	atpB	rbcL	matK	trnTLF	ITS	TOTAL
Total native tips	41	114	119	128	176	226
Total invasive tips	21	133	134	108	125	140
Total tips	62	247	253	236	301	366
Final sequence length	1451	1311	1456	881	646	5745
Matrix coverage	0.17%	0.67%	0.69%	0.64%	0.82%	0.60%

Table S1 Number of native and invasive species for each gene region, final sequence length, and percent of each gene partition in the final concatenated nucleotide matrix post cleaning (Decisivator version 0.55).