**Hansenula alni**, a New Heterothallic Species of Yeast from Exudates of Alder Trees

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A novel member of the yeast genus *Hansenula* was recovered three times in 1968 from slime exudates of *Alnus rubra* in the state of Washington and in the province of British Columbia. The new species is named *Hansenula alni* because of its specific habitat in exudates of alder trees. This species occurs naturally in the haploid condition. Upon mixing of compatible mating types, conjugation occurs, and up to four hat-shaped spores are produced in zygotes or in diploid cells arising from zygotes. *H. alni* resembles *H. canadensis* (syn.: *H. wingei*) but differs from it in habitat and in ability to grow at 37°C; furthermore, the deoxyribonucleic acid base composition of the former is approximately 1.5 mol% higher than that of the latter. The type strain of *H. alni* is UCD-FS&T 68-928A (= ATCC 36594 = CBS 6986). Its compatible mating type is UCD-FS&T 68-940 (= ATCC 36595 = CBS 6987).

Many tree species show the phenomenon of fluxing, the flowing of tree sap from a wound (4). Tree fluxes often persist for long periods of time, and the flowing sap becomes heavily colonized with yeasts and bacteria that are introduced by visiting insects such as drosophilae. The tree sap usually assumes a thick, slimy consistency (probably caused by microbial polysaccharide formation) and is therefore often referred to as a “slime flux.”

In 1972 Phaff et al. (4) reported the isolation of three yeast strains, tentatively identified as yeast resembling *Hansenula wingei* Wickerham, from slime fluxes of *Alnus rubra* Bongard in the Pacific Northwest of North America. The three strains were isolated from widely separated trees and represented haploid, heterothallic mating types that sporulated well upon mixing of appropriate strains. Doubts about the identity of the three strains with *H. wingei* arose when it was found that the mating types of the isolates mated poorly with those of the heterothallic *H. wingei* and no spores were evident (4). Furthermore, *H. wingei* occurs exclusively in insect frass of coniferous trees (mainly *Picea*), and the guanine-plus-cytosine (G+C) content of its nuclear deoxyribonucleic acid (DNA) was significantly lower than that of the strains from *Alnus* (2).

In the accompanying paper (2), we have shown that there is little complementarity between the DNA from *H. wingei* and that of the strains from alders. Consequently, we regard these strains as belonging to a new species of *Hansenula*, for which the name *H. alni* (L. noun *Alnus* generic name of the alder; L. gen. noun *alni* of alder) is proposed because its habitat is the exudate of alder trees.

**MATERIALS AND METHODS**

Samples of slime fluxes were collected during June and July of 1968 in various locations of Alaska, the Yukon Territory, British Columbia, Washington, Oregon, and California (4). The precise locations of the samples from *Alnus* which contained the new species of *Hansenula* are given in Table 1.

The samples were collected in new plastic vials or bags. Usually within 6 to 18 h after collection, a loopful of the slimy exudate was streaked directly on 5% malt agar acidified to pH 3.7 with a precalculated amount of 1 N HCl. The plates were stored at ambient temperatures (ca. 15 to 25°C) until colonies appeared. In spite of the low pH, various numbers of bacterial colonies usually appeared together with those of yeasts. After 3 to 6 days, the plates were inspected with a dissecting microscope, and, for identification purposes, one yeast colony of each morphological type was brought into pure culture by two successive platings on malt agar. For further details on the collecting areas, see reference 4.

Identification of the isolates involved characters determined by standard methods currently used in yeast taxonomy (8). DNA extraction and purification were done by a combination of the procedures of Marmur (3) and Bernardi et al. (1) as described by Price et al. (5). The G+C content of the DNA was calculated from buoyant density values in cesium chloride (6, 7) and was based on three separate determinations. *Micrococcus lysodeikticus* DNA, with a buoyant density of 1.7311 g/ml, was used as a reference. The buoyant density of the *M. lysodeikticus* DNA was derived from comparison with plasmid-free *Escherichia coli* K-12 DNA, whose buoyant density was taken to be 1.7100 g/ml.
RESULTS

Latin diagnosis of Hansenula alni sp. nov.

In extracto mali post dies 3 cellulae ovoideae, (2.4-4.6) × (3.0-5.4) μm, singulae, binae, aut in catenis brevis; sedimentum, annulus et pellicula tenuis, non-nitida formantur. Post dies 21 cellula crassa et sedimentum abundum.

Cultura in agaro mali post unem mensem (18°C) creamea, mollis, nitida et glabra, convexa, margine piloso.

In agaro farinae Zea mays post dies 10 pseudomycelium, blastoconidia et blastosporae abundat.

Species heterothallica, haploidea. Oriuntur asci ex conjugatione inter cellulas sexus contrariae. Pileiformae ascosporae formantur ad 4 in quoque asco; asci rumpunter post 5-7 dies.

Fermentatio glucosi nullum. Glucosum, maltosum, saccharum, celllobiosum, trehalosum, melitosum, amyllum solubile (exigue), D-xylom, L-rhamnun, ethanolum, glycerolum, D-mannitolum, D-glucitololum, α-methyl-D-glucosidum, salicinum, glucono-D-lactonum, kalium gluconatum (lente), acidum lacticum (exigue), acidum succinicum, et acidum citricum assimilantur. Acidum aceticum, acidum butyricum, acidovalericum, acidum imide per, acidum propionicum, acidum caprinum, acidum aceticum, acidum lacticum (exigue), acidum succinicum, et acidum citricum assimilantur. D-Galactose, L-sorbose, lactose, melibiose, raffinose, inulinum, L-mabinose, D-arabinose, D-ribose, salicinum, a-methyl-D-glucoside, salicinum, glucono-6-lactonum-, kalium gluconatum (lente), acidum lacticum (exigue), acidum succinicum, et acidum citricum assimilantur at non D-galactosum, L-sorbitosum, lactosum, melibiosum, raffinosum, inulinum, L-arabinosum, D-arabinosum, D-ribose, methanol, i-erythritolum, ribitolom, galactitolum, meso-inositol, et hexadecane are not assimilated.

Kalium nitricum, natrium nitrosum, et ethyl amonium assimilantur.

Ad crescentiam thiaminum et pyridoxinum necessaria sunt.

Crescere potest in 33°C; nullum incrementum calore 35°C. G+C acidi deoxyribonucleati 41.8-42.0 mol% (3 stirpes, vide tabulum 1).

Habitatio in exudati A. rubrae Canadensis isolata est.

Cultura auxiliaris UCS-FS&T 68-940 (= CBS 6986) ex exudato A. rubrae Canadensis isolata est.

TPUS: Stirps UCD-FS&T 68-928A

In collectione zymotica Centraalbureau voor Schimmelcultures, Delphi Batavorum sub no. 6986 deposita est.

Cultura in agaro malti post unem mensem (18°C) creamea, mollis, nitida et glabra, convexa, margine piloso. After 1 month, the streak culture is cream-colored to white, smooth, shiny, soft, and convex in cross-section; the border is fringed with pseudomycelia.

Dalmau plate culture on corn meal agar. After 9 days, a well-developed pseudomycelium with abundant blastospores and blastoconidia is present. True hyphae are lacking.

Formation of ascospores. Vegetative cells are haploid and heterothallic. Upon mixing of compatible strains, conjugation takes place; one of the two cells forming the zygote increases considerably in size (Fig. 1a to c). Ascospores usually develop in diploid cells arising from the zygote and rarely in the zygote itself. Two to four hat-shaped spores are produced per ascus (Fig. 1d to g); the asci gradually lyse and release the spores in groups of two, three, or four (Fig. 1e). Good sporulation was observed on malt agar after ca. 6 to 10 days. This yeast has not been isolated from nature in the diploid state.

Other characteristics.

Fermentation: Glucose is not fermented. Assimilation of carbon compounds: D-Glucose, maltose, sucrose, celllobiose, trehalose (latent), melezitose, soluble starch (weak), D-xylom, L-rhamnun, ethanol, glycerol, D-mannitol, D-glucitol, α-methyl-D-glucoside, salicin, glucono-D-lactone, potassium gluconate, DL-lactic acid (weak), succinic acid, and citric acid are assimilated. D-Galactose, L-sorbose, lactose, melibiose, raffinose, inulin, L-arabinose, D-arabinose, D-ribose, methanol, i-erythritolum, ribitol, galactitol, 2- and 5-ketogluconatum, meso-inositol, glucosamine, and hexadecane are not assimilated.

Assimilation of nitrogen compounds: Potassium nitratum, +; sodium nitritum, +; ethyl amine, +.

Vitamin requirements: Thiaminum et pyridoxinum essential for growth.

Growth on 50% (wt/wt) glucose–yeast extract agar: Absent.

Growth on 10% (wt/vol) sodium chloride–yeast extract agar: Absent.

Maximum temperature for growth: 33°C.


G+C content of the nuclear DNA: 41.8 to 42.0 mol% (range for three strains, see Table 1).

Type strain. The type strain, UCD-FS&T 68-928A, was isolated from the slime flux of an alder tree (A. rubra Bongard) along highway 7, 10 miles (ca. 16.1 km) west of Agassiz, British Columbia. The auxiliary strain giving the best sporulation (UCD-FS&T 68-940, of opposite sex
to 68-928A) was isolated from the slime flux of an alder tree 2 miles (ca. 3.2 km) south of Parksville, Vancouver Island, British Columbia. Strains 68-928A and 68-940 have been deposited in the collection of the Yeast Division of the Centraalbureau voor Schimmelmicrobiologie in Delft, The Netherlands as CBS 6986 and 6987, respectively, and in the American Type Culture Collection, Rockville, Md., as ATCC 36594 and 36595, respectively.

**Sources.** Table 1 lists the substrates, localities, mating types, and G+C contents of the nuclear DNAs of the three strains. It appears that *A. rubra* is the host tree for *H. alni*.

**TABLE 1.** Localities, sources, mating types, and G+C contents of nuclear DNAs of isolates of *Hansenula alni* and phenotypically similar species of *Hansenula*

<table>
<thead>
<tr>
<th>UCD-FS&amp;T(^c) strain designation</th>
<th>Habitat (substrate)</th>
<th>Mating type</th>
<th>G+C of nuclear DNA (mol% ± SD(^d))</th>
</tr>
</thead>
<tbody>
<tr>
<td><em>Hansenula alni</em> 68-928A(^a)</td>
<td>Slime flux from <em>Alnus</em>(^d)</td>
<td>α</td>
<td>42.0 ± 0.36</td>
</tr>
<tr>
<td><em>H. alni</em> 68-940</td>
<td>Slime flux from <em>Alnus</em>(^c)</td>
<td>a</td>
<td>41.9 ± 0.09</td>
</tr>
<tr>
<td><em>H. alni</em> 68-985</td>
<td>Slime flux from <em>Alnus</em>(^c)</td>
<td>α</td>
<td>41.8 ± 0.25</td>
</tr>
<tr>
<td><em>H. wingei</em> 71-80(^c)</td>
<td>Frass from dead conifer</td>
<td>Heterothallic</td>
<td>40.5 ± 0.25</td>
</tr>
<tr>
<td><em>H. canadensis</em> 74-71(^c)</td>
<td>Frass from <em>Pinus resinosa</em></td>
<td>Homothallic</td>
<td>40.8 ± 0.12</td>
</tr>
</tbody>
</table>

\(^a\) UCD-FS&T, University of California, Davis, Department of Food Science and Technology.

\(^b\) SD, Standard deviation.

\(^c\) Type strain.

\(^d\) Located on highway 7, 10 miles (ca 16.1 km) west of Agassiz, British Columbia.

\(^e\) Located 2 miles (ca. 3.2 km) south of Parksville, Vancouver Island, British Columbia.

\(^f\) Located along Hoh Road, Olympic National Park, state of Washington.

**Fig. 1.** (a) Mixture of *H. alni* mating types 68-928A and 68-940 on malt agar after 2 days. Note zygote formation. (b) Single zygote; one of the conjugated cells has increased in size. (c) Variously shaped zygotes. (d and f) Ascii containing four and three hat-shaped ascospores, respectively. (e) Ascus and four dehisced ascospores after 7 days on malt agar. (g) Lateral view of a single dehisced ascospore. Bar represents 4 μm in all cases.
DISCUSSION

When H. alni was first isolated (4), its colonies on malt agar were smooth and glistening. After storage on slants over a period of several years, a high proportion of the colonies had become mat and rugose, probably as the result of mutation. This phenomenon was minimal when the strains were stored in the freeze-dried state.

H. alni resembles H. canadensis in many of its phenotypic properties. It differs from that species in producing four, rather than two, spores per ascus; in occurring in nature as haploid, heterothallic cells rather than as diploid homothallic cells; in habitat (in exudates of a broad-leaf tree rather than in bark beetle frass of coniferous trees); and in that the G+C content of its nuclear DNA is about 1.5 mol% higher than that of H. canadensis (Table 1).

The three isolates of H. alni were initially regarded as possible strains of H. wingei (4). The molecular taxonomy studies reported in the accompanying paper (2) have shown that H. wingei and H. canadensis share close to 80% of their DNA sequences, and therefore their names are synonyms. H. alni, on the other hand, showed very low DNA complementarity with these organisms as well as with others considered closely related to H. canadensis by Wickerham (9). H. alni thus constitutes a valid new species, and a description of it is provided in this paper.

Because of the many phenotypic properties in common between H. alni and H. canadensis, it may be postulated that H. wingei or H. canadensis is a progenitor of H. alni and that the latter diverged by adapting to a new habitat, the sap of a deciduous tree, A. rubra. It is not known whether H. alni occurs in nature as haploid cells rather than in diploid form. The three haploid strains isolated came from exudates of three different alder trees located at least 80 km apart (Table 1).

Aside from the difference in nuclear DNA base composition, the only phenotypic properties that can be used to differentiate H. alni from H. canadensis (syn.: H. wingei) are habitat and inability to grow at 37°C. H. alni occurs in exudates of alder trees, and its maximum temperature for growth is ca. 33°C, whereas H. canadensis has been isolated only from beetle-infested coniferous trees and is capable of growing at 37°C.

ACKNOWLEDGMENTS

We are greatly indebted to Nancy Krauter for her technical assistance in determining the DNA base compositions.

This research was supported, in part, by Public Health Service grant GM-16307-08 from the National Institute of General Medical Sciences to H. J. Phaff.

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LITERATURE CITED