

Scenario 1

Inference validation

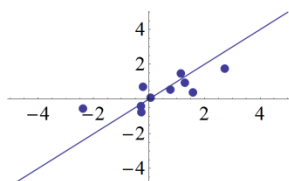
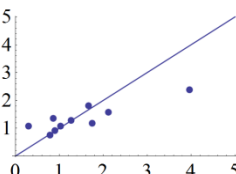
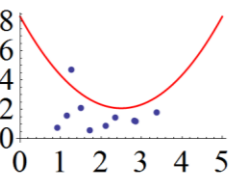
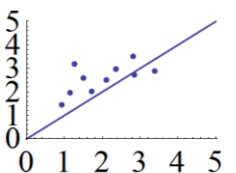
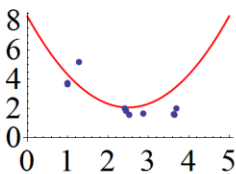
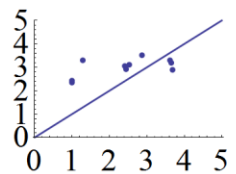
Model A:

sympatric speciation rate is diversity-dependent

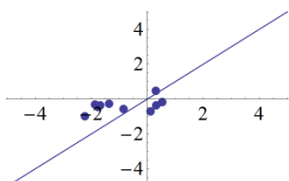
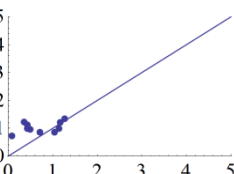
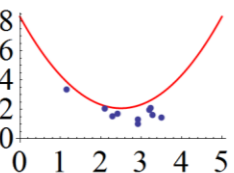
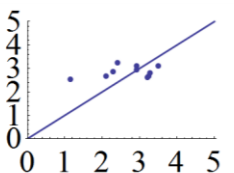
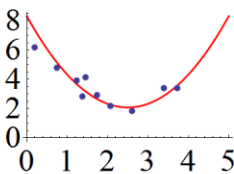
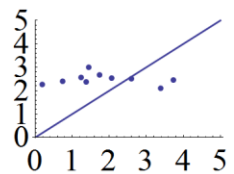
- NO PARAMETER HAS BEEN FIXED
- Time has been fixed to $T=8$ for simulating and estimating parameters
- 10 simulated data have been generated
- Sample size per generation = 400
- Generations = 5
- $q=0.5$
- Maximum number of lineages = 150

Regions

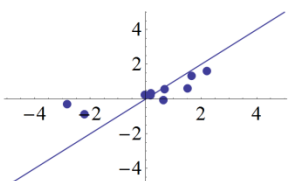
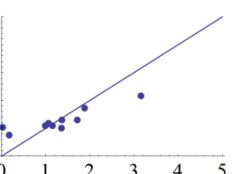
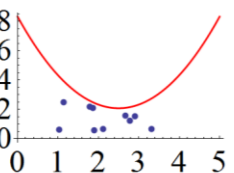
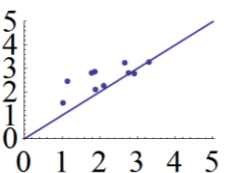
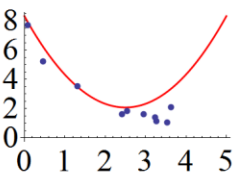
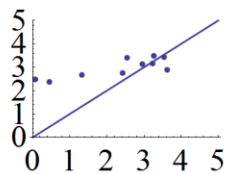
West



North



East



Sympatric
speciation (s)

Known
value
vs mean

Mean
Squared
error

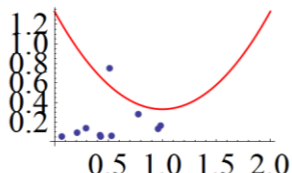
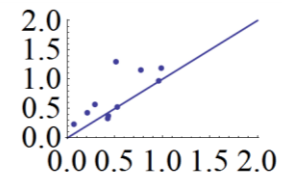
Known
value
vs mean

Mean
Squared
error

Ratio s:e

Net
diversification
s-e

Colonization



Allopatric
speciation

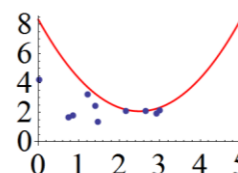
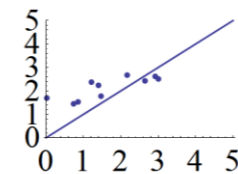


Figure SA1

Distribution of correlation between sympatric speciation and extinction (S..._E...)

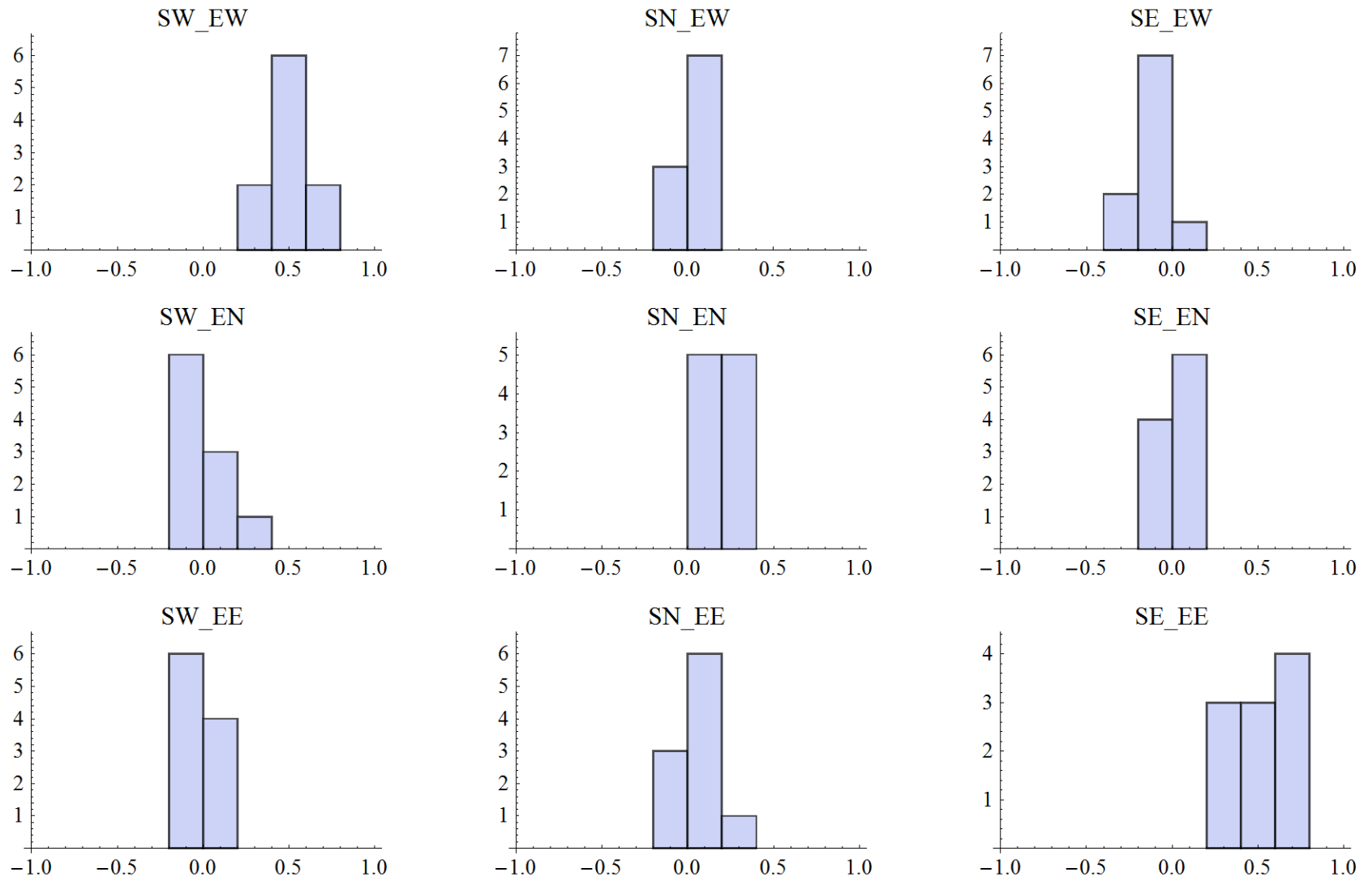


Figure SA2

Distribution of correlation between sympatric speciation (S..._S...), and between sympatric speciation and allopatric speciation (S..._A), and between extinction (E..._E...)

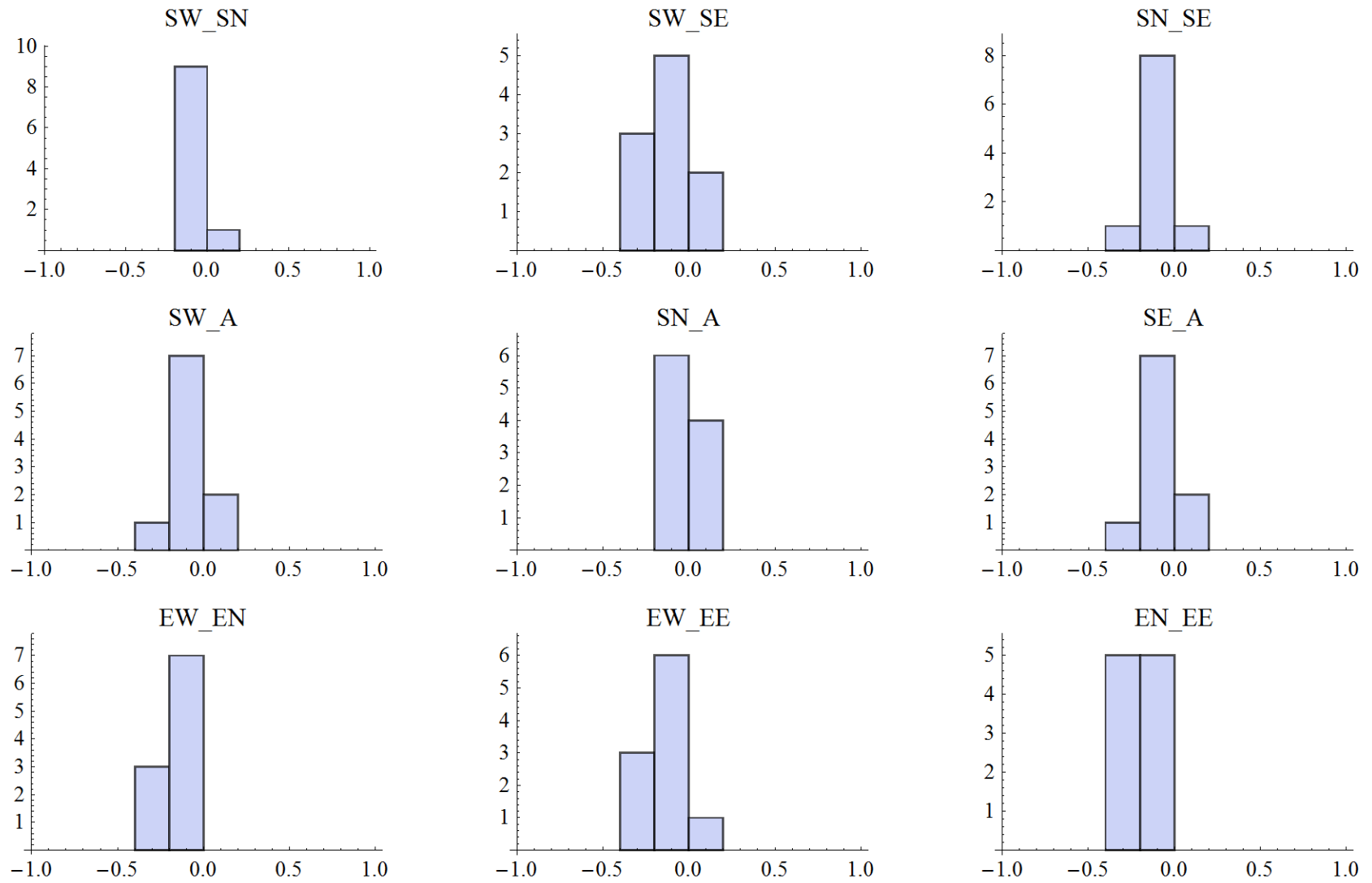


Figure SA3

Distribution of correlation between sympatric speciation and colonization (C_S...),
and between colonization and extinction (C_E...)

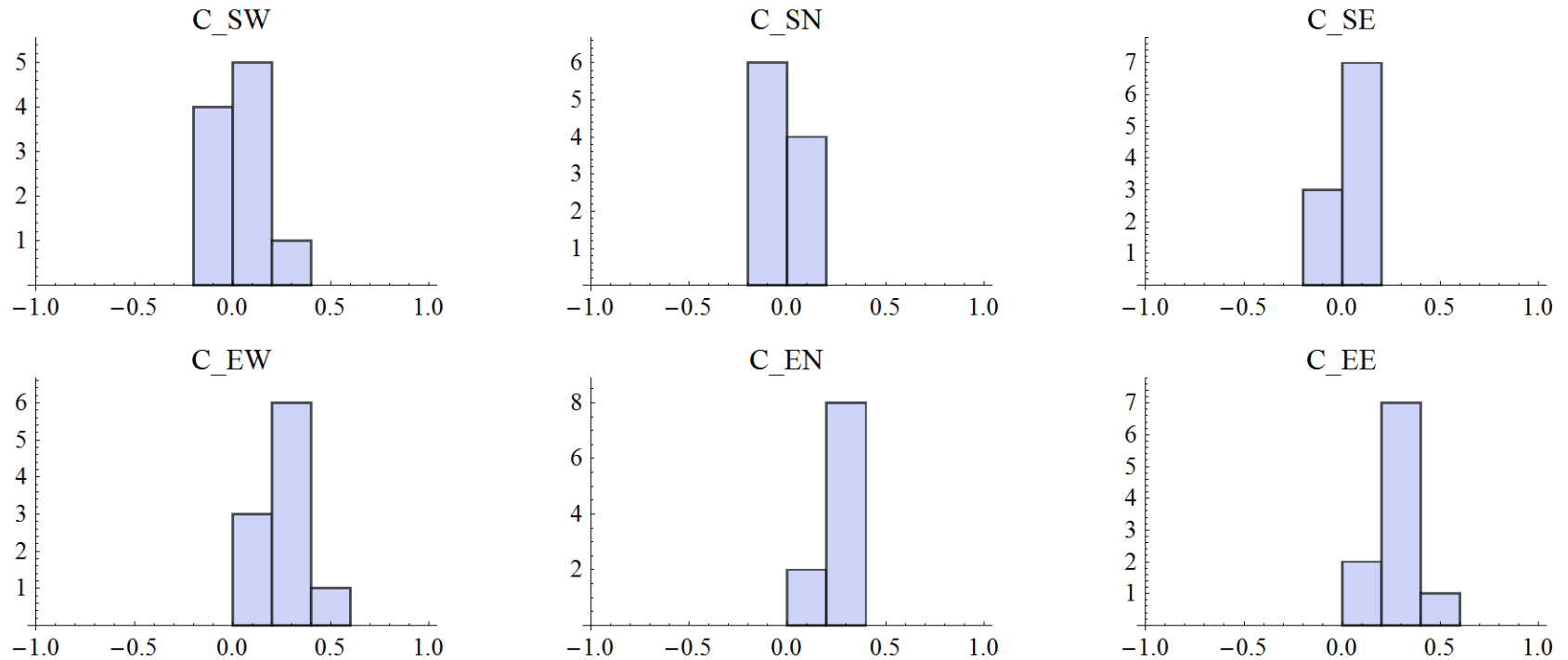


Figure SA4

Distribution of correlation between allopatric speciation and extinction (A_E...),
and between allopatric speciation and colonization (C_E...)

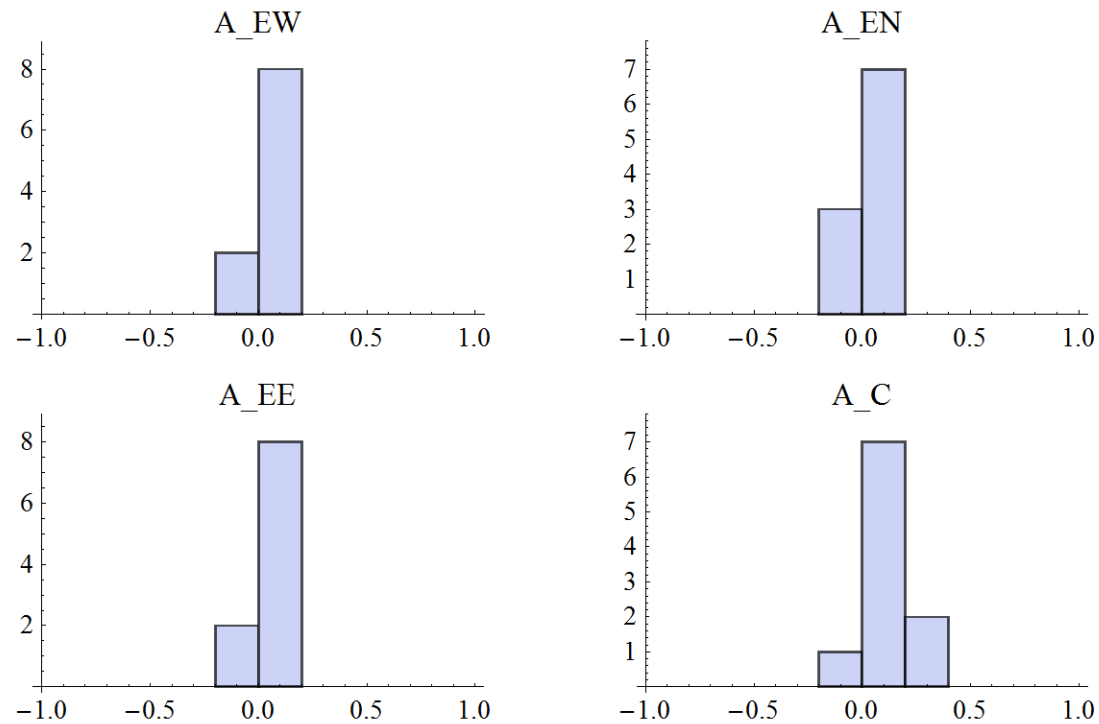


Figure SA5

Scenario 1

Estimates from Empirical data

Model A:

sympatric speciation rate is diversity-dependent

- NO PARAMETER HAS BEEN FIXED
- Estimating parameters for empirical data
- $T=8$
- Sample size = 400
- $q=0.5$
- Number of generation = 7
- Maximum number of lineages = 150

Generations

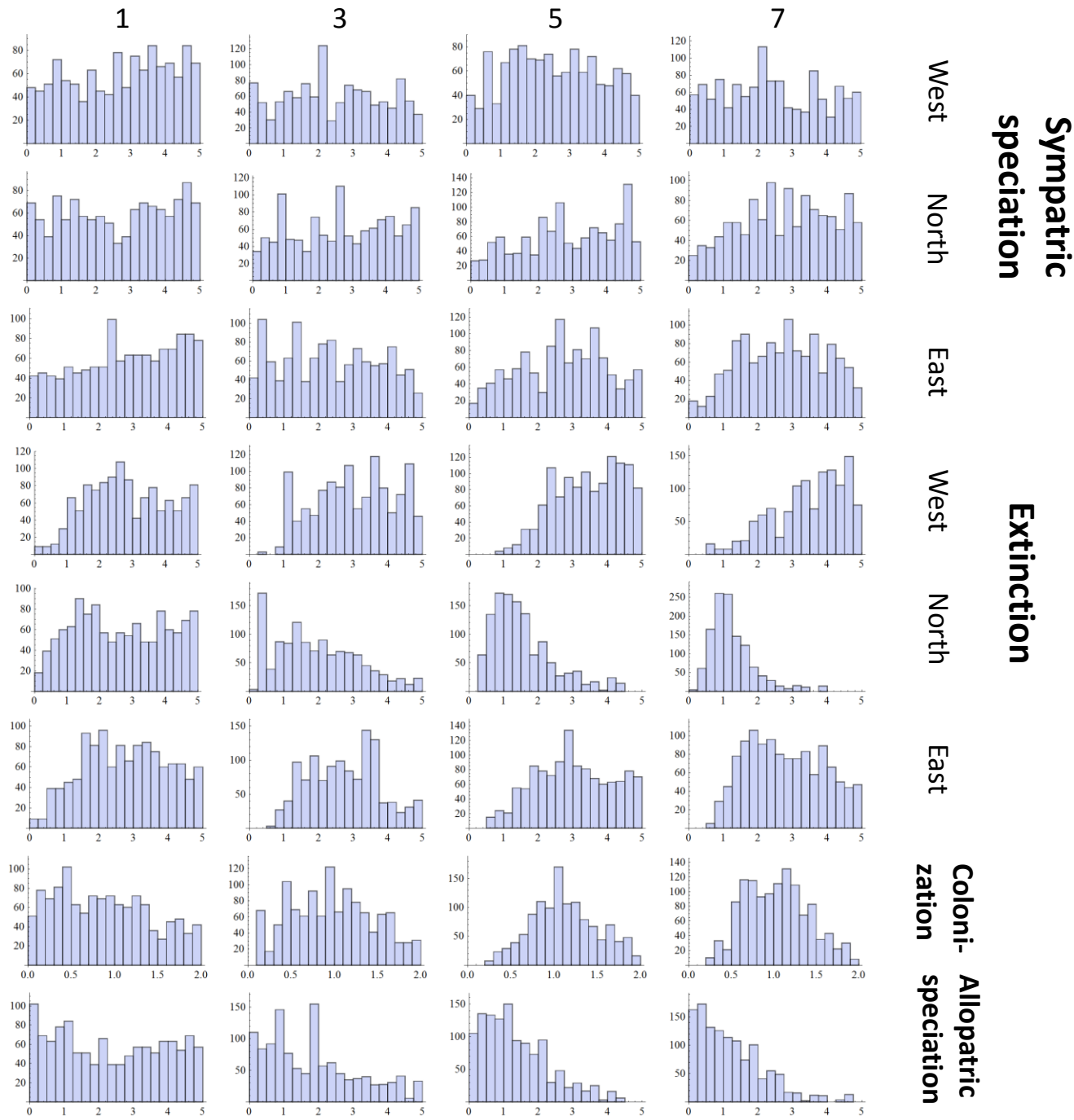


Figure SA6

Generations

1

3

5

7

West

North

East

West

North

East

Ratio
S:E

Net
diversification
S-E

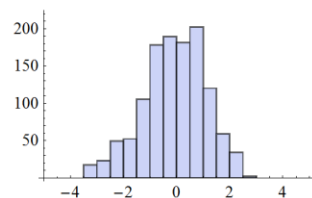
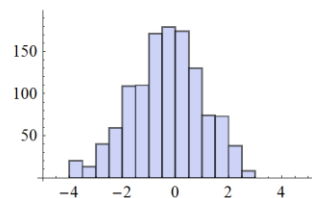
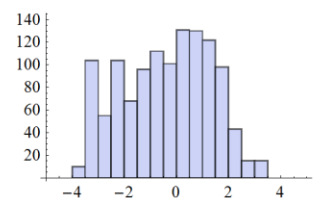
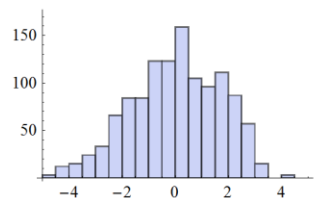
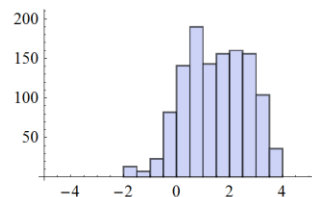
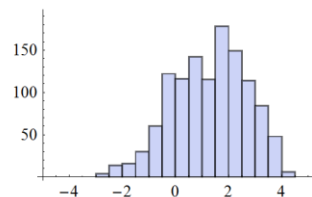
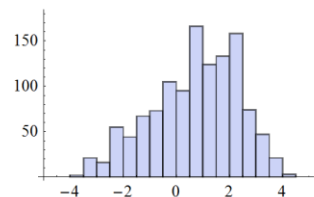
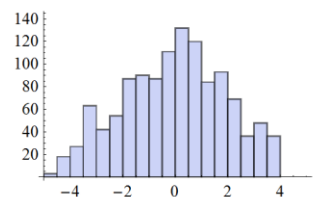
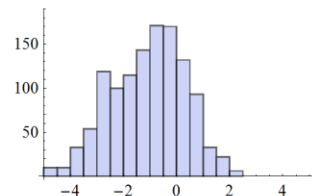
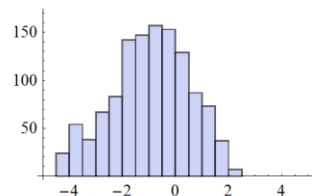
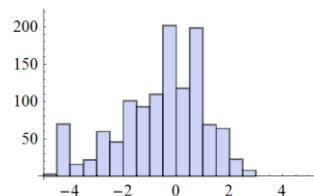
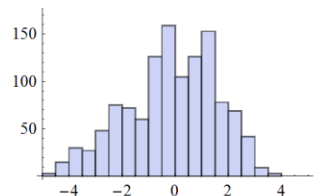
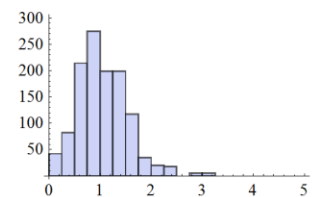
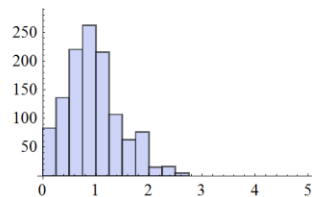
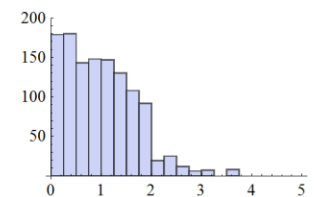
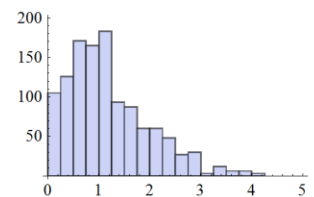
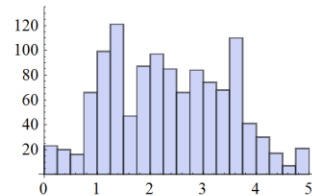
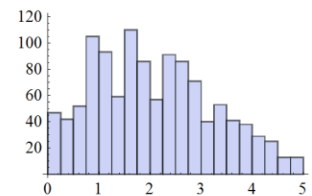
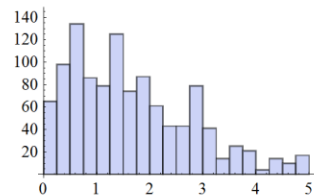
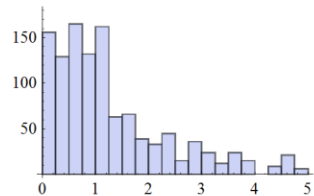
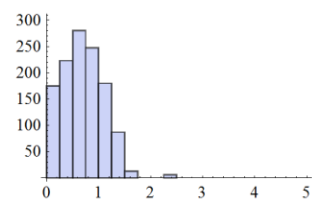
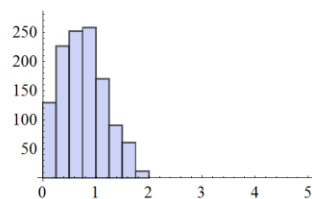
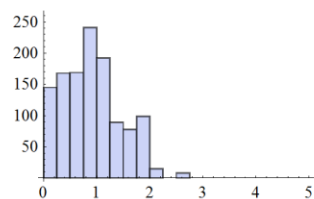
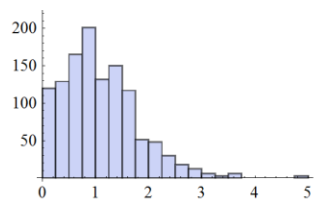


Figure SA7

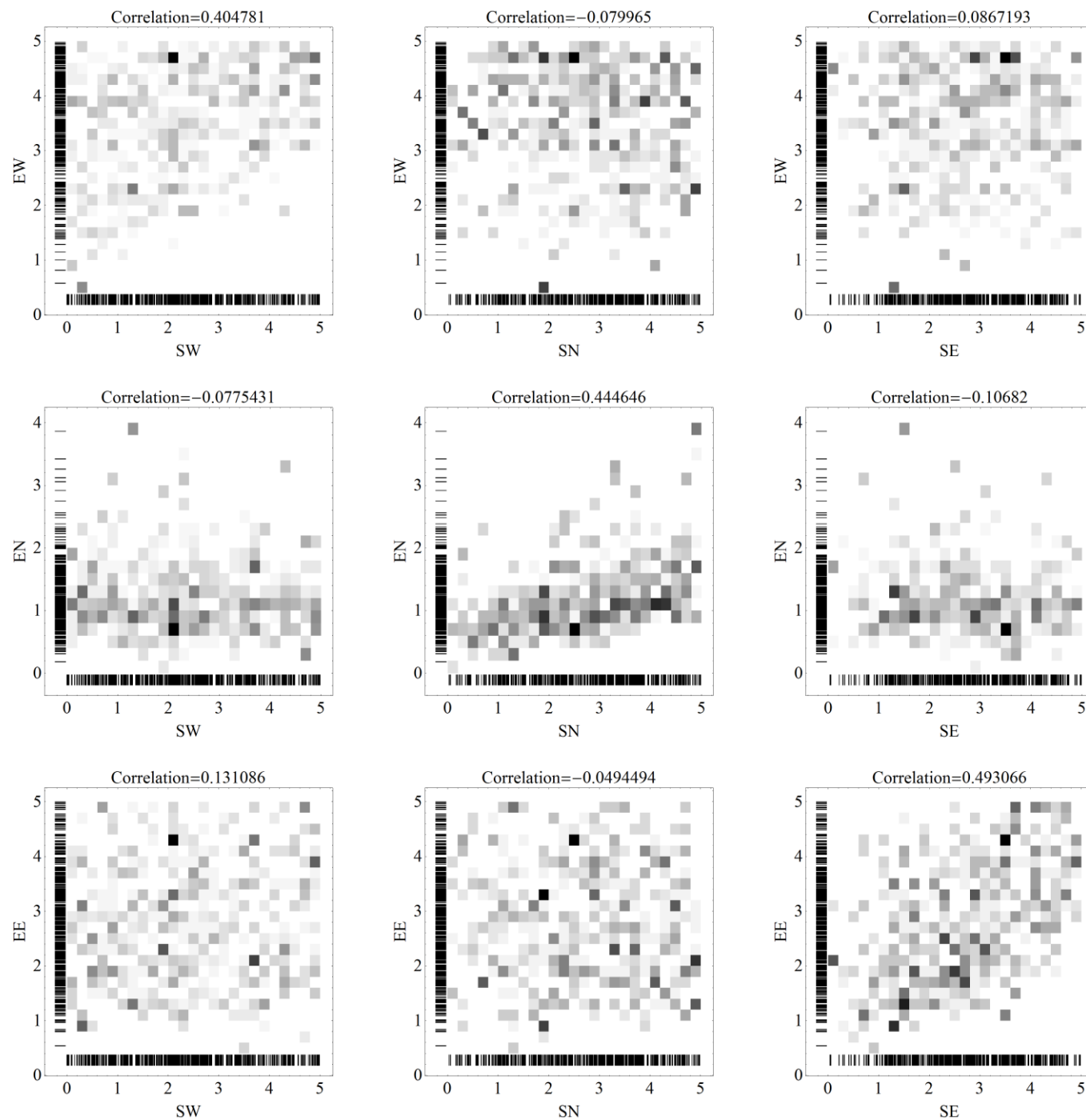


Figure SA8

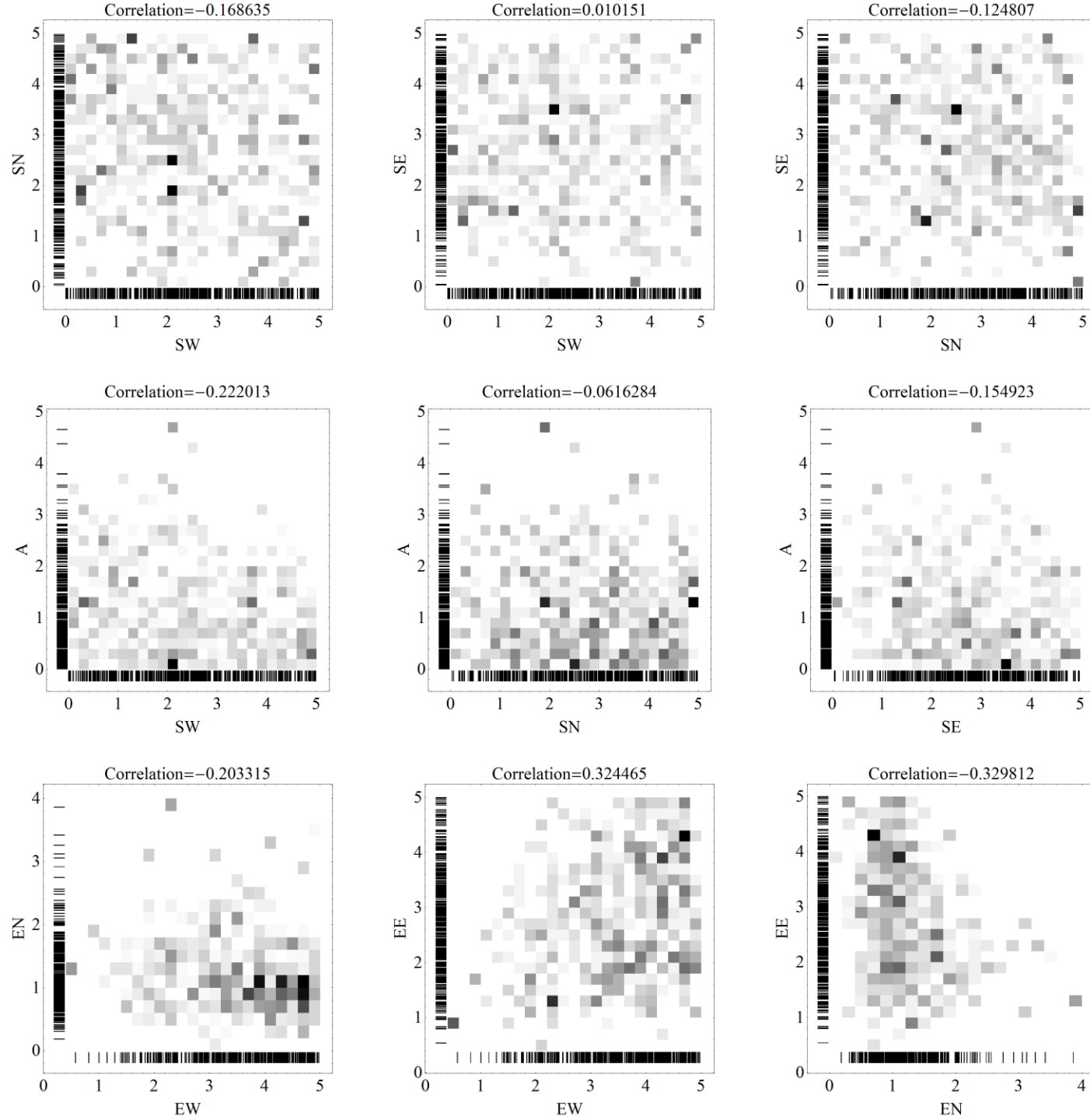


Figure SA9

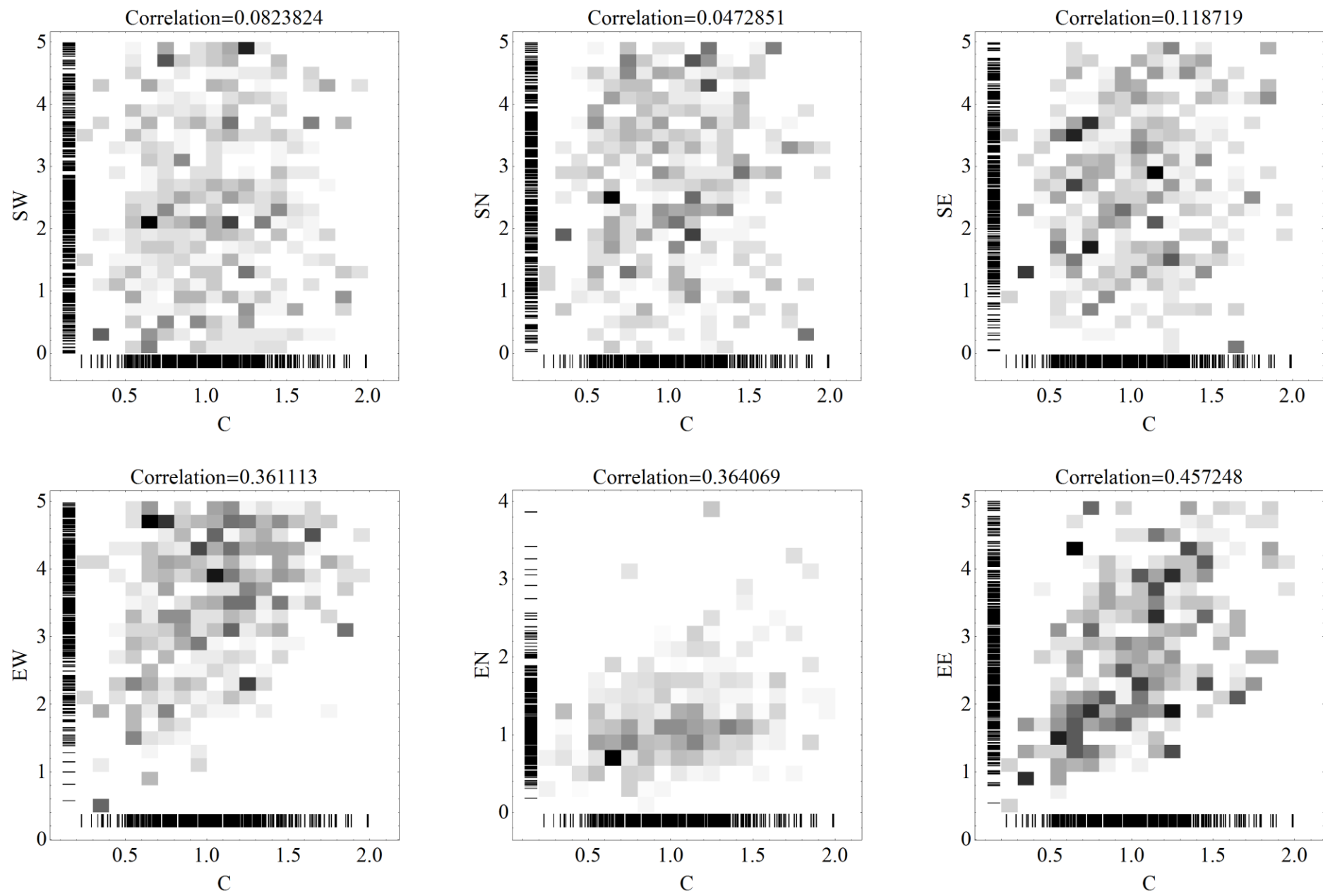


Figure SA10

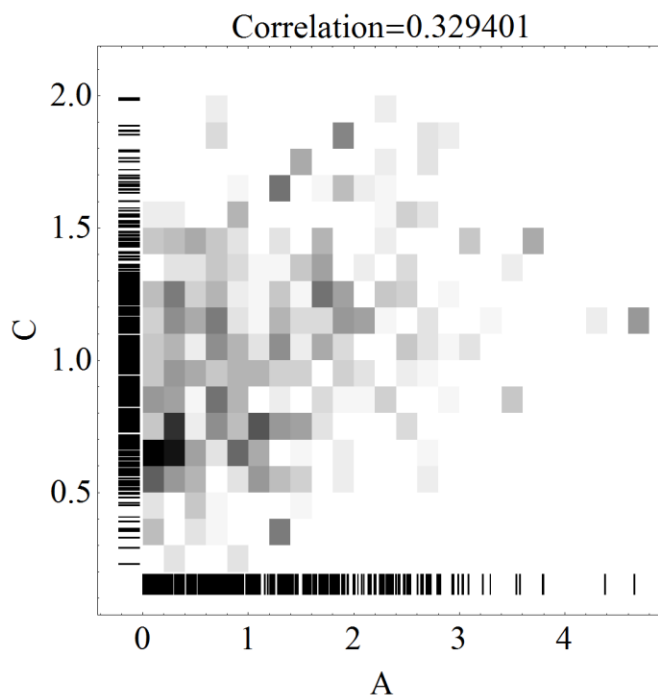
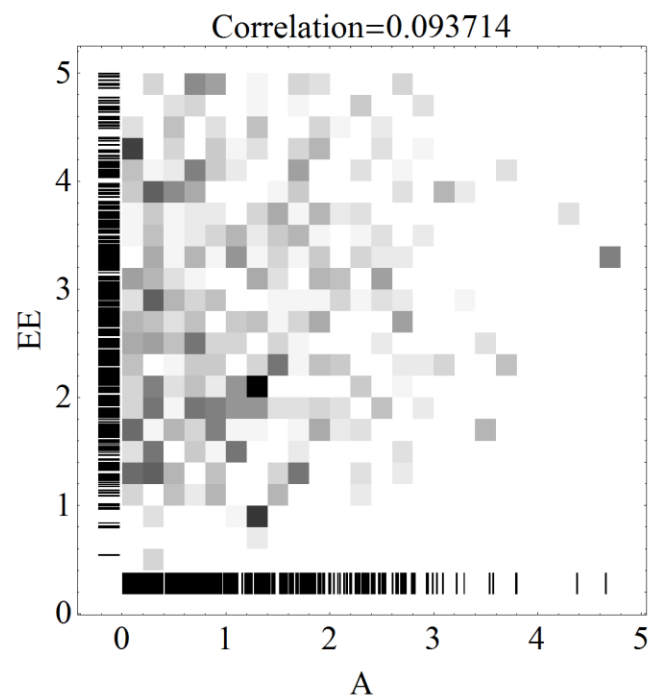
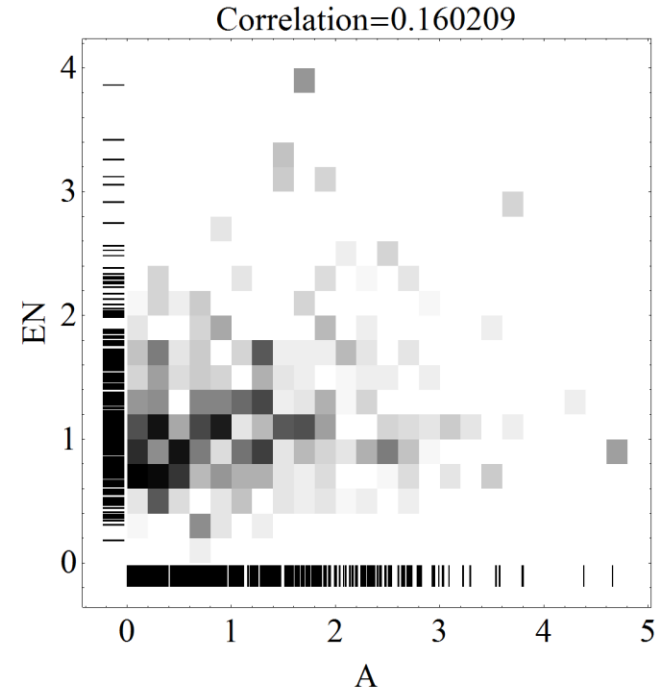
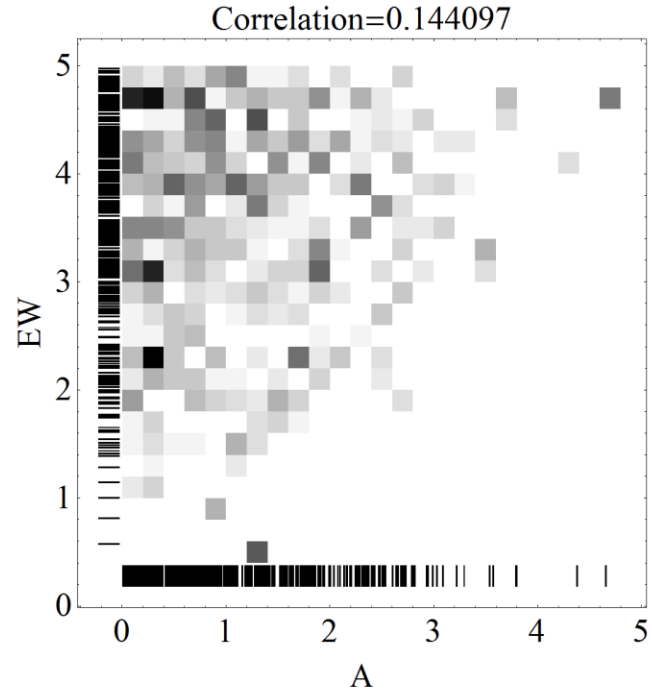


Figure SA11

Scenario 1

Inference validation

Model B:

extinction rate is diversity-dependent

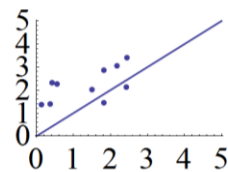
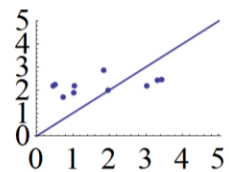
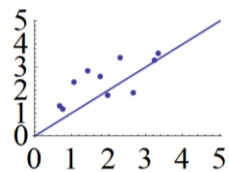
- NO PARAMETER HAS BEEN FIXED
- Time has been fixed to $T=8$ for simulating and estimating parameters
- 10 simulated data have been generated
- Sample size per generation = 400
- Generations = 5
- $q=0.5$
- Maximum number of lineages = 150

Regions

West

North

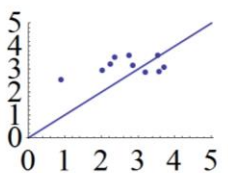
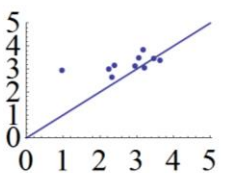
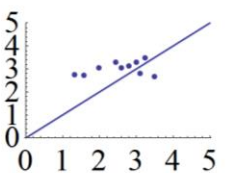
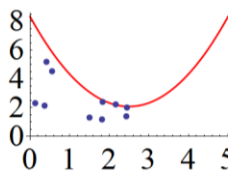
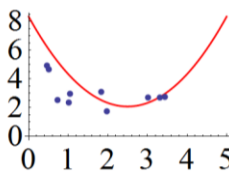
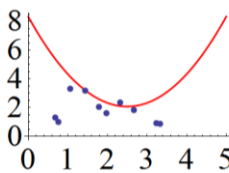
East



Known
value
vs mean

**Sympatric
speciation (s)**

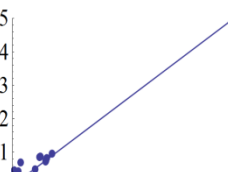
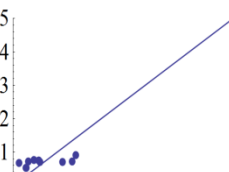
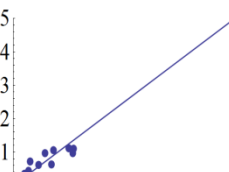
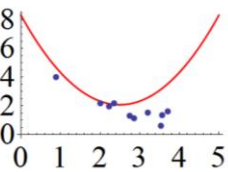
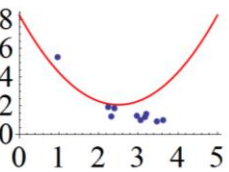
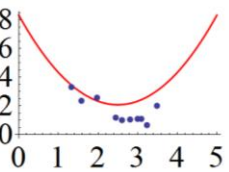
Mean
Squared
error



Known
value
vs mean

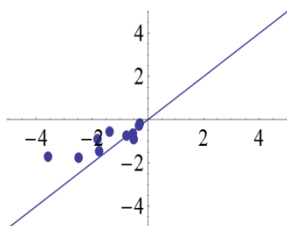
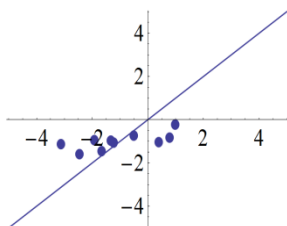
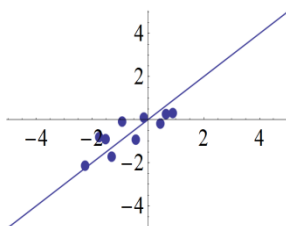
Extinction (e)

Mean
Squared
error



Ratio s:e

**Net
diversification
s-e**



Colonization

**Allopatric
speciation**

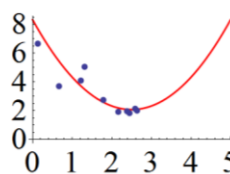
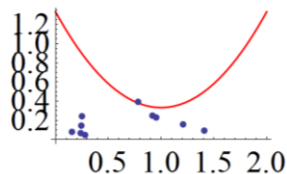
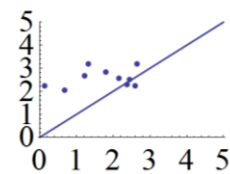
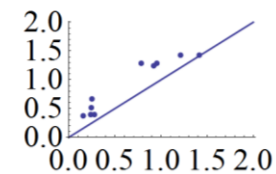


Figure SB1

Distribution of correlation between sympatric speciation and extinction (S..._E...)

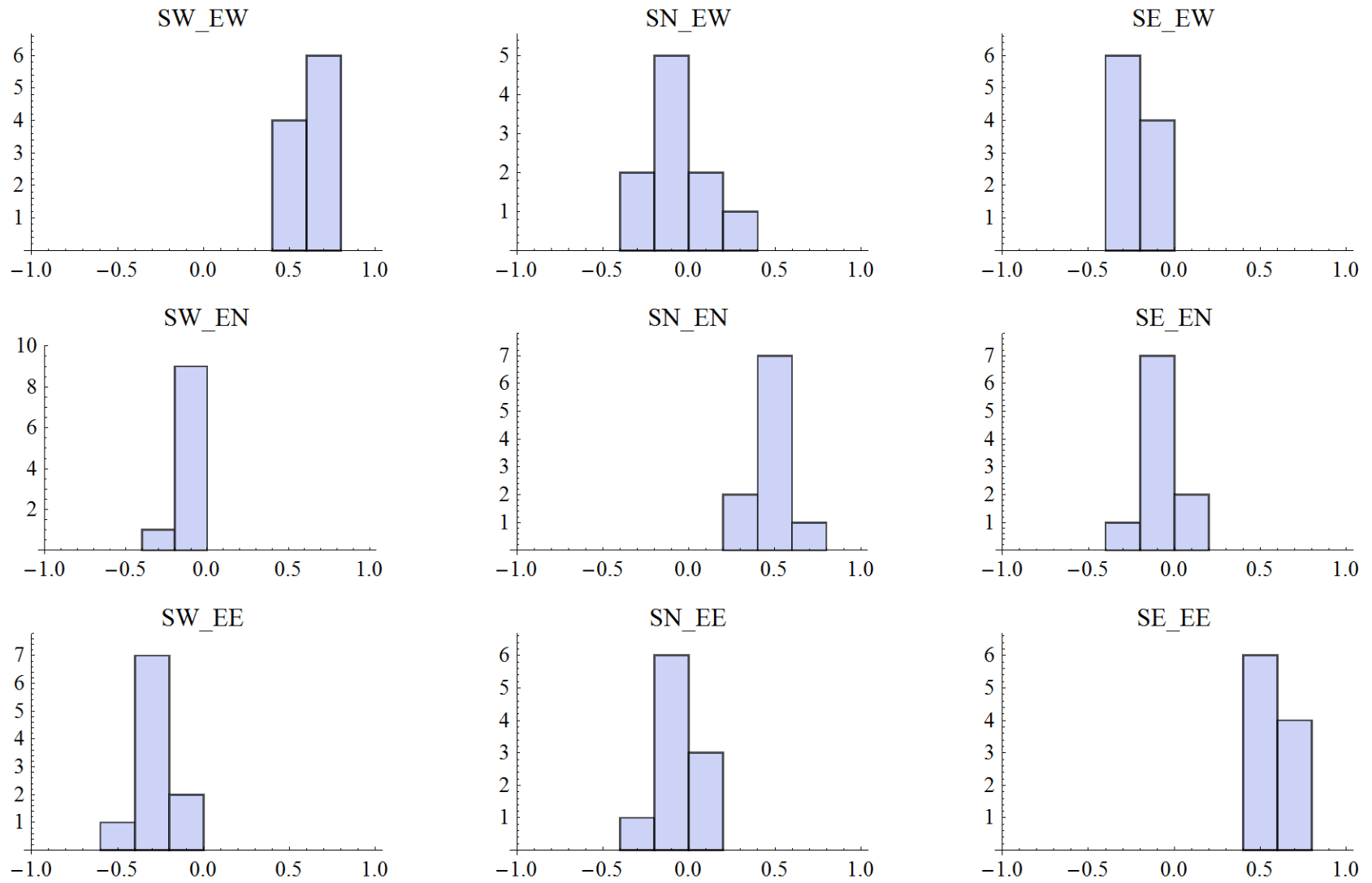


Figure SB2

Distribution of correlation between sympatric speciation (S..._S...), and between sympatric speciation and allopatric speciation (S..._A), and between extinction (E..._E...)

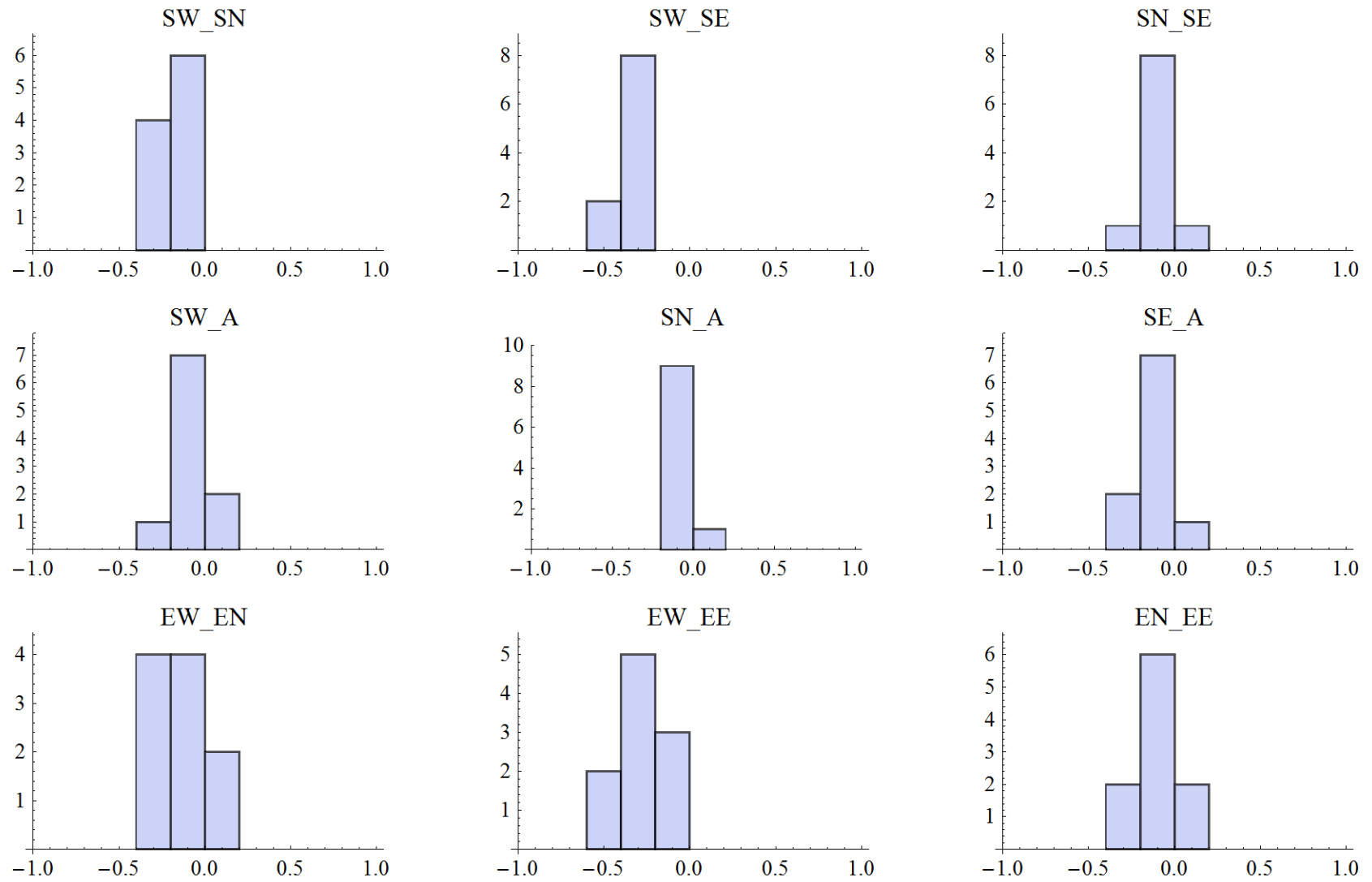


Figure SB3

Distribution of correlation between sympatric speciation and colonization (C_S...),
and between colonization and extinction (C_E...)

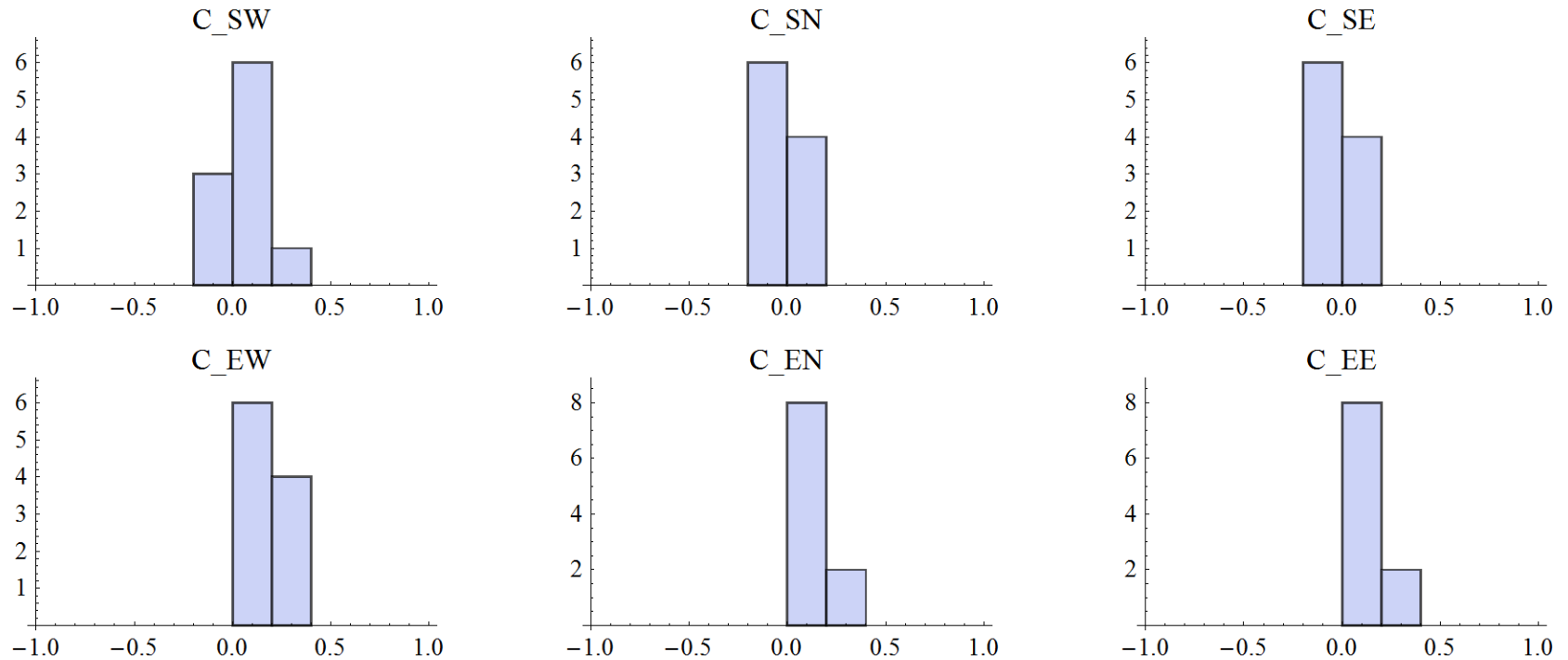


Figure SB4

Distribution of correlation between allopatric speciation and extinction (A_E...),
and between allopatric speciation and colonization (C_E...)

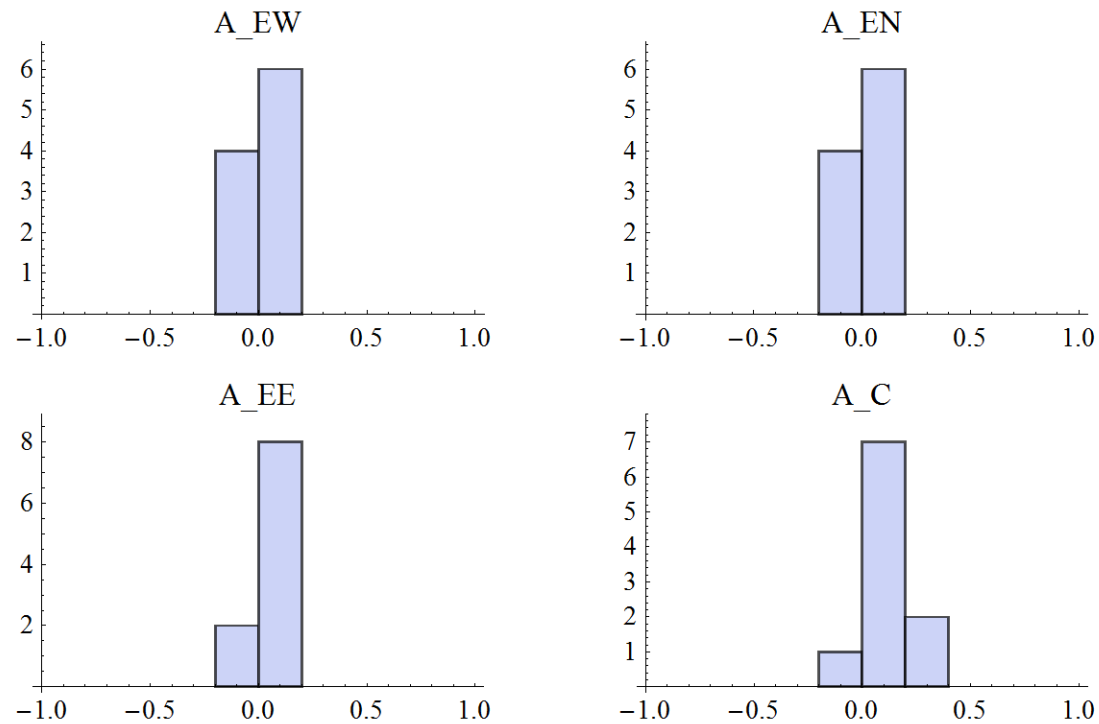


Figure SB5

Scenarion 1

Estimates from Empirical data

Model B: extinction rate is diversity-dependent

- NO PARAMETER HAS BEEN FIXED
- Estimating parameters for empirical data
- $T=8$
- Sample size = 400
- $q=0.5$
- Number of generation = 7
- Maximum number of lineages = 150

Generations

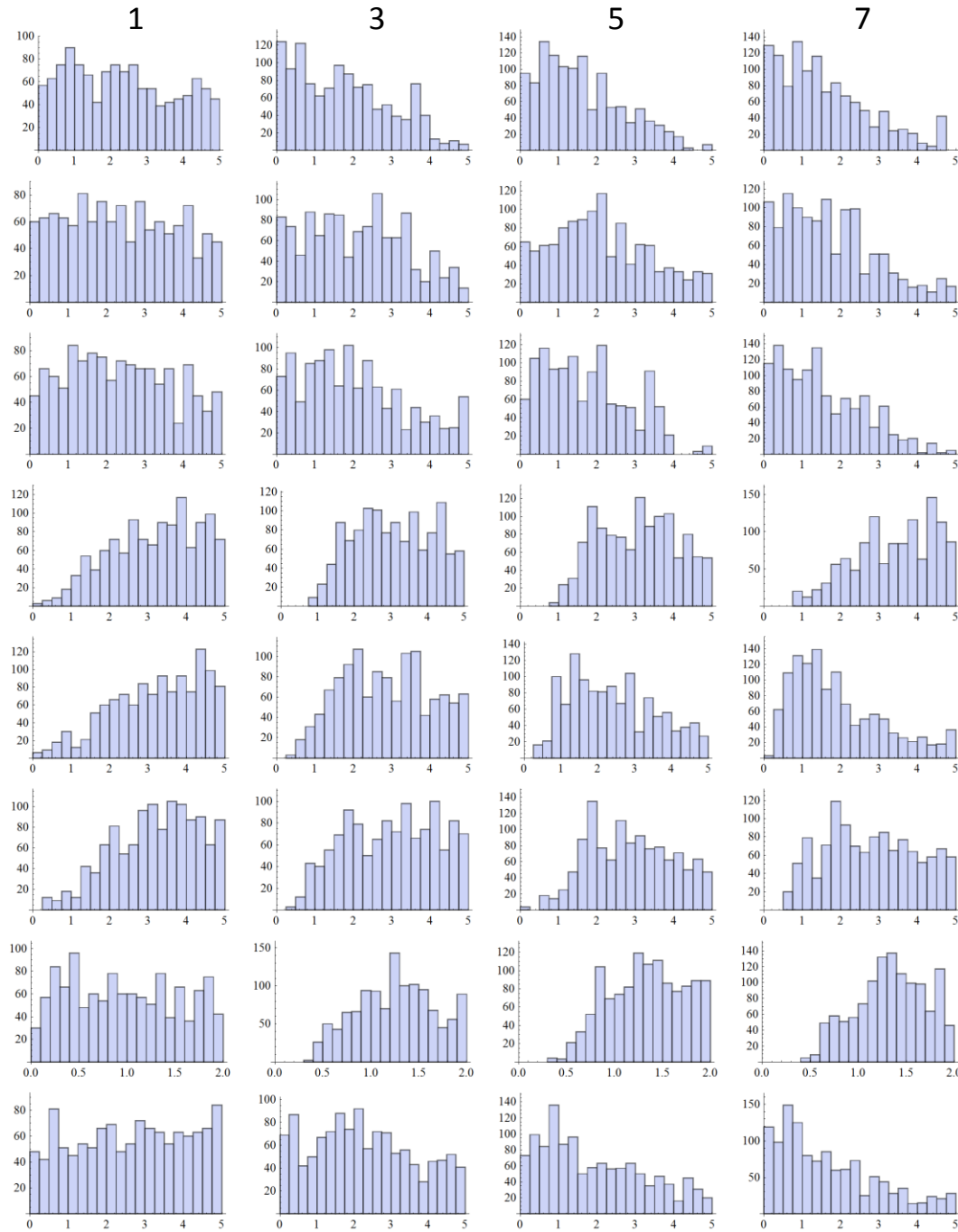


Figure SB6

Generations

1

3

5

7

West

North

East

West

North

East

Ratio
S:E

Net
diversification
S-E

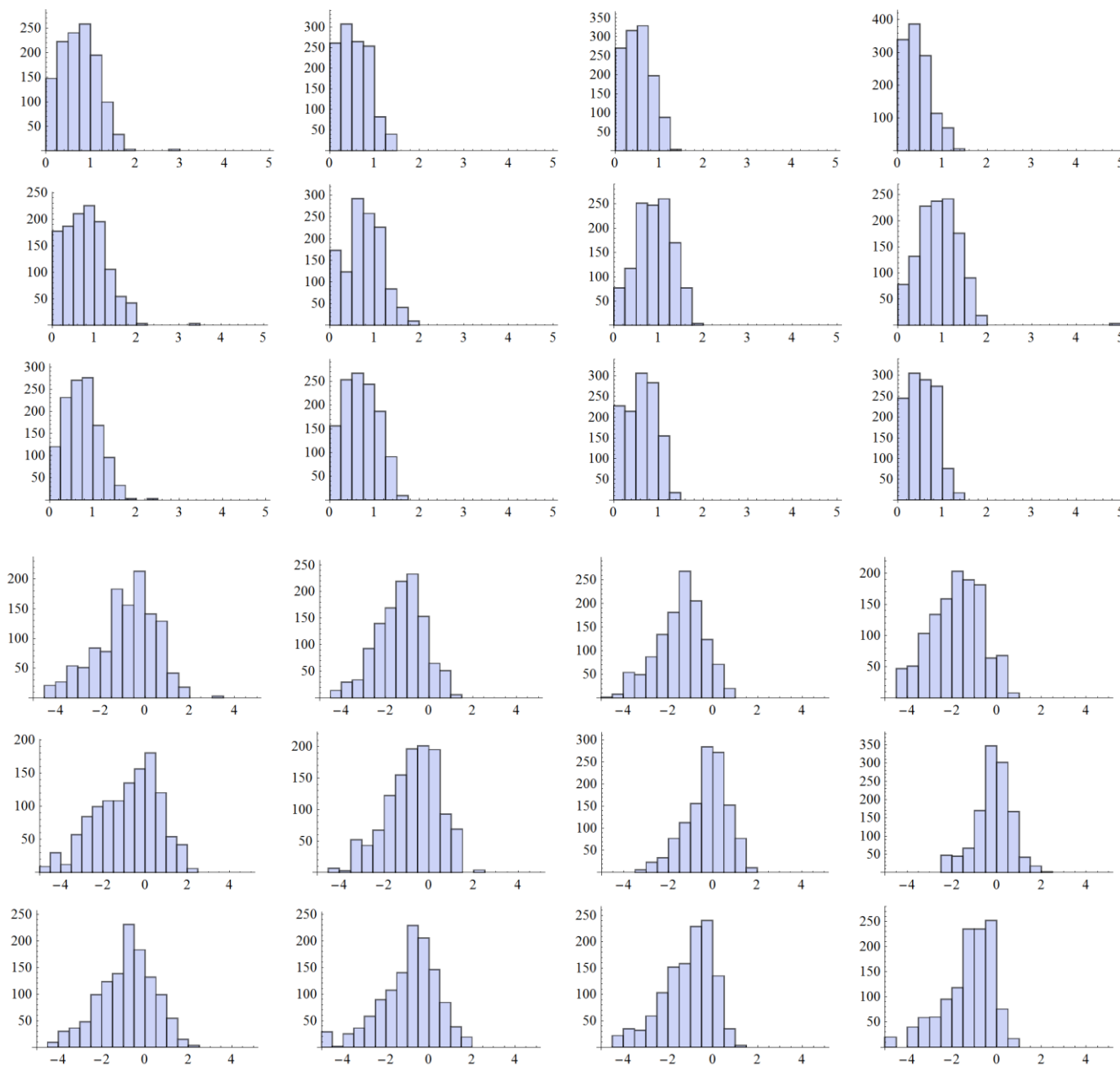


Figure SB7

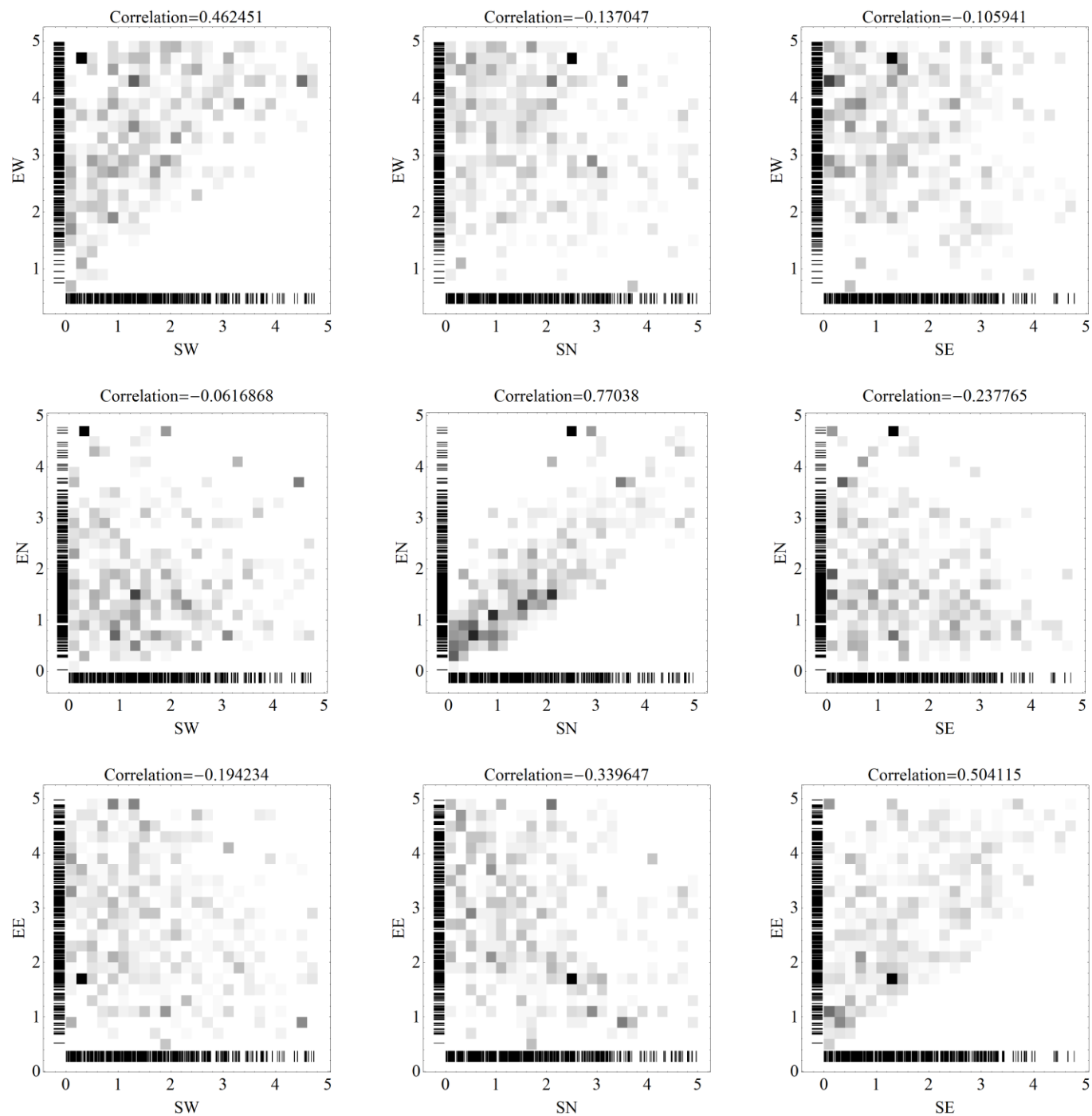


Figure SB8

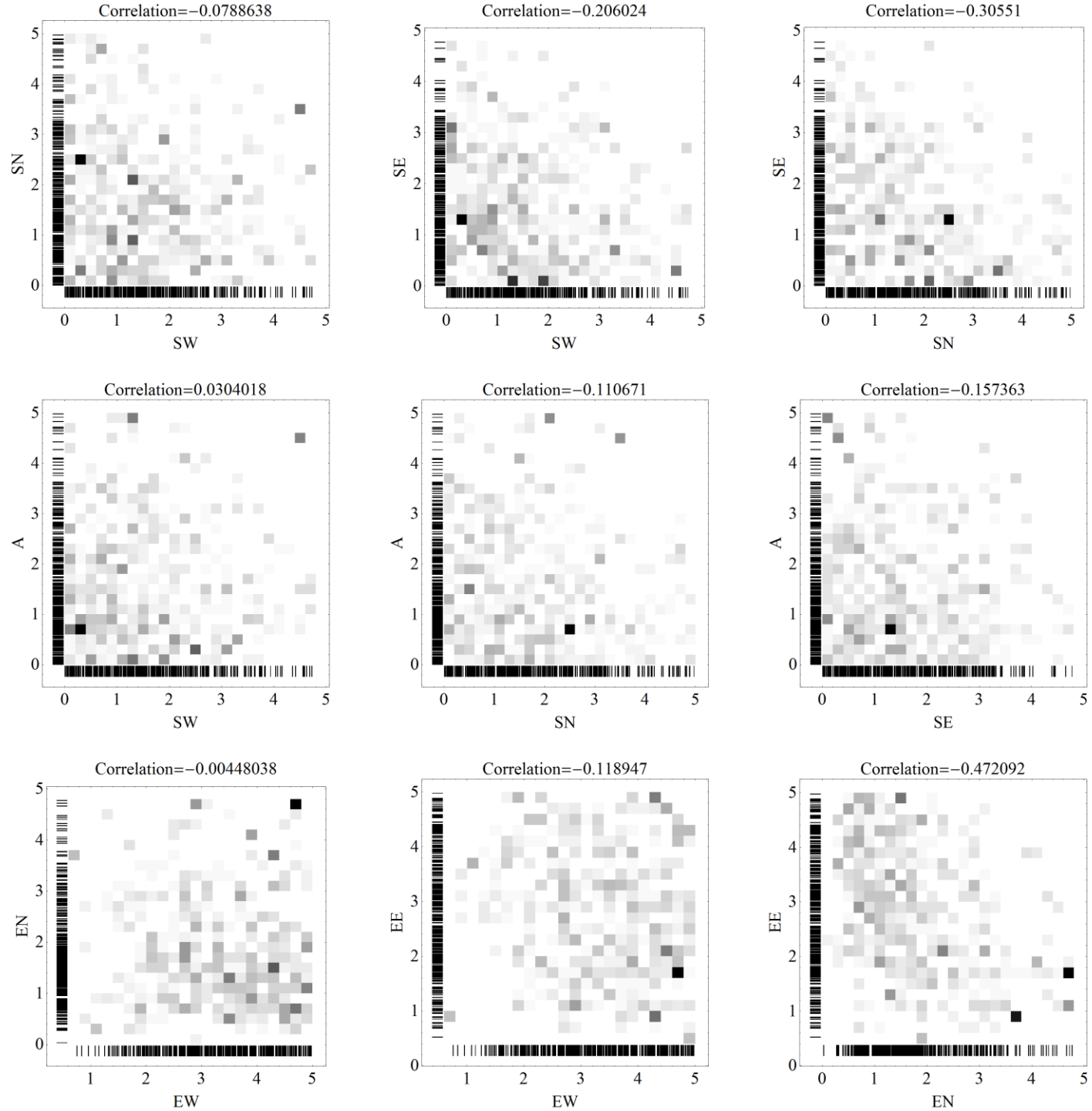


Figure SB9

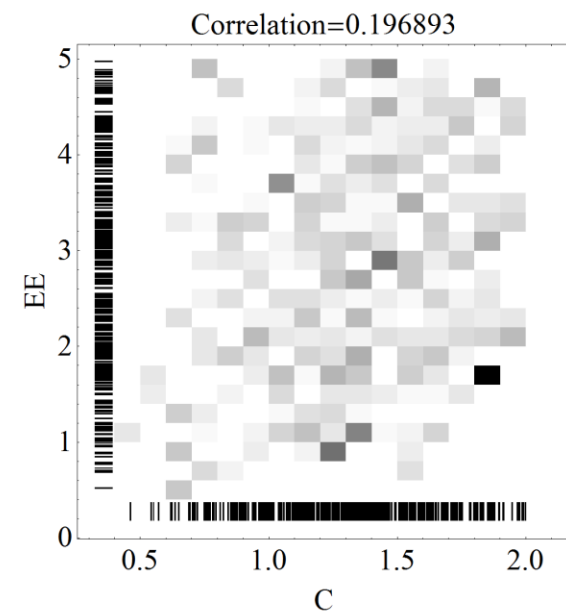
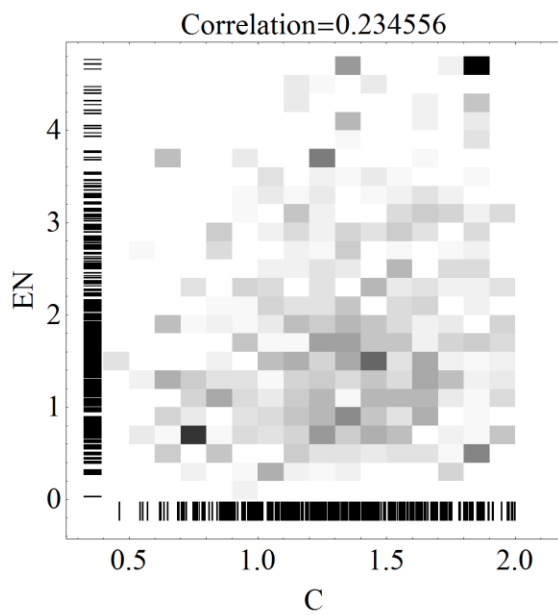
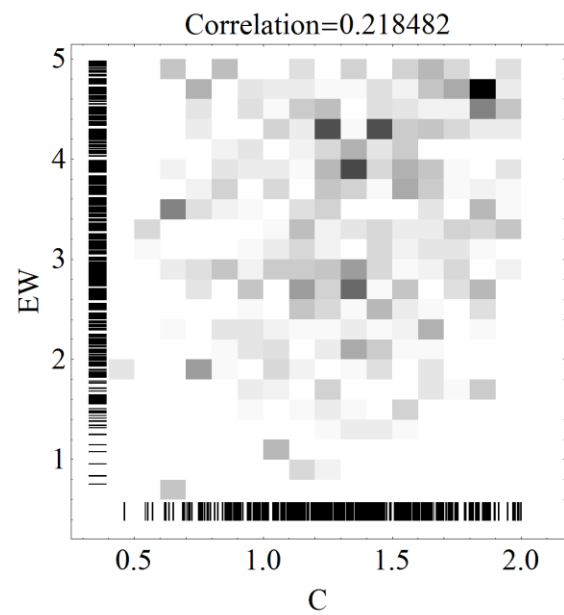
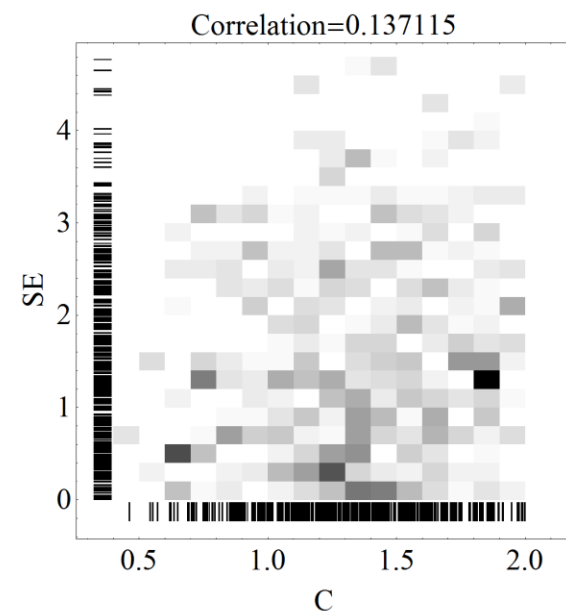
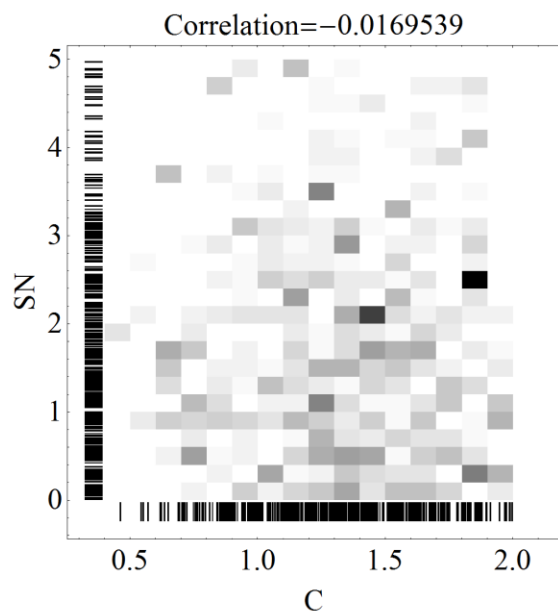
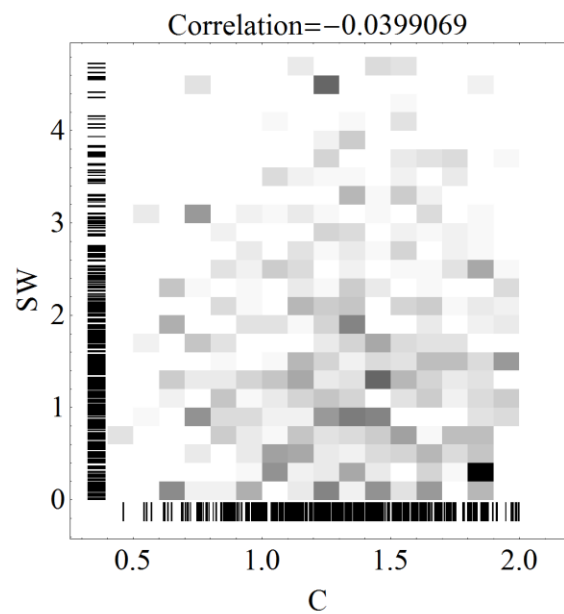


Figure SB10

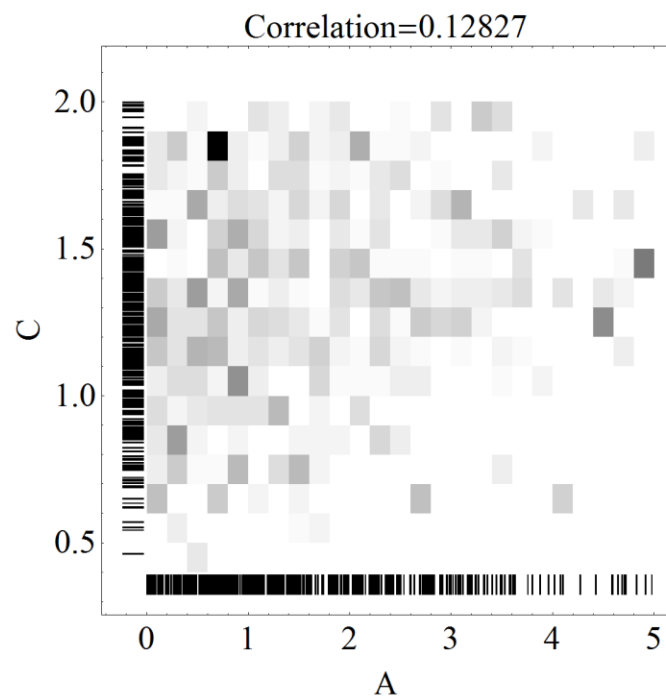
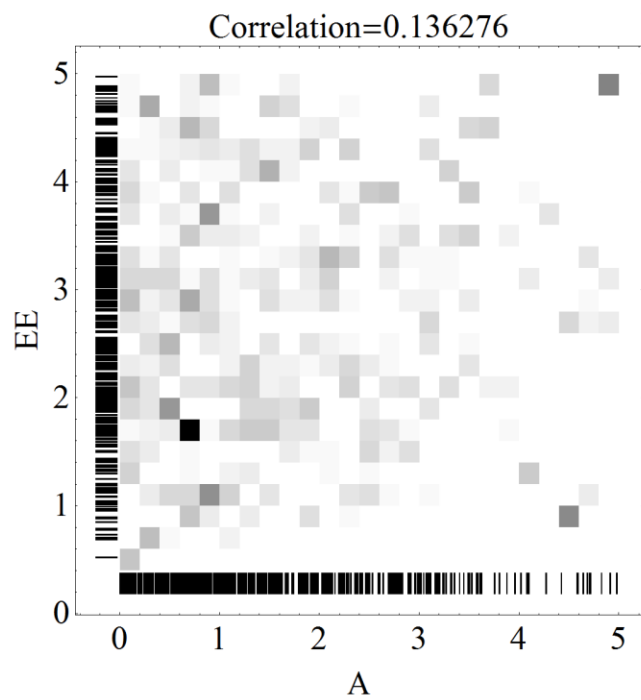
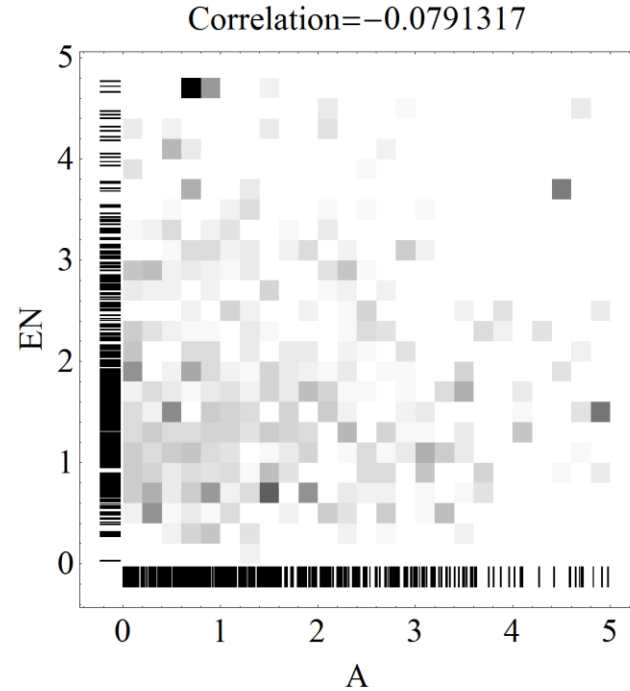
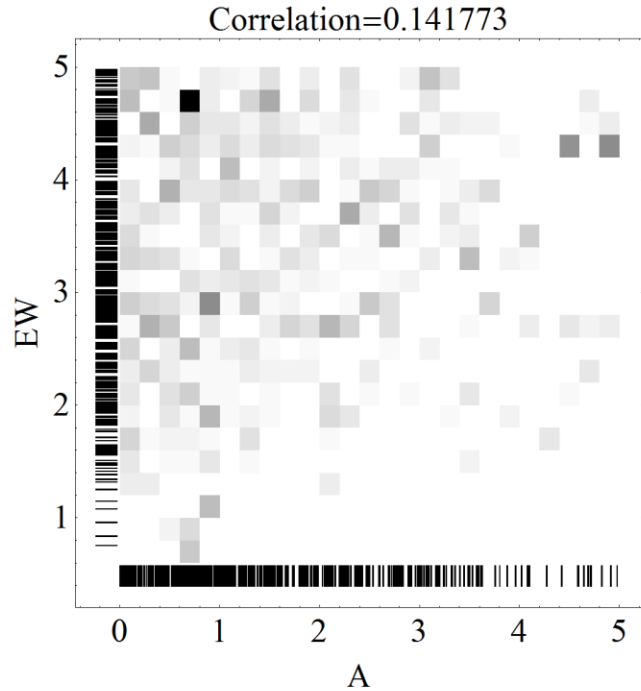


Figure SB11