Report: Testing hybrid vigor in the lab in response to Eimeria

Alice

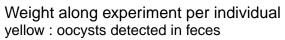
28 August 2018

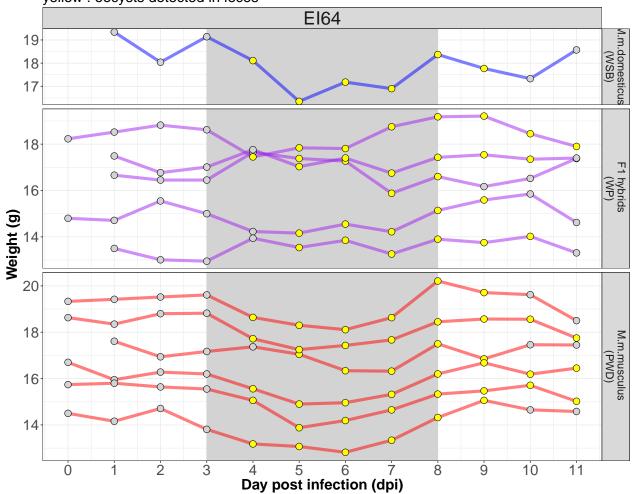
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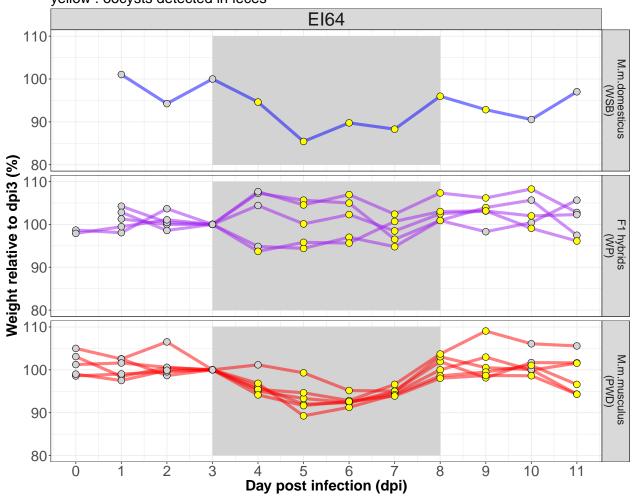
${\tt Expe_001},$ March 2017, Francisca's experiment. infection with E64 and Eflab

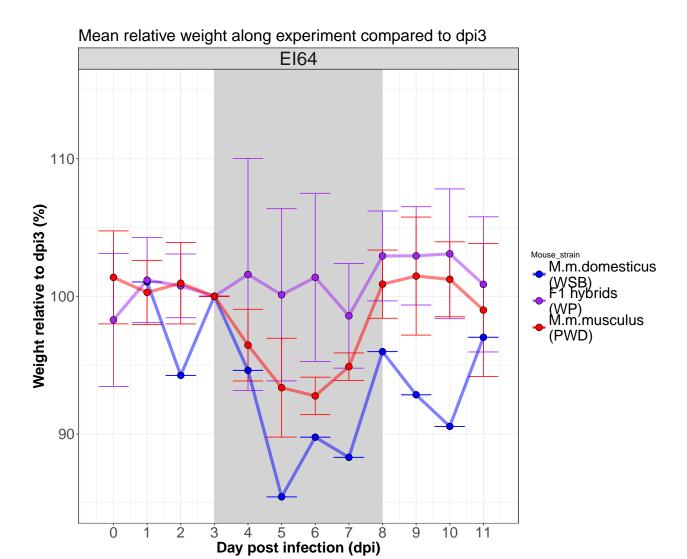
1. Weight loss





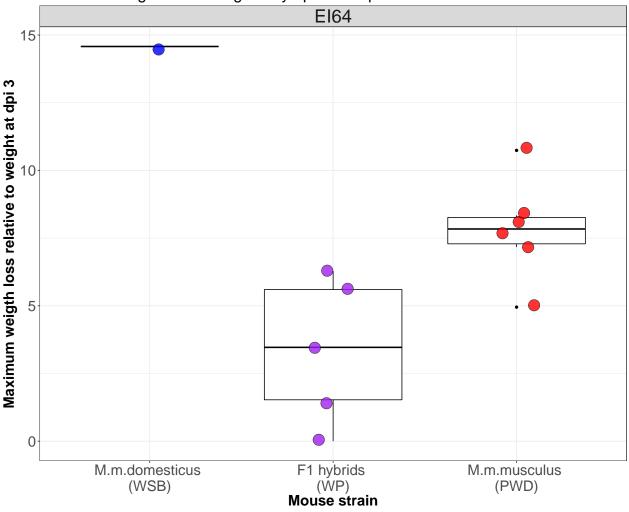
Relative weight along experiment compared to dpi3 yellow : oocysts detected in feces





For statistical analysis, we compare the maximum relative weight loss between the different groups. We limit our analysis to the period : dpi3 to dpi8 (symptomatic period for E64 strain).

Maximum weigth loss during the symptomatic phase

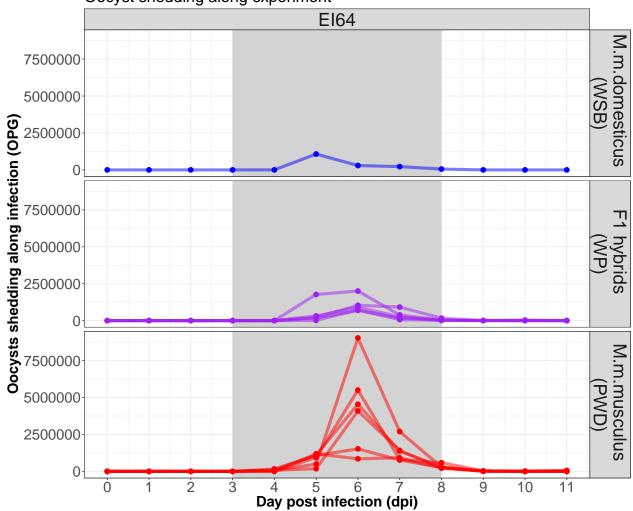


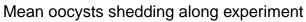
```
##
## lm(formula = relativeWeight ~ Mouse_strain, data = max.loss_001)
##
## Residuals:
##
      Min
               10 Median
                                3Q
                                      Max
  -2.9271 -0.9534 -0.0448 0.9445
                                   3.3754
##
## Coefficients:
##
                                      Estimate Std. Error t value Pr(>|t|)
                                                    1.008 95.836 7.44e-15
## (Intercept)
                                       96.625
## Mouse_strainM.m.domesticus \n(WSB)
                                      -11.201
                                                    2.470 -4.536 0.00141
                                                    1.365 -3.250 0.00999
                                        -4.437
## Mouse_strainM.m.musculus \n(PWD)
##
## (Intercept)
## Mouse_strainM.m.domesticus \n(WSB) **
## Mouse_strainM.m.musculus \n(PWD)
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 2.254 on 9 degrees of freedom
```

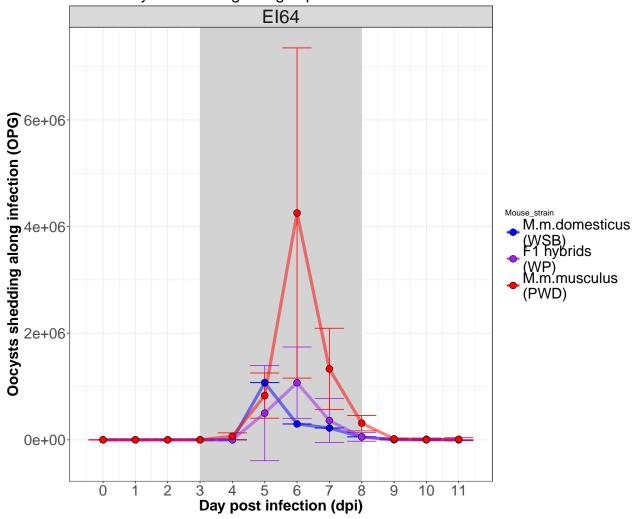
Multiple R-squared: 0.7311, Adjusted R-squared: 0.6714
F-statistic: 12.24 on 2 and 9 DF, p-value: 0.002711

2. Parasite shedding

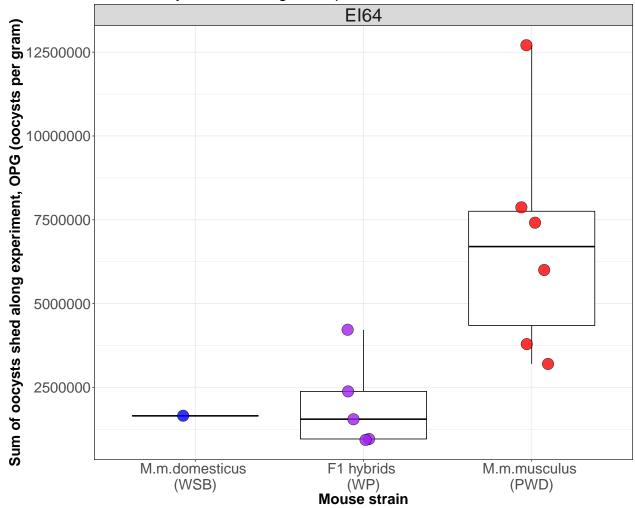
Oocyst shedding along experiment







Sum of oocysts shed along the experiment



```
##
## lm(formula = sum.oo ~ Mouse_strain, data = sum.oocysts_001)
##
## Residuals:
       Min
                  1Q
                      Median
                                    30
                                            Max
## -3628205 -1053694 -228126
                                693597
                                        5872235
##
## Coefficients:
##
                                      Estimate Std. Error t value Pr(>|t|)
                                                            1.653
                                                                    0.1328
## (Intercept)
                                       2007347
                                                  1214692
## Mouse_strainM.m.domesticus \n(WSB)
                                       -357449
                                                  2975376 -0.120
                                                                    0.9070
## Mouse_strainM.m.musculus \n(PWD)
                                       4820216
                                                  1644701
                                                            2.931
                                                                    0.0167 *
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 2716000 on 9 degrees of freedom
## Multiple R-squared: 0.5187, Adjusted R-squared: 0.4117
## F-statistic: 4.849 on 2 and 9 DF, p-value: 0.03724
```

3. Comparison host/parasite proxy

Relative weight loss compared to dpi3, during the symptomatic phase, vs parasite

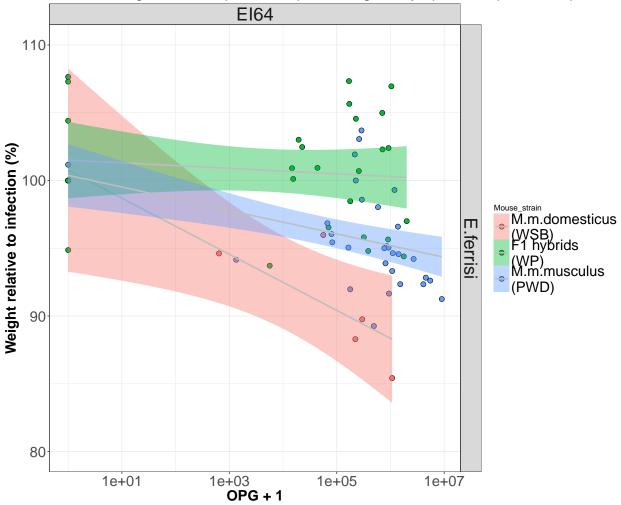
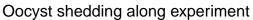


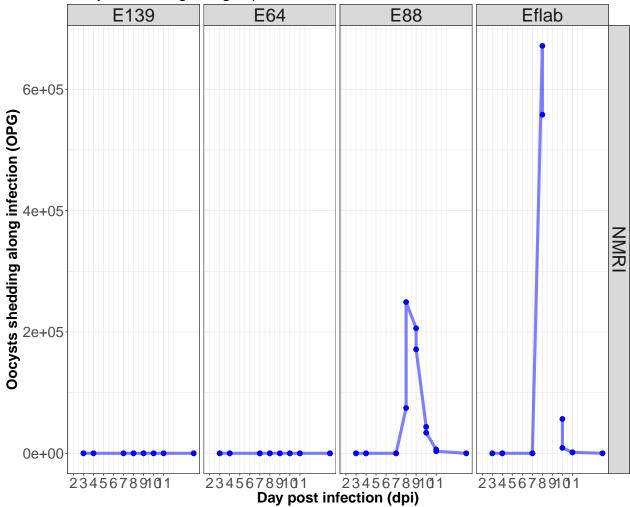
Figure 1: Weight as a function of OPG

Pass001: Nov 2017, passaging 4 isolates (some missing data)

(Eflab, E88, E139, E64) in NMRI. 2 mice per cage. Only OPG recorded

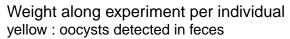
Parasite shedding

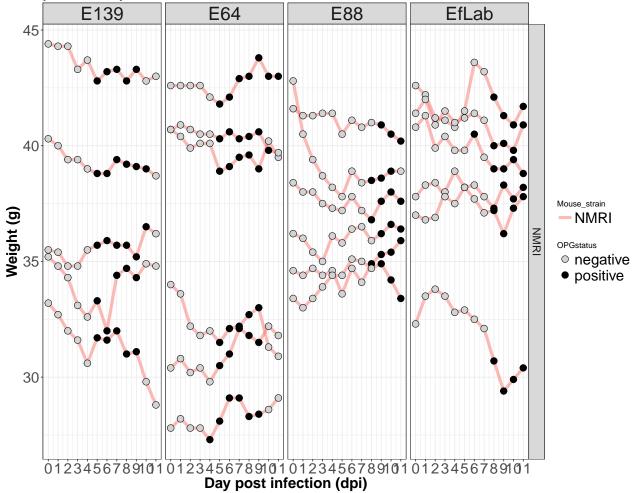




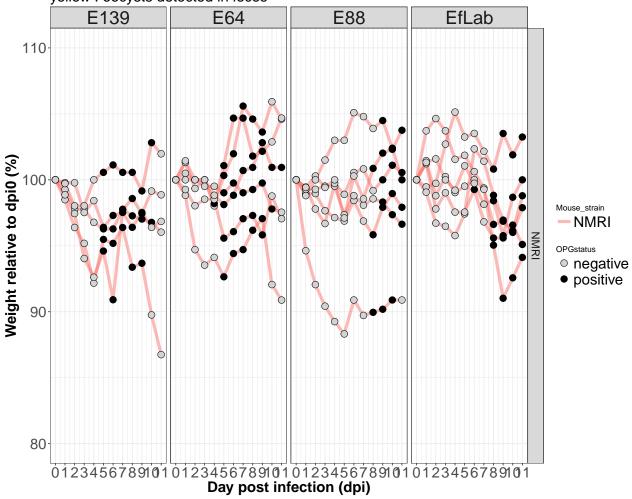
Expe_002: March 2018, NMRI mice infected with 4 *Eimeria* strains (Eflab, E88, E139, E64)

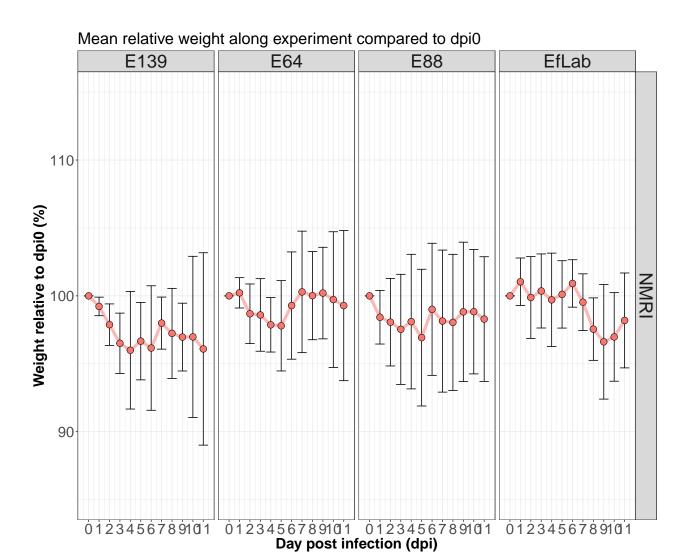
1. Weight loss



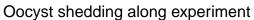


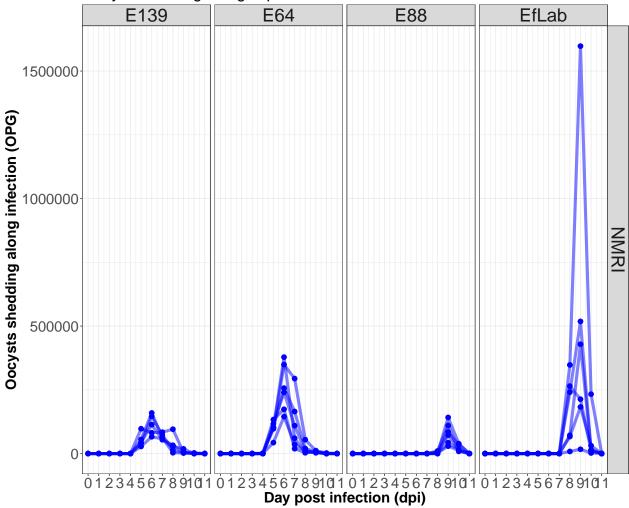
Relative weight along experiment compared to dpi0 yellow : oocysts detected in feces



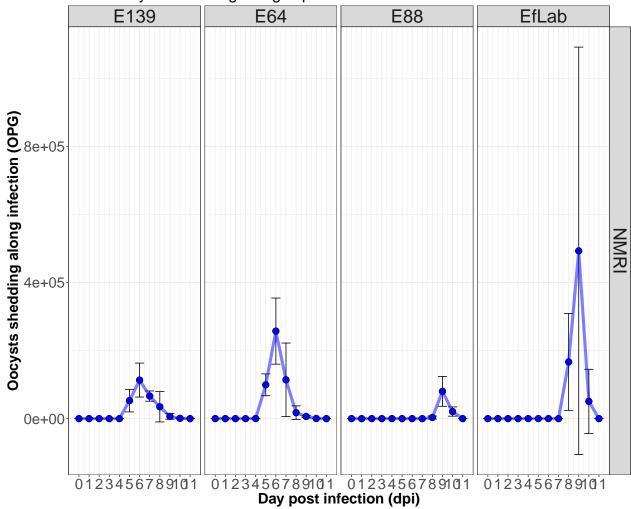


2. Parasite shedding









3. Comparison host/parasite proxy

Relative weight loss compared to dpi0, during the symptomatic phase, vs parasite

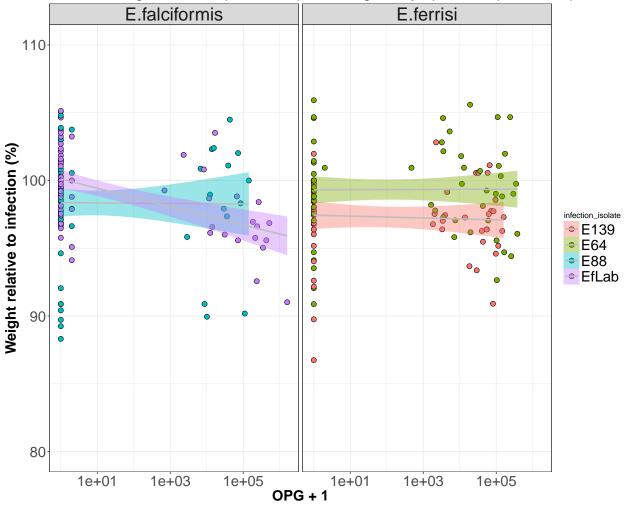
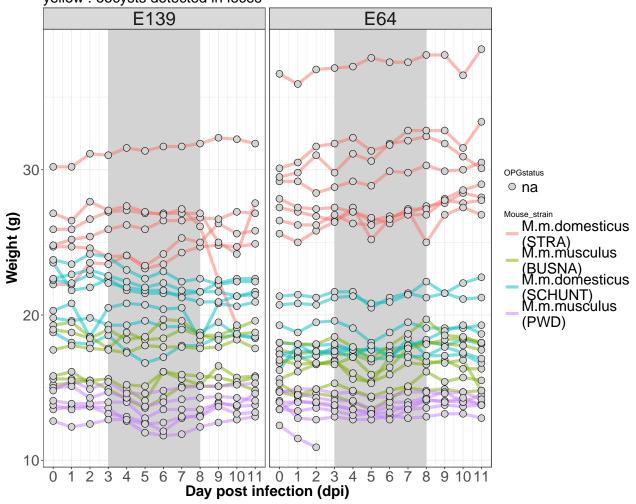


Figure 2: Weight as a function of OPG

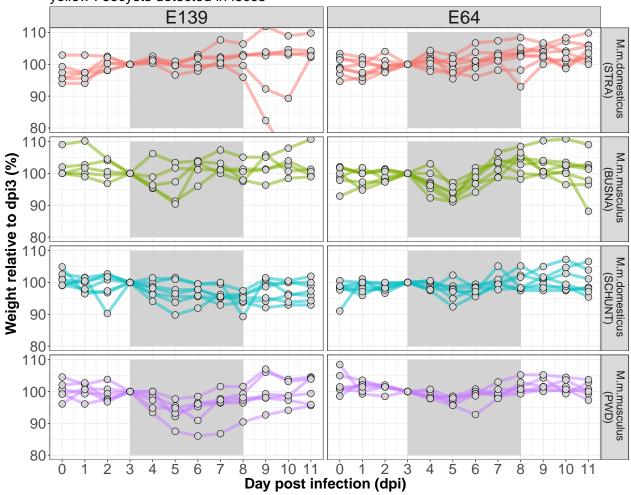
Expe_003 & Expe_004, April-May 2018, first batch Parental strains (F0) BUSNA, STRA, SCHUNT, PWD, infection with Eferrisi (E64 and E139) [2 batches]

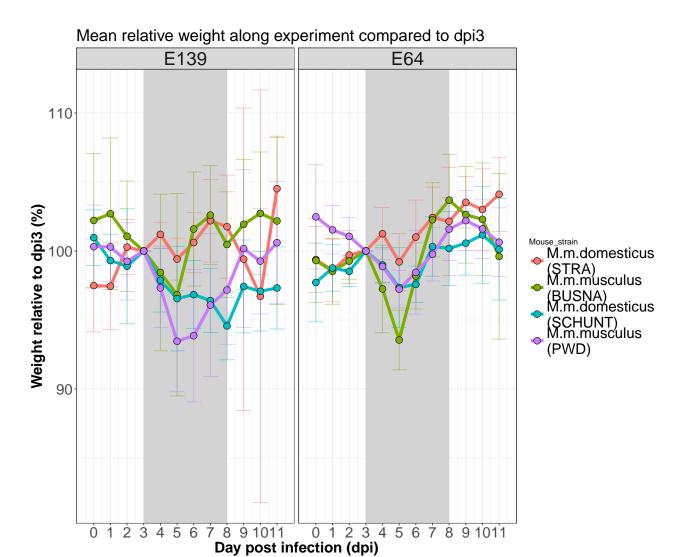
1. Weight loss

Weight along experiment per indiidual yellow: oocysts detected in feces



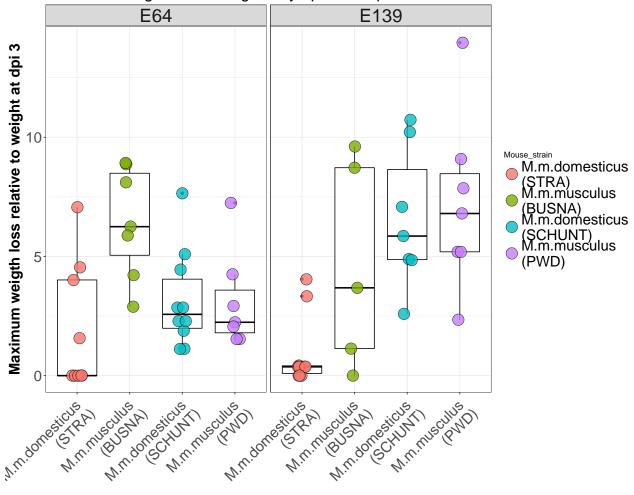
Relative weight along experiment compared to dpi3 yellow: oocysts detected in feces





For statistical analysis, we compare the maximum relative weight loss between the different groups. We limit our analysis to the period : dpi3 to dpi8 (symptomatic period for E64 strain).

Maximum weigth loss during the symptomatic phase



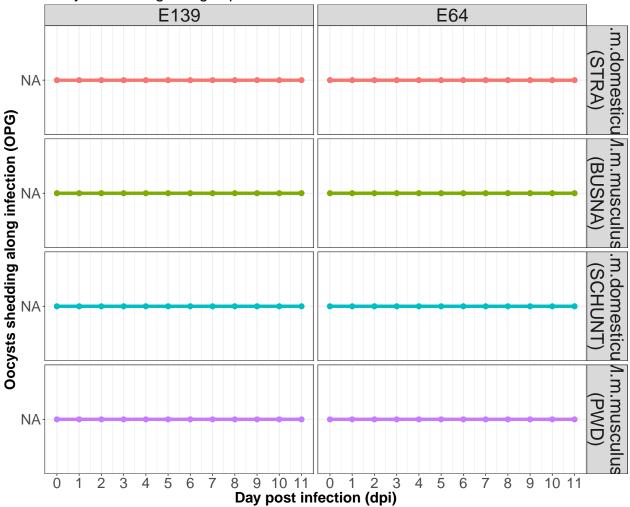
Mouse strain

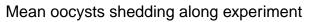
```
##
## Call:
##
  lm(formula = relativeWeight ~ Mouse_strain, data = max.loss_003_4)
##
## Residuals:
##
                1Q
                   Median
                                 3Q
                   0.9961
                           1.9204
##
   -8.8101 -2.3330
                                    5.6927
##
## Coefficients:
##
                                          Estimate Std. Error t value Pr(>|t|)
## (Intercept)
                                           98.6066
                                                       0.6760 145.870 < 2e-16
## Mouse_strainM.m.musculus \n(BUSNA)
                                           -4.2993
                                                       1.0865
                                                               -3.957 0.000209
## Mouse_strainM.m.domesticus \n(SCHUNT)
                                           -3.1843
                                                       0.9837
                                                               -3.237 0.001999
  Mouse_strainM.m.musculus \n(PWD)
                                           -3.7671
                                                       1.0378 -3.630 0.000601
##
##
## (Intercept)
## Mouse_strainM.m.musculus \n(BUSNA)
## Mouse_strainM.m.domesticus \n(SCHUNT)
## Mouse_strainM.m.musculus \n(PWD)
##
```

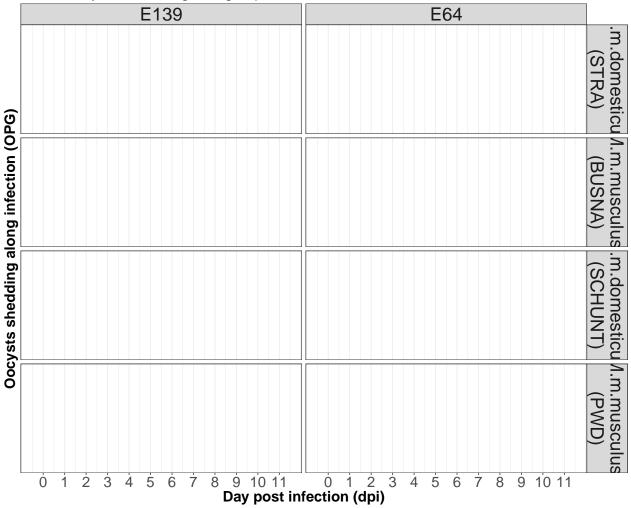
```
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 2.947 on 58 degrees of freedom
## Multiple R-squared: 0.2717, Adjusted R-squared: 0.234
## F-statistic: 7.212 on 3 and 58 DF, p-value: 0.0003403
```

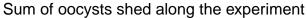
2. Parasite shedding

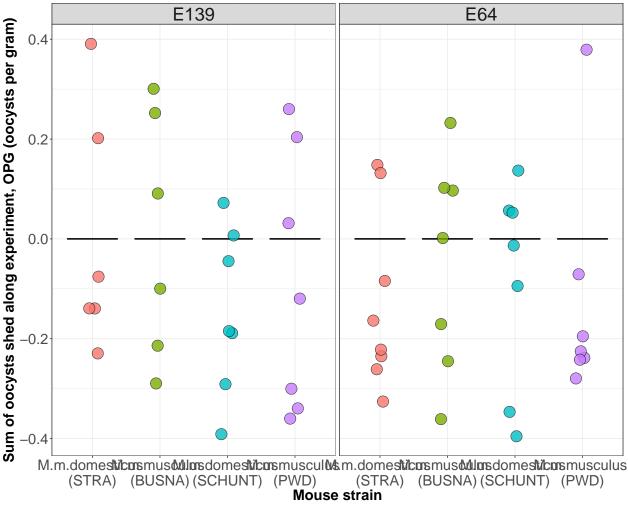
Oocyst shedding along experiment









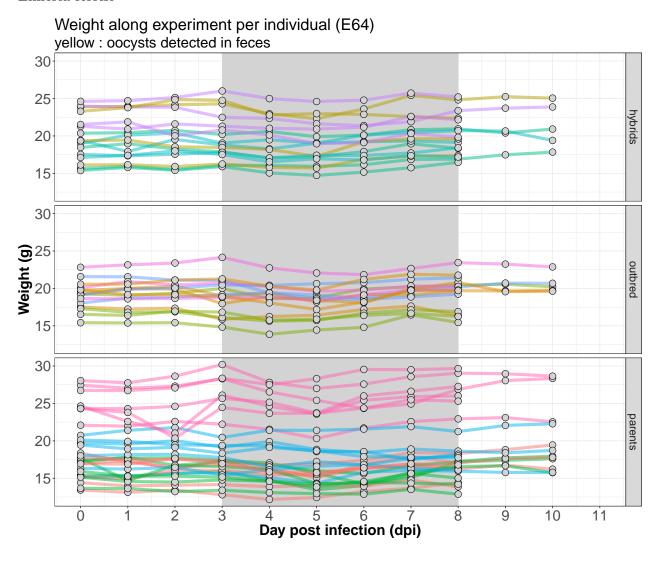


```
##
## lm(formula = sum.oo ~ Mouse_strain, data = sum.oocysts_003_4)
##
## Residuals:
      Min
##
              1Q Median
                             3Q
                                   Max
##
        0
               0
                      0
                             0
                                     0
##
  Coefficients:
##
##
                                          Estimate Std. Error t value Pr(>|t|)
## (Intercept)
                                                 0
                                                             0
                                                                    NA
                                                                             NA
## Mouse_strainM.m.musculus \n(BUSNA)
                                                 0
                                                             0
                                                                    NA
                                                                             NA
## Mouse_strainM.m.domesticus \n(SCHUNT)
                                                 0
                                                             0
                                                                    NA
                                                                             NA
## Mouse_strainM.m.musculus \n(PWD)
                                                                    NA
                                                                             NA
##
## Residual standard error: 0 on 51 degrees of freedom
## Multiple R-squared:
                         NaN, Adjusted R-squared:
                 NaN on 3 and 51 DF, p-value: NA
## F-statistic:
```

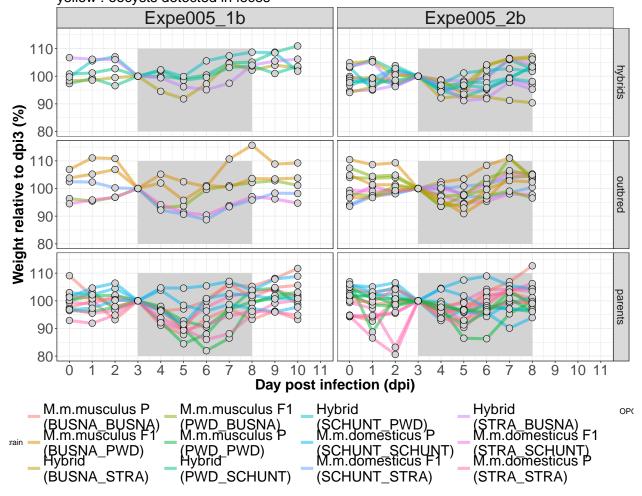
Expe_005, July 2018, FULL experiment (parents, intra specific and inter species hybrids) BUSNA, STRA, SCHUNT, PWD, infection with Eferrisi and Efalciformis (E64 and E88)

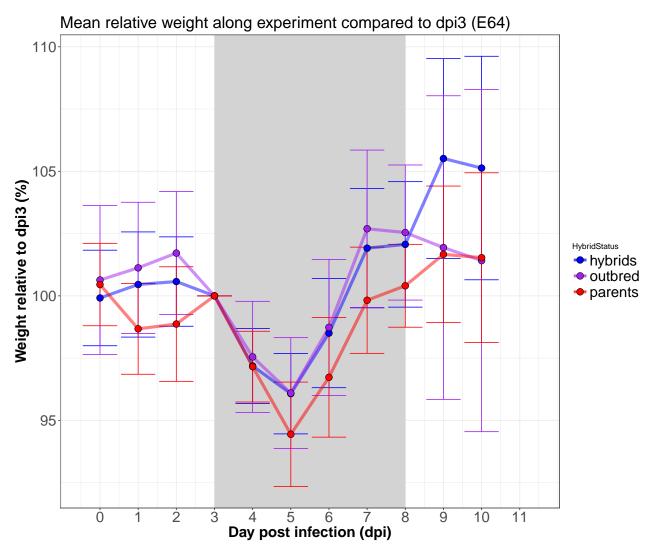
1. Weight loss

Eimeria ferrisi



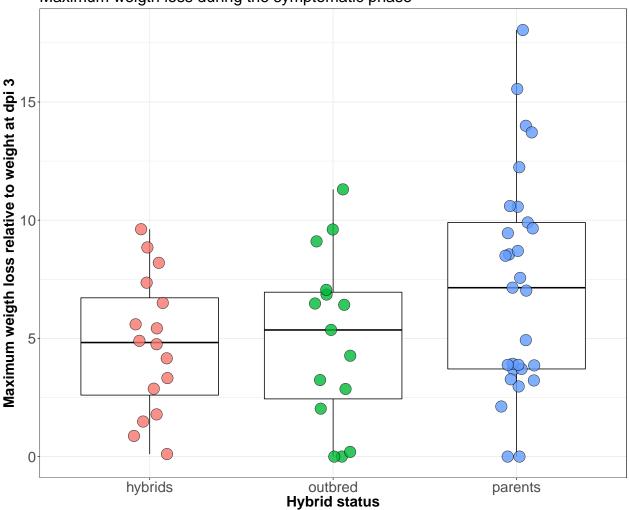
Relative weight along experiment compared to dpi3 (E64) yellow: oocysts detected in feces





For statistical analysis, we compare the maximum relative weight loss between the different groups. We limit our analysis to the period : dpi3 to dpi8 (symptomatic period for E64 strain).

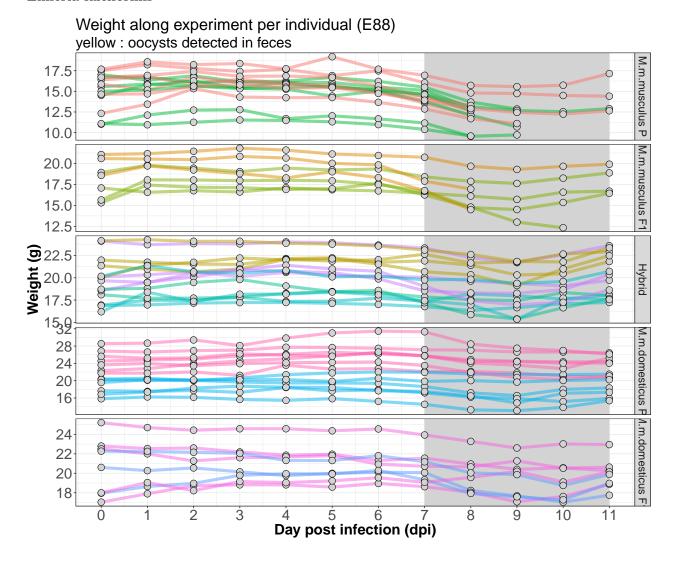


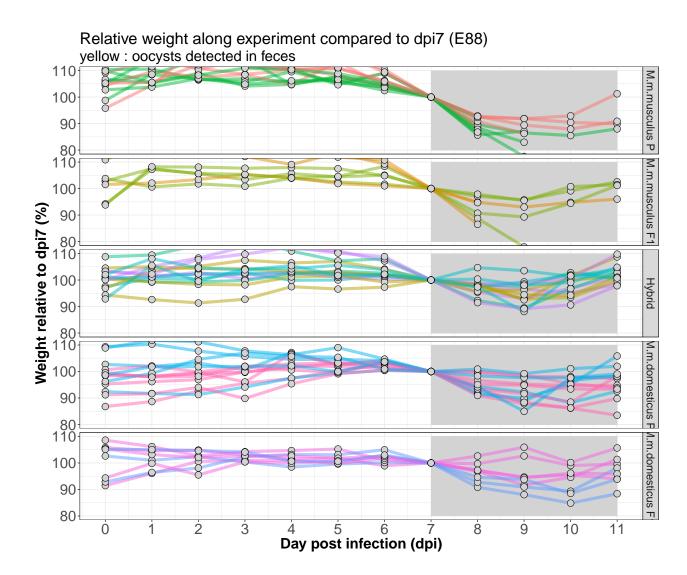


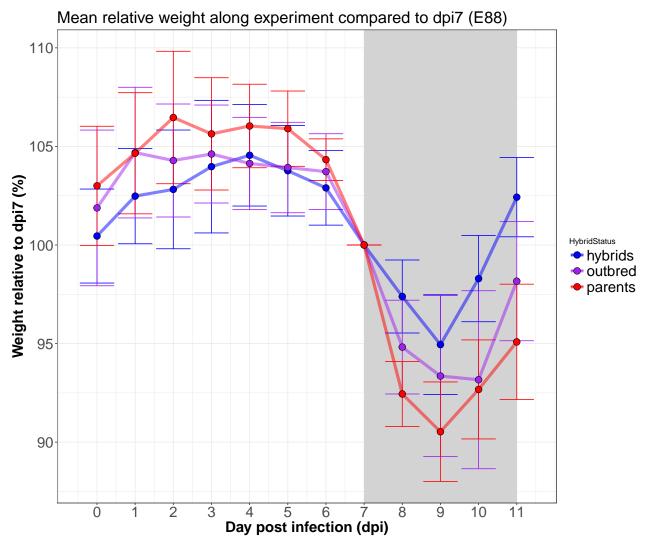
```
##
## lm(formula = relativeWeight ~ HybridStatus, data = max.loss_005_E64)
##
## Residuals:
##
      Min
               1Q Median
                               ЗQ
                                      Max
## -10.784 -2.451 -0.088
                            3.381
                                    7.262
##
## Coefficients:
##
                      Estimate Std. Error t value Pr(>|t|)
## (Intercept)
                       95.2644
                                   0.9968
                                          95.569
                                                    <2e-16 ***
## HybridStatusoutbred -0.2467
                                   1.4330 -0.172
                                                    0.8639
## HybridStatusparents -2.5264
                                   1.2417 -2.035
                                                    0.0466 *
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 3.987 on 57 degrees of freedom
## Multiple R-squared: 0.08785,
                                 Adjusted R-squared: 0.05585
## F-statistic: 2.745 on 2 and 57 DF, p-value: 0.07275
## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
```

```
## lmerModLmerTest]
## Formula: relativeWeight ~ HybridStatus + 1 | Mouse_strain
     Data: max.loss_005_E64
##
## REML criterion at convergence: 334.7
##
## Scaled residuals:
##
       Min 1Q
                    Median
                                  3Q
## -2.47724 -0.64423 -0.01634 0.87861 1.39148
##
## Random effects:
## Groups
                                   Variance Std.Dev. Corr
                Name
## Mouse_strain (Intercept)
                                   7.172e-13 8.469e-07
##
                HybridStatusoutbred 1.260e-12 1.123e-06 -1.00
##
                HybridStatusparents 5.504e+00 2.346e+00 -0.64 0.64
                                   1.465e+01 3.827e+00
## Residual
## Number of obs: 60, groups: Mouse_strain, 12
## Fixed effects:
              Estimate Std. Error df t value Pr(>|t|)
## (Intercept) 94.6748 0.6145 43.4600 154.1 <2e-16 ***
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## convergence code: 0
## unable to evaluate scaled gradient
## Model failed to converge: degenerate Hessian with 1 negative eigenvalues
```

Eimeria falciformis

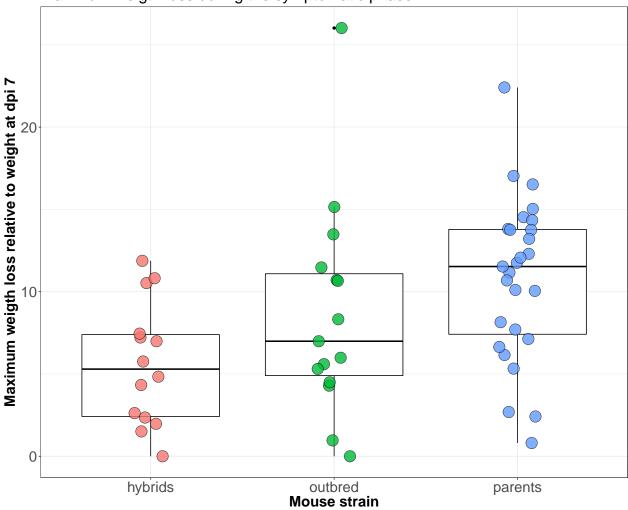






For statistical analysis, we compare the maximum relative weight loss between the different groups. We limit our analysis to the period : dpi7 to dpi11 (symptomatic period for E88 strain).





```
##
## lm(formula = relativeWeight ~ HybridStatus, data = max.loss_005_E88)
##
## Residuals:
##
       Min
                 1Q
                      Median
                                    3Q
  -17.3894 -2.9743 -0.0367
                               3.3960
                                        9.9730
##
## Coefficients:
##
                      Estimate Std. Error t value Pr(>|t|)
## (Intercept)
                        94.410
                                    1.369 68.976 < 2e-16 ***
## HybridStatusoutbred
                        -3.040
                                    1.903 -1.597 0.11617
## HybridStatusparents
                        -5.189
                                    1.687 -3.077 0.00331 **
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 5.121 on 53 degrees of freedom
## Multiple R-squared: 0.1524, Adjusted R-squared: 0.1204
## F-statistic: 4.763 on 2 and 53 DF, p-value: 0.01252
## [1] "Along all experiment:"
```