# Positive feedback measures

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#### Positive feedback model

The purpose of this document is to present and explore different measures of positive feedback. For this, we consider a model governed by the following three reactions:

$$\emptyset \stackrel{k_1}{\underset{k_2}{\rightleftharpoons}} A \xrightarrow[k_3]{} A + A$$

where  $k_1$  is the birth rate,  $k_2$  is the decay rate and  $k_3$  is the positive feedback strength. We want to study stochastic realisations of this model using the stochastic simulation algorithm which is implemented in the function "simple\_positive\_feedback.r", so first we must source the function.

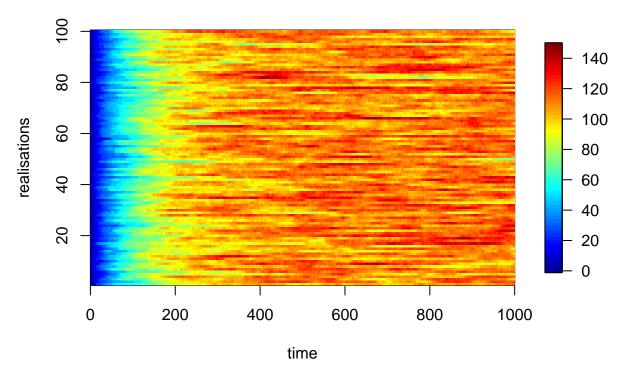
```
source('simple_positive_feedback.r')
```

Let's define parameters, run 100 trajectories of the model and obtain the output:

```
\begin{array}{l} k\_1 = 1 \\ k\_2 = 0.01; \\ k\_3 = 0.001; \\ maxtime = 1000; \\ timestep = 0.1; \\ number of realisations = 100; \\ output = simple\_positive\_feedback(k\_1,k\_2,k\_3,maxtime,timestep,number of realisations); \end{array}
```

We can also make a heatmap plot (you may have to install the "fields" package for this – install.packages("fields")):

```
time = seq(0,maxtime,by=timestep);
realisations = 1:numberofrealisations;
image.plot(time,realisations,output);
```

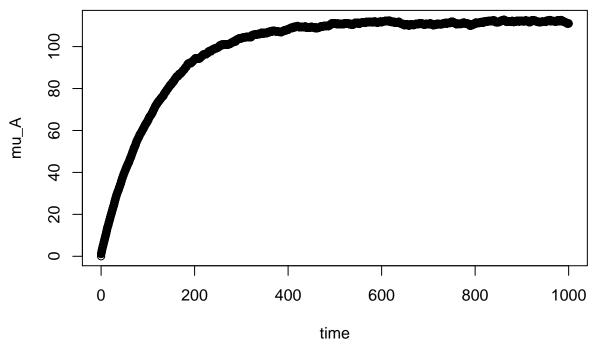


Now that we have a stochastic model with positive feedback strength as a parameter, we can try to find measures that are suitable for identifying positive feedbacks.

# Mean

The first and most obvious measure is the mean. The mean value of A should increase with the positive feedback strength,  $k_3$ . To study this, first we can plot the mean time series for a single value of  $k_3$ :

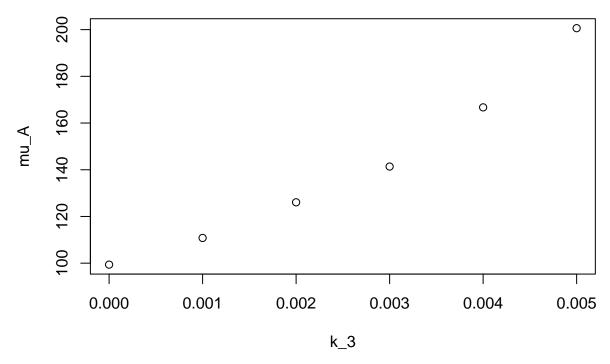
```
time = seq(0,maxtime,by=timestep);
mu_A = rowMeans(output);
plot(time,mu_A);
```



the value of  $k_3$  is small, the mean value ( $\mu$ ) of A takes a value just above the steady state with no positive feedback ( $\frac{k_1}{k_2}$ ). We can now look at how the final value of this time series varies with  $k_3$ .

Since

```
count = 0;
mu_A <- rep(0, 6);
for (k_3 in seq(0,0.005,by=0.001)){
  count = count + 1;
  output = simple_positive_feedback(k_1,k_2,k_3,maxtime,timestep,numberofrealisations);
  temp_mean_A = rowMeans(output);
  mu_A[count] = temp_mean_A[length(temp_mean_A)];
}
k_3 = seq(0,0.005,by=0.001);
plot(k_3,mu_A);</pre>
```



We can see that the mean value increases with positive feedback strength. The next measure we will inspect is the noise of A, specifically the Fano Factor.

#### **Fano Factor**

The Fano Factor (F) of A is defined as

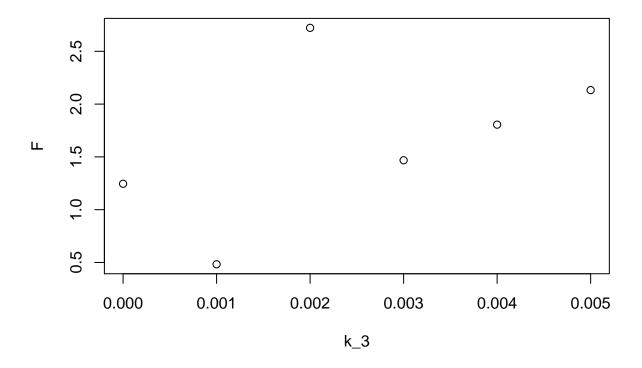
$$F = \frac{\sigma^2}{\mu}$$

where  $\sigma^2$  is the variance of A. To compute this, we need to define the variance, which can be done using the following function:

```
RowVar <- function(x) {
  rowSums((x - rowMeans(x))^2)/(dim(x)[2] - 1)
}</pre>
```

We can now examine the examine the relationship between the positive feedback strength and Fano Factor. Probably need more realisations to get a decent estimate of the Fano Factor.

```
numberofrealisations = 20;
count = 0;
F <- rep(0, 6);
for (k_3 in seq(0,0.005,by=0.001)){
   count = count + 1;
   output = simple_positive_feedback(k_1,k_2,k_3,maxtime,timestep,numberofrealisations);
   temp_mean_A = rowMeans(output);
   temp_var_A = RowVar(output);
   F[count] = temp_var_A[length(temp_var_A)]/temp_mean_A[length(temp_mean_A)];
}
k_3 = seq(0,0.005,by=0.001);
plot(k_3,F);</pre>
```



## Autocorrelation

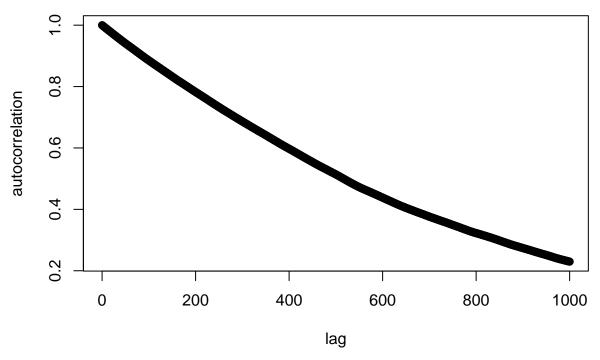
For a discrete process with known mean and variance for which we observe n observations  $\{X_1, X_2, \dots, X_n\}$ , an estimate of the autocorrelation may be obtained as

$$\hat{R}(k) = \frac{1}{(n-k)\sigma^2} \sum_{t=1}^{n-k} (X_t - \mu)(X_{t+k} - \mu)$$

We can plot an example autocorrelation for a single trajectory of the model and plot it.

```
output = simple_positive_feedback(1,0.01,0.001,1000,0.1,1);
  autocorr = acf(output,lag.max = 1000,plot = FALSE)
test <- min(abs(autocorr$acf-0.5))
half_autocorr = which(abs(autocorr$acf-0.5) == test)
lag = autocorr$lag;
autocorrelation = autocorr$acf;
plot(lag,autocorrelation,main=paste("Half autocorrelation = ", half_autocorr))</pre>
```

# Half autocorrelation = 517



Now we can examine how the half autocorrelation (which is displayed in the title of the previous plot) varies with  $k_3$ .

```
numberofrealisations = 10;
maxlag=5000;
count1 = 0;
half_autocorr <- matrix(0,nrow=6,ncol=numberofrealisations);</pre>
for (k_3 in seq(0,0.005,by=0.001)){
  count1 = count1 + 1;
  autocorr <- matrix(0,1,ncol=maxlag);</pre>
  for (count2 in seq(1,numberofrealisations,by=1)){
  output = simple_positive_feedback(1,0.01,k_3,1000,0.1,1);
  autocorr1 = acf(output,lag.max = maxlag,plot = FALSE);
  test <- min(abs(autocorr1$acf-0.5))</pre>
  autocorr=rbind(autocorr,as.vector(autocorr1$acf))
  half_autocorr[count1,count2] = which(abs(autocorr1$acf-0.5) == test);
  if (count1==1){
  plot(seq(1,maxlag,by=1),colMeans(autocorr),type="l",col=count1)
  else{
lines(seq(1,maxlag,by=1),colMeans(autocorr),type="l",col=count1)
#lines(lag, autocorrelation, col="green")
```

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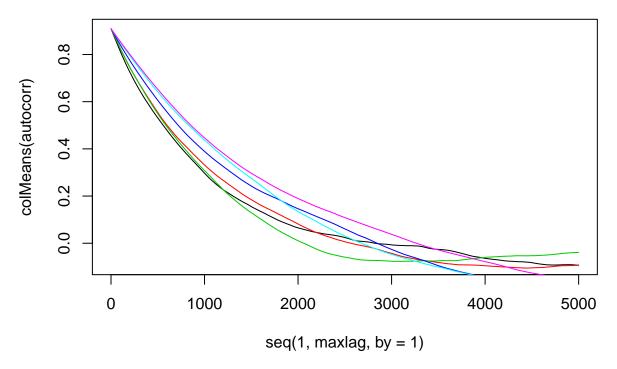
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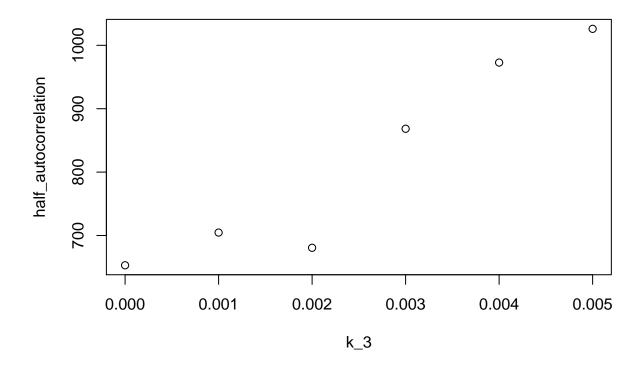
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```
k_3 = seq(0,0.005,by=0.001);
half_autocorrelation = rowMeans(half_autocorr)
plot(k_3,half_autocorrelation);
```



# Entropy

Named after Boltzmann's H-theorem, Shannon defined the entropy H (Greek letter H) of a discrete random variable X with possible values  $x_1, \ldots, x_n$  and probability mass function P(X) as:

$$H(X) = E[I(X)] = E[-\ln(P(X))]$$

.

Here E is the expected value operator, and I is the information content of X. I(X) is itself a random variable. The entropy can explicitly be written as

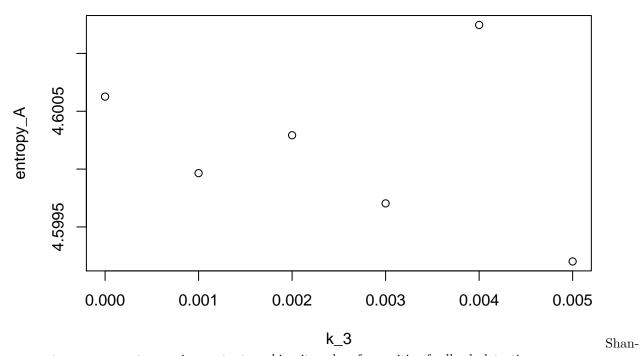
$$H(X) = \sum_{i=1}^{n} P(x_i) I(x_i) = -\sum_{i=1}^{n} P(x_i) \log_b P(x_i),$$

where b is the base of the logarithm used. Common values of b are 2, Euler's number e, and 10, and the unit of entropy is shannon for b = 2, nat for b = e, and hartley for b = 10. When b = 2, the units of entropy are also commonly referred to as bits.

We can see how the entropy varies with  $k_3$  by doing the following:

```
numberofrealisations = 100;
entropy_A <- rep(0,6);
count1 = 0;
for (k_3 in seq(0,0.005,by=0.001)){
   count1 = count1 + 1;
   A_end <- rep(0, numberofrealisations);
   for (count2 in seq(1,numberofrealisations,by=1)){
   output = simple_positive_feedback(1,0.01,k_3,1000,0.1,1);
   temp_A = output[length(output)]; #use final time series value
   A_end[count2] = temp_A;</pre>
```

```
}
entropy_A[count1] = entropy(A_end);
}
k_3 = seq(0,0.005,by=0.001);
plot(k_3,entropy_A);
```



non entropy appears to remain constant, making it useless for positive feedback detection.