

TABLE 6. Statistical comparison of geographic range size in taxonomic groups by sequence. (1) Sequence C1: Edenian; (2) Sequence C2: early Maysvillian; (3) Sequence C3: late Maysvillian; (4) Sequence C4: early Richmondian; (5) Sequence C5: middle Richmondian; (6) Sequence C6: late Richmondian. Analyses were conducted by defining *Strophomena* species as either native or invasive as indicated in text. Blocks color coded by statistical significance.

1. Sequence C1: Edenian

	Species: Area	Species: Linear	Genus: Area	Genus: Linear
<i>Strophomena</i> native	0.152	0.209	0.673	0.706
<i>Strophomena</i> invasive	0.152	0.209	0.219	0.233

2. Sequence C2: early Maysvillian

	Species: Area	Species: Linear	Genus: Area	Genus: Linear
<i>Strophomena</i> native	0.001	0.591	0.121	0.083
<i>Strophomena</i> invasive	0.001	0.591	0.108	0.098

3. Sequence C3: late Maysvillian

	Species: Area	Species: Linear	Genus: Area	Genus: Linear
<i>Strophomena</i> native	0.019	0.001	0.018	<0.001
<i>Strophomena</i> invasive	0.019	0.001	0.018	<0.001

4. Sequence C4: early Richmondian

	Species: Area	Species: Linear	Genus: Area	Genus: Linear
<i>Strophomena</i> native	0.014	0.011	0.038	0.100
<i>Strophomena</i> invasive	0.015	0.067	<0.001	<0.001

5. Sequence C5: middle Richmondian

	Species: Area	Species: Linear	Genus: Area	Genus: Linear
<i>Strophomena</i> native	0.092	0.174	0.007	0.045
<i>Strophomena</i> invasive	0.139	0.091	0.024	0.067

6. Sequence C6: late Richmondian

	Species: Area	Species: Linear	Genus: Area	Genus: Linear
<i>Strophomena</i> native	0.820	0.664	0.112	0.771
<i>Strophomena</i> invasive	0.204	0.744	0.072	0.619

p ≤ 0.050	p ≤ 0.100
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