool, and % volume of community pool
Fric <- read_csv(here("Data", "Sp_FE_Vol.csv"))

# create color palette for plotting
mypalette <- pnw_palette(name="Bay")
midpalette <- pnw_palette(name = "Bay", n = 30)
largepalette <- pnw_palette(name = "Bay", n = 100)
```

Process and Join data

```
# reduce metadata
meta <- meta %>%
  drop_na(lat) %>% # removes Maya's project sites (not being used for my study)
  left_join(turb)
```

```
## Joining, by = c("Location", "CowTagID", "del15N", "C_N", "N_percent")
```

```
# join dfs
Full_data <- Fric %>%
  left_join(meta) %>%
  mutate(meanRugosity = if_else(CowTagID == "VSEEP", 0.97, meanRugosity)) %>% # i'm a
  ninny and still haven't added the seep rugosity yet *sigh*
  left_join(chem) %>%
  select(-Location)
```

```
## Joining, by = "CowTagID"
## Joining, by = c("CowTagID", "Location", "lat", "lon")
```

```
myturb <- turb %>%
  pivot_longer(cols = del15N:N_percent, names_to = "Parameters", values_to = "Values") %
  >%
  right_join(Fric) %>%
  select(-Location) %>%
  relocate(Parameters, .after = Vol8D) %>%
  relocate(Values, .after = Parameters) %>%
  mutate(Season = "June2022")
```

```
## Joining, by = "CowTagID"
```

Visualize linear and nonlinear regressions: Richness and Volume

```

# Plotting function
# create ggplot function
plotfun <- function(mydata = Full_data, x, y, myfacet, myformula) {

  x<-enquo(x)
  y<-enquo(y)

  plot <- ggplot(data = mydata,
                 aes(y = !!y,
                     x = !!x,
                     color = Season)) +
  geom_point() +
  geom_smooth(method = "lm", formula = myformula) +
  facet_wrap(enquo(myfacet), scales = "free") +
  labs(y = paste(as_label(y)),
       x = paste(as_label(x))) +
  theme_bw()+
  theme(legend.direction = "horizontal",
        legend.position = "top")

  return(plot)
}

# pvalue Functions: LM and Polynomial
pvalLM <- function(mydata = Full_data, myparam, myseason){

  Param_data <- mydata %>%
    select(CowTagID, NbSp:Vol8D, Season, Parameters, {{myparam}}) %>%
    pivot_wider(names_from = Parameters, values_from = {{myparam}})

  ParamType <- as.character(myparam)

  p_df <- tibble(Parameter = as.character(),
                 ParamType = as.character(),
                 Dependent = as.character(),
                 Season = as.character(),
                 pvalue = as.numeric(),
                 r_squared = as.numeric(),
                 adj_r_squared = as.numeric())

  for(i in 2:6){ # 2:6 are our dependent variables NbSp, NbFEs, Vol8D

    for(j in 8:ncol(Param_data)){

      Param_data <- Param_data %>%
        filter(Season == myseason)
      Season <- myseason

      Parameter <- colnames(Param_data)[j]
      Dependent <- colnames(Param_data)[i] # select dependent parameter
    }
  }
}

```

```

    if(is.na(Param_data[1,j])){ # some variables have no data in the wet season, so skip these variables
      p_df <- p_df
    } else {

      mod <- lm(paste(Dependent, "~", Parameter), data = Param_data)
      pvalue <- summary(mod)[4]$coefficients[8]
      r_squared <- summary(mod)[8]$r.squared
      adj_r_squared <- summary(mod)[9]$adj.r.squared

      temp <- as_tibble(cbind(Parameter, ParamType, Dependent, Season,
                             pvalue, r_squared, adj_r_squared)) %>%
        mutate(pvalue = as.numeric(pvalue),
               r_squared = as.numeric(r_squared),
               adj_r_squared = as.numeric(adj_r_squared))

      p_df <- p_df %>%
        rbind(temp)
    }
  }
}
return(p_df)
}

```

```

pvalpoly <- function(mydata = Full_data, myparam, myseason){

  Param_data <- mydata %>%
    select(CowTagID, NbSp:Vol8D, Season, Parameters, {{myparam}}) %>%
    pivot_wider(names_from = Parameters, values_from = {{myparam}})

  ParamType <- as.character(myparam)

  p_df <- tibble(Parameter = as.character(),
                 ParamType = as.character(),
                 Dependent = as.character(),
                 Season = as.character(),
                 pvalue1 = as.numeric(),
                 pvalue2 = as.numeric(),
                 r_squared = as.numeric(),
                 adj_r_squared = as.numeric())

  for(i in 2:6){ # 2:6 are our dependent variables NbSp, NbFEs, Vol8D

    for(j in 8:ncol(Param_data)){

      Param_data <- Param_data %>%
        filter(Season == myseason)
      Season <- myseason

      Parameter <- colnames(Param_data)[j]
    }
  }
}

```

```

    Dependent <- colnames(Param_data)[i] # select dependent parameter

    if(is.na(Param_data[1,j])){ # some variables have no data in the wet season, so skip these variables
      p_df <- p_df
    } else {
      mod <- lm(paste(Dependent, "~ poly(", Parameter, ",2)"), data = Param_data)
      pvalue1 <- summary(mod)[4]$coefficients[11]
      pvalue2 <- summary(mod)[4]$coefficients[12]
      r_squared <- summary(mod)[8]$r.squared
      adj_r_squared <- summary(mod)[9]$adj.r.squared

      temp <- as_tibble(cbind(Parameter, ParamType, Dependent, Season,
                             pvalue1, pvalue2, r_squared, adj_r_squared)) %>%
        mutate(pvalue1 = as.numeric(pvalue1),
               pvalue2 = as.numeric(pvalue2),
               r_squared = as.numeric(r_squared),
               adj_r_squared = as.numeric(adj_r_squared))

      p_df <- p_df %>%
        rbind(temp)
    }
  }
}
return(p_df)
}

# pvalue plots
pvalplot <- function(mydata, season = "Dry"){

  mydata <- mydata %>%
    pivot_longer(cols = c(pvalue1, pvalue2), names_to = "p1_p2", values_to = "pvalue")
  # pvalue1 is the same in Linear and Poly2

  season <- enquos(season)

  plot1 <- mydata %>%
    filter(Dependent == "NbFEs" | Dependent == "NbSp" | Dependent == "Vol8D") %>%
    filter(Season == {{season}}) %>%
    ggplot(aes(x = Parameter, y = pvalue)) +
    geom_col(aes(fill = p1_p2),
            position = "dodge") +
    theme_bw() +
    geom_hline(yintercept = 0.05, color = "black", size = 1) +
    theme(axis.text.x = element_text(angle = 90),
          legend.position = "top") +
    facet_wrap(~Dependent, scales = "free")

  return(plot1)
}

```

```

# site characteristic regression plots
paramPlot <- function(mydata = Full_data, ParamType, PTname, myformula = "y~x"){

  # isolate the desired paramtype (ex. Max, Min, Mean)
  data <- mydata %>%
    select(-c(NbSp:Vol8D,lat, lon))

  myparamdata <- data %>%
    select(CowTagID, Season, Parameters, Maximum)# {{ParamType}}

  data <- data[,1:13]
  data <- full_join(data, myparamdata)

  data <- data %>%
    pivot_wider(names_from = Parameters, values_from = Maximum) %>% # {{ParamType}} %>%
    relocate(Season, .after = CowTagID)

  p <- list() # empty list for saving plots

  for(i in 3:ncol(data)){
    # isolate dependent variable
    myDep <- data %>%
      select(CowTagID, Season) %>% # will join back after pivoting remaining variables
      cbind(data[,i]) %>%
      drop_na()

    newdata <- data %>%
      select(-c(colnames(data[,i]))) %>% # remove dependent var to not pivot on itself
      pivot_longer(cols = 3:ncol(.), names_to = "Parameters", values_to = "Values") %>%
      right_join(myDep) # bring back dependent var

    yplot <- as.name(colnames(newdata[,5]))

    p[[i]] <- newdata %>%
      ggplot(aes(x = Values, y = !!yplot, color = Season)) +
      geom_point() +
      geom_smooth(method = "lm", formula = myformula) +
      theme_bw() +
      theme(legend.position = "top",
            legend.direction = "horizontal") +
      facet_wrap(~Parameters, scales = "free")
  }

  # Save all plots in a single pdf
  pdf(here("Output","Biogeochem_Regression",paste0("V_site_regressions_", PTname ,".pdf")), onefile = TRUE)
  for (i in 1:length(p)) {
    tplot <- p[[i]]
    print(tplot)
  }
  dev.off()
}

```



```
return(p)
}
```

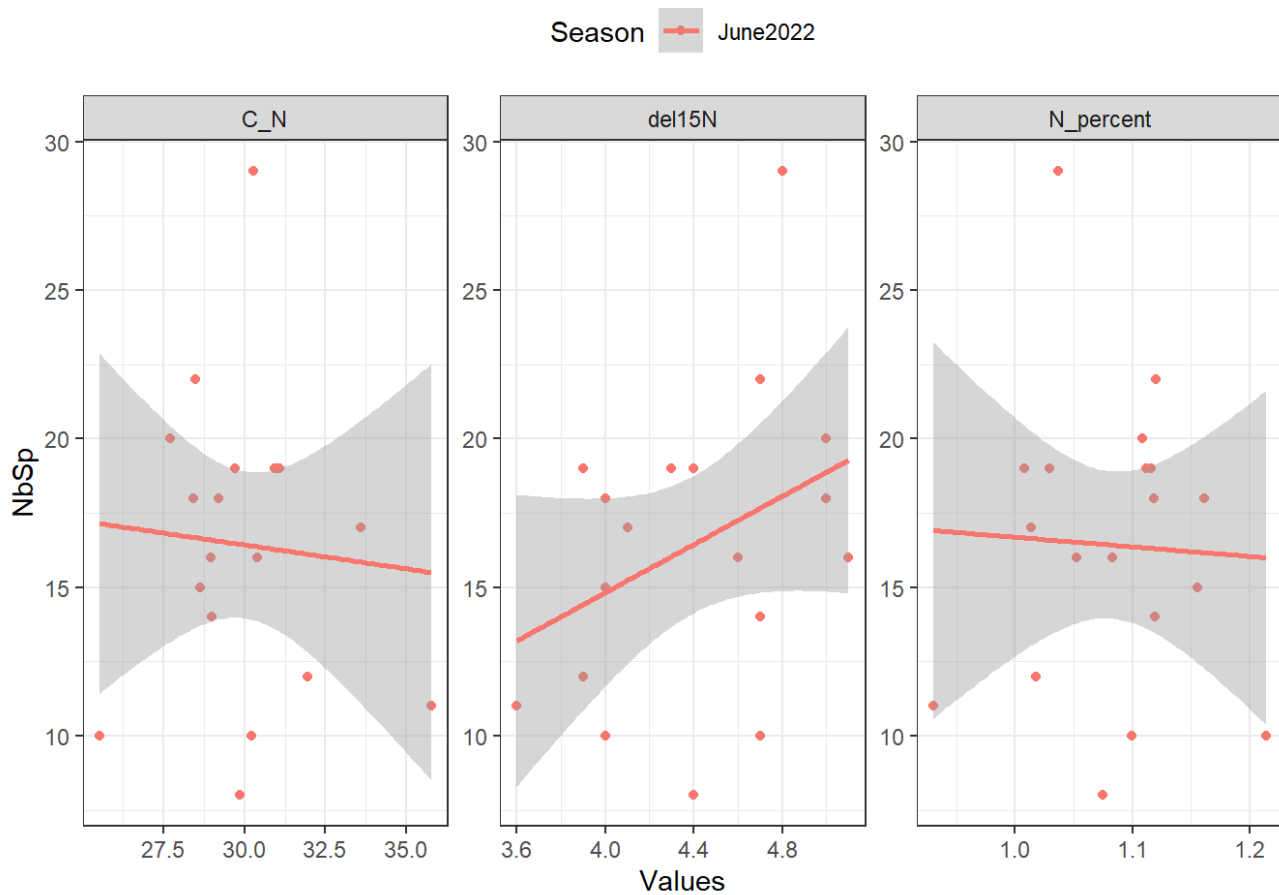
NOT INCLUDING VSEEP

Remove VSEEP from Full_data

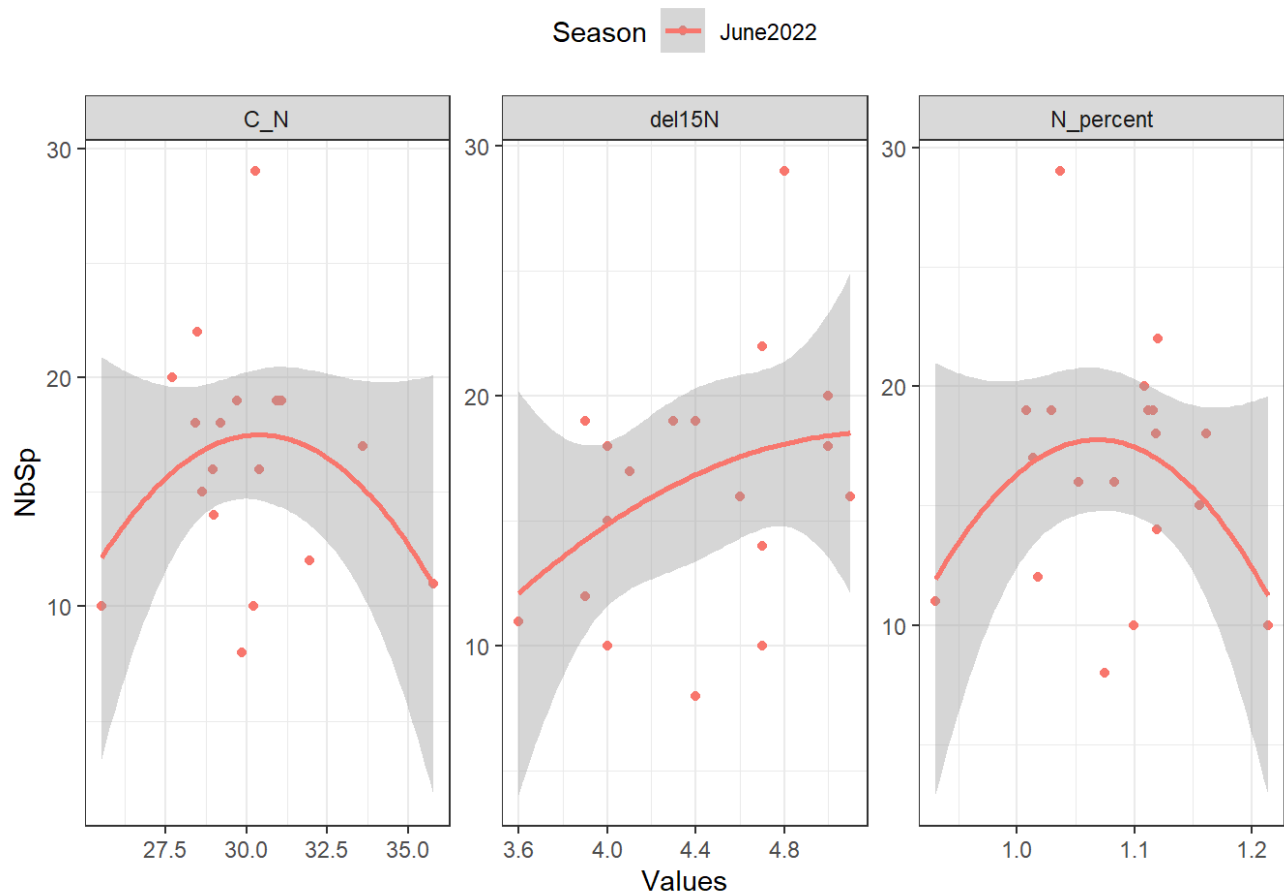
```
Full_data <- Full_data %>%
  filter(CowTagID!= "VSEEP")
myturb <- myturb %>%
  filter(CowTagID != "VSEEP")
```

TURBINARIA NUTRIENT LOADING

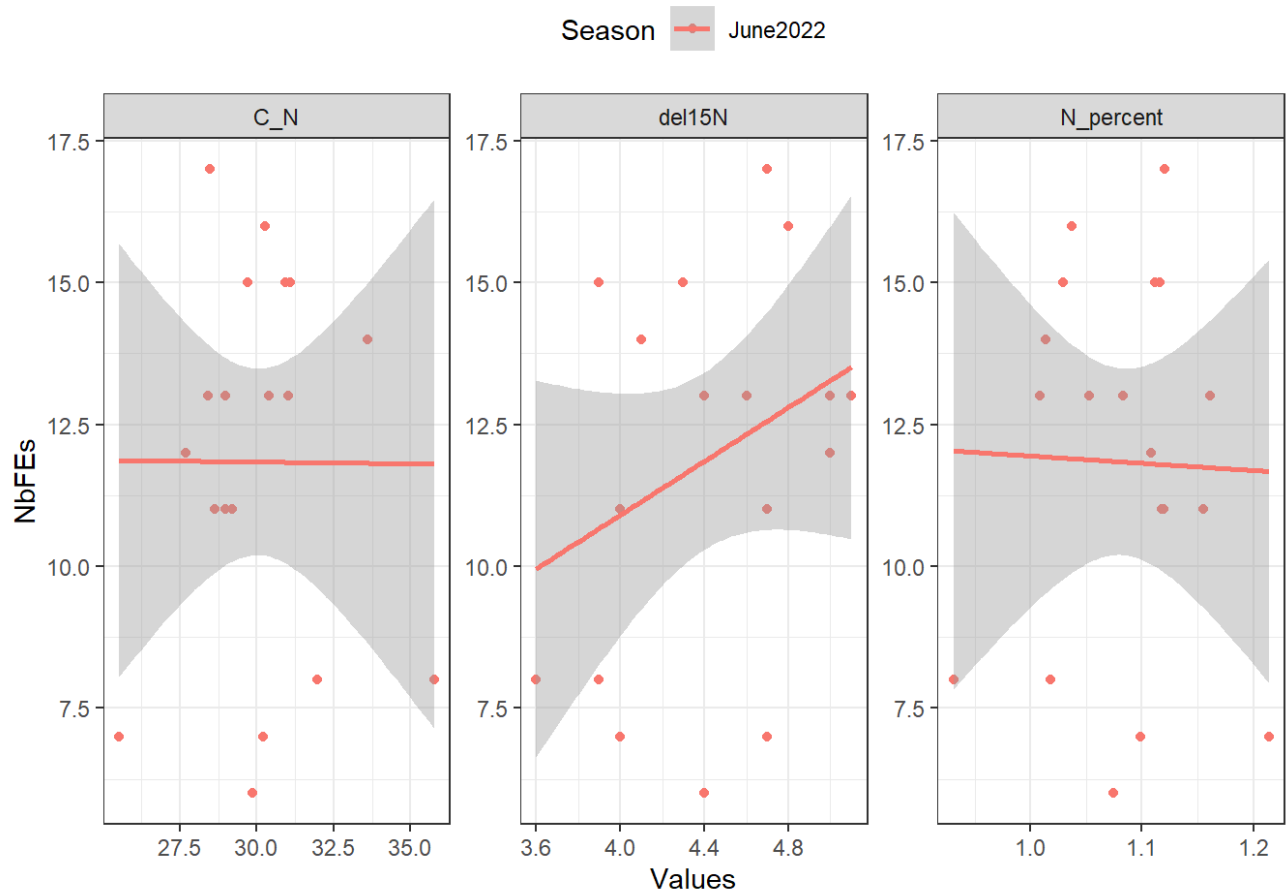
```
plotfun(y = NbSp, x = Values, myfacet = Parameters, myformula = "y~x", mydata = myturb)
```



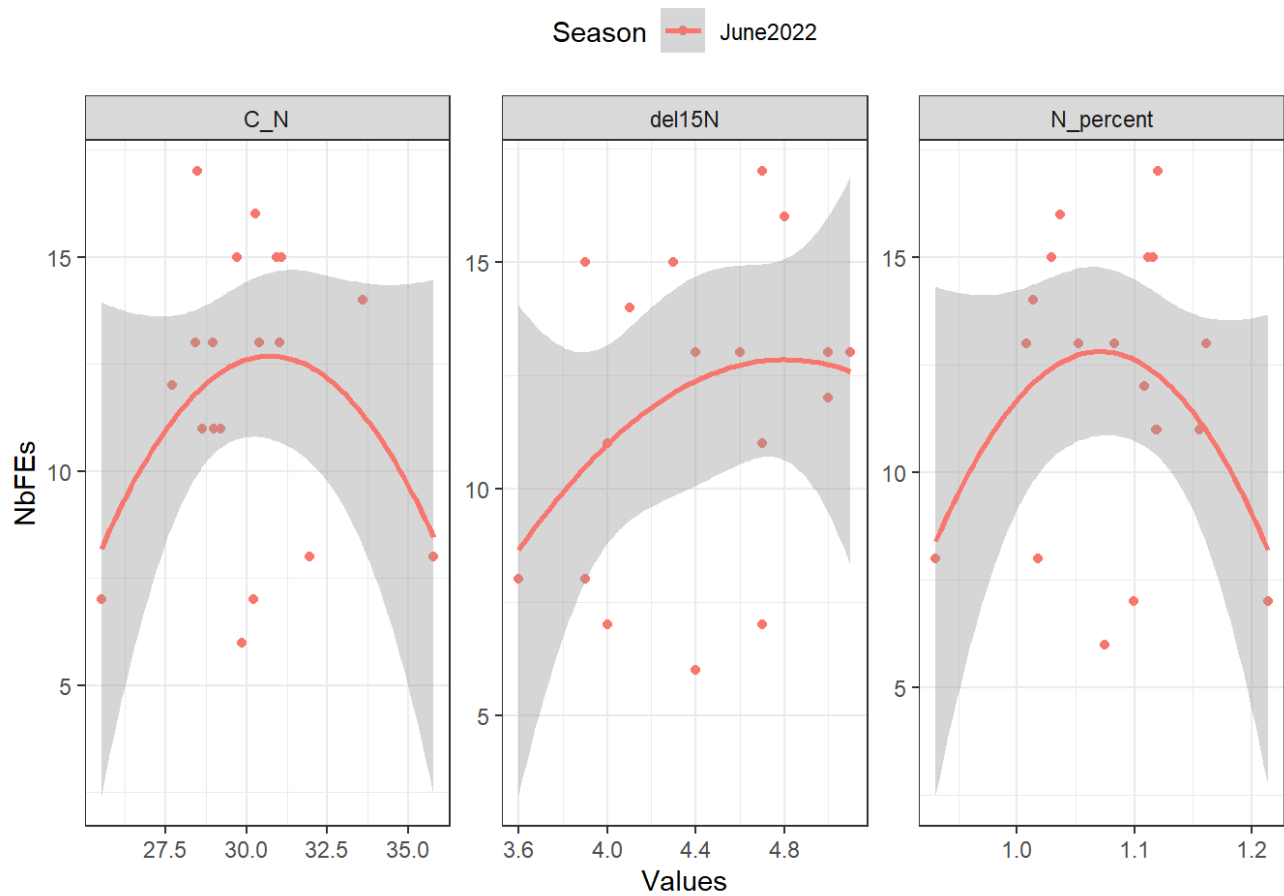
```
plotfun(y = NbSp, x = Values, myfacet = Parameters, myformula = "y~poly(x,2)", mydata =
myturb)
```



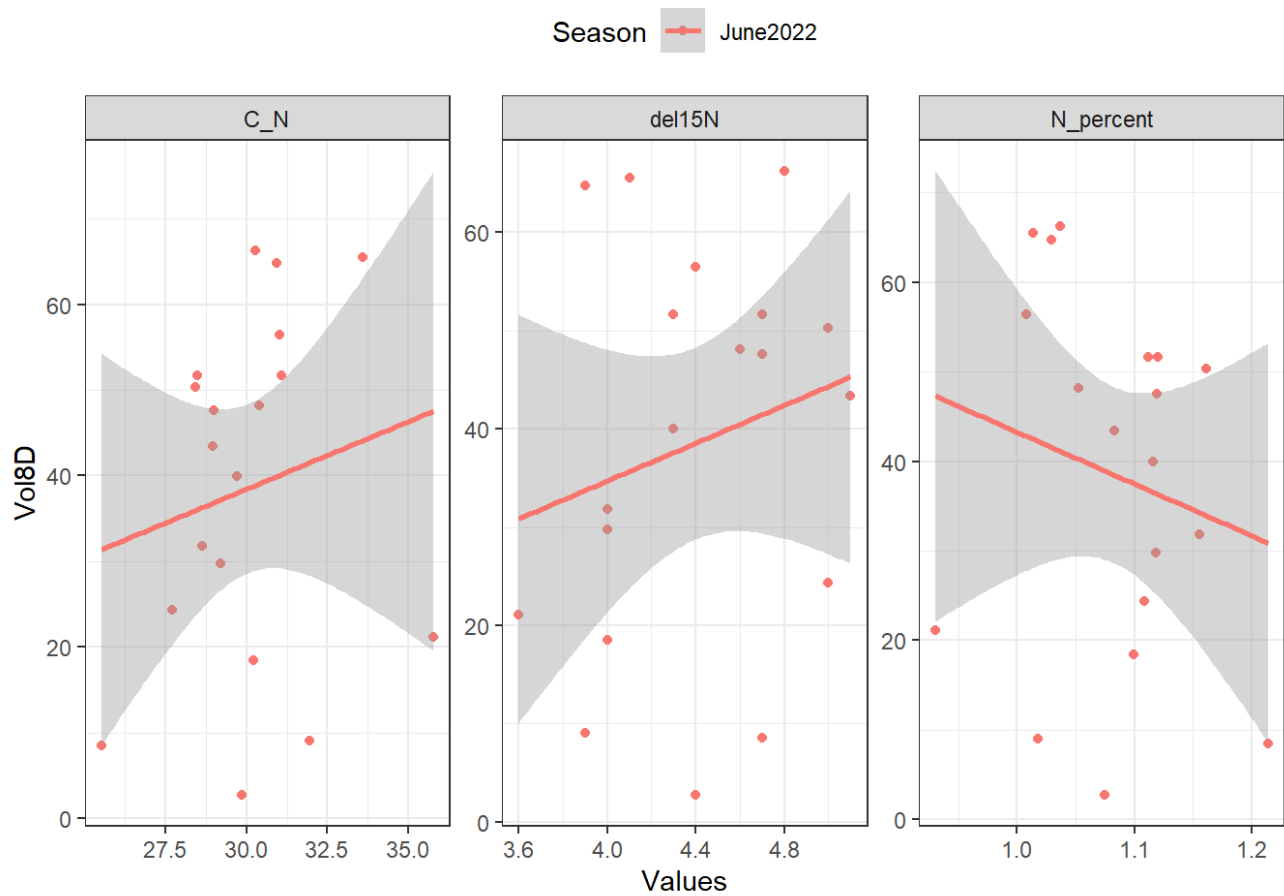
```
plotfun(y = NbFEs, x = Values, myfacet = Parameters, myformula = "y~x", mydata = myturb)
```



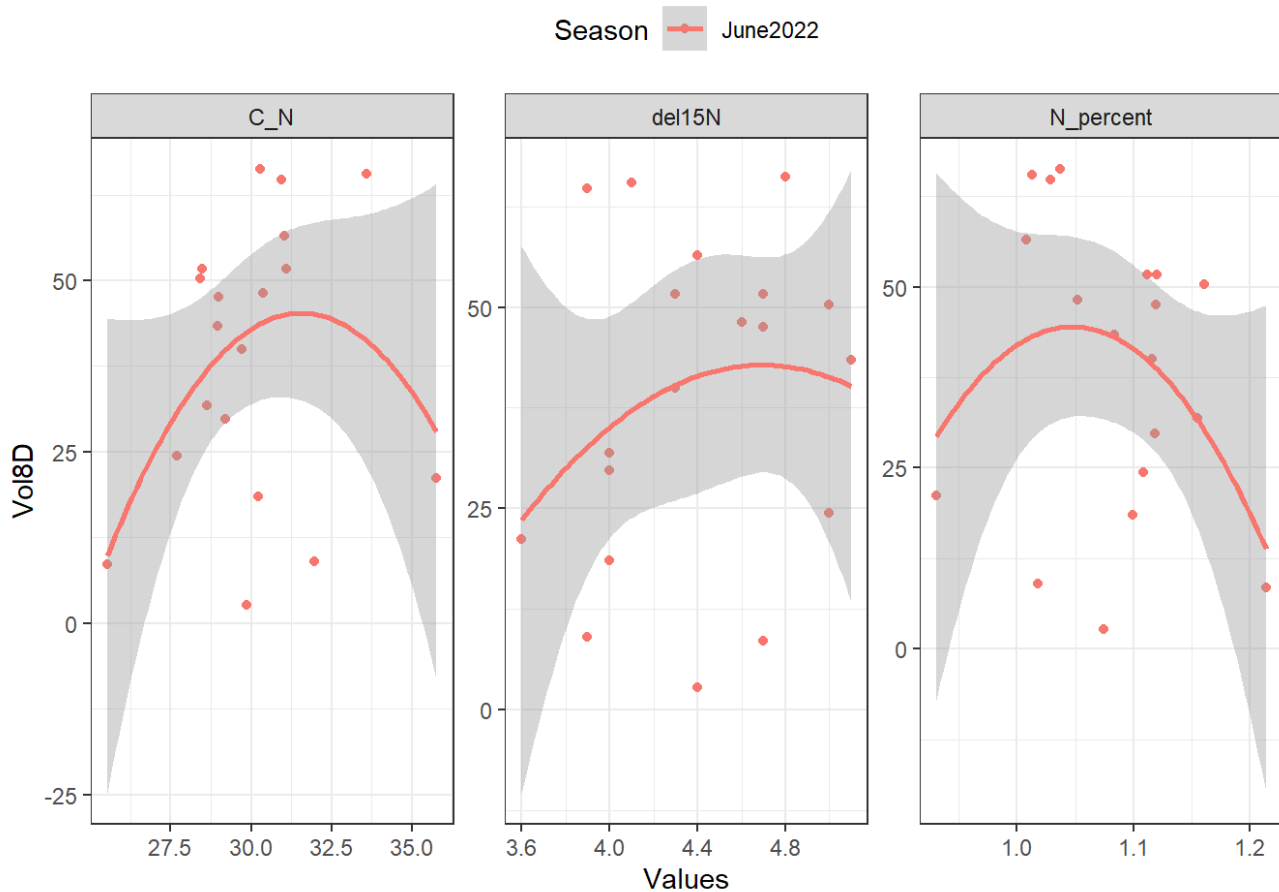
```
plotfun(y = NbFEs, x = Values, myfacet = Parameters, myformula = "y~poly(x,2)", mydata = myturb)
```



```
plotfun(y = Vol8D, x = Values, myfacet = Parameters, myformula = "y~x", mydata = myturb)
```



```
plotfun(y = Vol8D, x = Values, myfacet = Parameters, myformula = "y~poly(x,2)", mydata = myturb)
```



```
# get pvalues
```

```
pTO<-pvalpoly(mydata = myturb, myparam = "Values", myseason = "June2022")
```

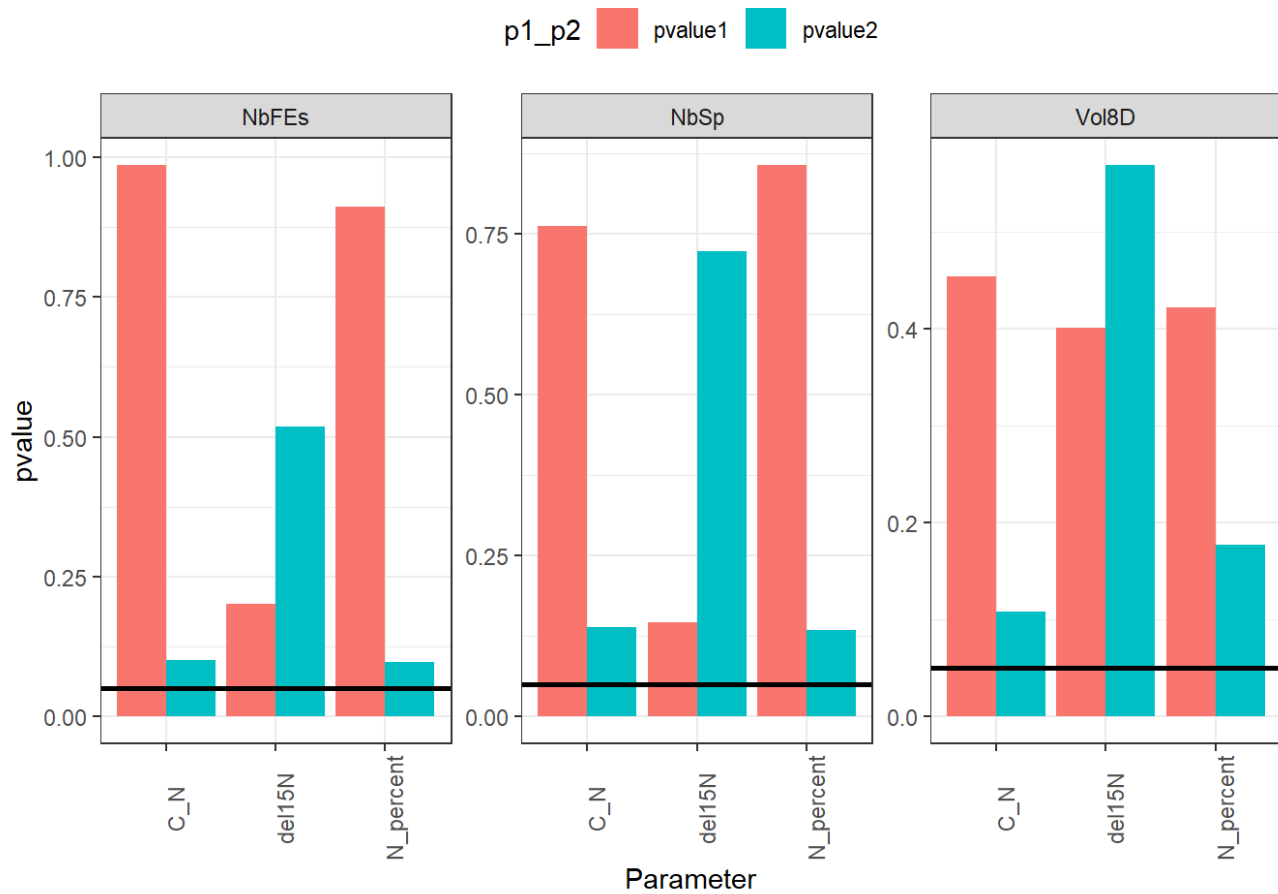
```
pTO
```

```
## # A tibble: 15 × 8
```

##	Parameter	ParamType	Dependent	Season	pvalue1	pvalue2	r_squared	adj_r_squa... ¹
##	<chr>	<chr>	<chr>	<chr>	<dbl>	<dbl>	<dbl>	<dbl>
## 1	del15N	Values	NbSp	June2022	0.146	0.722	0.133	0.0252
## 2	C_N	Values	NbSp	June2022	0.762	0.138	0.137	0.0286
## 3	N_percent	Values	NbSp	June2022	0.856	0.134	0.137	0.0286
## 4	del15N	Values	NbSpP	June2022	0.146	0.722	0.133	0.0252
## 5	C_N	Values	NbSpP	June2022	0.762	0.138	0.137	0.0286
## 6	N_percent	Values	NbSpP	June2022	0.856	0.134	0.137	0.0286
## 7	del15N	Values	NbFEs	June2022	0.201	0.518	0.122	0.0118
## 8	C_N	Values	NbFEs	June2022	0.986	0.100	0.160	0.0549
## 9	N_percent	Values	NbFEs	June2022	0.912	0.0968	0.163	0.0588
## 10	del15N	Values	NbFEsP	June2022	0.201	0.518	0.122	0.0118
## 11	C_N	Values	NbFEsP	June2022	0.986	0.100	0.160	0.0549
## 12	N_percent	Values	NbFEsP	June2022	0.912	0.0968	0.163	0.0588
## 13	del15N	Values	Vol8D	June2022	0.401	0.569	0.0633	-0.0538
## 14	C_N	Values	Vol8D	June2022	0.454	0.108	0.179	0.0761
## 15	N_percent	Values	Vol8D	June2022	0.422	0.177	0.143	0.0362

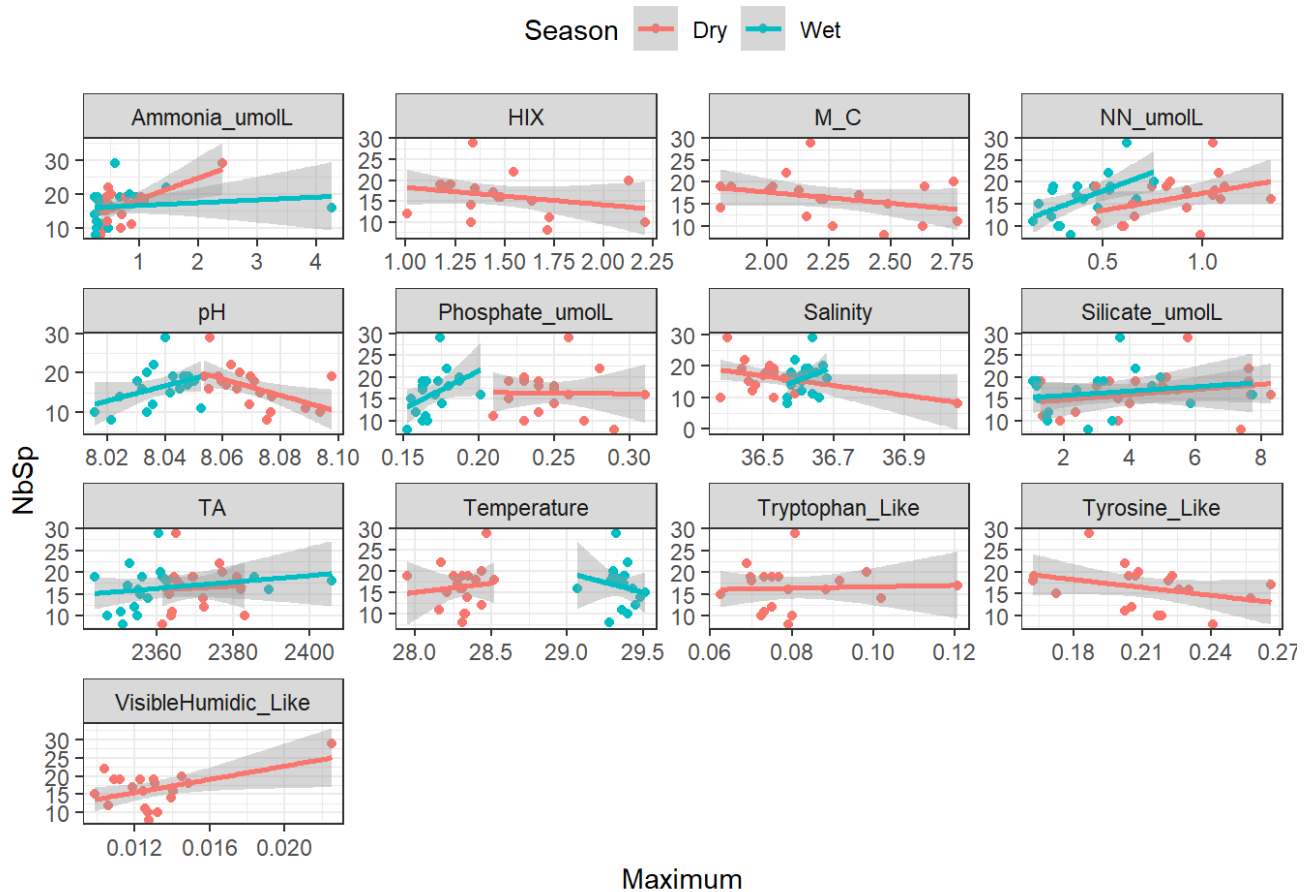
... with abbreviated variable name ¹adj_r_squared

```
pvalplot(mydata = pT0, season = "June2022")
```

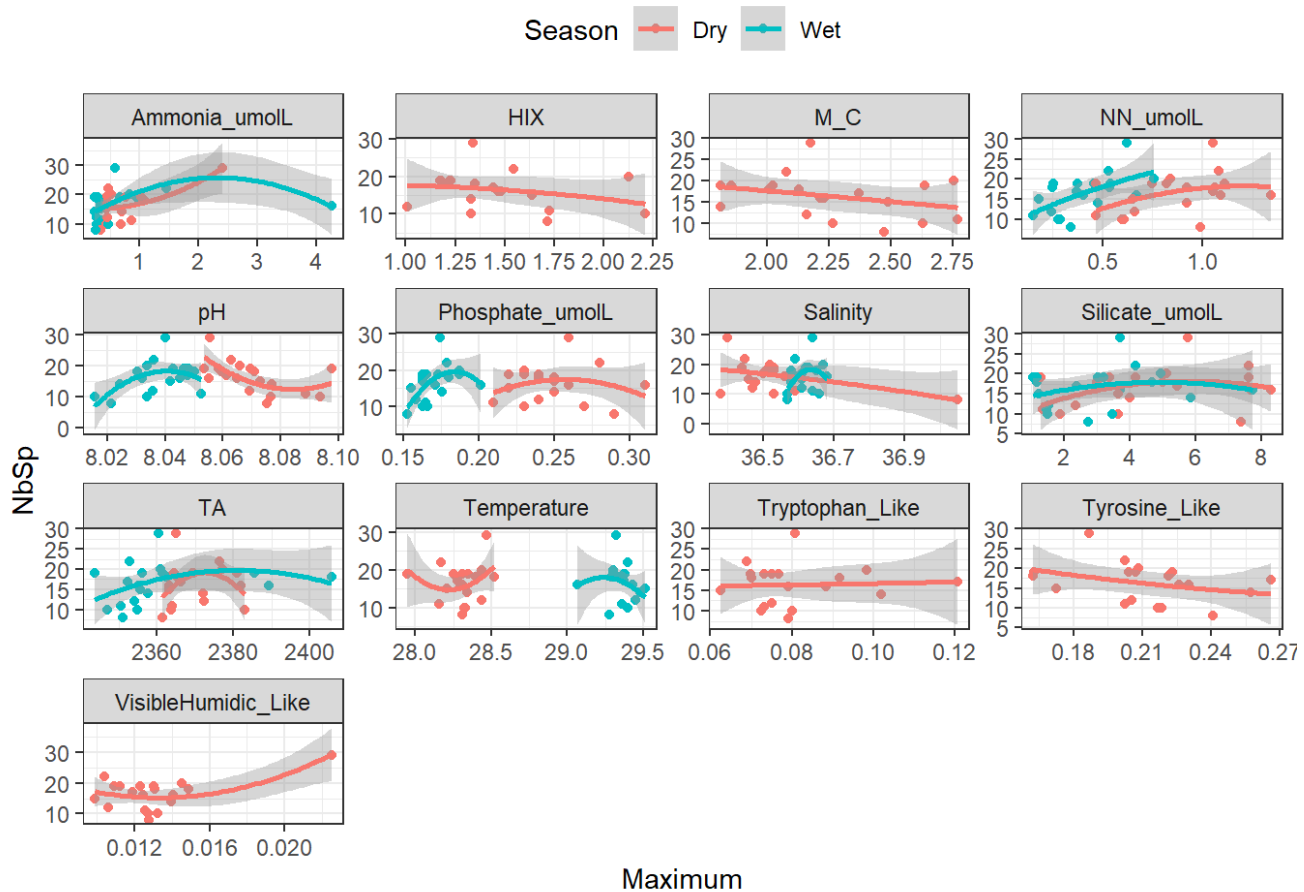


MAXIMUM VALUES

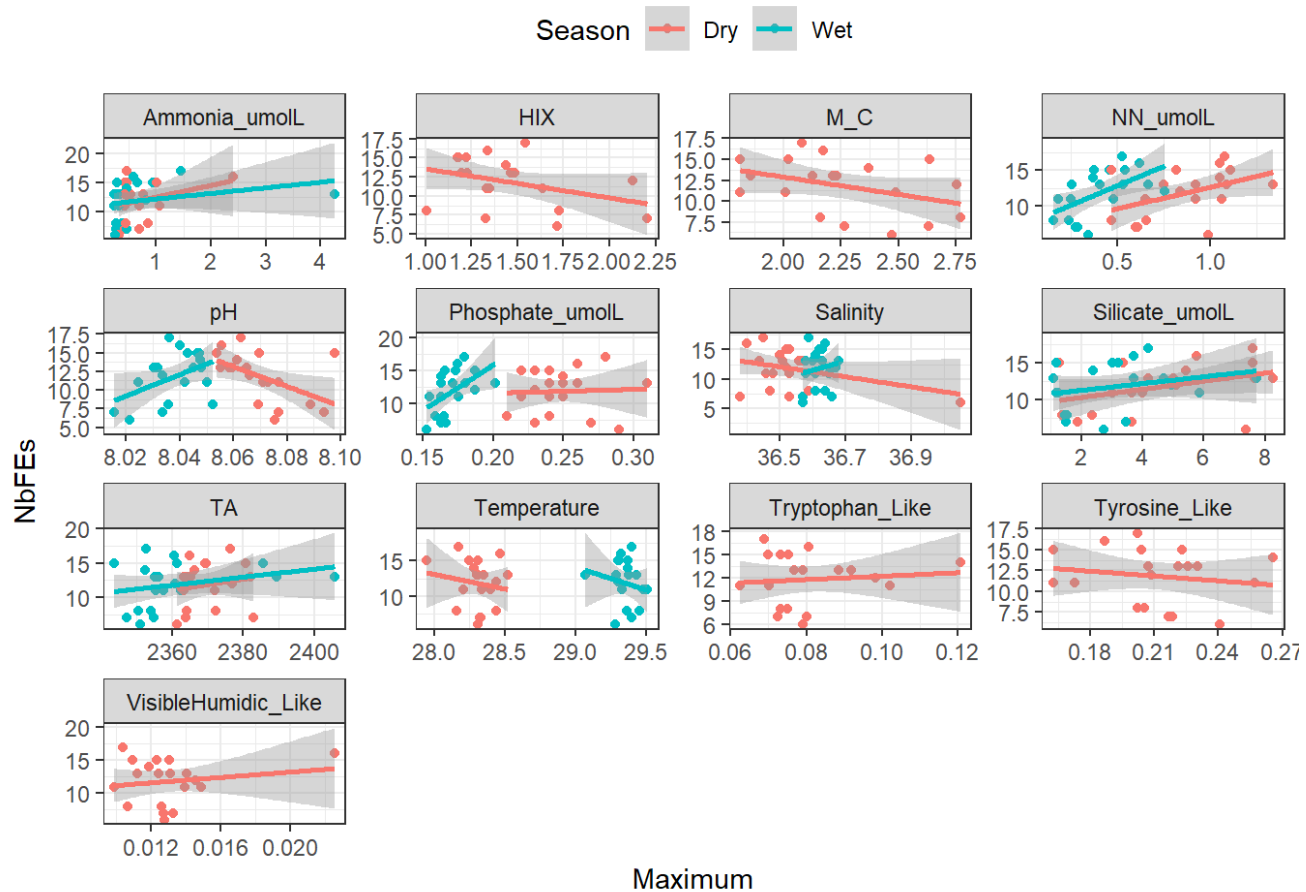
```
plotfun(y = NbSp, x = Maximum, myfacet = Parameters, myformula = "y~x")
```



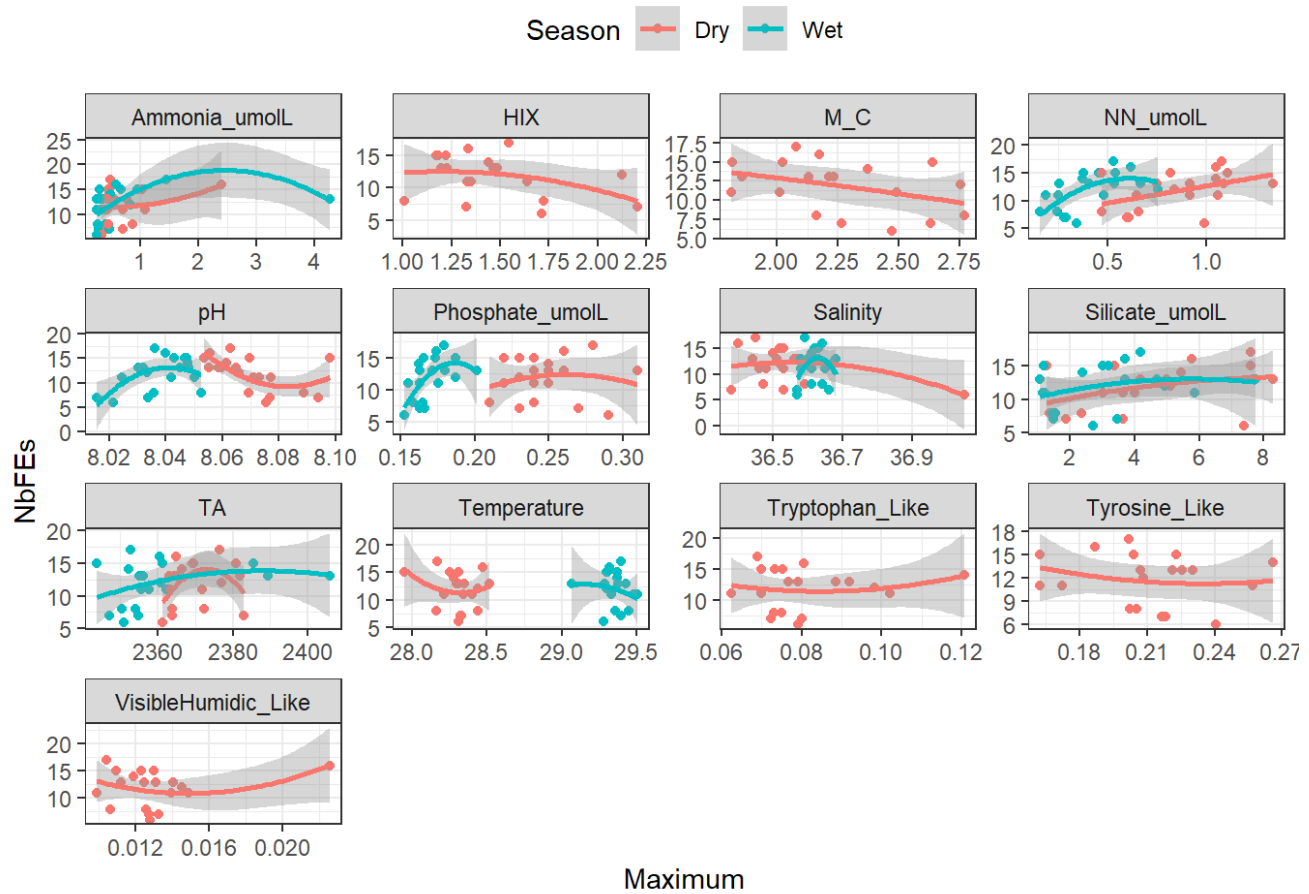
```
plotfun(y = NbSp, x = Maximum, myfacet = Parameters, myformula = "y~poly(x,2)")
```

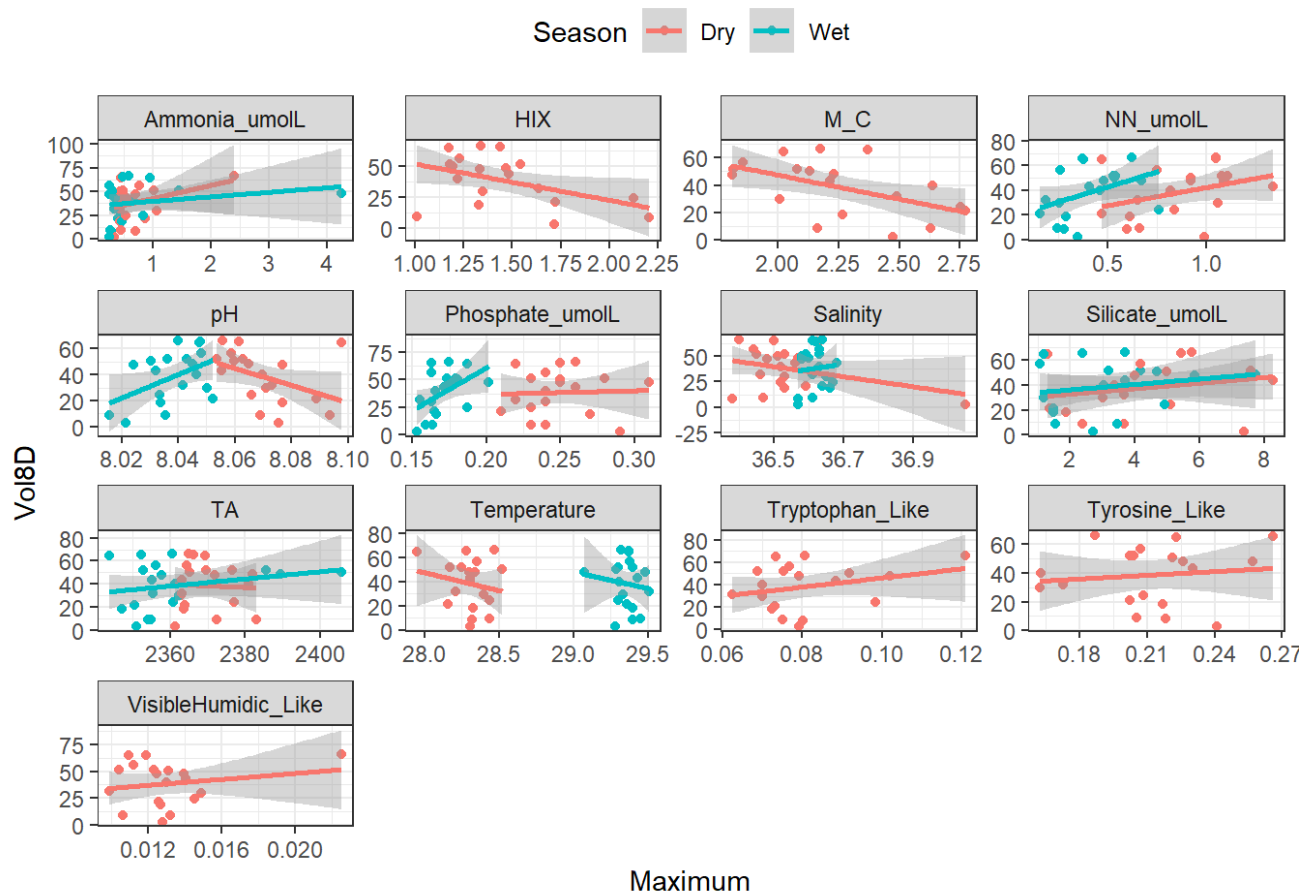
```
plotfun(y = NbFEs, x = Maximum, myfacet = Parameters, myformula = "y~x")
```



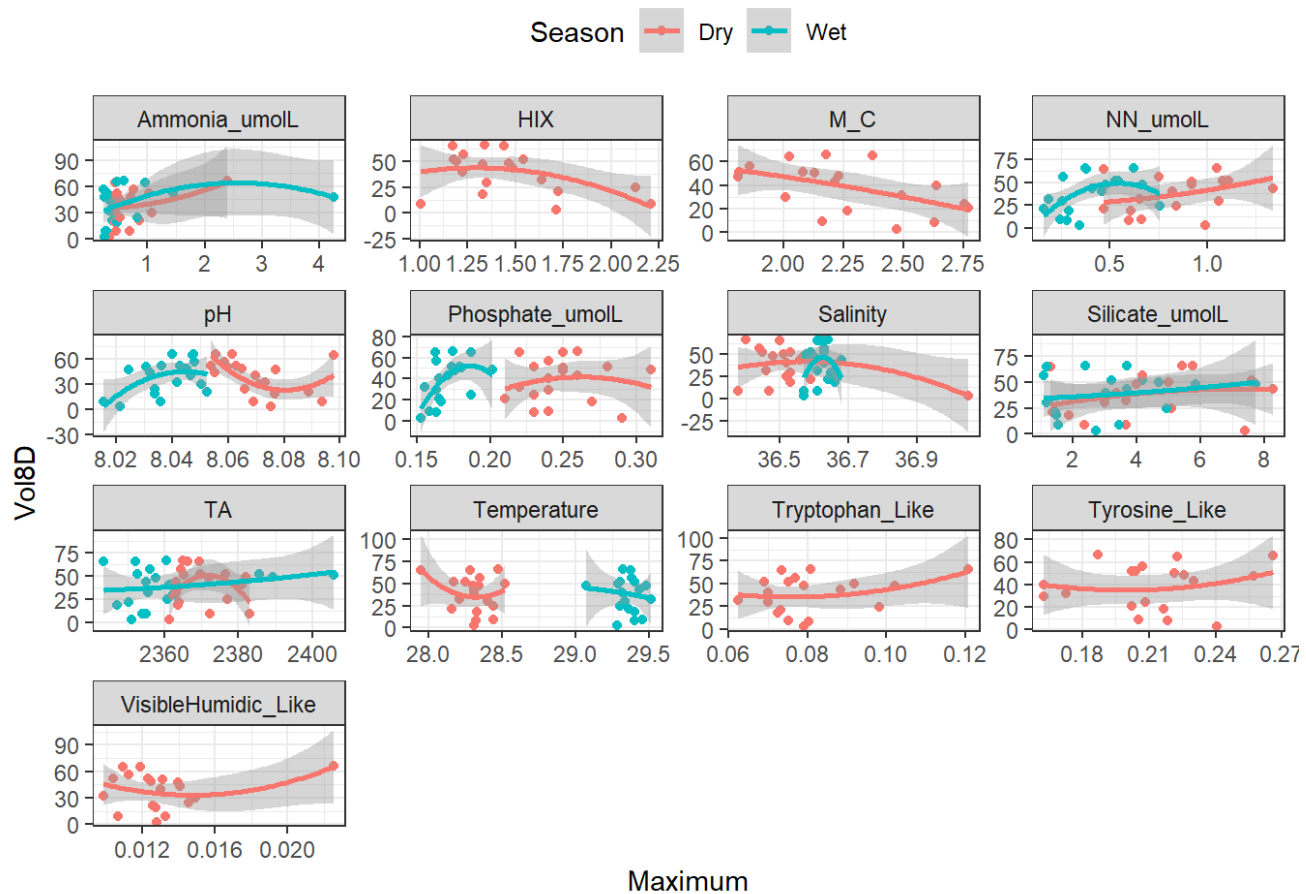
```
plotfun(y = NbFEs, x = Maximum, myfacet = Parameters, myformula = "y~poly(x,2)")
```



```
plotfun(y = Vol8D, x = Maximum, myfacet = Parameters, myformula = "y~x")
```



```
plotfun(y = Vol8D, x = Maximum, myfacet = Parameters, myformula = "y~poly(x,2)")
```



```
# get pvalues
```

```
Max3<-pvalpoly(myparam = "Maximum", myseason = "Dry")
```

```
Max4<-pvalpoly(myparam = "Maximum", myseason = "Wet")
```

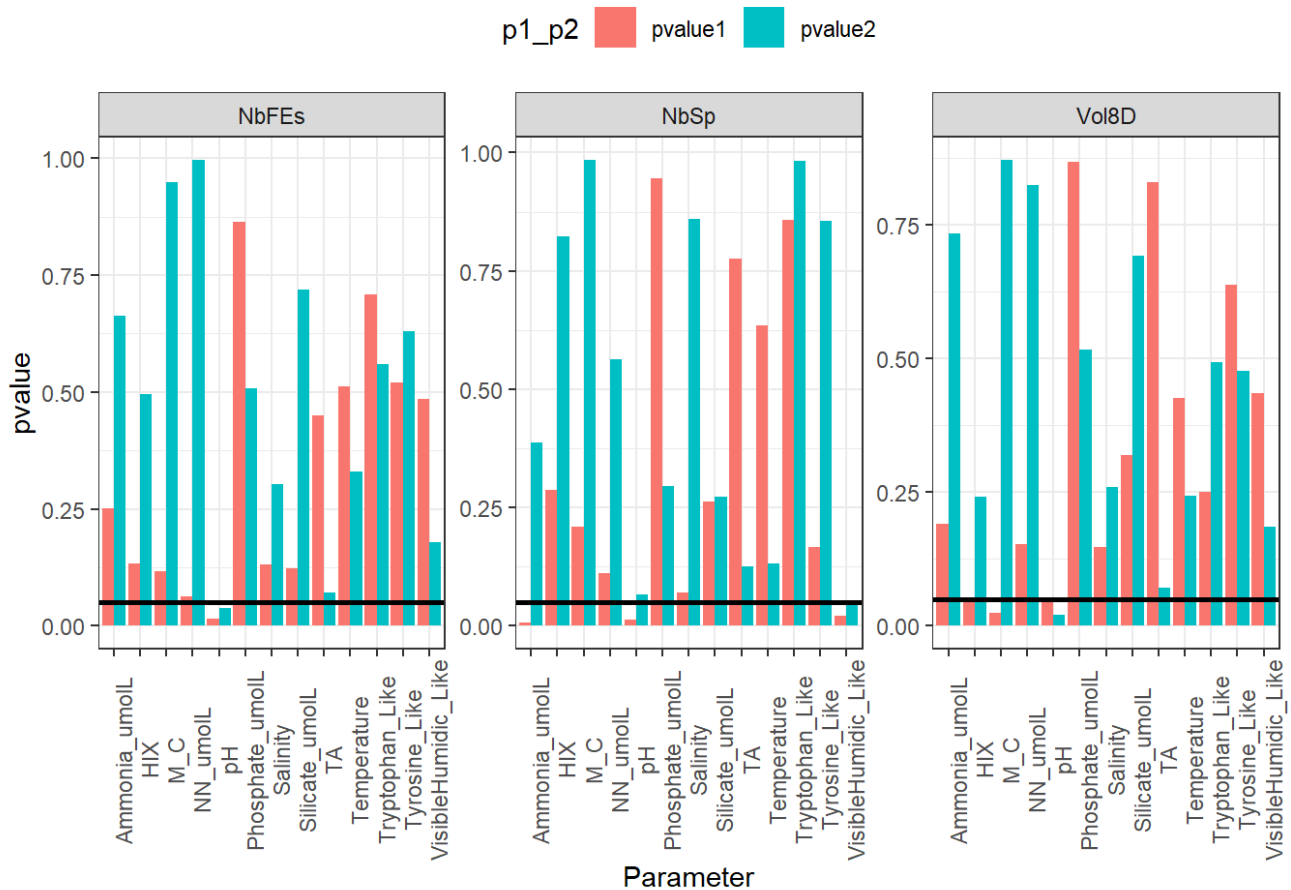
```
Max <- full_join(Max3, Max4)
```

```
## Joining, by = c("Parameter", "ParamType", "Dependent", "Season", "pvalue1",  
## "pvalue2", "r_squared", "adj_r_squared")
```

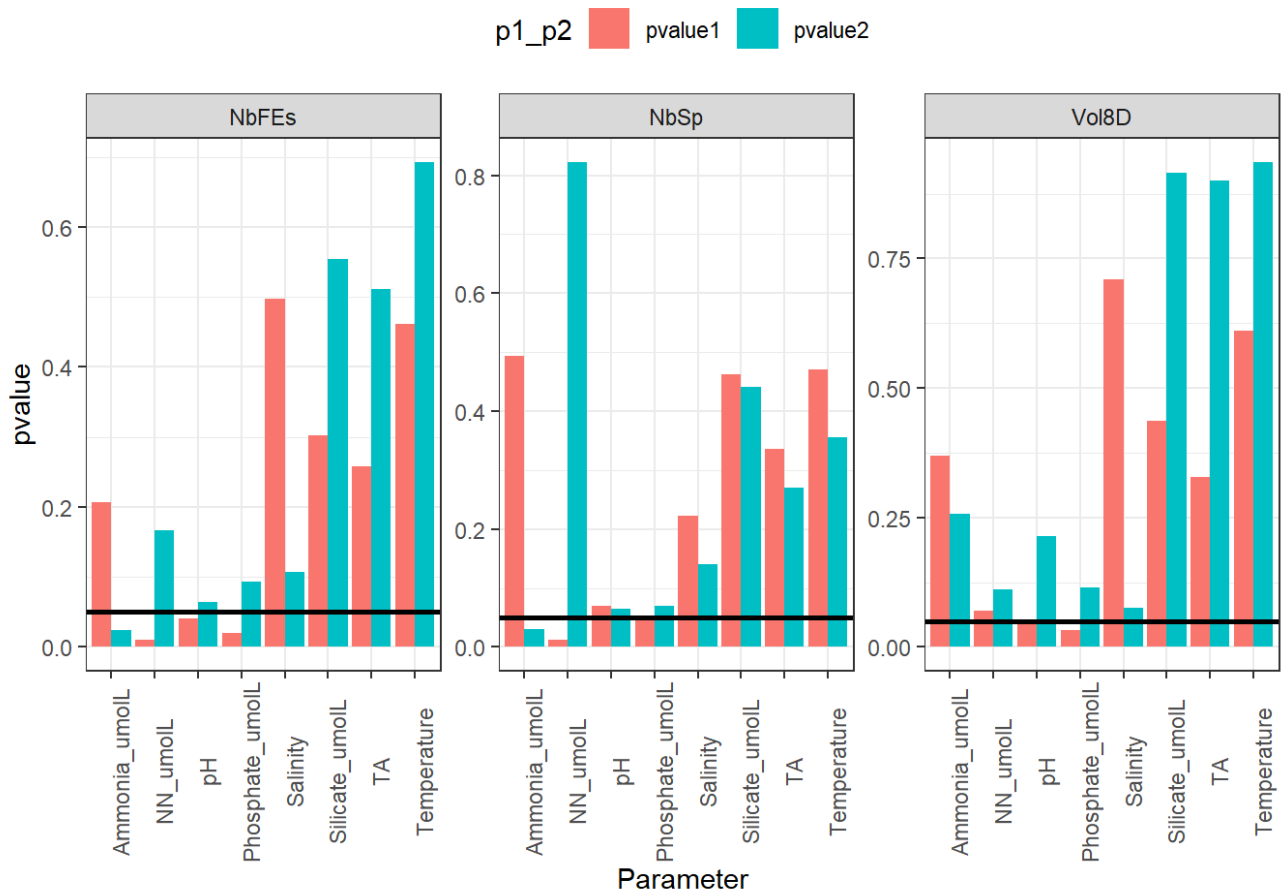
```
Max
```

```
## # A tibble: 105 × 8
##   Parameter      ParamType Dependent Season pvalue1 pvalue2 r_squared adj_r_...1
##   <chr>          <chr>      <chr>    <chr>    <dbl>   <dbl>   <dbl>   <dbl>
## 1 Salinity       Maximum    NbSp     Dry     0.0708  0.859   0.191   0.0900
## 2 Temperature    Maximum    NbSp     Dry     0.635   0.131   0.148   0.0410
## 3 TA             Maximum    NbSp     Dry     0.776   0.125   0.145   0.0378
## 4 pH             Maximum    NbSp     Dry     0.0123  0.0669  0.425   0.353
## 5 Phosphate_umolL Maximum    NbSp     Dry     0.947   0.295   0.0686 -0.0478
## 6 Silicate_umolL Maximum    NbSp     Dry     0.263   0.272   0.142   0.0343
## 7 NN_umolL       Maximum    NbSp     Dry     0.112   0.564   0.166   0.0615
## 8 Ammonia_umolL  Maximum    NbSp     Dry     0.00776 0.387   0.386   0.309
## 9 M_C            Maximum    NbSp     Dry     0.209   0.984   0.0966 -0.0163
## 10 HIX           Maximum    NbSp     Dry     0.288   0.822   0.0730 -0.0428
## # ... with 95 more rows, and abbreviated variable name 1adj_r_squared
```

```
pvalplot(mydata = Max, season = "Dry")
```



```
pvalplot(mydata = Max, season = "Wet")
```



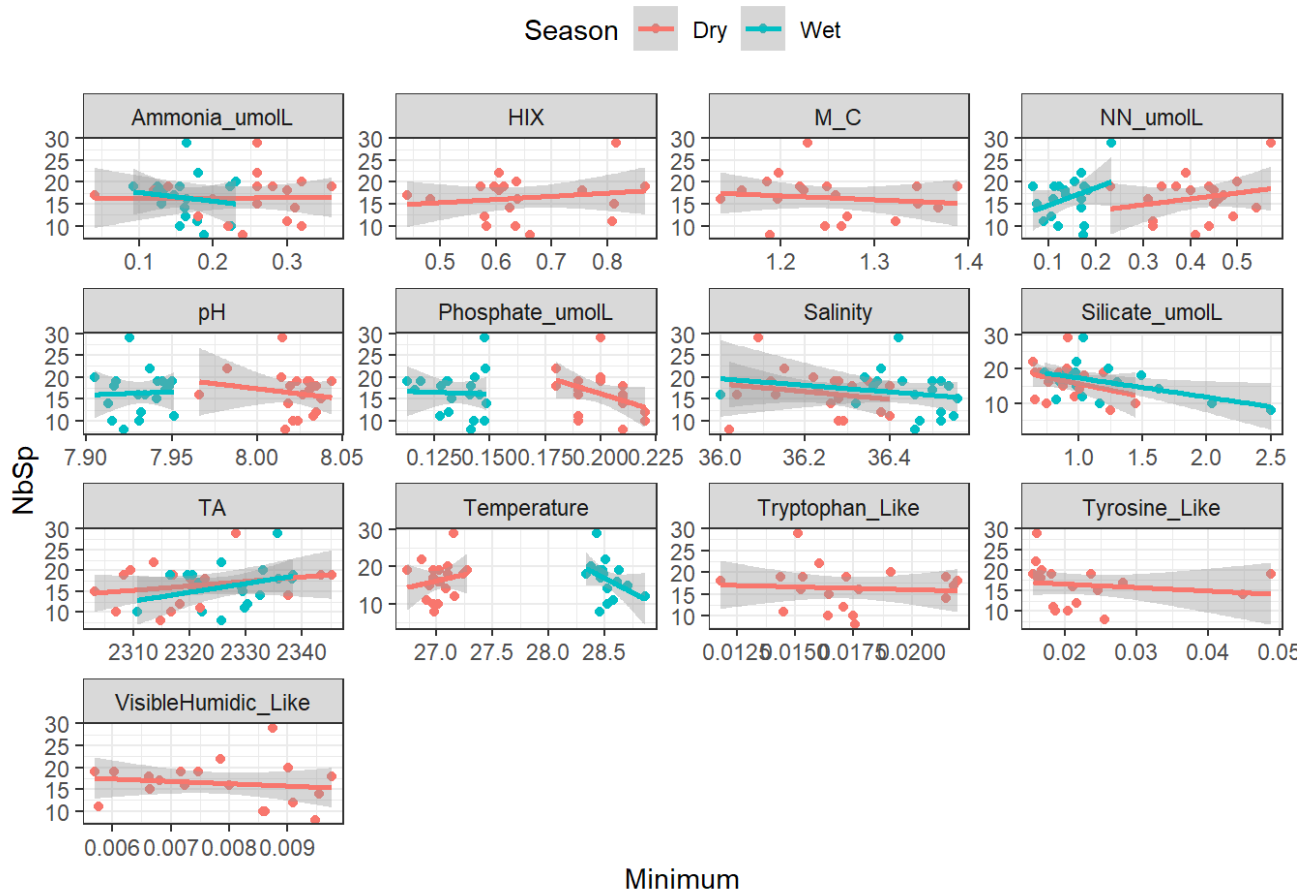
```
p1 <- paramPlot(mydata = Full_data, ParamType = Maximum, PTname = "Maximum")
```

```
## NULL
## NULL
```

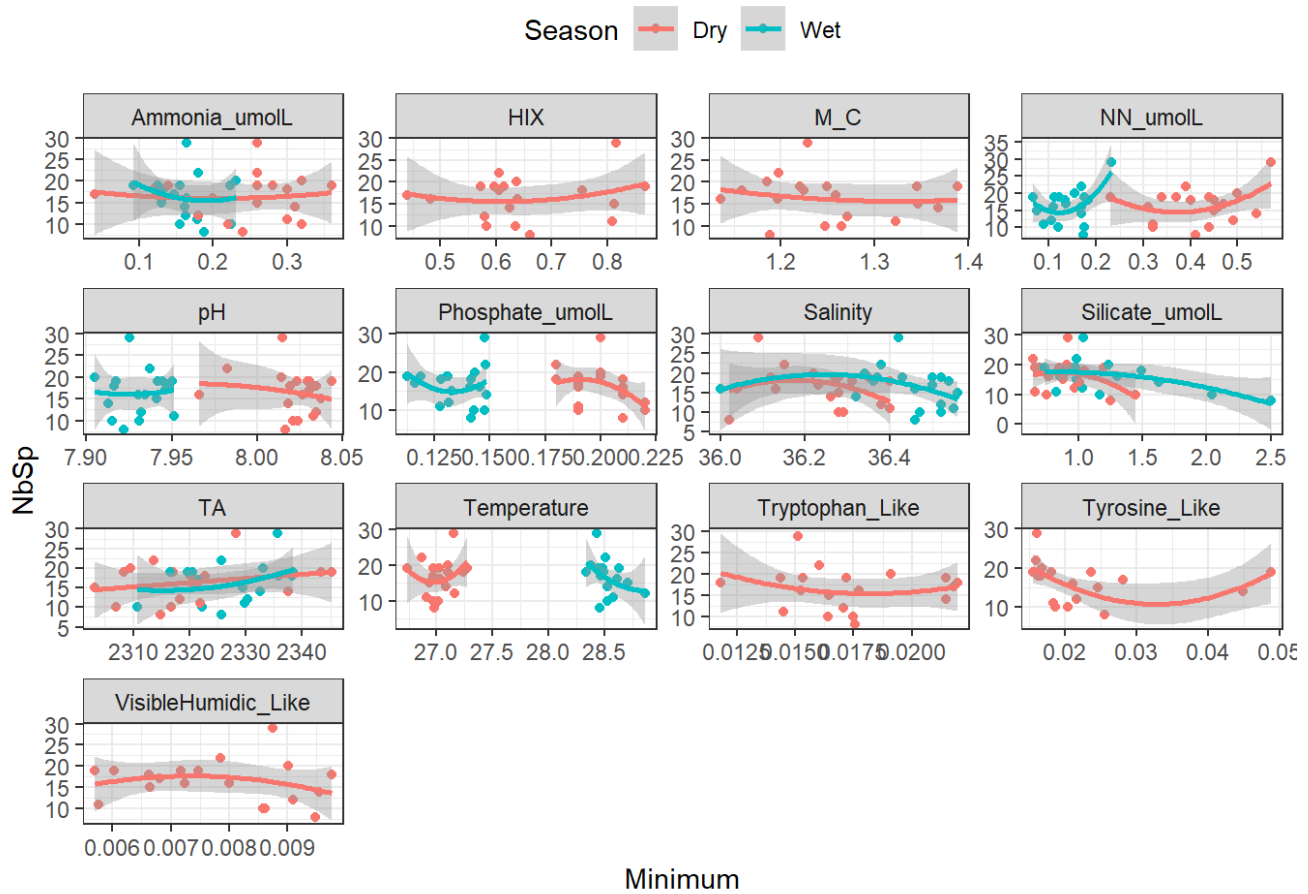
5/11/2023, 8:17 PM


```
## NULL
## NULL
```

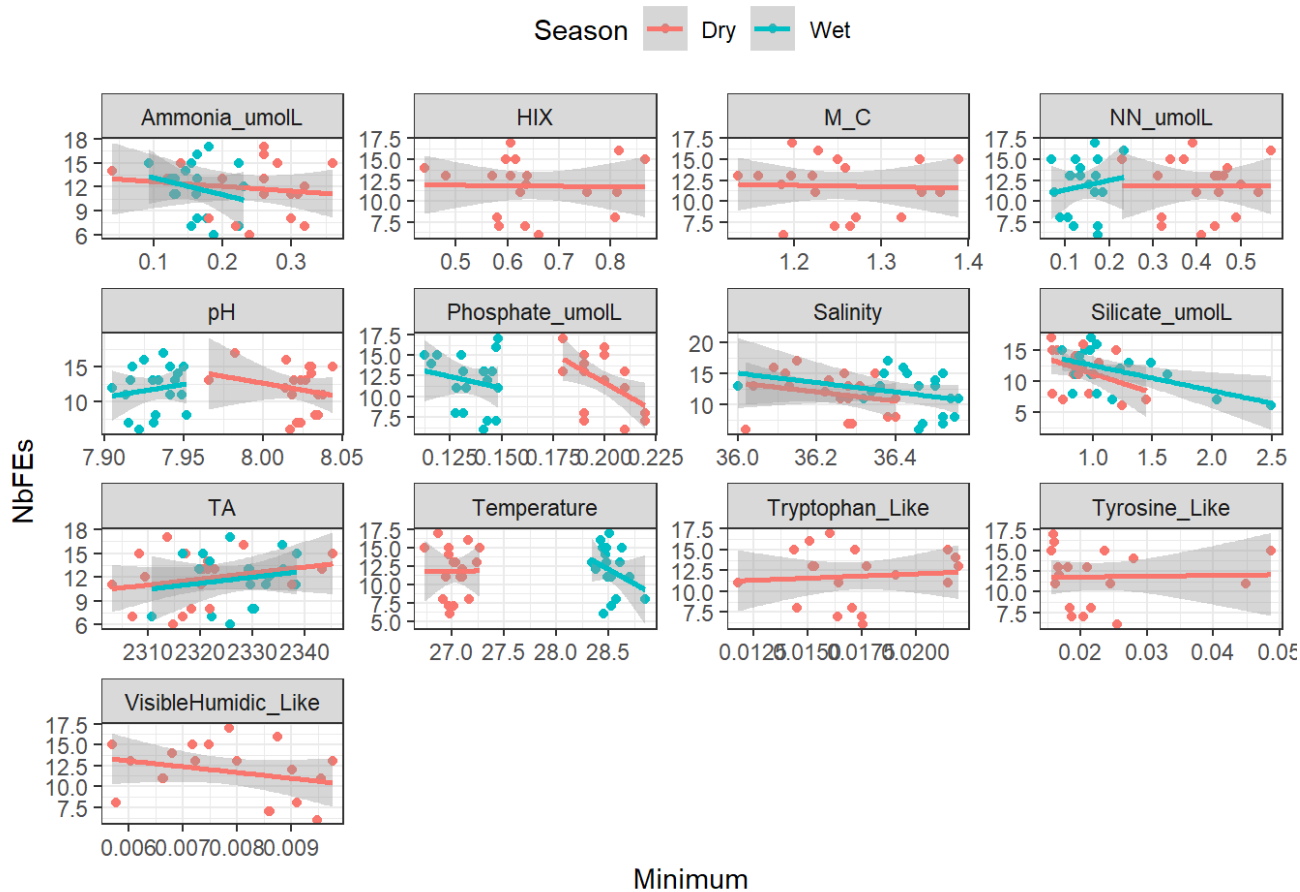
```
plotfun(y = NbSp, x = Minimum, myfacet = Parameters, myformula = "y~x")
```



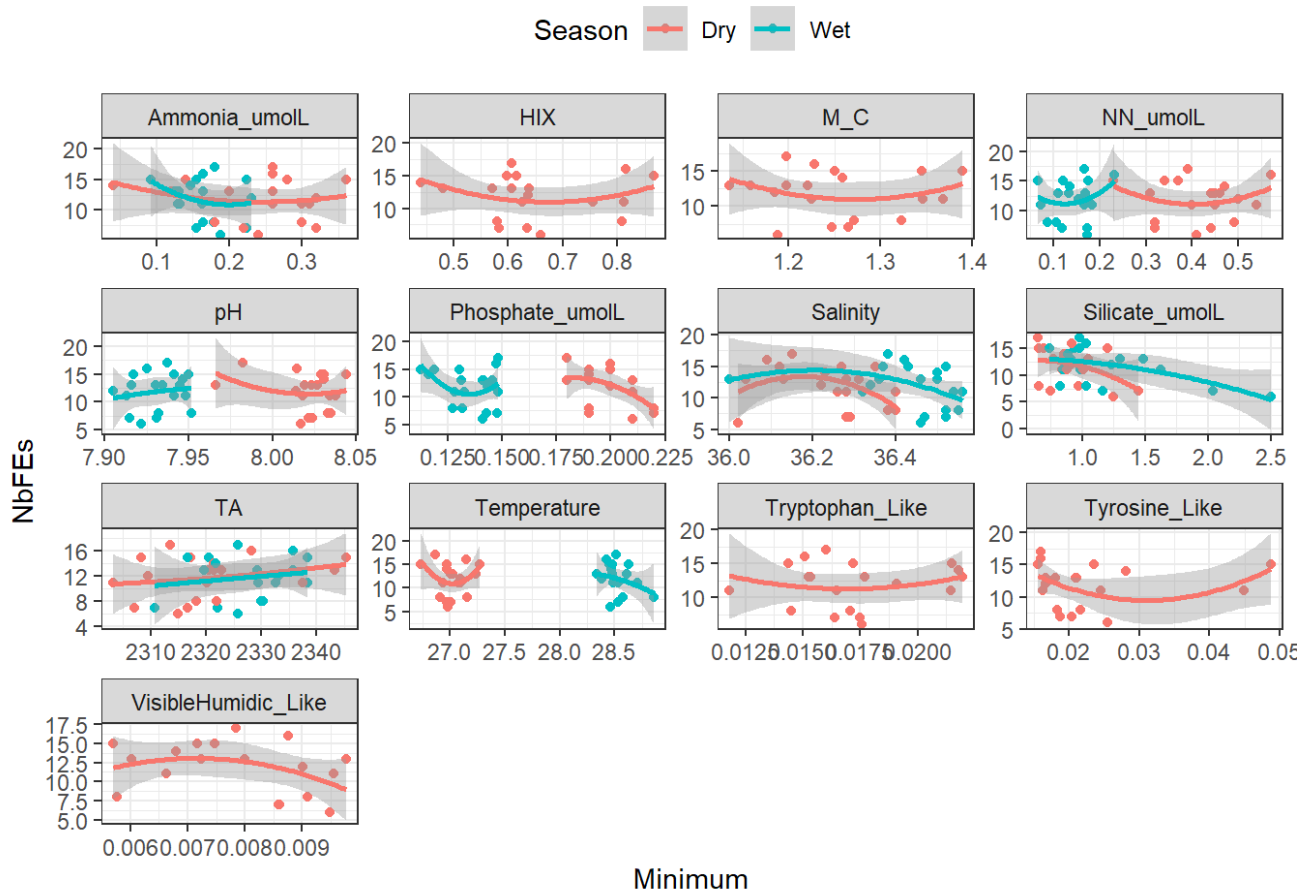
```
plotfun(y = NbSp, x = Minimum, myfacet = Parameters, myformula = "y~poly(x,2)")
```



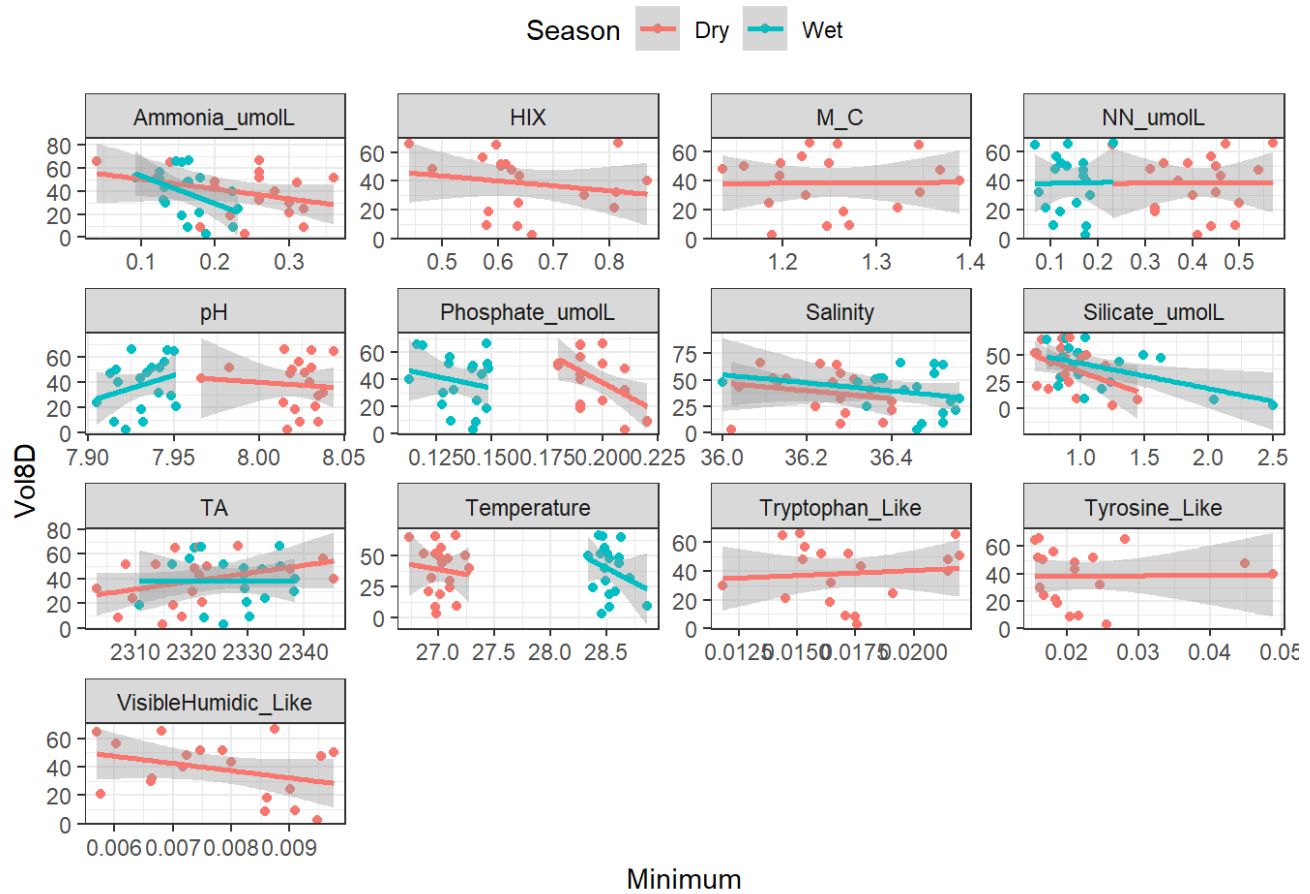
```
plotfun(y = NbFEs, x = Minimum, myfacet = Parameters, myformula = "y~x")
```



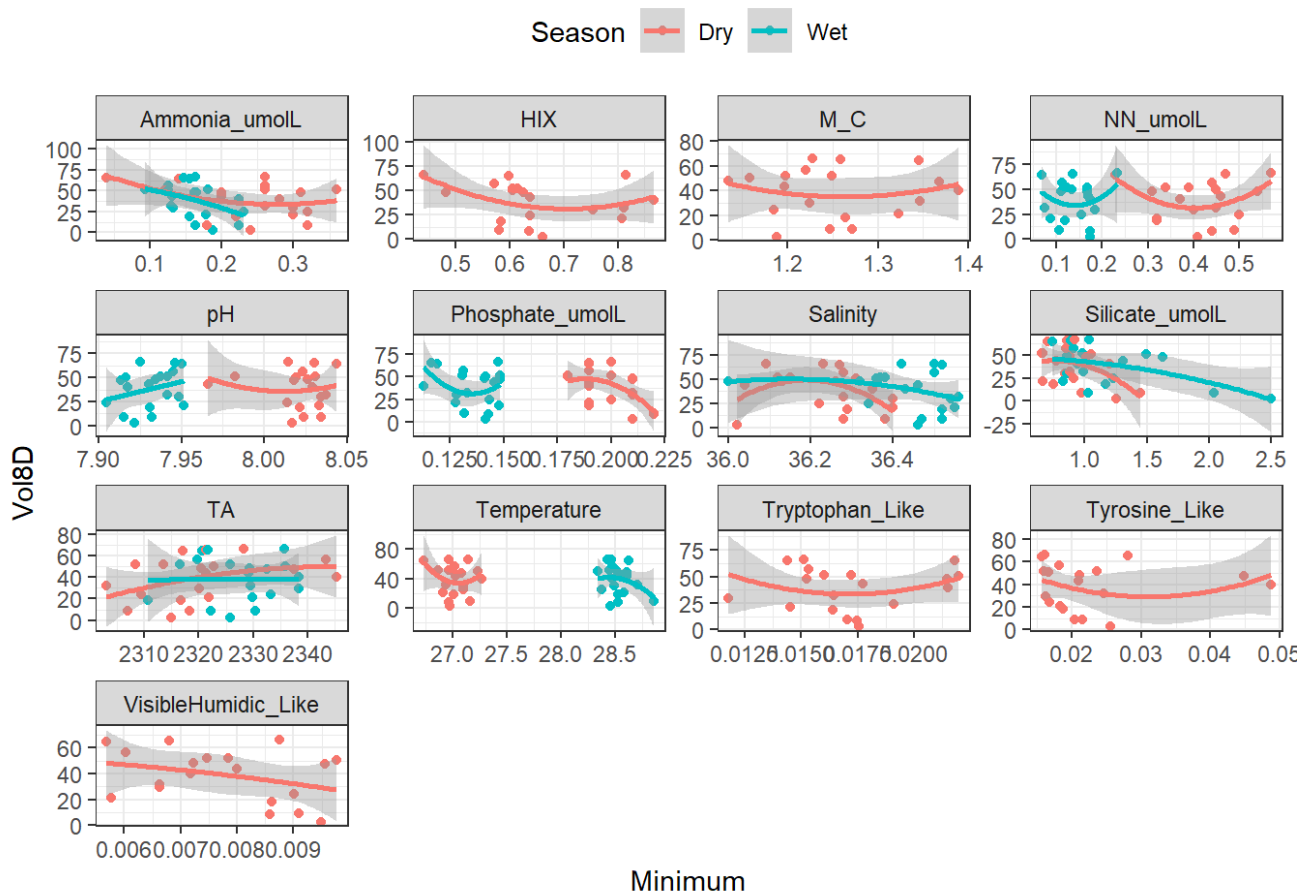
```
plotfun(y = NbFEs, x = Minimum, myfacet = Parameters, myformula = "y~poly(x,2)")
```



```
plotfun(y = Vol8D, x = Minimum, myfacet = Parameters, myformula = "y~x")
```



```
plotfun(y = Vol8D, x = Minimum, myfacet = Parameters, myformula = "y~poly(x,2)")
```



```
# get pvalues
```

```
Min3<-pvalpoly(myparam = "Minimum", myseason = "Dry")
```

```
Min4<-pvalpoly(myparam = "Minimum", myseason = "Wet")
```

```
Min <- full_join(Min3, Min4)
```

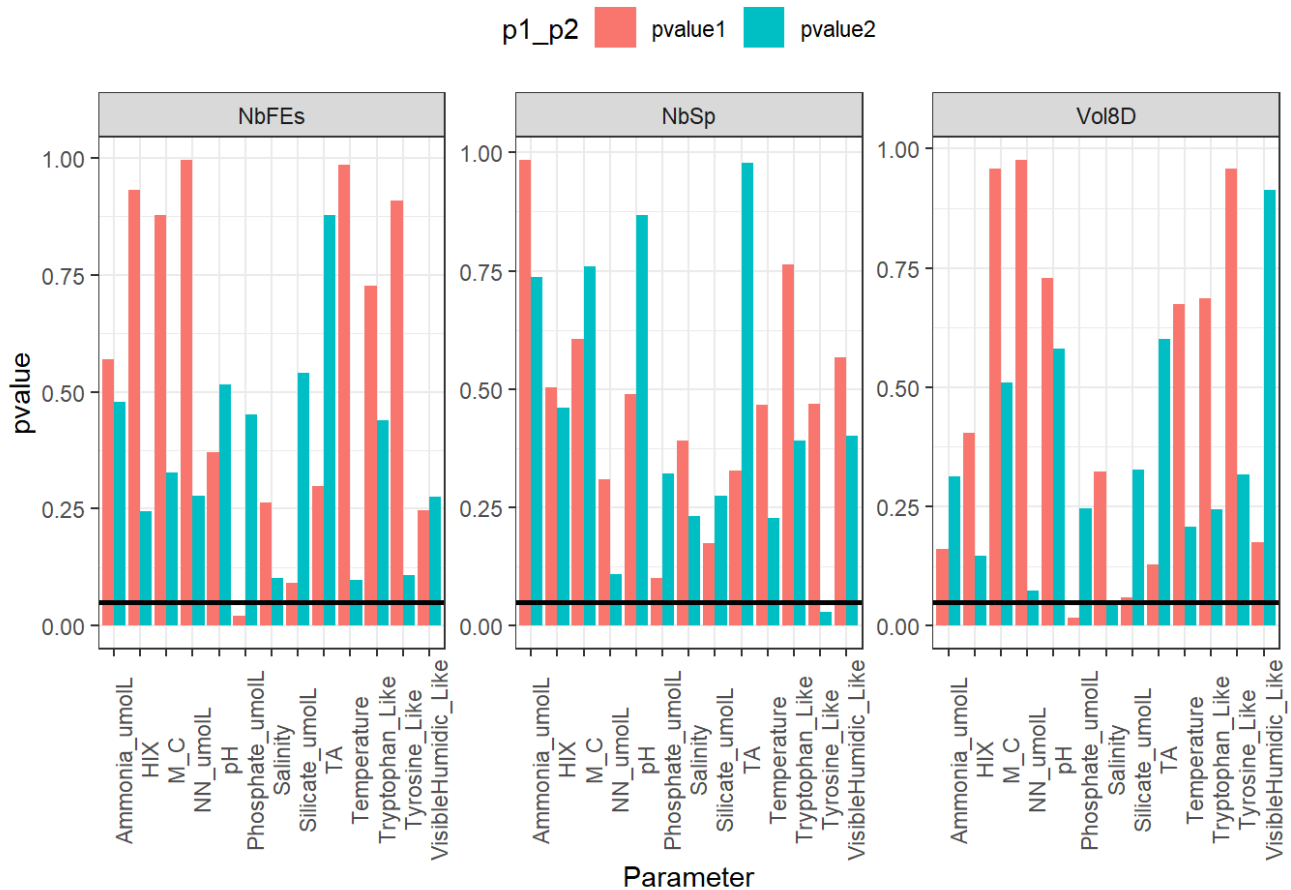
```
## Joining, by = c("Parameter", "ParamType", "Dependent", "Season", "pvalue1",
```

```
## "pvalue2", "r_squared", "adj_r_squared")
```

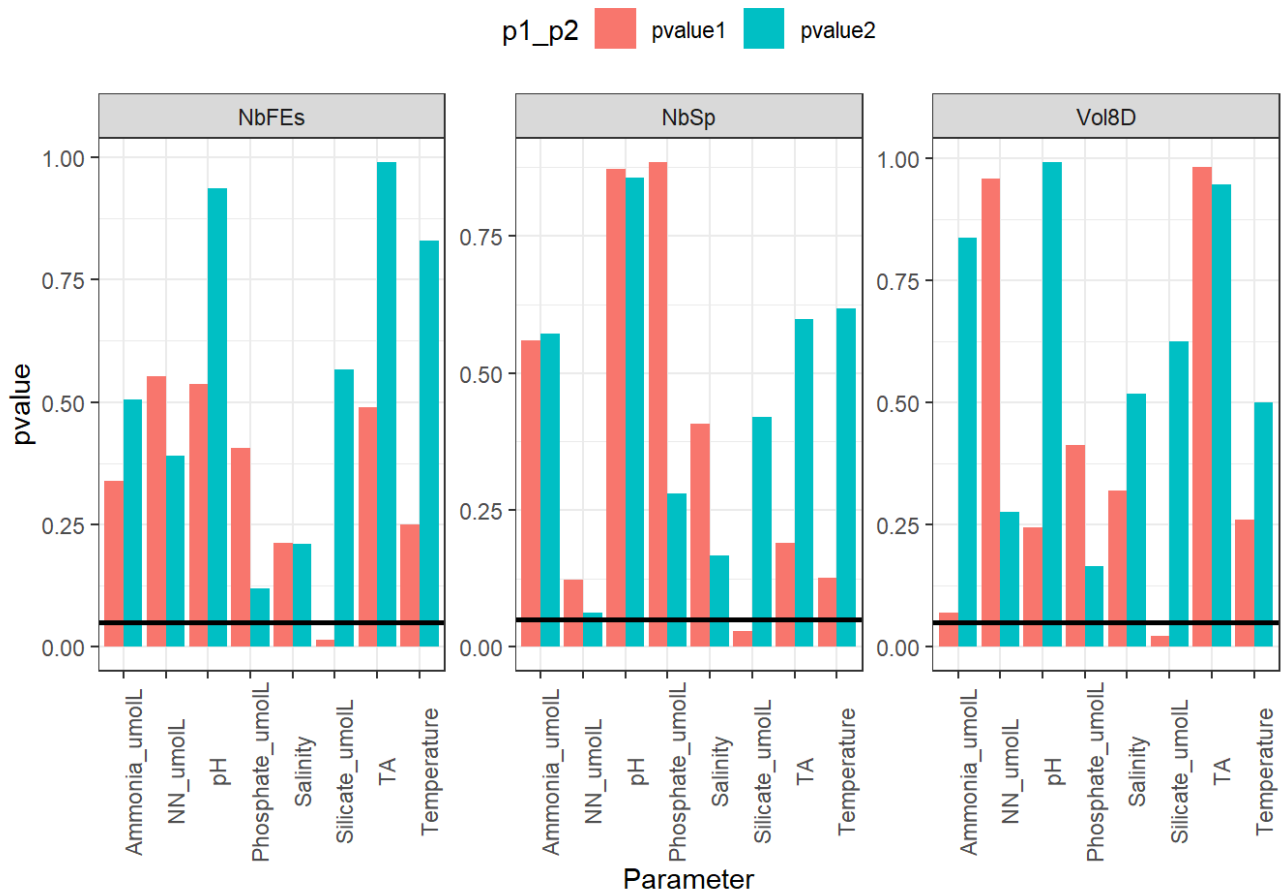
```
Min
```

```
## # A tibble: 105 × 8
##   Parameter      ParamType Dependent Season pvalue1 pvalue2 r_squared adj_r_...1
##   <chr>          <chr>      <chr>    <chr>    <dbl>   <dbl>   <dbl>   <dbl>
## 1 Salinity       Minimum    NbSp     Dry      0.392   0.232   0.127   0.0176
## 2 Temperature    Minimum    NbSp     Dry      0.467   0.228   0.117   0.00693
## 3 TA             Minimum    NbSp     Dry      0.328   0.978   0.0598  -0.0577
## 4 pH            Minimum    NbSp     Dry      0.490   0.867   0.0320  -0.0890
## 5 Phosphate_umolL Minimum    NbSp     Dry      0.101   0.322   0.203   0.103
## 6 Silicate_umolL Minimum    NbSp     Dry      0.174   0.275   0.171   0.0674
## 7 NN_umolL       Minimum    NbSp     Dry      0.310   0.110   0.198   0.0982
## 8 Ammonia_umolL  Minimum    NbSp     Dry      0.984   0.736   0.00730 -0.117
## 9 M_C           Minimum    NbSp     Dry      0.606   0.759   0.0229  -0.0993
## 10 HIX           Minimum    NbSp     Dry      0.504   0.460   0.0611  -0.0563
## # ... with 95 more rows, and abbreviated variable name 1adj_r_squared
```

```
pvalplot(mydata = Min, season = "Dry")
```



```
pvalplot(mydata = Min, season = "Wet")
```

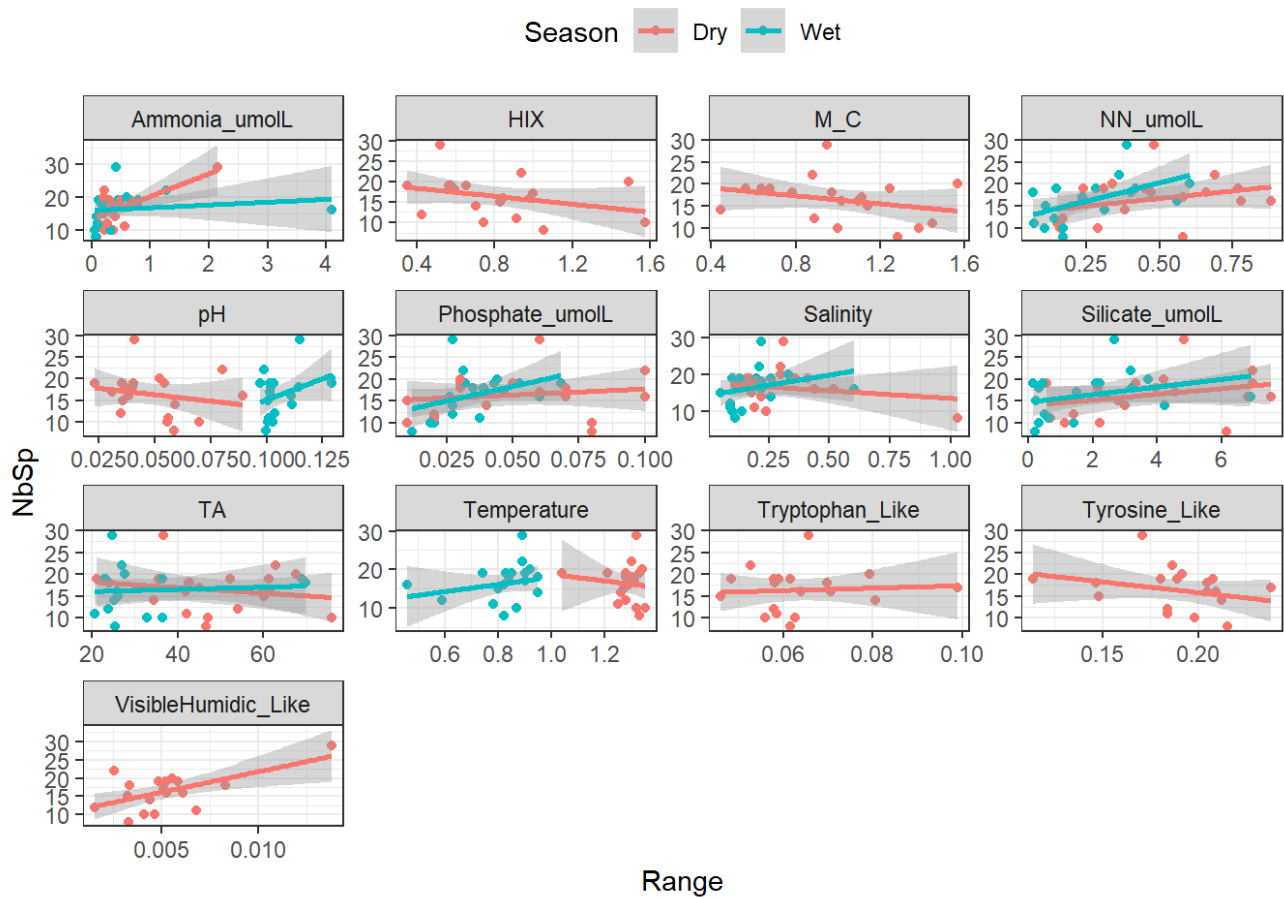
```
p2 <- paramPlot(mydata = Full_data, ParamType = Minimum, PTname = "Minimum")
```

```
## NULL
## NULL
```

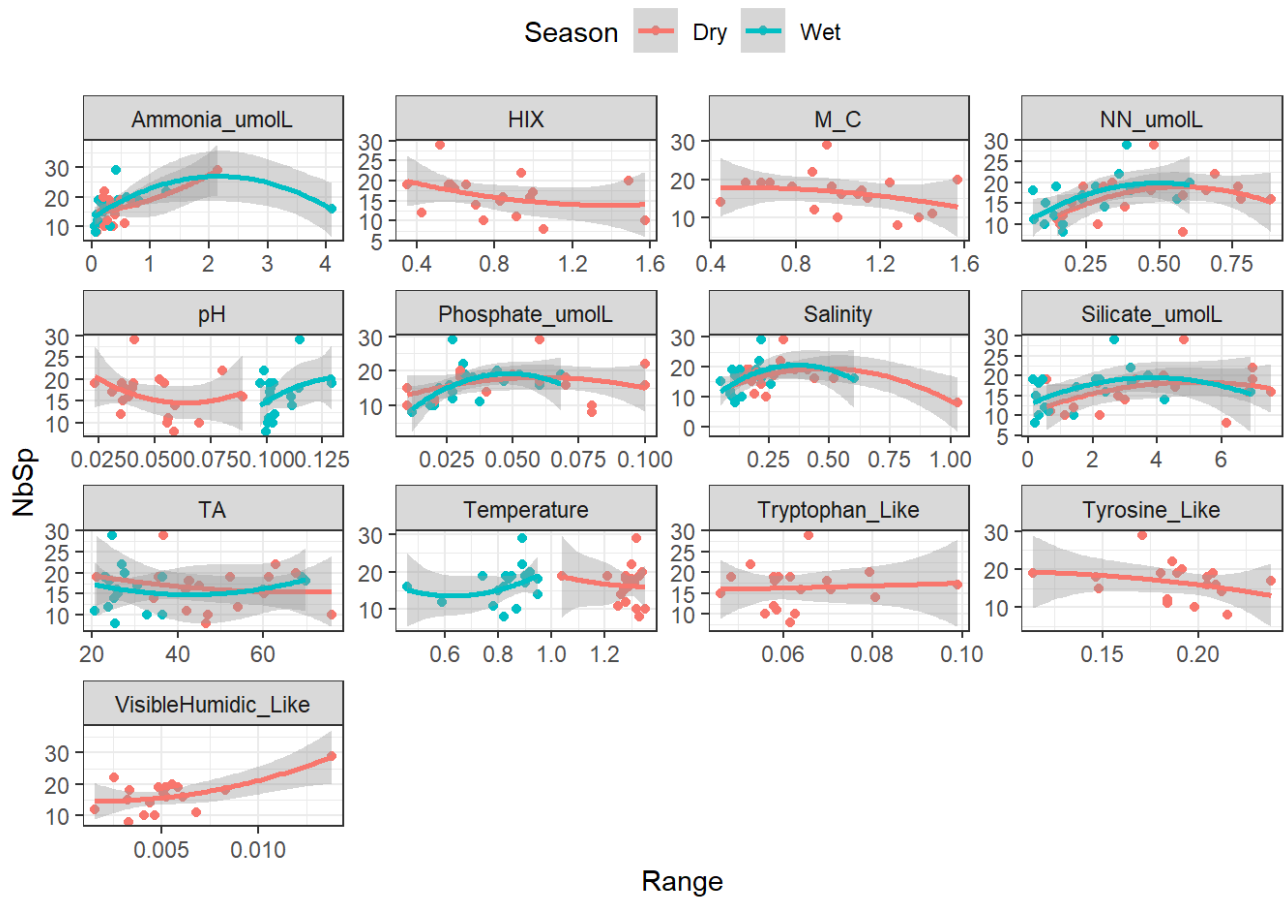
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```
## NULL
## NULL
```

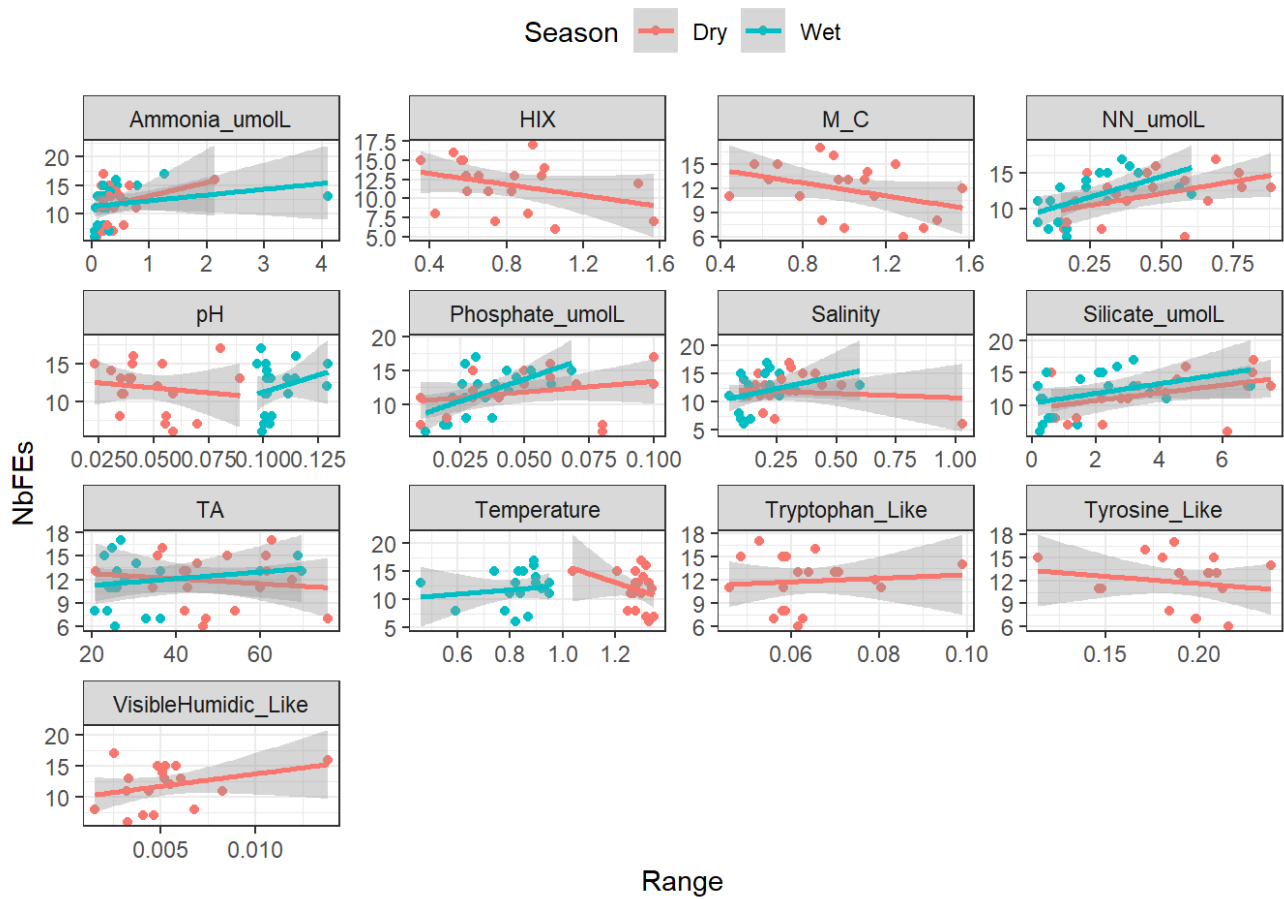
```
plotfun(y = NbSp, x = Range, myfacet = Parameters, myformula = "y~x")
```



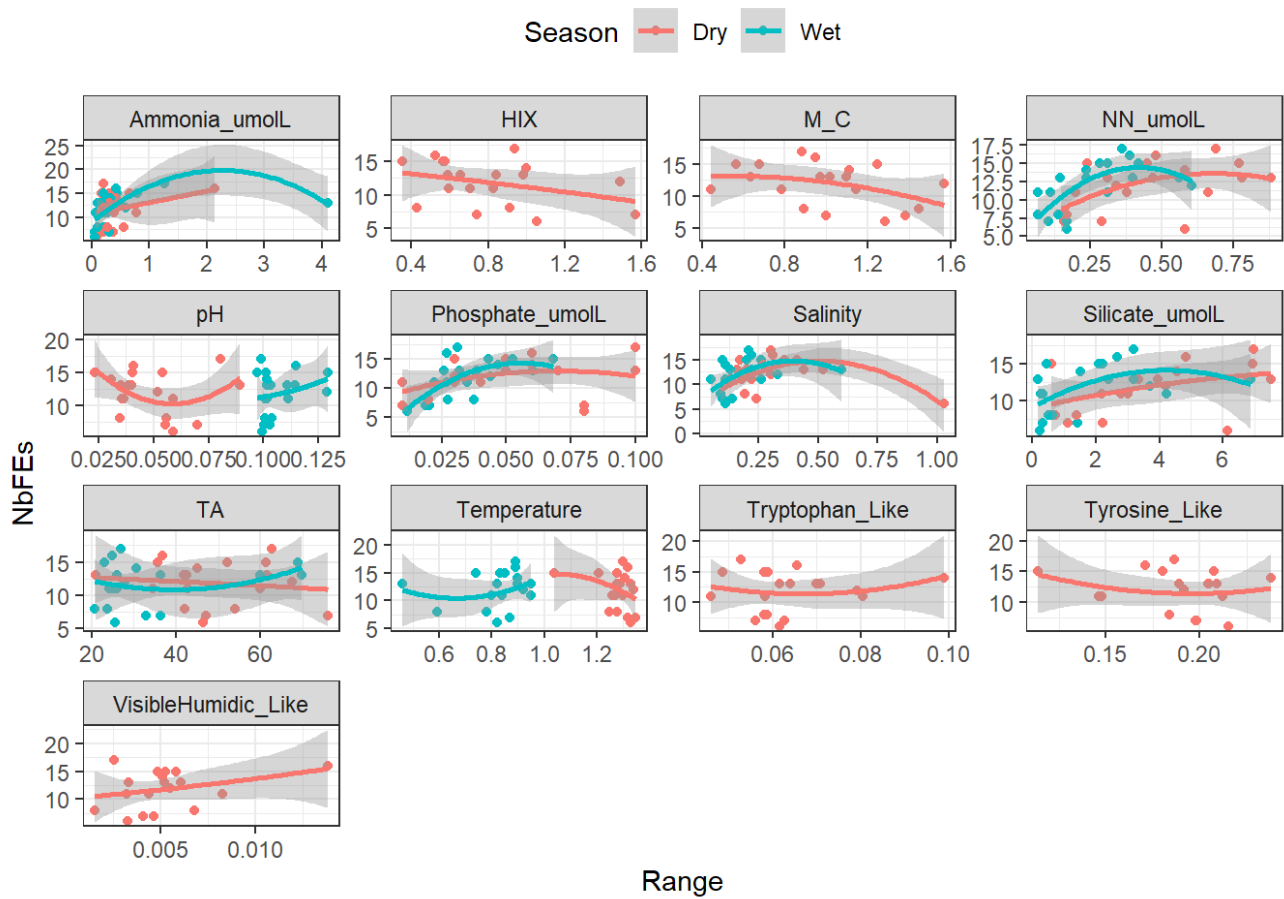
```
plotfun(y = NbSp, x = Range, myfacet = Parameters, myformula = "y~poly(x,2)")
```



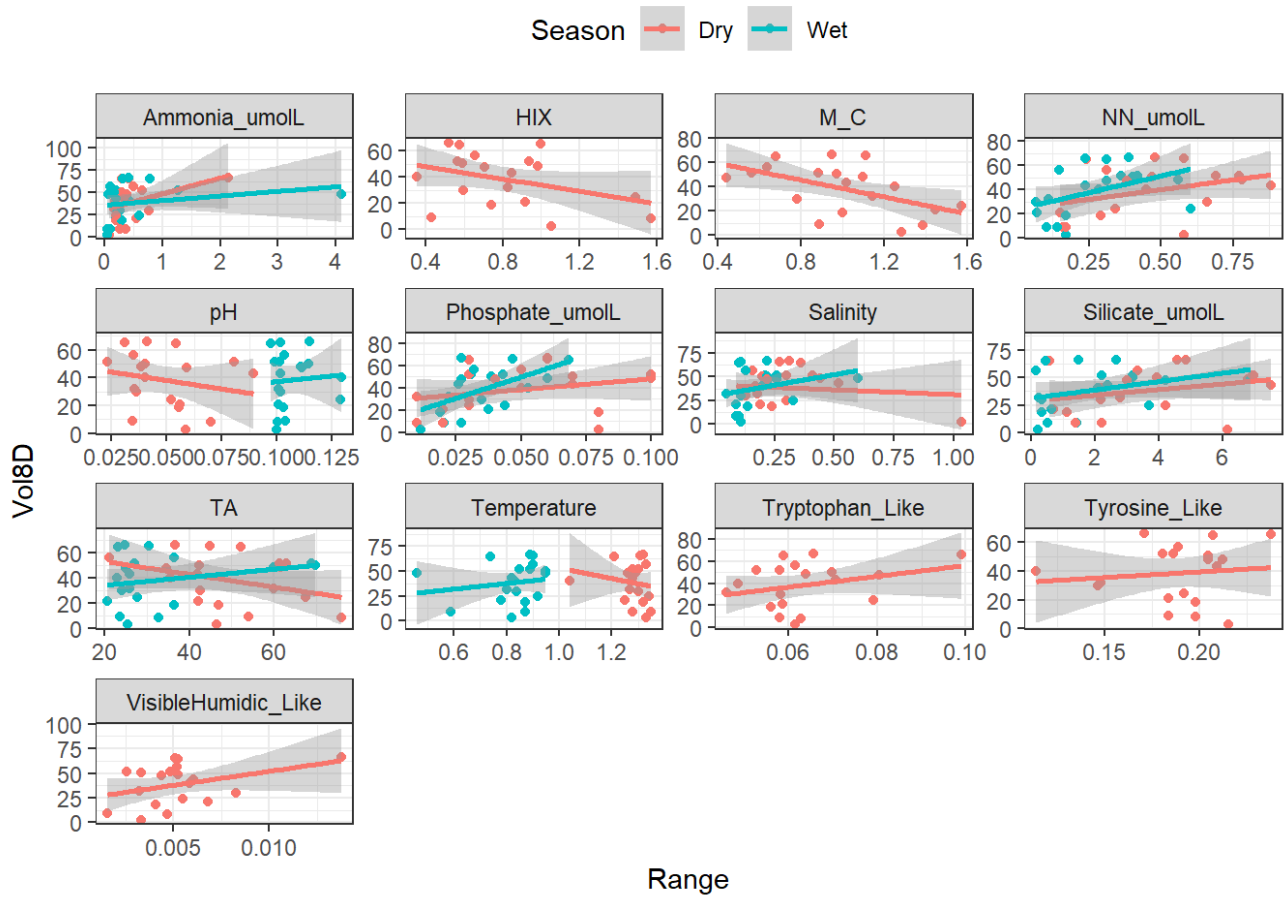
```
plotfun(y = NbFEs, x = Range, myfacet = Parameters, myformula = "y~x")
```



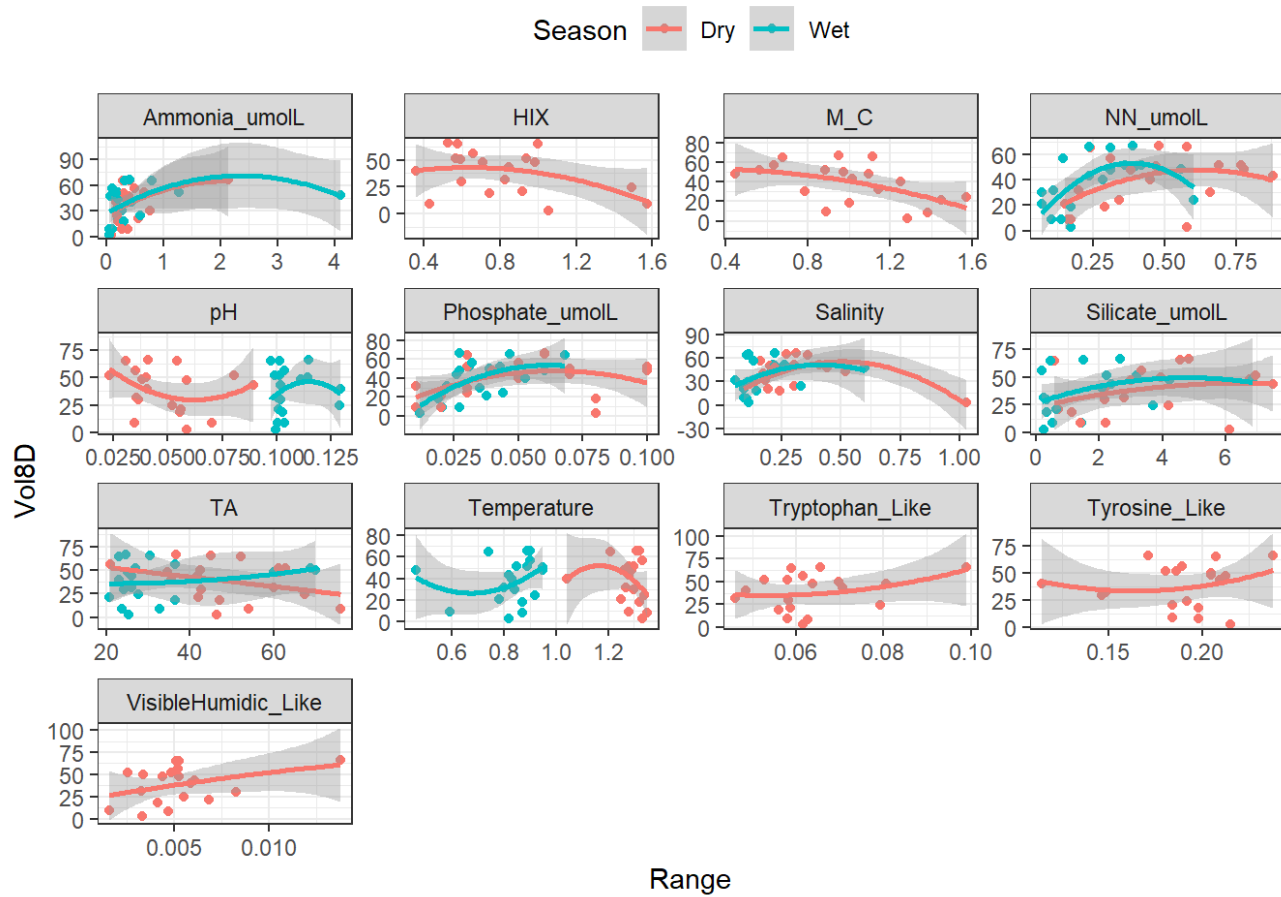
```
plotfun(y = NbFEs, x = Range, myfacet = Parameters, myformula = "y~poly(x,2)")
```



```
plotfun(y = Vol8D, x = Range, myfacet = Parameters, myformula = "y~x")
```



```
plotfun(y = Vol8D, x = Range, myfacet = Parameters, myformula = "y~poly(x,2)")
```

```
# get pvalues
```

```
Ran3<-pvalpoly(myparam = "Range", myseason = "Dry")
```

```
Ran4<-pvalpoly(myparam = "Range", myseason = "Wet")
```

```
Ran <- full_join(Ran3, Ran4)
```

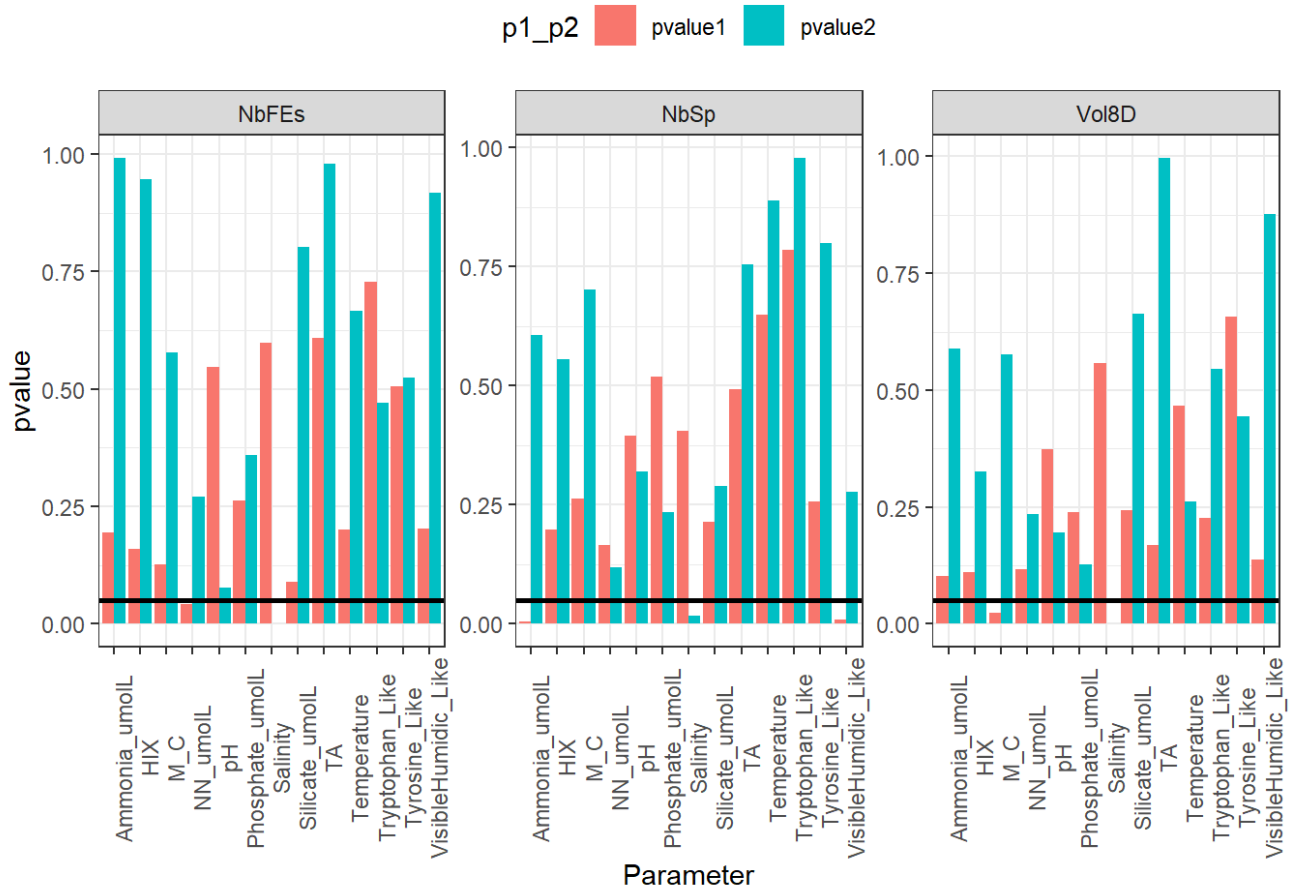
```
## Joining, by = c("Parameter", "ParamType", "Dependent", "Season", "pvalue1",
```

```
## "pvalue2", "r_squared", "adj_r_squared")
```

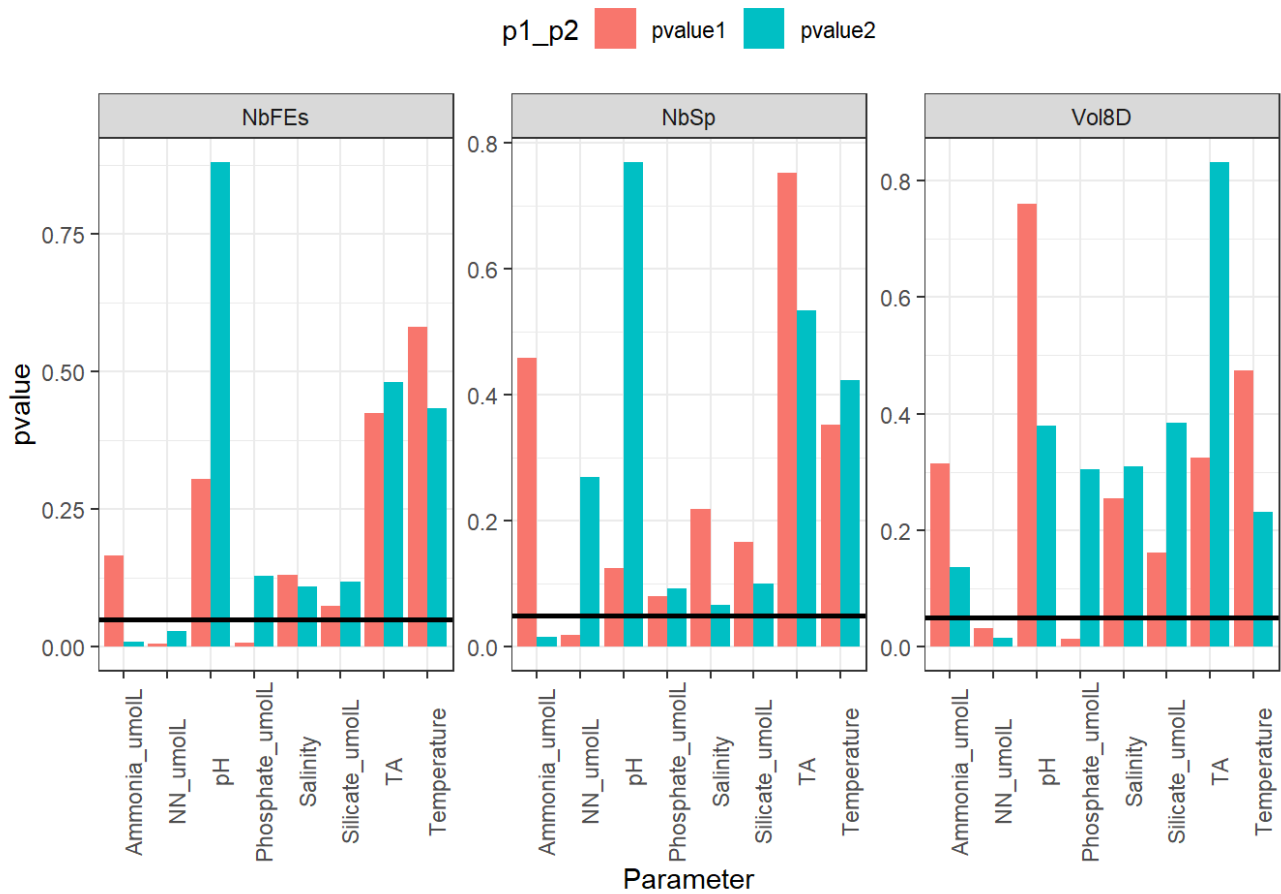
```
Ran
```

```
## # A tibble: 105 × 8
##   Parameter      ParamType Dependent Season pvalue1 pvalue2 r_squared adj_r_...1
##   <chr>          <chr>      <chr>    <chr>    <dbl>   <dbl>   <dbl>   <dbl>
## 1 Salinity       Range      NbSp     Dry     0.406   0.0161   0.332   0.249
## 2 Temperature    Range      NbSp     Dry     0.648   0.888   0.0146 -0.109
## 3 TA             Range      NbSp     Dry     0.493   0.755   0.0357 -0.0848
## 4 pH             Range      NbSp     Dry     0.396   0.320   0.102   -0.0103
## 5 Phosphate_umolL Range      NbSp     Dry     0.520   0.234   0.109   -0.00223
## 6 Silicate_umolL Range      NbSp     Dry     0.214   0.289   0.152   0.0462
## 7 NN_umolL       Range      NbSp     Dry     0.166   0.119   0.231   0.135
## 8 Ammonia_umolL  Range      NbSp     Dry     0.00567 0.607   0.395   0.320
## 9 M_C            Range      NbSp     Dry     0.262   0.702   0.0859 -0.0283
## 10 HIX           Range      NbSp     Dry     0.197   0.556   0.120   0.00966
## # ... with 95 more rows, and abbreviated variable name 1adj_r_squared
```

```
pvalplot(mydata = Ran, season = "Dry")
```



```
pvalplot(mydata = Ran, season = "Wet")
```



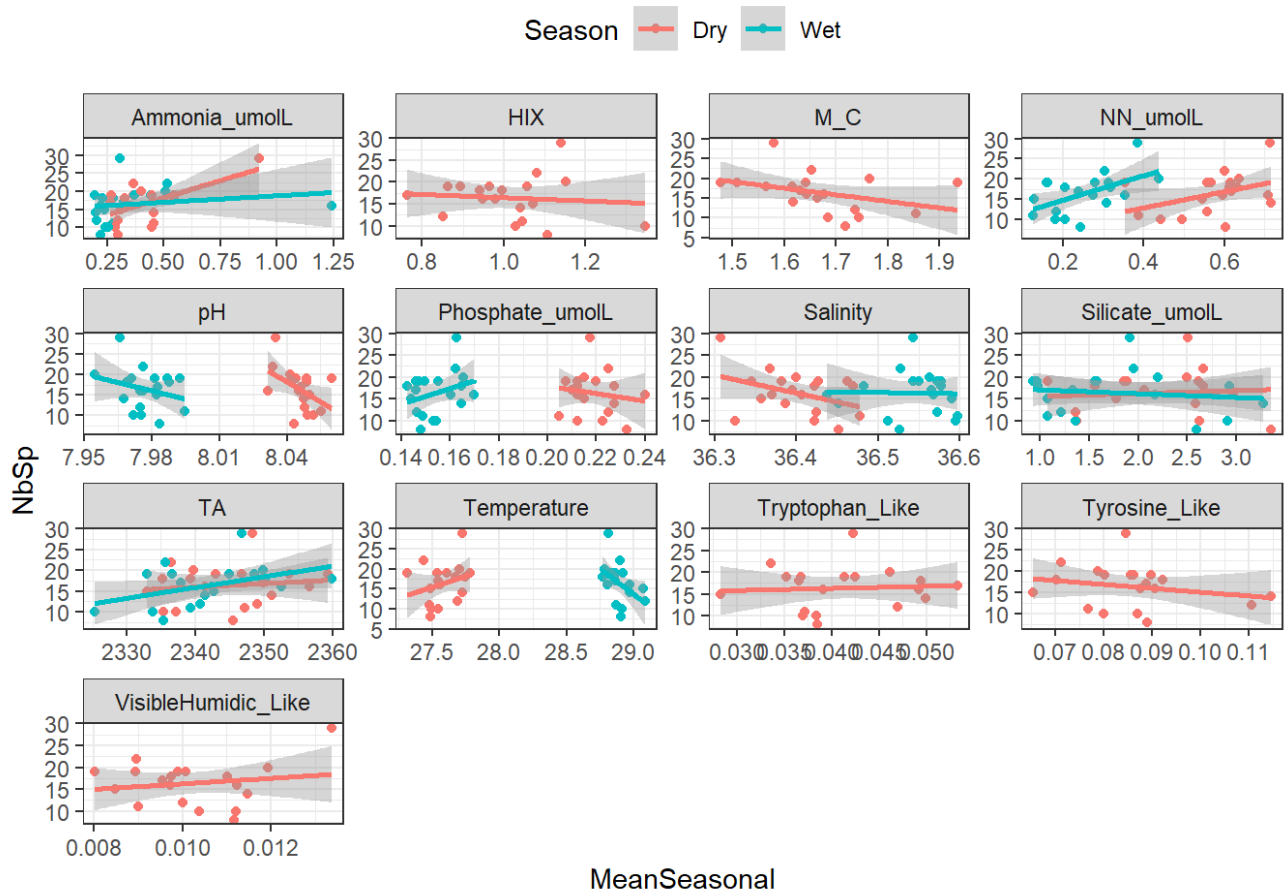
```
p3 <- paramPlot(mydata = Full_data, ParamType = Range, PTname = "Range")
```

```
## NULL
## NULL
```

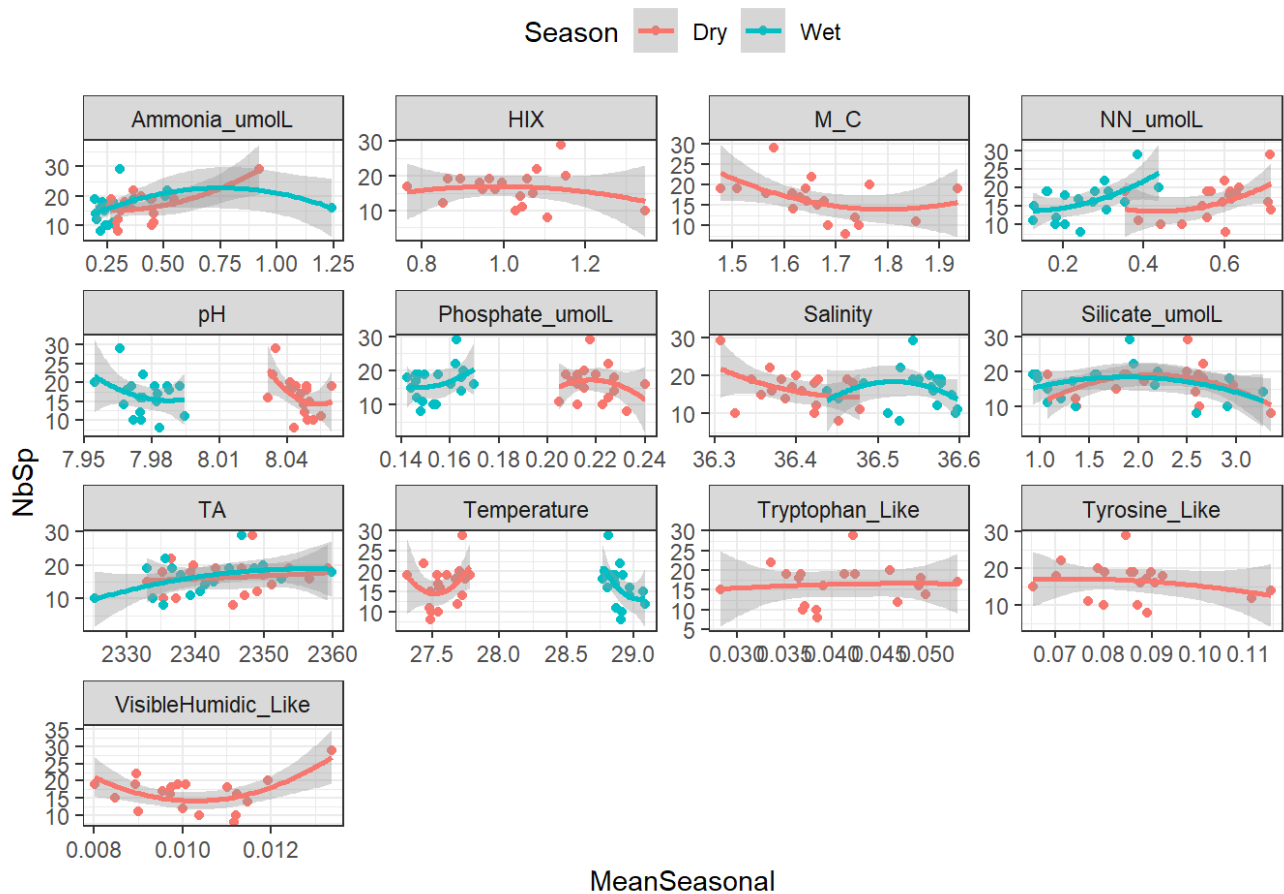
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```
## NULL
## NULL
```

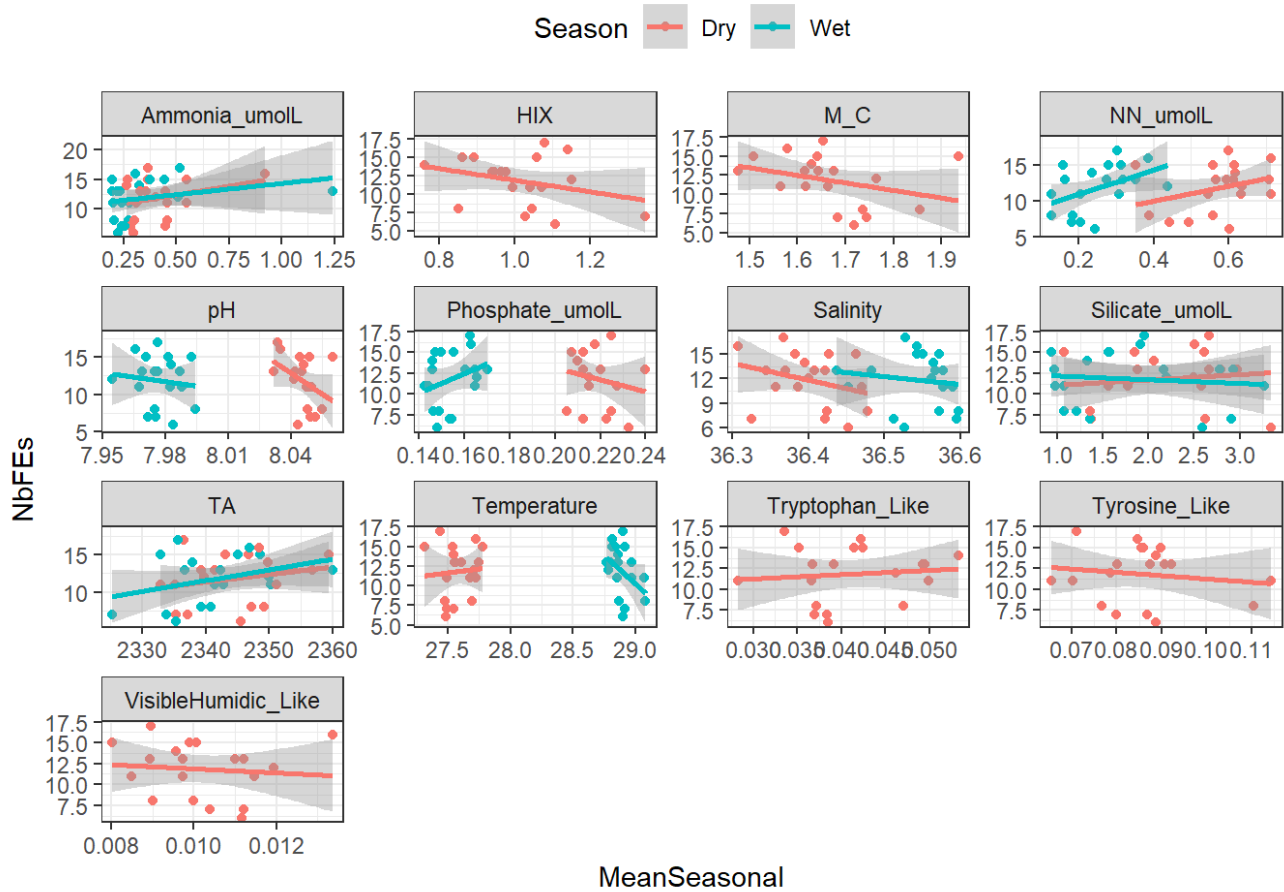
```
plotfun(y = NbSp, x = MeanSeasonal, myfacet = Parameters, myformula = "y~x")
```



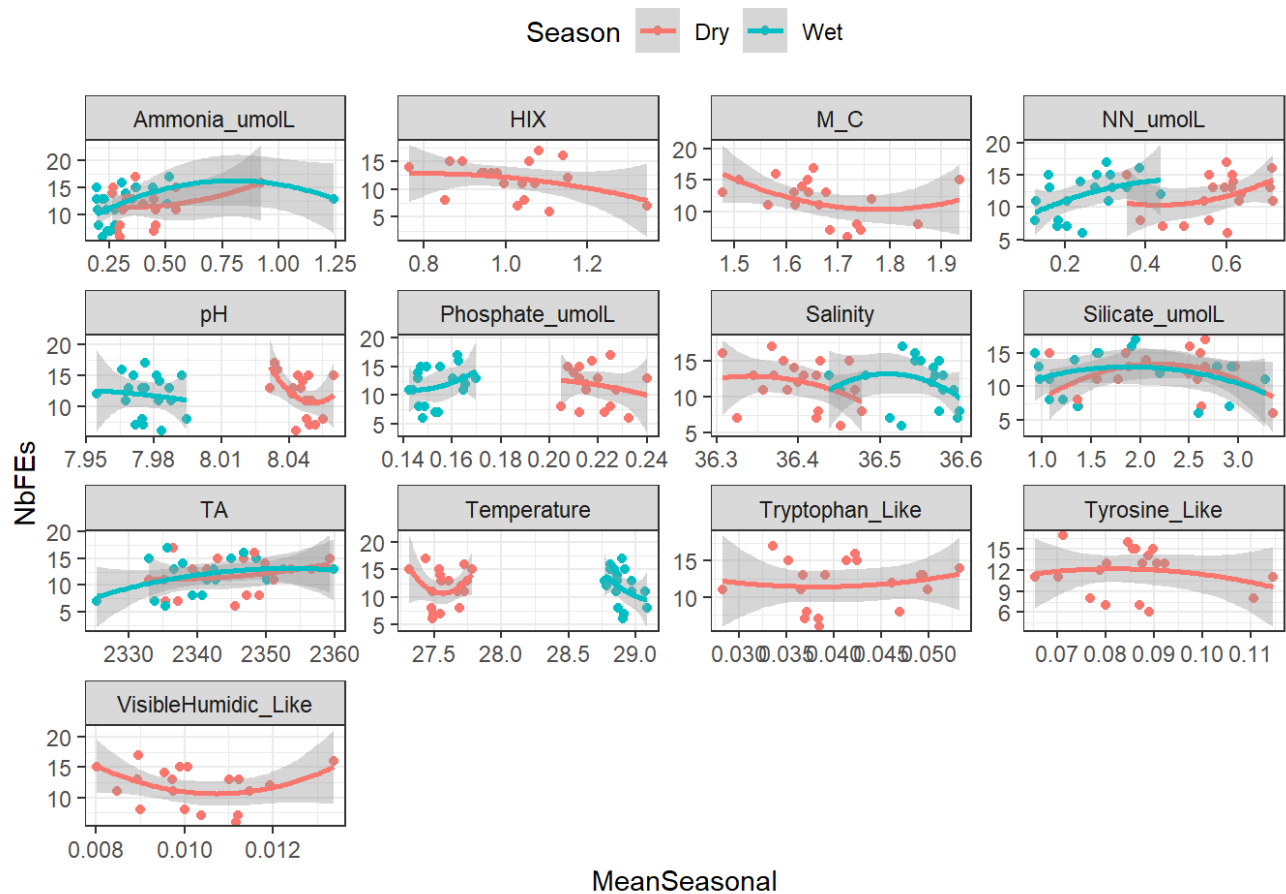
```
plotfun(y = NbSp, x = MeanSeasonal, myfacet = Parameters, myformula = "y~poly(x,2)")
```



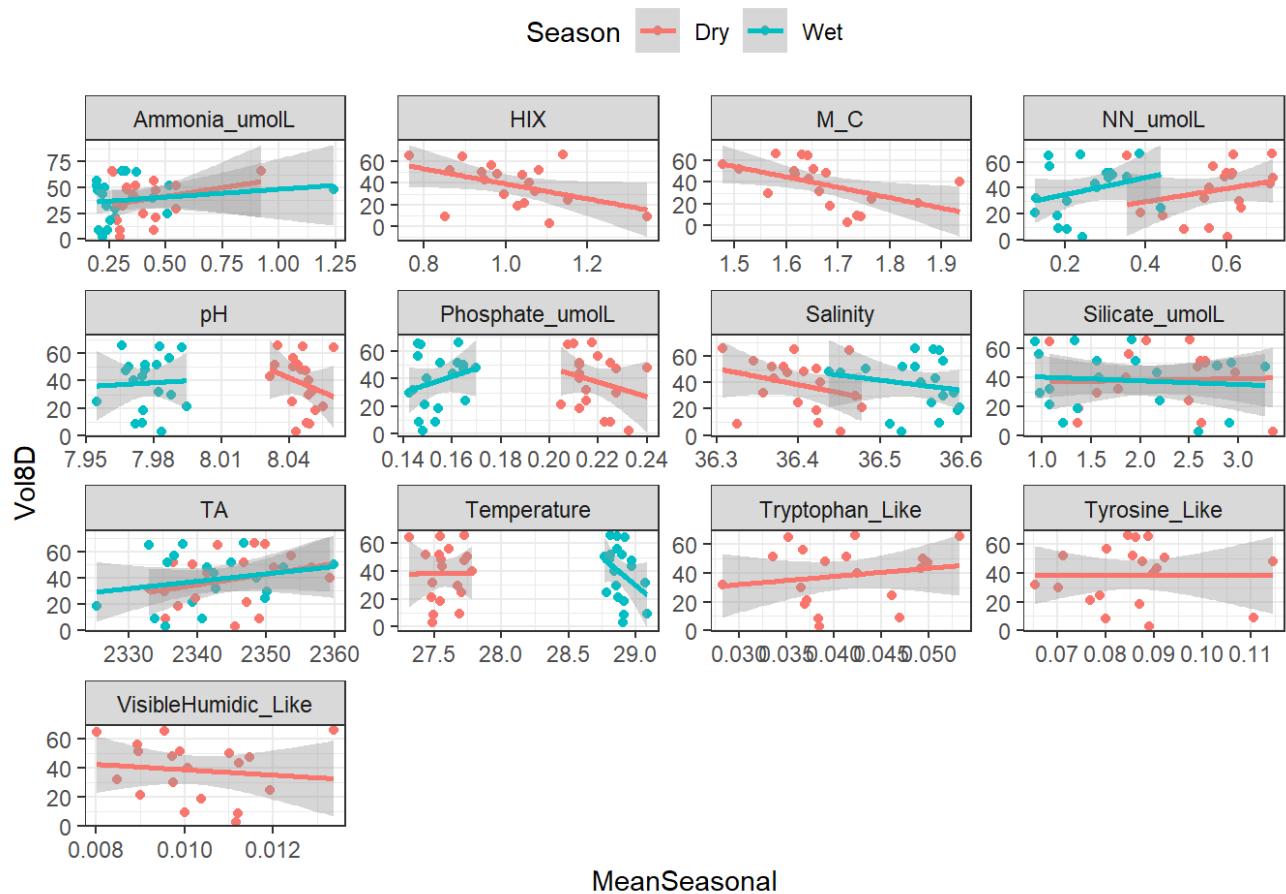
```
plotfun(y = NbFEs, x = MeanSeasonal, myfacet = Parameters, myformula = "y~x")
```



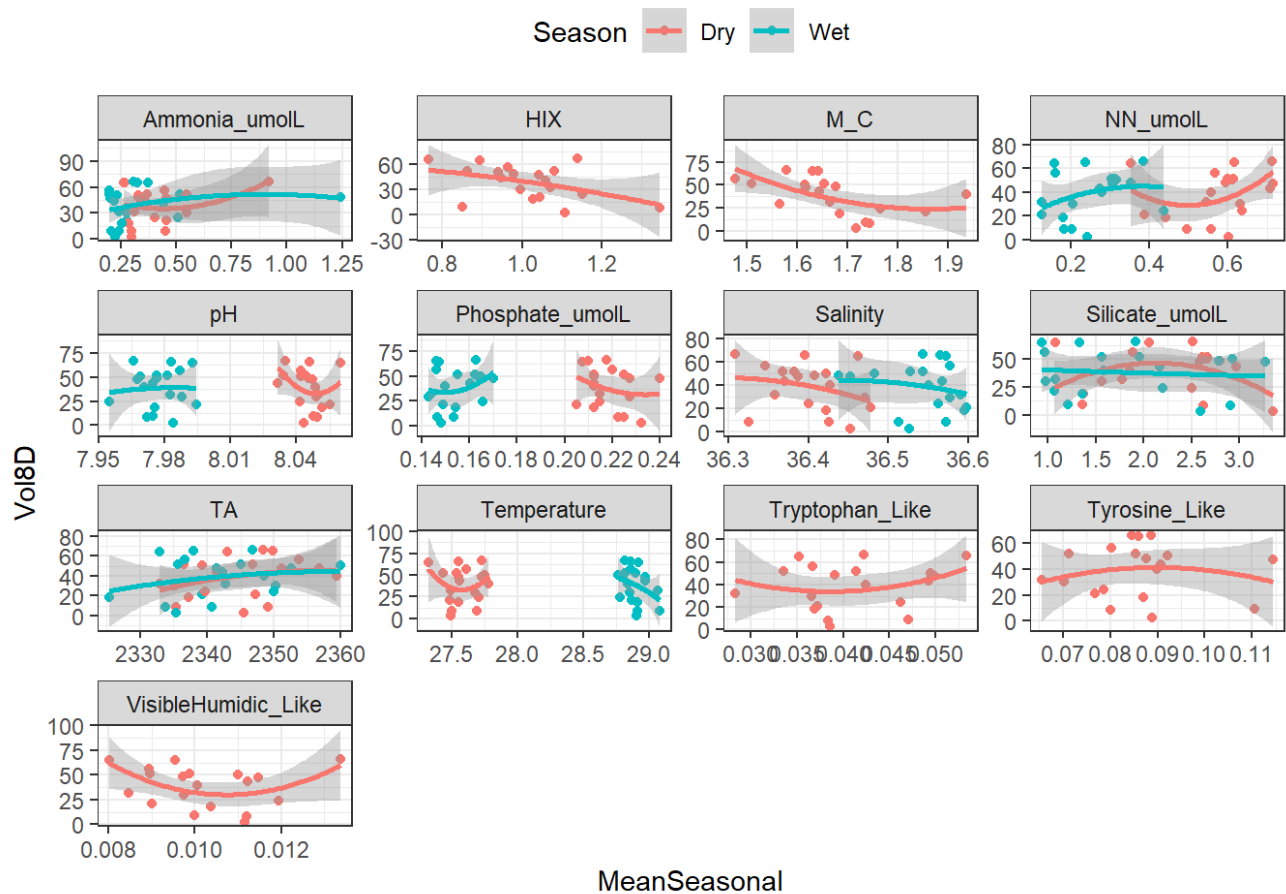
```
plotfun(y = NbFEs, x = MeanSeasonal, myfacet = Parameters, myformula = "y~poly(x,2)")
```

```
plotfun(y = Vol8D, x = MeanSeasonal, myfacet = Parameters, myformula = "y~x")
```



```
plotfun(y = Vol8D, x = MeanSeasonal, myfacet = Parameters, myformula = "y~poly(x,2)")
```



```
# get pvalues
```

```
MeanS3<-pvalpoly(myparam = "MeanSeasonal", myseason = "Dry")
```

```
MeanS4<-pvalpoly(myparam = "MeanSeasonal", myseason = "Wet")
```

```
MeanS <- full_join(MeanS3, MeanS4)
```

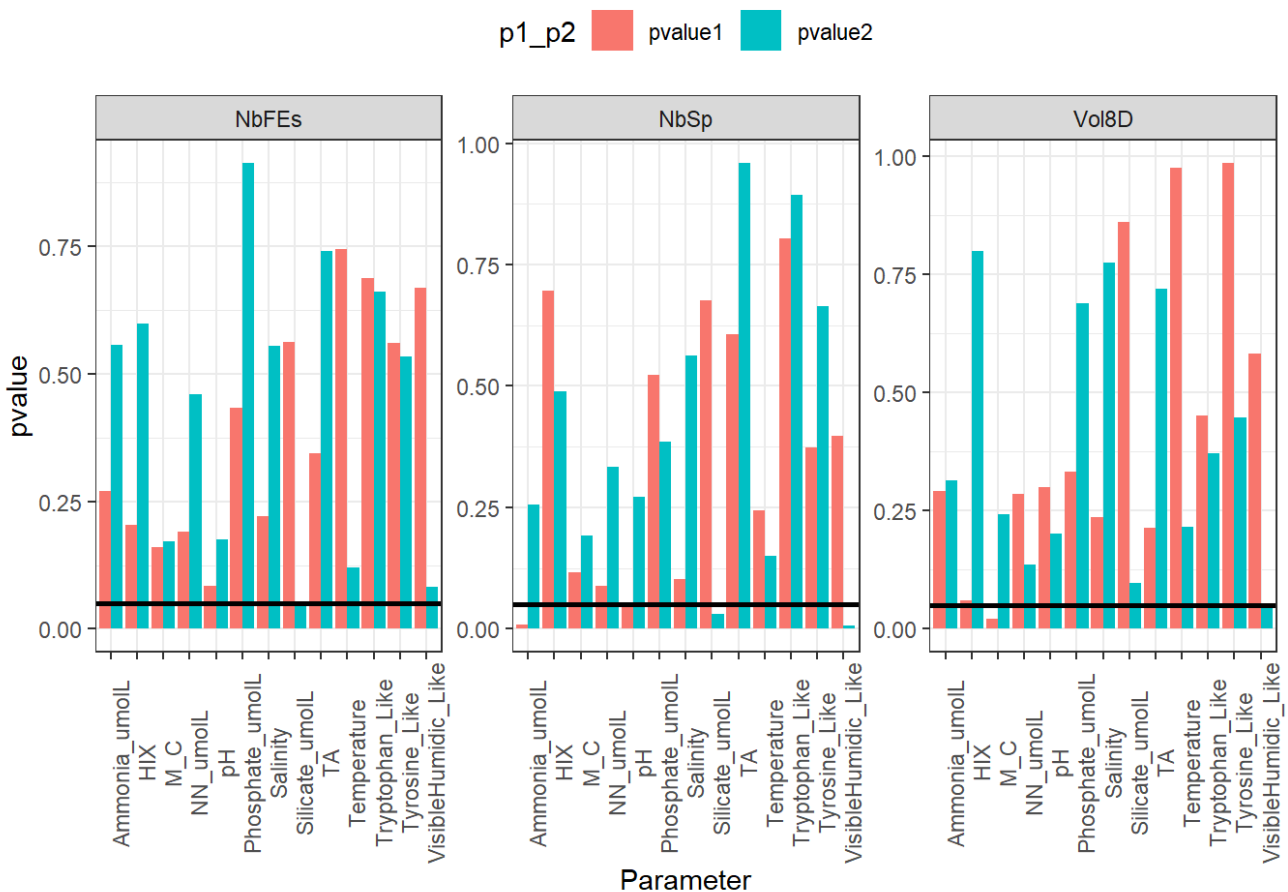
```
## Joining, by = c("Parameter", "ParamType", "Dependent", "Season", "pvalue1",
```

```
## "pvalue2", "r_squared", "adj_r_squared")
```

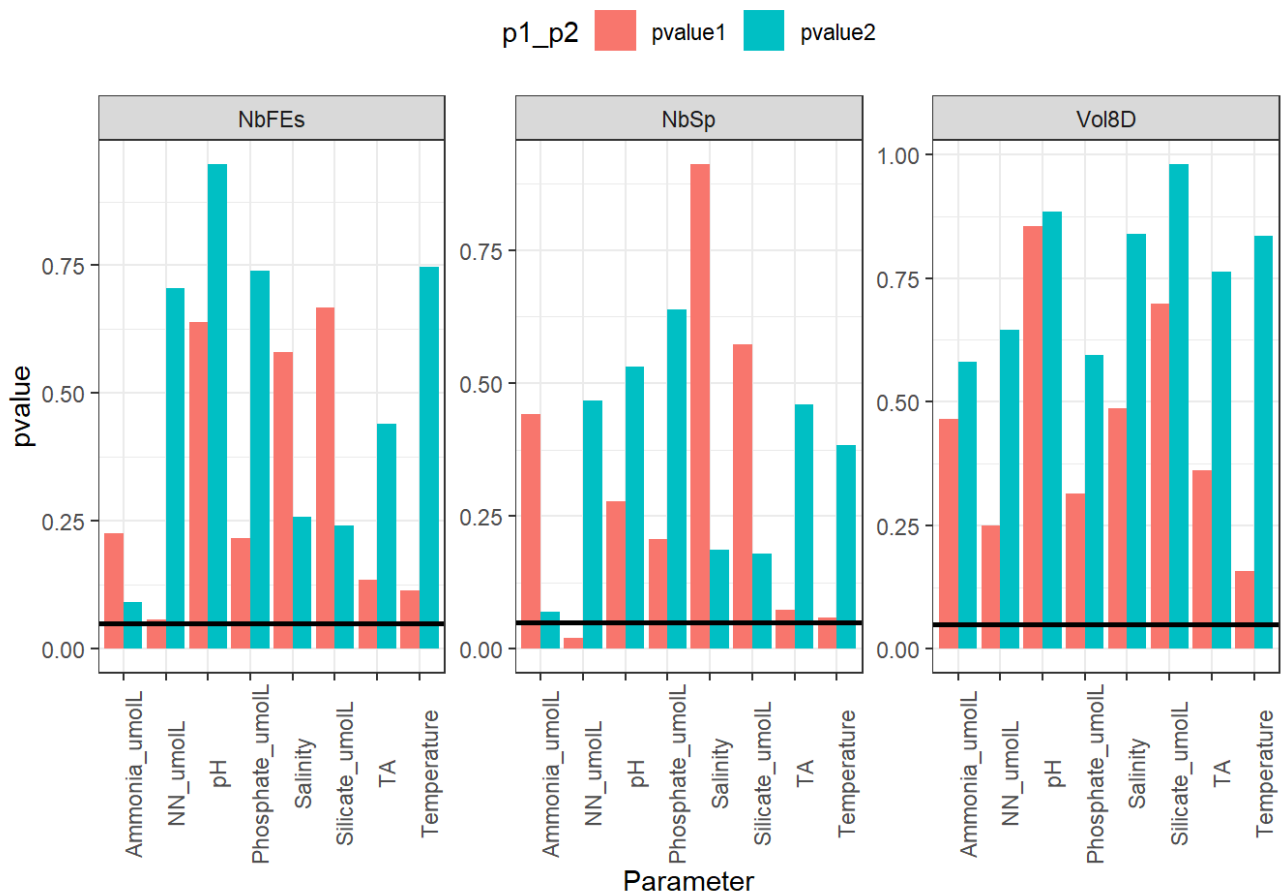
```
MeanS
```

```
## # A tibble: 105 × 8
##   Parameter      ParamType   Dependent Season pvalue1 pvalue2 r_squ...1 adj_r...2
##   <chr>          <chr>       <chr>    <chr>    <dbl>  <dbl>  <dbl>    <dbl>
## 1 Salinity       MeanSeasonal NbSp      Dry      0.102   0.562   0.173   0.0701
## 2 Temperature    MeanSeasonal NbSp      Dry      0.244   0.151   0.190   0.0883
## 3 TA             MeanSeasonal NbSp      Dry      0.606   0.959   0.0172  -0.106
## 4 pH             MeanSeasonal NbSp      Dry      0.0445  0.272   0.274   0.184
## 5 Phosphate_umolL MeanSeasonal NbSp      Dry      0.524   0.386   0.0709 -0.0453
## 6 Silicate_umolL  MeanSeasonal NbSp      Dry      0.676   0.0301  0.268   0.176
## 7 NN_umolL       MeanSeasonal NbSp      Dry      0.0890  0.334   0.211   0.112
## 8 Ammonia_umolL   MeanSeasonal NbSp      Dry      0.00863 0.256   0.392   0.317
## 9 M_C            MeanSeasonal NbSp      Dry      0.116   0.192   0.224   0.127
## 10 HIX            MeanSeasonal NbSp      Dry      0.696   0.488   0.0397 -0.0804
## # ... with 95 more rows, and abbreviated variable names 1r_squared,
## # 2adj_r_squared
```

```
pvalplot(mydata = MeanS, season = "Dry")
```



```
pvalplot(mydata = MeanS, season = "Wet")
```



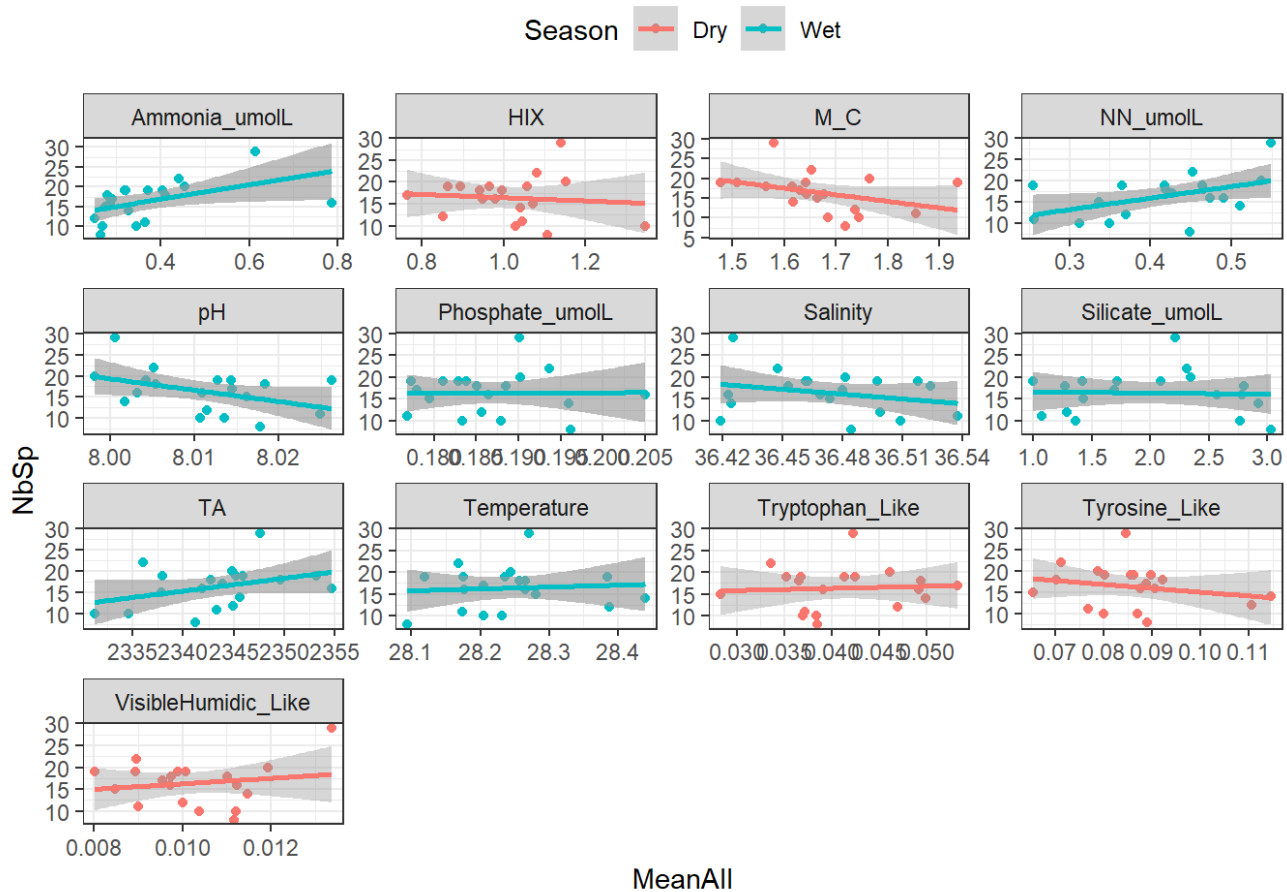
```
p4 <- paramPlot(mydata = Full_data, ParamType = MeanSeasonal, PTname = "MeanSeasonal")
```

```
## NULL
## NULL
```

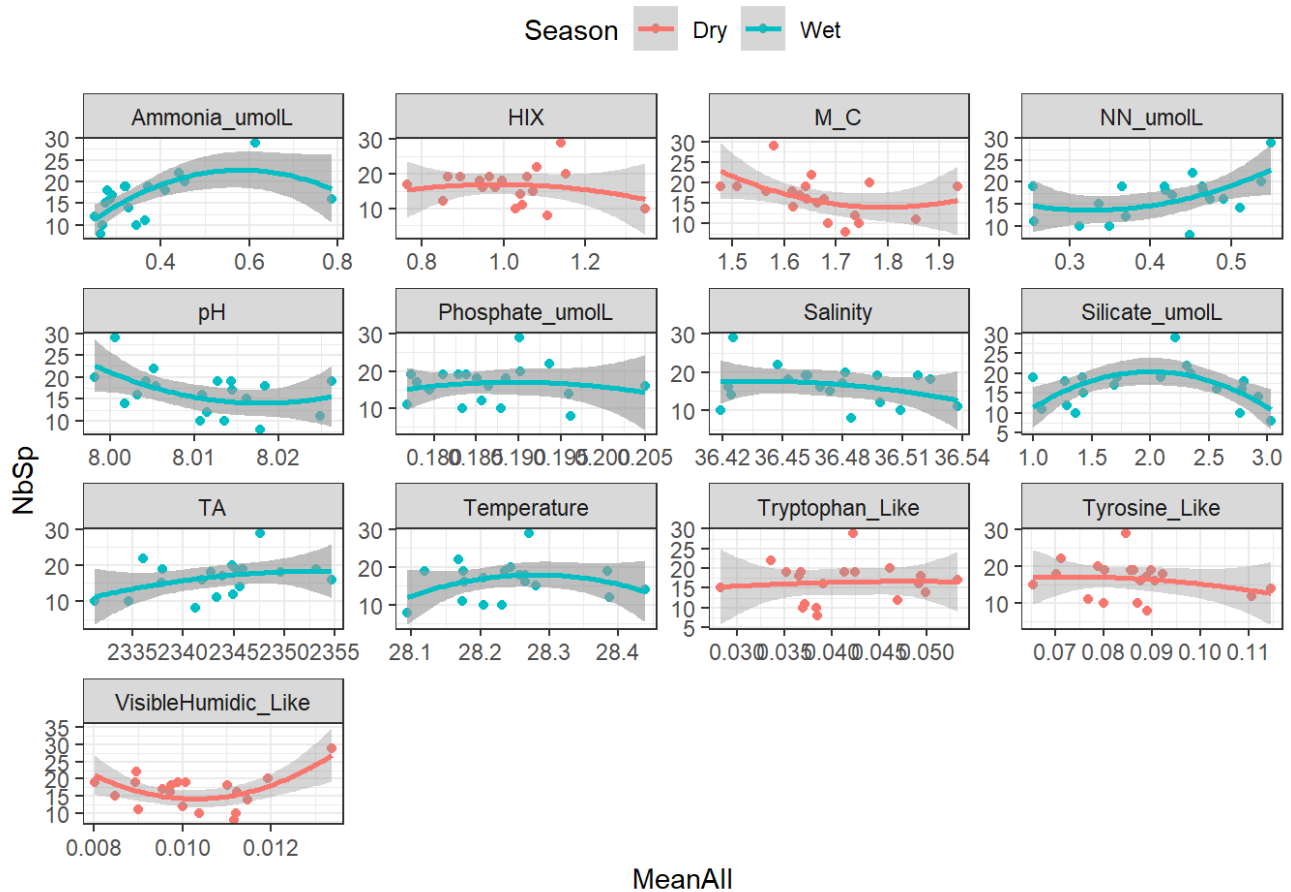
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```
## NULL
## NULL
```

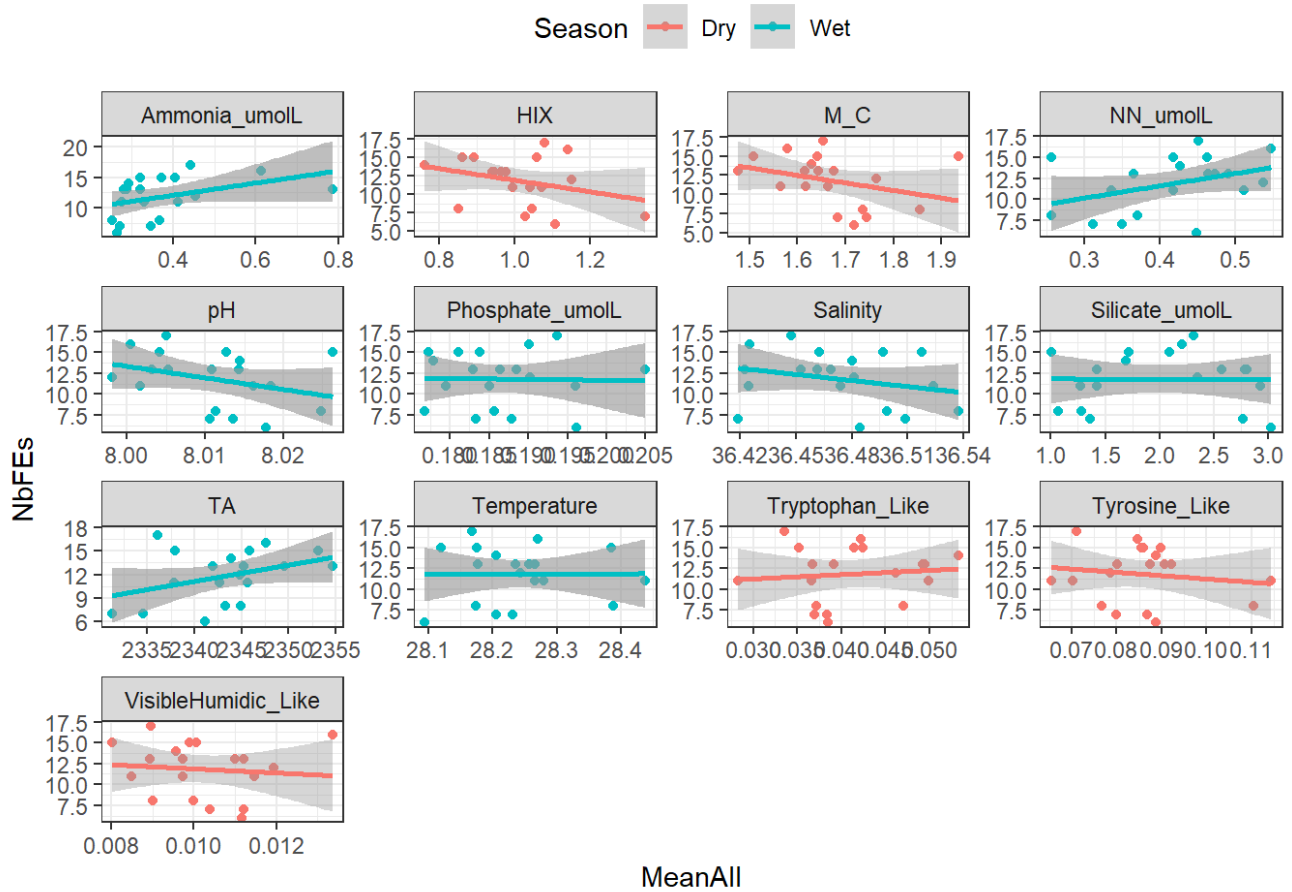
```
plotfun(y = NbSp, x = MeanAll, myfacet = Parameters, myformula = "y~x")
```



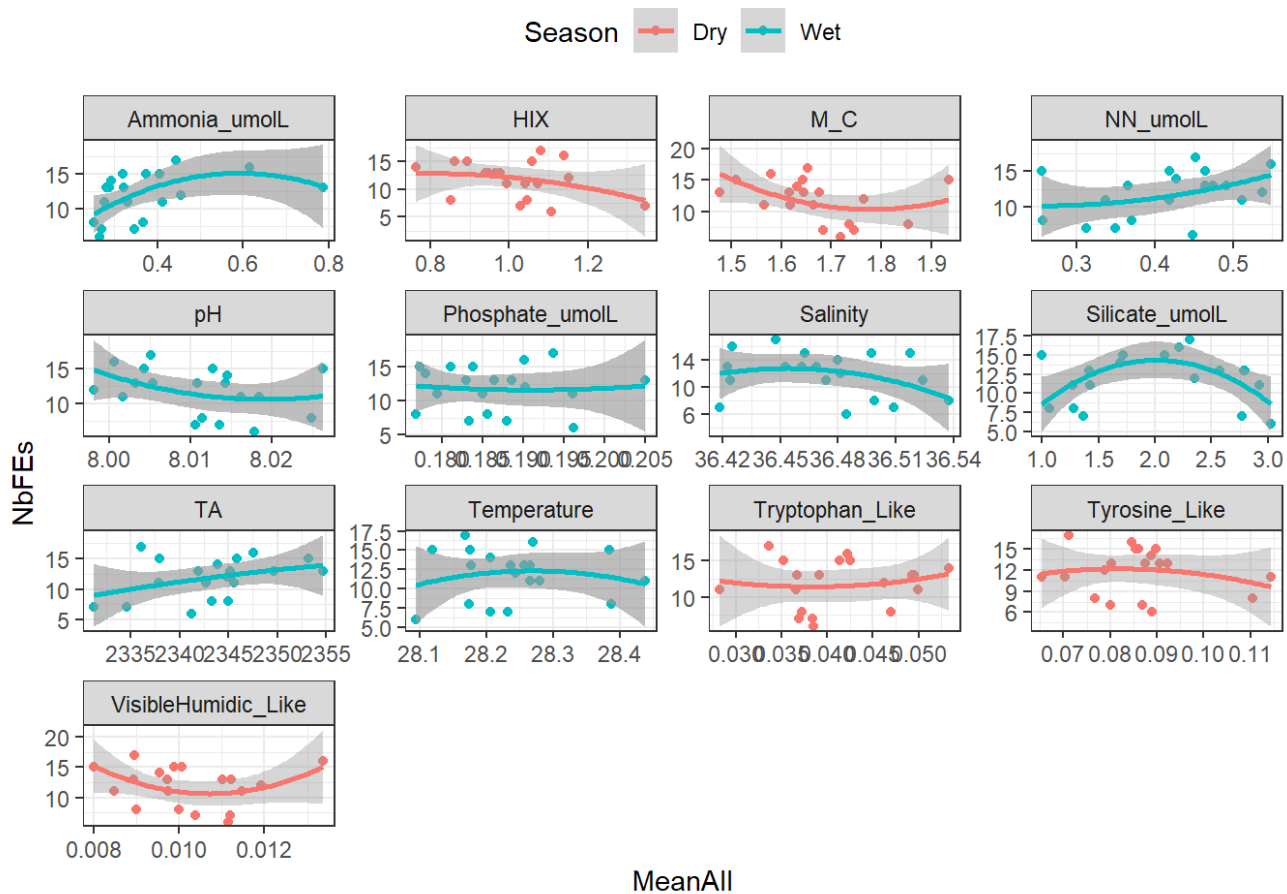
```
plotfun(y = NbSp, x = MeanAll, myfacet = Parameters, myformula = "y~poly(x,2)")
```

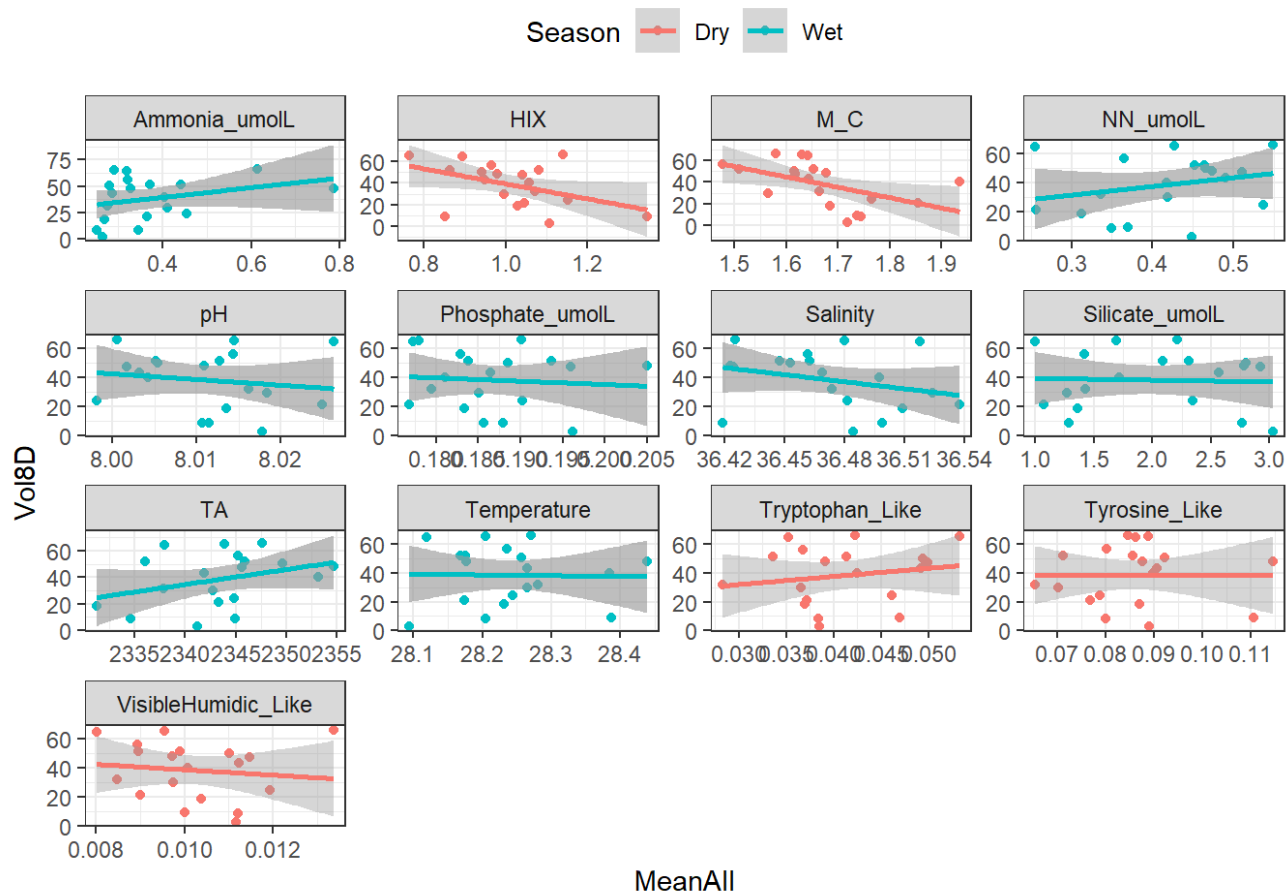
```
plotfun(y = NbFEs, x = MeanAll, myfacet = Parameters, myformula = "y~x")
```



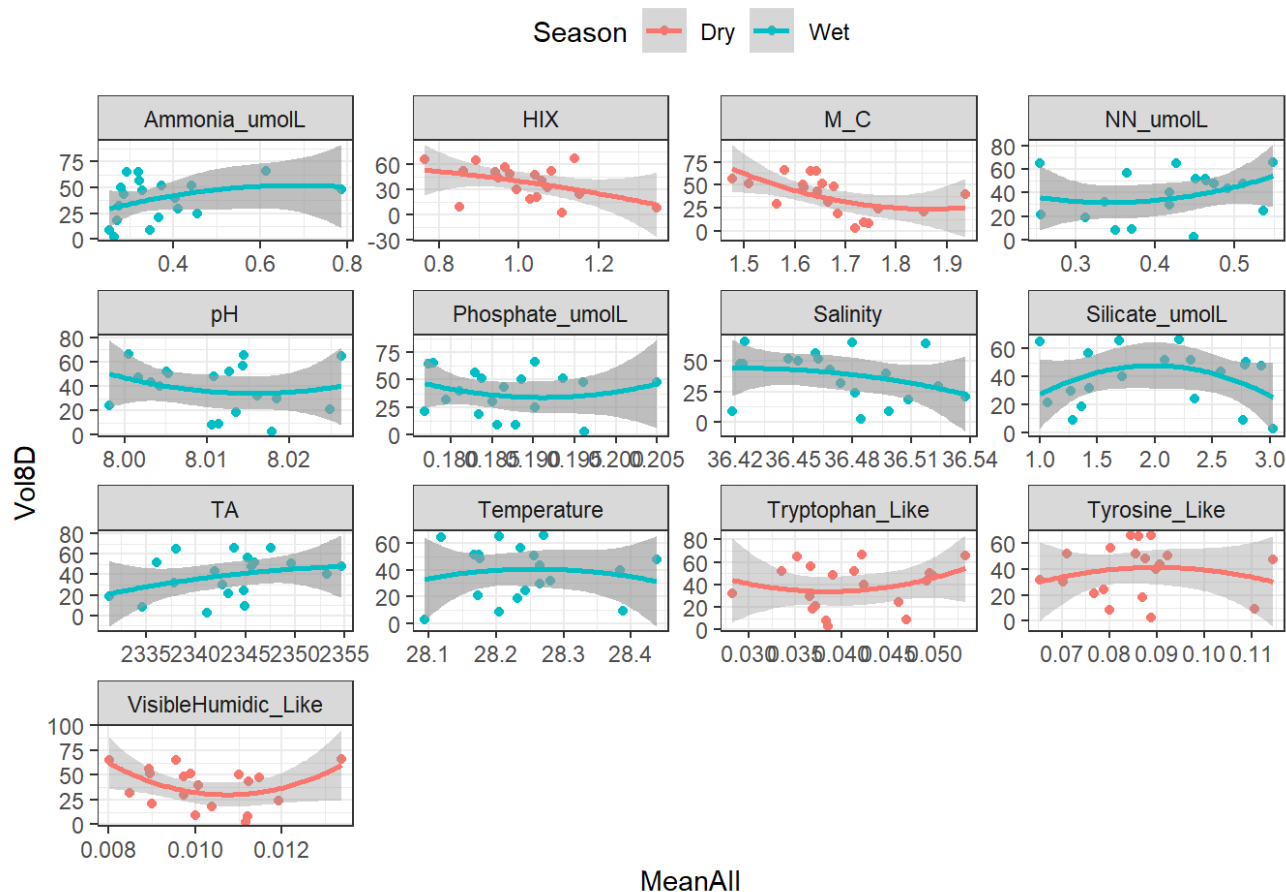
```
plotfun(y = NbFEs, x = MeanAll, myfacet = Parameters, myformula = "y~poly(x,2)")
```



```
plotfun(y = Vol8D, x = MeanAll, myfacet = Parameters, myformula = "y~x")
```



```
plotfun(y = Vol8D, x = MeanAll, myfacet = Parameters, myformula = "y~poly(x,2)")
```



```
# get pvalues
```

```
MeanA3<-pvalpoly(myparam = "MeanAll", myseason = "Dry")
```

```
MeanA4<-pvalpoly(myparam = "MeanAll", myseason = "Wet")
```

```
MeanA <- full_join(MeanA3, MeanA4)
```

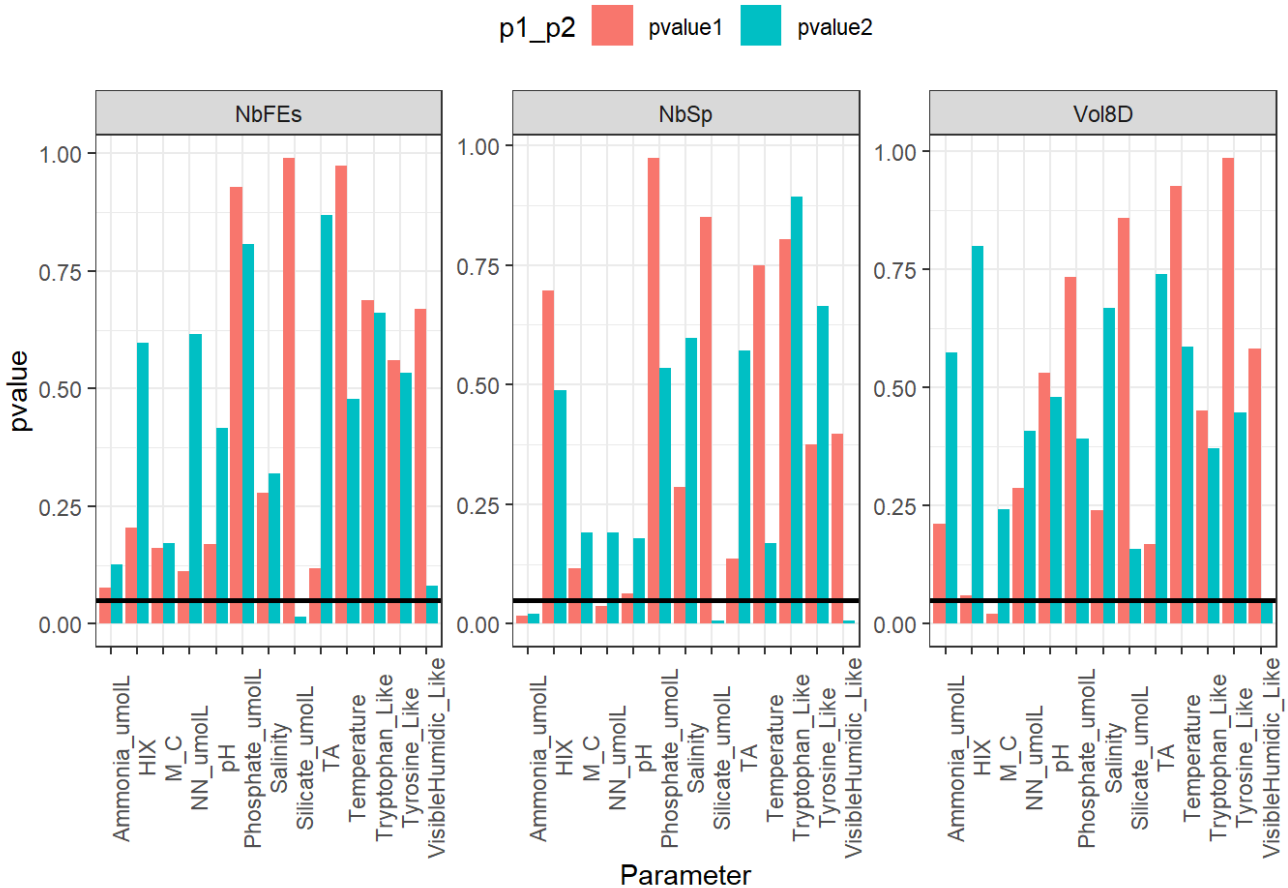
```
## Joining, by = c("Parameter", "ParamType", "Dependent", "Season", "pvalue1",
```

```
## "pvalue2", "r_squared", "adj_r_squared")
```

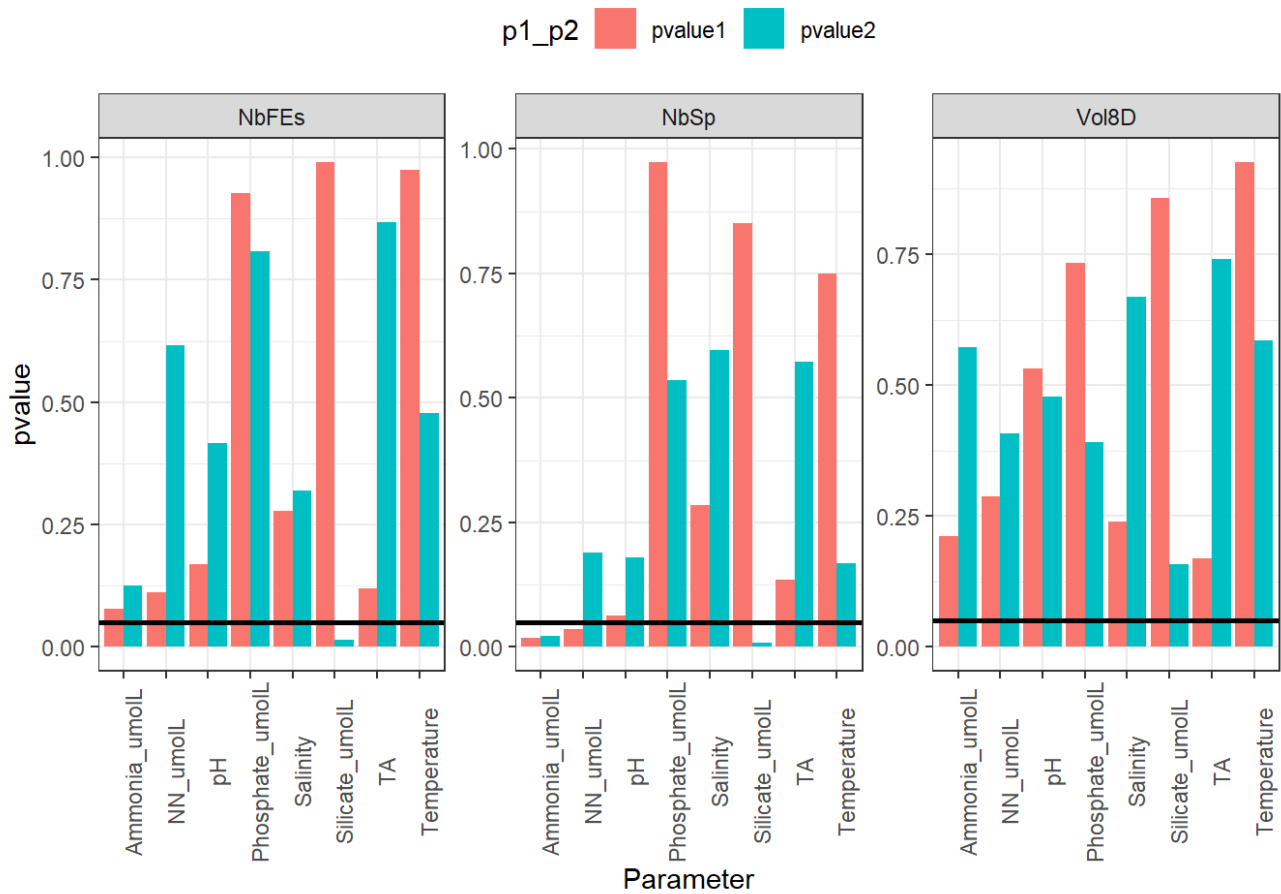
```
MeanA
```

```
## # A tibble: 105 × 8
##   Parameter      ParamType Dependent Season pvalue1 pvalue2 r_squared adj_r_...1
##   <chr>          <chr>      <chr>    <chr>    <dbl>   <dbl>   <dbl>   <dbl>
## 1 Salinity       MeanAll    NbSp     Dry      0.285   0.597    0.0865  -0.0277
## 2 Temperature    MeanAll    NbSp     Dry      0.749   0.168    0.120    0.0104
## 3 TA             MeanAll    NbSp     Dry      0.136   0.572    0.149    0.0425
## 4 pH             MeanAll    NbSp     Dry      0.0635  0.180    0.271    0.180
## 5 Phosphate_umolL MeanAll    NbSp     Dry      0.973   0.535    0.0246  -0.0973
## 6 Silicate_umolL  MeanAll    NbSp     Dry      0.851   0.00769  0.368    0.289
## 7 NN_umolL       MeanAll    NbSp     Dry      0.0363  0.190    0.307    0.221
## 8 Ammonia_umolL   MeanAll    NbSp     Dry      0.0177  0.0219   0.456    0.388
## 9 M_C            MeanAll    NbSp     Dry      0.116   0.192    0.224    0.127
## 10 HIX            MeanAll    NbSp     Dry      0.696   0.488    0.0397  -0.0804
## # ... with 95 more rows, and abbreviated variable name 1adj_r_squared
```

```
pvalplot(mydata = MeanA, season = "Dry")
```



```
pvalplot(mydata = MeanA, season = "Wet")
```



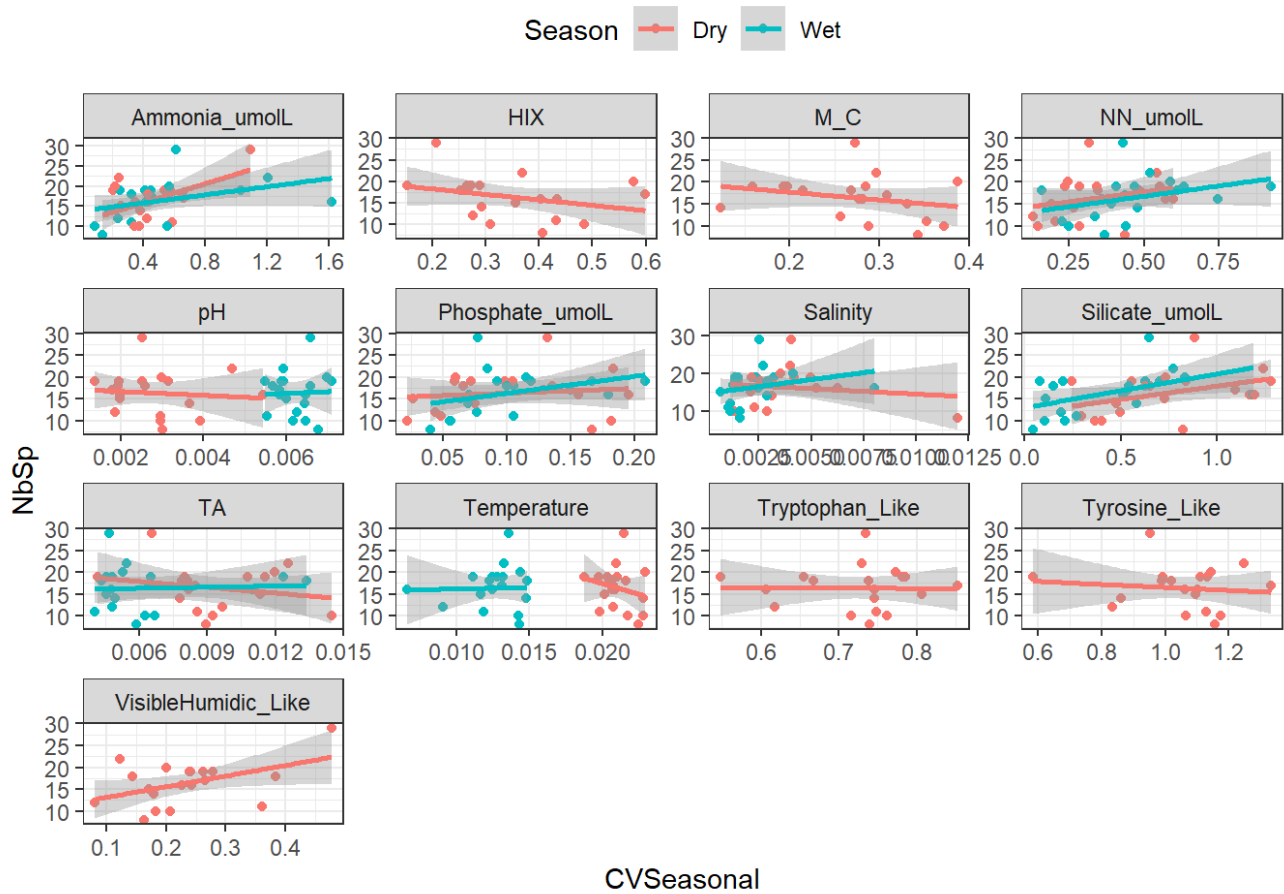
```
p5 <- paramPlot(mydata = Full_data, ParamType = MeanAll, PTname = "MeanAll")
```

```
## NULL
## NULL
```

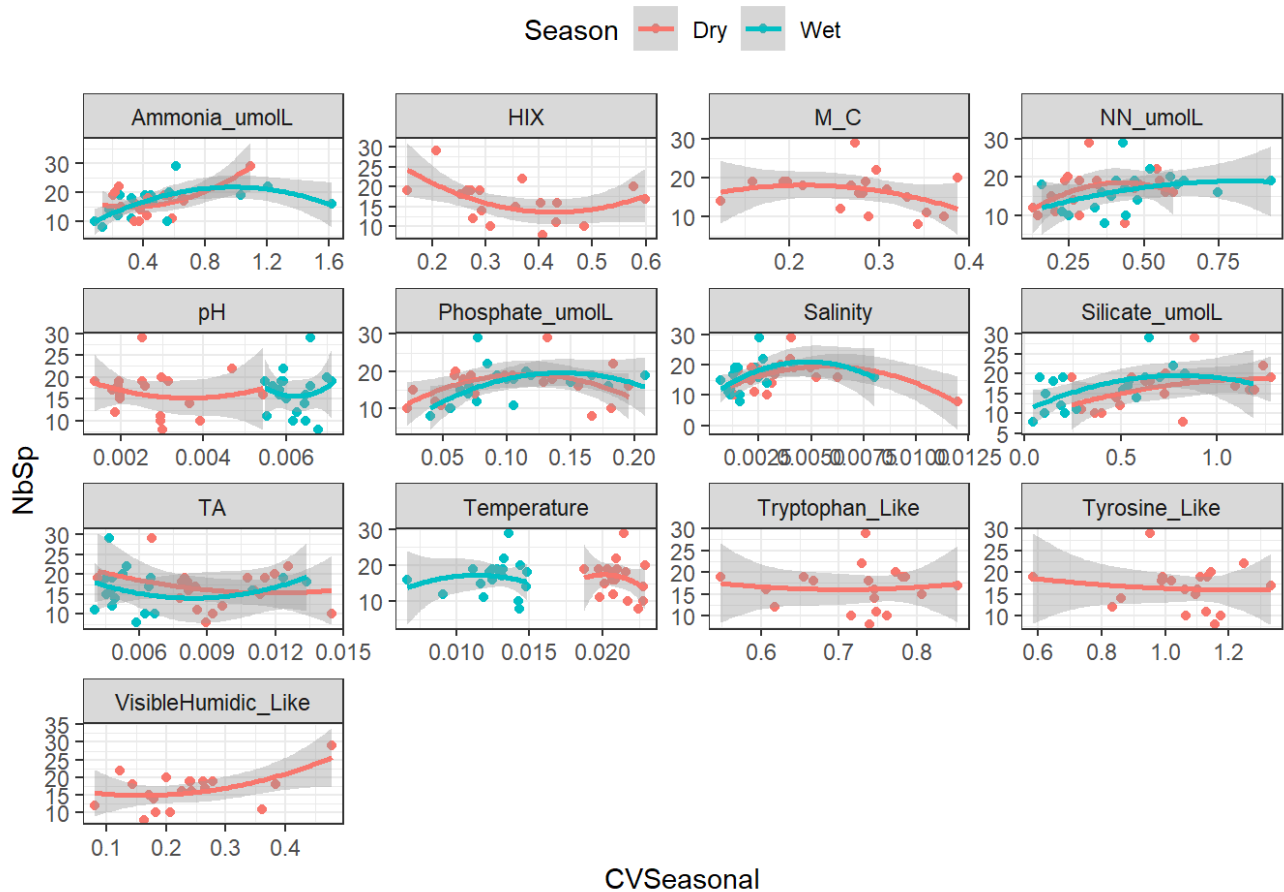
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```
## NULL
## NULL
```

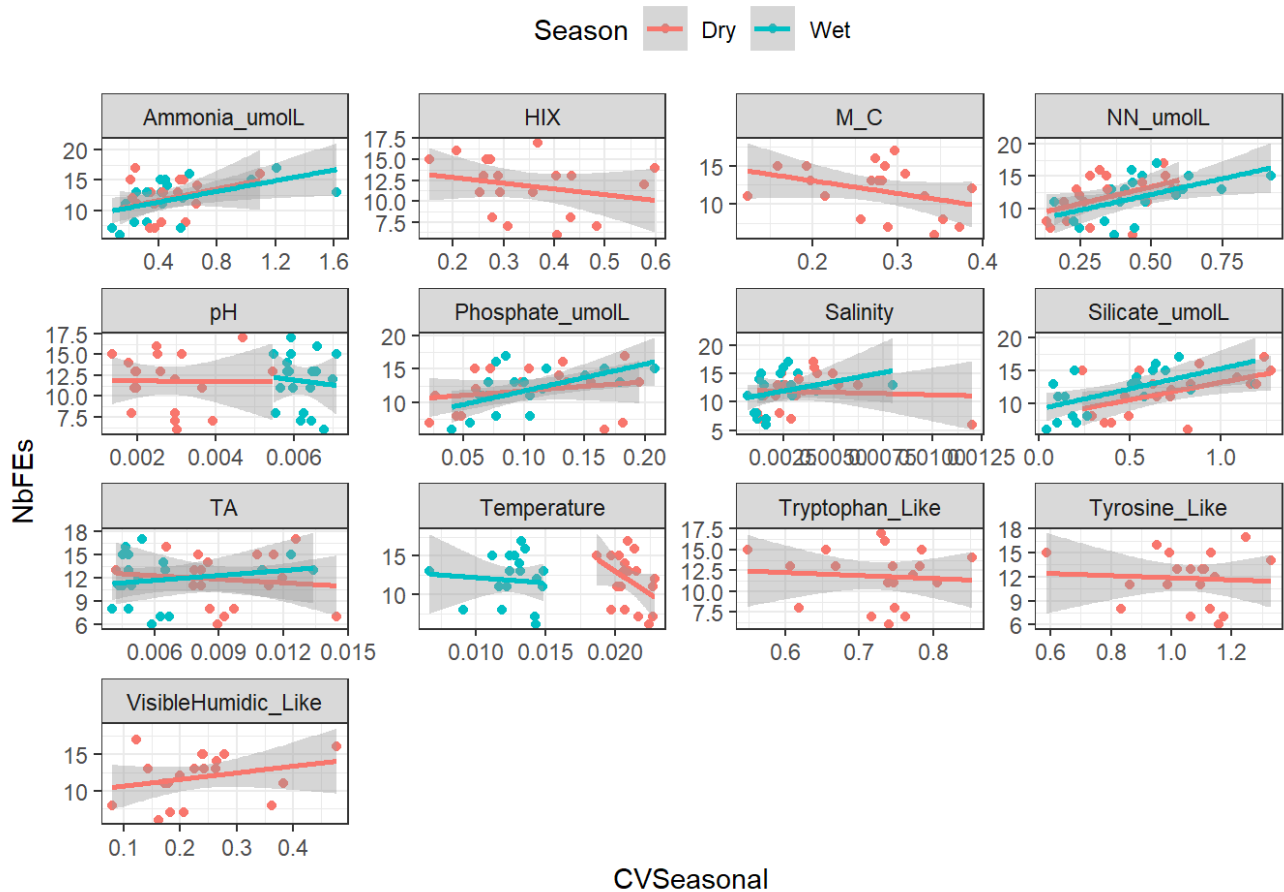
```
plotfun(y = NbSp, x = CVSeasonal, myfacet = Parameters, myformula = "y~x")
```



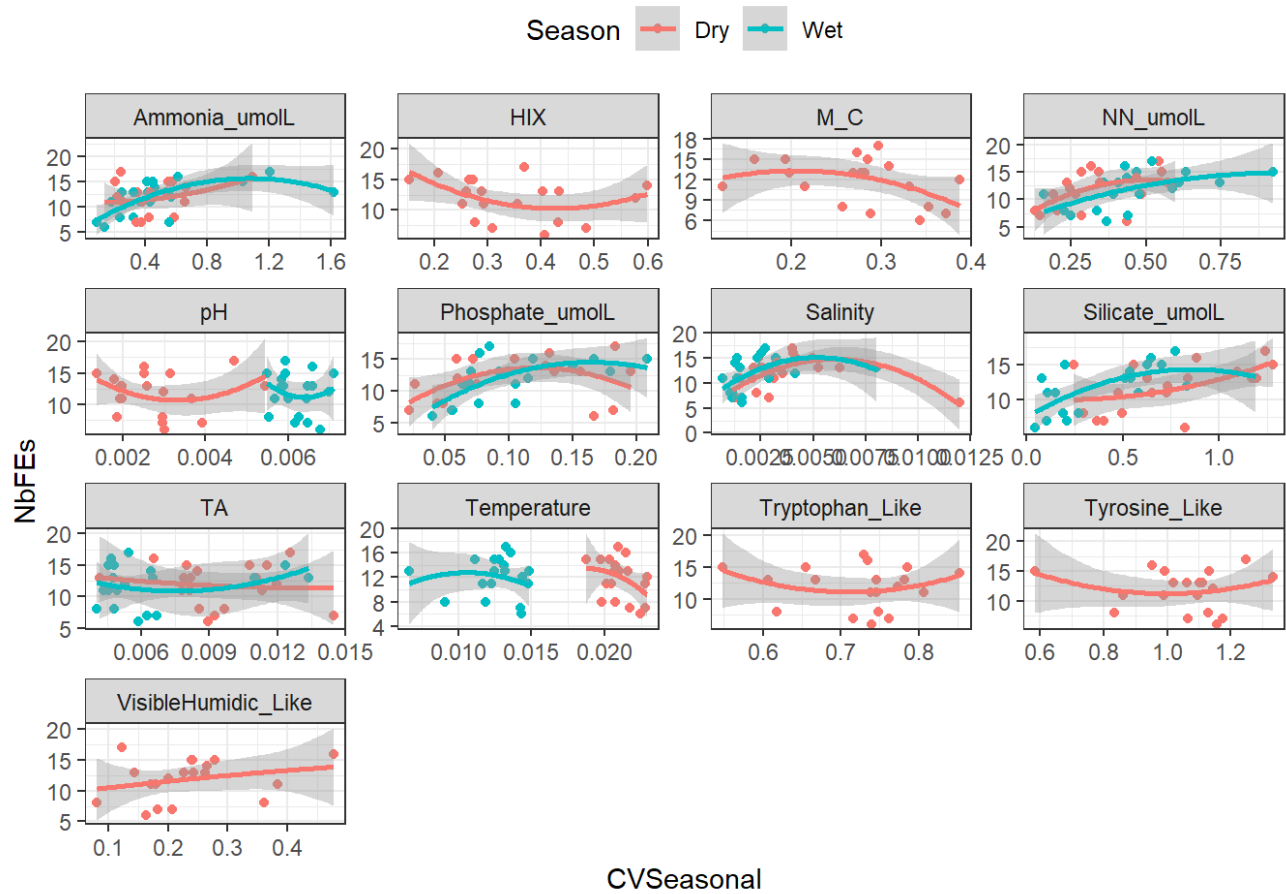
```
plotfun(y = NbSp, x = CVSeasonal, myfacet = Parameters, myformula = "y~poly(x,2)")
```



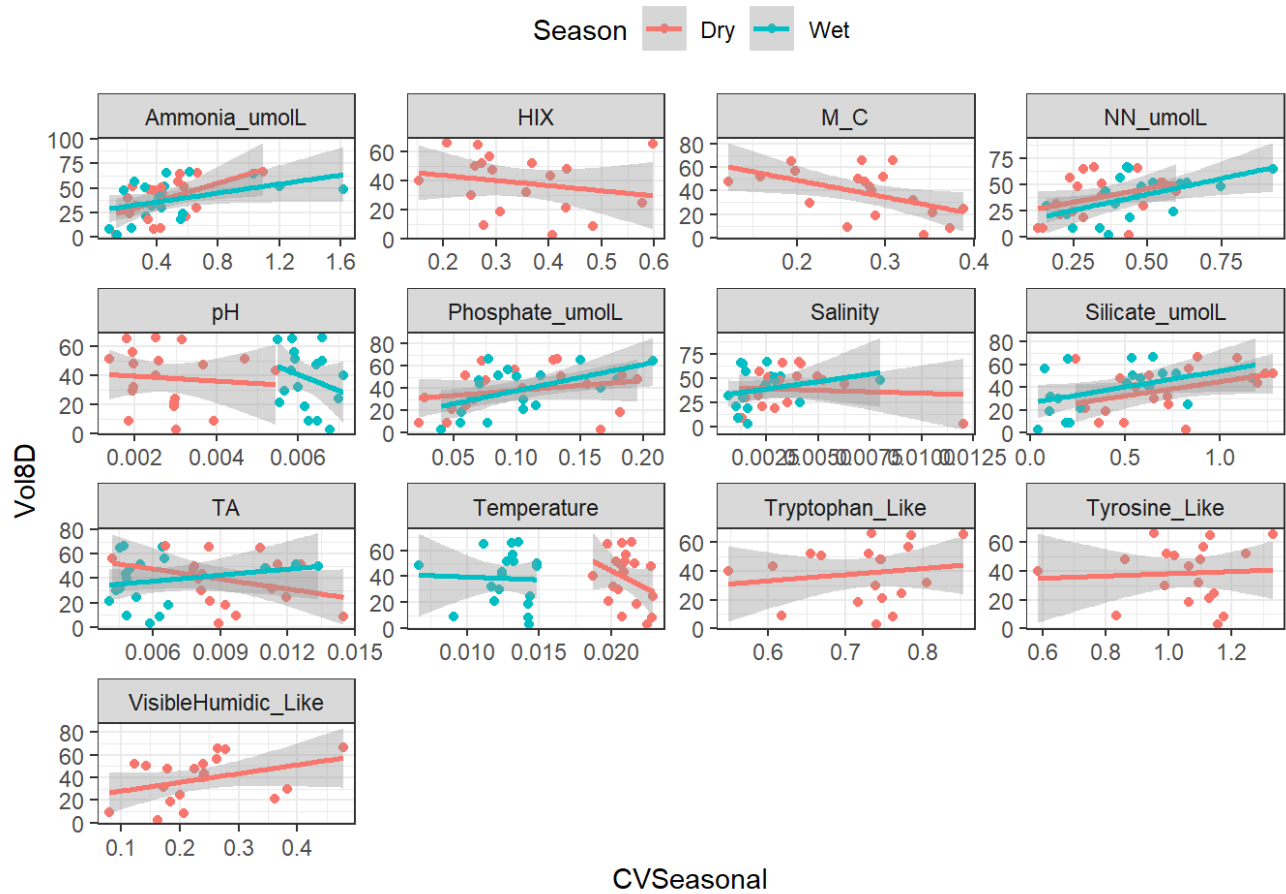
```
plotfun(y = NbFEs, x = CVSeasonal, myfacet = Parameters, myformula = "y~x")
```



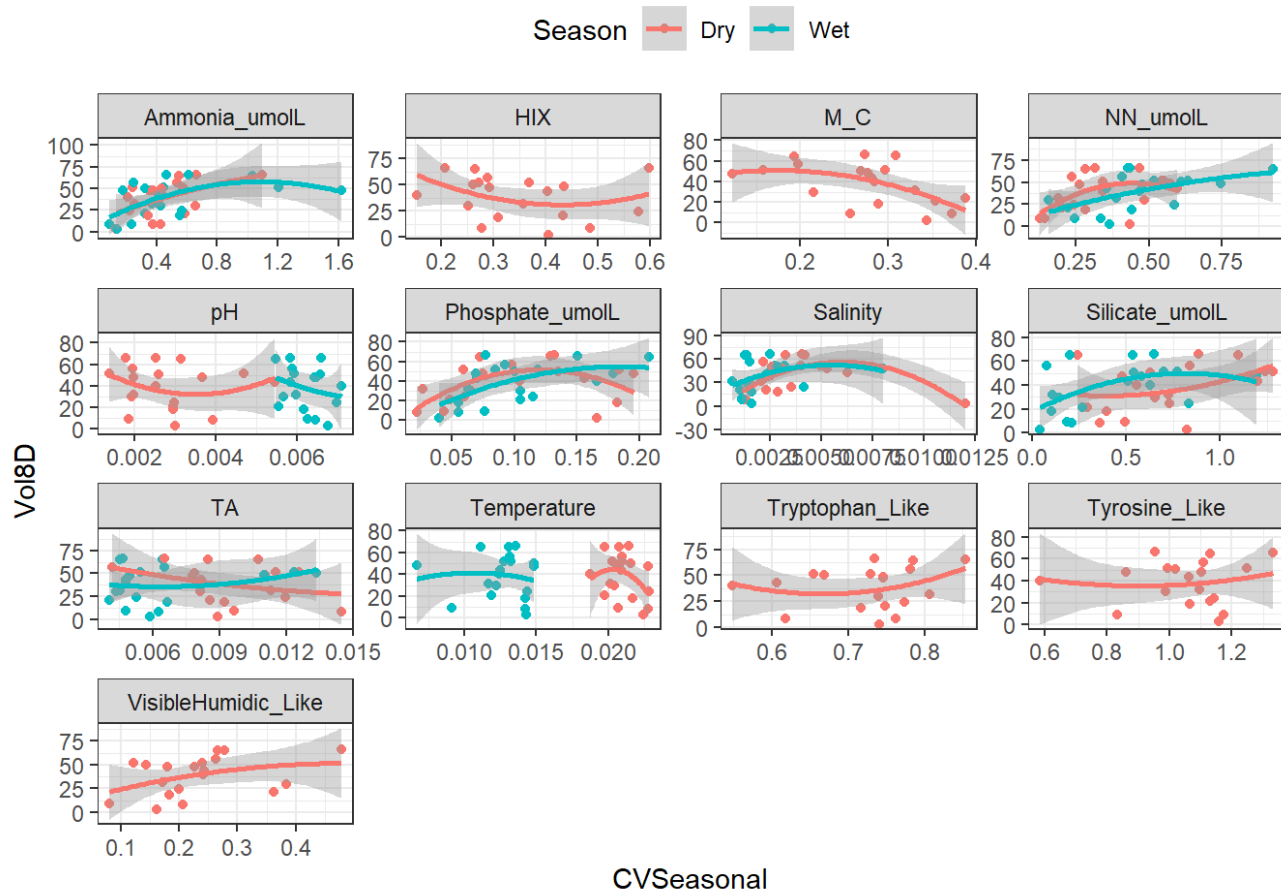
```
plotfun(y = NbFEs, x = CVSeasonal, myfacet = Parameters, myformula = "y~poly(x,2)")
```



```
plotfun(y = Vol8D, x = CVSeasonal, myfacet = Parameters, myformula = "y~x")
```



```
plotfun(y = Vol8D, x = CVSeasonal, myfacet = Parameters, myformula = "y~poly(x,2)")
```



```
# get pvalues
```

```
CVS3<-pvalpoly(myparam = "CVSeasonal", myseason = "Dry")
```

```
CVS4<-pvalpoly(myparam = "CVSeasonal", myseason = "Wet")
```

```
CVS <- full_join(CVS3, CVS4)
```

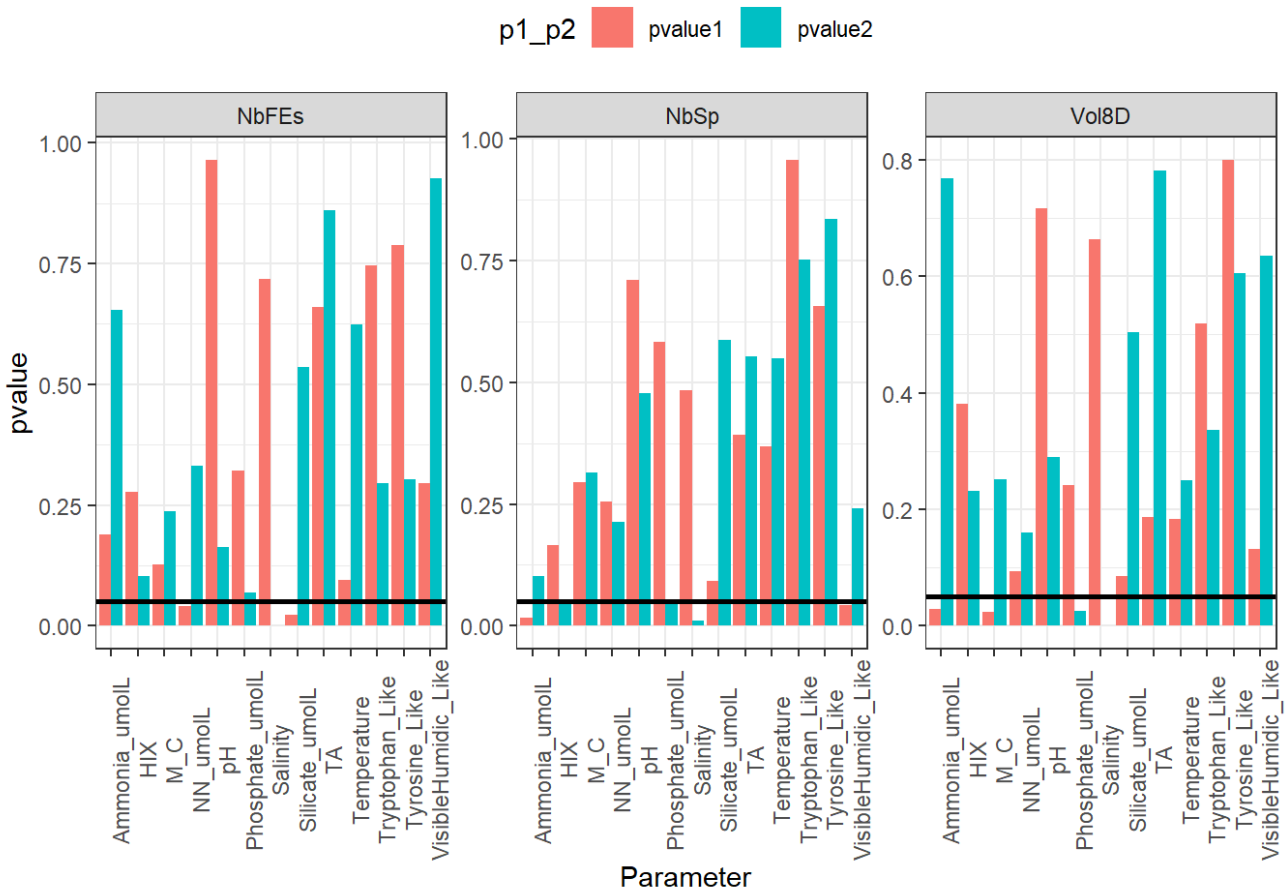
```
## Joining, by = c("Parameter", "ParamType", "Dependent", "Season", "pvalue1",
```

```
## "pvalue2", "r_squared", "adj_r_squared")
```

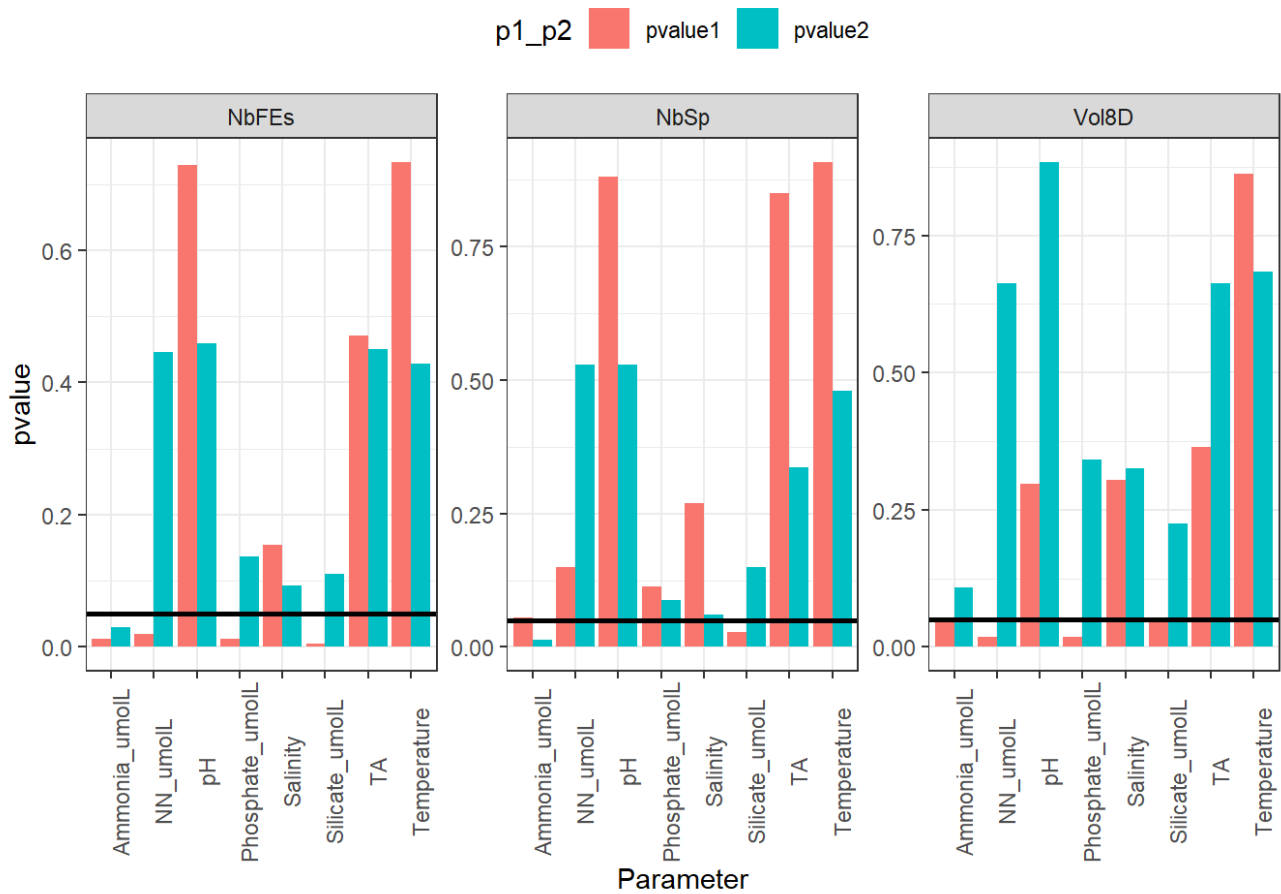
```
CVS
```

```
## # A tibble: 105 × 8
##   Parameter      ParamType Dependent Season pvalue1 pvalue2 r_squared adj_r...1
##   <chr>          <chr>      <chr>    <chr>    <dbl>   <dbl>   <dbl>   <dbl>
## 1 Salinity       CVSeasonal NbSp      Dry      0.484   0.0101   0.360   0.280
## 2 Temperature    CVSeasonal NbSp      Dry      0.368   0.548   0.0716  -0.0444
## 3 TA             CVSeasonal NbSp      Dry      0.393   0.553   0.0664  -0.0504
## 4 pH             CVSeasonal NbSp      Dry      0.710   0.477   0.0404  -0.0796
## 5 Phosphate_umolL CVSeasonal NbSp      Dry      0.583   0.0499   0.231   0.135
## 6 Silicate_umolL  CVSeasonal NbSp      Dry      0.0915  0.586   0.181   0.0784
## 7 NN_umolL       CVSeasonal NbSp      Dry      0.255   0.212   0.161   0.0566
## 8 Ammonia_umolL   CVSeasonal NbSp      Dry      0.0162  0.101   0.390   0.314
## 9 M_C            CVSeasonal NbSp      Dry      0.294   0.316   0.123   0.0136
## 10 HIX            CVSeasonal NbSp      Dry      0.166   0.0471  0.296   0.208
## # ... with 95 more rows, and abbreviated variable name 1adj_r_squared
```

```
pvalplot(mydata = CVS, season = "Dry")
```



```
pvalplot(mydata = CVS, season = "Wet")
```

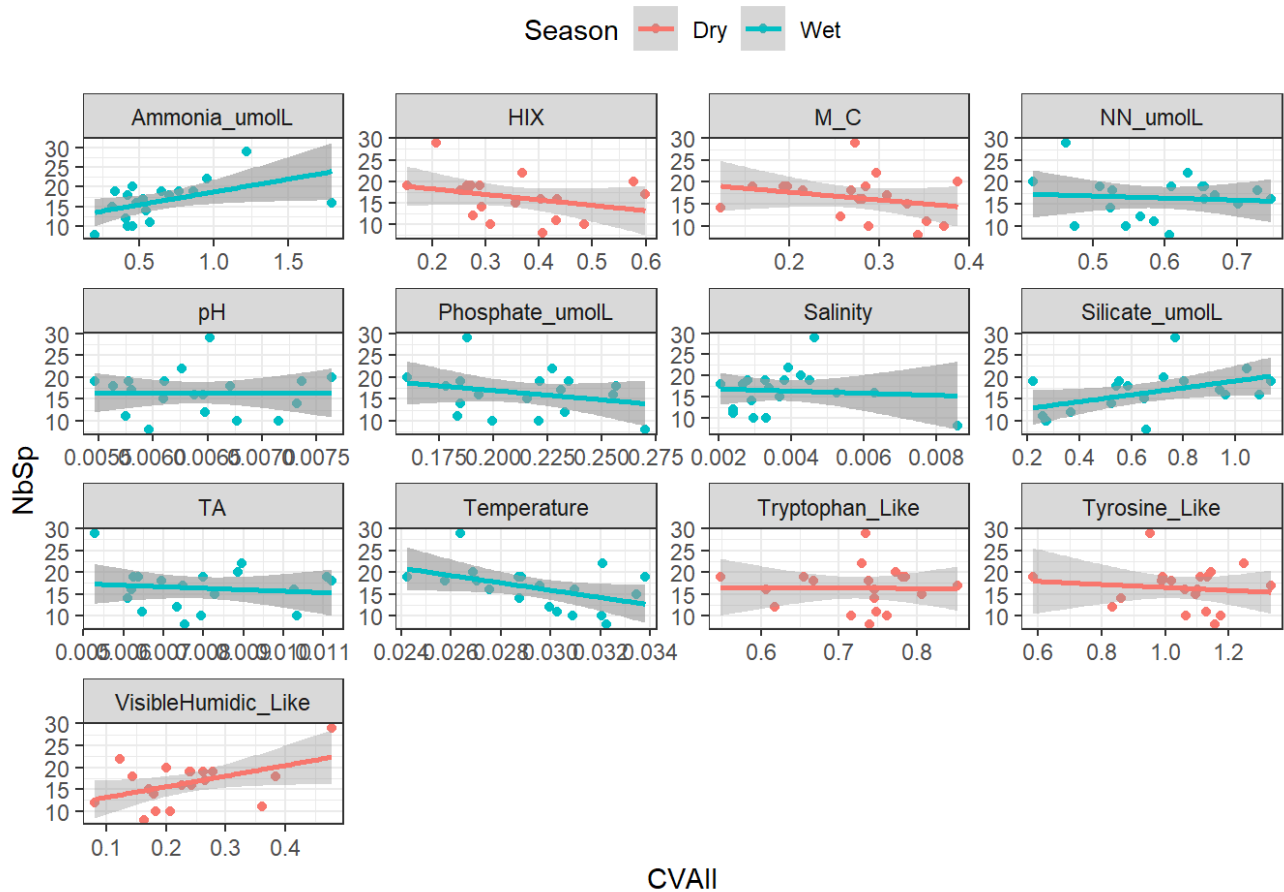
```
p6 <- paramPlot(mydata = Full_data, ParamType = CVSeasonal, PTname = "CVSeasonal")
```

```
## NULL
## NULL
```

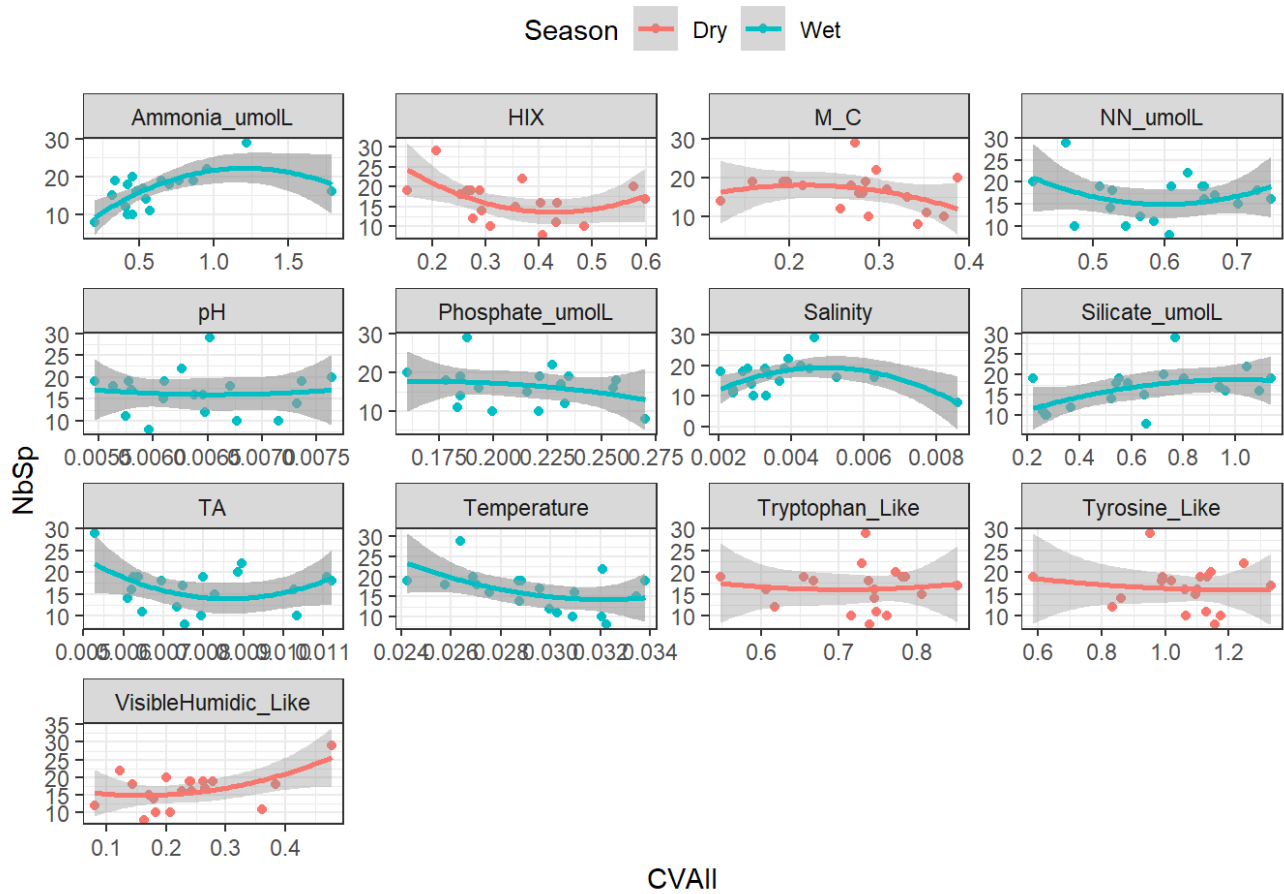
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```
## NULL
## NULL
```

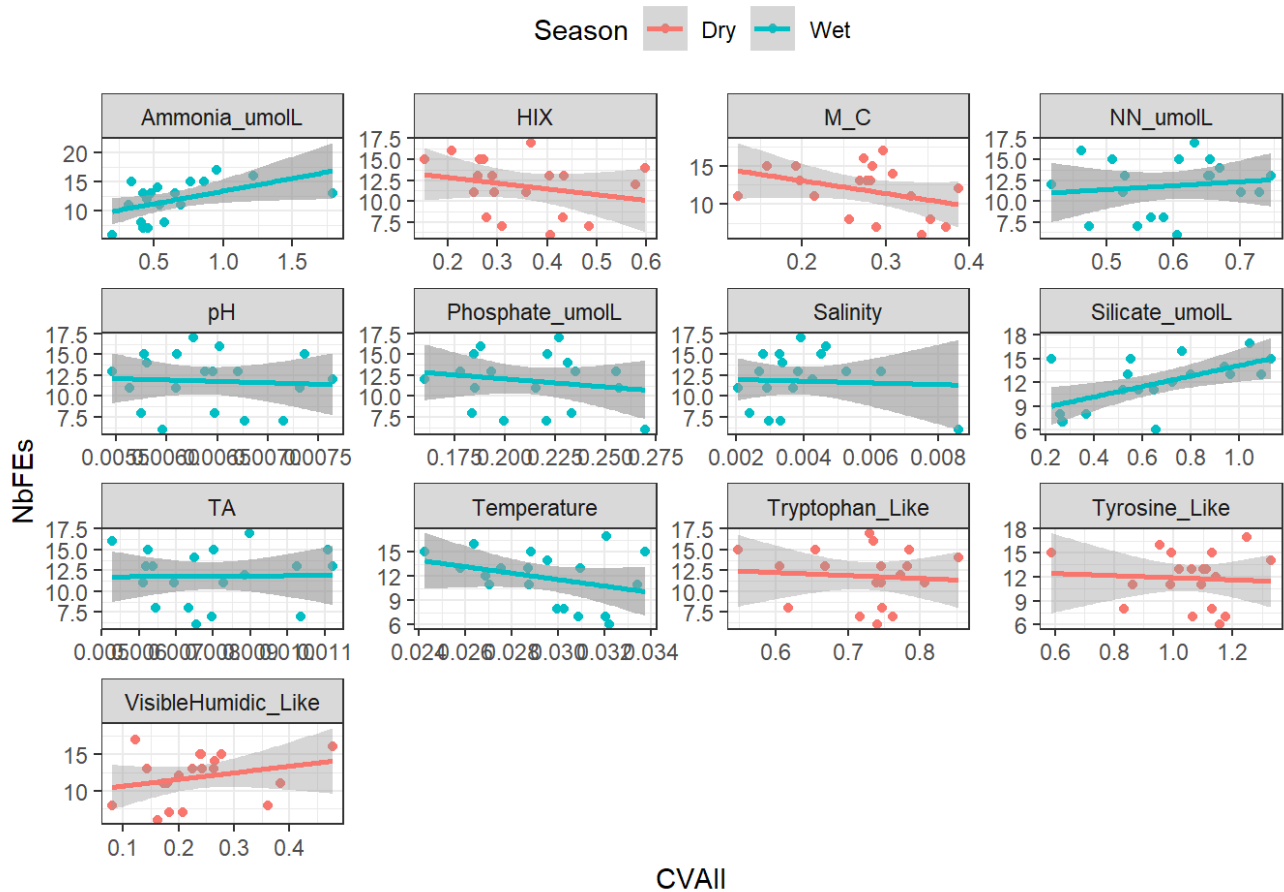
```
plotfun(y = NbSp, x = CVA11, myfacet = Parameters, myformula = "y~x")
```



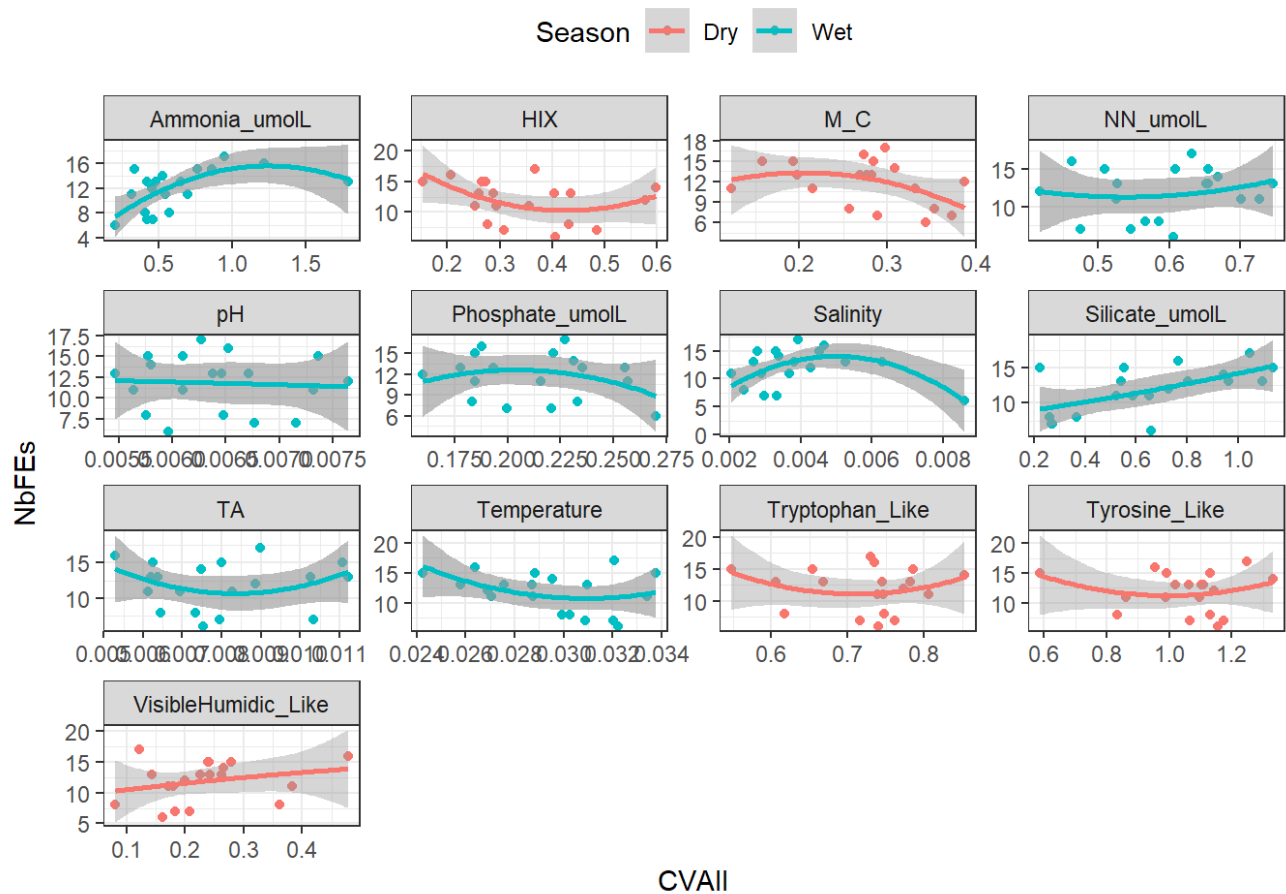
```
plotfun(y = NbSp, x = CVAII, myfacet = Parameters, myformula = "y~poly(x,2)")
```



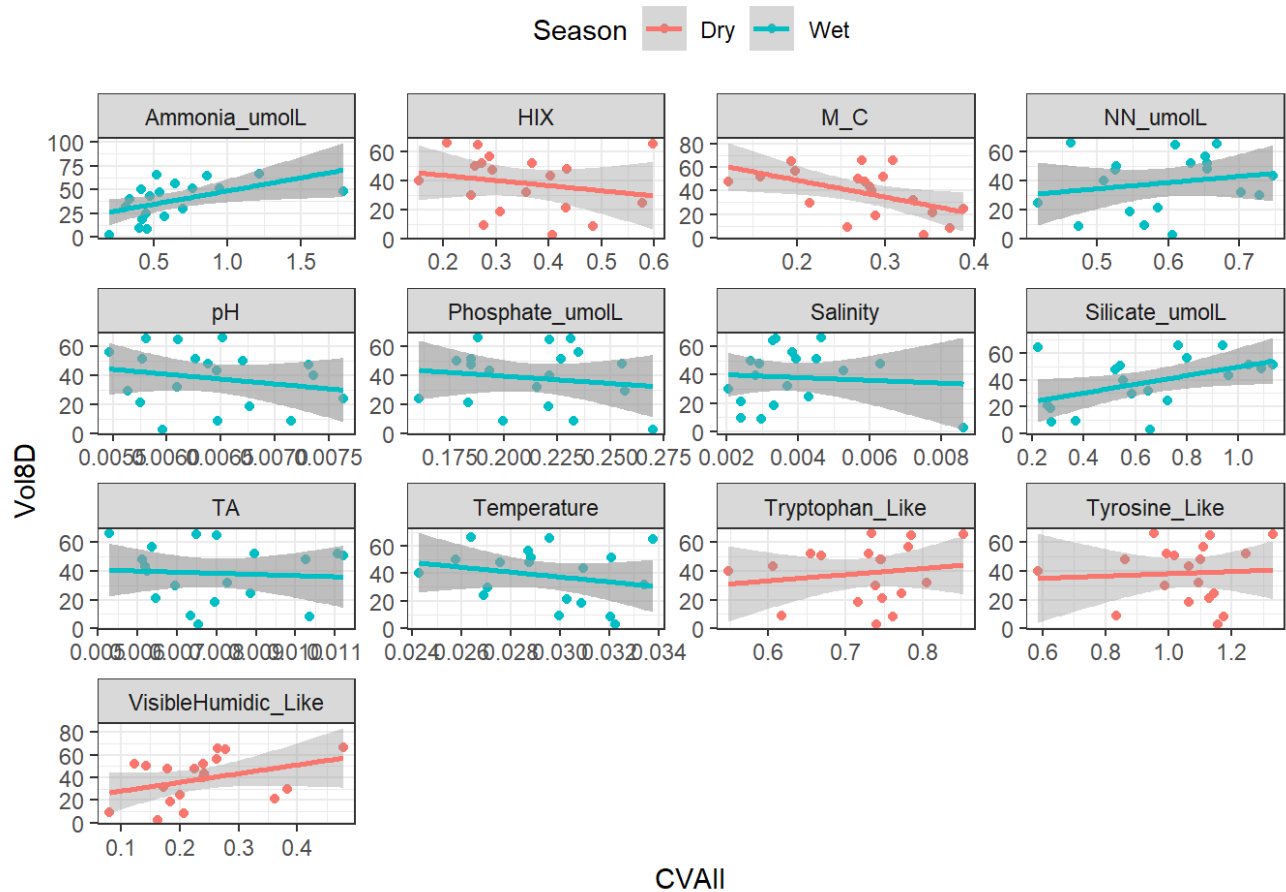
```
plotfun(y = NbFEs, x = CVAII, myfacet = Parameters, myformula = "y~x")
```



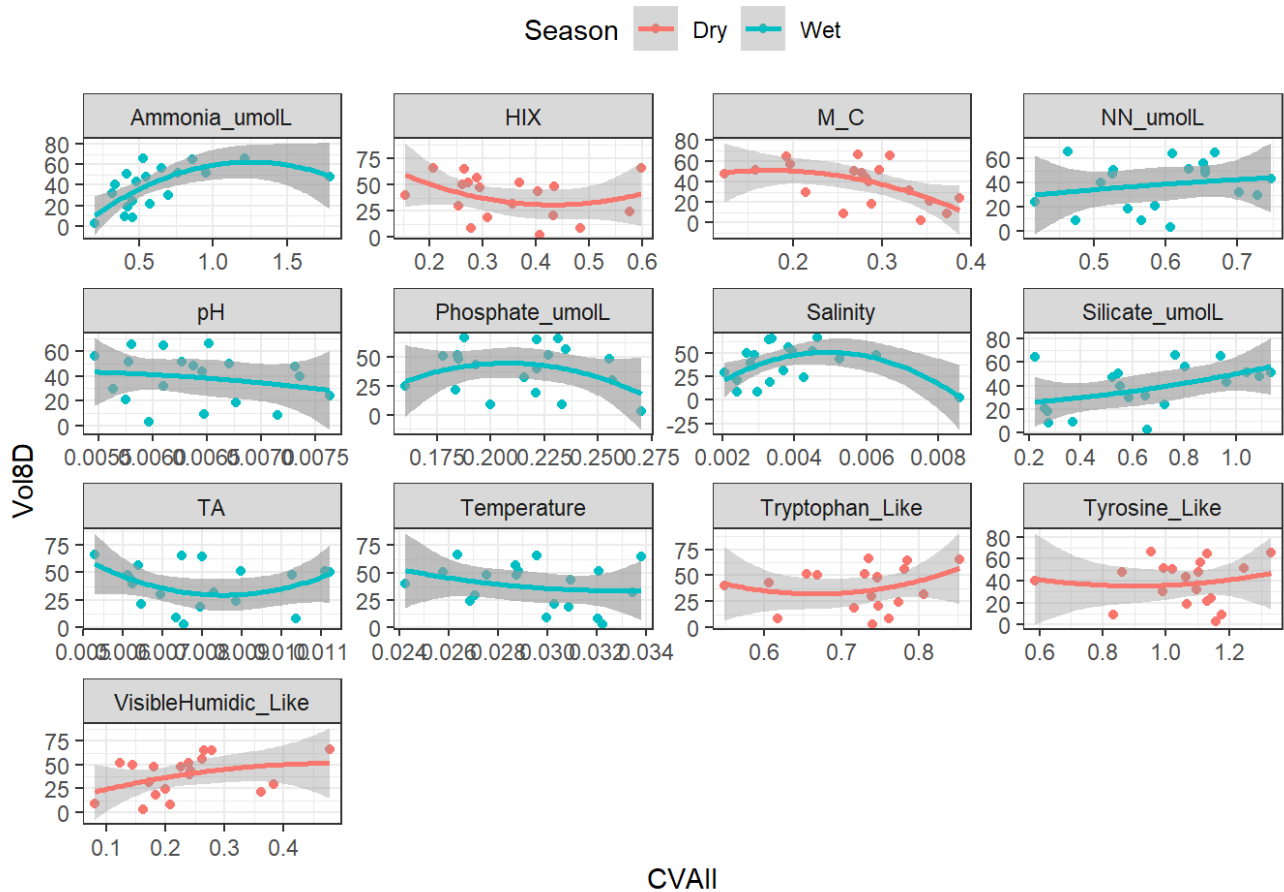
```
plotfun(y = NbFEs, x = CVAII, myfacet = Parameters, myformula = "y~poly(x,2)")
```



```
plotfun(y = Vol8D, x = CVAII, myfacet = Parameters, myformula = "y~x")
```



```
plotfun(y = Vol8D, x = CVAII, myfacet = Parameters, myformula = "y~poly(x,2)")
```

```
# get pvalues
```

```
CVA3<-pvalpoly(myparam = "CVA11", myseason = "Dry")
```

```
CVA4<-pvalpoly(myparam = "CVA11", myseason = "Wet")
```

```
CVA <- full_join(CVA3, CVA4)
```

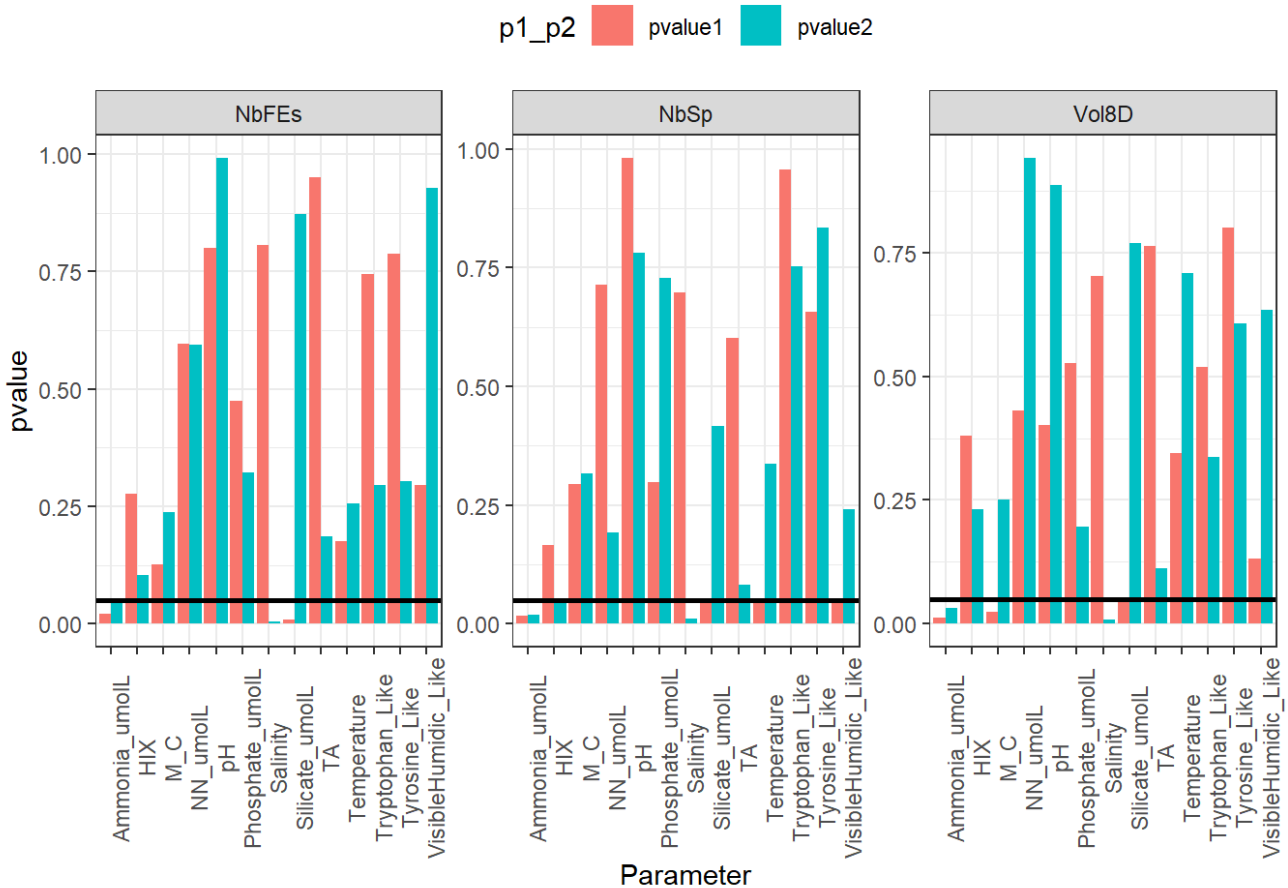
```
## Joining, by = c("Parameter", "ParamType", "Dependent", "Season", "pvalue1",
```

```
## "pvalue2", "r_squared", "adj_r_squared")
```

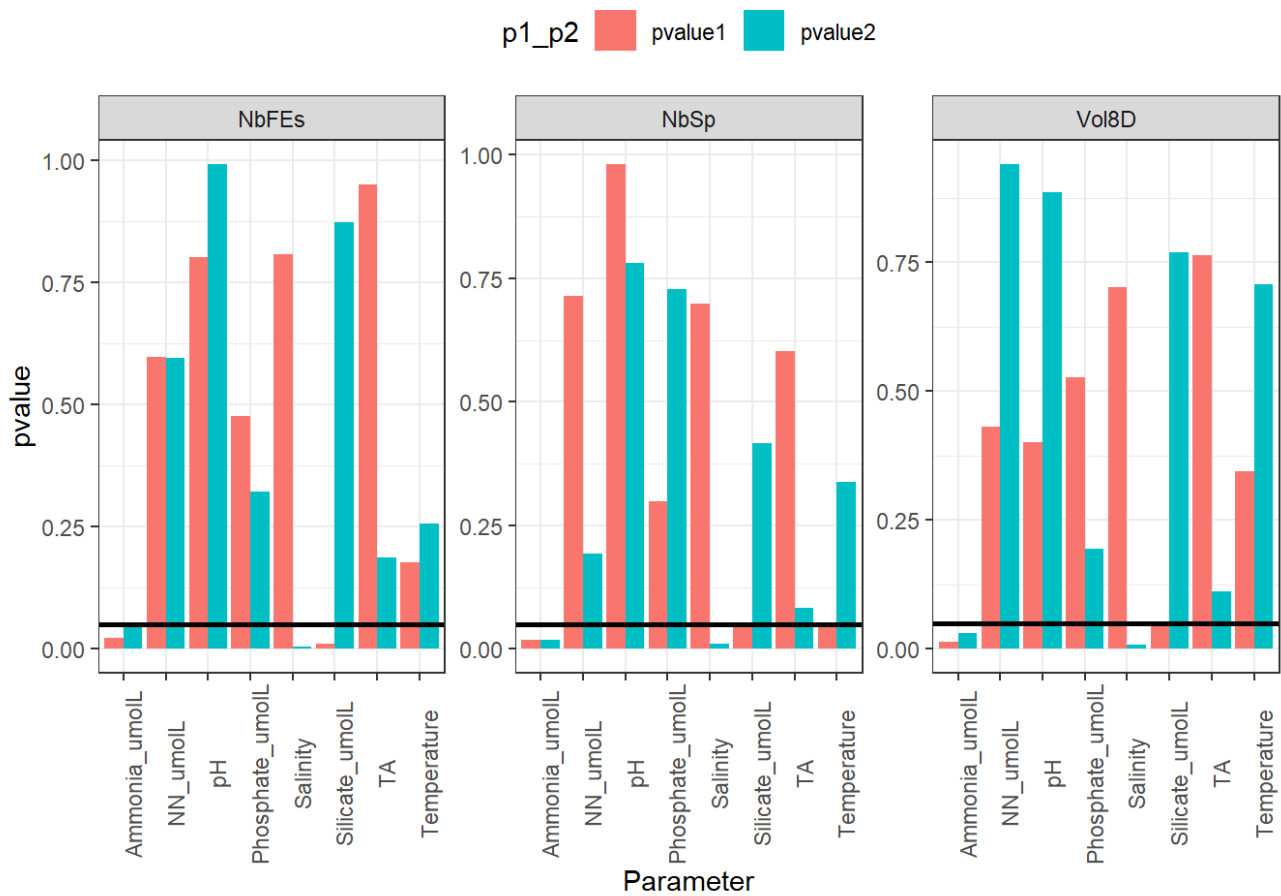
```
CVA
```

```
## # A tibble: 105 × 8
##   Parameter      ParamType Dependent Season pvalue1 pvalue2 r_squared adj_r_...1
##   <chr>          <chr>      <chr>    <chr>    <dbl>   <dbl>   <dbl>   <dbl>
## 1 Salinity       CVA11      NbSp     Dry      0.698   0.0114   0.342   2.60e-1
## 2 Temperature    CVA11      NbSp     Dry      0.0535  0.338    0.250   1.56e-1
## 3 TA             CVA11      NbSp     Dry      0.602   0.0829   0.188   8.65e-2
## 4 pH             CVA11      NbSp     Dry      0.981   0.781    0.00500 -1.19e-1
## 5 Phosphate_umolL CVA11      NbSp     Dry      0.299   0.728    0.0740  -4.18e-2
## 6 Silicate_umolL  CVA11      NbSp     Dry      0.0466  0.416    0.251   1.57e-1
## 7 NN_umolL       CVA11      NbSp     Dry      0.714   0.193    0.110   -8.27e-4
## 8 Ammonia_umolL   CVA11      NbSp     Dry      0.0177  0.0189   0.463   3.96e-1
## 9 M_C            CVA11      NbSp     Dry      0.294   0.316    0.123   1.36e-2
## 10 HIX            CVA11      NbSp     Dry      0.166   0.0471   0.296   2.08e-1
## # ... with 95 more rows, and abbreviated variable name 1adj_r_squared
```

```
pvalplot(mydata = CVA, season = "Dry")
```



```
pvalplot(mydata = CVA, season = "Wet")
```



```
p7 <- paramPlot(mydata = Full_data, ParamType = CVA11, PTname = "CVA11")
```

```
## NULL
## NULL
```

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```
## NULL
## NULL
```