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Table 5. Sequence length for the *trnH-psbA* spacer in the 50 families encompassing 72 genera and 83 species sampled from Plummers Island flora

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Family (number of taxa sampled: genera/species)	<i>trnH-psbA</i> spacer length, bp
Acanthaceae (1/1)	289
Amaranthaceae (2/4)	138–201
Apiaceae (3/3)	185–188
Apocynaceae (3/3)	230–361
Asteraceae (9/15)	204–400
Balsaminaceae (1/1)	240
Betulaceae (1/1)	200
Bignoniaceae (1/1)	342
Boraginaceae (1/1)	376/
Brassicaceae (2/2)	291
Cannabaceae (1/1)	288
Caprifoliaceae (2/3)	334–386
Caryophyllaceae (2/2)	242
Celastraceae (1/1)	424
Commelinaceae (1/1)	743
Convolvulaceae (1/1)	240

Crassulaceae (1/1)	412
Cucurbitaceae (1/1)	172
Cyperaceae (2/4)	497–788
Dioscoreaceae (1/1)	279
Euphorbiaceae (1/1)	441
Fabaceae (1/1)	296
Hydrangeaceae (1/1)	338
Hypericaceae (1/1)	487
Lamiaceae (2/2)	324–349
Lobeliaceae (1/1)	337
Lythraceae (1/1)	238
Malvaceae (2/2)	385–637
Melanthiaceae (1/1)	1,094
Molluginaceae (1/1)	234
Oleaceae (1/1)	446
Onagraceae (2/2)	283–431
Oxalidaceae (1/1)	324
Passifloraceae (1/1)	220
Phytolaccaceae (1/1)	461
Plantaginaceae (1/1)	260
Poaceae (2/2)	534
Polemoniaceae (1/1)	407
Polygonaceae (1/1)	265
Pontederiaceae (1/1)	559
Primulaceae (1/1)	414
Ranunculaceae (2/2)	119–189
Rosaceae (1/1)	277

Rubiaceae (1/1)	371
Sapindaceae (1/1)	403
Scrophulariaceae (1/1)	323
Solanaceae (1/1)	417
Urticaceae (1/1)	200
Verbenaceae (1/1)	274
Violaceae (1/1)	375

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