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### *Using DNA Metabarcoding to Identify Floral Visitation by Pollinators*

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**Table S2:** Recommended primer sequences used to amplify the *rbcl* and ITS2 barcode regions. A 6N sequence is added between the forward template specific primer and the universal tail to improve clustering on the Illumina MiSeq.

Primer name	Universal Tail	6N Sequence	Primer sequence
<i>rbclaf</i> [1]	<b>ACACTCTTTCCTACACGACGCTCTTCCGATCT</b>	NNNNNN	ATGTCACCACAAACAGAGACTAAAGC
<i>rbclr506</i> [2]	<b>GTGACTGGAGTTCAGACGTGTGCTCTTCCGATCT</b>		AGGGGACGACCATACTTGTTC
ITS2F [3]	<b>ACACTCTTTCCTACACGACGCTCTTCCGATCT</b>	NNNNNN	ATGCGATACTTGGTGTGAAT
UniPlantR [4]	<b>GTGACTGGAGTTCAGACGTGTGCTCTTCCGATCT</b>		CCCGHYTGAYTGRGGTDC

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