

IPD REFCARD

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Imports

from game import *
from ipd import *
from strategies import *

Game

Creation

ipd_scores = [(3, 3), (0, 5), (5, 0), (1, 1)]
ipd = game.Game(ipd_scores, ["C", "D"])

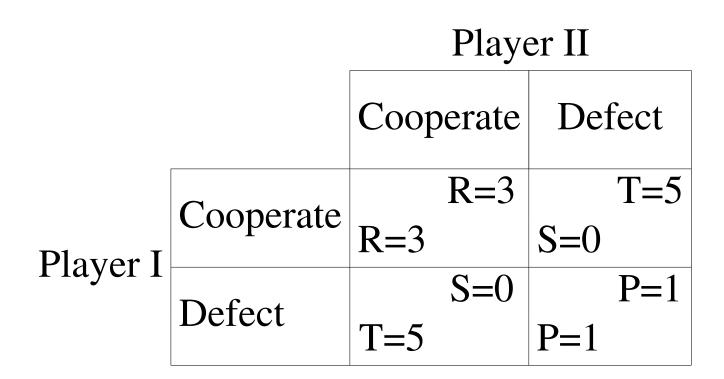
Display

ipd.prettyPrint()

Nash and Pareto optima

ipd.getNash() # provide index
ipd.getPareto() # provide index

The classical matrix



Meetings

Execution

sA = Tft()
sB = All_D()
m = Meeting(g,sA,sB,20) # default 1000
m.run()

Score de la premiere

m.s1_score

Affichage

m.prettyPrint(20)

Tournament

Execution

t = Tournament(g, getMem(1,1), 20) # default 1000 t.run()

Python display limits

pd.set_option('display.max_rows', None)

Display all the score matrix

t.matrix

Display only the results

t.matrix['Total']

Display the 10 best

t.matrix['Total'][0:10]

Get one of the winners(ex æquo possibles)

t.matrix.index[0]

Get the best score

t.matrix['Total'][0]

Ecological competitions

Execution

e = Ecological(g,getMem(1,1), 20) # default 1000
e.run()

Python display limits

pd.set_option('display.max_rows', None)

History of the evolutions

e.historic[-1:]

Display the 3 bests

e.historic.iloc[-1][0:3]

Get the best score

e.historic.iloc[-1][0]

Display all those still alive

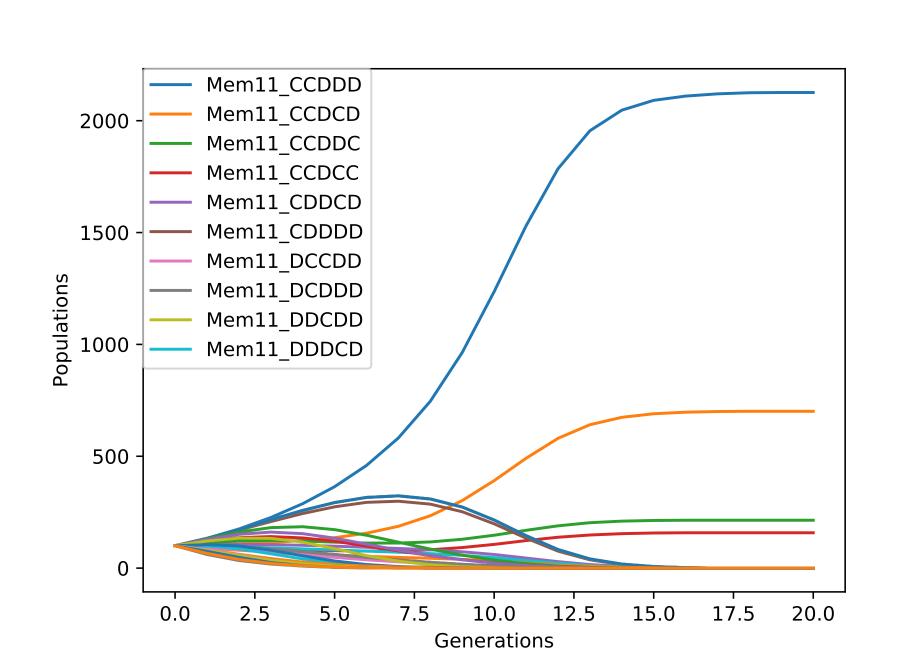
e.historic.iloc[-1][e.historic.iloc[-1]>0]

Graphical aspects

e.drawPlot() # on-screen display
e.drawPlot(save='fig.pdf') # save in a file
el.drawPlot(5,None) # only 5 lines in legend

Genotype Mem(1,2) - plot Mem(1,1)





Classical equivalences

```
Mem(0,0,'C','allc')
Mem(0,0,'D','alld')
Mem(1,0,'cDC','percd')
Mem(1,0,'dDC','perdc')
Mem(0,1,'cCD','tft')
Mem(0,1,'dCD','mistrust')
Mem(1,1,'cCDDD','spiteful')
Mem(1,1,'cCDDC','pavlov')
Mem(0,2,'ccCCCD','tf2t')
Mem(0,2,'ccCCDDD','hard_tft')
Mem(1,2,'ccCCCDDDD','slow_tft')
Mem(1,2,'ccCDCDDDDDD','spiteful_cc')
```

Tailles

Name	Size
memory(0,1)	$2^1 * 2^2 = 8$
memory(1,0)	$2^1 * 2^2 = 8$
memory(1,1)	$2^1 * 2^4 = 32$
memory(2,0)	$2^2 * 2^4 = 64$
memory(1,2)	$2^2 * 2^8 = 1024$
memory(2,1)	$2^2 * 2^8 = 1024$
memory(2,2)	$2^2 * 2^{16} = 262144$