## Evolutionary results - enzyme efficiency

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## Fitness landscapes accounting for possible toxic effects

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1) First model: linear cost, with f = \Phi - \tau[S] where \tau is the linear toxicity rate.
The following set of parameters was used:
###5bis.Plotting itness with a linear penalization
###Yields problems because of the linearity of the cost
N_reso=25
#Defining parameters
Vm_set=c(10^-6)\#,10^-3) #unit: M.s^-1, with Vm=5.10^-5pmol.s^-1.cell^-1 and Vcell=50 micrometers^3
KT_set=c(5*10^-5)#,10^-3)#
log10kf_var<-c(0,10,2) #forward preferred to 1 as forward is already an emerging constant from diffusion
log10kcat_var<-c(-4,6,2) #catalytic preferred to -1 for same reasons
kf<-10^seq(log10kf_var[1],log10kf_var[2],length.out=N_reso)
kcat<-10^seq(log10kcat_var[1],log10kcat_var[2],length.out=N_reso)</pre>
E tot conc=10^-3##c(10^-7,10^-4) Both values need to be studied as what matters most is kf[Etot]
#Best possible values for the first reaction
kf1 set < -c(10^2, 10^10)
kcat1_set < -c(10^-2, 10^-6)
#kr1_set<-c(10^5,10^6,10^3,10^4)
#kinh1_set<-c(10^9,10^10,10^7,10^8)
#Ki<-10^-6
eta_d_set<-c(10^-6)#degradation rate #Sensitivity study to be done relatively to Vm
delta_t_set<-c(10^-6,10^-3)#Toxicity penality
kf2_set<-kf
kcat2_set<-kcat
kr2_set<-c(10^3)##Testing with 2 values, but only one should be used later on in the paper
E1_conc<-E_tot_conc
E2_conc<-E_tot_conc
2)Second model: sigmoid cost (Rausher, 2007) representing the fact that increasing concentrations can
interfere with other reactions and significantly alter fitness. It is computed through f = \Phi(1 - \frac{[S]}{[S] + T_{1/2}}),
where T_{1/2} represents the half toxicity constant (substrate concentration for which the fitness given by a flux
is half the value of the flux, because substrate disrupts another reaction dividing its yield by this factor.
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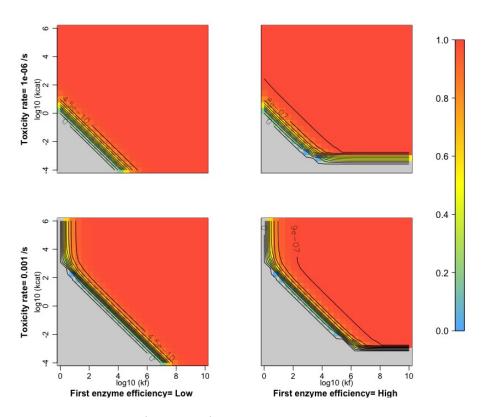


Figure 1: Flux (and fitness) values obtained with linear toxicity rate