

First records of the entomopathogenic fungus
Ophiocordyceps variabilis (Petch) G.H. Sung, J.M. Sung,
Hywel-Jones et Spatafora from Siberia

Первые сведения об энтомопаразитическом грибе
Ophiocordyceps variabilis (Petch) G.H. Sung, J.M. Sung,
Hywel-Jones et Spatafora в Сибири

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Abstract. The entomopathogenic fungus *Ophiocordyceps variabilis* (Petch) G.H. Sung, J.M. Sung, Hywel-Jones et Spatafora is recorded from Siberia for the first time. The fungus was found in the mixed forests of Novosibirsk city (54.8° N, 83.1° E) and in the Yugansky Reserve (59.2–60.3° N, 73.5–74.0° E) on larvae of Xylophagidae (Diptera) in dead fallen wood. The fungi were genotyped using *TEF-1α* gene sequence analysis and they were 99–100 % identical to the North American *O. variabilis*.

Резюме. Впервые в Сибири обнаружен вид энтомопаразитического гриба *Ophiocordyceps variabilis* (Petch) G.H. Sung, J.M. Sung, Hywel-Jones et Spatafora. Грибы обнаружены в смешанных лесах окр. г. Новосибирск (54,4° с.ш., 83,6° в.д.) и в Юганском заповеднике (59,2–60,3° с.ш., 73,5–74,0° в.д.) на личинках мух ксилофагид (*Xylophagidae*, *Diptera*) в валежной древесине. Анализ последовательностей региона 52 *TEF-1α* показал 99–100 % сходство Сибирских изолятов *O. variabilis* с экземплярами из Северной Америки.

Entomopathogenic ascomycetes are natural resource for the development of bioinsecticidal and pharmacological products. The species diversity of these organisms in Russia, and especially in Siberia, is poorly studied. This lack of information is related to the locality of their distribution, which is caused by the continental climate and is not optimal for the development of these fungi. That have been sporadic studies of these fungi

in Siberia [Ogarkov, Ogarkova, 2000; Gorbunova et al., 2011; Kryukov et al., 2011, 2018].

We found ascomycetes *Ophiocordyceps variabilis* in the territory of Novosibirsk Academic town and the Yugansky Reserve from 2009 to 2016. The morphological characteristics of the collected samples conformed to those previously described for this species [Beug et al., 2014] (Fig. 1). In Novosibirsk Academic Town, the fungi were found in July and August in mixed forest with a predominance of spruce, aspen, birch and pine in the overstory and *Pteridium aquilinum* (L.) Kuhn and *Aegopodium* L. in the grass layer. In the Yugansky Reserve, these ascomycetes were found in September in mature mixed coniferous forest with relatively old, falling aspen trees. The insects infected by the fungi were located in dead wood (*Populus* sp.) that was overgrown with mosses. The hosts of the fungi were xylophagid flies (Diptera, Xylophagidae).

The sample identification was carried out by sequencing the region of the gene for the elongation factor (*TEF-1α*) [Sung et al., 2007]. The total DNA was isolated from fresh stromata. The isolation of the DNA, amplification and purification of the product were carried out as previously described [Kryukov et al., 2018]. The phylogenetic analysis was based on a comparison (700 bp) of the Siberian samples (Ak-2, Ak-4), a sample collected in the Far East (Obh) and sequences from

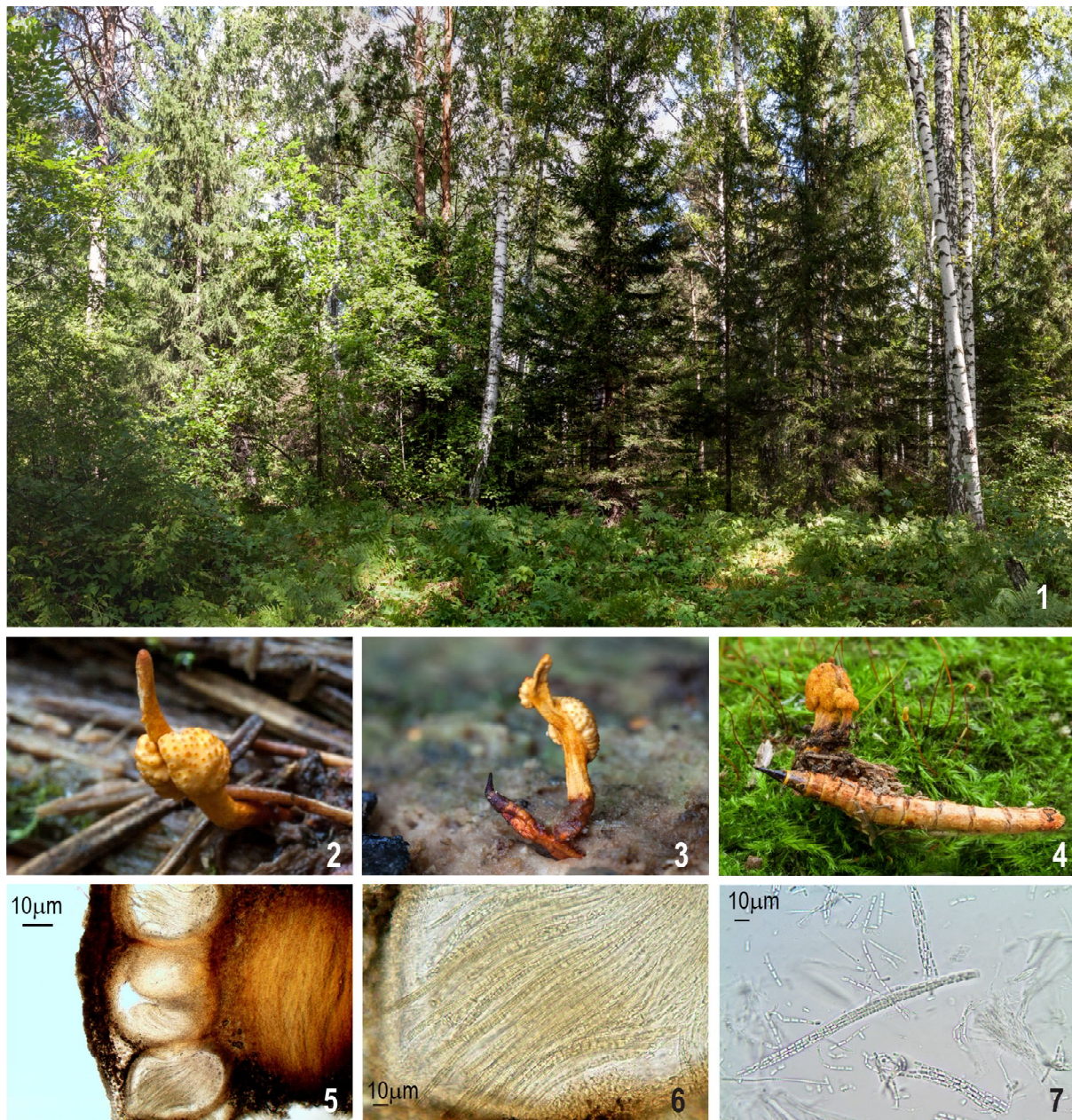


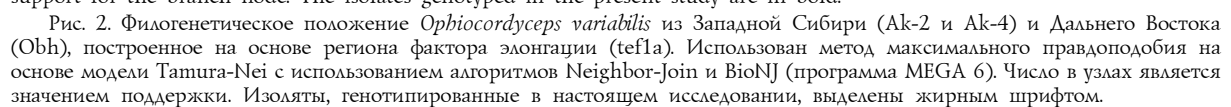
Fig. 1. Entomopathogenic fungus *Ophiocordyceps variabilis*. 1 — habitat (Novosibirskaya Oblast), 2–4 — stromata, 5–6 — perithecia with ascospores, 7 — ascospores. Photos by D.V. Ageev, and T.M. Bulyonkova.

Рис. 1. Энтомопатогенный гриб *Ophiocordyceps variabilis*. 1 — биоценоз (окр. г. Новосибирск), 2–4 — стромы, 5–6 — перитеции с аскоспорами, 7 — аскоспоры. Фотографии Д.В. Агеева и Т.М. Бульонковой.

GenBank. The Siberian and Far East samples were divided into two haplotypes with an identity of 99 % (Fig. 2). The Siberian samples were 100 % identical to the North American strain ARSEF 5365. The Far East sample was 99.39 % identical to the strain OSC 111003, which was also collected in North America [R. Kepler, personal communication].

O. variabilis was first described by Petch in 1937 as *Cordyceps variabilis*. The fungus is a highly specialized parasite of Xylophagidae [Hodge et al., 1998]. Previously, some authors indicated that the larvae of beetles were the hosts of this pathogen [Petch, 1937; Mains,

1958]. However, by checking collected samples, it was determined that the hosts are diptera larvae [Hodge et al., 1998]. The fungus anamorph was described by Hodge and coauthors [1998], and it was classified as *Syngliocladium*. The authors isolated this anamorph from both *O. variabilis* (*C. variabilis*) ascospores and from infected insects (*Tetanops myopaeformis*) collected in the field. However, this description was based only on morphocultural analysis. We were unable to isolate this fungus in a culture and obtain its anamorphic stage. Further study be focused on interaction between *O. variabilis* and hosts.



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