Vojtěchová K., Kobrlová L., Schönswetter P. & Duchoslav M. (2023) Disentangling the taxonomic structure of the *Allium paniculatum* species complex in central and eastern Europe using molecular, cytogenetic and morphological tools. – Preslia 95: 119–163.

Supplementary Table S5. – Absolute genome size (AGS), DNA base content (AT, CG, in %) of selected plants of the taxonomic groups studied. For detailed information on localities see Supplementary Table S1.

Taxonomic group	Population ID	Ploidy (2n)	AGS (mean±SD; pg)	AT (%)	CG (%)
A. fuscum	6	2x	30.86±0.14	63.87	36.13
A. fuscum	7	2x	32.17±0.15	62.06	37.94
A. marginatum	8	2x	30.36±0.09	63.47	36.53
A. marginatum	17	2x	28.54±0.21	63.45	36.55
A. podolicum	18	3x	40.00±0.25	63.01	36.99
A. podolicum	20	2x	27.47±0.04	63.69	36.31
A. paniculatum	21	2x	28.25±0.04	63.40	36.60
A. paniculatum	22	2x	29.38±0.17	62.78	37.22