Scenario 2

Inference validation

Model A: sympatric speciation rate is diversity-dependent

- Time has been fixed to T=8 for simulating and estimating parameters
- 10 simulated data have been generated by fixing extinction rate in the North to 1
- Sample size per generation = 400
- Generations = 7
- q=0.5
- Maximum number of species allowed 150

Regions West North East 4 3 2 1 0 4 3 2 1 4 3 2 1 vs mean Known value speciation (s) Sympatric Colonization **Allopatric** speciation 2.0 1.5 Mean Squared error 1.0 0.5 0.0 0.5 1.0 1.5 2.0 4 3 2 1 0 4 3 2 1 Known value vs mean Extinction (e) 1.0 1.5 2.0 0.5 Mean Squared error Ratio s:e Net diversification s-e

Figure SA1

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

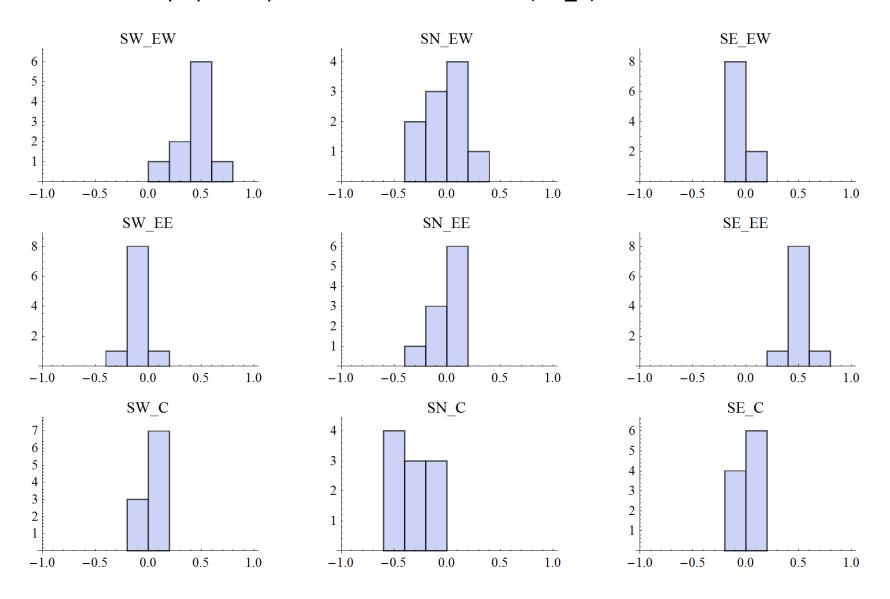


Figure SA2

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

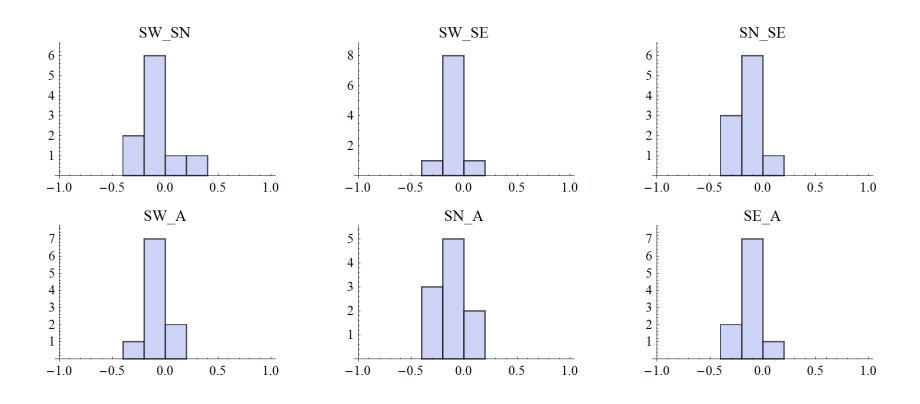
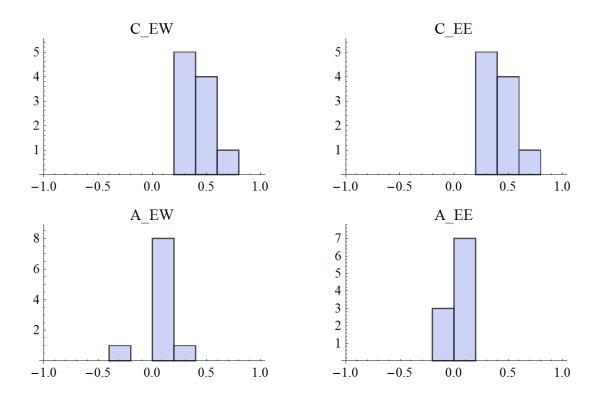
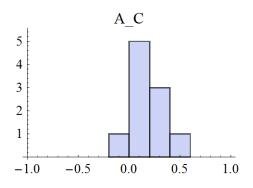


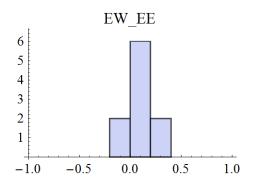
Figure SA3

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



Distribution of correlation between allopatric speciation and colonization (A_C), and between extinction (EW_EE).





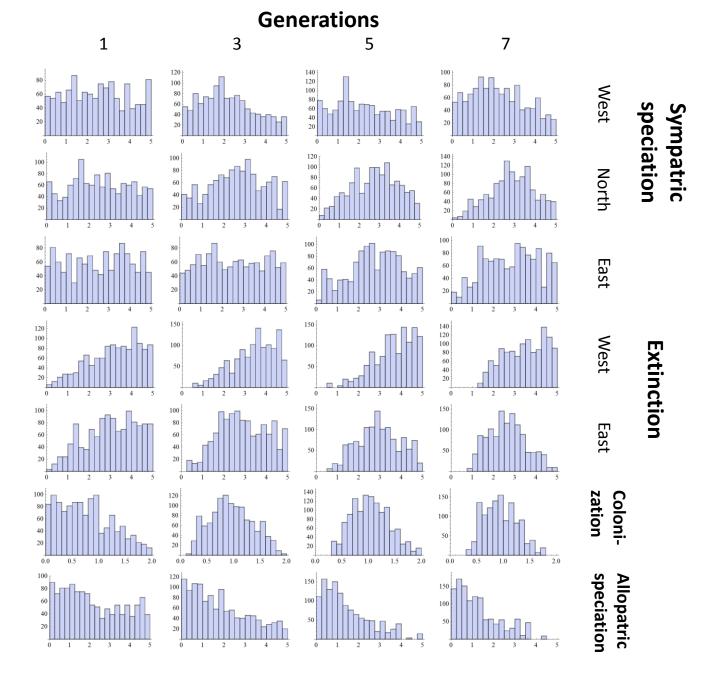
Scenario 2

Estimates from Empirical data

Model A:

sympatric speciation rate is diversity-dependent

- Estimating parameters for empirical data while fixing extinction rate in the north to 1
- T=8
- Sample size = 400
- q=0.5
- Number of generation = 7
- Maximum number of species = 150



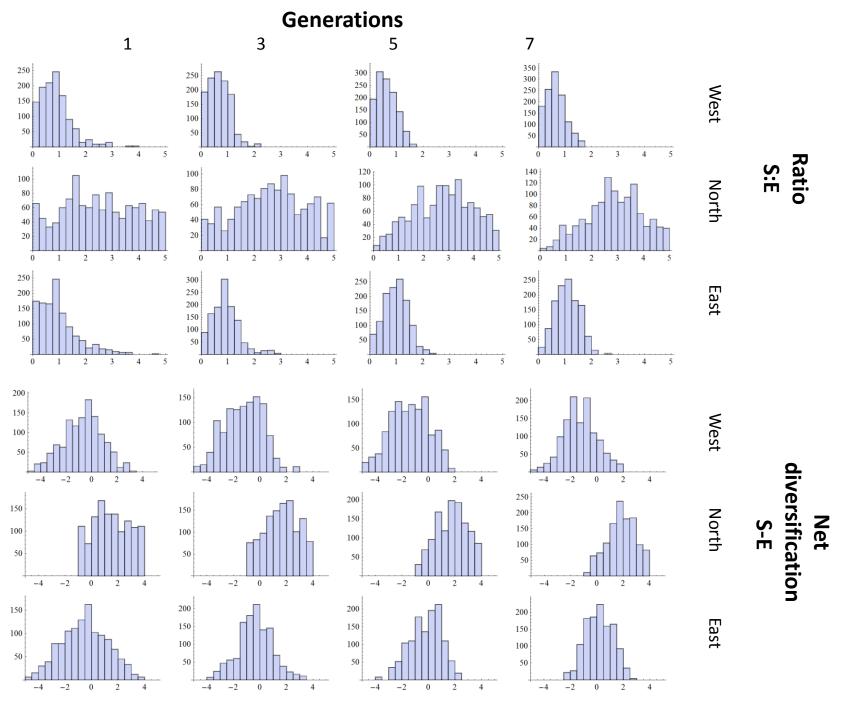


Figure SA7

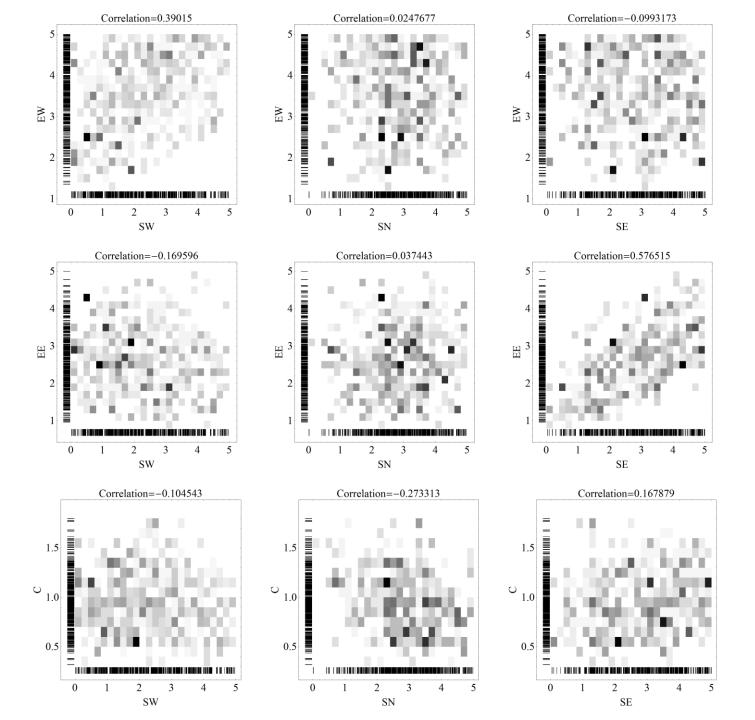
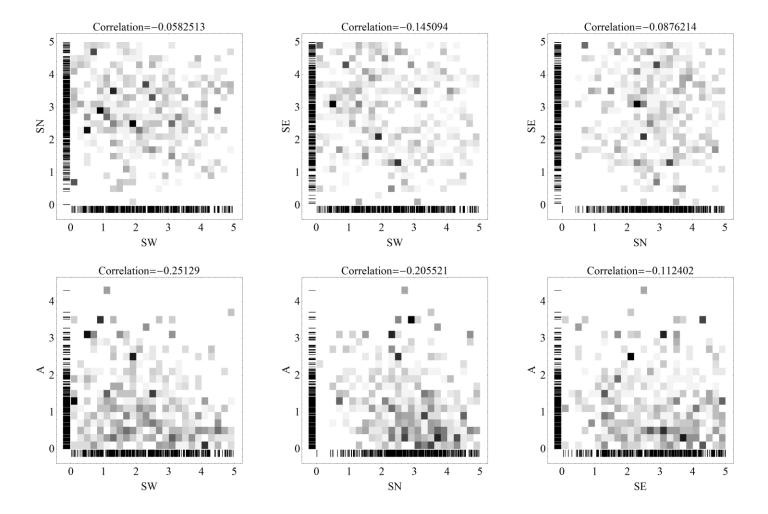


Figure SA8



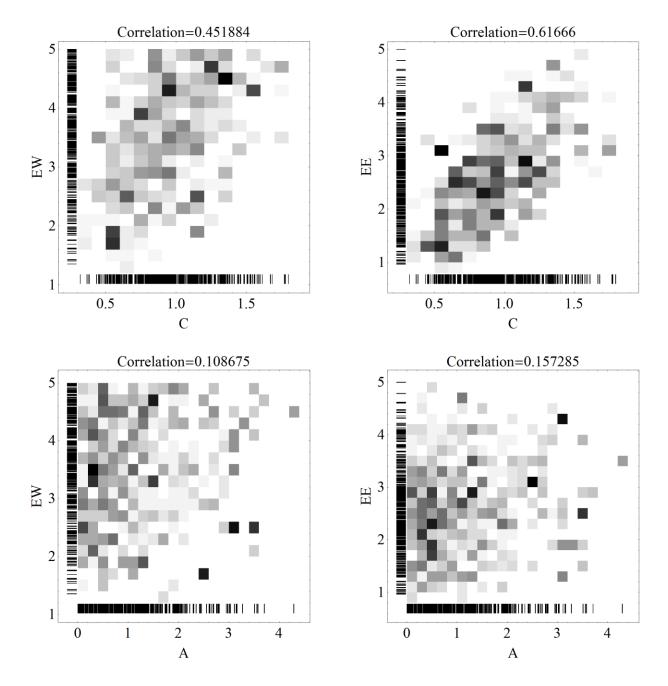
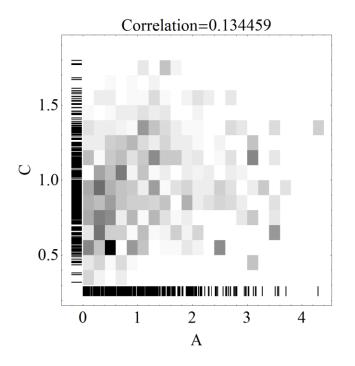
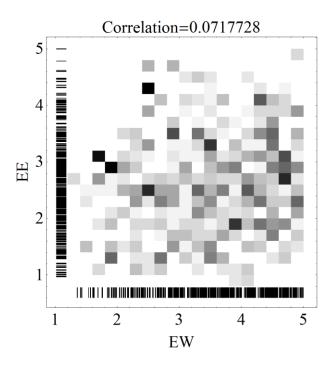


Figure SA10





Scenario 2

Inference validation

Model B: extinction rate is diversity-dependent

- Time has been fixed to T=8 for simulating and estimating parameters
- 10 simulated data have been generated by fixing extinction rate in the North to 1
- Sample size per generation = 400
- Generations = 7
- q=0.5
- Maximum number of species allowed 150

Regions West North East 4 3 2 1 vs mean Known value speciation (s) Sympatric Colonization **Allopatric** speciation 2.0 1.5 Mean Squared error 1.0 0.5 0.0 0.5 1.0 1.5 2.0 4 3 2 1 0 0 vs mean Known value Extinction (e) ٠. 1.0 1.5 2.0 0.5 Mean Squared error 4 3 2 Ratio s:e Net diversification s-e

Figure SB1

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

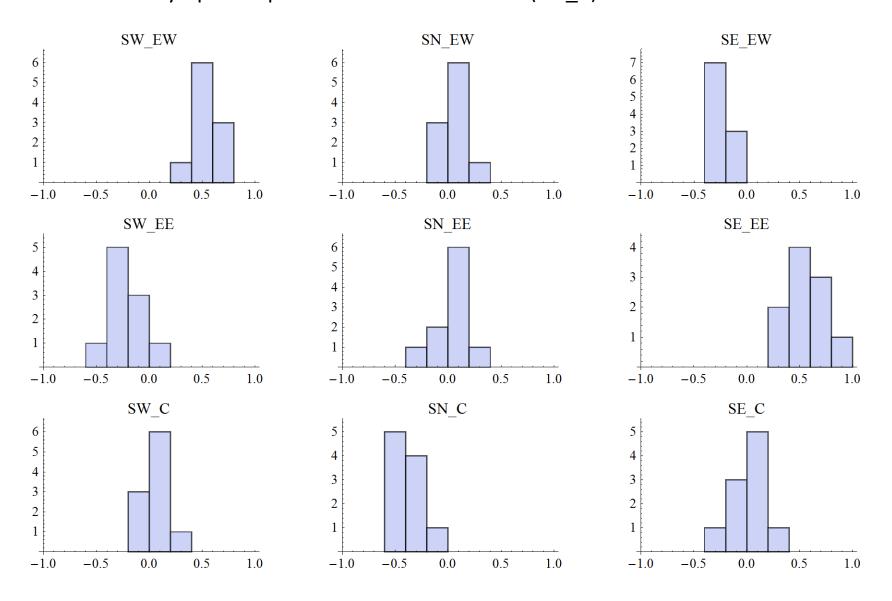


Figure SB2

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

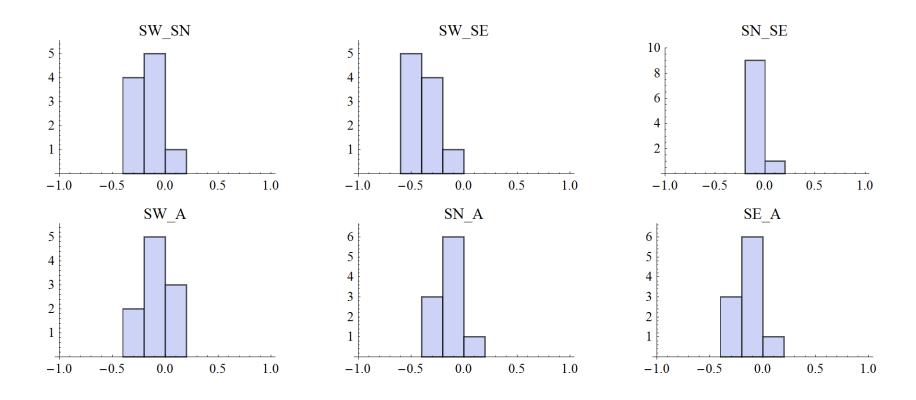
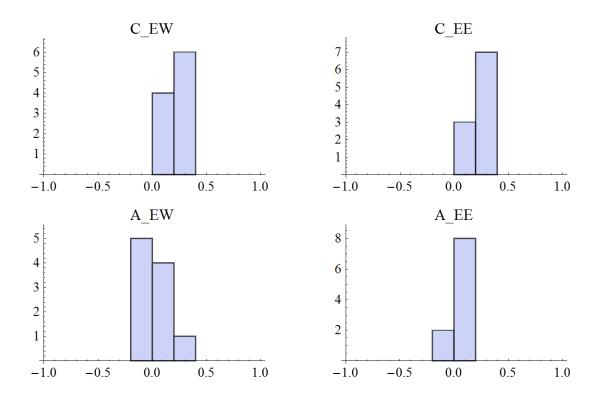
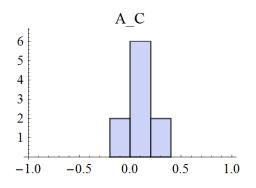


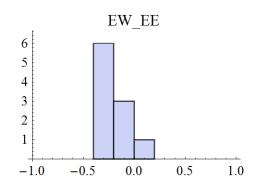
Figure SB3

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



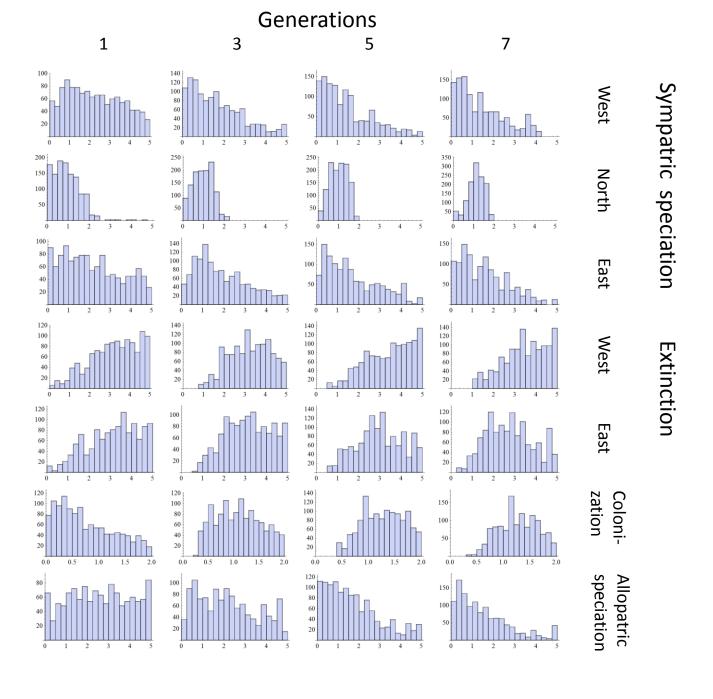
Distribution of correlation between allopatric speciation and colonization (A_C), and between extinction (EW_EE).





Model B: extinction rate is diversity-dependent

- Estimating parameters for empirical data while fixing extinction rate in the north to 1
- T=8
- Sample size = 400
- q=0.5
- Number of generation = 7
- Maximum number of species = 150



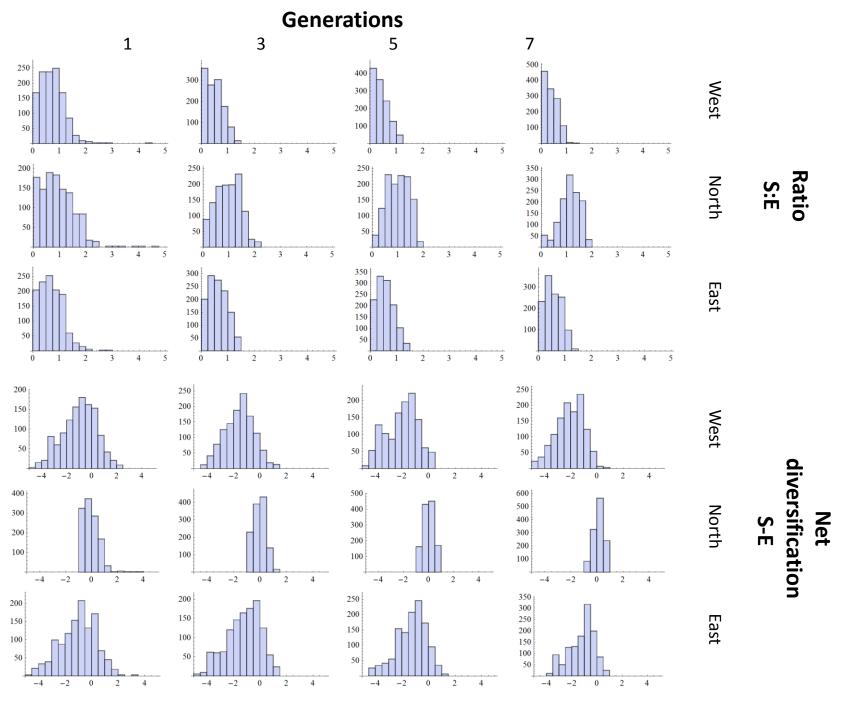


Figure SB7

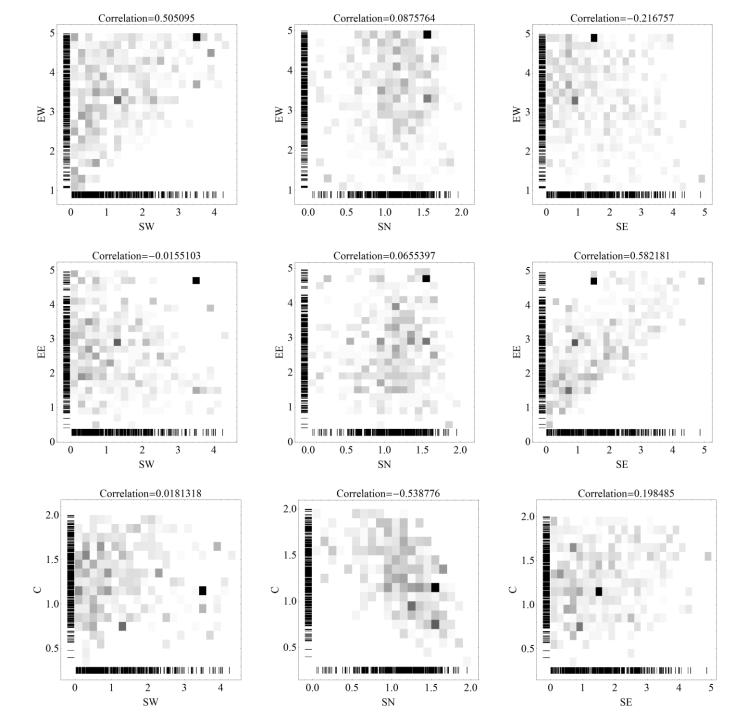
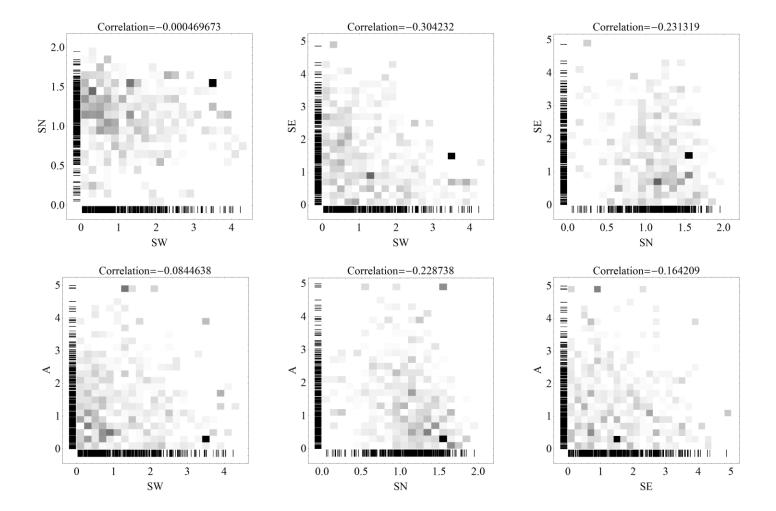


Figure SB8



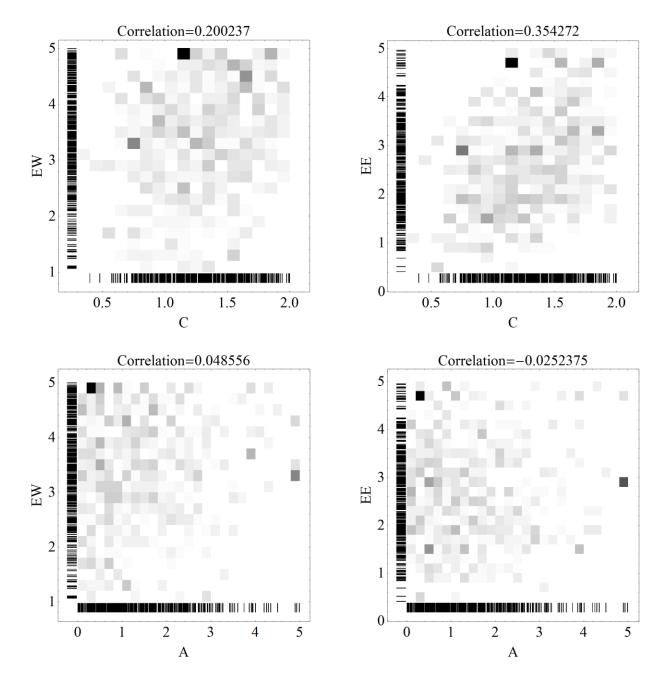


Figure SB10

