

# 3D model related to the publication: New data on the Miocene dormouse *Simplomys* García-Paredes, 2009 from the peri-alpin basins of Switzerland and Germany: palaeodiversity of a rare genus in Central Europe

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### **Abstract**

This contribution contains the 3D model of the holotype of *Simplomys hugi*, the new dormouse species from the locality of Glovelier described and figured in the following publication: New data on the Miocene dormouse *Simplomys* García-Paredes, 2009 from the peri-alpin basins of Switzerland and Germany: palaeodiversity of a rare genus in Central Europe. https://doi.org/10.1007/s12549-018-0339-y

**Keywords:** Early Miocene, Gliridae, Maxilla, Simplomys, Switzerland

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# **Specimen**

### Description

Simplomys hugi (MJSN-GLM017-0001)

Left maxilla with DP4, P4, M1 and M2 teeth

**Table 1.** The new glirid species (collection: Jurassica Museum).

# INTRODUCTION

The genus Simplomys, a dormouse with a simple dental morphology, was erected by García-Paredes in García-Paredes et al. (2009). The 3D model presented here has been described by Prieto et al. (2018), and is the holotype of the newly described taxon Simplomys hugi, from the Early Miocene of Glovelier, Switzerland. The left Maxilla MJSN-GLM017-0001 (Fig. 1) shows the decidual fourth premolar (DP4) with the unerupted permanent premolar (P4) just below, and first and second molars (M1 and M2); it is deposited in the JURASSICA Museum in Porrentruy (Canton of Jura, Switzerland). This new mediumsized Simplomys is unique by its weak precingulum which is always present on M1 and never found in any other glirids. This genus shows a continuous evolution in Spain during the early Miocene, at the end of the Ramblian and up to the middle Aragonian (García-Paredes et al. 2009, 2016), but its record is in contrast extremely scarce in the corresponding interval in central Europe. Thus, this specimen is of particular importance for the evolution of Simplomys, but also for the understanding of faunal exchanges in Europe.

## **METHODS**

The 3D model was produced by using the Bruker Skyscan 2211 CT scanner in University of Fribourg. It was CT-scanned with beam energy of 180 kV and a flux of 90  $\mu$ A at a detector resolution of 2.5  $\mu$ m per pixel using a 360° rotation with a step

size of  $0.2^{\circ}$  and an aluminum filter. The 3D reconstruction was produced in Amira 6.0 (FEI).

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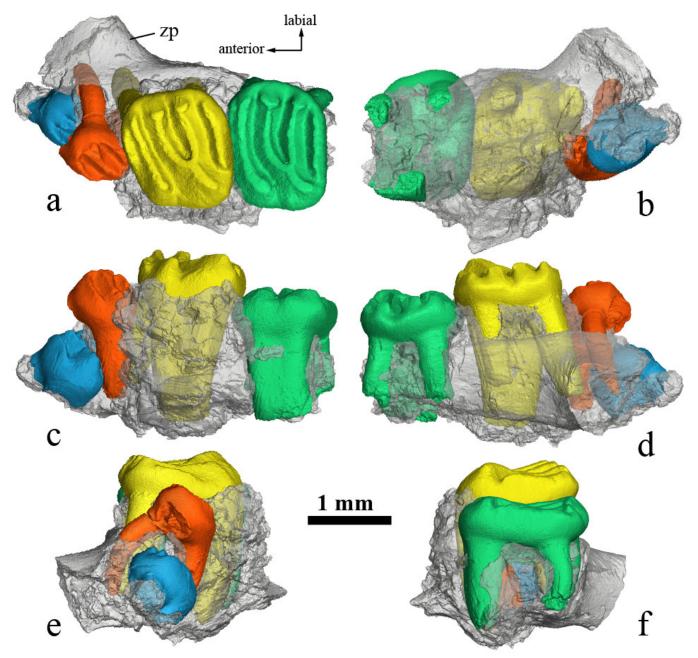
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**Figure 1.** Maxilla MJSN-GLM017-0001, the holotype of *Simplomys hugi* with DP4 and P4-M2 in occlusal (a), dorsal (b), lingual (c), labial (d), anterior (e) and posterior (f) views. (blue, P4; red, DP4; yellow, M1; green, M2; zp, zygomatic process)

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