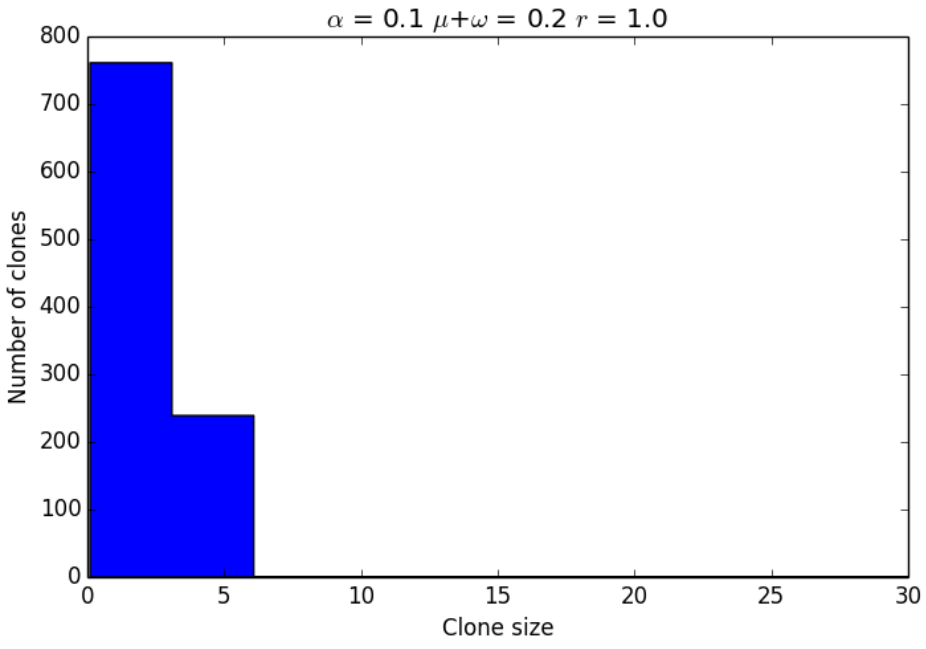
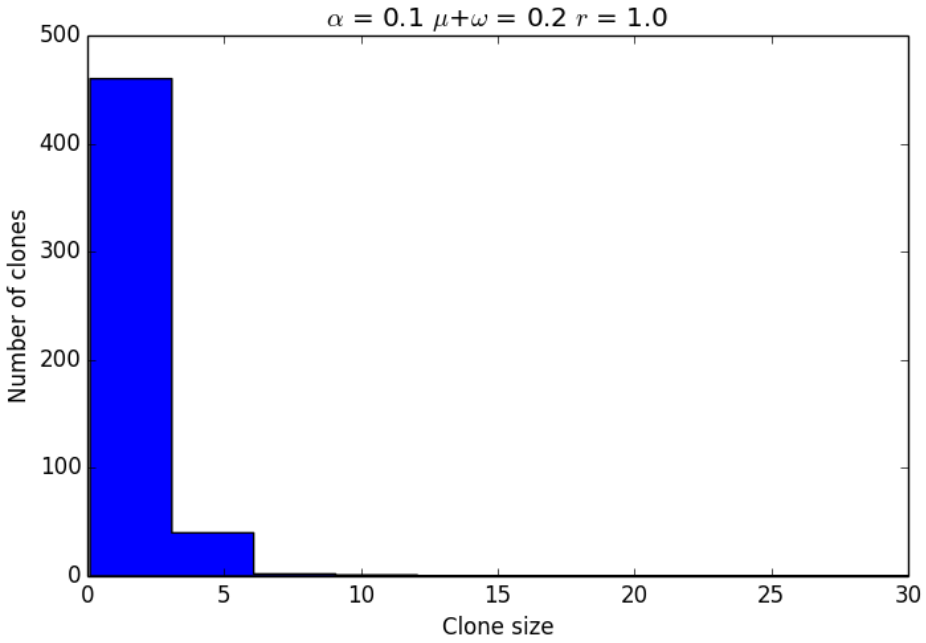
Actually bone marrow capacity is 1000, rather than 500.

Clone-size distributions

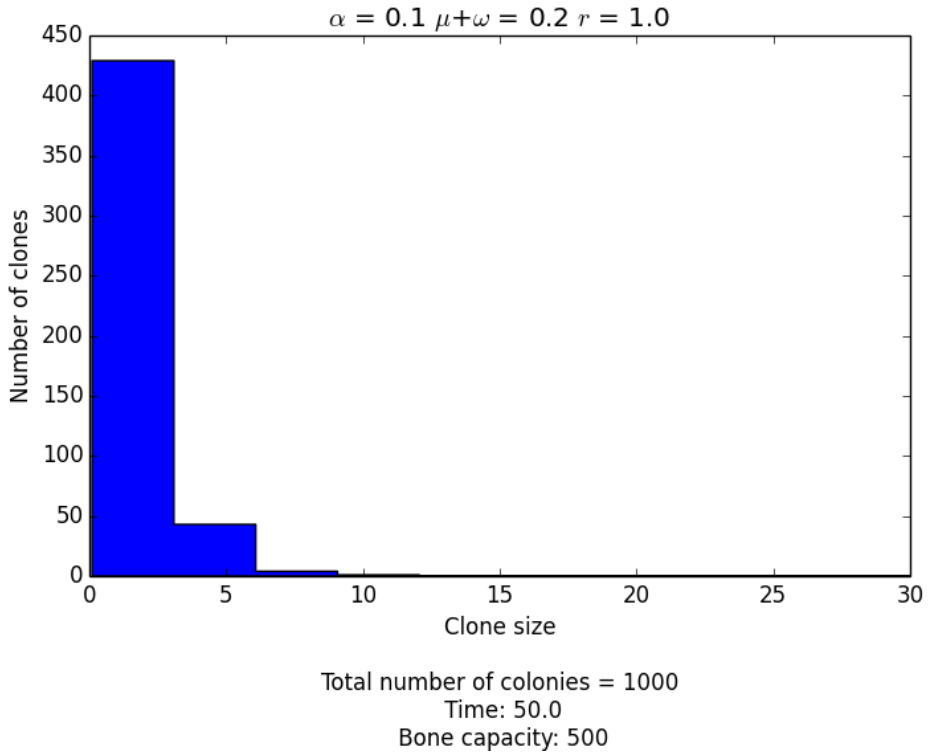
1) First parameters set



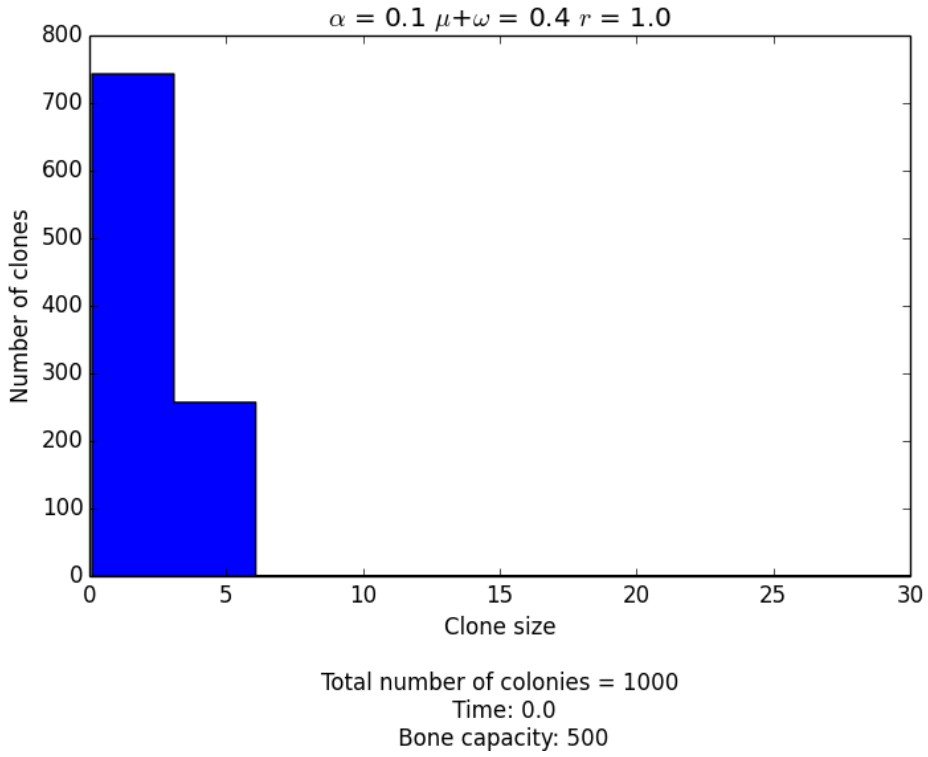
Total number of colonies = 1000
Time: 0.0
Bone capacity: 500

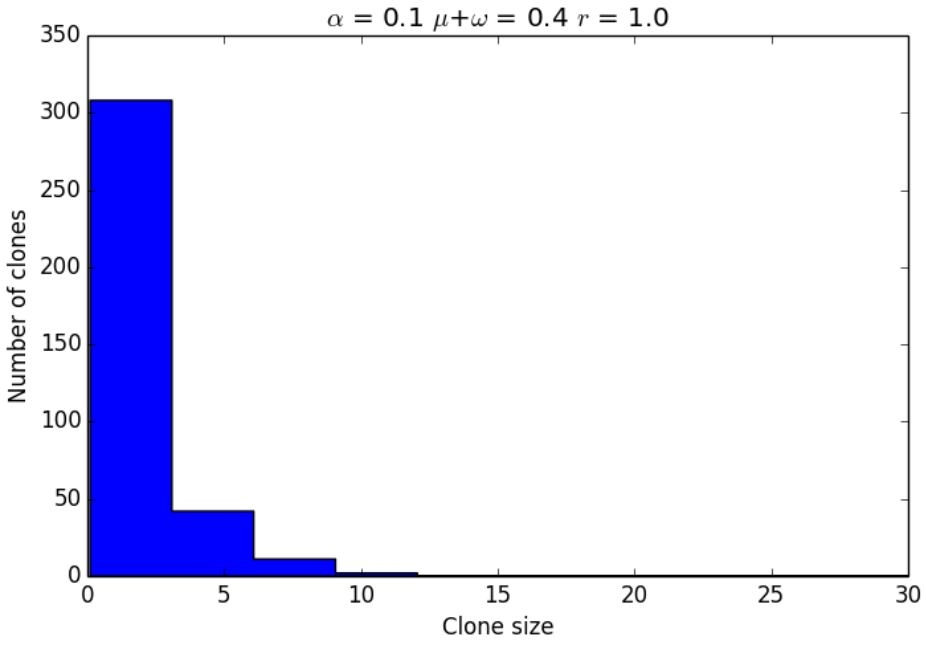


Total number of colonies = 1000
Time: 24.4
Bone capacity: 500

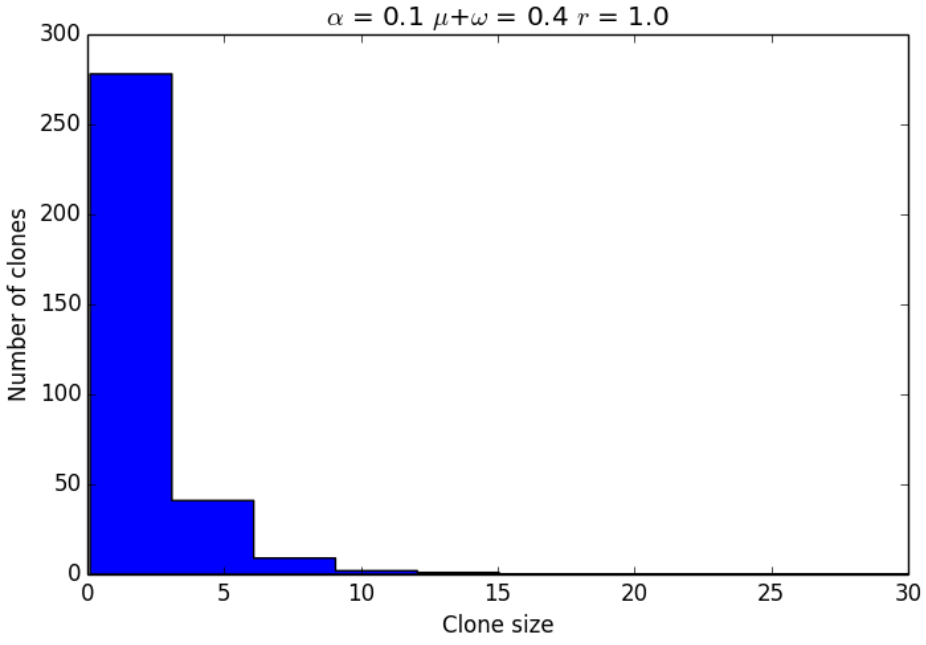


2) Second parameters set



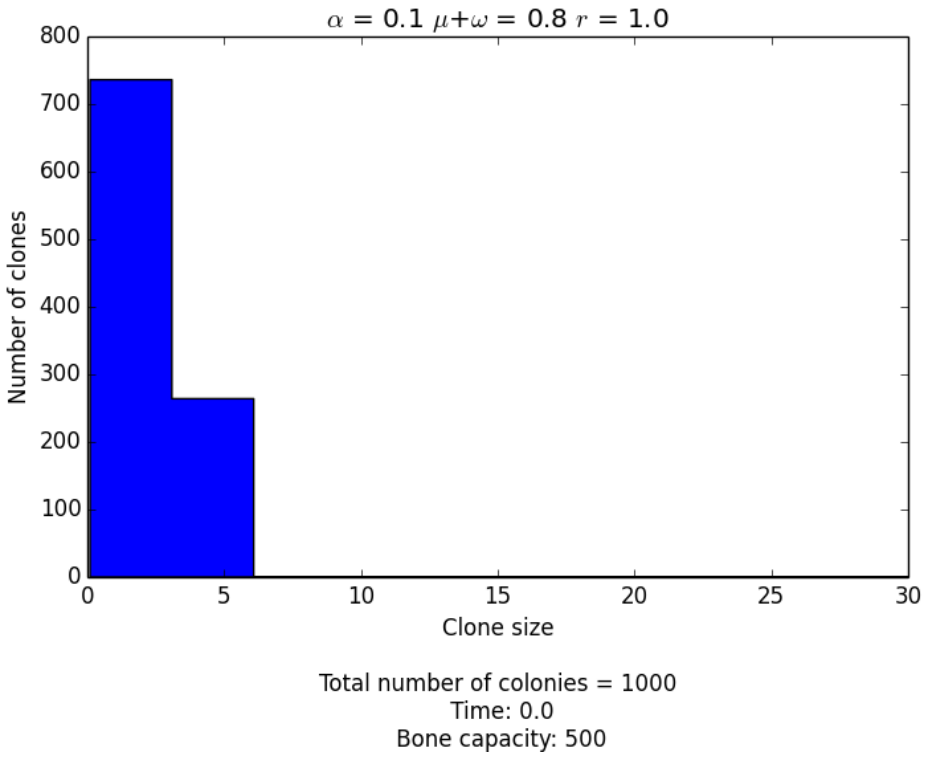


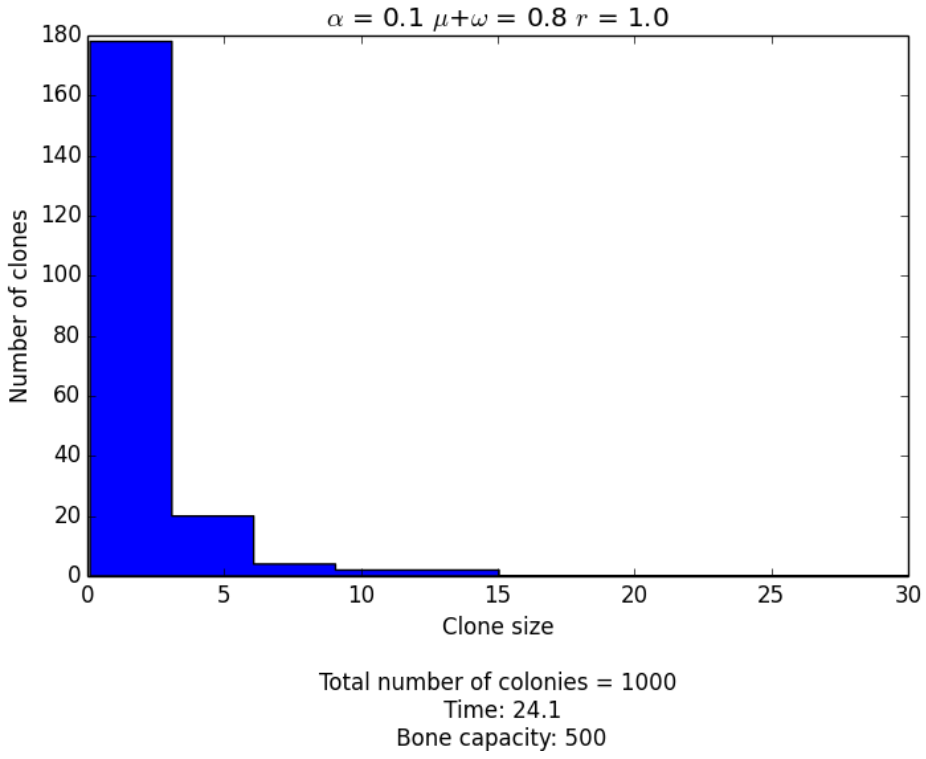
Total number of colonies = 1000
Time: 24.5
Bone capacity: 500

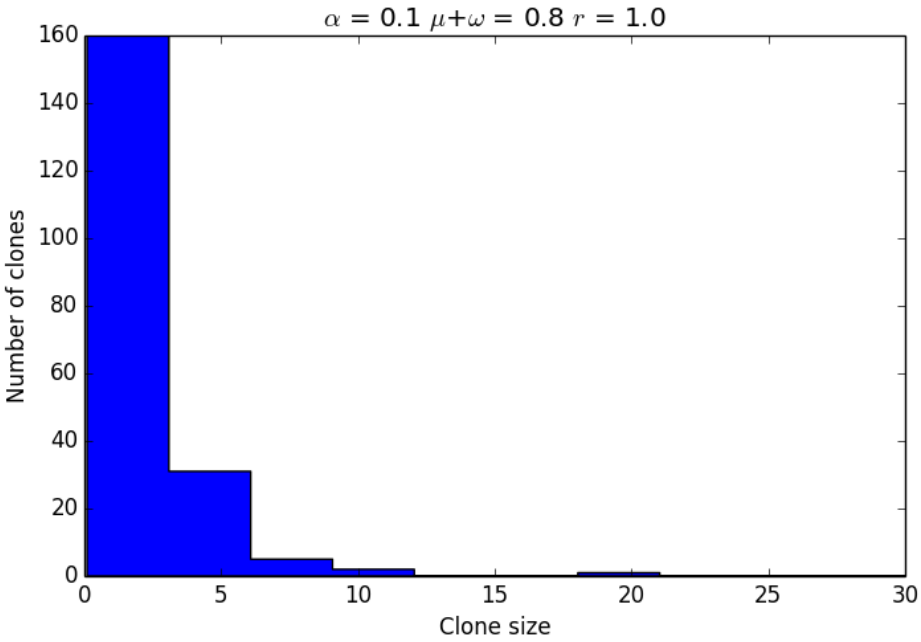


Total number of colonies = 1000
Time: 50.0
Bone capacity: 500

3) Third parameters set



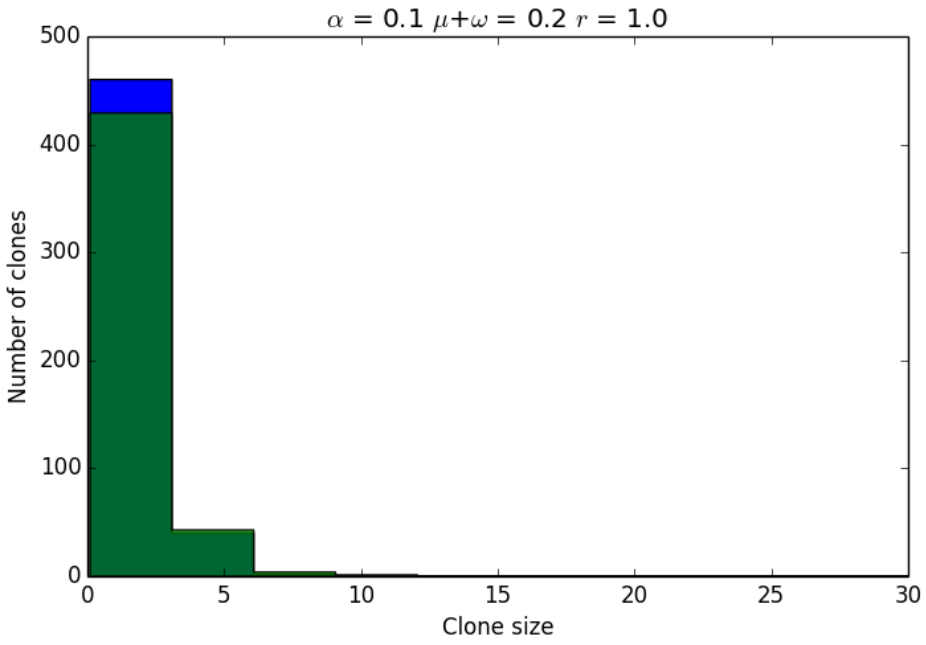




Total number of colonies = 1000
Time: 50.0
Bone capacity: 500

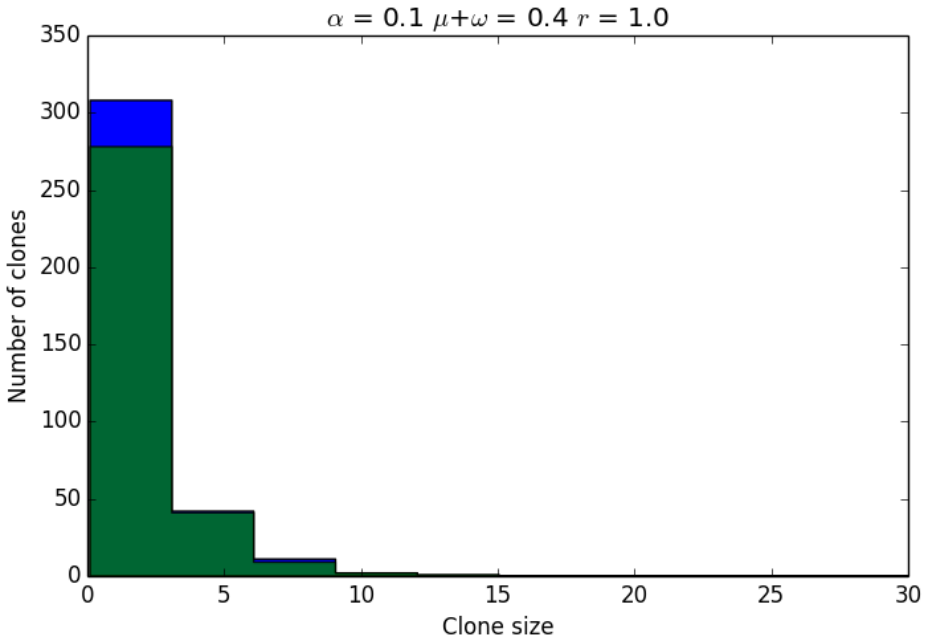
Join middle and end time clone-size distributions

1) First parameters set



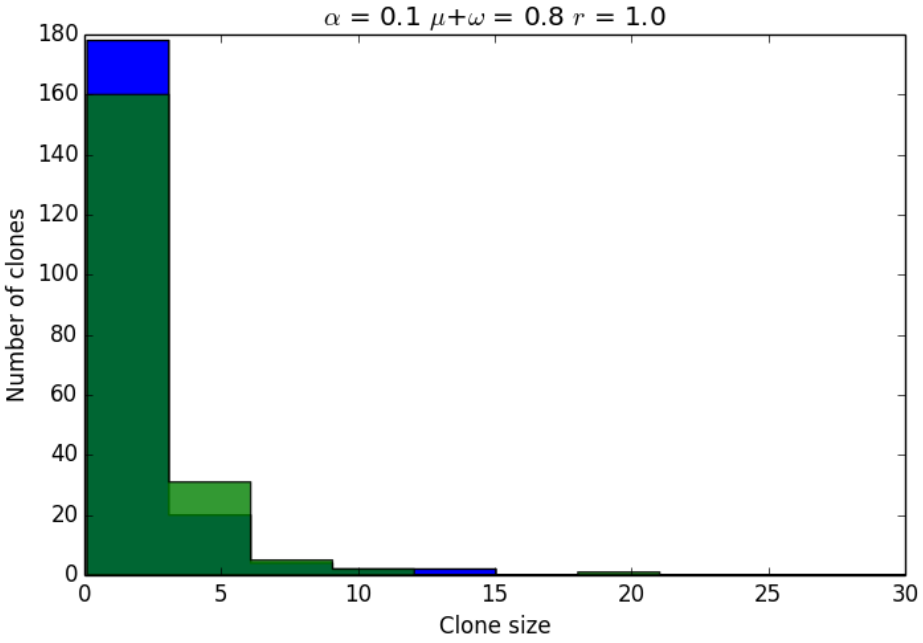
Total number of colonies = 1000
Time: 24.4
Bone capacity: 500

2) Second parameters set



Total number of colonies = 1000
Time: 24.5
Bone capacity: 500

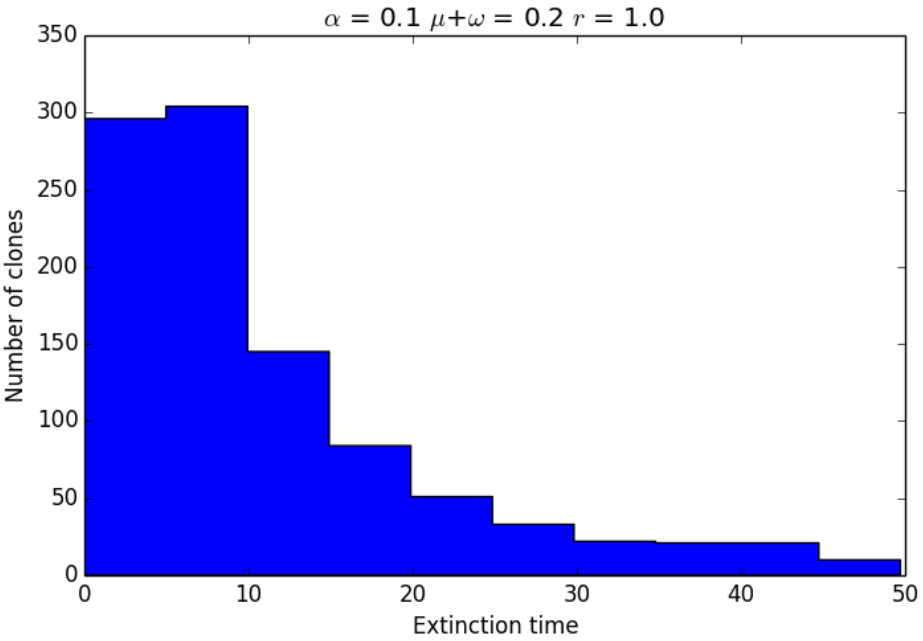
3) Third parameters set



Total number of colonies = 1000
Time: 24.1
Bone capacity: 500

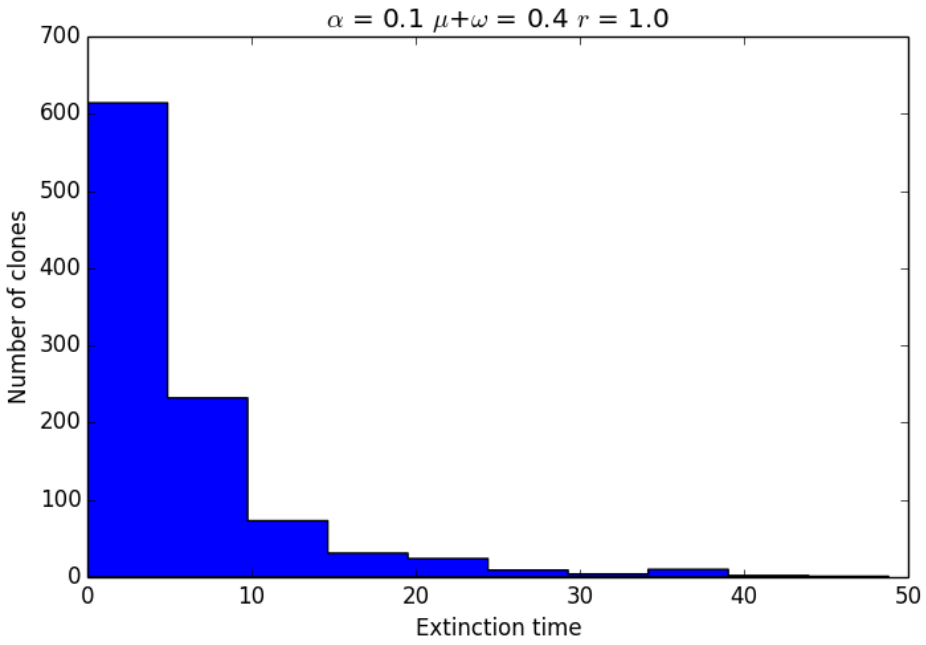
Extinction time

1) First parameters set



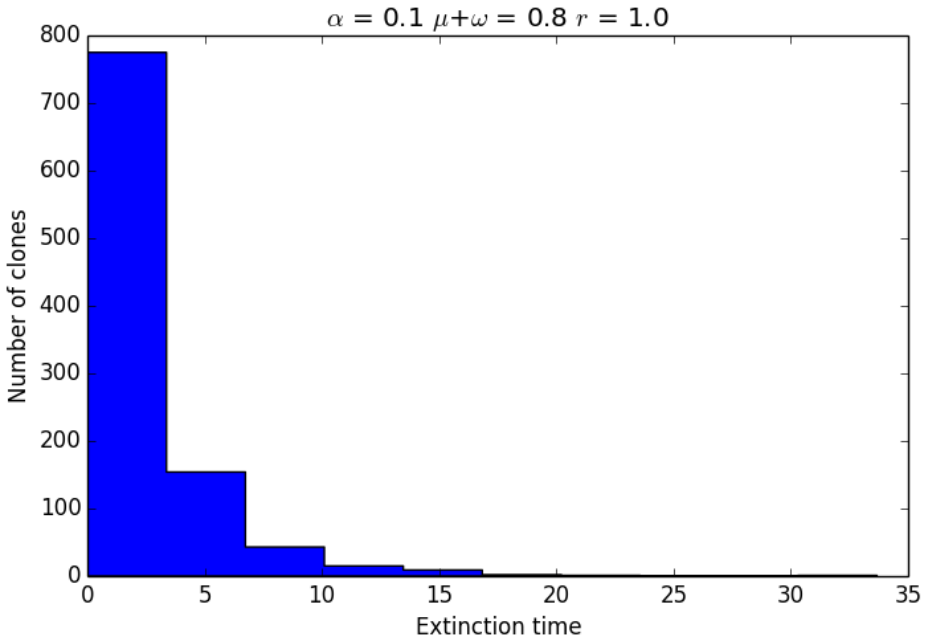
Total number of colonies = 1000
Time: 24.4
Bone capacity: 500

2) Second parameters set



Total number of colonies = 1000
Time: 24.5
Bone capacity: 500

3) Third parameters set



Total number of colonies = 1000
Time: 24.1
Bone capacity: 500