



Weed model example using a lattice

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Meetup

You should keep in mind



Weed models

- Weeds are plants
- Process based crop models
- Life cycle

 Competition of plant species can be modelled with lattice models

Lattice



Cells State of the cells, here 0 or 1 (TRUE/FALSE)

Lattice models

- Cells, state of the cells (0 or 1)
- Discrete in space and discrete in time
- State update depends on the neighbourhood

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Lattice models

- Cells, state of the cells (0 or 1)
- State update depends on the neighbourhood
- Conway's Game of Life
- Cellular automata (automaton)

Cellular automata in biology



Cellular automata

- Simple lattice model
- Cells, state of the cells (0 or 1)
- State update depends on the neighbourhood

Neighbourhood in the lattice

- Moore (Chebyshev distance)
- von Neumann (Manhattan distance)



We can generalise

Manhattan Distance



Chebyshev Distance



 $|x_1 - x_2| + |y_1 - y_2|$ max $(|x_1 - x_2|, |y_1 - y_2|)$

Stucture squares vs hexagons

Civ1 vs Civ5



Prehaps the hexagons have clearer neighbourhood propeties, but programming skills

Edge of the lattice





Create a torus – no edge effect

Lattice size

- Rule of thumb: bigger is better
- BUT CPU time

- Sensitivity analysis
 - to find a lattice size have no effect on the output

Moore nhood - R

field <- matrix(sample(0:1,64,replace=T),ncol=8)
this is only 0s and 1s
field
field[5,4] # a particular cell
1:5 # easy way to do a sequence in R
field[4:6,3:5] # can be used in []</pre>

How many 1s do we have in the **Moore** nhood of this cell?

```
sum(field[4:6,3:5])-field[5,4]
```

Moore nhood - R

```
# Lets assume we have crop cells (C), weed cells (W)
and cells without either of this two (else, E)
field2 <- matrix(sample(c("C", "W", "E"), 64,</pre>
      replace=T), ncol=8)
field2
field2[6,3]# a particular crop cell
field2[5:7,2:4] # the nhood
# How many weed cells do we have in Moore nhood of
this cell?
sum(field2[5:7,2:4]=="W")
```

Toroidal arrangement - R

Instead of adding and subtracting we define a new vector of going South (down) in the lattice #goingS should be (2,3,4,...,n,1) goingS <- c(2:8,1) # and going North (up) #goingN should be c(n,1,2,3...,n-1) goingN <- c(8,1:7)</pre>

```
#same for East and West
goingW <- goingS
goingE <- goingN</pre>
```

Toroidal arrangement - R

Thus, the number of weed cells in Moore nhood of a
cell [8,2]:

sum(field2[c(goingN[8],8,goingS[8]),c(goingE[2],2,go
ingW[2])]=="W")