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Mitogenome Announcement

Complete mitochondrial genome of the larch hawk moth, *Sphinx morio* (Lepidoptera: Sphingidae)

Min Jee Kim, Sei-Woong Choi & Iksoo Kim ✉

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Abstract

The larch hawk moth, Sphinx morio, belongs to the lepidopteran family Sphingidae that has long been studied as a family of model insects in a diverse field. In this study, we describe the complete mitochondrial genome (mitogenome) sequences of the species in terms of general genomic features and characteristic short repetitive sequences found in the ... typical set of genes (1... the major non-coding A... The 316-bp-long seque... served seque... y, the A+T-rich regi... are rarely found in ... andem repeat, and six nearly identical 5-6 bp long repeat sequences.

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