

**Urfusová R., Urfus T., Krahulec F., Jarolímová V., Kopecký D. & Mahelka V. (2024)**  
**Morphological, flow cytometric and cytogenetic evidence of a hybrid swarm between**  
***Elymus hispidus* and *E. repens* in the Bílé Karpaty Mts, Czech Republic. – Preslia 95:**  
**97–122.**

**Supplementary Table S3.** Correlation of characters with the first and second principal components axes in the combined dataset for central-European and Čertoryje populations.

<b>Character</b>	<b>PC1</b>	<b>PC2</b>
logLSTwA	0.165516	-0.85218
logLSTwtA	-0.1212	-0.84077
logL2IR	-0.48784	-0.7277
logLSwA	-0.45145	-0.68122
logL4IR	-0.62222	-0.66708
logLG1	0.738012	-0.56807
logWST	0.046762	-0.51032
logL2UI	-0.40985	-0.3997
logLSTwtA.LSTwA	-0.41444	-0.35988
logW2L	-0.05379	-0.33131
logLSTwA.WST	0.11348	-0.28084
logSL	0.653115	-0.25226
logLAL.0.5	0.549709	-0.21935
logLAG.0.5	0.760775	-0.21566
logLSTwtA.WST	-0.18792	-0.2101
logNS	-0.06079	-0.19013
logSG	0.888465	-0.17759
logMRS.0.5.5	-0.00108	-0.13893
logHRS.0.5.5	0.308726	-0.11095
logNL	0.352851	-0.10221
logHL.0.5.5	-0.10251	-0.05442
logLG1.LSTwtA	0.877406	-0.05153
logLSwtA.LSwA	-0.14245	-0.02008
logDG1.2	0.230504	-0.01122
logLG1.LSTwA	0.794741	0.027202
logHALS.0.5.5	-0.01535	0.076542
logLC.0.5	-0.81264	0.077239
logNoLSwC.0.5	-0.88115	0.087675
logNoC.0.5	-0.88634	0.12831
logLG1.L4IR	0.920753	0.142942
logCSM.0.5	-0.86777	0.159949
logLSTwA.L4IR	0.762941	0.185325
logGL.0.5.5	-0.44337	0.292087