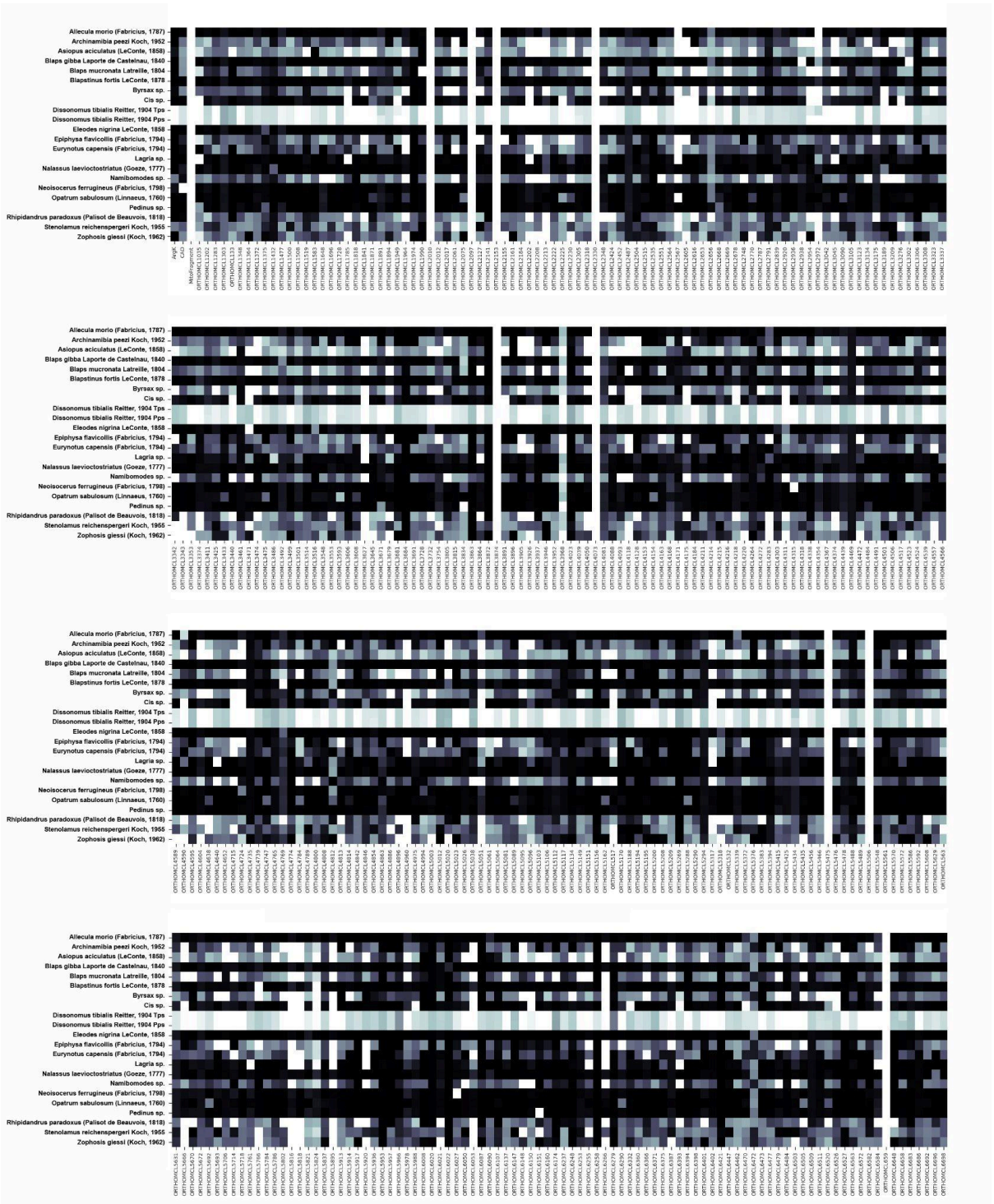
Supplementary data for Kamiński *et al*. (2024): Dissonomini Medvedev, 1968: the eighth tribe of the subfamily Blaptinae (Coleoptera: Tenebrionidae).

Figure S1. Heatmap of gene recovery efficiency. Rows represent processed samples, while columns loci. Color represents the percentage of recovered sequences (black - full recovery, white - loci not recovered).



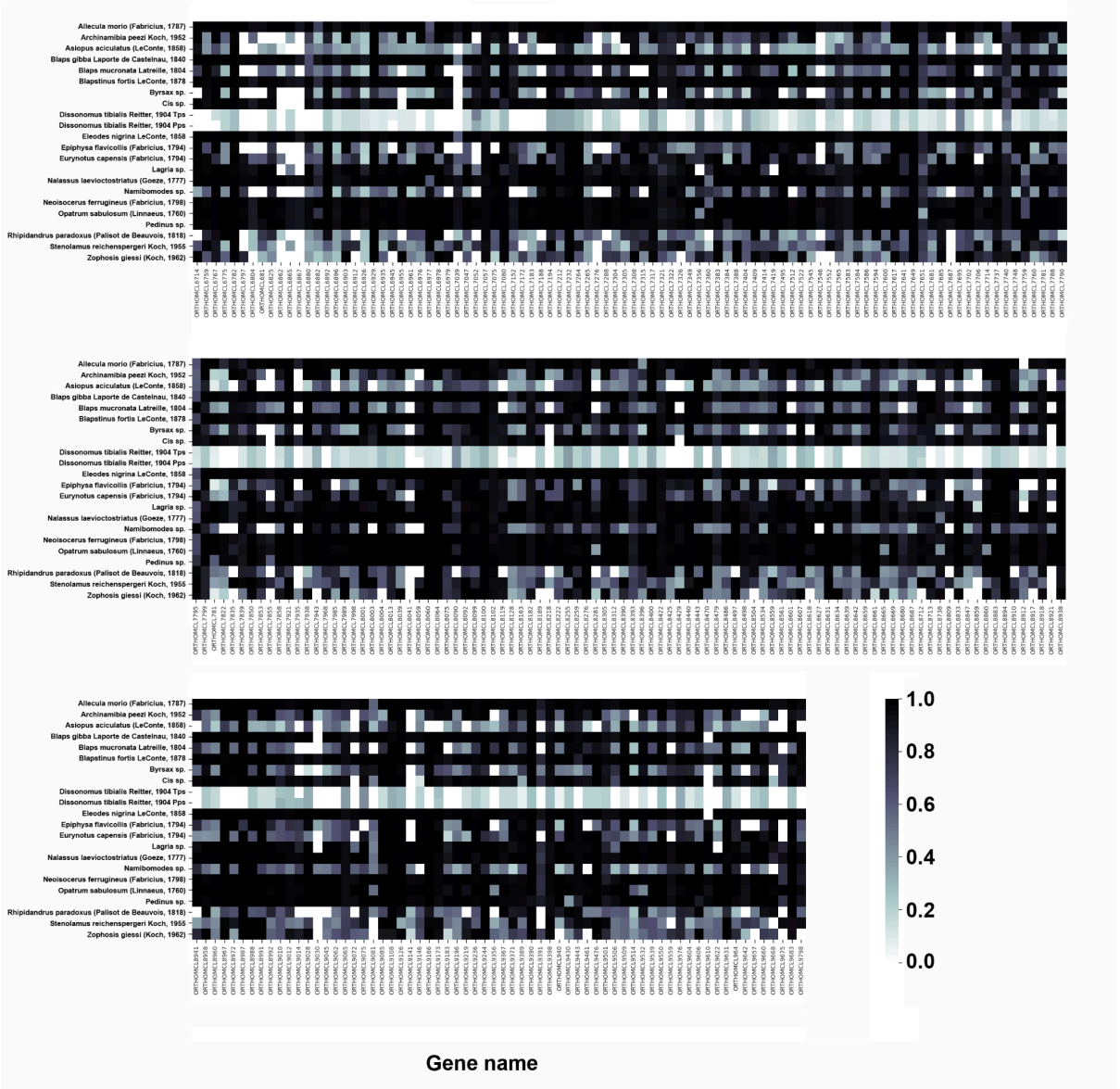


Table S1. List of recovered loci for the museum sample of *Dissonomus tibialis* Reitter, 1904. Tps – sample mapped to Tenebrionidae probeset sequences; Pps – sample mapped to Pimeliinae probeset sequences.

Gene code	**Annotation**	**No. of recovered amino acids (*Tps*)**	**No. of recovered amino acids (*Pps*)**
OrthoMCL4214	Short transient receptor potential channel	163	163
OrthoMCL4501	Heterogeneous nuclear ribonucleoprotein K	173	169
OrthoMCL4765	EGF region	98	98
OrthoMCL6279	expressed protein	175	171

OrthoMCL7052	Hexokinase-2	140	138
OrthoMCL7356	hypothetical protein	140	140
OrthoMCL7740	protocadherin 11	103	101
OrthoMCL7943	Protein Smaug	109	104
OrthoMCL8090	Ras-related protein Rab-4B	117	113
OrthoMCL8255	phosphoglucomutase 2	106	102
OrthoMCL8400	Nucleolar protein 5A	137	137
OrthoMCL8479	LUC7	113	111
OrthoMCL8559	Regulator of G-protein signaling 4	97	97
OrthoMCL9065	Rhomboid-related protein 1	160	160
OrthoMCL9075	conserved hypothetical protein	216	211