

Spacek et al., 2021, Figure 6-Supplement 1

**Figure 6-S1a<sub>1</sub>**

Comparing RMI during suppression against 0

```
# Fixed effect intercept only,
# random intercept for neurons,
# random intercept for experiments, nested within series
lmer.a1.1 = lmer(suppressionrmi ~ 1 + (1 | uid) + (1 | sid/eid),
                  data = tbw %>% drop_na(suppressionrmi))

display(lmer.a1.1)

## lmer(formula = suppressionrmi ~ 1 + (1 | uid) + (1 | sid/eid),
##       data = tbw %>% drop_na(suppressionrmi))
## coef.est  coef.se
##      0.20    0.09
##
## Error terms:
##   Groups     Name        Std.Dev.
##   uid        (Intercept) 0.07
##   eid:sid   (Intercept) 0.02
##   sid        (Intercept) 0.24
##   Residual           0.09
##   ---
##   number of obs: 57, groups: uid, 37; eid:sid, 9; sid, 7
##   AIC = -58.8, DIC = -74.8
##   deviance = -71.8
```

Mean firing rate RMI

Suppression: RMI =  $0.20 \pm 0.19$   
n = 37 neurons from 4 mice

## Figure 6-S1a<sub>2</sub>

### Comparing RMI during suppression against 0

```
# LMM gives singular fits, even with a single random intercept for neurons.  
# We do ordinary regression instead:  
lm.a2.1 = lm(suppressionrmi ~ 1, data = tbw %>% drop_na(suppressionrmi))  
  
display(lm.a2.1)  
  
## lm(formula = suppressionrmi ~ 1, data = tbw %>% drop_na(suppressionrmi))  
##             coef.est  coef.se  
## (Intercept) -0.12      0.04  
## ---  
## n = 51, k = 1  
## residual sd = 0.30, R-Squared = 0.00
```

### Mean burst ratio RMI

Suppression: RMI =  $-0.12 \pm 0.08$   
n = 34 neurons from 3 mice

## Figure 6-S1b<sub>1</sub>

### Slope of regression line

```
# Random intercept for neurons,  
# random intercept for experiments  
lmer.b1.1 = lmer(runfmi ~ sitfmi + (1 | uid) + (1 | eid),  
                  data = tbw %>% drop_na(runfmi, sitfmi))  
  
display(lmer.b1.1)  
  
## lmer(formula = runfmi ~ sitfmi + (1 | uid) + (1 | eid), data = tbw %>%  
##       drop_na(runfmi, sitfmi))  
##             coef.est  coef.se  
## (Intercept) 0.04      0.03  
## sitfmi      0.52      0.09  
##  
## Error terms:  
##   Groups    Name        Std.Dev.  
##   uid       (Intercept) 0.09  
##   eid       (Intercept) 0.06  
##   Residual            0.12  
## ---  
## number of obs: 57, groups: uid, 37; eid, 9  
## AIC = -35.7, DIC = -62.2  
## deviance = -53.9  
anova(lmer.b1.1)  
  
## Type III Analysis of Variance Table with Satterthwaite's method  
##           Sum Sq Mean Sq NumDF DenDF F value    Pr(>F)  
## sitfmi 0.45992 0.45992     1  47.435 33.468 5.525e-07 ***  
## ---  
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Slope of  $0.52 \pm 0.18$

n = 37 neurons from 4 mice

## Figure 6-S1b<sub>2</sub>

### Slope of regression line

```
# Random intercept for neurons,
# random intercept for experiments, nested in series
lmer.b2.1 = lmer(runfmi ~ sitfmi + (1 | uid) + (1 | sid/eid),
                 data = tbw %>% drop_na(runfmi, sitfmi))

display(lmer.b2.1)

## lmer(formula = runfmi ~ sitfmi + (1 | uid) + (1 | sid/eid), data = tbw %>%
##       drop_na(runfmi, sitfmi))
##       coef.est coef.se
## (Intercept) -0.28      0.16
## sitfmi       0.52      0.16
##
## Error terms:
##   Groups     Name     Std.Dev.
##   uid        (Intercept) 0.21
##   eid:sid   (Intercept) 0.11
##   sid        (Intercept) 0.27
##   Residual           0.19
## ---
## number of obs: 51, groups: uid, 33; eid:sid, 7; sid, 5
## AIC = 36.2, DIC = 16.1
## deviance = 20.2
anova(lmer.b2.1)

## Type III Analysis of Variance Table with Satterthwaite's method
##          Sum Sq Mean Sq NumDF DenDF F value    Pr(>F)
## sitfmi 0.36662 0.36662     1  30.117 10.076 0.003451 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Slope of 0.52 ± 0.33
n = 33 neurons from 3 mice
```

## Figure 6-S1c<sub>1</sub>

### Slope of regression line

```
# Random intercept for neurons,  
# random intercept for experiments  
lmer.c1.1 = lmer(fmi ~ rmi + (1 | uid) + (1 | eid),  
                  data = tbw %>% drop_na(fmi, rmi))  
  
display(lmer.c1.1)  
  
## lmer(formula = fmi ~ rmi + (1 | uid) + (1 | eid), data = tbw %>%  
##       drop_na(fmi, rmi))  
##             coef.est  coef.se  
## (Intercept) 0.01      0.05  
## rmi         0.18      0.14  
##  
## Error terms:  
##   Groups     Name        Std.Dev.  
##   uid        (Intercept) 0.19  
##   eid        (Intercept) 0.09  
##   Residual            0.09  
## ---  
## number of obs: 57, groups: uid, 37; eid, 9  
## AIC = -16.2, DIC = -39.3  
## deviance = -32.8  
anova(lmer.c1.1)  
  
## Type III Analysis of Variance Table with Satterthwaite's method  
##           Sum Sq  Mean Sq NumDF DenDF F value Pr(>F)  
## rmi 0.013502 0.013502     1  38.356  1.6842 0.2021
```

Slope of  $0.18 \pm 0.27$

$n = 37$  neurons from 4 mice

## Figure 6-S1c<sub>2</sub>

### Slope of regression line

```
# Random intercept for neurons,
# random intercept for experiments, nested in series
lmer.c2.1 = lmer(fmi ~ rmi + (1 | uid) + (1 | sid/eid),
                 data = tbw %>% drop_na(fmi, rmi))

display(lmer.c2.1)

## lmer(formula = fmi ~ rmi + (1 | uid) + (1 | sid/eid), data = tbw %>%
##       drop_na(fmi, rmi))
##             coef.est  coef.se
## (Intercept) -0.38      0.09
## rmi          0.25      0.05
##
## Error terms:
##   Groups     Name     Std.Dev.
##   uid        (Intercept) 0.14
##   eid:sid   (Intercept) 0.14
##   sid        (Intercept) 0.15
##   Residual           0.10
## ---
## number of obs: 51, groups: uid, 34; eid:sid, 7; sid, 5
## AIC = -10, DIC = -36.3
## deviance = -29.2
anova(lmer.c2.1)

## Type III Analysis of Variance Table with Satterthwaite's method
##   Sum Sq Mean Sq NumDF DenDF F value    Pr(>F)
## rmi 0.27033 0.27033     1 30.557  24.893 2.294e-05 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Slope of  $0.25 \pm 0.10$

n = 34 neurons from 3 mice