Table S7. ANOVA outputs of predictor 'Data type', which represents the neighbor treatment predictor with two levels: with neighbors, without neighbors. Separate models were fit for each scale and (mis)alignment category. NAs indicate there were no instances in our data, thus comparisons are not possible.

| Scale level | (Mis)alignment | Predictor | Chi.squared | Df | P-value |
|-------------|---------------------------|-----------|-------------|----|---------|
| plot | i. aligned present | Intercept | 77.445 | 1 | 0.000 |
| plot | i. aligned present | Data type | 43.856 | 1 | 0.000 |
| plot | ii. sink | Intercept | 4.649 | 1 | 0.031 |
| plot | ii. sink | Data type | 16.316 | 1 | 0.000 |
| plot | iii. dispersal limitation | Intercept | 73.827 | 1 | 0.000 |
| plot | iii. dispersal limitation | Data type | 19.423 | 1 | 0.000 |
| plot | iv. aligned absent | Intercept | 19.730 | 1 | 0.000 |
| plot | iv. aligned absent | Data type | 17.664 | 1 | 0.000 |
| grid | i. aligned present | Intercept | 0.004 | 1 | 0.948 |
| grid | i. aligned present | Data type | 16.322 | 1 | 0.000 |
| grid | ii. sink | Intercept | 0.605 | 1 | 0.437 |
| grid | ii. sink | Data type | 15.032 | 1 | 0.000 |
| grid | iii. dispersal limitation | Intercept | 24.113 | 1 | 0.000 |
| grid | iii. dispersal limitation | Data type | 0.000 | 1 | 0.993 |
| grid | iv. aligned absent | Intercept | 6.181 | 1 | 0.013 |
| grid | iv. aligned absent | Data type | 1.005 | 1 | 0.316 |
| site | i. aligned present | Intercept | 0.404 | 1 | 0.525 |
| site | i. aligned present | Data type | 8.011 | 1 | 0.005 |
| site | ii. sink | Intercept | 0.404 | 1 | 0.525 |
| site | ii. sink | Data type | 8.011 | 1 | 0.005 |
| site | iii. dispersal limitation | Intercept | NA | NA | NA |

Table S7. ANOVA outputs of predictor 'Data type', which represents the neighbor treatment predictor with two levels: with neighbors, without neighbors. Separate models were fit for each scale and (mis)alignment category. NAs indicate there were no instances in our data, thus comparisons are not possible.

| Scale level | (Mis)alignment | Predictor | Chi.squared | Df | P-value |
|-------------|---------------------------|-----------|-------------|----|---------|
| site | iii. dispersal limitation | Data type | NA | NA | NA |
| site | iv. aligned absent | Intercept | NA | NA | NA |
| site | iv. aligned absent | Data type | NA | NA | NA |