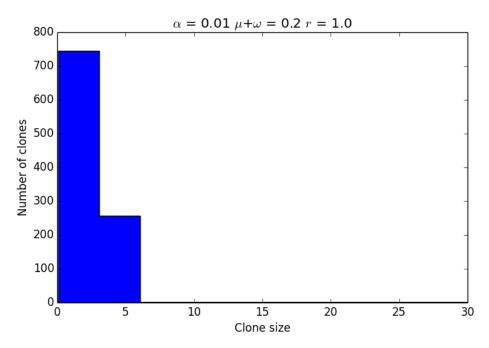
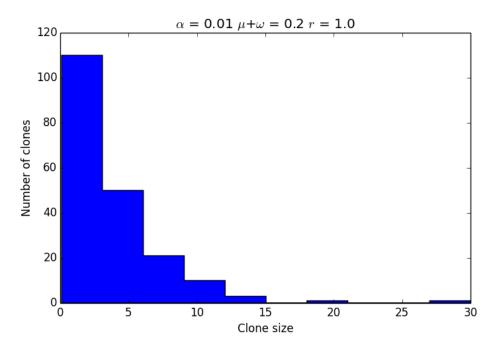
Clone-size distributions

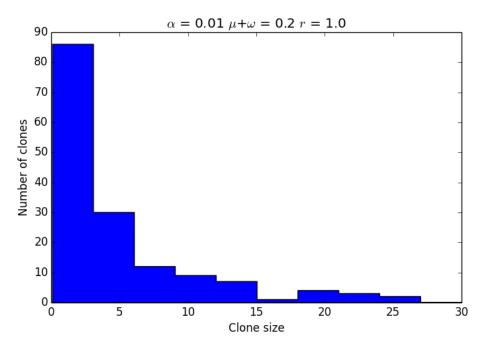
1) First parameters set



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

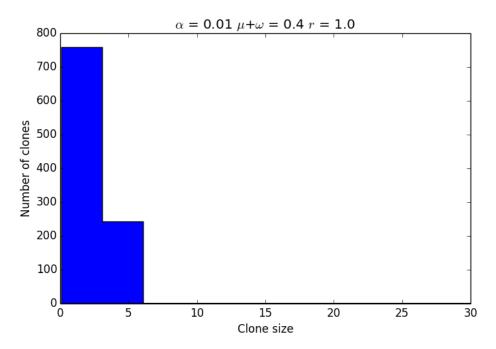


Total number of colonies = 1000 Time: 24.7 Bone capacity: 500

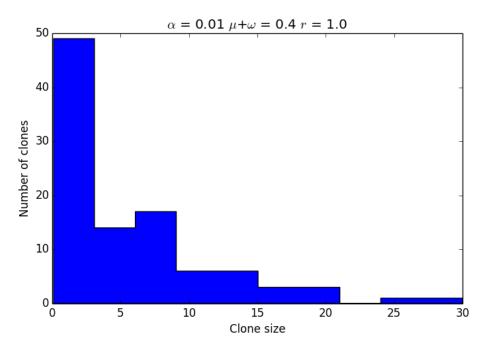


Total number of colonies = 1000 Time: 50.0 Bone capacity: 500

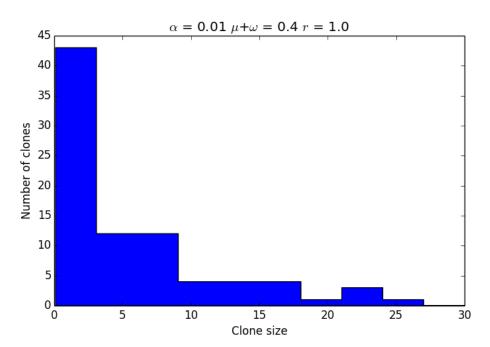
2) Second parameters set



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

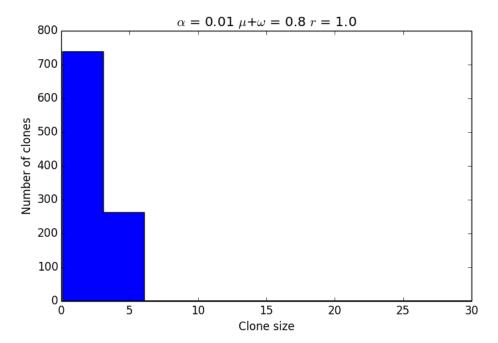


Total number of colonies = 1000 Time: 24.5 Bone capacity: 500

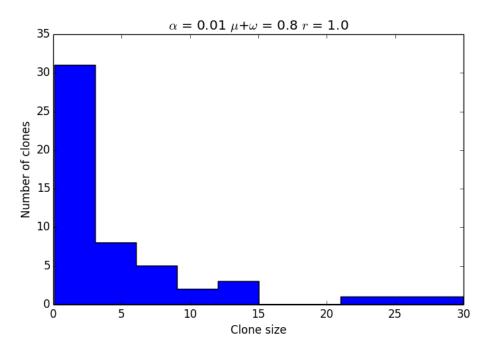


Total number of colonies = 1000 Time: 50.0 Bone capacity: 500

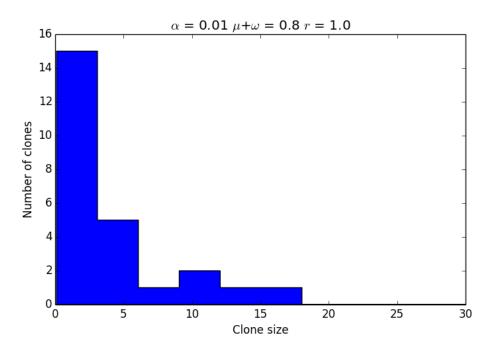
3) Third parameters set



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



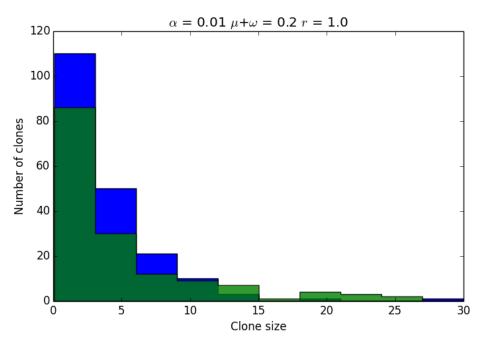
Total number of colonies = 1000 Time: 22.2 Bone capacity: 500



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

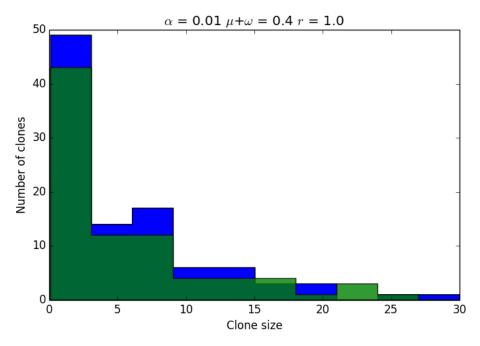
Join middle and end time clone-size distributions

1) First parameters set



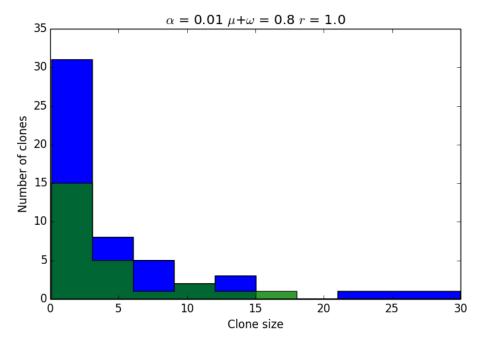
Total number of colonies = 1000 Time: 24.7 Bone capacity: 500

2) Second parameters set



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

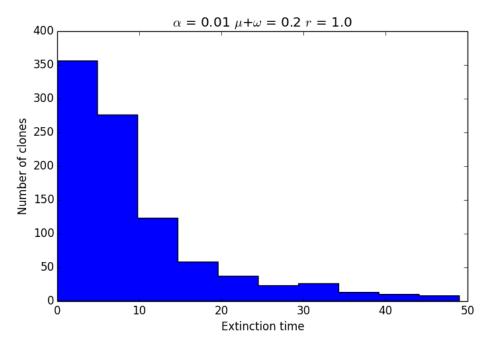
3) Third parameters set



Total number of colonies = 1000 Time: 22.2 Bone capacity: 500

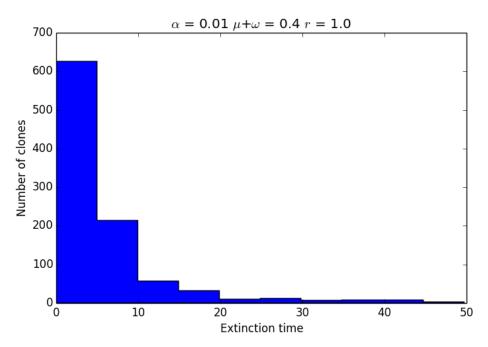
Extinction time

1) First parameters set



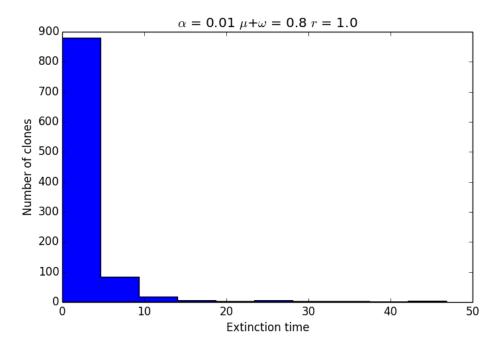
Total number of colonies = 1000 Time: 24.7 Bone capacity: 500

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: 22.2 Bone capacity: 500