

Complexity in social dynamics : from the micro to the macro

Laboratory 4

Franco Bagnoli

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# 1 Laboratory 4

## Evolution and game theory

1. Evolution on a flat landscape: mutations.
2. Evolution on a smooth fitness landscape: quasispecies and the Red Queen.
3. Evolution of a sharp landscape: the error threshold.
4. Niches and coexistence in fitness landscapes.
5. Competition: a stabilizing force.
6. Evolution of cooperation: direct reciprocity.

## Evolutionary models

- An individual is modeled as a vector of  $L$  genes (that take value 0 and 1).
- The phenotype  $f$  is just the sum of the genes ( $0 \leq f \leq L$ ).
- The fitness is a function of the phenotype and the phenotype distribution of other individuals (for competition).
- For modeling an evolutionary population (fixed size: `quasispecies.f90`), we just compare the fitness of two individuals. That with lower fitness tends to disappear, replaced by a copy of the opponent, with eventual mutations.

```
quasispecies.f90
module fit
contains
  function pheno(g)
    integer :: pheno
    integer*1 :: g(:)
    integer :: L,i

    L = size(g)
    pheno = 0
    do i=1,L
      pheno = pheno + g(i)
    end do
  end function

  function fitness(f, fd)
    implicit none
    real :: fitness
    integer :: f ! phenotype
    integer :: fd(0:) ! phenotype distribution
    integer :: LL, i
    real :: d,h
    integer :: NN

    LL = size(fd)
    LL = LL-1
    NN = sum(fd)

    !fitness=1 ! neutral evolution

    ! a sharp-peaked landscape
```

```

!if (f == 0) then
! fitness = 1
!else
! fitness = 0
!end if
! smooth landscape
!fitness = (real(LL-f)/LL)**.1
! smooth landscape
fitness = abs(real(LL/2-f)/LL)**1.01

!competition
! h = 0
! do i=0,LL
!     d = abs(i-f) ! phenotypic distance
!     h = h - 1*fd(i)*exp(-.01*d)/NN
! end do
! fitness = exp(h)
end function
end module

program quasispecies
use fit
implicit none
integer, parameter :: N=5000 !population size
integer, parameter :: L=200 ! genome size
integer :: i, j, k, kk, t
integer :: TMAX=10000
integer*1 :: x(N,L) !population
real :: A(N) ! fitness
integer :: f(N) ! phenotype
integer :: fd(0:L) ! phenotype distribution
real :: r , AA
real :: mu=0.0001 !mutation prob per gene
real :: Temp=0.1
character*200 :: str

call random_seed()
!initialization
fd = 0
do i=1, N
do j=1, L
call random_number(r)
x(i,j) = floor(r+0.5)
end do
!x(i,:) = 0
f(i) = pheno(x(i,:))
fd(f(i)) = fd(f(i))+1
end do

! compute fitness
do i=1, N
A(i) = fitness(f(i), fd)
end do

call gnuplotOpen("gnuplot")

```

```

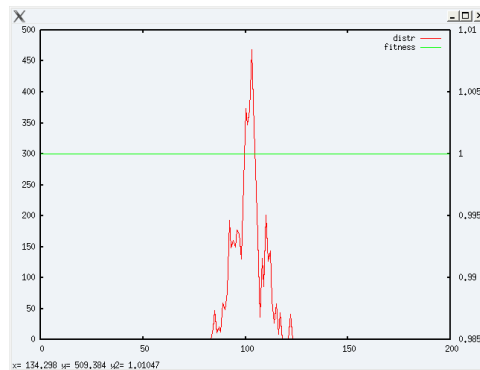
call gnuplotExecute("set term x11; set mouse;");//&
    "set ytics nomirror; set y2tics")

! evolution
do t=1, TMAX
  do kk=1, N ! a MonteCarlo step
    ! pick two individuals at random
    call random_number(r)
    i = floor(r*N)+1
    call random_number(r)
    j = floor(r*N)+1
    ! selection
    call random_number(r)
    if (r < 1/(1+exp((A(i)-A(j))/Temp))) then
      k=i; i=j; j=k ! exchange i and j
    end if
    ! i duplicates into j
    fd(f(j)) = fd(f(j)) -1 ! remove j
    ! clone i into j
    x(j,:) = x(i,:)
    !mutations: up to 1 mutation at random
    call random_number(r)
    if (r< mu*L) then
      call random_number(r)
      k = floor(r*L)+1
      x(j,k) = 1-x(j,k)
    end if
    ! compute phenotype and fitness
    f(j) = pheno(x(j,:))
    fd(f(j)) = fd(f(j)) +1
    A(j) = fitness(f(j), fd)
  end do
  AA = sum(A)/N ! average fitness
  print *, AA
  !plot
  call gnuplotExecute("plot '-' t 'distr' w l, "//&
    "-'" axis x1y2 t 'fitness' w l")
  do i=0,L
    write(str, *) i, fd(i)
    call gnuplotExecute(str)
  end do
  call gnuplotExecute("end")
  do i=0,L
    write(str, *) i, fitness(i, fd)
    call gnuplotExecute(str)
  end do
  call gnuplotExecute("end")
  call gnuplotFlush()
end do
end program

```

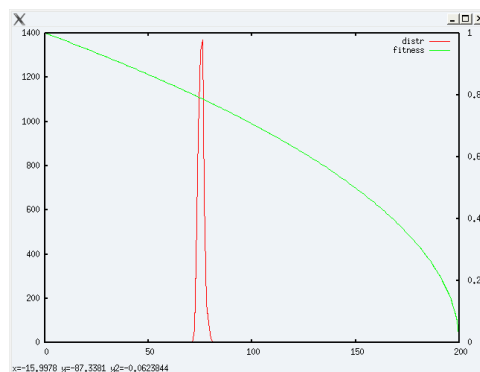
## Evolution on a flat landscape

- Without selection (flat fitness landscape), mutations (random drift) tend to favor the intermediate phenotype (0.5). The asymptotic distribution approximates a binomial one.
- Notice that for a genome length  $L$  sufficiently high, the binomial distribution is so sharp that, for finite populations, the extreme values (say, genotype 0, 0, 0, ... never appear.



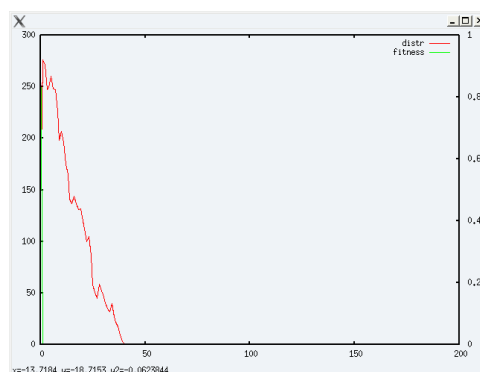
### Evolution on smooth landscapes

- In smooth landscapes, there is a competition between fitness (order) and mutations (disorder).
- you can try to explore the relationship between fitness shape, mutations and position and width of quasispecies.



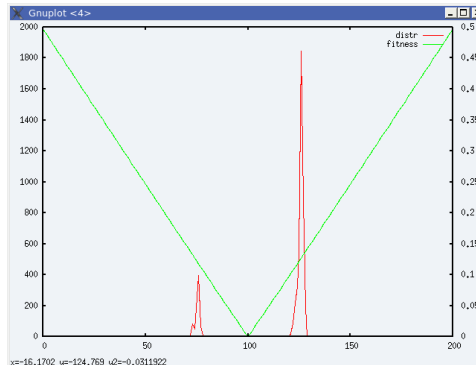
### Evolution on sharp landscapes: error threshold

- For a sharp landscape and finite populations, the asymptotic population is a quasispecies centered around the master sequence (here the sequence 0, 0, ...).
- it may happen that the master sequence is lost.
- Since the fitness landscape is flat (except for the master sequence), the evolution is here just a random search of a point in a high-dimensional space: no hope of finding it.



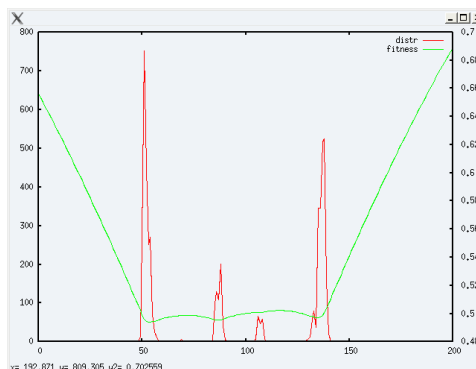
## Speciation and coexistence on smooth landscapes

- The allopatric speciation theory identifies speciations with the “discovering” of niches.
- Since niches (fitness maxima) are separated by valleys, one needs “hopeful monsters” that accumulates mutations. This is easier for smaller populations, and therefore in isolated islands.
- However, coexistence is fragile: random fluctuations may bring species into extinction.



## Competition

- Competition arises when one individual uses some resource correlated to its phenotype (example: seeds of a diameter related to its beak size).
- In principle, one should simulate at least two species (a prey and a predator), but we can use an “effective” competition term.
- In the presence of competition, more species can occupy the same niche, even if random drift “pushes” phenotypes towards the intermediate one.



## Evolutionary game theory

- In evolutionary game theory, the fitness landscape is replaced by direct interactions among individuals.
- We simulate here the evolution of cooperation by direct reciprocity (`cooperation.f90`). A population composed by TIT-FOR-TAT (0) and ALL-D (1) strategies is engaged in a round-robin tournament, and accumulates payoff.
- After that, selection takes part, as in `quasispecies.f90`.

```

program cooperation
  implicit none
  integer, parameter :: N=200 !population size
  integer :: i, j,ii, k,kk, t, ll
  integer :: TMAX=10
  integer*1 :: x(N) !population 1=tit for tat, 2=allD
  real :: A(N) ! fitness:average payoff
  real :: r , AA, xx
  real :: Temp=0.001 !selection temperature
  real :: w = 0.5 ! end tournament probability
  character*200 :: str
  real :: payoff(0:1,0:1)
  integer :: D1, D2, DD1
  real :: b = 5 ! benefit
  real :: c = 2.5 ! cost
  real :: length, ww
  real :: x0=.01 ! initial fraction of defectors

  payoff(0,0) = b-c ! CC cooperation
  payoff(0,1) = -c ! CD
  payoff(1,0) = b ! DC exploitation
  payoff(1,1) = 0 ! DD

  call random_seed()

  !initialization
  do i=1, N
    call random_number(r)
    x(i)=floor(r*x0)
  end do

  xx = 0
  do i=1, N
    xx = xx + x(i)
  end do
  xx = xx/N
  print *, "initially x=", xx

  ! evolution
  do t=1, TMAX
    A = 0
    length=0
    do k=1, N
      do kk=1, N ! complete tournament
        ! pick opponent at random
        !call random_number(r)
        !j = floor(r*N)+1
        i = k
        j = kk
        ! tournament
        ! first deal of TFT is cooperate, ALLD always defects
        D1 = x(i)
        D2 = x(j)
        A(i) = A(i) + payoff(D1,D2)
        A(j) = A(j) + payoff(D2,D1)
      end do
    end do
  end do

```

```

ll = 1
!   print *, "-----"
!   print *, ll,",", i, "(", x(i), ") plays ",d1, " and scores ", payoff(D1,D2)
!   print *, "vs ", j, "(", x(j), ") plays ",d2, " and scores ", payoff(D2,D1)
do while (.true.)
    call random_number(r)
    if (r < w) exit !end tournament
    ll = ll + 1
    if (x(i) == 0) then
        DD1 = D2 ! TFT repeats last move of opponent
    else
        DD1 = 1
    end if
    if (x(j) == 0) then
        D2 = D1
    else
        D2 = 1
    end if
    D1 = DD1
!   print *, ll,",", i, "(", x(i), ") plays ",d1, " and scores ", payoff(D1,D2)
!   print *, "vs ", j, "(", x(j), ") plays ",d2, " and scores ", payoff(D2,D1)

    A(i) = A(i) + payoff(D1,D2)
    A(j) = A(j) + payoff(D2,D1)
end do
length=length + ll
end do
end do
length = length / N**2
ww=1/length
! selection
do k=1, N
    i = k
    ! pick an opponent at random
    call random_number(r)
    j = floor(r*N)+1
!   print *, i, a(i),"vs ", j, a(j)
    ! and select
    call random_number(r)
    if (r < 1/(1+exp((A(i)-A(j))/Temp))) then
        ii=i; i=j; j=ii ! exchange i and j
    end if
!   print *, i, x(i), " wins ", j, x(j)
    ! clone i into j
    x(j) = x(i)
end do
AA = sum(A) /N
xx = 0
do i=1, N
    xx = xx + x(i)
end do
!   print *, t,xx/N, c/b, ww, ww/(2-ww),ww/(3-2*ww)
end do
print *, "after ", tmax, "rounds, x=",xx/N," c/b=", c/b, &
"; ESS (w)=", ww, "; RD (w/(2-w))=", ww/(2-ww),&
"; AD (w/(3-2w))=", ww/(3-2*ww)

```



```
end program
```

### Direct reciprocity

- Depending on the ration  $c/b$  of cost with respect to benefit, and the expected number of rounds  $1/w$ , TIT-FOR-TAT (TFT) may be evolutionary stable (ESS: cannot be invaded by a single mutant, but can be invaded by a large group of defeaters), robust (RD: a random initial condition with a small majority of TFT leads to a homogeneous population) or advantageous (AD: even in small population, a fraction of mutants larger than  $1/N$  cannot overcome).
- Try to compare the results with the mean-field approximation by Novak.