

Anexos - Relatório 1

Anexo 1

```
simul.summary()

Metabolites: 1877
Reactions: 2712
Genes: 1516
```

Anexo 2

Reaction ID	Flux rate
0 EX_lac__L_e	0.0
1 EX_lac__D_e	0.0

Reaction ID	Flux rate
0 EX_glc_e	0.0
1 EX_glc__D_e	-10.0
2 EX_glc_e	0.0
3 EX_2ddglc_e	0.0
4 EX_glc_e	0.0
5 EX_glc_e	0.0
6 EX_udpglc_e	0.0
7 EX_5dglc_e	0.0
8 EX_2dglc_e	0.0
9 EX_metglc_e	0.0

Reaction ID	Flux rate
0 BIOMASS_Ec_iML1515_core_75p37M	0.876997
1 BIOMASS_Ec_iML1515_WT_75p37M	0.000000

Reaction ID	Flux rate
0 EX_co2_e	24.003293
1 EX_h2o2_e	0.000000
2 EX_no2_e	0.000000
3 EX_so2_e	0.000000
4 EX_o2_e	-22.131763
5 EX_o2s_e	0.000000

Anexo 3

```
const = {'EX_glc__D_e' : (-15.0, 100000.0),
        'EX_o2_e' : (0,10000)}

print(simul.FVA(reactions=["EX_lac__L_e"], constraints = const))
print(simul.FVA(reactions=["EX_lac__D_e"], constraints = const))

{'EX_lac__L_e': [0.0, 1.114181818181784]}
{'EX_lac__D_e': [0.0, 12.255999999999567]}
```

Anexo 4

```
print(best_val_l)
print(best_val_d)

[]
[('s0001', 6.86)]
```

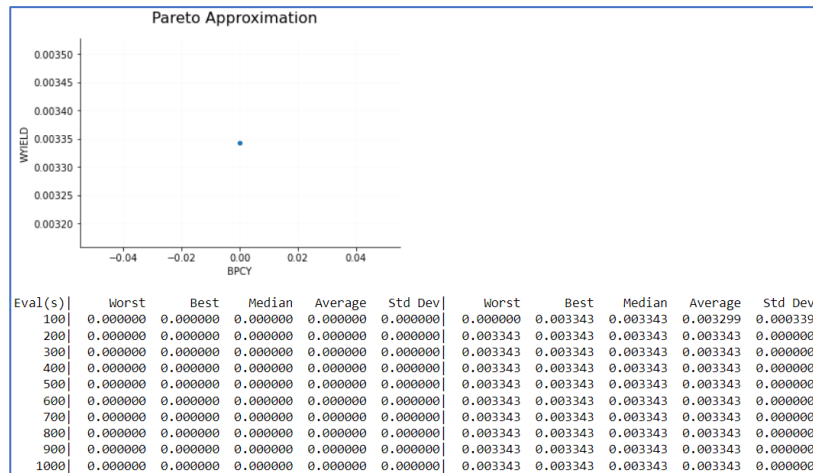
Anexo 5

```
from mewpy.problems import GKOPProblem

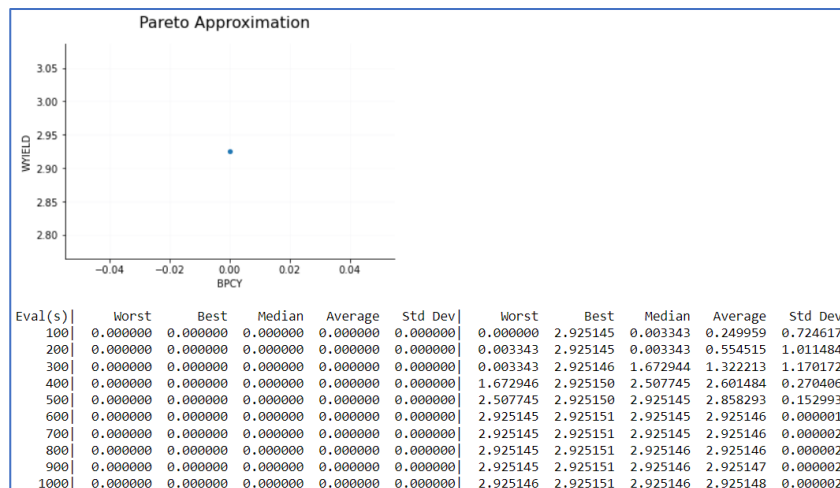
problem = GKOPProblem(model, [])
c = problem.solution_to_constraints({'s0001':0})
result = simul.simulate(method="pFBA", constraints = c)
result.fluxes["BIOMASS_Ec_iML1515_core_75p37M"]

0.0
```

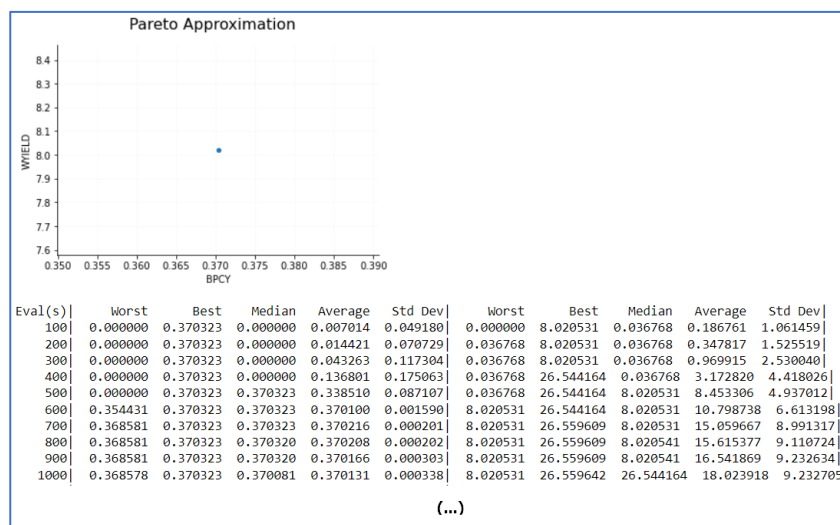
Anexo 6



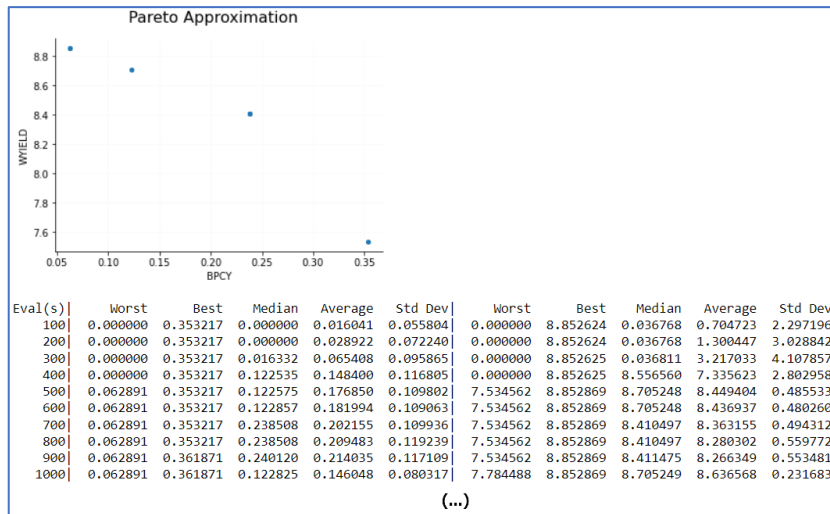
Anexo 7



Anexo 8



Anexo 9



Anexo 10

```
[0.3700803194305872, 26.544171601337318];{'ACKr': 0, 'ACALD': 0}
```

Anexo 11

```
const = {'ACKr':(0,0), 'ACALD':(0,0), 'EX_glc_D_e' : (-15.0, 100000.0), 'EX_o2_e' : (0,10000)}
result = simul.simulate(method="pFBA", constraints = const)

print(result.find("EX_lac_D_e"))
print()
print(simul.FVA(reactions=["EX_lac_D_e"], constraints = const))
print()
print()
print(result.find("BIOMASS_Ec_iML1515_core_75p37M"))
print()
print(simul.FVA(reactions=["BIOMASS_Ec_iML1515_core_75p37M"], constraints = const))

Reaction ID Flux rate
0 EX_lac_D_e 26.668039

{'EX_lac_D_e': [11.570203860940747, 27.230339294122086]}

Reaction ID Flux rate
0 BIOMASS_Ec_iML1515_core_75p37M 0.208159

{'BIOMASS_Ec_iML1515_core_75p37M': [0.18734351937956611, 0.20815946597729315]}
```

Anexo 12

Gene Reaction Rule:
b2296 or b1849 or b3115

Genes:
b3115 (tdcD)
b1849 (purT)
b2296 (ackA)

Anexo 13

Gene Reaction Rule:
b0351 or b1241

Genes:
b1241 (adhE)
b0351 (mhpF)

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