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# Chemotaxonomy of the genus Talaromyces

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### Abstract

Species of the ascomycetous genus *Talaromyces* have been examined for profiles of secondary metabolites on TLC. The greatest number of specific metabolites were produced on oatmeal-, malt extract- and yeast-extract sucrose agars. Profiles of intracellular secondary metabolites produced on oatmeal agar were specific for each species and provided a means of simple differentiation of the taxa. Examination of the most important species using high performence liquid chromatography (HPLC) allowed to solve some taxonomic problems. Known mycotoxins are produced by *T. stipitatus* (duclauxin, talaromycins, botryodiploidin), *T. stipitatus* chemotype II (emodin), *T. panasenkoi* (spiculisporic acid), *T. trachyspermus* (spiculisporic acid), *T. macrosporus* (duclauxin) and *T. wortmannii* (rugulosin). Wortmannin is produced by an atypical strain of *T. flavus* but not *T. wortmannii*. Several other secondary metabolites were discovered for the first time in the following species: Glauconic acid is produced by *T. panasenkoi*, *T. ohiensis* and *T. trachyspermus*; vermiculine by *T. ohiensis*; duclauxin by *T. flavus* var. *macrosporus* and the mitorubrins by *T. flavus* and *T. udagawae*. The profiles of secondary metabolites support the established taxonomy of the species based on morphology, showing the genetic stability of profiles of secondary metabolites in *Talaromyces*. Two new taxa are proposed: *T. macrosporus* comb. nov. (stat. anam. *Penicillium macrosporum* stat. nov.), and *Penicillium vonarxii*, sp. nov. for the anamorph of *T. luteus*.

### Introduction

Since the monographic treatment by Stolk & Samson (1972) the species delimitation in the ascomycetous genus *Talaromyces* has been generally accepted, except for the classification of certain species in either *Talaromyces*, *Hamigera* or *Byssochlamys* (Stolk & Samson 1971 and 1972; Samson & Abdel-Fattah 1978; Pitt 1980; Pitt & Hocking 1979; von Arx 1986). These genera contain species with heat-resistant ascospores (Hocking & Pitt 1984) and some species are known mycotoxin producers (Frisvad 1986). The stability of *Talaromyces* taxonomy contrasts with the different views in *Eupen*- *icillium* and *Penicillium* taxonomy (Stolk & Samson 1983; Pitt 1980; Ramirez 1982; Frisvad 1986). Species of *Talaromyces* should therefore provide good test material for the hypothesis that profiles of secondary metabolites are fundamental taxonomic criteria in filamentous fungi.

In this paper the results of the analysis of secondary metabolites of the genus are presented and the taxonomic implications discussed.

### Materials and methods

Isolates in good condition of all species in Talaro-

*myces* (Table 2) were grown on Czapek yeast-extract agar (CYA), malt extract agar (MEA), yeastextract sucrose agar (YES) agar, and oatmeal agar (OA) (for formulations see Samson & Pitt 1985; Frisvad & Filtenborg 1983). Trace metals were added to all media (Frisvad & Filtenborg 1983) and the yeast extract and peptone ingredients were from Difco. The fungi were incubated at 25° C and 30° C in the dark and examined after 1 and 2 weeks.

Representative and authentic strains of all species in Talaromyces were examined for secondary metabolites by simple thin-layer chromatographic techniques (Filtenborg & Frisvad 1980; Filtenborg et al. 1983): Extracellular metabolites were determined by applying of superimposed agar plugs of YES and MEA agar; intracellular metabolites were determined by applying of superimposed extracted mycelial plugs of YES plus CYA agar, MEA plus OA or either MEA or OA agar alone on silica gel plates. The plates were developed in toluene: ethylacetate: 90% formic acid (5/4/1, v/v/v) (TEF) and chloroform: acetone: 2-propanol (85/ 15/20, v/v/v) (CAP) with griseofulvin as an external standard. All elutions were allowed to migrate 15 cm from the application line. Plates eluted in TEF were treated with cold 48% sulphuric acid and afterwards with anisaldehyde spray (Frisvad & Filtenborg 1983) and heated at 130°C for 8 minutes. Plates eluted in CAP were treated with cerium sulphate spray (Filtenborg et al. 1983). The plates were examined in daylight and under UV light (254 and 366 nm) before and after all treatments. Liquid chromatography (HPLC) was performed according to Frisvad & Thrane (1987) using a Hewlett Packard diode array detector (DAD) to obtain UV-VIS spectra of all eluting compounds and an alkylphenone retention index system. A series of standards were used to confirm the identity of some of the secondary metabolites detected. These standards are listed in Table 1. For HPLC analysis the contents of five 9cm diam Petri dishes each of MEA and OA, with 14 day old cultures grown at 25°C, were placed in a plastic bag together with 100 ml chloroform/methanol (2:1, v/v) and extracted for 3 min in a Colworth Stomacher 400. After filtration through a hydrophobic filter (Whatman 1PS), the remaining agar and mycelium was reextracted with 100 ml ethyl acetate containing 1 ml 85% phosphoric acid. The filtered water free organic phases were combined and evaporated in vacuo. Further details of the procedure are described by Frisvad & Thrane (1987).

## **Results and discussion**

### Production of secondary metabolites

All species in *Talaromyces* produced high numbers of secondary metabolites making a direct identification of each species based on TLC profiles of coloured spots possible. The HPLC results showed that each species produced chromophore families

*Table 1.* Retention indices for secondary metabolites standards in the study of *Talaromyces*.

Secondary metabolite	Retention index (RI)
Alternariol	937
Alternariol monomethylether	1070
Catenarin	1191
Chrysophanol	1240
Citreoviridin	1028
Dipicolinic acid	670
Duclauxin	1133
Emodin	1130
Erythroglaucin	1439
Flavoglaucin	1538
Glauconic acid	910
Helminthosporin	1325
Islandicin	1354
Luteoskyrin	1240
Mitorubrin	1083
Mitorubrinic acid	924
Mitorubrinol	928
Mitorubrinolacetate	1052
Monorden	917
Naphthalic anhydride	1432
Purpurogenone	1184
Rubratoxin B	1071
Rugulosin	1123
Rugulovasine A	716
Secalonic acid D	1165
Skyrin	1333
Stipitatic acid	676
Vermiculine	834
Vermicelline	962
Wortmannin	938

specific for one taxon and some chromophore families which were shared by two or more taxa.

All species of Talaromyces produced many yellow, orange and red metabolites. Some of these metabolites were members of the known anthraquinone chemosyndrome (biosynthetic family) centered around emodin, skyrin, rugulosin, catenarin and erythroglaucin (T. wortmannii and chemotype II of T. stipitatus). The mitorubrin chemosyndrome was restricted to T. flavus, T. macrosporus, T. udagawae, T. mimosinus and T. wortmannii. The variety of members of the mitorubrin biosynthetic family differed from species to species and were most diverse in T. wortmannii. Other chromophore families with UV-vis spectra containing strong absorptions from 400 to 450 nm were major coloured products of species like T. helicus and T. luteus, so the macroscopically visible yellow pigments in the mycelium and ascomata of these species are caused by different biosynthetic families of compounds. Weakly coloured secondary metabolites such as duclauxin are present in only two taxa: T. stipitatus and T. flavus var. macrosporus, while the closely related bacillosporins were only present in another heat-resistant fungus T. bacillisporus. These three heat-resistant species characteristically contained these closely related phenalenones in large amounts and as dominant secondary metabolites, indicating that they may be a major protective principle in heat resistance.

While stipitatic acid was only produced by T. stipitatus, glauconic acid was detected for the first time in T. panasenkoi, T. ohiensis and T. trachyspermus indicating a close chemotaxonomic relationship between the latter three species. This confirms the morphological observations by Stolk and Samson (1972), who found that these species produce similar ascomatal initials. Vermiculine was found in T. ohiensis and the production of this secondary metabolite was confirmed for T. flavus, but the production of vermiculine in T. wortmannii (Jones et al. 1984) could not be confirmed. All the secondary metabolites mentioned above have also been found in one or more species of Penicillium subgenus Biverticillium (Turner 1971; Turner & Aldridge 1983; Frisvad 1986; Mantle 1987), and this supports the close taxonomic relationship between *Talaromyces* and the anamorphic *Penicillium* subgenus *Biverticillium*.

The profiles of intracellular metabolites produced on OA, as seen on TLC plates using the agar plug method, were specific and consistent for each taxon, but with TLC alone it is difficult to assess whether these metabolites represent one or more biosynthetic families in each taxon. T. stipitatus and T. wortmannii (Table 2, see also Turner 1971; Turner & Aldridge 1983) are at least 4 to 5 known biosynthetic families and our HPLC results strongly indicate that all Talaromyces taxa produce at least four, often eight to ten chromophore families on MEA and OA. The profiles of intracellular metabolites produced on MEA were less specific as evaluated using TLC, but the technique was useful in showing which metabolites were common in different taxa. The most conspicous of these metabolites was a yellow coloured (both in daylight and UV light before and after spraying with sulphuric acid, relative Rf value to griseofulvin 0.98) substance produced by T. flavus, T. macrosporus, T. mimosinus, T. udagawae, T. wortmannii and several species in Penicillium subgenus Biverticillium. A partial spectroscopic characterisation (UV-vis and NMR) showed that this metabolite was mitorubrinic acid and this was confirmed by comparison with an authentic standard. T. rotundus, T. trachyspermus, T. assiutensis and T. ohiensis also have some metabolites in common, viz. glauconic and glaucanic acid seen on TLC (Table 2 and 3).

Mycotoxin production by Talaromyces species is of interest because these species may occur in pasteurized fruit juices (Hocking & Pitt 1984; Baggerman & Samson 1988) and other food products. T. flavus could be a potential producer of wortmannin in food, but this species is much less heat-resistant than T. flavus var. macrosporus, a producer of the mycotoxin duclauxin, Wortmannin, however, is also reported from food-borne species such as Penicillium funiculosum, Myrothecium roridum and Fusarium oxysporum (Abbas & Mirocha 1988). Very little is known on the toxicity of the metabolites of the other species in Talaromyces but emodin, cited as a toxin by Wells et al. (1975) is produced by an atypical strain of T. stipitatus (see discussion under Taxonomic implications). Other

# 182

Table 2. Species in Talaromyces, anamorph names, isolates examined and secondary metabolite production.

Teleomorph / anamorph	Isolates	Secondary metabolites
T. flavus (Klocker) Stolk & Samson /	NRRL 2098 (NT) (b)	Mitorubrin (+ acid) (a), Vermicellin, vermiculine
P. dangeardii Pitt	CBS 261.55	Mitorubrin (+ acid), Vermicellin, vermiculine
	CBS 284.58	Mitorubrin (+ acid), Vermicellin, vermiculine
	CBS 582.72A	Mitorubrin (+ acid), Vermicellin, vermiculine
	CBS 387.47	Mitorubrin (+ acid), Vermicellin, vermiculine
	Fravel 282.3	Mitorubrin (+ acid), Vermicellin, vermiculine
T macrosporus (Stolk & Samson) Frisvad	CBS 317.63 (T)	Mitorubrin (+ acid), Duclauxin
Stolk & Samson chemotype I /	CBS 117.72	Mitorubrin (+ acid), Duclauxin
P macrosporum Frisvad. Stolk & Samson	CBS 353.72	Mitorubrin (+ acid), Duclauxin
	CBS 580.72	Mitorubrin (+ acid), Duclauxin
	CBS 130.89	Mitorubrin (+ acid), Duclauxin
T macrosporus chemotype II	CBS 350.72	
T helicus (Raper & Fennell) C R Benjamin /	CBS 335 48 (T) (poor)	
<i>P</i> snirillum Pitt	CBS 137.65	
1. <i>Spiniani</i> 1 ici	CBS 760.68	
	CBS 550.72B	
	CBS 562 72	
	CBS 585 72	
T stipitatus (Thom) C B Benjamin	CBS 375 48 (T)	Stipitatic acid, Duclauxin
chemotype I / P emmonsii Pitt	CBS 292.53	Stipitatic acid, Duclauxin
enemotype 171. enumental 1 ne	CBS 189 72	Stipitatic acid, Duclauxin
	CBS 227 72	Stipitatic acid, Duclauxin
T stipitatus chemotype II	CBS 349 72	Catenarin emodin erythroglaucin
T. nanasankoj Pitt / P. nanasankoj Pitt	CBS 583 72C	Glauconic acid
1. panasenkoi 1 ht / 1. panasenkoi 1 ht	NRRI 2103	Glauconic acid
T lutaus (Zukal) C. P. Benjamin /	CBS 348 51 (NT)	Gladeonie acid
P. vonannii Erisund & Samson	CBS 533 50	
	CDS 355.57	
	CDS 005.71	
Tudagewas Stall & Somson / P. udagewas	CBS 730.74 CBS 570.72 (T)	Mitorubrin (+ acid) mitorubrinol acetate
Stolk & Samson	CBS 579.72 (1)	Mitorubini (+ acid), intorubinioi acetate
T. wortmannii (Klocker) C.R. Benjamin / P. kloeckeri Pitt	CBS 391.48 (T)	Mitorubrin (+ acid), mitorubrinol (+ acetate) Rugulosin, skyrin
1 . Riocekent i ite	CBS 235 38	Mitorubrin (+ acid) mitorubrinol (+ acetate)
		Rugulosin
	CBS 293 63	Mitorubrin $(+ \text{ acid})$ mitorubrinol $(+ \text{ acetate})$
	000 275.00	Rugulosin skyrin
	CBS 319 63	Mitorubrin (+ acid) mitorubrinol (+ acetate)
	000 515.05	Rugulosin skyrin
	CBS 387 67	Mitorubrin (+ acid) mitorubrinol (+ acetate)
	CD3 567.67	Rugulosin skyrine
	CBS 533 72	Mitorubrin (+ acid) mitorubrinol (+ acetate)
	000 555.72	Rugulosine skyrine
T rotundus (Donor & Fonnall)	CPS 360 48 (T)	Ruguloshie, skyllie
C. P. Benjamin / P. snhaerum Pitt	CBS 587 72	
T. trachysnarmus (Shear) Stolk & Samson /	CBS 373 48 (T)	Glauconic acid
P lehmanii Pitt	CBS 346 54	Glauconic acid
	CBS 282 58	Glauconic acid
	CBS 112 64	Glauconic acid
T assistancis Samson & Abdel-Fattab	CBS 147 78 (T)	Glauconic acid
(-T) assume that $T = T$ assume that $T = T$	CBS 645 80	Glauconic acid
(- 1. gossyph 1 m) / 1. ussuuense Samson & Abdel-Fattab		Saucome acia
T objensis Pitt / P objense Huana & Smith	CBS 162 67 (T)	Vermiculine, Glauconic acid
1. Onicidio Fitti I. Onicide Huang & Smith	CD0 102.07 (1)	, onnounne, Olaucome acia

Table 2. Continued.

Teleomorph / anamorph	Isolates	Secondary metabolites
T. galapagensis Samson & Mahoney /	CBS 751.74 (T)	
P. galapagense Samson & Manoney		
T. mimosinus Hocking / P. mimosinum Hocking	CBS 659.80 (T)	Mitorubrin (+ acid)
T. intermedius (Apinis) Stolk & Samson /	CBS 152.65 (T)	
P. intermedium Stolk & Samson		
T. derxii Takada & Udagawa /	NHL 2981	
P. derxii Takada & Udagawa	NHL 2982	
T. bacillisporus (Swift) C.R. Benjamin / G. swiftii Pitt	CBS 296.48 (T)	
T. byssochlamydoides Stolk & Samson /	CBS 413.71 (T)	
Paecilomyces byssochlamydoides Stolk & Samson	CBS 533.71	
T. emersonii Stolk / Penicillium emersonii Stolk	CBS 393.64 (T)	
	CBS 814.70	
	IMI 116825ii	
	CBS 204 75	
	CBS 373 75	
T thermonhilus Stolk / P. dupontii	CBS 236 58 (T)	
Griffin & Maublanc	CBS 116 72	
T numuraus (E. Millor & Deche Aug) Stallt &	CDS 110.72 CDS 475 71 (T)	
Samson / P. purpureum Stolk & Samson	CBS 475.71 (1)	
T. leycettanus Evans et Stolk / Paecilomyces leycettanus (Evans & Stolk) Samson & Evans	CBS 398.68 (T)	

(a) T = type culture, NT = neotype culture

(b) (+ acid) = mitorubrinic acid, (+ acetate) = mitorubrinol acetate.

mycotoxin producers in *Talaromyces* are *T. stipitatus*, producing duclauxin (Mantle 1987) talaromycins (Phillips et al. 1987), and botryodiploidin (Fuska et al. 1988), *T. macrosporus* producing duclauxin (this paper) and *T. trachyspermus* producing spiculisporic acid (Fujimoto et al. 1988)

Strains of *T. flavus* have been applied as biological control agents of *Verticillium* wilt of potato (Davis et al. 1986; Fravel et al. 1986). One of the isolates (Fravel 282.3) was examined and found to produce mitorubrin, mitorubrinic acid, and vermicellin, typical products of *T. flavus*. Fig. 1 shows a HPLC trace of this strain grown on MEA and OA. Kim et al. (1988) identified glucose oxidase as a metabolite that mediated inhibition of *Verticillium dahliae*. However, other secondary metabolites detected by us could play also a role (vermicellin). Another metabolite (retention time 28.6 min), produced in quite large amounts on MEA and OA shown in Fig. 1 seems to agree well with talaron, a fungal inhibitor (Mizuno et al. 1974). All known species in *Talaromyces* are homothallic, with the exception of a recently described heterothallic taxon, *T. derxii* Takada & Udagawa (1988). The extract of this fungus was green, (like that of *T. bacillisporus*), and both A and a strains and mated A + a strains produced the same characteristic profile of secondary metabolites. The major products in this fungus has not been found in any other species of *Talaromyces* or the related anamorphs *Penicillium* of subgenus *Biverticillium*.

#### Taxonomic implications

Table 3 lists the species accepted in *Talaromyces*, their anamorphic states and representative cultures chosen for the profile of secondary metabolites.

Raper & Thom (1949) placed NRRL 2103 hesistantly in *T. luteus*. The TLC trace of this strain was typical of *T. panasenkoi* and our morphological reexamination of NRRL 2103 showed that this

184
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Species	Specific profile of spots on TLC plates (a)	Production of secondary metabolites including mycotoxins
T. flavus	++	I: Mitorubrin (c,d), mitorubrinic acid (c,d)
5		II: Vermicellin (Fuska et al. 1979) (c)
		III: Vermiculine (Fuska et al. 1972) (c)
		IV: Vermistatine (Fuska et al. 1986)
		V: Talarone (Mizuno et al. 1974)
		VI: Wortmannin, deoxywortmannin (Simpson et al. 1979;
		MacMillan et al. 1972), wortmannolone (only seen in IMI 44277)
T. helicus	++	
T. macrosporus chemotype I	++	I: Mitorubrin (c,d), mitorubrinic acid (c,d) II: Duclauxin (c,d)
T. macrosporus chemotype II	++	
T. stipitatus chemotype I	++	I: Stipitatic acid (c), stipitatonic acid (c), stipitalide,
		ethylstipitate, stipitaldehydic acid (Dewar 1945; Segal 1959; Divekar et al. 1961; Bryant & Lieht 1974)
		II. 3-methyltriacetic acid lactone, triacetic acid lactone
		tetraacetic acid lactone (Acker et al. 1966: Bentley & Zwitkovitz
		1967: Scott et al. 1971)
		III: Botryodiploidin (Fuska et al. 1988)
		IV: Duclauxin (c) (Kuhr et al. 1973)
		V: Talaromycins (Phillips et al. 1987)
T. stinitatus chemotype II	++	I: Catenarin (c), emodin (c), erythroglaucin (c) (van Eijk 1973)
T. panasenkoi	+	I: Spiculisporic acid (Fujimoto et al. 1988)
		II: Glauconic acid (c.d)
T. luteus	++	I: Luteic acid (Turner & Aldridge 1983)
T. udagawae	<b>+</b> +	I: Mitorubrin (c,d), mitorubrinic acid (c,d), mitorubrinol acetate
		(c,d)
T. wortmannii	++	I: Mitorubrin (c), mitorubrinol (c), mitorubrinic acid (c),
		mitorubrinol acetat (c), wortmin (Turner & Aldridge 1983)
		II: Flavomannin (Atherton et al. 1968), skyrin (c), chrysophanol
		(c) (Turner 1971)
		III: Rugulosin (c) (Breen et al. 1955)
T. rotundus	++	<b>-</b> .,, .
T. trachyspermus	++	I: Spiculisporic acid, (-) decyl citric acid (Gatenbeck & Mahlen
		1968; Tabuchi et al. 1977)
		II: Glauconic acid (c,d)
T. assiutensis	++	II: Glauconic acid (c,d)
T. ohiensis	++	I: Vermiculine (c,d)
		II: Glauconic acid (c,d)
T. galapagensis	+	
T. mimosinus	++	I: Mitorubrin (c,d), mitorubrinic acid (c,d)
T. intermedius	+	
T. purpureus	++	
T. thermophilus	++	
T. derxii	++	
T. bacillisporus	++	I: Bacillosporin A, B & C (Turner & Aldridge 1983)
		II: Pinselin, pinselic acid (Turner & Aldridge 1983)
T. emersonii	+	
T. byssochlamydoides	++	
T. leycettanus	++	

Table 3. Biosynthetic families of secondary metabolites produced by species of Talaromyces.

(a) ++: A very distinct pattern of more than 5 spots before spraying on a TLC plate developed in TEF, +: 2-5 spots on the TLC plates before and after spraying with sulphuric acid and heating.

(b) The known and newly found secondary metabolites were ordered in biosynthetic families and given roman numbers.

(c) Confirmed by TLC and HPLC-DAD and using authentic standards (see table 1).

(d) New record.



Fig. 1. HPLC traces of an extract of T. flavus (Fravel 282.3). Note the production of mitorubrinic acid, vermicellin and possibly talaron (see UV spectrum in upper right corner).

strain was indeed *T. panasenkoi*. The TLC traces also showed that *T. helicus* var. *major* is not sufficiently distinct from var. *helicus*.

Stolk & Samson (1972) recognized two varieties in *T. flavus:* var. *flavus* and var. *macrosporus*. The strains of both varieties only had one metabolite family in common (the mitorubrins), but both taxa produced a conspicous profile of blue, violet, yellow and orange metabolites in TLC plates. This observation was confirmed by HPLC of cultures grown on MEA and OA (Fig. 3). Three isolates of var. *macrosporus* (CBS 353.72, CBS 317.63, and CBS 130.89) all produced great amounts of duclauxin and other specific metabolites, while



Fig. 2. HPLC traces of *Talaromyces macrosporus* CBS 353.72 and two isolates of *T. flavus* (CBS 387.48 and CBS 284.58). Members of the mitorubrinic biosynthetic family were the only common metabolites.



Fig. 3. HPLC traces of Talaromyces udagawae (CBS 579.72) and two isolates of T. luteus (CBS 865.71 and 348.51) to show the different profiles of secondary metabolites.

strains of *T. flavus* produced vermiculine, vermicelline and other unknown secondary metabolites not found in var. *macrosporus*. Beuchat (1988) compared several isolates of *T. flavus* and found that small-spored strains are less heat-resistant than large-spored strains. Among the large-spored strains he included CBS 317.63, the type culture of *T. flavus* var. *macrosporus*, isolated as a heat resistant strain from pasteurized apple juice by Van der Spuy et al. (1975). Based on these differences in ascospore size, heat-resistance and profile of secondary metabolites we propose that the two taxa should be considered as separate species.

# Talaromyces macrosporus (Stolk & Samson) Frisvad, Samson & Stolk, comb. nov.

Basionym: *Talaromyces flavus* (Klöcker) Stolk & Samson var. *macrosporus* Stolk & Samson, Stud. Mycol., Baarn 2: 15, 1972.

Status conidialis *Penicillium macrosporum* stat. nov.

Descriptio stat. anam. in Stolk & Samson, Stud. Mycol., Baarn, 2: 15, 1972. Metabolica: Duclauxin, mitorubrin, mitorubrinic acid. Typus: CBS 317.63. Stolk & Samson (1972) separated *T. udagawae* and *T. luteus* on the basis of different ascoma initials and smaller ascopores with different ornamentation. Pitt (1980) found that these differences were not sufficient and synonymized both taxa. Our TLC and HPLC results, however, showed that *T. udagawae* is a well-defined species, and that the two taxa have no secondary metabolites in common (metabolites having the same retention time in the two species were different as they had different UV-VIS spectra (Fig. 3).

In recognizing *T. luteus* and *T. udagawae* as two different species, there is no name available for the anamorph of *T. luteus*, as *P. luteum* was described inclusive of the teleomorph. We therefore propose the following name for the anamorph of *T. luteus*:

### Penicillium vonarxii Frisvad & Samson, sp. nov.

Status anamorphosis *Talaromycetis lutei*. Conidiophora portata ex hyphis aeriis, stipites 15–35  $(-100) \times 2.0-3 \mu m$ , parietibus levibus, interdum incrustatis, penicilli biverticillati et monoverticillati, metulae  $10-20 \times 2-3 \mu m$ , phialides acerosae,  $9-15 \times 2-3 \mu m$ , collulis longis, apicibus interdum viridibus, conidia ellipsoidea vel pyriformia,  $2.5-4 \times 2-3.2 \,\mu$ m, parietibus levibus, viridibus. In agaro CYA, coloniae aetate unius hebdomadis 15– 20 mm diam. Metabolica absorb. max. 232 nm, 281 nm, 320 nm, 330 nm, 404 nm. Typus: CBS 579.72.

Samson & Abdel-Fattah (1978) described *T. assiutensis*. A morphological and chemical comparison of the type cultures of *T. assiutensis* and *T. gossypii* Pitt (1980) showed that these species are conspecific.

In their monograph Stolk & Samson (1972) mentioned deviating strains of T. flavus var. macrosporus (CBS 350.72) (with more ellipsoidal ascospores, showing more variation in size than other isolates of var. macrosporus) and T. stipitatus (CBS 349.72) (with slightly larger ascospores and different mycelial colours than T. stipitatuson all media) respectively. In our studies of the secondary metabolites these strains produce completely different profiles of secondary metabolites and they might represent new taxa in Talaromyces. As neither of the two aberrant strains have any metabolites in common with the other species of the genus they cannot be regarded as mutants or varieties of those species. However, before warranting their description as new species we would like to examine more than one isolate. For the time being we call them T. macrosporus chemotype II for CBS 350.72 and T. stipitatus chemotype II for CBS 349.72, following the recommendations by Pitt & Hawksworth (1985) for naming chemical variants.

Our comparative studies of the strains of Talaromyces and Penicillium subgenus Biverticillium showed that T. wortmannii and the anamorph P. variabile have many secondary metabolites in common, which may indicate that the latter may be the anamorph of the former fungus. This is further supported by similar growth rates and the fact that Raper & Thom (1949) reported the production of a few ascomata in P. variabile NRRL 2025 with ascospores like those of T. wortmannii. Both T. wortmannii and P. variabile have, however, each some specific secondary metabolites too and at present we prefer to keep these species separate.

The results reported here indicate the great value of profiles of secondary metabolites as taxonomic characters in *Talaromyces*. While the morphological treatment of the genus *Talaromyces* by Stolk & Samson (1972) and Pitt (1980) is upheld, the simple TLC technique can also help in solving problems where morphological characters alone leave uncertainty. The profiles of secondary metabolites in *Talaromyces* and *Penicillium* subgenus *Biverticillium* indicate a close relationship between these taxa, but no *Biverticillium* taxa could be regarded as anamorphs of any *Talaromyces* species.

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