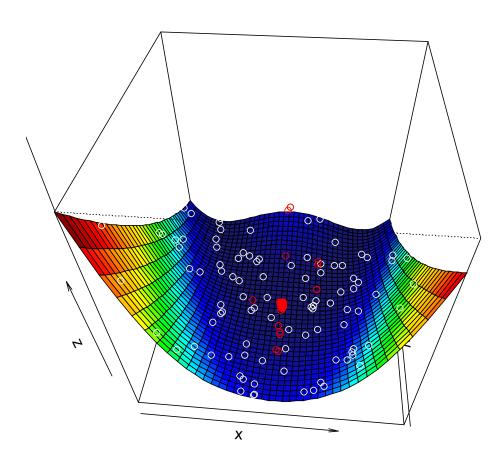
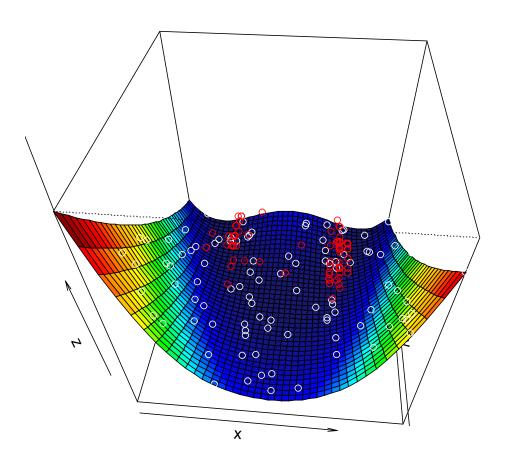
In the following examples white points represent points from initial population and red points represent chromosomes from Nrg population.

Function: RosenbrockValley; Nrg = 20

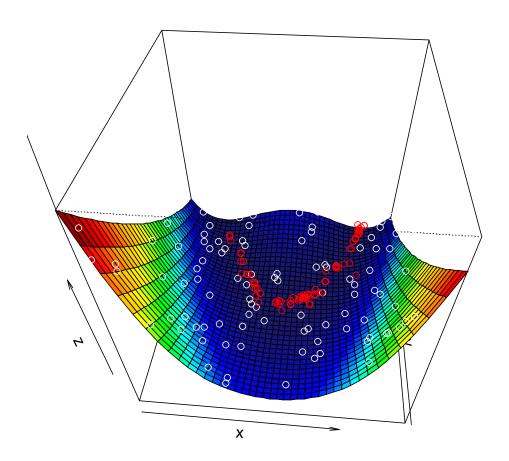
Alg = Clearing + FortuneWheelSelection

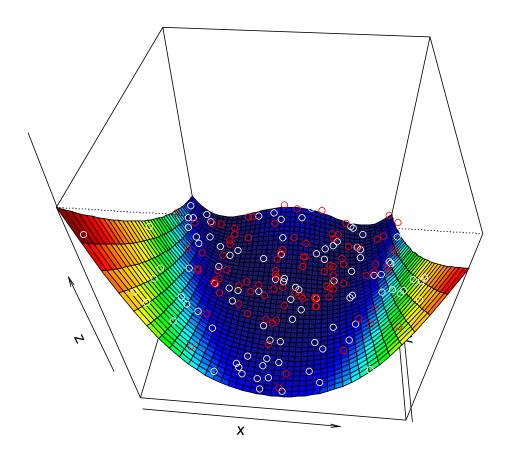


Alg = Clearing + TournamentSelection



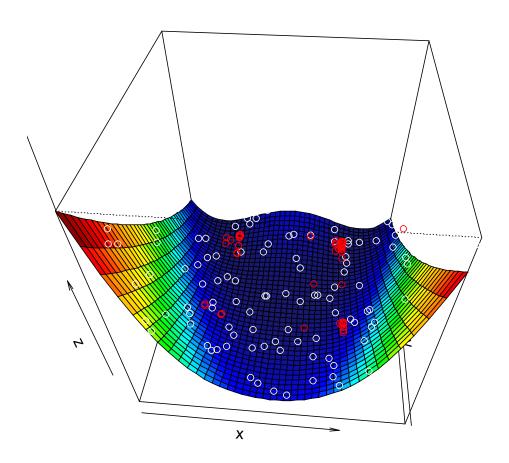
Alg = Deterministic Crowding

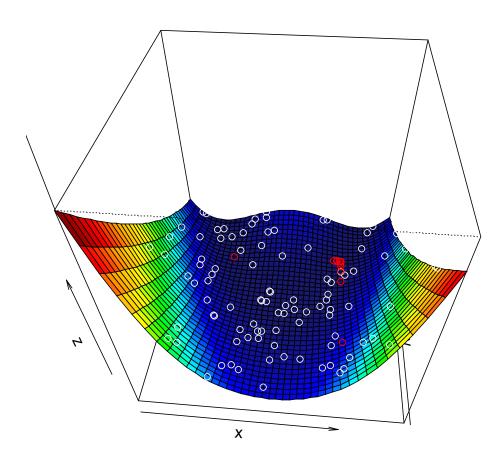


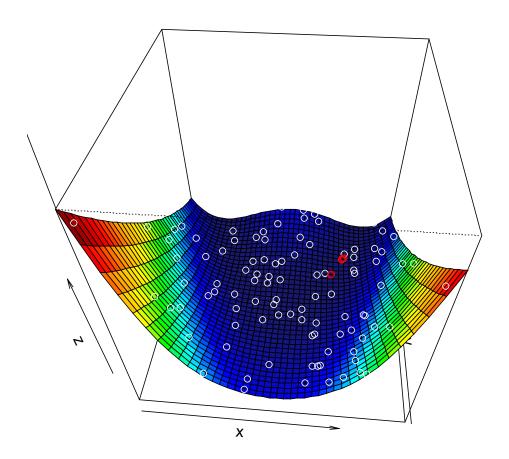


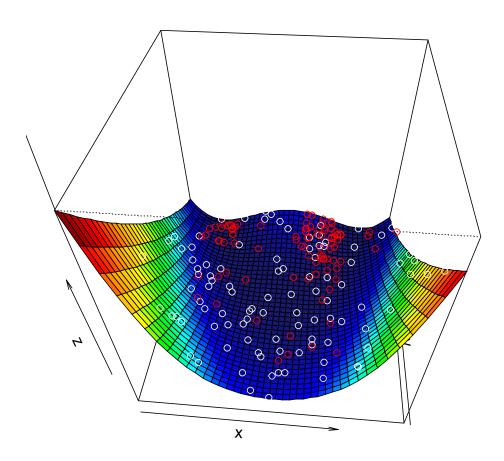
Function: RosenbrockValley; Nrg = 200

Alg = Clearing + FortuneWheelSelection





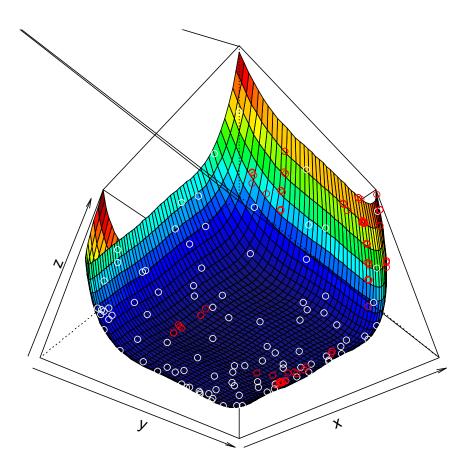




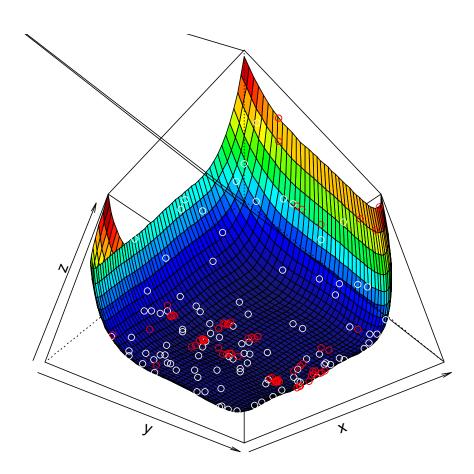
Further in order to reduce number of graphs I'll register for each function and algorithm different number of generations accordingly to the speed of algorithm convergence.

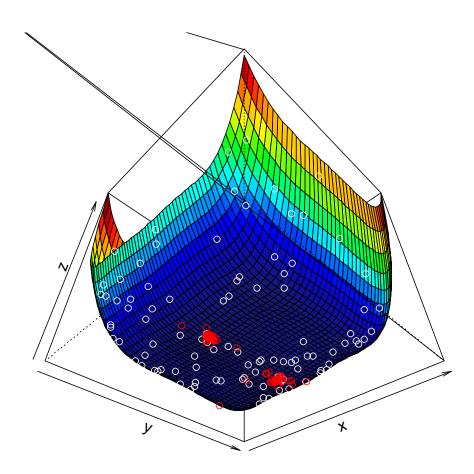
Function: Six-hump camel back;

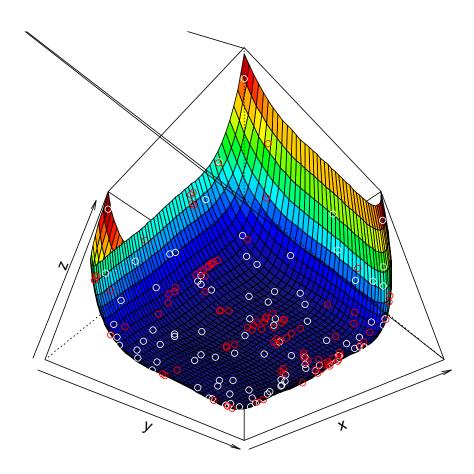
Alg = Clearing + FortuneWheel (nrg=100)



Alg = Clearing + Tournament (nrg=50)

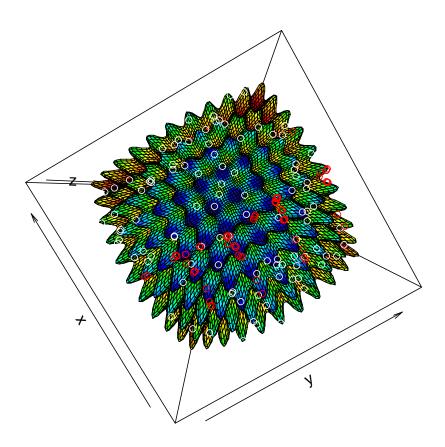


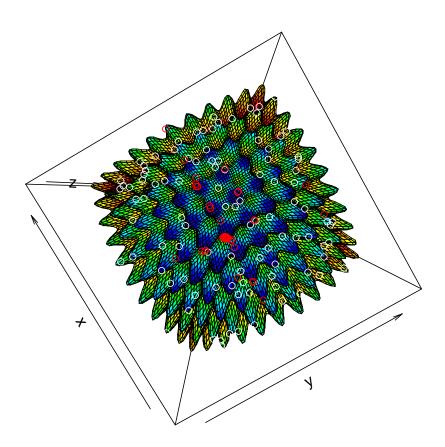


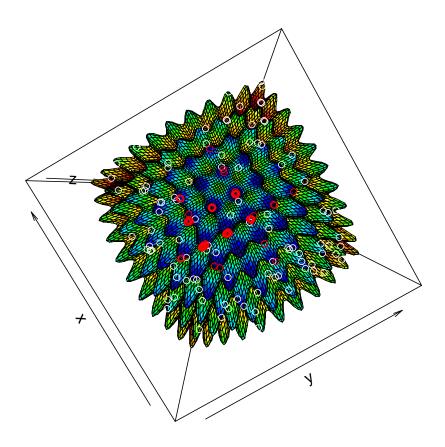


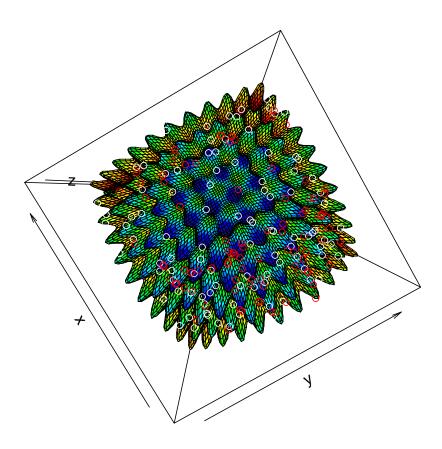
Function Rastrigin:

Alg= Clearing + FortuneWheel (nrg=200)



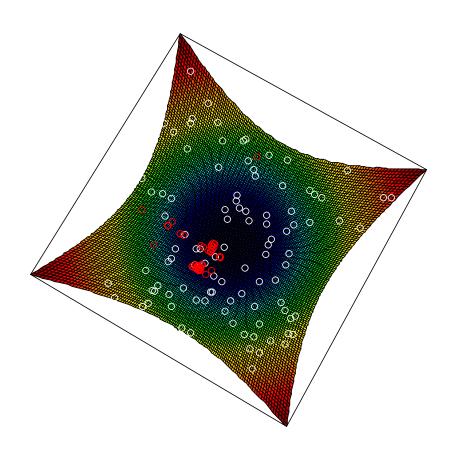


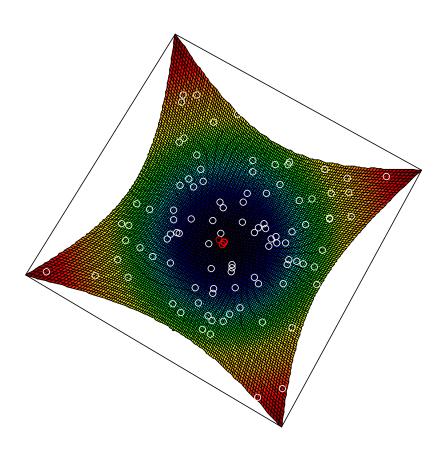


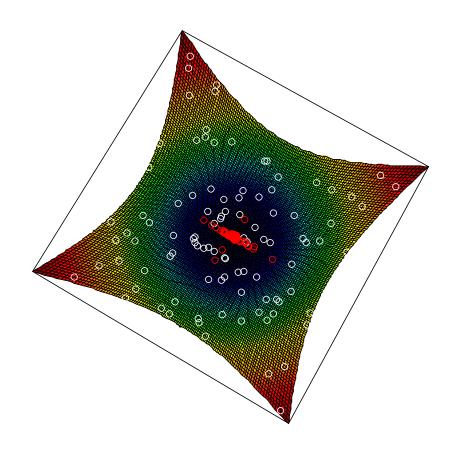


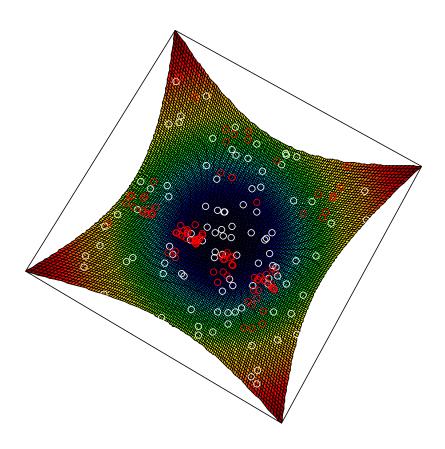
Function: Griewang

Alg = Clearing + FortuneWheel (nrg=200)









There is no need to draw histograms with points on X and Y axis as we can see clearly from images how points are distributed in space.