Ácidos orgánicos

Contents

Acidos orgánicos en peso fresco

Concentración del perfíl de ácidos orgánicos a distintos estados de Madurez

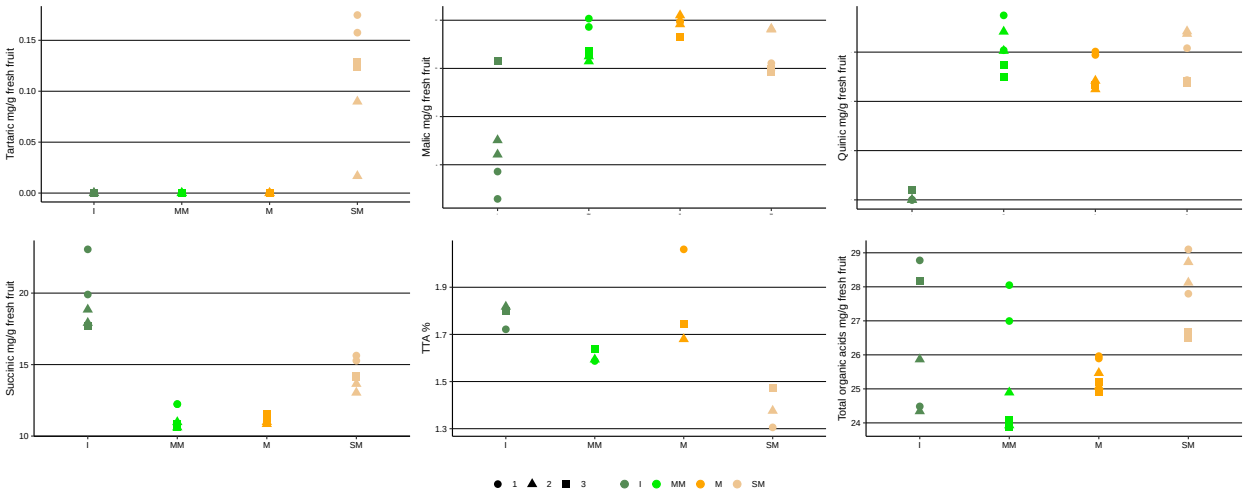


Tabla descriptiva

CAR	MAD	N	CONF	sd	se	ci
Tartárico	I	6	0.000000000	0.000000000	0.000000000	0.000000000
Tartárico	MM	6	0.000000000	0.000000000	0.000000000	0.000000000
Tartárico	M	6	0.000000000	0.000000000	0.000000000	0.000000000
Tartárico	SM	6	0.115256167	0.056458865	0.023049235	0.059249945
Málico	I	6	7.387488667	2.390397202	0.975875571	2.508568017
Málico	MM	6	10.975546167	0.732827314	0.299175498	0.769055101
Málico	M	6	11.759538667	0.362052545	0.147807333	0.379950845
Málico	SM	6	10.536866833	0.856124888	0.349511522	0.898447970
Quínico	I	6	0.066569333	0.103128768	0.042102143	0.108227004
Quínico	MM	6	3.076024167	0.452481056	0.184724617	0.474849746
Quínico	M	6	2.545536167	0.339503763	0.138601831	0.356287349
Quínico	SM	6	2.849149333	0.498337071	0.203445257	0.522972683
Succínico	I	6	19.183774833	2.084913315	0.851162297	2.187982339
Succínico	MM	6	11.252415500	0.774289201	0.316102243	0.812566683
Succínico	M	6	11.112724833	0.255429687	0.104278733	0.268057017
Succínico	SM	6	14.317510333	0.977555462	0.399085346	1.025881542
ATT	I	3	1.779200000	0.050798425	0.029328484	0.126190284
ATT	MM	3	1.606400000	0.027896953	0.016106313	0.069299874
ATT	M	3	1.828266667	0.203906384	0.117725406	0.506531538
ATT	SM	3	1.384533333	0.083527560	0.048224659	0.207493963
TOTALac	I	6	26.637833026	1.990422196	0.812586459	2.088819991
TOTALac	MM	6	25.303985619	1.788299304	0.730070134	1.876705024
TOTALac	M	6	25.417799690	0.434146910	0.177239734	0.455609240
TOTALac	SM	6	27.818782698	1.061812720	0.433483228	1.114304111
NA	I	6	45.629847833	4.811537003	1.964301756	5.049398414
NA	MM	6	36.768222667	5.141035970	2.098819146	5.395186373
NA	M	6	39.159253500	2.599418000	1.061207955	2.727921892
NA	SM	6	53.119642833	1.842284749	0.752109600	1.933359275

Evolución del perfíl de ácidos orgánicos

```
## Error in `palette()`:
## ! Insufficient values in manual scale. 6 needed but only 4 provided.
```

Acidos orgánicos Totales

Concentración de ácidos orgánicos totales

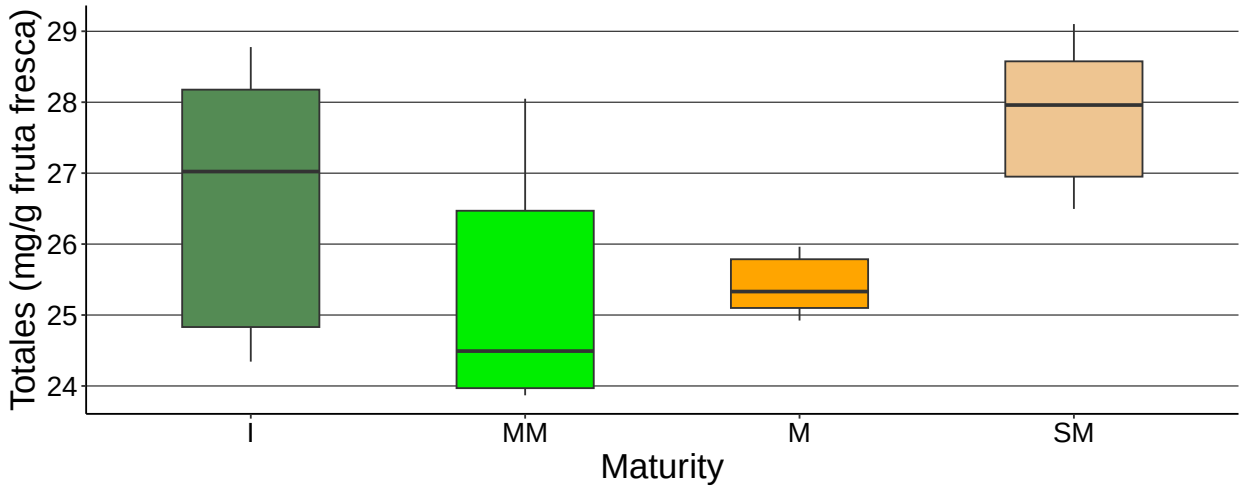
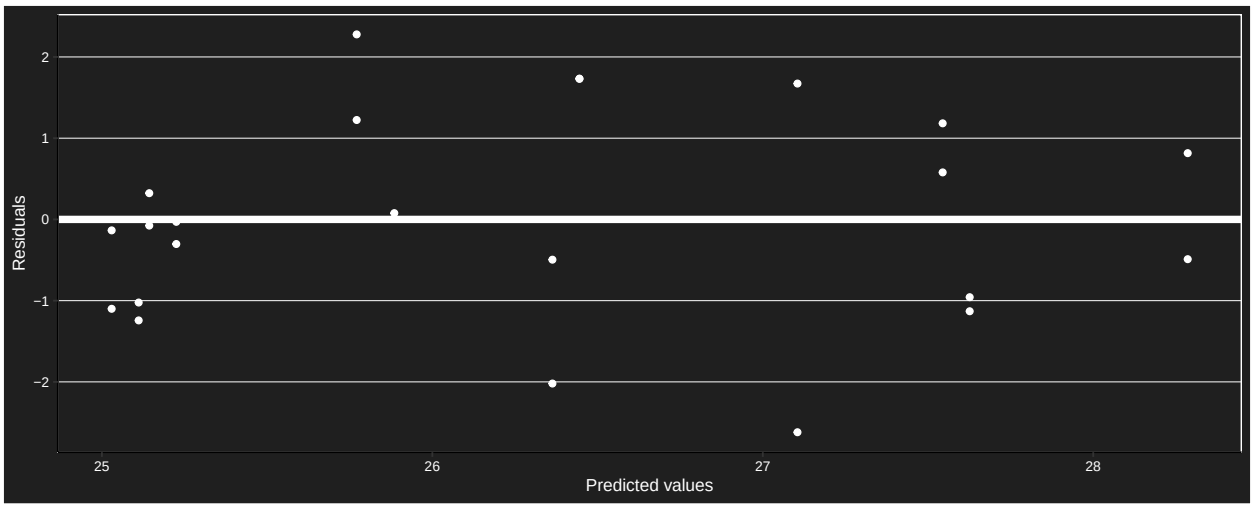
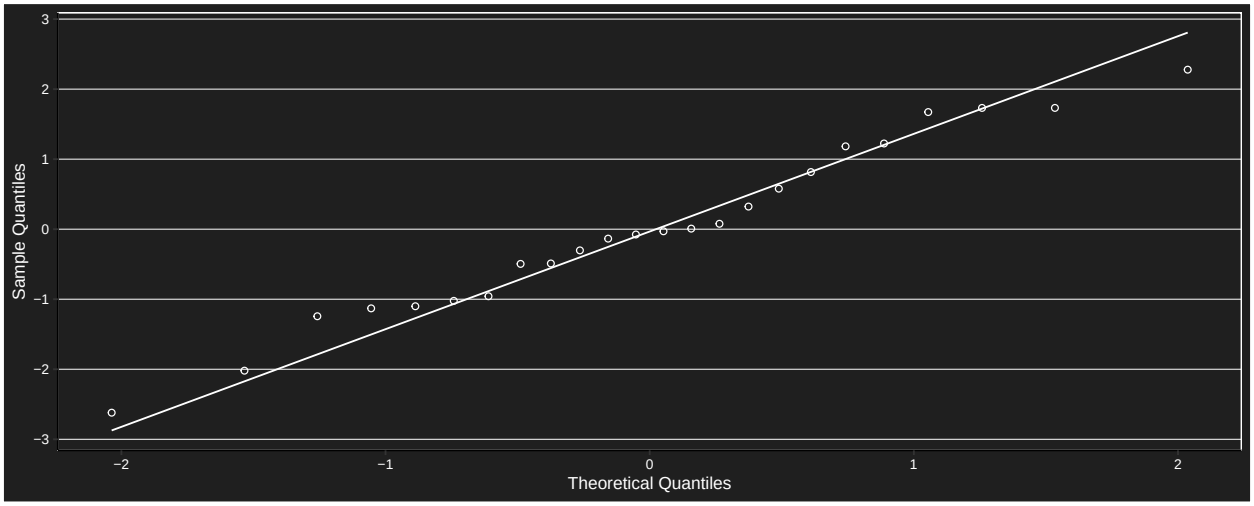


Tabla descriptiva totales

CAR	MAD	N	TOTALF	sd	se	ci
ACIDS	I	6	26.63783300	1.990422128	0.812586431	2.088819919
ACIDS	MM	6	25.30398550	1.788299193	0.730070088	1.876704908
ACIDS	M	6	25.41779983	0.434147222	0.177239861	0.455609568
ACIDS	SM	6	27.81878267	1.061812544	0.433483156	1.114303926
CATIONS	I	3	3.59063333	1.325520714	0.765289741	3.292775993
CATIONS	MM	3	2.56066667	0.313536989	0.181020665	0.778869058
CATIONS	M	3	2.60383333	0.308997643	0.178399872	0.767592698
CATIONS	SM	3	2.21436667	0.394508331	0.227769491	0.980013023
STAT	I	3	1.72112248	0.263331774	0.152034671	0.654152390
STAT	MM	3	1.44765169	0.116290944	0.067140608	0.288882719
STAT	M	3	1.54171590	0.132691311	0.076609364	0.329623489
STAT	SM	3	1.90950428	0.007733365	0.004464861	0.019210744
SUGARS	I	6	45.62984817	4.811537421	1.964301927	5.049398853
SUGARS	MM	6	36.76822233	5.141036221	2.098819248	5.395186636
SUGARS	M	6	39.15925367	2.599418252	1.061208058	2.727922157
SUGARS	SM	6	53.11964317	1.842284986	0.752109696	1.933359523

```
## Linear mixed-effects model fit by REML
##   Data: dataAT
##   Log-restricted-likelihood: -39.0996772
##   Fixed: TOTALF ~ MAD
## (Intercept)       MADMM        MADM       MADSM
## 26.63783300 -1.33384750 -1.22003317  1.18094967
##
## Random effects:
##  Formula: ~1 | REP
##         (Intercept)   Residual
## StdDev: 0.544428081 1.37176539
##
## Number of Observations: 24
## Number of Groups: 3
```






```
##
##  Shapiro-Wilk normality test
##
## data:  e
## W = 0.9776689, p-value = 0.849326
```

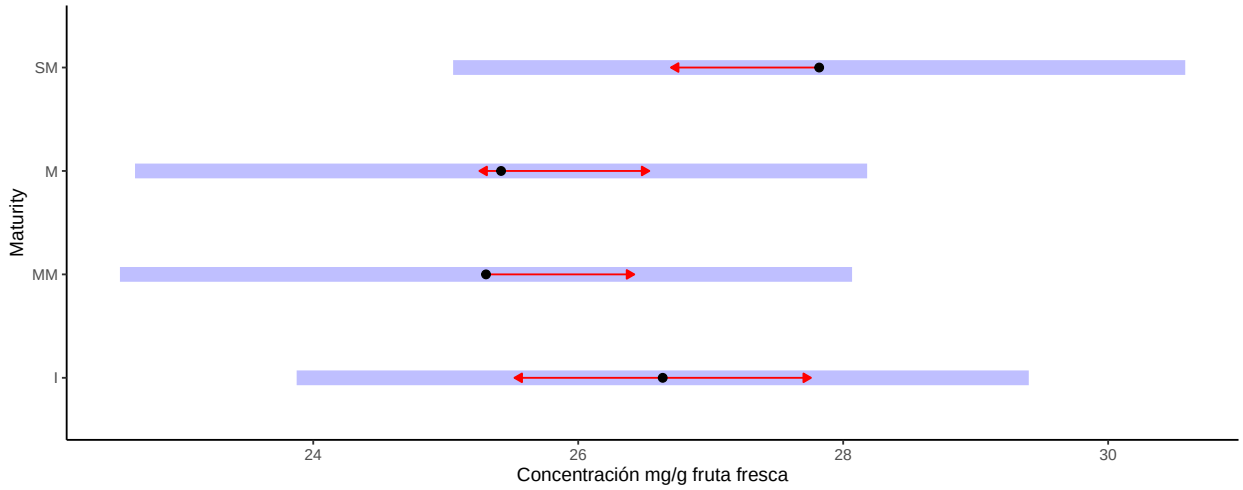
Anova

```
##             numDF denDF    F-value p-value
## (Intercept)     1    18 3901.69682  <.0001
## MAD             3    18    4.45443  0.0165
```

Test de Tukey

```
## $emmeans
##  MAD emmean    SE df lower.CL upper.CL
##  I     26.6 0.642  2     23.9     29.4
##  MM    25.3 0.642  2     22.5     28.1
```

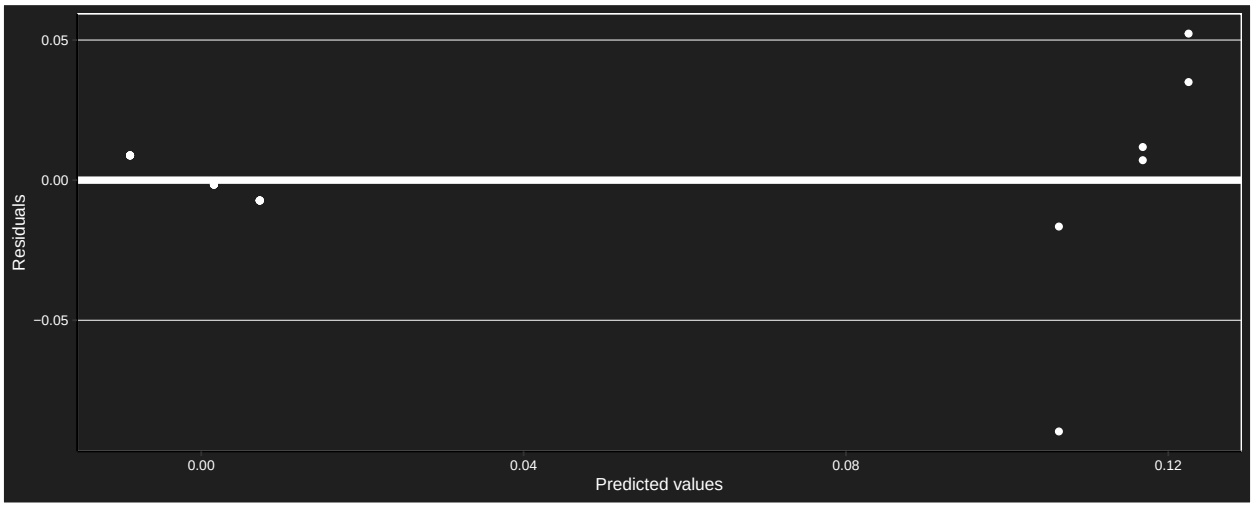
```
##  M     25.4 0.642  2     22.7     28.2
##  SM    27.8 0.642  2     25.1     30.6
##
## Degrees-of-freedom method: containment
## Confidence level used: 0.95
##
## $contrasts
##  contrast estimate    SE df t.ratio p.value
##  I - MM      1.334 0.792 18   1.684  0.3602
##  I - M       1.220 0.792 18   1.540  0.4354
##  I - SM     -1.181 0.792 18  -1.491  0.4629
##  MM - M     -0.114 0.792 18  -0.144  0.9989
##  MM - SM    -2.515 0.792 18  -3.175  0.0246
##  M - SM     -2.401 0.792 18  -3.032  0.0331
##
## Degrees-of-freedom method: containment
## P value adjustment: tukey method for comparing a family of 4 estimates
```

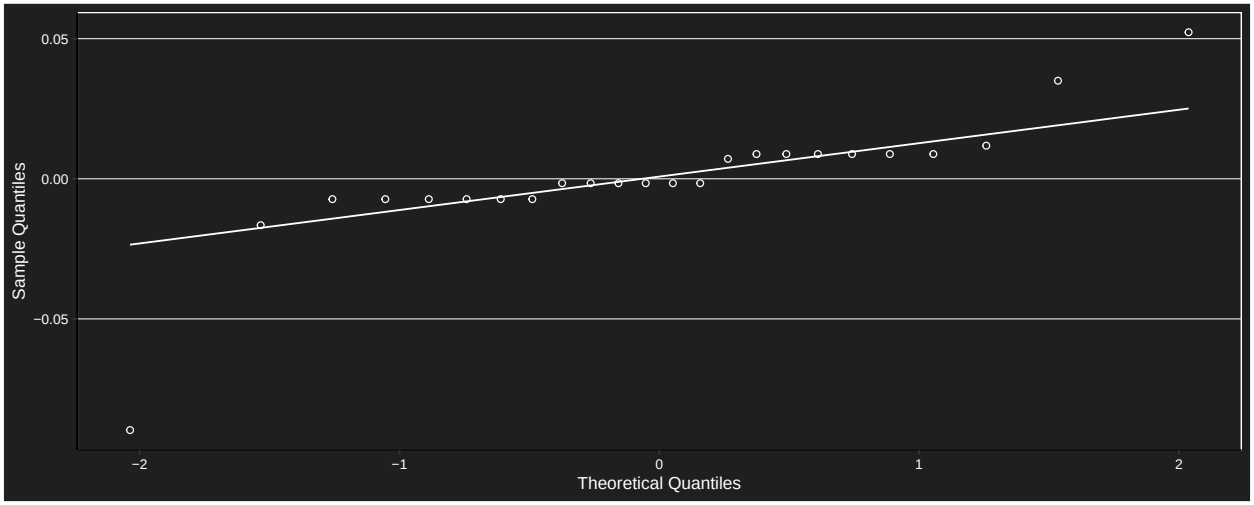


Ácido Tartárico

Modelo y supuestos

```
## Linear mixed-effects model fit by REML
##   Data: tar
##   Log-restricted-likelihood: 39.7876038
##   Fixed: CONF ~ MAD
##     (Intercept)           MADMM            MADM           MADSM
## -1.16339963e-17 -1.91612264e-18  1.38777878e-17  1.15256167e-01
##
## Random effects:
##  Formula: ~1 | REP
##          (Intercept)     Residual
## StdDev: 0.0108045226 0.0265232023
##
## Number of Observations: 24
## Number of Groups: 3
```






```
##
##  Shapiro-Wilk normality test
##
## data:  e
## W = 0.7268705, p-value = 2.33095e-05
```

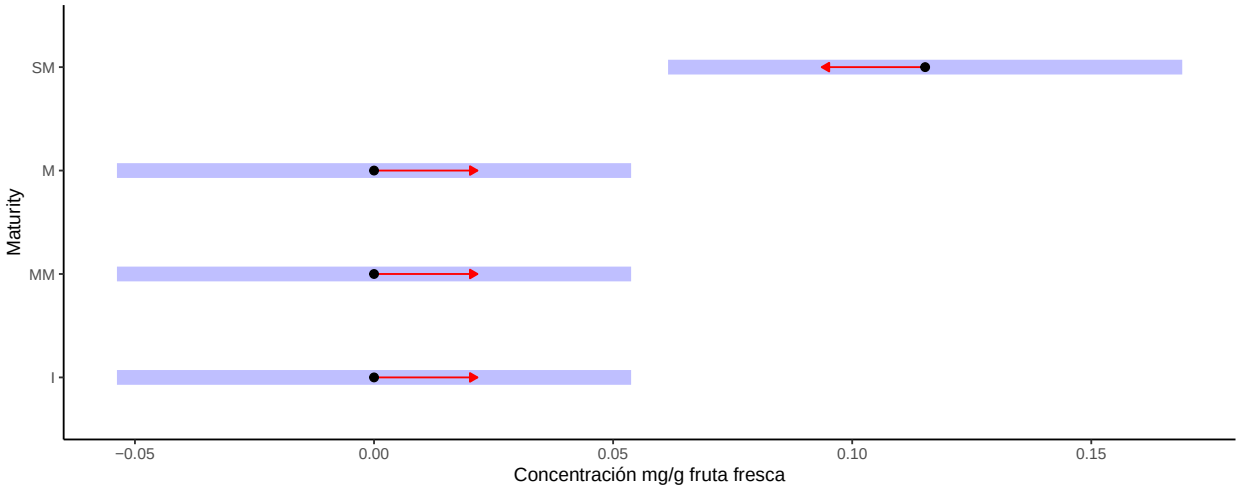
Anova

```
##             numDF denDF    F-value p-value
## (Intercept)     1    18 12.1694124  0.0026
## MAD             3    18 28.3248543  <.0001
```

Test de Tukey

```
## $emmeans
##  MAD emmean     SE df lower.CL upper.CL
##  I    0.000 0.0125  2  -0.0538   0.0538
```

```
##  MM   0.000 0.0125  2  -0.0538   0.0538
##  M    0.000 0.0125  2  -0.0538   0.0538
##  SM   0.115 0.0125  2   0.0615   0.1690
##
## Degrees-of-freedom method: containment
## Confidence level used: 0.95
##
## $contrasts
##  contrast estimate     SE df t.ratio p.value
##  I - MM      0.000 0.0153 18   0.000  1.0000
##  I - M       0.000 0.0153 18   0.000  1.0000
##  I - SM     -0.115 0.0153 18  -7.527  <.0001
##  MM - M      0.000 0.0153 18   0.000  1.0000
##  MM - SM    -0.115 0.0153 18  -7.527  <.0001
##  M - SM     -0.115 0.0153 18  -7.527  <.0001
##
## Degrees-of-freedom method: containment
## P value adjustment: tukey method for comparing a family of 4 estimates
```

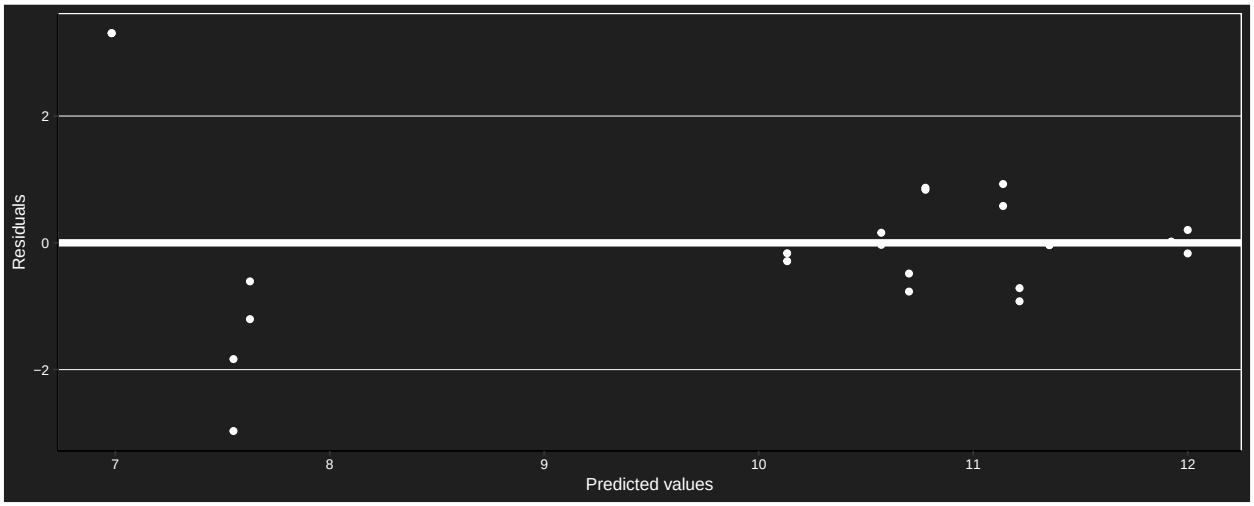


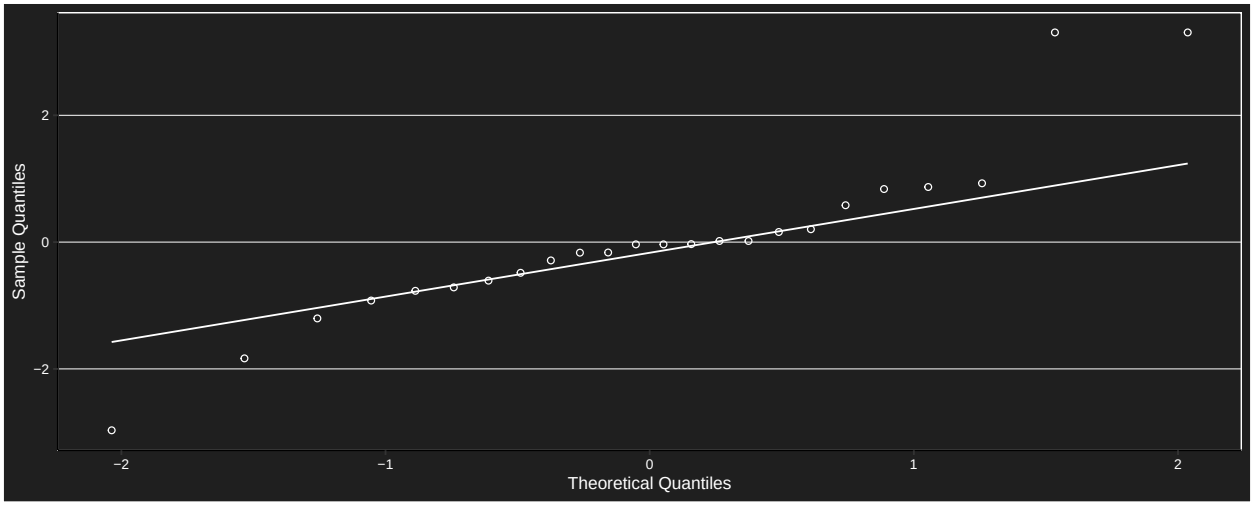
Ácido málico

Modelo y supuestos

```
## Linear mixed-effects model fit by REML
##   Data: mal
##   Log-restricted-likelihood: -26.5820545
##   Fixed: CONF ~ MAD
## (Intercept)       MADMM        MADM       MADSM
##  7.38748867  3.58805750  4.37205000  3.14937817
##
## Random effects:
##  Formula: ~1 | REP
##         (Intercept)   Residual
## StdDev: 0.365216528 2.68012348
##
```

```
## Variance function:
##  Structure: Different standard deviations per stratum
##  Formula: ~1 | MAD
##  Parameter estimates:
##            I            M           MM           SM
## 1.0000000000 0.0552668662 0.2702105687 0.2600800859
## Number of Observations: 24
## Number of Groups: 3
```






```
##
##  Shapiro-Wilk normality test
##
## data:  e
## W = 0.8897338, p-value = 0.0131242
```

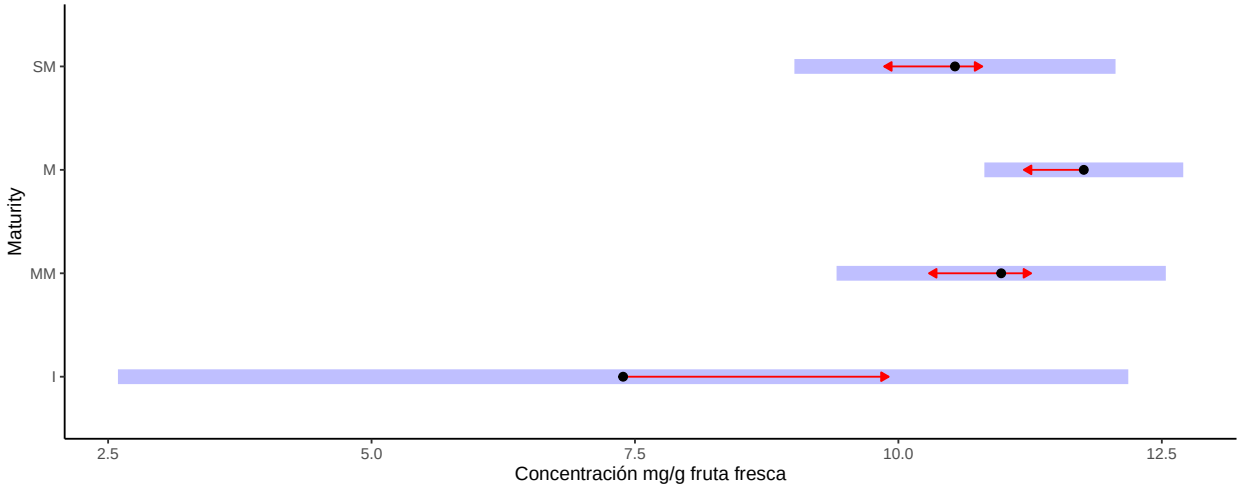
Anova

```
##             numDF denDF     F-value p-value
```

```
## (Intercept)     1    18 2846.522875  <.0001
## MAD             3    18   12.960439   1e-04
```

Test de Tukey

```
## $emmeans
##  MAD emmean    SE df lower.CL upper.CL
##  I     7.39 1.114  2     2.59     12.2
##  MM   10.98 0.363  2     9.41     12.5
##  M    11.76 0.219  2    10.82     12.7
##  SM   10.54 0.354  2     9.01     12.1
##
## Degrees-of-freedom method: containment
## Confidence level used: 0.95
##
## $contrasts
##  contrast estimate    SE df t.ratio p.value
##  I - MM     -3.588 1.133 18  -3.166  0.0251
##  I - M      -4.372 1.096 18  -3.990  0.0043
##  I - SM     -3.149 1.131 18  -2.786  0.0542
##  MM - M     -0.784 0.302 18  -2.598  0.0780
##  MM - SM     0.439 0.410 18   1.069  0.7121
##  M - SM      1.223 0.291 18   4.203  0.0027
##
## Degrees-of-freedom method: containment
## P value adjustment: tukey method for comparing a family of 4 estimates
```

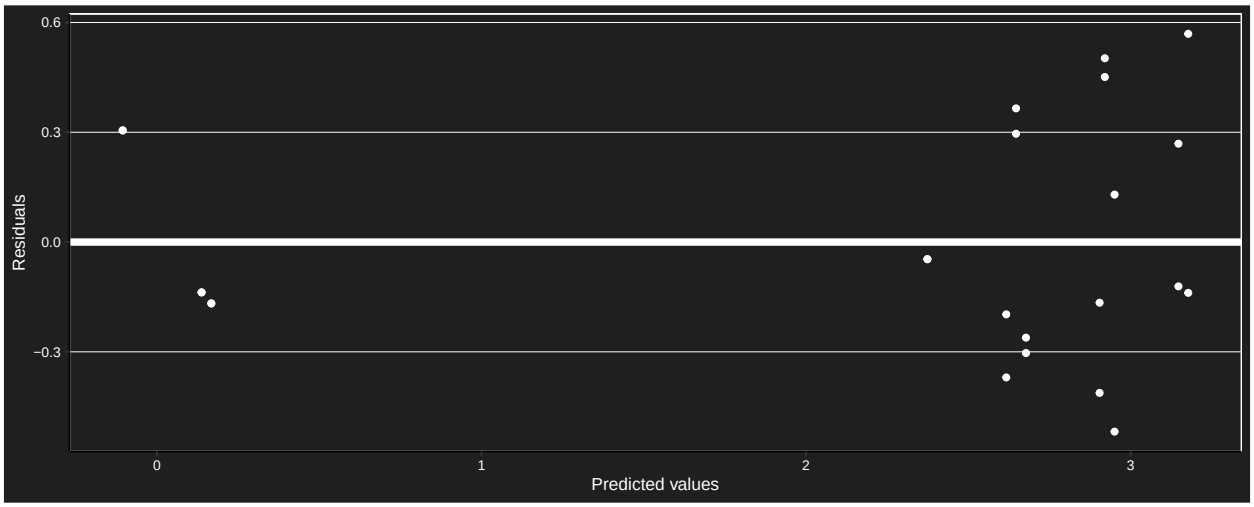


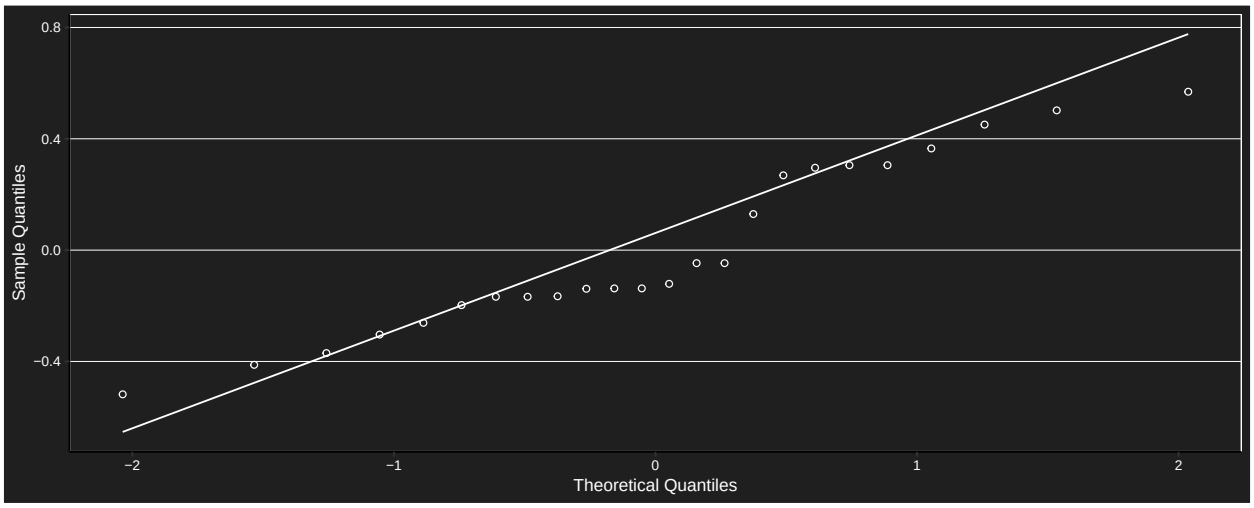
Ácido quínico

Modelo y supuestos

```
## Linear mixed-effects model fit by REML
##   Data: qui
##   Log-restricted-likelihood: -11.8097603
```

```
##   Fixed: CONF ~ MAD
##  (Intercept)        MADMM         MADM        MADSM
## 0.0665693333 3.0094548333 2.4789668333 2.7825800000
##
## Random effects:
##  Formula: ~1 | REP
##         (Intercept)   Residual
## StdDev: 0.180571585 0.34446664
##
## Number of Observations: 24
## Number of Groups: 3
```






```
##
##  Shapiro-Wilk normality test
##
## data:  e
## W = 0.9324113, p-value = 0.110376
```

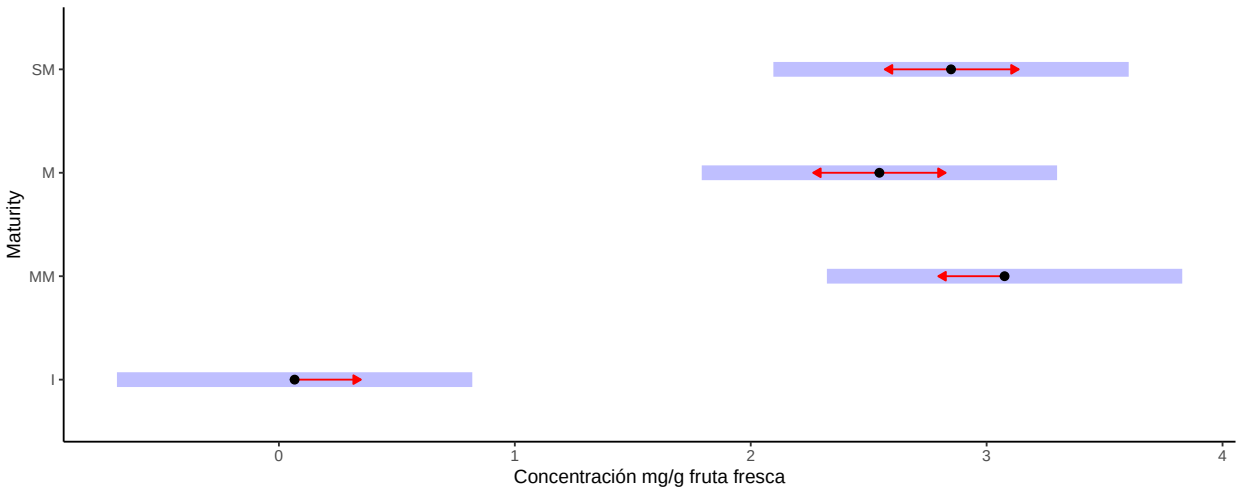
```
## Levene's Test for Homogeneity of Variance (center = median)
##       Df F value   Pr(>F)
## group  3 3.05613 0.052027 .
##       20
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Anova

```
##             numDF denDF     F-value p-value
## (Intercept)     1    18 288.0789394  <.0001
## MAD             3    18  98.4765528  <.0001
```

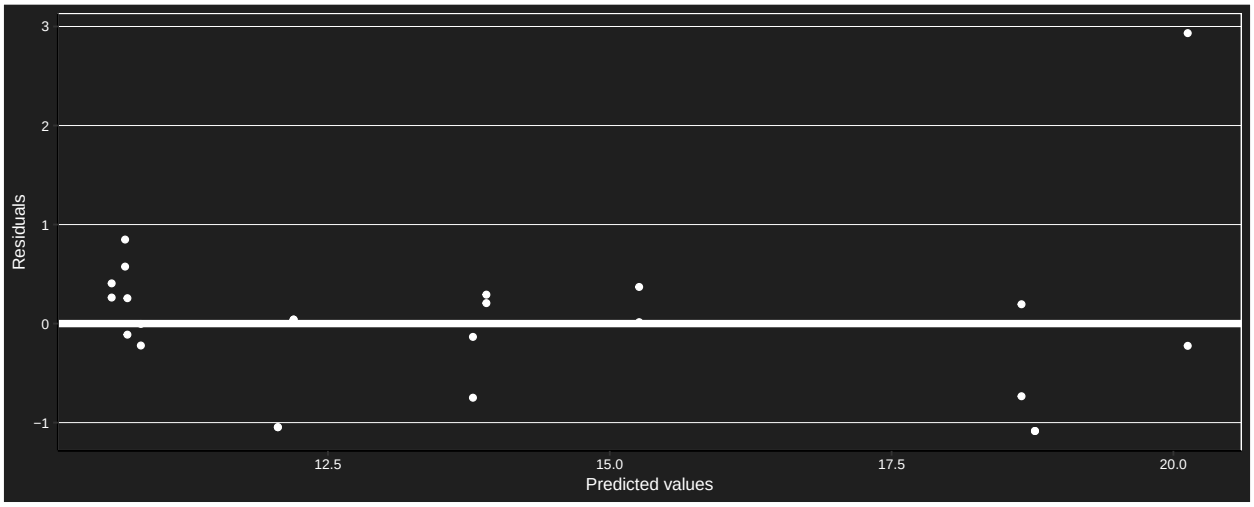
Test de Tukey

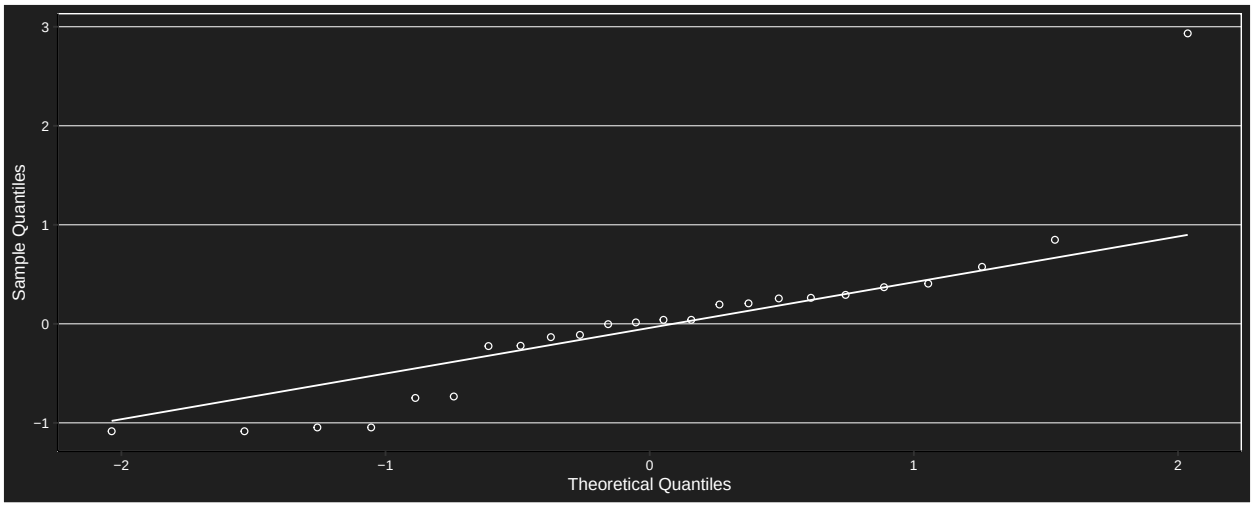
```
## $emmeans
##  MAD emmean    SE df lower.CL upper.CL
##  I   0.0666 0.175  2   -0.687     0.82
##  MM  3.0760 0.175  2    2.323     3.83
##  M   2.5455 0.175  2    1.792     3.30
##  SM  2.8491 0.175  2    2.096     3.60
##
## Degrees-of-freedom method: containment
## Confidence level used: 0.95
##
## $contrasts
##  contrast estimate    SE df t.ratio p.value
##  I - MM     -3.009 0.199 18 -15.132  <.0001
##  I - M      -2.479 0.199 18 -12.465  <.0001
##  I - SM     -2.783 0.199 18 -13.991  <.0001
##  MM - M      0.530 0.199 18   2.667  0.0683
##  MM - SM     0.227 0.199 18   1.141  0.6700
##  M - SM     -0.304 0.199 18  -1.527  0.4431
##
## Degrees-of-freedom method: containment
## P value adjustment: tukey method for comparing a family of 4 estimates
```



Ácido succinico

Modelo y supuestos






```
##
##  Shapiro-Wilk normality test
##
## data:  e
## W = 0.8253994, p-value = 0.000789836
```

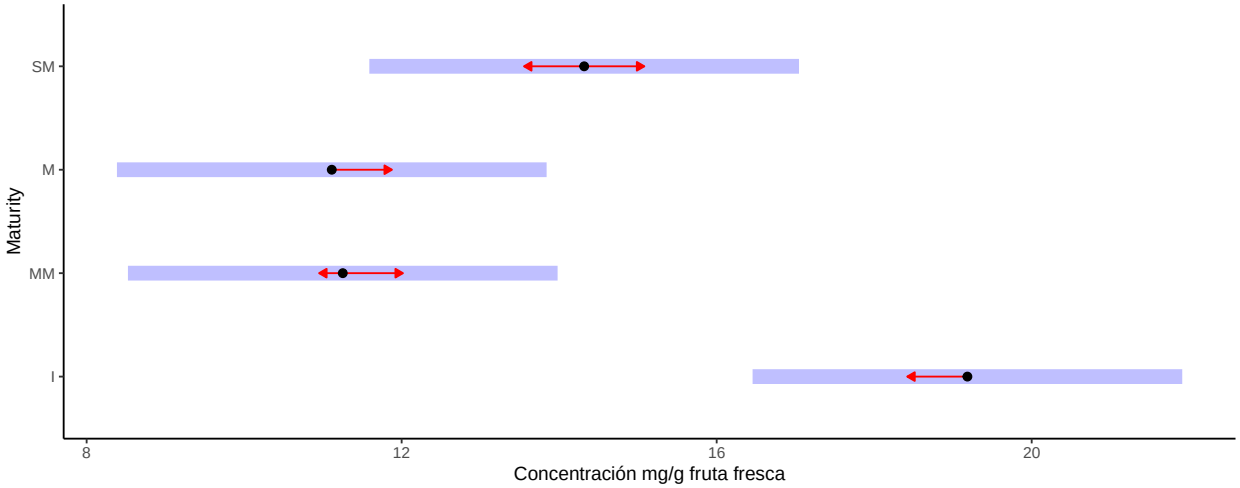
```
## Levene's Test for Homogeneity of Variance (center = median)
##       Df F value Pr(>F)
## group  3 1.89009 0.1638
##       20
```

Anova

```
##             numDF denDF    F-value p-value
## (Intercept)     1    18 667.582635  <.0001
## MAD             3    18  97.574430  <.0001
```

Test de Tukey

```
## $emmeans
##  MAD emmean    SE df lower.CL upper.CL
##  I     19.2 0.634  2    16.46     21.9
##  MM    11.3 0.634  2     8.52     14.0
##  M     11.1 0.634  2     8.38     13.8
##  SM    14.3 0.634  2    11.59     17.0
##
## Degrees-of-freedom method: containment
## Confidence level used: 0.95
##
## $contrasts
##  contrast estimate    SE df t.ratio p.value
##  I - MM       7.93 0.541 18  14.658  <.0001
##  I - M        8.07 0.541 18  14.916  <.0001
##  I - SM       4.87 0.541 18   8.993  <.0001
##  MM - M       0.14 0.541 18   0.258  0.9938
##  MM - SM     -3.07 0.541 18  -5.665  0.0001
##  M - SM      -3.20 0.541 18  -5.923  0.0001
##
## Degrees-of-freedom method: containment
## P value adjustment: tukey method for comparing a family of 4 estimates
```



Acidos orgánicos en peso seco

Concentración del perfíl de ácidos orgánicos

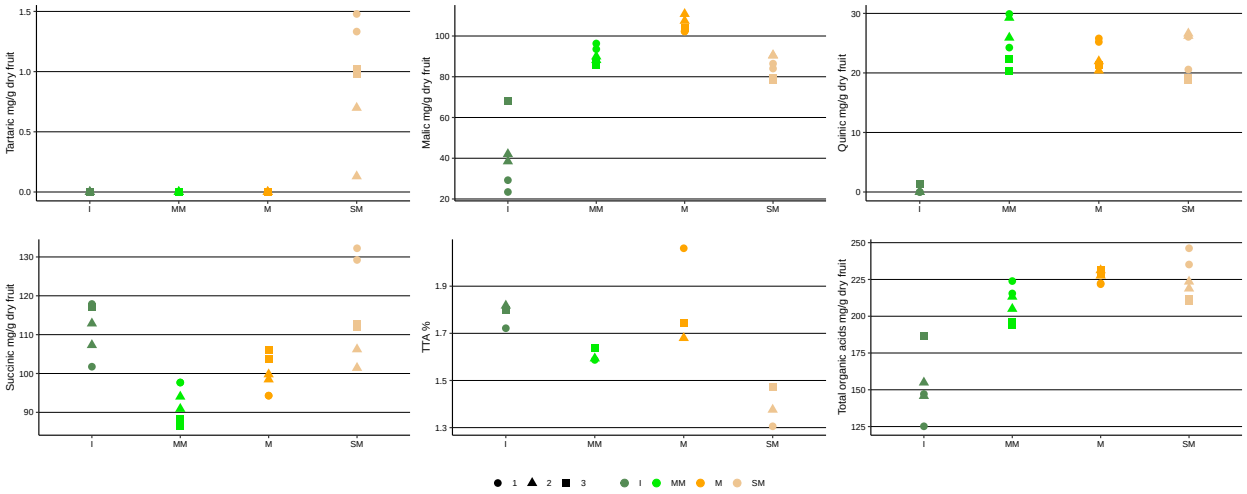


Tabla descriptiva

	CAR	MAD	N	CONS	sd	se	ci
1	Tartárico	I	6	0.000000000	0.0000000000	0.0000000000	0.0000000000
2	Tartárico	MM	6	0.000000000	0.0000000000	0.0000000000	0.0000000000
3	Tartárico	M	6	0.000000000	0.0000000000	0.0000000000	0.0000000000
4	Tartárico	SM	6	0.941135667	0.4830674796	0.1972114727	0.5069482296
5	Málico	I	6	44.902134000	19.1523848051	7.8189283550	20.0991952034
6	Málico	MM	6	90.189531667	4.0220151138	1.6419807944	4.2208460046
7	Málico	M	6	105.147489667	3.3635751544	1.3731738066	3.5298556445
8	Málico	SM	6	84.794599167	5.3680332445	2.1914903952	5.6334054030
9	Quínico	I	6	0.440663333	0.6826727005	0.2786999629	0.7164210623
10	Quínico	MM	6	25.317606333	3.8140013880	1.5570595465	4.0025489871
11	Quínico	M	6	22.688808167	2.2426216210	0.9155464429	2.3534870559
12	Quínico	SM	6	22.920424500	3.7568053163	1.5337093480	3.9425253911
13	Succínico	I	6	112.333486500	6.5345265736	2.6677093027	6.8575650762
14	Succínico	MM	6	92.481592333	4.7805090422	1.9516346440	5.0168365657
15	Succínico	M	6	99.437239833	4.8240138759	1.9693954180	5.0624920887
16	Succínico	SM	6	115.655151333	12.4187930831	5.0699510458	13.0327240659
17	ATT	I	3	1.779200000	0.0507984252	0.0293284844	0.1261902837
18	ATT	MM	3	1.606400000	0.0278969532	0.0161063135	0.0692998736
19	ATT	M	3	1.828266667	0.2039063837	0.1177254055	0.5065315375
20	ATT	SM	3	1.384533333	0.0835275603	0.0482246594	0.2074939626
21	TOTALac	I	6	157.676283833	24.4147956495	9.9672985859	25.6217566954
22	TOTALac	MM	6	207.988730333	11.6569574255	4.7589329410	12.2332265750
23	TOTALac	M	6	227.273537667	4.2846255274	1.7491910468	4.4964387320
24	TOTALac	SM	6	224.311310667	13.9874059778	5.7103345784	14.6788823428
25	<NA>	I	6	266.880791667	8.4819431471	3.4627387896	8.9012534341
26	<NA>	MM	6	301.680354667	34.6167224669	14.1322177686	36.3280222931
27	<NA>	M	6	350.625756167	31.1823867560	12.7301560857	32.7239079987
28	<NA>	SM	6	428.368083167	26.7527791069	10.9217763355	28.0753198610

Evolución del perfíl de ácidos orgánicos

```
## Error in `palette()`:
## ! Insufficient values in manual scale. 6 needed but only 4 provided.
```

Ácidos orgánicos totales

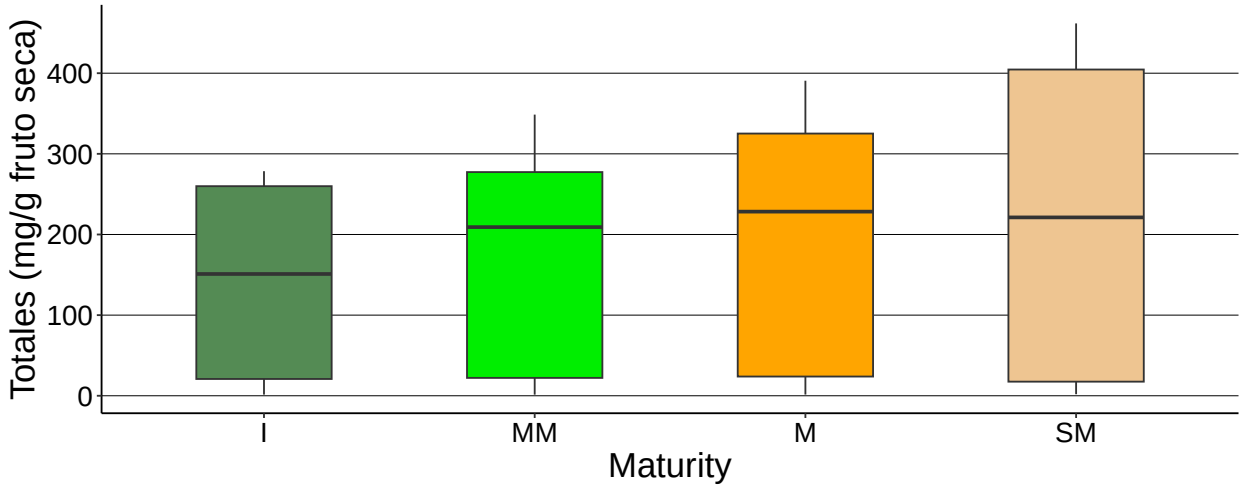


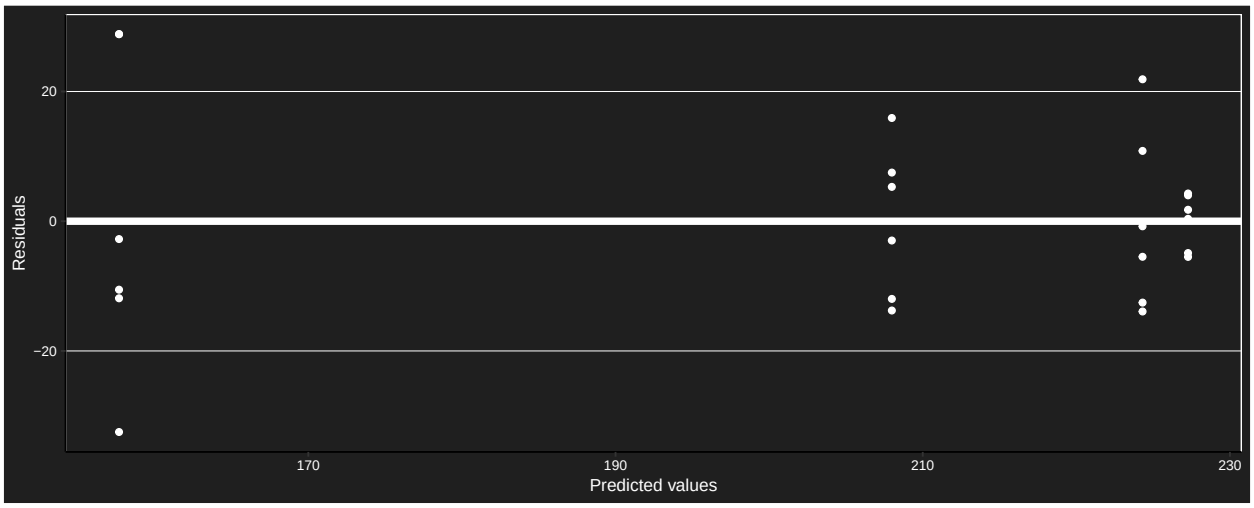
Tabla descriptiva totales

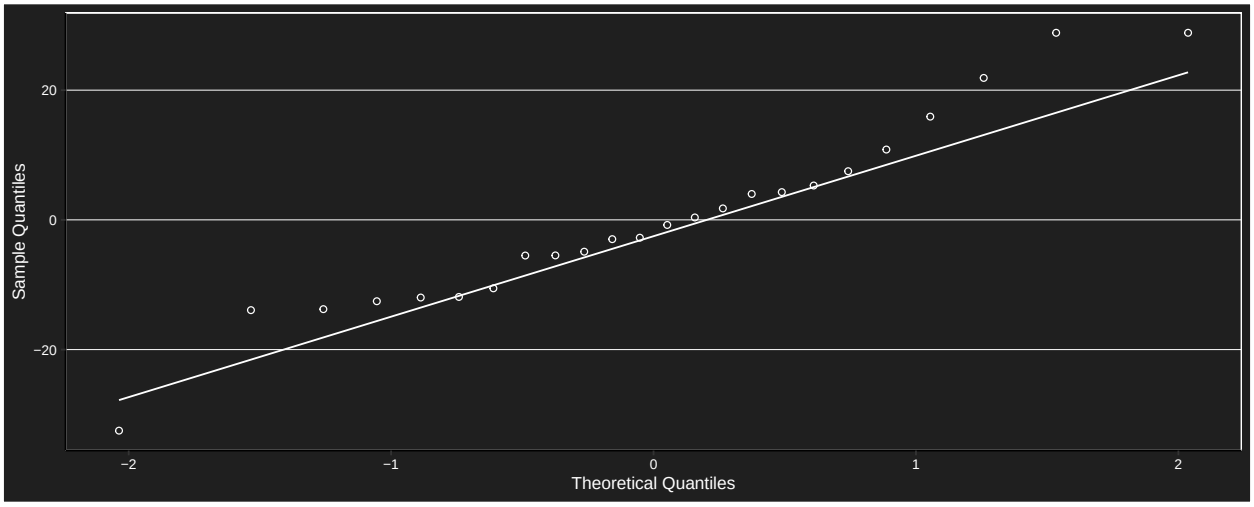
```
##        CAR MAD N       TOTALS             sd             se             ci
## 1    ACIDS   I 6 157.67628383 24.41479564951  9.96729858594 25.6217566954
## 2    ACIDS  MM 6 207.98873017 11.65695733461  4.75893290386 12.2332264797
## 3    ACIDS   M 6 227.27353783  4.28462560986  1.74919108050  4.4964388186
## 4    ACIDS  SM 6 224.31131067 13.98740580960  5.71033450979 14.6788821663
## 5  CATIONS   I 3  20.97175533  7.74194818235  4.46981586713 19.2320654422
## 6  CATIONS  MM 3  21.05341000  2.57785325998  1.48832427358  6.4037424985
## 7  CATIONS   M 3  23.27046433  2.76151278503  1.59436014981  6.8599780508
## 8  CATIONS  SM 3  17.82636733  3.17591985185  1.83361818139  7.8894222735
## 9     STAT   I 3   1.72112247  0.26333177284  0.15203466994  0.6541523876
## 10    STAT  MM 3   1.44765169  0.11629091852  0.06714059311  0.2888826562
## 11    STAT   M 3   1.54171591  0.13269129972  0.07660935761  0.3296234617
## 12    STAT  SM 3   1.90950427  0.00773336868  0.00446486249  0.0192107528
## 13  SUGARS   I 6 266.88079200  8.48194327733  3.46273884278  8.9012535708
## 14  SUGARS  MM 6 301.68035450 34.61672214508 14.13221763719 36.3280219554
## 15  SUGARS   M 6 350.62575617 31.18238675596 12.73015608570 32.7239079987
## 16  SUGARS  SM 6 428.36808300 26.75277856794 10.92177611552 28.0753192954
```

Modelo y supuestos

```
## Linear mixed-effects model fit by REML
##   Data: dataAT
##   Log-restricted-likelihood: -86.6219141
##   Fixed: TOTALS ~ MAD
## (Intercept)       MADMM        MADM       MADSM
## 157.6762838  50.3124463  69.5972540  66.6350268
##
## Random effects:
##  Formula: ~1 | REP
##         (Intercept)   Residual
## StdDev:  0.11875108 15.3782993
```

```
##
## Number of Observations: 24
## Number of Groups: 3
```





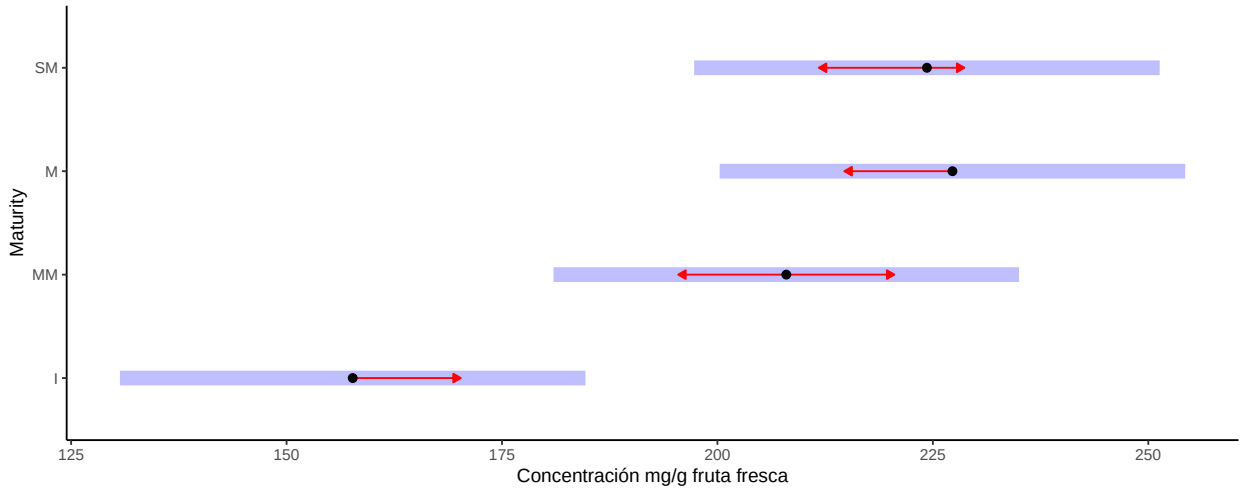

```
##
##  Shapiro-Wilk normality test
##
## data:  e
## W = 0.9558794, p-value = 0.361338
```

Anova

```
##             numDF denDF    F-value p-value
## (Intercept)     1    18 4234.25715  <.0001
## MAD             3    18   26.34856  <.0001
```

Test de Tukey

```
## $emmeans
##  MAD emmean   SE df lower.CL upper.CL
##  I      158 6.28  2      131      185
##  MM     208 6.28  2      181      235
##  M      227 6.28  2      200      254
##  SM     224 6.28  2      197      251
##
## Degrees-of-freedom method: containment
## Confidence level used: 0.95
##
## $contrasts
##  contrast estimate   SE df t.ratio p.value
##  I - MM     -50.31 8.88 18  -5.667  0.0001
##  I - M      -69.60 8.88 18  -7.839  <.0001
##  I - SM     -66.64 8.88 18  -7.505  <.0001
##  MM - M     -19.28 8.88 18  -2.172  0.1689
##  MM - SM    -16.32 8.88 18  -1.838  0.2886
##  M - SM       2.96 8.88 18   0.334  0.9868
##
## Degrees-of-freedom method: containment
## P value adjustment: tukey method for comparing a family of 4 estimates
```

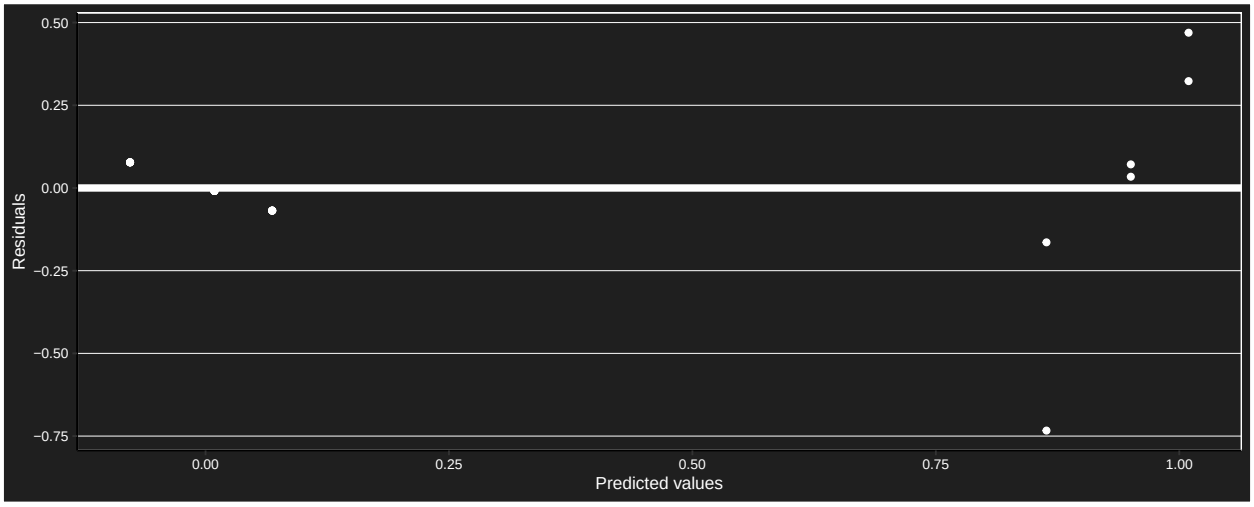


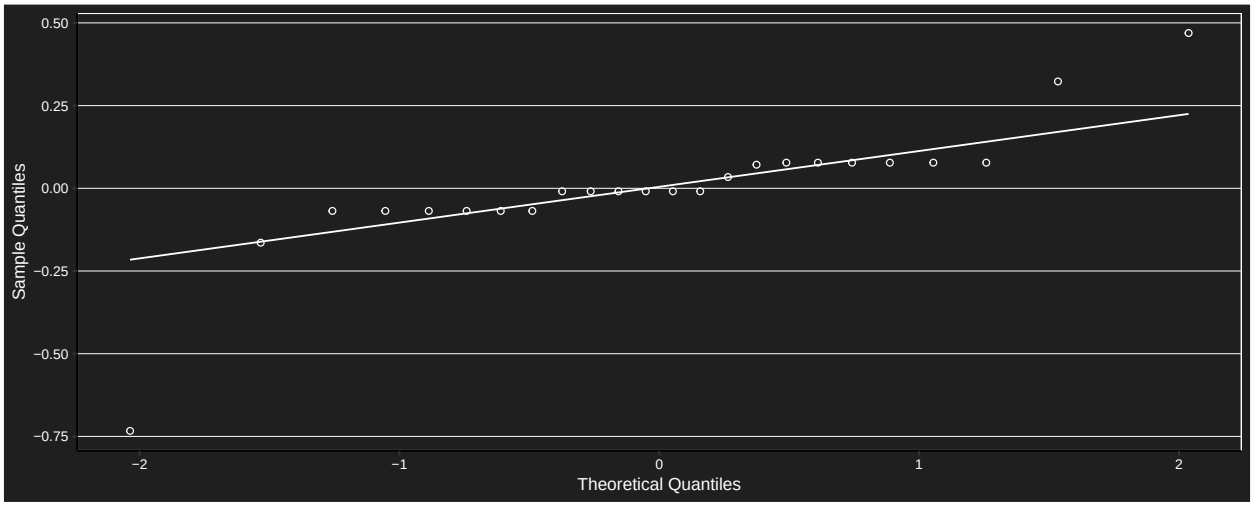
Ácido tartárico

Modelo y supuestos

```
## Linear mixed-effects model fit by REML
##   Data: tar
##   Log-restricted-likelihood: -3.09688774
##   Fixed: CONS ~ MAD
##     (Intercept)           MADMM            MADM           MADSM
##  8.70989721e-17 -1.26351284e-16 -2.22044605e-16  9.41135667e-01
##
## Random effects:
##  Formula: ~1 | REP
##         (Intercept)    Residual
```

```
## StdDev:  0.09563016 0.225881487
##
## Number of Observations: 24
## Number of Groups: 3
```





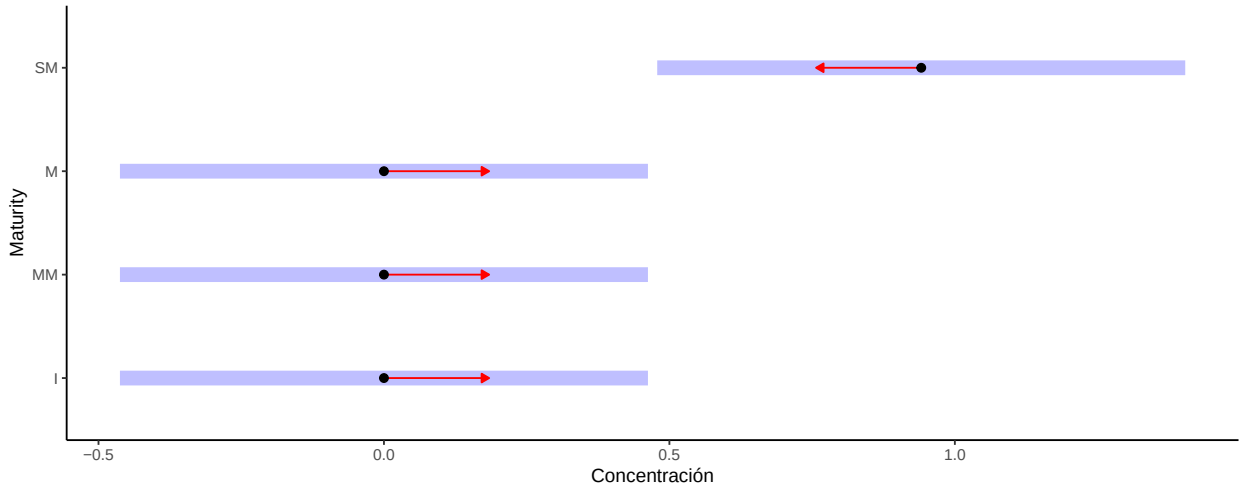

```
##
##  Shapiro-Wilk normality test
##
## data:  e
## W = 0.7503611, p-value = 5.0413e-05
```

Anova

```
##             numDF denDF    F-value p-value
## (Intercept)     1    18 10.6987231  0.0042
## MAD             3    18 26.0396084  <.0001
```

Test de Tukey

```
## $emmeans
##  MAD emmean    SE df lower.CL upper.CL
##  I    0.000 0.107  2   -0.462    0.462
##  MM   0.000 0.107  2   -0.462    0.462
##  M    0.000 0.107  2   -0.462    0.462
##  SM   0.941 0.107  2    0.479    1.404
##
## Degrees-of-freedom method: containment
## Confidence level used: 0.95
##
## $contrasts
##  contrast estimate   SE df t.ratio p.value
##  I - MM      0.000 0.13 18   0.000  1.0000
##  I - M       0.000 0.13 18   0.000  1.0000
##  I - SM     -0.941 0.13 18  -7.217  <.0001
##  MM - M      0.000 0.13 18   0.000  1.0000
##  MM - SM    -0.941 0.13 18  -7.217  <.0001
##  M - SM     -0.941 0.13 18  -7.217  <.0001
##
## Degrees-of-freedom method: containment
## P value adjustment: tukey method for comparing a family of 4 estimates
```

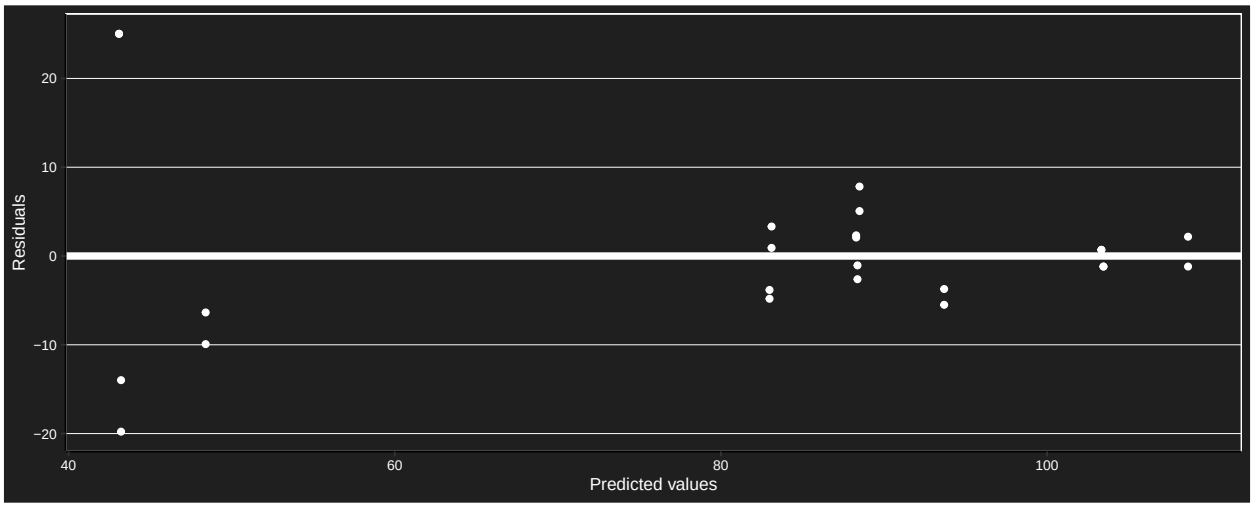


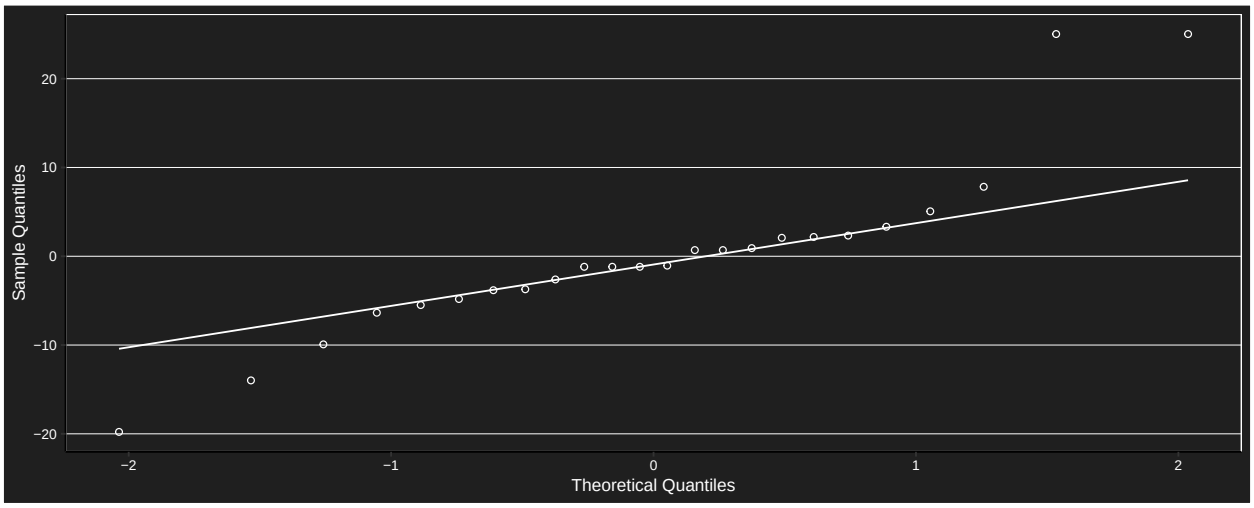
Ácido málico

Modelo y supuestos

```
## Linear mixed-effects model fit by REML
##   Data: mal
##   Log-restricted-likelihood: -66.5285047
##   Fixed: CONS ~ MAD
## (Intercept)       MADMM        MADM       MADSM
##  44.9021340  45.2873977  60.2453557  39.8924652
##
## Random effects:
##  Formula: ~1 | REP
```

```
##         (Intercept)   Residual
## StdDev:  3.18440631 19.9143832
##
## Variance function:
##  Structure: Different standard deviations per stratum
##  Formula: ~1 | MAD
##  Parameter estimates:
##            I            M           MM           SM
## 1.0000000000 0.0829966037 0.2681538614 0.1785082895
## Number of Observations: 24
## Number of Groups: 3
```





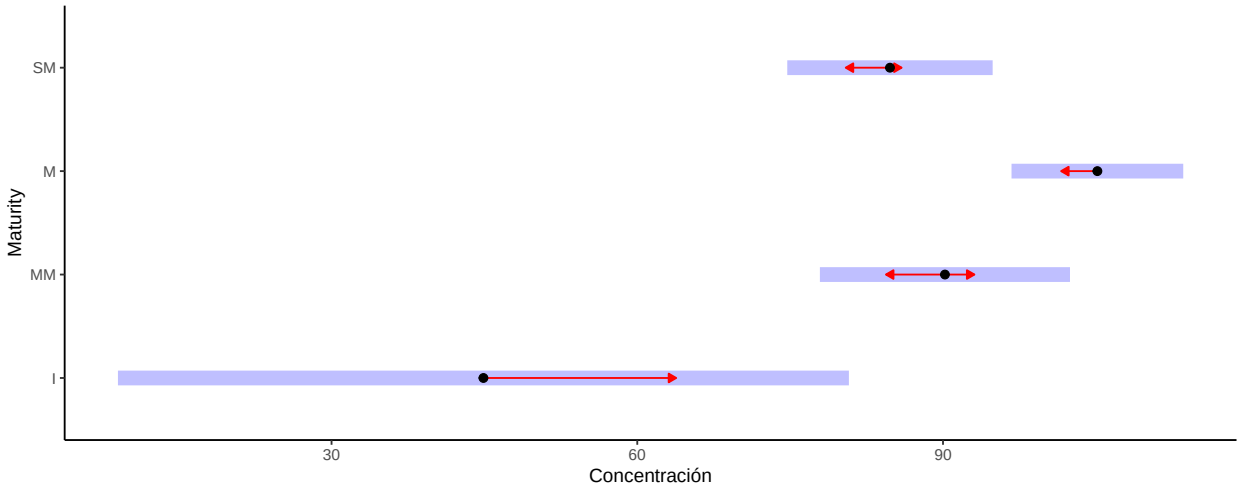

```
##
##  Shapiro-Wilk normality test
##
## data:  e
## W = 0.872026, p-value = 0.00576551
```

Anova

```
##             numDF denDF    F-value p-value
## (Intercept)     1    18 2706.36602  <.0001
## MAD             3    18   77.88670  <.0001
```

Test de Tukey

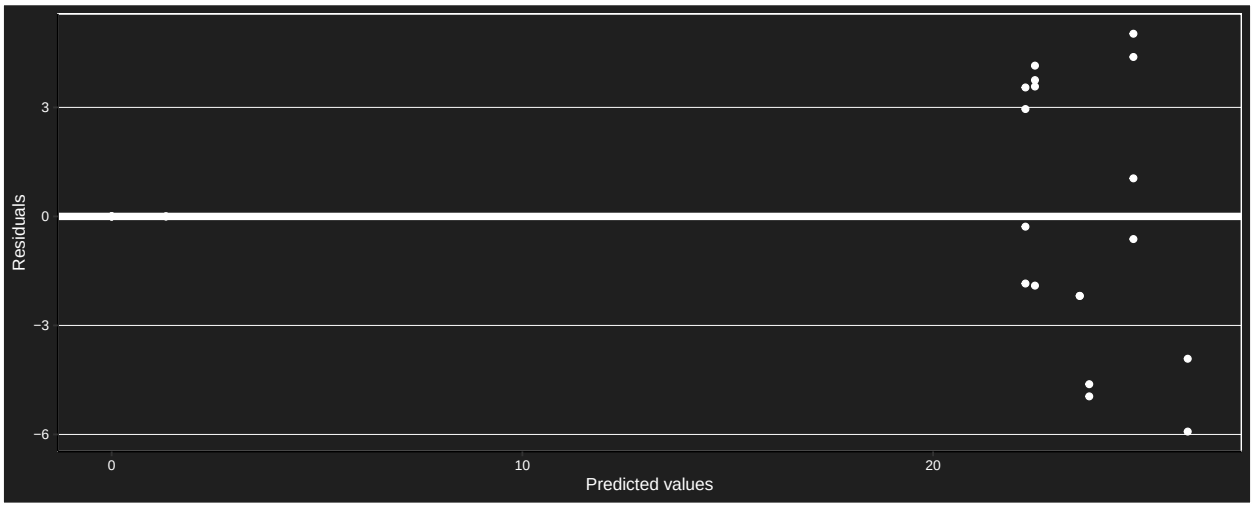
```
## $emmeans
##  MAD emmean   SE df lower.CL upper.CL
##  I     44.9 8.34  2     9.04     80.8
##  MM    90.2 2.85  2    77.92    102.5
##  M    105.1 1.96  2    96.72    113.6
##  SM    84.8 2.34  2    74.72     94.9
##
## Degrees-of-freedom method: containment
## Confidence level used: 0.95
##
## $contrasts
##  contrast estimate   SE df t.ratio p.value
##  I - MM     -45.29 8.42 18  -5.380  0.0002
##  I - M      -60.25 8.16 18  -7.385  <.0001
##  I - SM     -39.89 8.26 18  -4.830  0.0007
##  MM - M     -14.96 2.28 18  -6.554  <.0001
##  MM - SM      5.39 2.62 18   2.060  0.2038
##  M - SM      20.35 1.60 18  12.717  <.0001
##
## Degrees-of-freedom method: containment
## P value adjustment: tukey method for comparing a family of 4 estimates
```

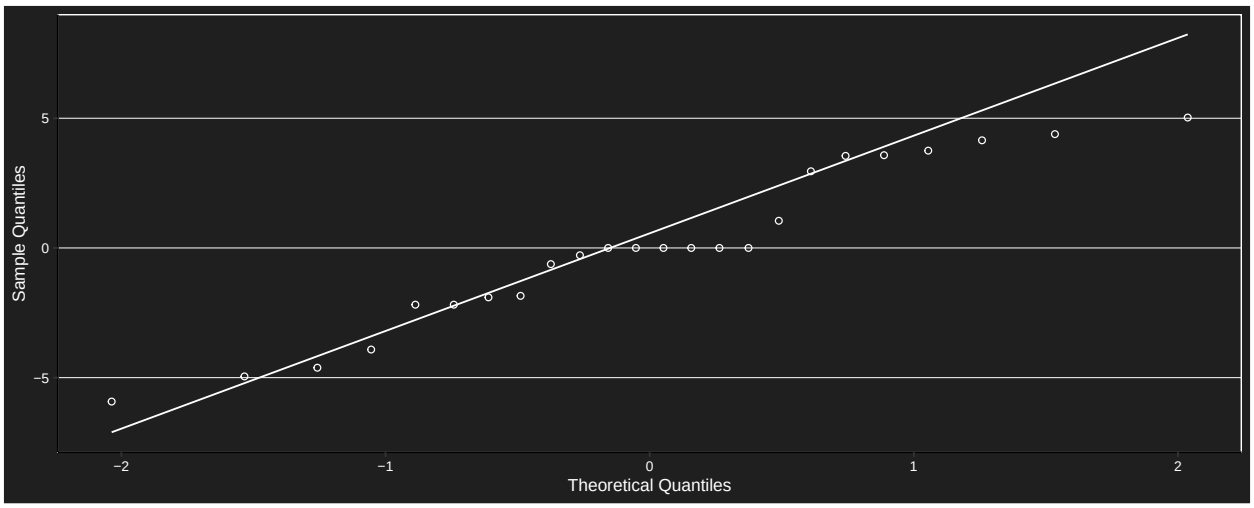


Ácido quínico

Modelo y supuestos

```
## Linear mixed-effects model fit by REML
##   Data: qui
##   Log-restricted-likelihood: 58.1548863
##   Fixed: CONS ~ MAD
##  (Intercept)        MADMM         MADM        MADSM
##  0.374677755 24.876943000 22.248144833 22.479761167
##
## Random effects:
##  Formula: ~1 | REP
##         (Intercept)       Residual
## StdDev: 0.671027165 1.17940857e-16
##
## Variance function:
##  Structure: Different standard deviations per stratum
##  Formula: ~1 | MAD
##  Parameter estimates:
##              I              M             MM             SM
## 1.00000000e+00 2.22331623e+16 3.72325836e+16 3.66668232e+16
## Number of Observations: 24
## Number of Groups: 3
```





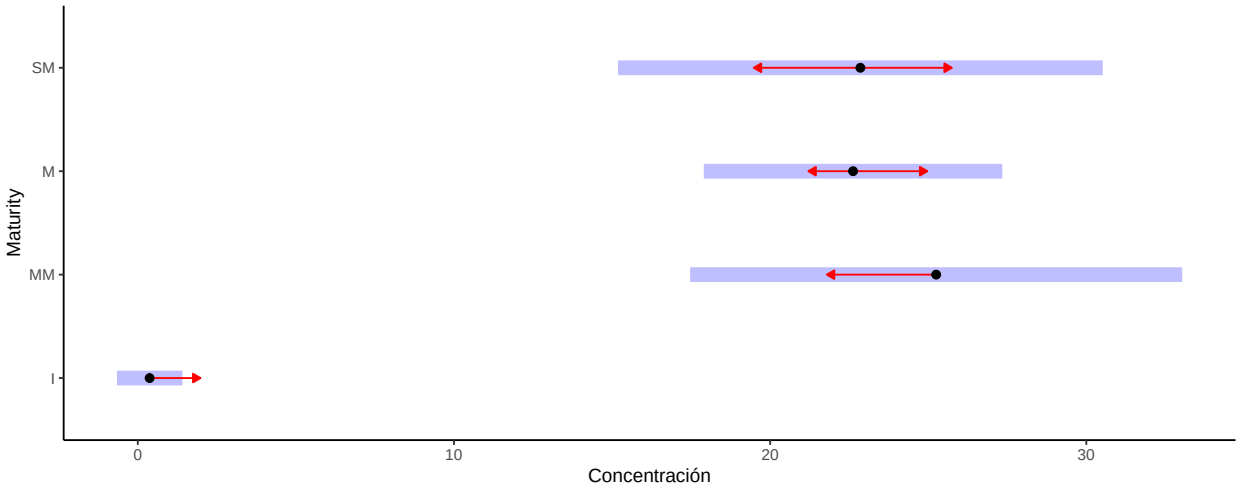

```
##
##  Shapiro-Wilk normality test
##
## data:  e
## W = 0.9457795, p-value = 0.219116
```

Anova

```
##             numDF denDF     F-value p-value
## (Intercept)     1    18   2.4280693  0.1366
## MAD             3    18 262.2049802  <.0001
```

Test de Tukey

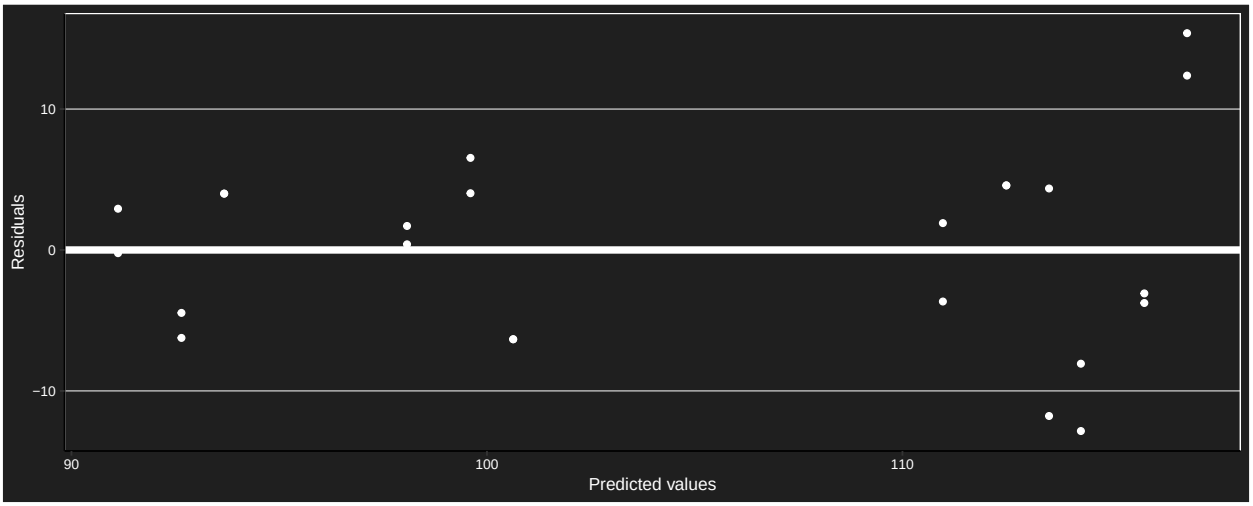
```
## $emmeans
##  MAD emmean   SE df lower.CL upper.CL
##  I    0.375 0.24  2    -0.66     1.41
##  MM  25.252 1.81  2    17.47    33.03
##  M   22.623 1.10  2    17.90    27.34
##  SM  22.854 1.78  2    15.19    30.52
##
## Degrees-of-freedom method: containment
## Confidence level used: 0.95
##
## $contrasts
##  contrast estimate   SE df t.ratio p.value
##  I - MM    -24.877 1.79 18 -13.877  <.0001
##  I - M     -22.248 1.07 18 -20.783  <.0001
##  I - SM    -22.480 1.77 18 -12.733  <.0001
##  MM - M      2.629 2.09 18   1.259  0.5992
##  MM - SM     2.397 2.52 18   0.953  0.7772
##  M - SM     -0.232 2.06 18  -0.112  0.9995
##
## Degrees-of-freedom method: containment
## P value adjustment: tukey method for comparing a family of 4 estimates
```

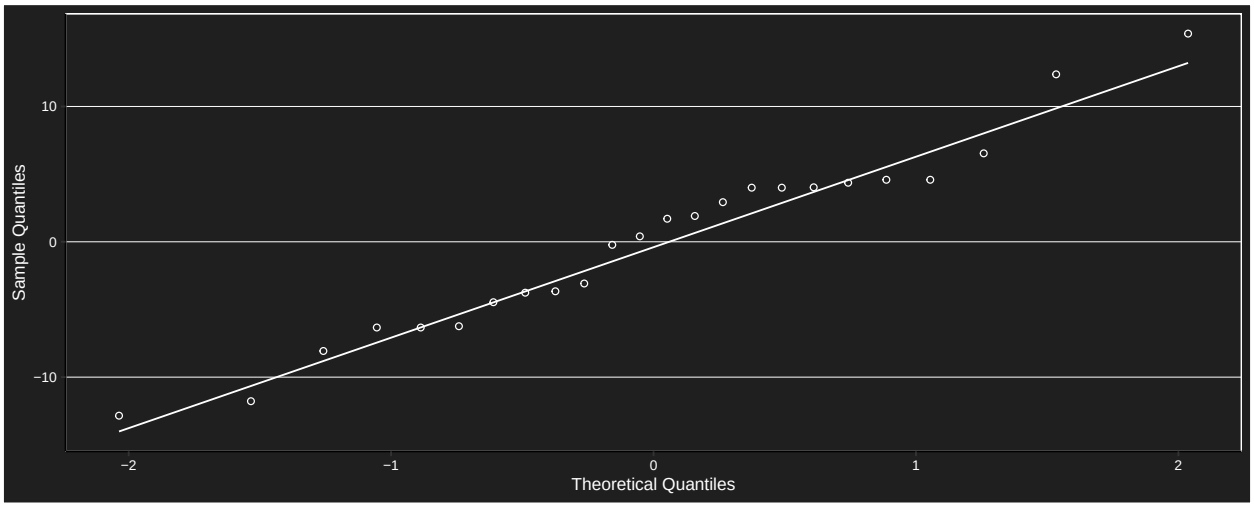


Ácido succinico

Modelo y supuestos

```
## Linear mixed-effects model fit by REML
##   Data: suc
##   Log-restricted-likelihood: -72.9162935
##   Fixed: CONS ~ MAD
##  (Intercept)        MADMM         MADM        MADSM
## 112.33348650 -19.85189417 -12.89624667   3.32166483
##
## Random effects:
##  Formula: ~1 | REP
##         (Intercept)   Residual
## StdDev:  2.08906489 7.56778496
##
## Number of Observations: 24
## Number of Groups: 3
```






```
##
##  Shapiro-Wilk normality test
##
## data:  e
## W = 0.9675281, p-value = 0.606483

## Levene's Test for Homogeneity of Variance (center = median)
##       Df F value  Pr(>F)
## group  3 1.48182 0.24974
##       20
```

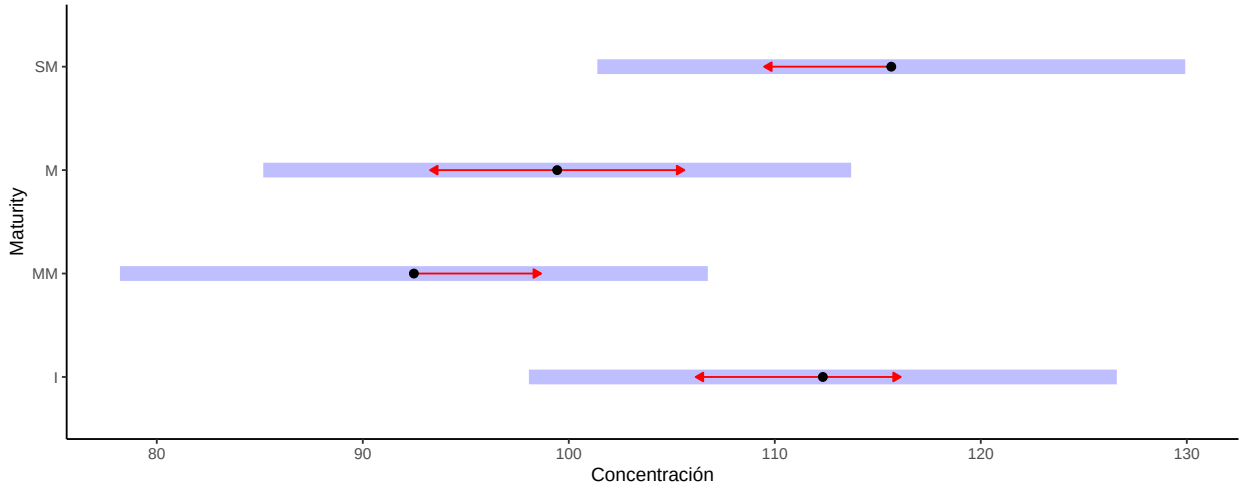
Anova

```
##             numDF denDF     F-value p-value
## (Intercept)     1    18 2869.053469  <.0001
## MAD             3    18   12.395896   1e-04
```

Test de Tukey

```
## $emmeans
##  MAD emmean   SE df lower.CL upper.CL
##  I    112.3 3.32  2     98.1      127
##  MM    92.5 3.32  2     78.2      107
##  M     99.4 3.32  2     85.2      114
##  SM   115.7 3.32  2    101.4      130
##
## Degrees-of-freedom method: containment
## Confidence level used: 0.95
##
## $contrasts
##  contrast estimate   SE df t.ratio p.value
##  I - MM      19.85 4.37 18   4.544  0.0013
##  I - M       12.90 4.37 18   2.952  0.0389
##  I - SM      -3.32 4.37 18  -0.760  0.8711
##  MM - M      -6.96 4.37 18  -1.592  0.4076
```

```
##  MM - SM    -23.17 4.37 18  -5.304  0.0003
##  M - SM     -16.22 4.37 18  -3.712  0.0079
##
## Degrees-of-freedom method: containment
## P value adjustment: tukey method for comparing a family of 4 estimates
```



Relación de ácidos orgánicos y acidez total titulable ATT.

	CAR	MAD	N	CONS	sd	se	ci
1	Tartárico	I	6	0.000000000	0.0000000000	0.0000000000	0.0000000000
2	Tartárico	MM	6	0.000000000	0.0000000000	0.0000000000	0.0000000000
3	Tartárico	M	6	0.000000000	0.0000000000	0.0000000000	0.0000000000
4	Tartárico	SM	6	0.941135667	0.4830674796	0.1972114727	0.5069482296
5	Málico	I	6	44.902134000	19.1523848051	7.8189283550	20.0991952034
6	Málico	MM	6	90.189531667	4.0220151138	1.6419807944	4.2208460046
7	Málico	M	6	105.147489667	3.3635751544	1.3731738066	3.5298556445
8	Málico	SM	6	84.794599167	5.3680332445	2.1914903952	5.6334054030
9	Quínico	I	6	0.440663333	0.6826727005	0.2786999629	0.7164210623
10	Quínico	MM	6	25.317606333	3.8140013880	1.5570595465	4.0025489871
11	Quínico	M	6	22.688808167	2.2426216210	0.9155464429	2.3534870559
12	Quínico	SM	6	22.920424500	3.7568053163	1.5337093480	3.9425253911
13	Succínico	I	6	112.333486500	6.5345265736	2.6677093027	6.8575650762
14	Succínico	MM	6	92.481592333	4.7805090422	1.9516346440	5.0168365657
15	Succínico	M	6	99.437239833	4.8240138759	1.9693954180	5.0624920887
16	Succínico	SM	6	115.655151333	12.4187930831	5.0699510458	13.0327240659
17	ATT	I	3	1.779200000	0.0507984252	0.0293284844	0.1261902837
18	ATT	MM	3	1.606400000	0.0278969532	0.0161063135	0.0692998736
19	ATT	M	3	1.828266667	0.2039063837	0.1177254055	0.5065315375
20	ATT	SM	3	1.384533333	0.0835275603	0.0482246594	0.2074939626
21	TOTALac	I	6	157.676283833	24.4147956495	9.9672985859	25.6217566954
22	TOTALac	MM	6	207.988730333	11.6569574255	4.7589329410	12.2332265750
23	TOTALac	M	6	227.273537667	4.2846255274	1.7491910468	4.4964387320
24	TOTALac	SM	6	224.311310667	13.9874059778	5.7103345784	14.6788823428
25	<NA>	I	6	266.880791667	8.4819431471	3.4627387896	8.9012534341
26	<NA>	MM	6	301.680354667	34.6167224669	14.1322177686	36.3280222931

```
## 27      <NA>   M 6 350.625756167 31.1823867560 12.7301560857 32.7239079987
## 28      <NA>  SM 6 428.368083167 26.7527791069 10.9217763355 28.0753198610
## 29     ACIDS   I 6 157.676283833 24.4147956495  9.9672985859 25.6217566954
## 30     ACIDS  MM 6 207.988730167 11.6569573346  4.7589329039 12.2332264797
## 31     ACIDS   M 6 227.273537833  4.2846256099  1.7491910805  4.4964388186
## 32     ACIDS  SM 6 224.311310667 13.9874058096  5.7103345098 14.6788821663
```

Concentración del ratio azúcares totales / ácidos orgánicos totales a distintos estados.

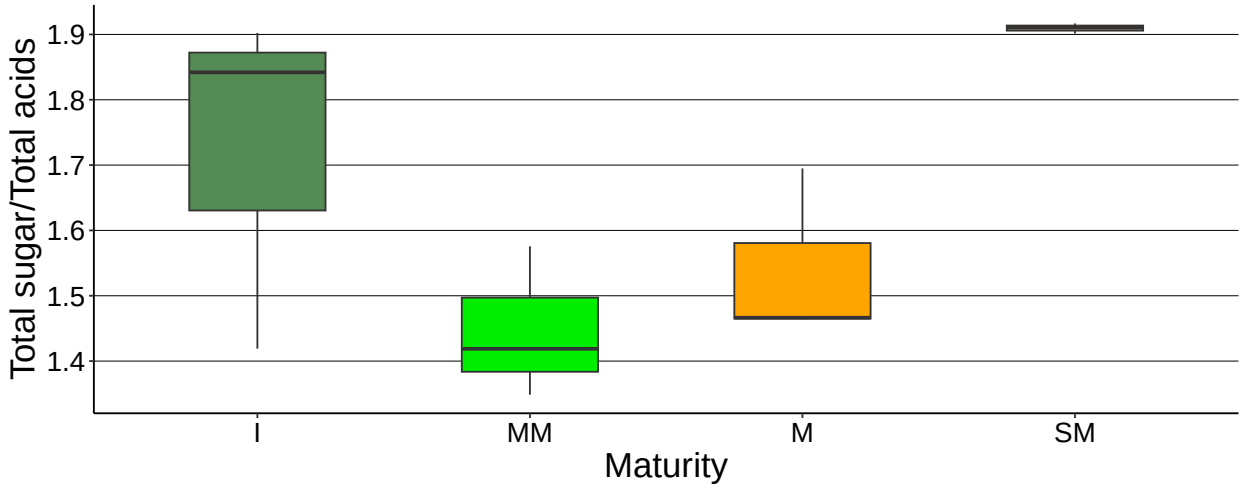
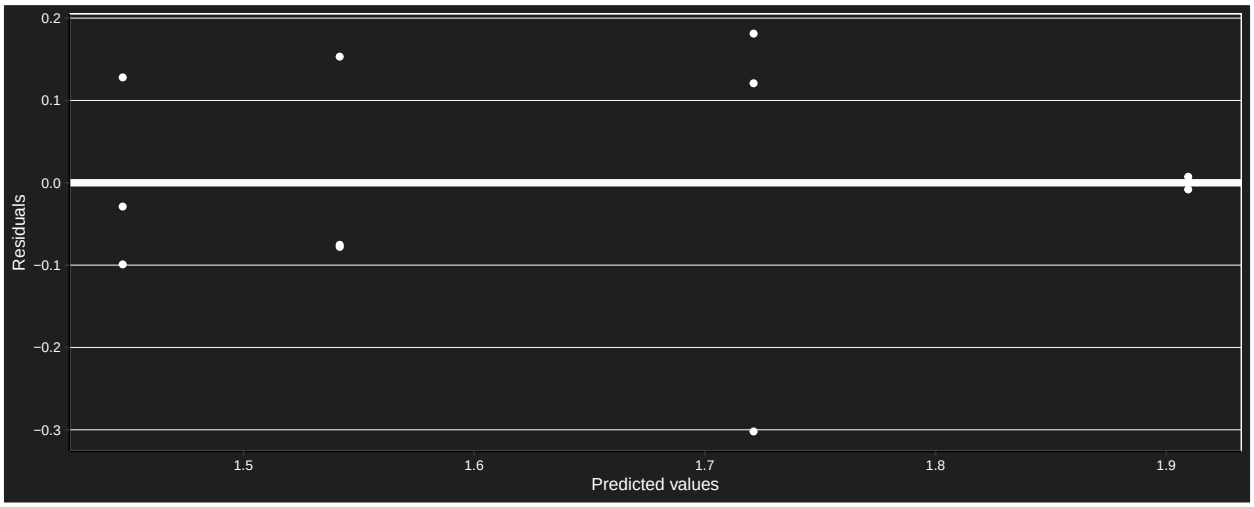


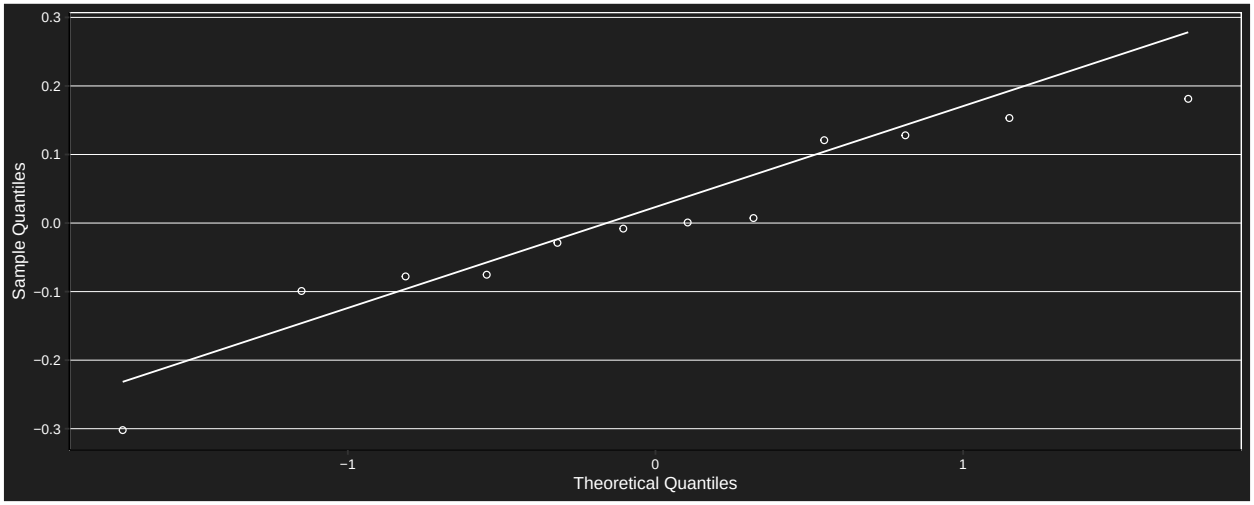
Tabla descriptiva totales

```
##   MAD N     TOTALS            sd            se           ci
## 1   I 3 1.72112247 0.26333177284 0.15203466994 0.6541523876
## 2  MM 3 1.44765169 0.11629091852 0.06714059311 0.2888826562
## 3   M 3 1.54171591 0.13269129972 0.07660935761 0.3296234617
## 4  SM 3 1.90950427 0.00773336868 0.00446486249 0.0192107528
```

Relación ST/AT (azúcares totales / ácidos totales)

```
## Linear mixed-effects model fit by REML
##   Data: dataSTAT
##   Log-restricted-likelihood: 1.18539612
##   Fixed: TOTALS ~ MAD
##  (Intercept)        MADMM         MADM        MADSM
##  1.721122474 -0.273470784 -0.179406567  0.188381795
##
## Random effects:
##  Formula: ~1 | REP
##            (Intercept)    Residual
## StdDev: 0.000999703726 0.158534031
##
## Number of Observations: 12
## Number of Groups: 3
```






```
##
##  Shapiro-Wilk normality test
##
## data:  e
## W = 0.9262267, p-value = 0.341827
```

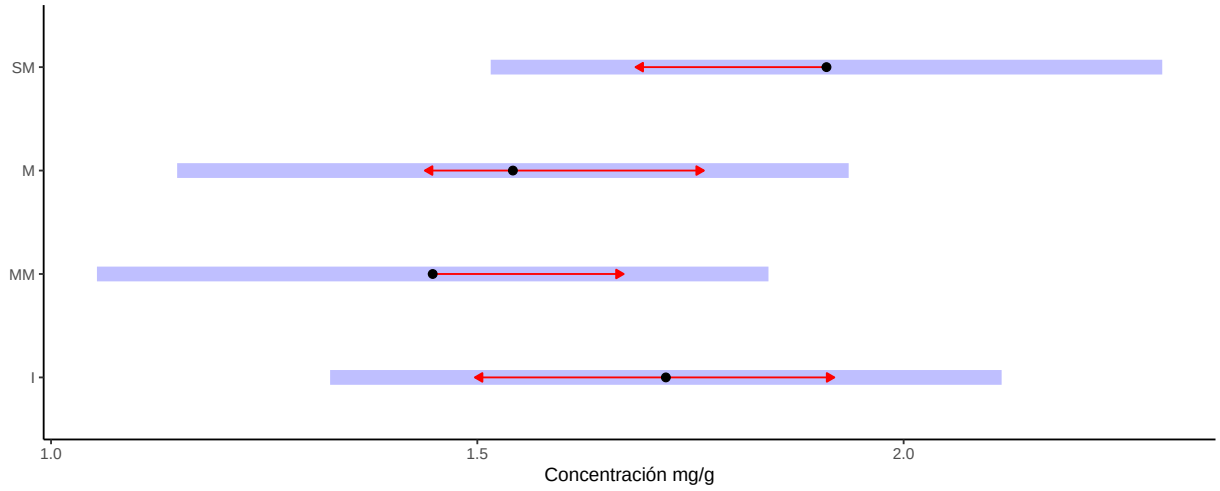
Anova

```
##             numDF denDF     F-value p-value
## (Intercept)     1     6 1307.562391  <.0001
## MAD             3     6    4.972388  0.0457
```

Test de Tukey

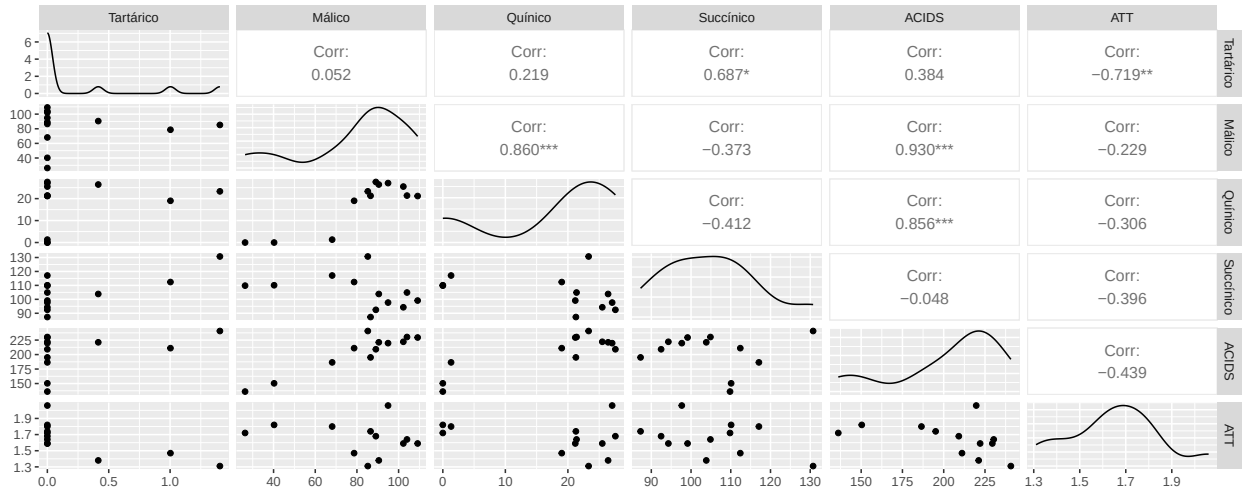
```
## $emmeans
##  MAD emmean     SE df lower.CL upper.CL
##  I     1.72 0.0915  2     1.33     2.11
```

```
##  MM    1.45 0.0915  2     1.05     1.84
##  M     1.54 0.0915  2     1.15     1.94
##  SM    1.91 0.0915  2     1.52     2.30
##
## Degrees-of-freedom method: containment
## Confidence level used: 0.95
##
## $contrasts
##  contrast estimate    SE df t.ratio p.value
##  I - MM     0.2735 0.129  6   2.113  0.2498
##  I - M      0.1794 0.129  6   1.386  0.5497
##  I - SM    -0.1884 0.129  6  -1.455  0.5142
##  MM - M    -0.0941 0.129  6  -0.727  0.8831
##  MM - SM   -0.4619 0.129  6  -3.568  0.0443
##  M - SM    -0.3678 0.129  6  -2.841  0.1041
##
## Degrees-of-freedom method: containment
## P value adjustment: tukey method for comparing a family of 4 estimates
```



Correlaciones

Correlaciones de Pearson.




```
##
##  Pearson's product-moment correlation
##
## data:  FACO$Málico and FACO$Quínico
## t = 5.329902, df = 10, p-value = 0.000333007
## alternative hypothesis: true correlation is not equal to 0
## 95 percent confidence interval:
##  0.564971048 0.960065608
## sample estimates:
##       cor
## 0.860021264
```

```
##
##  Pearson's product-moment correlation
##
## data:  FACO$Tartárico and FACO$Succínico
## t = 2.989547, df = 10, p-value = 0.0135842
## alternative hypothesis: true correlation is not equal to 0
## 95 percent confidence interval:
##  0.186681215 0.904338855
## sample estimates:
##       cor
## 0.686981893
```

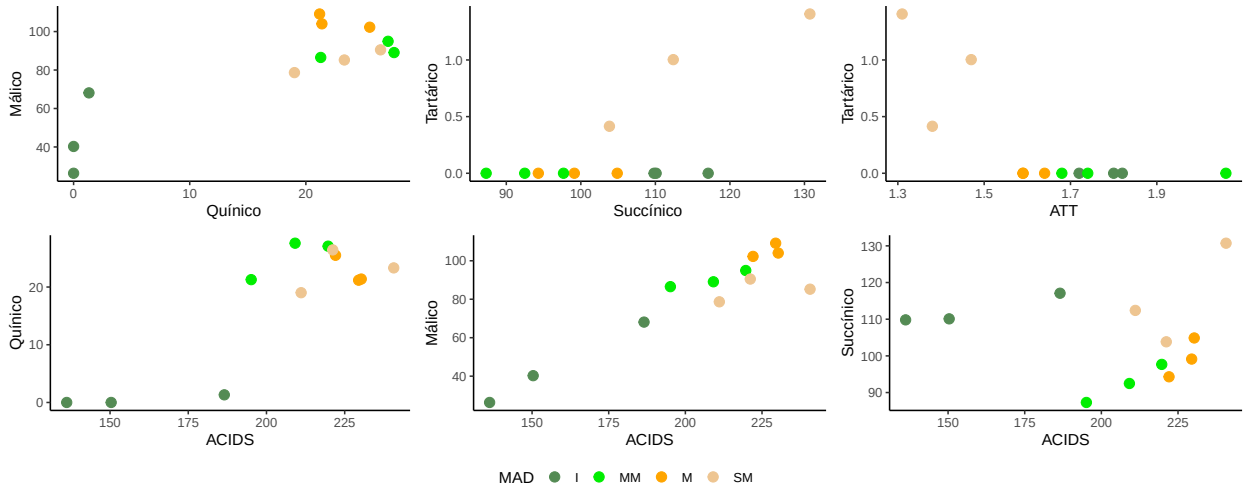
```
##
##  Pearson's product-moment correlation
##
## data:  FACO$ATT and FACO$Tartárico
## t = -3.268633, df = 10, p-value = 0.00844982
## alternative hypothesis: true correlation is not equal to 0
## 95 percent confidence interval:
##  -0.915141012 -0.246455115
## sample estimates:
##       cor
## -0.71870271
```

```
##
```

```
##  Pearson's product-moment correlation
##
## data:  FACO$ACIDS and FACO$Quínico
## t = 5.230734, df = 10, p-value = 0.000383888
## alternative hypothesis: true correlation is not equal to 0
## 95 percent confidence interval:
##  0.553901763 0.958784719
## sample estimates:
##       cor
## 0.855767597
```

```
##
##  Pearson's product-moment correlation
##
## data:  FACO$ACIDS and FACO$Málico
## t = 8.006896, df = 10, p-value = 1.16864e-05
## alternative hypothesis: true correlation is not equal to 0
## 95 percent confidence interval:
##  0.763990213 0.980578157
## sample estimates:
##       cor
## 0.930089316
```

Gráficos de correlación detallados por estado.



- Correlaciones: Se evidenciaron relaciones lineales entre los ácidos orgánicos, entre el ácido málico y el ácido quínico con un coeficiente de correlación (r) de 0.8600213 y un valor de p=0.000333, y entre el ácido tartárico y el ácido succínico con un r=0.6869819 y un p-valor=0.01358. La acidez titulable total (TTA) mostró una asociación lineal significativa únicamente con el ácido tartárico, con un r=-0.7187027 y un p-valor=0.00845. Sin embargo, esta asociación inversa está vinculada al hecho de que el ácido tartárico solo aparece en cantidades mínimas en frutas muy maduras. La concentración total de ácidos con ácido quínico presentó una correlación de 0.8557676 con un p-valor=0.0003839. Mientras tanto, el ácido málico mostró un r=0.9300893 y un p-valor=1.169e-05. En ambos casos, estos ácidos explican el aumento en la concentración total de ácidos a lo largo del proceso de maduración de la fruta.