Additional datasets from this study can be found in the following repository:

Wöhner, T., Emeriewen, O. F.,Wittenberg, A. H. J., Nijbroek, K.,Wang, R. P., Blom, Q5

E.-J., et al. (2023). Data set: the structure of the tetraploid sour cherry ‘Schattenmorelle’

(Prunus cerasus L.) genome reveals insights into its segmental allopolyploid nature.

OpenAgrar Repository. doi: 10.5073/20230324-105730-0