

# User Input of Scripts for Article Figures: Version 75

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May 24, 2023

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## 1 Figure 2

### 1.1 Figure 2a:

From “Data Analysis” directory run the function,  
*masterPsychometricFunctionPlot('approachavoid')*.

For user inputs please enter the following inputs:

- i) Enter genotype: CRL: Long Evans
- ii) Do you want to analyze only approach trials? (y/n) n
- iii) Enter tasktypedone (or enter "all" for all task types): P2L1
- iv) Which health types do you want to analyze?  
(enter multiple values separated by comma and a space or type 'all' for all types): N/A
- v) Start date? 06/16/2022
- vi) End date? 06/23/2022
- vii) Do you want to split the graph by gender? (y/n) y

### 1.2 Figure 2d:

From “Data Analysis” directory run the function,  
*masterPsychometricFunctionPlot('entrytime')*.

For user inputs please enter the following inputs:

- i) Enter genotype: CRL: Long Evans
- ii) Do you want to analyze only approach trials? (y/n) y

For rest of the user inputs please enter the same inputs as in **Figure 2a**

### 1.3 Figure 2e:

From “Data Analysis” directory run the function,  
*masterPsychometricFunctionPlot('distanceaftertoneuntillimitingtimestamp')*.

For user inputs please enter the same inputs as **Figure 2a**

### 1.4 Figure 2f:

From “Data Analysis” directory run the function,  
*masterPsychometricFunctionPlot('stoppingpts\_per\_unittravel\_method6')*.

For user inputs please enter the same inputs as **Figure 2a**

### 1.5 Figure 2g:

From “Data Analysis” directory run the function,  
*masterPsychometricFunctionPlot('bigaccelerationperunittravel')*.

For user inputs please enter the same inputs as **Figure 2a**

## 1.6 Figure 2h:

From “Data Analysis” directory run the function,  
*masterPsychometricFunctionPlot('passingcentralzonerejectinitialpresence')*.  
For user inputs please enter the same inputs as **Figure 2a**

## 2 Figure 5

### 2.1 Figure 5a:

Step 1: Get ‘Sans food dep’ figure.

For ‘Sans food dep’ From “Data Analysis” directory run the function,  
*masterPsychometricFunctionPlot('approachavoid')*.

For user inputs please enter the following inputs:

- i) Enter genotype: CRL: Long Evans
- ii) Do you want to analyze only approach trials? (y/n) n
- iii) Enter tasktypedone (or enter "all" for all task types): P2L1
- iv) Which health types do you want to analyze?  
(enter multiple values separated by comma and a space or type 'all' for all types): N/A
- v) Start date? 06/16/2022
- vi) End date? 06/23/2022
- vii) Do you want to split the graph by gender? (y/n) n
- viii) Do you want to a graph for specific animal? (y/n) n

Step2: Get ‘Food dep’ figure.

For ‘Food dep’ From “Data Analysis” directory run the function,  
*masterPsychometricFunctionPlot('approachavoid')*.

For user inputs please enter the following inputs:

- i) Enter genotype: CRL: Long Evans
- ii) Do you want to analyze only approach trials? (y/n) n
- iii) Enter tasktypedone (or enter "all" for all task types): P2L1
- iv) Which health types do you want to analyze?  
(enter multiple values separated by comma and a space or type 'all' for all types): Food Deprivation
- v) Start date? 08/23/2022
- vi) End date? 08/25/2022
- vii) Do you want to split the graph by gender? (y/n) n
- viii) Do you want to a graph for specific animal? (y/n) n

Step3: Overlay ‘Sans food dep’ and ‘Food dep’ figures.

- i) Open *mergePlots.m* from 'Plots' directory
- ii) Paste the figures obtained in step 1 and 2 for 'f1' and 'f2'
- iii) Comment out rest of the figure names since we don't want to overlay more
- iv) Run the script

## 2.2 Figure 5c:

Step 1: Get 'Sans food dep' figure.

For 'Sans food dep' From "Data Analysis" directory run the function, *masterPsychometricFunctionPlot('entrytime')*. For user inputs please enter the following inputs:

- i) Enter genotype: CRL: Long Evans
- ii) Do you want to analyze only approach trials? (y/n) y

For rest of the user inputs please enter the same inputs as Step1 in **Figure 5a**

Step2: Get 'Food dep' figure.

For 'Food dep' From "Data Analysis" directory run the function, *masterPsychometricFunctionPlot('entrytime')*.

For user inputs please enter the following inputs:

- i) Enter genotype: CRL: Long Evans
- ii) Do you want to analyze only approach trials? (y/n) y

For rest of the user inputs please enter the same inputs as Step2 in **Figure 5a**

Step3: Overlay 'Sans food dep' and 'Food dep' figures.

Please follow the same steps as Step3 in **Figure 5a**

## 2.3 Figure 5d:

Step 1: Get 'Sans food dep' figure.

For 'Sans food dep' From "Data Analysis" directory run the function, *masterPsychometricFunctionPlot('distanceaftertoneuntillimitingtimestamp')*.

For user inputs please enter the same inputs as Step1 in **Figure 5a**

Step2: Get 'Food dep' figure.

For 'Food dep' From "Data Analysis" directory run the function, *masterPsychometricFunctionPlot('distanceaftertoneuntillimitingtimestamp')*.

For user inputs please enter the same inputs as Step2 in **Figure 5a**

Step3: Overlay 'Sans food dep' and 'Food dep' figures.

Please follow the same steps as Step3 in **Figure 5a**

## 2.4 Figure 5e:

Step 1: Get 'Sans food dep' figure.

For 'Sans food dep' From "Data Analysis" directory run the function,  
*masterPsychometricFunctionPlot('stoppingpts\_per\_unittravel\_method6')*.

For user inputs please enter the same inputs as Step1 in **Figure 5a**

Step2: Get 'Food dep' figure.

For 'Food dep' From "Data Analysis" directory run the function,  
*masterPsychometricFunctionPlot('stoppingpts\_per\_unittravel\_method6')*.

For user inputs please enter the same inputs as Step2 in **Figure 5a**

Step3: Overlay 'Sans food dep' and 'Food dep' figures.

Please follow the same steps as Step3 in **Figure 5a**

## 2.5 Figure 5f:

Step 1: Get 'Sans food dep' figure.

For 'Sans food dep' From "Data Analysis" directory run the function,  
*masterPsychometricFunctionPlot('bigaccelerationperunittravel')*.

For user inputs please enter the same inputs as Step1 in **Figure 5a**

Step2: Get 'Food dep' figure.

For 'Food dep' From "Data Analysis" directory run the function,  
*masterPsychometricFunctionPlot('bigaccelerationperunittravel')*.

For user inputs please enter the same inputs as Step2 in **Figure 5a**

Step3: Overlay 'Sans food dep' and 'Food dep' figures.

Please follow the same steps as Step3 in **Figure 5a**

## 2.6 Figure 5g:

Step 1: Get 'Sans food dep' figure.

For 'Sans food dep' From "Data Analysis" directory run the function,  
*masterPsychometricFunctionPlot('passingcentralzonerejectinitialpresence')*.

For user inputs please enter the same inputs as Step1 in **Figure 5a**

Step2: Get 'Food dep' figure.

For 'Food dep' From "Data Analysis" directory run the function,  
*masterPsychometricFunctionPlot('passingcentralzonerejectinitialpresence')*.

For user inputs please enter the same inputs as Step2 in **Figure 5a**

Step3: Overlay 'Sans food dep' and 'Food dep' figures.

Please follow the same steps as Step3 in **Figure 5a**

## 3 Figure 6

### 3.1 Figure 6b (Left):

From “Data Analysis” directory run the function,  
*alcoholPsychometricFunctionPlot('approachavoid')*.

For user inputs please enter the following inputs:

- i) Enter genotype: CRL: Long Evans
- ii) Do you want to analyze only approach trials? (y/n) n
- iii) Enter tasktypedone (or enter "all" for all task types): P2A
- iv) Which health types do you want to analyze?  
(enter multiple values separated by comma and a space or type 'all' for all types): N/A
- v) Start date? 09/16/2022
- vi) End date? 10/03/2022
- vii) Do you want to split the graph by gender? (y/n) y

### 3.2 Figure 6b (Right):

From “Data Analysis” directory run the function,  
*alcoholPsychometricFunctionPlot('approachavoid')*.

For user inputs please enter the following inputs:

- i) Enter genotype: lg\_boost, lg\_etoh
- ii) Do you want to analyze only approach trials? (y/n) n
- iii) Enter tasktypedone (or enter "all" for all task types): P2A
- iv) Which health types do you want to analyze?  
(enter multiple values separated by comma and a space or type 'all' for all types): N/A
- v) Start date? 11/02/2022
- vi) End date? 12/01/2022
- vii) Do you want to split the graph by gender? (y/n) y

### 3.3 Figure 6d (Left):

From “Data Analysis” directory run the function,  
*alcoholPsychometricFunctionPlot('entrytime')*

For user inputs please enter the following inputs:

- i) Enter genotype: CRL: Long Evans
  - ii) Do you want to analyze only approach trials? (y/n) y
- For rest of the user inputs please enter the same inputs as in **Figure 6b (Left)**

### 3.4 Figure 6d (Right):

From “Data Analysis” directory run the function,  
*alcoholPsychometricFunctionPlot('entrytime')*

For user inputs please enter the following inputs:

i) Enter genotype: lg\_boost, lg\_etoh

ii) Do you want to analyze only approach trials? (y/n) y

For rest of the user inputs please enter the same inputs as in **Figure 6b (Right)**

### 3.5 Figure 6e:

From “Data Analysis” directory run the function,  
*oxyPsychometricFunctionPlot('approachavoid')*.

For user inputs please enter the following inputs:

i) Which data do you want to analyze? Print "Oxycodon" or "Incubation"

Oxycodon

ii) Do you want to analyze only approach trials? (y/n) n

iii) Do you want to split the graph by gender? (y/n) y

### 3.6 Figure 6f:

From “Data Analysis” directory run the function,

*oxyPsychometricFunctionPlot('stoppingpts\_per\_unittravel\_method6')*.

For user inputs please enter the same inputs as **Figure 6e**

### 3.7 Figure 6g:

From “Data Analysis” directory run the function,

*oxyPsychometricFunctionPlot('approachavoid')*.

For user inputs please enter the following inputs:

i) Which data do you want to analyze? Print "Oxycodon" or "Incubation"

Incubation

ii) Do you want to analyze only approach trials? (y/n) n

iii) Do you want to split the graph by gender? (y/n) y

### 3.8 Figure 6h:

From “Data Analysis” directory run the function,

*oxyPsychometricFunctionPlot('stoppingpts\_per\_unittravel\_method6')*.

For user inputs please enter the same inputs as **Figure 6g**



### 3.9 Figure 6k:

From “Data Analysis” directory run the function, *barPlotOfOxy.m*.

## 4 Supplemental Figure 6

### 4.1 Supplemental Figure 6a:

From “Data Analysis” directory run the function,  
*masterPsychometricFunctionPlot('approachavoid')*.

For user inputs please enter the following inputs:

- i) Enter genotype: CRL: Long Evans
- ii) Do you want to analyze only approach trials? (y/n) n
- iii) Enter tasktypedone (or enter "all" for all task types): P2L1
- iv) Which health types do you want to analyze?  
(enter multiple values separated by comma and a space or type 'all' for all types): Food Deprivation
- v) Start date? 08/23/2022
- vi) End date? 08/25/2022
- vii) Do you want to split the graph by gender? (y/n) y

### 4.2 Supplemental Figure 6b:

From “Data Analysis” directory run the function,  
*masterPsychometricFunctionPlot('distanceaftertoneuntillimitingtimestamp')*.  
For user inputs please enter the same inputs as **Supplemental Figure 6a**.

### 4.3 Supplemental Figure 6c:

From “Data Analysis” directory run the function,  
*masterPsychometricFunctionPlot('stoppingpts\_per\_unittravel\_method6')*.  
For user inputs please enter the same inputs as **Supplemental Figure 6a**.

### 4.4 Supplemental Figure 6d:

From “Data Analysis” directory run the function,  
*masterPsychometricFunctionPlot('bigaccelerationperunittravel')*.  
For user inputs please enter the same inputs as **Supplemental Figure 6a**.

## 4.5 Supplemental Figure 6e:

From “Data Analysis” directory run the function,  
*masterPsychometricFunctionPlot('passingcentralzonerejectinitialpresence')*.  
For user inputs please enter the same inputs as **Supplemental Figure 6a**.

## 5 Supplemental Figure 7

### 5.1 Supplemental Figure 7a (Left):

From “Data Analysis” directory run the function,  
*alcoholPsychometricFunctionPlot('bigaccelerationperunittravel')*.  
For user inputs please enter the same inputs as **Figure 6b (Left)**.

### 5.2 Supplemental Figure 7a (Right):

From “Data Analysis” directory run the function,  
*alcoholPsychometricFunctionPlot('bigaccelerationperunittravel')*.  
For user inputs please enter the same inputs as **Figure 6b (Right)**.

### 5.3 Supplemental Figure 7b (Left):

From “Data Analysis” directory run the function,  
*alcoholPsychometricFunctionPlot('distanceaftertoneuntillimitingtimestamp')*.  
For user inputs please enter the same inputs as **Figure 6b (Left)**.

### 5.4 Supplemental Figure 7b (Right):

From “Data Analysis” directory run the function,  
*alcoholPsychometricFunctionPlot('distanceaftertoneuntillimitingtimestamp')*.  
For user inputs please enter the same inputs as **Figure 6b (Right)**.

### 5.5 Supplemental Figure 7d:

From “Data Analysis” directory run the function,  
*oxyPsychometricFunctionPlot('entrytime')*.  
For user inputs please enter the following inputs:  
i) Which data do you want to analyze? Print "Oxycodon" or "Incubation"  
Oxycodon  
ii) Do you want to analyze only approach trials? (y/n) y  
iii) Do you want to split the graph by gender? (y/n) y

## 5.6 Supplemental Figure 7e:

From “Data Analysis” directory run the function,  
*oxyPsychometricFunctionPlot('passingcentralzonerejectinitialpresence')*.  
For user inputs please enter the same inputs as **Figure 6e**.

## 5.7 Supplemental Figure 7f:

From “Data Analysis” directory run the function,  
*oxyPsychometricFunctionPlot('bigaccelerationperunittravel')*.  
For user inputs please enter the same inputs as **Figure 6e**.

## 5.8 Supplemental Figure 7g:

From “Data Analysis” directory run the function,  
*oxyPsychometricFunctionPlot('distanceaftertoneuntillimitingtimestamp')*.  
For user inputs please enter the same inputs as **Figure 6e**.

## 5.9 Supplemental Figure 7h:

From “Data Analysis” directory run the function,  
*oxyPsychometricFunctionPlot('entrytime')*.  
For user inputs please enter the following inputs:  
i) Which data do you want to analyze? Print "Oxycodon" or "Incubation"  
Incubation  
ii) Do you want to analyze only approach trials? (y/n) y  
iii) Do you want to split the graph by gender? (y/n) y

## 5.10 Supplemental Figure 7i:

From “Data Analysis” directory run the function,  
*oxyPsychometricFunctionPlot('bigaccelerationperunittravel')*.  
For user inputs please enter the same inputs as **Figure 6g**.

## 5.11 Supplemental Figure 7j:

From “Data Analysis” directory run the function,  
*oxyPsychometricFunctionPlot('distanceaftertoneuntillimitingtimestamp')*.  
For user inputs please enter the same inputs as **Figure 6g**.

### 5.12 Supplemental Figure 7k:

From “Data Analysis” directory run the function,  
*oxyPsychometricFunctionPlot('passingcentralzonerejectinitialpresence')*.  
For user inputs please enter the same inputs as **Figure 6g**.