

Table S2.

Comparison of genome size of selected specimens of *V. pubifolia*, *V. palustris*, hybrids *V. epipsila* × *V. palustris*, and *V. epipsila*.

Taxon	Mean	
	2C DNA (pg) (±SD)	1C/Cx (pg)
<i>V. pubifolia</i> (N=39)	4.14 (±0.14) <sup>a</sup>	2.07/0.52
<i>V. epipsila</i> (N= 3)	2.52* (±0.03) <sup>b</sup>	1.26/0.63
<i>V. palustris</i> (N= 4)	4.26* (±0.02) <sup>a</sup>	2.13/0.53
Hybrid (N=15) ( <i>V. epipsila</i> × <i>V. palustris</i> )	3.42* (±0.06) <sup>c</sup>	1.71/0.57

Average values followed by the same letter do not differ significantly at  $P \leq 0.05$  in Kruskal-Wallis test

\*Data taken from Žabicka et al. 2020 [30].