

This textbook collects all the figures generated for the manuscript.

First take care of the path

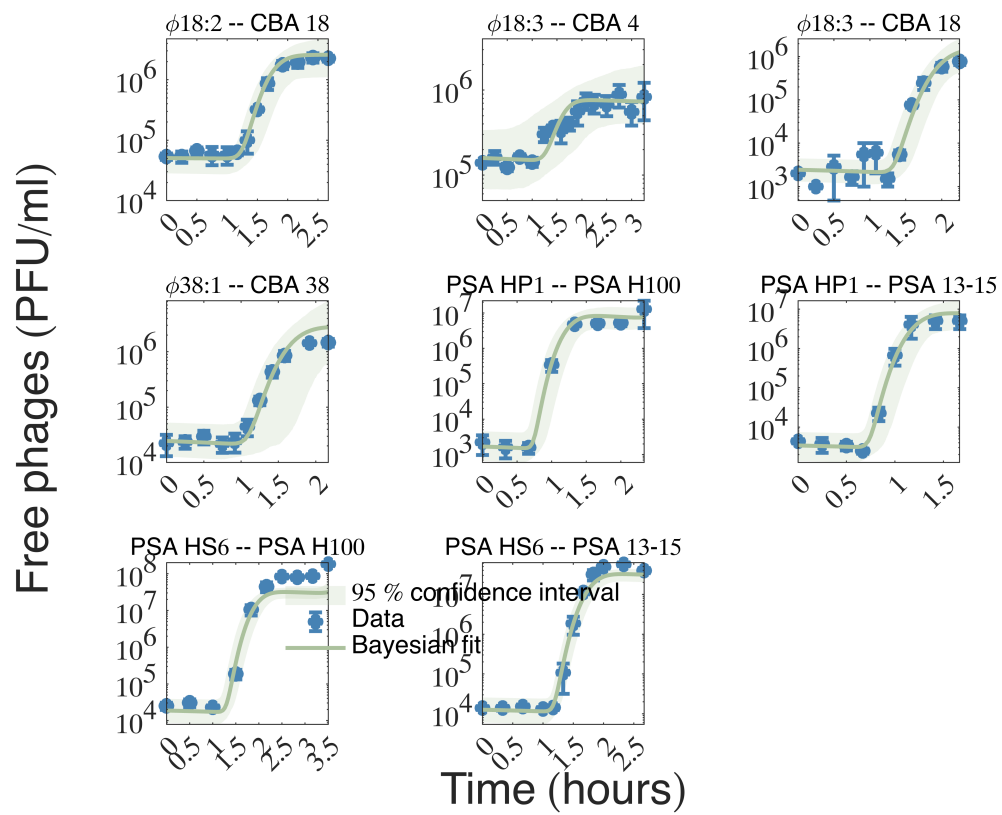
```
addpath(genpath('./..'))
```

Here is Figure1

```
cd('./figure1/')  
fig1_parameters;  
cd ..
```

Here is Fig 1 all the onesteps

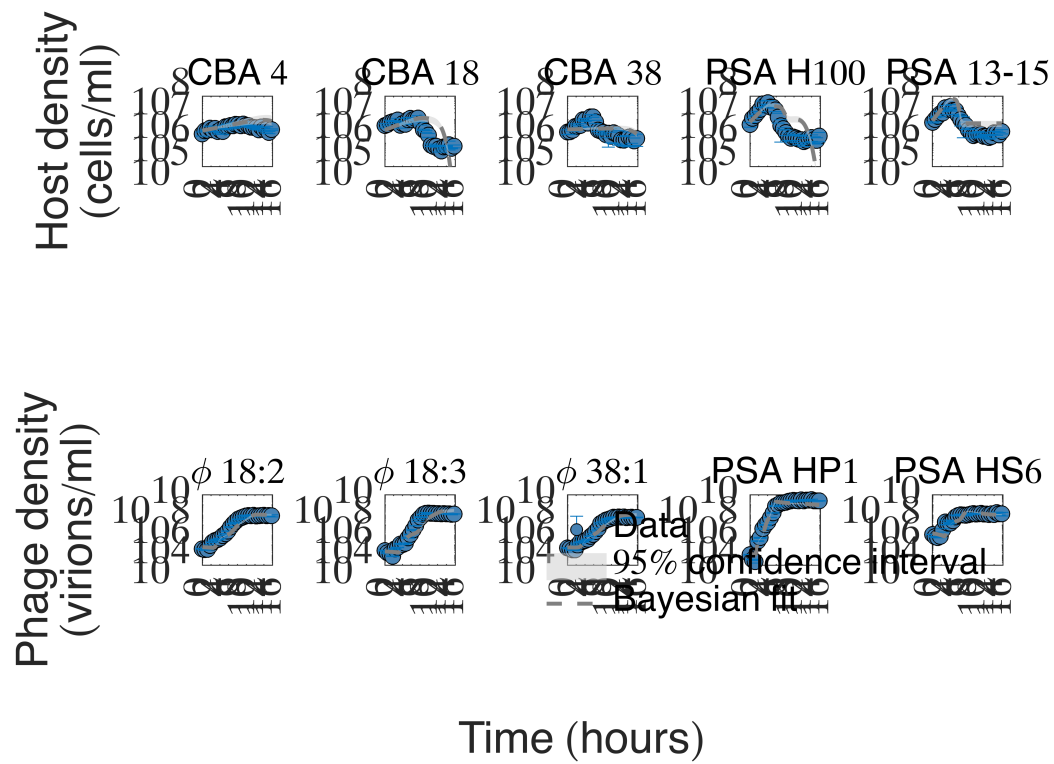
```
cd('./figure1/')  
figure1;
```



```
cd ..
```

Here is Figure 3

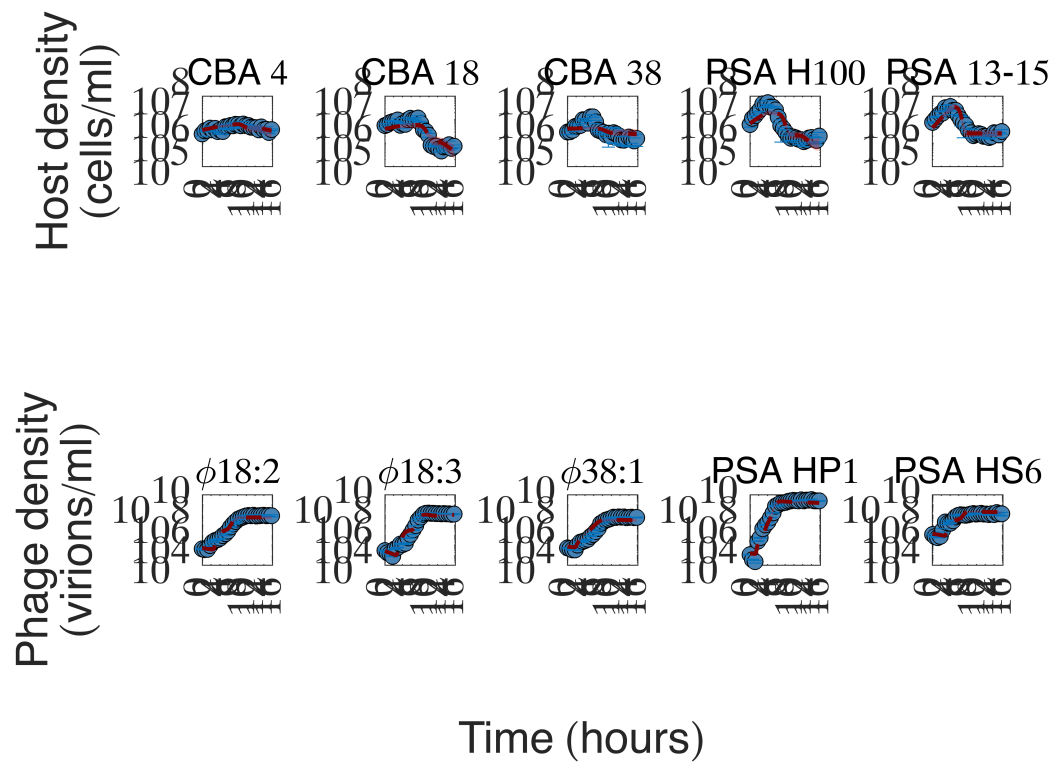
```
cd('./figure3');  
fig3_condifence;
```



```
cd(' ./.. ');
```

Here is Figure 4

```
cd(' ./figure4')
fig4_condifence;
```

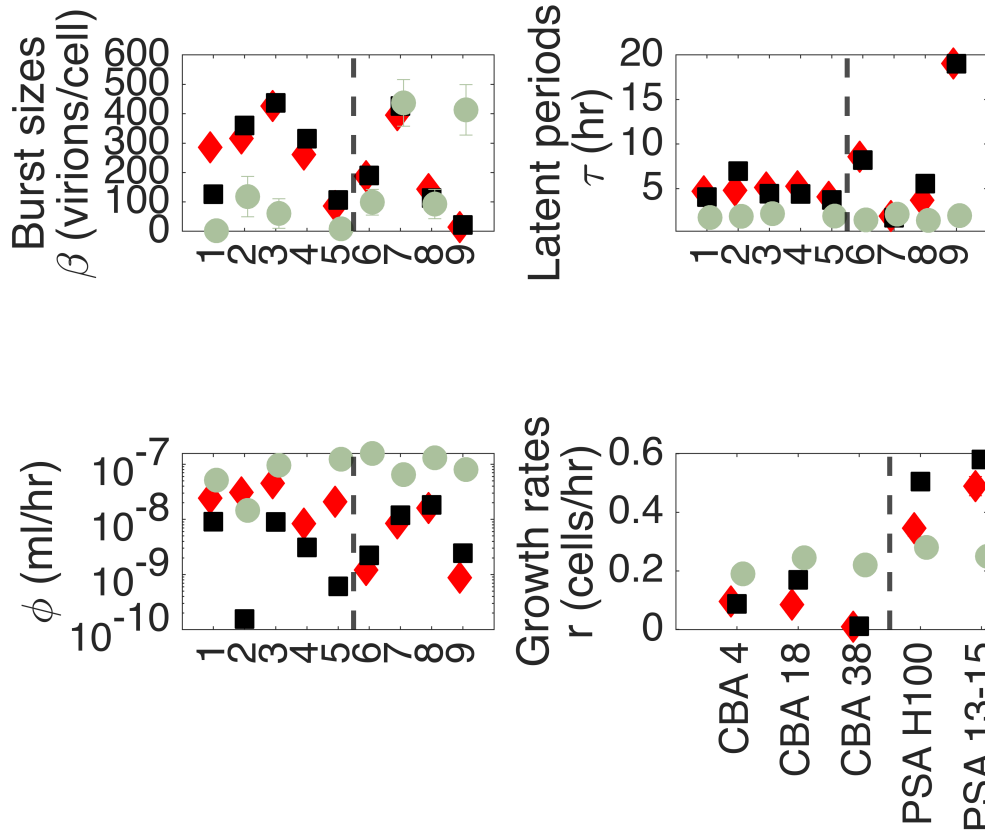


```
cd('../..');
```

Figure 5

```
cd('../figure5');
life_history_compare_new;
```

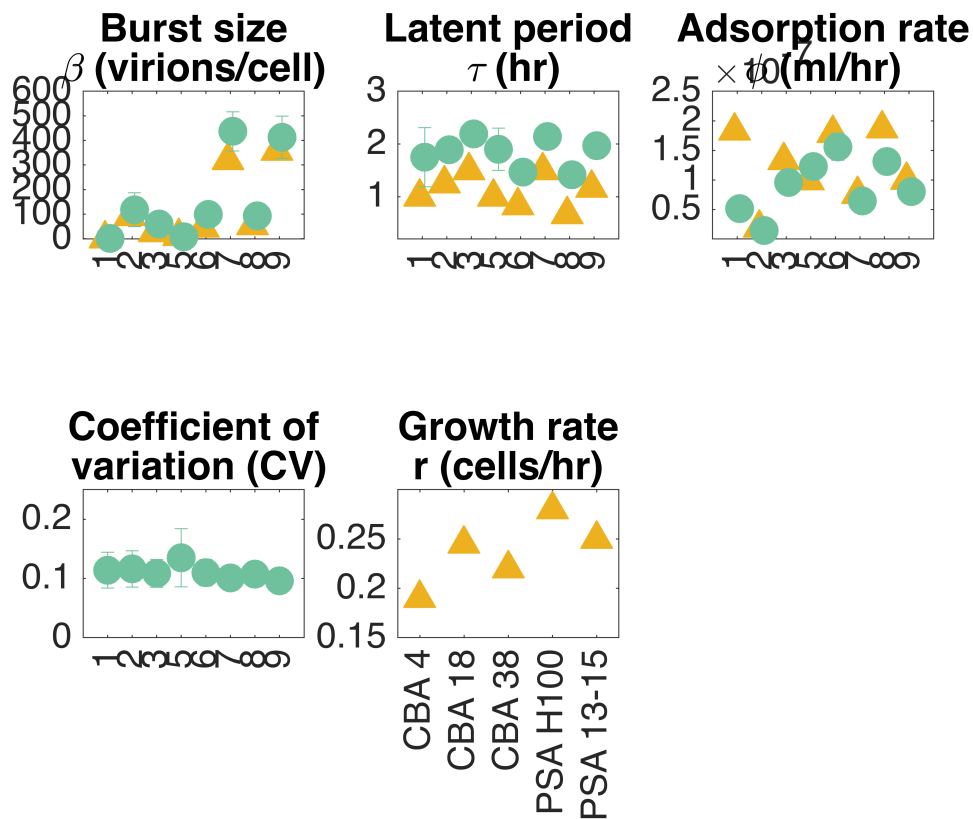
```
Warning: Could not find appropriate function
on path loading function handle /storage/coda1/p-jweitz3/0/rdey33/VIMIM0/
finalizing_script.m>@(theta,data)ssfun(theta,data,pars2,mcmcpars,model,lambda)
Warning: Could not find appropriate function
on path loading function handle /storage/coda1/p-jweitz3/0/rdey33/VIMIM0/
finalizing_script.m>@(theta,data)ssfun(theta,data,pars2,mcmcpars,model,lambda)
Warning: Could not find appropriate function on path loading function handle /storage/coda1/p-
jweitz3/0/rdey33/VIMIM0/finalizing_script.m>@(chain)median(chain)
```



```
cd(' ./... ');
```

Here is the life history comparison from Fig 1

```
fig1_parameters
```



For Figure 2, use this (takes a long time to run)

```
cd './figure2/';
figure3;
```

```
hf =
    Figure (16) with properties:
        Number: 16
        Name: ''
        Color: [9.4000e-01 9.4000e-01 9.4000e-01]
        Position: [476 460 560 420]
        Units: 'pixels'

    Show all properties
dummy_beta =
    1
dummy_beta =
    51
dummy_beta =
    101
dummy_beta =
    151
dummy_beta =
    201
dummy_beta =
    251
dummy_beta =
    301
dummy_beta =
    351
dummy_beta =
```

```

401
dummy_beta =
451
dummy_beta =
501
dummy_beta =
551
dummy_beta =
601
dummy_beta =
651
dummy_beta =
701
dummy_beta =
751
dummy_beta =
801
dummy_beta =
851
dummy_beta =
901

```

