

Random Covers

Problem:

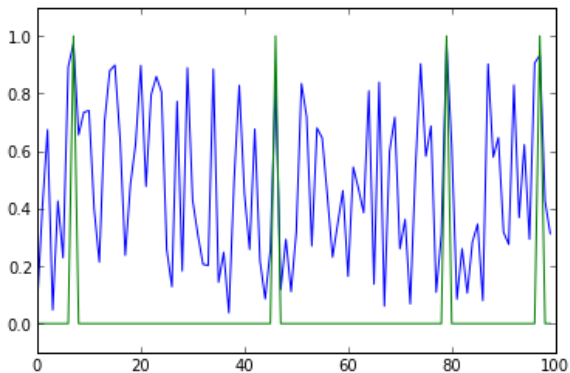
With only local information passing and processors at random locations, divide a 1D, 2D, or 3D region into subregions of approximately equal size.

Covering Protocol:

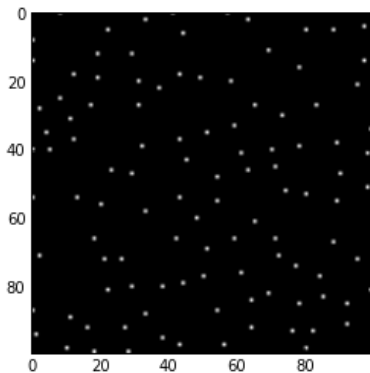
Goal: divide processors in a n -dimensional space into equal sized connected regions of size approximately d .

- ▶ every processor picks a random number
- ▶ processors broadcast their number to all neighbors within distance d
- ▶ processors become cell centers if they picked the maximum value among all their neighbors

```
1 a = rand(100)
2 b = filters.maximum_filter(a,20)
3 ylim((-0.1,1.1)); plot(a); plot(a==b)
```



```
1 a = rand(100,100)
2 b = filters.maximum_filter(a,10)
3 imshow(a==b)
```



Problems:

- ▶ everything is noisy
- ▶ can't make perfect comparisons
- ▶ can't make a binary decision based on a threshold
- ▶ only nearest neighbor communications

Thresholding by Logistic Growth

Thresholding:

Idea: use logistic regression and growth rate r

$$\dot{u} = ru(1 - u)$$

If $r > 0$, this converges to 1 .

If $r < 0$, this converges to 0 .

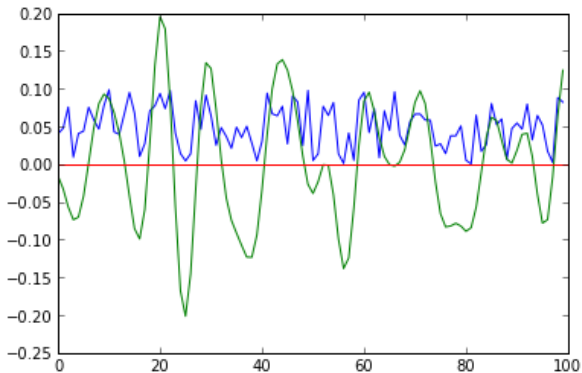
We can make r itself a variable.

Laplacian of Gaussian (or Difference of Gaussian):

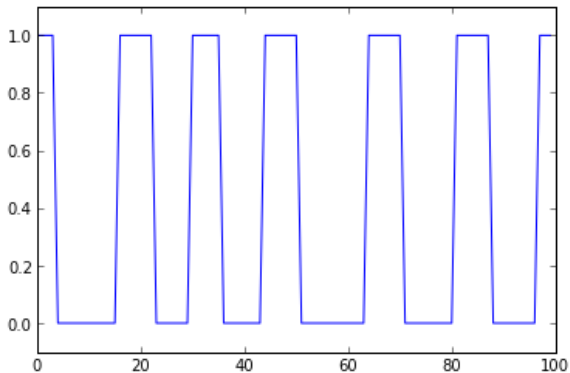
Use as the growth rate the Laplacian of Gaussian (LoG).

This will give rise to regions of positive and negative growth, spaced by about the size of the Gaussian.

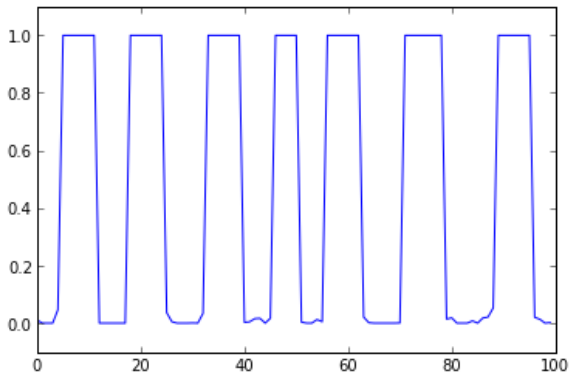
```
1 a = 0.1*rand(100)
2 r = -100*filters.gaussian_laplace(a,3.0)
3 plot(a); plot(r); plot([0,100],[0,0])
```



```
1 a = 0.1*rand(100)
2 for t in range(1001):
3     r = -100*filters.gaussian_laplace(a,3.0)
4     a = clip(a+r*a,0.001,0.999)
5 ylim((-0.1,1.1)); plot(a)
```

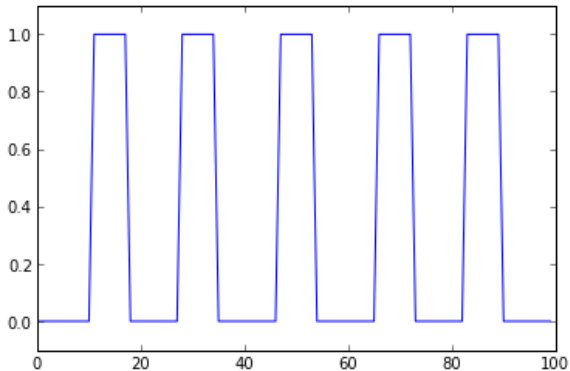


```
1 a = 0.1*rand(100)
2 for t in range(1001):
3     r = -100*filters.gaussian_laplace(a,3.0)
4     a = clip(a+r*a+0.03*rand(100),0.001,0.999)
5 ylim((-0.1,1.1)); plot(a)
```

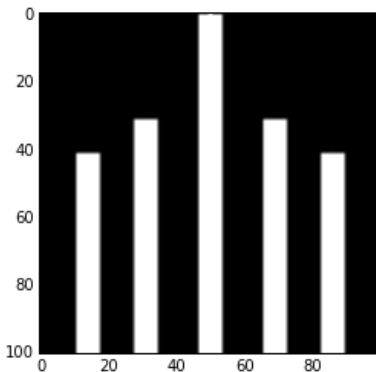


Wavefront Propagation

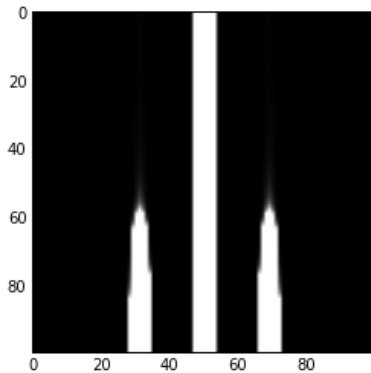
```
1 a = zeros(100)
2 a[len(a)//2] = 1
3 for t in range(10001):
4     r = -50*filters.gaussian_laplace(a,3.0)
5     a = clip(a+r*a,0.001,0.999)
6 ylim((-0.1,1.1)); plot(a)
```



```
1 a = zeros(100)
2 a[len(a)//2] = 1
3 result = []
4 for t in range(10001):
5     r = -50*filters.gaussian_laplace(a,3.0)
6     a = clip(a+r*a,0.001,0.999)
7     result.append(a)
8 result = array(result)
9 imshow(result[:,100])
```



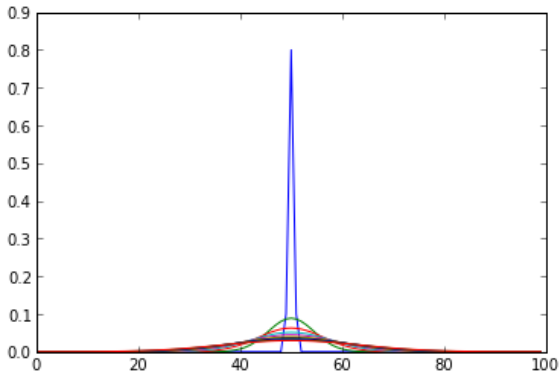

```
1 imshow(result[3100:3200,:])
```



Smoothing by Diffusion

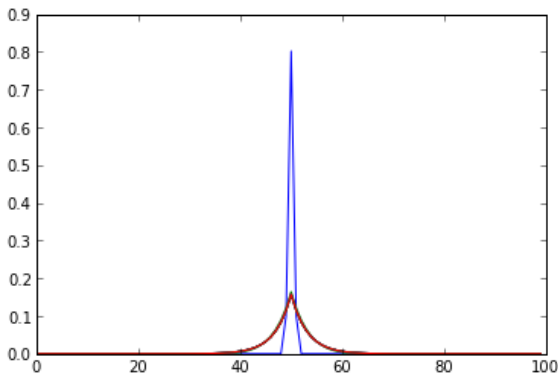
As we have seen, simple diffusion is like smoothing with larger and larger Gaussians over time.

```
1 a = zeros(100); a[len(a)//2] = 1
2 for t in range(1000):
3     a += 0.1*laplacian(a)
4     if t%100==0: plot(a)
```



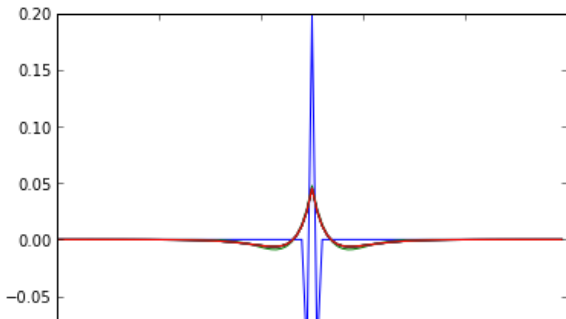
For the kind of center-surround inhibition we want for pattern formation, we need to construct a fixed degree of smoothing. We use the heat equation.

```
1 a = zeros(100); a[len(a)//2] = 1; b = a.copy()
2 for t in range(1000):
3     b += 0.1*laplacian(b)
4     b = 0.99*b+0.01*a
5     if t%100==0: plot(b)
```



We can put together two of these now to obtain a center-surround filter.

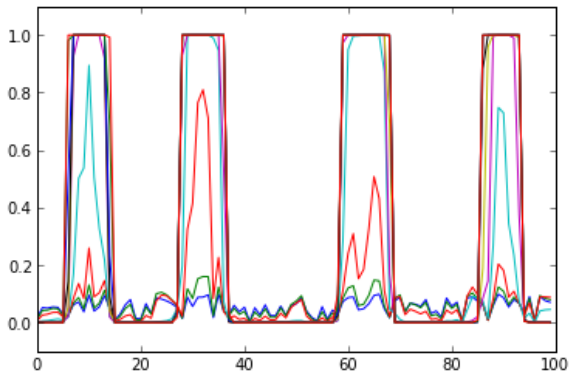
```
1 a = zeros(100)
2 a[len(a)//2] = 1
3 b = a.copy()
4 c = a.copy()
5 for t in range(1000):
6     b += 0.1*laplacian(b)
7     b = 0.99*b+0.01*a
8     c += 0.2*laplacian(c)
9     c = 0.99*c+0.01*a
10    if t%100==0: plot(b-c)
```



Complete Reaction-Diffusion

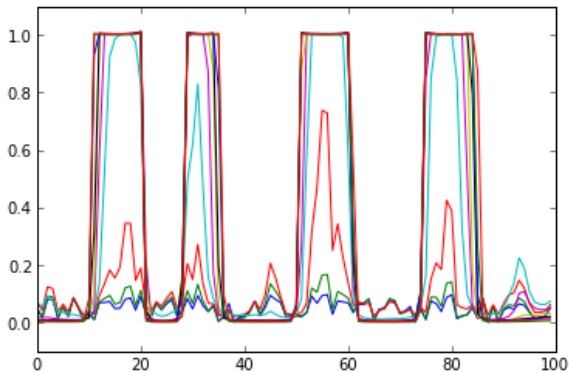
Now we use the two diffusions together with the logistic growth equation to arrive at a pattern formation equation.

```
1 a = 0.1*rand(100); b = a.copy(); c = a.copy()
2 for t in range(100000):
3     b += 0.1*laplacian(b); b = 0.99*b+0.01*a
4     c += 0.2*laplacian(c); c = 0.99*c+0.01*a
5     a += 0.01*(b-c)*a*(1-a)
6     if t%10000==0: ylim((-0.1,1.1)); plot(a)
```



As before, we can add a bit of random growth to cause the stripes to shift closer together over time.

```
1 a = 0.1*rand(100); b = a.copy(); c = a.copy()
2 for t in range(100000):
3     b += 0.1*laplacian(b); b = 0.99*b+0.01*a
4     c += 0.2*laplacian(c); c = 0.99*c+0.01*a
5     da = 0.01*(b-c)*a*(1-a); a += da + rand(len(a))*amax(da)*0.1
6     if t%10000==0: ylim((-0.1,1.1)); plot(a)
```



Mathematically, our reaction-diffusion equation now looks as follows.

$$\dot{b} = r_b \nabla^2 b + \lambda_b a - \gamma_b b$$

$$\dot{c} = r_c \nabla^2 c + \lambda_c a - \gamma_c c$$

$$\dot{a} = \gamma_a (b - c) a (1 - a) + \nu_a \epsilon$$

Here, ν_a and ϵ is an additional noise term (stochastic differential equation).

If the $\gamma_a \ll \gamma_b, \gamma_c$, then this behaves as originally designed:

- ▶ b and c are approximately smoothed versions of a
- ▶ the difference between b and c is a Laplacian-like center-surround operator
- ▶ each location tends towards 1 or 0 depending on whether it and its nearby growth rate is bigger or smaller than the intermediate average

Notes for the implementation:

- ▶ diffusion can be implemented with completely local communication
- ▶ diffusion can be implemented by randomly passing around a number of discrete tokens
- ▶ the exact form of the center-surround operator doesn't matter
- ▶ when γ_a is similar in magnitude to γ_b , we get complex dynamic behaviors (see in a minute)
- ▶ in most actual versions, variables b and c are combined

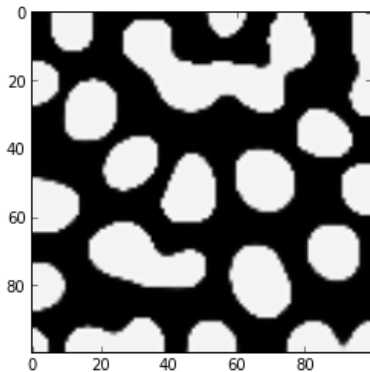
2D Pattern Formation

Everything works exactly the same way in 2D.

```
1 a = 0.1*rand(100,100); b = a.copy(); c = a.copy()
2 for t in range(100000):
3     b += 0.1*laplacian(b); b = 0.99*b+0.01*a
4     c += 0.2*laplacian(c); c = 0.99*c+0.01*a
5     da = 0.01*(b-c)*a*(1-a); a += da + rand(len(a))*amax(da)*0.1
```

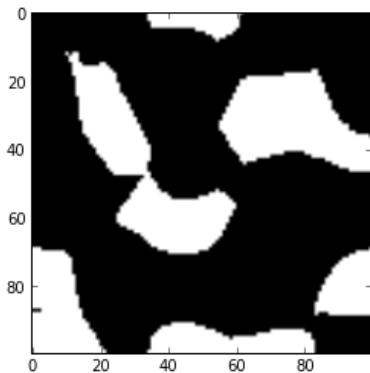
In 2D, we get spots (leopards) or stripes (zebras).

```
1 imshow(a)
```



Since we motivated pattern formation in reaction-diffusion systems as LoG smoothing followed by thresholding, what does that look like?

```
1 noise = randn(100,100)
2 gl = filters.gaussian_laplace(noise,12.0)
3 imshow(gl>scoreatpercentile(gl,70))
```



But, actually, we are effectively iterating the thresholding and smoothing multiple times, giving rise to this kind of iteration. This looks indistinguishable from reaction-diffusion patterns (for different parameters than above).

```
1 noise = randn(100,100)
2 gl = noise.copy()
3 for i in range(100):
4     gl = filters.gaussian_laplace(gl,8.0,mode='wrap')
5     gl = 1.0*(gl>scoreatpercentile(gl,50))
6 imshow(gl)
```



Summary

Reaction-Diffusion Systems:

Reaction-diffusion systems generate patterns by first smoothing noise, then amplifying and thresholding it.