

HIV Model

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```
In [69]: # Configure Jupyter so figures appear in the notebook
%matplotlib inline

# Configure Jupyter to display the assigned value after an assignment
%config InteractiveShell.ast_node_interactivity='last_expr_or_assign'

# import functions from the modsim library
from modsim import *

print('If this cell runs successfully, it produces no output other than this message')
```

If this cell runs successfully, it produces no output other than this message.

```
In [70]: init = State(r=200, l=0, e=0, v=100)
```

Out[70]:

values	
r	200
l	0
e	0
v	100

```
In [71]: system = System(tau = 0.2,
                        gamma = 1.36,
                        mu = 0.00136,
                        beta = 0.00027,
                        alpha = 0.036,
                        delta = 0.33,
                        pi = 100,
                        sigma = 2,
                        rho = 0.1,
                        dt = .5,
                        t0 = 0)
```

Out[71]:

	values
tau	0.20000
gamma	1.36000
mu	0.00136
beta	0.00027
alpha	0.03600
delta	0.33000
pi	100.00000
sigma	2.00000
rho	0.10000
dt	0.50000
t0	0.00000

In [72]: `def update_func(init, system, t):`



```
    r, l, e, v = init
    unpack(system)

    dr = (gamma * tau) - (mu * r) - (beta * r * v)
    dl = (rho * beta * r * v) - (mu * l) - (alpha * l)
    de = ((1-rho) * beta * r * v) + (alpha * l) - (delta * e)
    dv = (pi * e) - (sigma * v)

    r += dr*dt
    l += dl*dt
    e += de*dt
    v += dv*dt

    return State(r=r, l=l, e=e, v=v)
```

In [73]: `def run_simulation(update_func, system):`

`"""Runs a simulation of the system.`

`system: System object`
 `update_func: function that updates state`

`returns: TimeFrame`
 `"""`

`unpack(system)`

`frame = TimeFrame(columns=init.index)`
 `frame.row[t0] = init`



`for t in linrange(t0, 120, dt):`
 `frame.row[t+dt] = update_func(frame.row[t], system, t)`

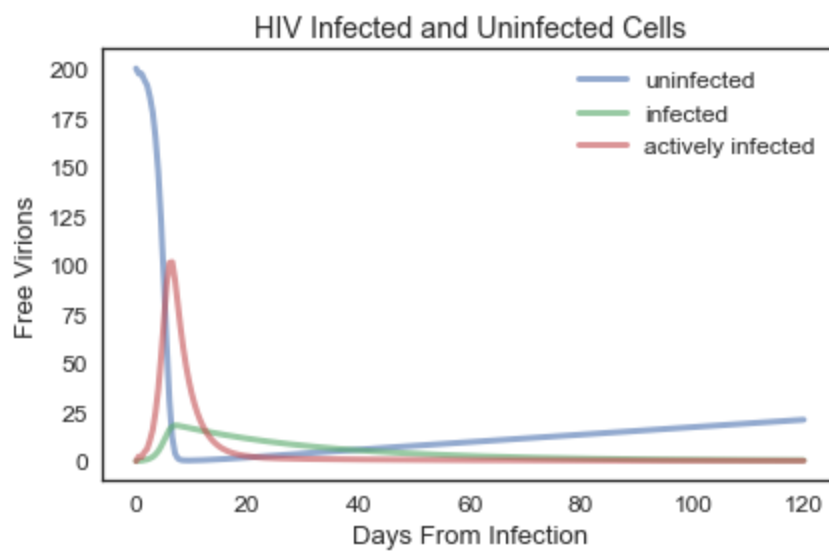


```
return frame
```

```
In [74]: data = run_simulation(update_func, system)
print()
```

```
In [79]: plot(data.r, label='uninfected')
plot(data.i, label='infected')
plot(data.e, label='actively infected')
# plot(data.v)

decorate(title='HIV Infected and Uninfected Cells',
          xlabel='Days From Infection',
          ylabel='Free Virions',
          loc='upper right')
```



```
In [ ]:
```