$TXRNA seq_LassoRegressions$

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Load libraries

```
library(tidyverse)
library(glmnet)
library(glmnetUtils)
library(caret)
library(Hmisc)
options(max.print=99999)
```

Open files and prep for analysis

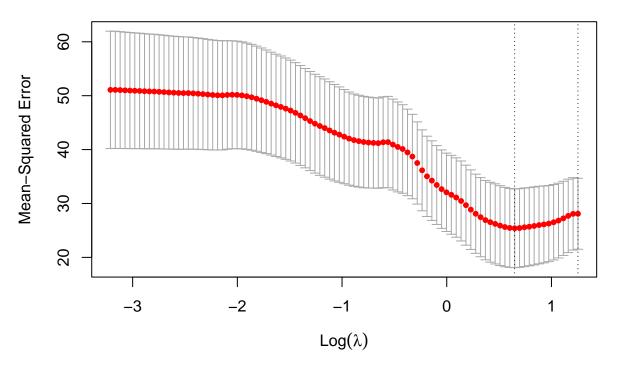
```
#log2fc gene data and plant response data
lfc <- read.csv(file="all_DEGs_log2fc.csv",</pre>
                header=TRUE, row.names=1, stringsAsFactors=FALSE)
plant <- read.csv(file="RNAseqPlantRespData.csv", header=TRUE, row.names=1, stringsAsFactors=FALSE)</pre>
# limit plant data to variables and replicates where RNA was extracted
colnames(plant)
                     "FunCode"
                                                           "Repl"
## [1] "Fungus"
                                 "PhysCode" "Water"
                                                                       "rgr"
   [7] "wilt"
                     "death"
                                 "waterloss" "totbiom"
                                                           "leafcol"
plant_rna <- plant %>%
  dplyr::select(!leafcol) %>%
  dplyr::filter(!is.na(totbiom)) %>%
  dplyr::filter(Fungus != "Alt")
# calculate averages for d calcs
ConD <- plant_rna %>%
  dplyr::filter(Fungus == "Con" & Water == "Drought")
ConD wilt <- mean(ConD$wilt)</pre>
ConD_waterloss <- mean(ConD$waterloss)</pre>
ConW <- plant_rna %>%
  dplyr::filter(Fungus == "Con" & Water == "WellWatered")
ConW_wilt <- mean(ConW$wilt)</pre>
ConW_waterloss <- mean(ConW$waterloss)</pre>
# calculate d values for treatments
plant_rna_d <- plant_rna %>%
  dplyr::filter(Fungus != "Con") %>%
  dplyr::mutate(wilt_d = base::ifelse(Water == "Drought", wilt-ConD_wilt, wilt-ConW_wilt)) %%
 dplyr::mutate(waterloss_d = base::ifelse(Water == "Drought", waterloss-ConD_waterloss, waterloss-ConW
# check
all(rownames(plant_rna_d) == colnames(lfc))
## [1] TRUE
# transpose
lfc <- t(lfc)</pre>
all(rownames(plant_rna_d) == rownames(lfc))
## [1] TRUE
lfc <- as.data.frame(lfc)</pre>
# merge
alldata <- merge(plant_rna_d, lfc, by = "row.names", all=TRUE)
alldata <- as.data.frame(alldata)</pre>
#export
write.csv(alldata, "TXRNAseq_LassoData.csv")
```

Lasso regressions

Create x variable

```
x <- alldata %>%
 dplyr::select(dplyr::starts_with("PhHal"))
anyNA(x)
## [1] FALSE
x <- as.matrix(x)
Wilt
# assign y
y <- alldata$wilt_d
anyNA(y)
## [1] FALSE
set.seed(123)
# use cross-validation (leave-one-out) to select lambda (minimum)
cvfit_wilt <- glmnet::cv.glmnet(x, y,</pre>
                      alpha = 1, nfolds = nrow(x), grouped=FALSE)
cvfit_wilt
##
## Call: glmnet::cv.glmnet(x = x, y = y, nfolds = nrow(x), grouped = FALSE, alpha = 1)
## Measure: Mean-Squared Error
##
##
      Lambda Index Measure
                              SE Nonzero
## min 1.913 14 25.40 7.299
              1 28.09 6.583
## 1se 3.502
r2_wilt <- cvfit_wilt$glmnet.fit$dev.ratio[which(cvfit_wilt$glmnet.fit$lambda == cvfit_wilt$lambda.min)
r2_wilt
## [1] 0.3895734
plot(cvfit_wilt)
```

29 30 30 29 29 27 23 25 23 21 17 9 5 3 1



```
lam_min <- cvfit_wilt$lambda.min
coef(cvfit_wilt, s=lam_min)</pre>
```

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## (Intercept)
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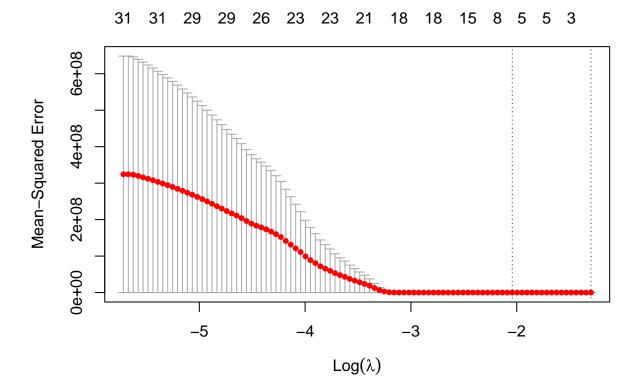
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\#capture.output(print(cvfit\_wilt), file = "TXRNAseq\_Lasso\_wilt.txt")
#capture.output(print(r2_wilt), file = "TXRNAseq_Lasso_wilt.txt", append=TRUE)
\#capture.output(coef(cvfit\_wilt, s=lam\_min), file="TXRNAseq\_Lasso\_wilt.txt", append=TRUE)
Water loss
```

```
# assign y
y <- alldata$waterloss_d
anyNA(y)
## [1] FALSE
set.seed(123)
# use cross-validation (leave-one-out) to select lambda (minimum)
cvfit_waterloss <- glmnet::cv.glmnet(x, y,</pre>
                       alpha = 1, nfolds = nrow(x), grouped=FALSE)
cvfit_waterloss
## Call: glmnet::cv.glmnet(x = x, y = y, nfolds = nrow(x), grouped = FALSE,
                                                                                  alpha = 1)
## Measure: Mean-Squared Error
##
       Lambda Index Measure
                                SE Nonzero
## min 0.1297 17 0.1628 0.03398
## 1se 0.2730 1 0.1862 0.03567
                                          0
r2_waterloss <- cvfit_waterloss$glmnet.fit$dev.ratio[which(cvfit_waterloss$glmnet.fit$lambda == cvfit_w
r2_waterloss
```

[1] 0.5265281



```
lam_min <- cvfit_waterloss$lambda.min
coef(cvfit_waterloss, s=lam_min)</pre>
```

```
## 2575 x 1 sparse Matrix of class "dgCMatrix"
##
## (Intercept)
                  -0.068448448
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- ## PhHAL.3G108600
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- ## PhHAL.9G028600 .
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- ## PhHAL.7G083900 .
- ## PhHAL.1G355400 .
- ## PhHAL.4G158300 .
- ## PhHAL.3G138900 .
- ## PhHAL.3G050200 . ## PhHAL.4G012300 .
- ## PhHAL.5G045600 .

- ## PhHAL.3G390800 .
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- ## PhHAL.2G396500 ## PhHAL.2G396600
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- ## PhHAL.5G410000
- ## PhHAL.4G155800 .
- ## PhHAL.9G043250
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- ## PhHAL.2G452300
- ## PhHAL.9G197400 .
- ## PhHAL.4G248700
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- ## PhHAL.6G201000 .
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- ## PhHAL.9G061700 .
- ## PhHAL.2G215200 ## PhHAL.2G257000
- ## PhHAL.9G444900
- ## PhHAL.3G407600 .
- ## PhHAL.9G179400 .
- ## PhHAL.5G083500
- ## PhHAL.9G589100
- ## PhHAL.2G042300
- ## PhHAL.3G163400
- ## PhHAL.4G317150 .
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## PhHAL.2G365600
## PhHAL.3G192900
## PhHAL.6G192400
## PhHAL.7G344200
## PhHAL.1G333400
## PhHAL.8G123800
## PhHAL.3G237000
## PhHAL.1G012200
## PhHAL.5G347550
## PhHAL.5G460600
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## PhHAL.9G458400 .
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- ## PhHAL.9G078900
- ## PhHAL.1G032000 .
- ## PhHAL.8G106900 .
- ## PhHAL.1G077400 .
- ## PhHAL.3G290400 ## PhHAL.9G271100
- ## PhHAL.9G219100
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- ## PhHAL.2G441900 ## PhHAL.1G364500
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- ## PhHAL.2G465100 .
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- ## PhHAL.9G503200 .
- ## PhHAL.9G503300 .
- ## PhHAL.3G270700 .
- ## PhHAL.1G234400
- ## PhHAL.5G032900
- ## PhHAL.7G312600 .
- ## PhHAL.9G526200 .
- ## PhHAL.5G536200 .
- ## PhHAL.2G247300 .
- ## PhHAL.9G264800 .
- ## PhHAL.9G300100 .
- ## PhHAL.9G268580 .
- ## PhHAL.9G268600 . ## PhHAL.9G269000 .
- ## PhHAL.9G652400 .

- ## PhHAL.4G011500 .
- ## PhHAL.2G256500
- ## PhHAL.1G309400
- ## PhHAL.2G265500
- ## PhHAL.5G490900
- ## PhHAL.5G491000
- ## IMME:00451000
- ## PhHAL.9G271200
- ## PhHAL.7G307600
- ## PhHAL.3G080100
- ## PhHAL.8G259900
- ## PhHAL.4G163500
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- ## PhHAL.1G263400
- ## I IIIIAL: IG200400
- ## PhHAL.2G183900
- ## PhHAL.6G227900
- ## PhHAL.8G004100
- ## PhHAL.4G072900
- ## PhHAL.9G050500
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- ## PhHAL.9G118800
- ## PhHAL.2G369100 .
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- ## PhHAL.1G387200
- ## PhHAL.4G337300 .
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- ## I IIIAL: 4004000
- ## PhHAL.4G343700 .
- ## PhHAL.3G180800
- ## PhHAL.1G022000
- ## PhHAL.3G245000 ## PhHAL.1G151800
- ## PhHAL.5G045700
- ## PhHAL.5G045800
- ## PhHAL.1G419500 .
- ## PhHAL.5G147400
- ## PhHAL.1G001400 .
- ## PhHAL.4G029700
- ## PhHAL.9G216600
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- ## PhHAL.8G232300
- ## PhHAL.8G047600 .
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- ## PhHAL.7G340900 .
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- ## PhHAL.5G461200 .
- ## PhHAL.4G056700 .
- ## PhHAL.5G216700 .
- ## PhHAL.2G387400 .
- ## PhHAL.3G454400 .
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- ## PhHAL.3G168500 .
- ## PhHAL.5G209200
- ## PhHAL.9G069200 .

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- ## PhHAL.5G479500
- ## PhHAL.7G217700
- ## PhHAL.7G246100
- ## PhHAL.9G035400
- ## PhHAL.5G261700
- ## PhHAL.9G152800
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- ## PhHAL.7G223000
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- ## PhHAL.7G310400
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- ## PhHAL.8G142700
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- ## PhHAL.9G487200
- ## PhHAL.3G209800
- ## PhHAL.6G008000
- ## PhHAL.4G321600
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- ## PhHAL.9G064200 .
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- ## PhHAL.3G070100
- ## PhHAL.9G625300
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- ## PhHAL.6G212100
- ## PhHAL.7G161600
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- ## PhHAL.2G251600
- ## PhHAL.6G021600
- ## PhHAL.5G392900 .
- ## PhHAL.9G059400 .
- ## PhHAL.9G059500
- ## PhHAL.9G624100
- ## PhHAL.9G155100
- ## PhHAL.2G061000
- ## PhHAL.6G286400
- ## PhHAL.7G035000 .
- ## PhHAL.1G269000
- ## PhHAL.2G439500 .
- ## PhHAL.3G154100 .
- ## PhHAL.3G310800 .
- ## PhHAL.6G155200
- ## PhHAL.5G283000
- ## PhHAL.1G081100
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- ## PhHAL.9G488500 .
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- ## PhHAL.3G468800
- ## PhHAL.4G028800
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- ## PhHAL.2G378800 .
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- ## PhHAL.7G292600 .
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- ## PhHAL.6G032300
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- ## PhHAL.1G002500
- ## PhHAL.3G197900 ## PhHAL.5G471600
- ## PhHAL.7G196800
- ## PhHAL.5G159500
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- ## PhHAL.3G242000
- ## PhHAL.3G125000 ## PhHAL.2G333100
- ## PhHAL.3G227400
- ## PhHAL.7G317000
- ## PhHAL.3G059700 ## PhHAL.7G201300
- ## PhHAL.7G162200
- ## PhHAL.3G158900
- ## PhHAL.7G333600
- ## PhHAL.9G070200
- ## PhHAL.5G121000
- ## PhHAL.4G046300
- ## PhHAL.8G250000
- ## PhHAL.8G250100
- ## PhHAL.8G250200
- ## PhHAL.4G115700
- ## PhHAL.2G123400
- ## PhHAL.2G123500
- ## PhHAL.7G226600
- ## PhHAL.4G361200
- ## PhHAL.7G176200
- ## PhHAL.1G443300
- ## PhHAL.3G433300 ## PhHAL.4G359800
- ## PhHAL.5G178900
- ## PhHAL.9G521700

- ## PhHAL.9G080600 .
- ## PhHAL.7G216200 .
- ## PhHAL.9G469100
- ## PhHAL.6G268000
- ## PhHAL.7G068600
- ## PhHAL.3G407000 .
- ## PhHAL.9G634200
- ## PhHAL.7G148500 .
- ## PhHAL.3G162000
- ## PhHAL.5G090600
- ## I IIIAE.0000000
- ## PhHAL.5G091600
- ## PhHAL.8G116800
- ## PhHAL.9G478800
- ## PhHAL.9G545500 ## PhHAL.2G355400
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- ## PhHAL.2G355500 .
- ## PhHAL.5G196700 . ## PhHAL.5G196750 .
- ## PhHAL.9G472700 .
- ## PhHAL.6G036000 .
- ## PhHAL.8G189500 .
- ## PhHAL.6G174000 .
- ## PhHAL.4G158100 .
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- ## PhHAL.1G163900
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- ## PhHAL.9G459100
- ## PhHAL.9G417400
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- ## PhHAL.9G595900
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- ## PhHAL.6G171700
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- ## PhHAL.1G381400
- ## PhHAL.1G339700 .
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- ## PhHAL.2G310900 .
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- ## PhHAL.9G456600
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- ## PhHAL.9G652600
- ## PhHAL.1G351800
- ## PhHAL.5G010800
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- ## PhHAL.4G086600
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- ## PhHAL.6G289200
- ## PhHAL.4G003600
- ## PhHAL.8G201975
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- ## PhHAL.2G030500
- ## PhHAL.4G163600
- ## PhHAL.6G007300
- ## PhHAL.9G543600
- ## PhHAL.2G377800
- ## PhHAL.4G055900 ## PhHAL.9G629100
- ## PhHAL.9G194200

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- ## PhHAL.5G058200
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- ## PhHAL.3G330400
- ## PhHAL.3G278600
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- ## PhHAL.3G215000
- ## PhHAL.3G215100
- ## PhHAL.3G215200
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- ## PhHAL.9G116600
- ## PhHAL.2G340000
- ## PhHAL.2G426200
- ## PhHAL.2G032600 ## PhHAL.5G069400
- ## PhHAL.9G444300
- ## PhHAL.9G463300
- ## PhHAL.4G021000 ## PhHAL.4G021100
- ## PhHAL.1G322000 ## PhHAL.4G069400
- ## PhHAL.9G240000
- ## PhHAL.9G183000
- ## PhHAL.1G124000
- ## PhHAL.5G232800
- ## PhHAL.3G380700
- ## PhHAL.1G339300
- ## PhHAL.5G286900
- ## PhHAL.5G306400
- ## PhHAL.9G608100
- ## PhHAL.1G417300
- ## PhHAL.6G079700
- ## PhHAL.9G641400
- ## PhHAL.4G350800
- ## PhHAL.5G139600
- ## PhHAL.4G036500
- ## PhHAL.2G357200
- ## PhHAL.9G402800
- ## PhHAL.4G086000
- ## PhHAL.5G190900
- ## PhHAL.1G285800
- ## PhHAL.8G125200 ## PhHAL.2G371600
- ## PhHAL.6G115100
- ## PhHAL.3G390600

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## PhHAL.9G209400
## PhHAL.5G180400
## PhHAL.7G029600
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## PhHAL.9G522600
## PhHAL.9G534800
## PhHAL.6G086400
## PhHAL.8G112300
## PhHAL.9G438900
## PhHAL.5G178200
## PhHAL.7G106200
## PhHAL.3G039500
## PhHAL.4G366900
## PhHAL.3G079700
## PhHAL.9G158000
## PhHAL.3G433500
## PhHAL.5G012850
## PhHAL.9G205166
## PhHAL.9G584900
\#capture.output(print(cufit\_waterloss), file = "TXRNAseq\_Lasso\_waterloss.txt")
#capture.output(print(r2_waterloss), file = "TXRNAseq_Lasso_waterloss.txt", append=TRUE)
\#capture.output(coef(cvfit\_waterloss, s=lam\_min), file="TXRNAseq\_Lasso\_waterloss.txt", append=TRUE)
```

Linear regression for DEGs identified in Lasso

re-open data file if needed

```
alldata <- read.csv("TXRNAseq_LassoData.csv")
```

wilt

```
#PhHAL.7G251400
summary(lm(wilt_d ~ PhHAL.7G251400, data=alldata))
##
## Call:
## lm(formula = wilt_d ~ PhHAL.7G251400, data = alldata)
##
## Residuals:
               1Q Median
                               ЗQ
                                      Max
## -8.4918 -1.0633 -0.0716 1.9053 7.6896
## Coefficients:
##
                 Estimate Std. Error t value Pr(>|t|)
## (Intercept)
                 -1.3708
                              0.8032 -1.707 0.09894 .
## PhHAL.7G251400 -3.0438
                              0.7207 -4.223 0.00023 ***
## ---
```

```
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 4.148 on 28 degrees of freedom
## Multiple R-squared: 0.3891, Adjusted R-squared: 0.3673
## F-statistic: 17.84 on 1 and 28 DF, p-value: 0.0002303
#PhHAL.2G280300
summary(lm(wilt_d ~ PhHAL.2G280300, data=alldata))
##
## Call:
## lm(formula = wilt_d ~ PhHAL.2G280300, data = alldata)
## Residuals:
    Min
             1Q Median
                           30
                                 Max
## -7.674 -1.974 -0.284 2.919 6.112
##
## Coefficients:
##
                 Estimate Std. Error t value Pr(>|t|)
## (Intercept)
                  -0.2498
                              0.8414 - 0.297
                                                0.769
## PhHAL.2G280300 -2.4406
                              0.4933 -4.948 3.2e-05 ***
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
## Residual standard error: 3.877 on 28 degrees of freedom
## Multiple R-squared: 0.4665, Adjusted R-squared: 0.4474
## F-statistic: 24.48 on 1 and 28 DF, p-value: 3.2e-05
#PhHAL.2G424800
summary(lm(wilt_d ~ PhHAL.2G424800, data=alldata))
##
## Call:
## lm(formula = wilt_d ~ PhHAL.2G424800, data = alldata)
##
## Residuals:
      Min
               1Q Median
                               3Q
                                      Max
## -8.7694 -3.5266 0.4438 2.8907 9.2037
##
## Coefficients:
##
                 Estimate Std. Error t value Pr(>|t|)
                  -2.3163
                              0.8073 -2.869 0.00774 **
## (Intercept)
                              0.4193 -3.537 0.00143 **
## PhHAL.2G424800 -1.4829
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 4.413 on 28 degrees of freedom
## Multiple R-squared: 0.3088, Adjusted R-squared: 0.2841
## F-statistic: 12.51 on 1 and 28 DF, p-value: 0.001432
#PhHAL.2G319800
summary(lm(wilt_d ~ PhHAL.2G319800, data=alldata))
```

```
## Call:
## lm(formula = wilt_d ~ PhHAL.2G319800, data = alldata)
## Residuals:
##
      Min
              1Q Median
                               3Q
                                      Max
## -7.9042 -2.7121 -0.6451 3.2927 6.5473
##
## Coefficients:
##
                 Estimate Std. Error t value Pr(>|t|)
## (Intercept)
                  0.1456
                              0.9679 0.150 0.881527
                              0.4600 -4.326 0.000174 ***
## PhHAL.2G319800 -1.9901
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 4.109 on 28 degrees of freedom
## Multiple R-squared: 0.4006, Adjusted R-squared: 0.3792
## F-statistic: 18.71 on 1 and 28 DF, p-value: 0.0001744
water loss
#PhHAL.9G036800
summary(lm(waterloss_d ~ PhHAL.9G036800, data=alldata))
##
## Call:
## lm(formula = waterloss d ~ PhHAL.9G036800, data = alldata)
## Residuals:
##
       Min
                 1Q Median
                                   3Q
                                           Max
## -0.47163 -0.28922 0.01046 0.17136 0.76237
##
## Coefficients:
##
                 Estimate Std. Error t value Pr(>|t|)
## (Intercept)
                 -0.03176
                             0.05952 -0.534
                                               0.598
## PhHAL.9G036800 0.21025
                             0.04636
                                      4.535 9.88e-05 ***
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 0.3259 on 28 degrees of freedom
## Multiple R-squared: 0.4235, Adjusted R-squared: 0.4029
## F-statistic: 20.57 on 1 and 28 DF, p-value: 9.88e-05
#PhHAL.6G027300
summary(lm(waterloss_d ~ PhHAL.6G027300, data=alldata))
##
## Call:
## lm(formula = waterloss_d ~ PhHAL.6G027300, data = alldata)
## Residuals:
```

##

```
1Q
                    Median
                                  3Q
## -0.52790 -0.26863 -0.03991 0.25304 0.73866
##
## Coefficients:
##
                 Estimate Std. Error t value Pr(>|t|)
                 -0.15007
                            0.06924 -2.167 0.038866 *
## (Intercept)
                            0.02970 3.911 0.000534 ***
## PhHAL.6G027300 0.11616
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.3452 on 28 degrees of freedom
## Multiple R-squared: 0.3533, Adjusted R-squared: 0.3302
## F-statistic: 15.29 on 1 and 28 DF, p-value: 0.0005336
#PhHAL.7G113600
summary(lm(waterloss_d ~ PhHAL.7G113600, data=alldata))
##
## Call:
## lm(formula = waterloss_d ~ PhHAL.7G113600, data = alldata)
## Residuals:
##
       Min
                 1Q Median
                                  3Q
                                          Max
## -0.62014 -0.20444 0.02908 0.21650 0.57189
##
## Coefficients:
                 Estimate Std. Error t value Pr(>|t|)
##
                 -0.03328
                            0.06093 -0.546 0.589297
## (Intercept)
## PhHAL.7G113600 -0.23110
                            0.05398 -4.281 0.000197 ***
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
##
## Residual standard error: 0.3337 on 28 degrees of freedom
## Multiple R-squared: 0.3957, Adjusted R-squared: 0.3741
## F-statistic: 18.33 on 1 and 28 DF, p-value: 0.0001967
#PhHAL.9G394700
summary(lm(waterloss_d ~ PhHAL.9G394700, data=alldata))
##
## Call:
## lm(formula = waterloss_d ~ PhHAL.9G394700, data = alldata)
##
## Residuals:
##
       Min
                 1Q Median
                                  3Q
                                          Max
## -0.43672 -0.25117 -0.03596 0.20987 0.74042
##
## Coefficients:
##
                 Estimate Std. Error t value Pr(>|t|)
                 ## (Intercept)
## PhHAL.9G394700 0.21330
                            0.04608 4.629 7.65e-05 ***
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
```

```
##
## Residual standard error: 0.323 on 28 degrees of freedom
## Multiple R-squared: 0.4335, Adjusted R-squared: 0.4133
## F-statistic: 21.43 on 1 and 28 DF, p-value: 7.646e-05
#PhHAL.2G014400
summary(lm(waterloss_d ~ PhHAL.2G014400, data=alldata))
##
## Call:
## lm(formula = waterloss_d ~ PhHAL.2G014400, data = alldata)
## Residuals:
##
       Min
                 1Q Median
                                   30
                                          Max
## -0.93490 -0.16902 0.01814 0.25319 0.53366
##
## Coefficients:
##
                 Estimate Std. Error t value Pr(>|t|)
                 -0.26341
                            0.09604 -2.743 0.0105 *
## (Intercept)
## PhHAL.2G014400 0.33755
                                     3.264 0.0029 **
                             0.10343
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
##
## Residual standard error: 0.3653 on 28 degrees of freedom
## Multiple R-squared: 0.2756, Adjusted R-squared: 0.2497
## F-statistic: 10.65 on 1 and 28 DF, p-value: 0.002897
#PhHAL.1G151400
summary(lm(waterloss_d ~ PhHAL.1G151400, data=alldata))
##
## Call:
## lm(formula = waterloss_d ~ PhHAL.1G151400, data = alldata)
## Residuals:
       Min
                 1Q Median
                                   3Q
## -0.67035 -0.27527 0.05551 0.26581 0.71055
## Coefficients:
##
                 Estimate Std. Error t value Pr(>|t|)
                -0.06530 0.06559 -0.996 0.32796
## (Intercept)
## PhHAL.1G151400 0.11178
                             0.03158 3.540 0.00142 **
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 0.3568 on 28 degrees of freedom
## Multiple R-squared: 0.3091, Adjusted R-squared: 0.2845
## F-statistic: 12.53 on 1 and 28 DF, p-value: 0.001422
#PhHAL.9G551200
summary(lm(waterloss_d ~ PhHAL.9G551200, data=alldata))
```

```
##
## Call:
## lm(formula = waterloss_d ~ PhHAL.9G551200, data = alldata)
## Residuals:
               1Q Median
                                  3Q
##
       Min
                                          Max
## -0.49299 -0.33291 -0.01092 0.27231 0.56237
## Coefficients:
##
                 Estimate Std. Error t value Pr(>|t|)
## (Intercept) -0.05169
                            0.06425 -0.804 0.427904
## PhHAL.9G551200 0.25403
                            0.06841 3.713 0.000902 ***
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
## Residual standard error: 0.3513 on 28 degrees of freedom
## Multiple R-squared: 0.3299, Adjusted R-squared: 0.306
## F-statistic: 13.79 on 1 and 28 DF, p-value: 0.0009017
```

Function of genes in lasso

open gene list file and annotation file

```
genelist <- read.csv("TXRNAseq_LassoGenes.csv", header=TRUE)
anno <- read.delim("PhalliiHAL_anno.txt")
gene_anno <- dplyr::left_join(genelist, anno, by="locusName")
write.csv(gene_anno, "TXRNAseq_LassoGeneFunctions.csv")</pre>
```

Session Info

```
sessionInfo()
```

```
## R version 4.2.2 (2022-10-31 ucrt)
## Platform: x86_64-w64-mingw32/x64 (64-bit)
## Running under: Windows 10 x64 (build 19044)
##
## Matrix products: default
##
## locale:
## [1] LC_COLLATE=English_United States.utf8
## [2] LC_CTYPE=English_United States.utf8
## [3] LC_MONETARY=English_United States.utf8
## [4] LC_NUMERIC=C
## [5] LC_TIME=English_United States.utf8
##
## attached base packages:
```

```
## [1] stats
                 graphics grDevices utils
                                                datasets methods
                                                                     base
##
## other attached packages:
   [1] Hmisc_4.7-2
                          Formula_1.2-4
                                             survival_3.4-0
                                                                caret_6.0-93
##
   [5] lattice_0.20-45
                          glmnetUtils_1.1.8 glmnet_4.1-6
                                                                Matrix_1.5-3
  [9] forcats 0.5.2
                          stringr_1.5.0
                                                                purrr 1.0.1
##
                                             dplyr 1.0.10
                                                                ggplot2 3.4.0
## [13] readr 2.1.3
                          tidyr_1.2.1
                                             tibble_3.1.8
## [17] tidyverse_1.3.2
##
## loaded via a namespace (and not attached):
  [1] googledrive_2.0.0
                              colorspace_2.1-0
                                                   deldir_1.0-6
   [4] ellipsis_0.3.2
                              class_7.3-20
                                                   htmlTable_2.4.1
## [7] base64enc_0.1-3
                              fs_1.6.0
                                                   rstudioapi_0.14
## [10] listenv_0.9.0
                             prodlim_2019.11.13
                                                   fansi_1.0.4
## [13] lubridate_1.9.0
                             xm12_1.3.3
                                                   codetools_0.2-18
## [16] splines_4.2.2
                             knitr_1.42
                                                   jsonlite_1.8.4
## [19] pROC_1.18.0
                             broom_1.0.3
                                                   cluster_2.1.4
                             png_0.1-8
## [22] dbplyr 2.3.0
                                                   compiler 4.2.2
## [25] httr_1.4.4
                             backports_1.4.1
                                                   assertthat_0.2.1
## [28] fastmap 1.1.0
                             gargle_1.3.0
                                                   cli 3.6.0
## [31] htmltools_0.5.4
                             tools_4.2.2
                                                   gtable_0.3.1
## [34] glue_1.6.2
                             reshape2_1.4.4
                                                   Rcpp 1.0.10
## [37] cellranger_1.1.0
                                                   nlme_3.1-161
                             vctrs 0.5.2
## [40] iterators 1.0.14
                             timeDate_4022.108
                                                   gower 1.0.1
## [43] xfun 0.36
                              globals 0.16.2
                                                   rvest_1.0.3
## [46] timechange_0.2.0
                             lifecycle_1.0.3
                                                   googlesheets4_1.0.1
## [49] future_1.31.0
                             MASS_7.3-58.1
                                                   scales_1.2.1
## [52] ipred_0.9-13
                             hms_1.1.2
                                                   parallel_4.2.2
## [55] RColorBrewer_1.1-3
                             yaml_2.3.6
                                                   gridExtra_2.3
## [58] rpart_4.1.19
                             latticeExtra_0.6-30
                                                   stringi_1.7.12
## [61] highr_0.10
                              foreach_1.5.2
                                                   checkmate_2.1.0
## [64] hardhat_1.2.0
                             lava_1.7.1
                                                   shape_1.4.6
## [67] rlang_1.0.6
                             pkgconfig_2.0.3
                                                   evaluate_0.20
## [70] recipes_1.0.4
                             htmlwidgets_1.6.1
                                                   tidyselect_1.2.0
## [73] parallelly 1.34.0
                             plvr 1.8.8
                                                   magrittr 2.0.3
## [76] R6_2.5.1
                                                   DBI_1.1.3
                             generics_0.1.3
## [79] pillar 1.8.1
                             haven 2.5.1
                                                   foreign_0.8-84
## [82] withr_2.5.0
                             nnet_7.3-18
                                                   future.apply_1.10.0
## [85] modelr_0.1.10
                              crayon_1.5.2
                                                   interp_1.1-3
## [88] utf8_1.2.2
                             tzdb_0.3.0
                                                   rmarkdown_2.20
## [91] jpeg_0.1-10
                             grid 4.2.2
                                                   readxl 1.4.1
## [94] data.table 1.14.6
                             ModelMetrics_1.2.2.2 reprex_2.0.2
## [97] digest_0.6.31
                             stats4_4.2.2
                                                   munsell 0.5.0
```