from Roman catacombs

Agromyces italicus sp. nov., Agromyces humatus sp. nov. and Agromyces lapidis sp. nov., isolated

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A polyphasic study was carried out to clarify the taxonomic positions of three Gram-positive isolates from the Catacombs of Domitilla, Rome (Italy). 16S rRNA gene sequence comparisons placed these strains within the genus *Agromyces*. The morphological and chemotaxonomic characteristics of these isolates were consistent with the description of the genus *Agromyces*. The three isolates could be readily distinguished from one another and from representatives of all *Agromyces* species with validly published names by a broad range of phenotypic characteristics and DNA–DNA relatedness studies. Therefore, these isolates are proposed to represent three novel species of the genus *Agromyces*, *Agromyces italicus* sp. nov. (type strain CD1^T = HKI 0325^T = DSM 16388^T = NCIMB 14011^T), *Agromyces humatus* sp. nov. (type strain CD5^T = HKI 0327^T = DSM 16389^T = NCIMB 14012^T) and *Agromyces lapidis* sp. nov. (type strain CD55^T = HKI 0324^T = DSM 16390^T = NCIMB 14013^T).

Strains of the genus Agromyces are common inhabitants of different soils ranging from fertile meadows to deserts [Agromyces ramosus Gledhill and Casida 1969, Agromyces cerinus subsp. cerinus Zgurskaya et al. 1992, Agromyces cerinus subsp. nitratus Zgurskaya et al. 1992, Agromyces fucosus Zgurskaya et al. 1992 emend. Ortiz-Martinez et al. 2004, Agromyces hippuratus (Zgurskava et al. 1992) Ortiz-Martinez et al., 2004, Agromyces mediolanus Suzuki et al. 1996 and Agromyces aurantiacus Li et al. 2003]. The rhizosphere of plants harbours a wide diversity of these organisms, as has been shown by Takeuchi & Hatano (2001) with the description of three species, Agromyces bracchium, Agromyces luteolus and Agromyces rhizospherae, from mangrove rhizosphere soil. Dorofeeva et al. (2003) isolated Agromyces albus from the leaves and inflorescences of members of the Primulaceae, while Agromyces ulmi was isolated from the decayed stump of an elm tree, Ulmus nigra (Rivas et al., 2004). Recently, two novel species of this genus, Agromyces salentinus and Agromyces neolithicus, have been isolated from the Grotta dei Cervi, a cave in southern Italy with singular neolithic rock art paintings (Jurado et al., 2005).

Strains CD1^{T} and CD5^{T} were obtained from the wall of a tomb located in the Little Apostle cubicle, Domitilla Catacombs, Rome, Italy (Sanchez-Moral *et al.*, 2004). Strain CD55^{T} was isolated by touching the stone wall of a tomb (first arcosolium on the left side behind the entrance of Domitilla Catacombs) with a sterile cotton swab and suspending the attached bacteria in 1:10 diluted organic medium 79 (OM79) (Prauser & Falta, 1968; Jurado *et al.*, 2005). The strains were isolated on nutrient agar (Difco) or, in the case of strain CD55^{T} , on peptone/yeast extract/brain heart infusion agar (Yokota *et al.*, 1993), using a standard dilution plate procedure.

All the methods used in this study have been recently described by Jurado *et al.* (2005). Range of pH for growth was established using liquid OM79 medium adjusted to initial pH values between 4 and 11 with either 1 M HCl or 20 % (w/v) Na₂CO₃ solution and incubated at 28 °C for up to 10 days. The following type strains were used as references for comparative studies: *A. albus* VKM Ac-1800^T, *A. cerinus* subsp. *cerinus* IMET 11525^T, *A. fucosus* IMET 11529^T, *A. mediolanus* VKM Ac-1388^T, *A. neolithicus* DSM 16197^T, *A. ramosus* IMET 11027^T and *A. salentinus* DSM 16198^T. Morphological and physiological traits are summarized in the species descriptions and Table 1.

Differences referring to the compositions of whole-cell sugars, menaquinones and polar lipids are shown in Table 2. Cell-wall amino acids in strains CD1^T, CD5^T and CD55^T

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The GenBank/EMBL/DDBJ accession numbers for the 16S rRNA gene sequences of strains $CD1^{T}$, $CD5^{T}$ and $CD55^{T}$ are AY618215, AY618216 and AY618217, respectively.

Table 1. Characteristics that differentiate strains $CD1^{T}$, $CD5^{T}$ and $CD55^{T}$ and their closest relatives within the genus Agromyces

Strains: 1, *A. italicus* sp. nov. $CD1^{T}$; 2, *A. humatus* sp. nov. $CD5^{T}$; 3, *A. lapidis* sp. nov. $CD55^{T}$; 4, *A. albus* VKM Ac-1800^T; 5, *A. cerinus* subsp. *cerinus* IMET 11525^T; 6, *A. fucosus* IMET 11529^T; 7, *A. mediolanus* VKM Ac-1388^T; 8, *A. neolithicus* DSM 16197^T; 9, *A. ramosus* IMET 11027^T; 10, *A. salentinus* DSM 16198^T. Data for *A. cerinus* subsp. *cerinus* and *A. ramosus* were taken from Groth *et al.* (1996). –, Negative; +, positive; (+) weakly positive; V, variable; ND, not determined. Strains $CD1^{T}$, $CD5^{T}$ and $CD55^{T}$ share the following properties. They produce acid from starch, arbutin, D-fructose, glycogen, maltose, D-mannose, but not from adonitol, D-arabitol, L-arabitol, dulcitol, erythritol, D-fucose, gluconate, 2-ketogluconate, 5-ketogluconate, β -gentiobiose, inositol, D-lyxose, melibiose, melezitose, sorbitol, L-sorbose; D-tagatose, L-xylose, methyl β -xyloside or xylitol. They produce alkaline phosphatase, esterase (C1), esterase lipase (C8), leucine arylamidase, valine arylamidase, cystine arylamidase, lipase (C14), trypsin or α -fucosidase. They are sensitive to chloramphenicol (30 µg), imipenem (10 µg), ofloxacin (10 µg), oxytetracycline hydrochloride (30 µg), rifampicin (5 µg) and vancomycin hydrochloride (30 µg), but not to kanamycin (30 µg), lincomycin (2 µg), norfloxacin (10 µg) or sulfonamide (200 µg). They hydrolyse aesculin and starch, but not adenine or Tween 80. Voges–Proskauer, methyl red and indole tests give negative results. They are positive for H₂S production.

Characteristic	1	2	3	4	5	6	7	8	9	10
DNA G+C content (mol%)	70.8	70.6	70.4	69.0	70.5	70.6	72.3	65.3	68.9	72.3
Growth at 10 °C	(+)	_	(+)	(+)	(+)	+	(+)	_	+	+
Growth in 4.0% NaCl (w/v)	+	_	+	(+)	_	+	+	_	(+)	(+)
Microaerophilic growth	+	_	+	—	_	+	+	+	+	(+)
Decomposition or hydrolysis of:										
Casein	+	+	+	+	_	+	+	+	_	+
Gelatin	+	+	V	+	_	+	_	+	_	_
Hippurate	+	+	+	+	_	+	+	+	+	+
Hypoxanthine	+	(+)	_	_	(+)	+	+	_	_	_
Tyrosine	+	+	+	_	+	+	+	+	_	+
Urea	_	_	_	+	_	_	_	_	_	+
Xanthine	+	_	_	_	_	+	+	_	_	_
Biochemical tests										
Nitrate reduction	+	+	+	_	_	_	_	+	V	_
Catalase reaction	+	+	+	+	+	+	+	+	_	+
Oxidase test	+	_	V	+	+	+	V	V	_	V
Acid production from (API 50 CH B/E):										
Amygdalin	(+)	_	+	_	+	+	+	_	+	(+)
D-Arabinose	+	_	+	+*	+	+	$(+)^{*}$	+	+	+
L-Arabinose	+	+	_	+	_	+	_	+	+	+
Cellobiose	+	V	+	_	+	+	+	+	_	+
L-Fucose	+	_	+	_	(+)	+	_	+	+	+
Galactose	+	+	+	(+)	+	+	+	+	_	+
D-Glucose	+	+	+	(+)	+	+	+	+	_	+
Glycerol	+	+	+	_	+	+	+	+	+	+
Inulin	_	+	+	_	_	(+)	_	_	+	+
Lactose	(+)	_	_	_	+	_	+	_	_	_
Methyl α-D-mannoside	_	_	+	_	_	+	_	_	_	_
Methyl α-D-glucoside	_	_	_	_	_	+	_	_	_	_
Mannitol	_	+	_	+	_	_	_	+	(+)	_
N-Acetylglucosamine	_	V	+	_	_	+	_	$(+)^{*}$	+	_
D-Raffinose	_	+	(+)	+*	_	+	_	+	+	(+)
Rhamnose	_	+	_	+	+	+*	+	_	+	+
Ribose	_	_	_	+*	_	_	+	_	_	+
Salicin	+	_	+	_	+	+	(+)	_	_	+
Sucrose	_	+	+	(+)	_	+	+	(V)	+	+
Trehalose	_	_	_	_	_	(+)*	+	_	_	+
D-Turanose	_	_	_	_	_	_	(+)	_	_	+
D-Xylose	+	_	_	+*	_	_	_	_	_	+
Utilization of:										
Aconitate	_	_	_	_	+	_	_	_	_	_

	Tabl	e 1.	cont.
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Characteristic	1	2	3	4	5	6	7	8	9	10
Citrate	_	_	_	_	+	_	_	_	-	_
Malate	_	+	—	+	+	_	_	_	+	_
Succinate	_	+	_	+	_	_	_	_	+	_
Enzyme assays (API ZYM):										
α-Chymotrypsin	_	_	+	_	_	_	_	+	_	_
α-Galactosidase	_	_	_	_	+	_	_	+	_	_
β -Galactosidase	+	+	+	+	+	+	+	_	_	+
β -Glucuronidase	_	+	_	_	_	_	_	+	_	_
N -Acetyl- β -glucosaminidase	_	_	_	+	_	+	+	+	_	+
α-Mannosidase	_	+	_	_	_	_	_	+	_	_
Antibiotic susceptibility:										
Ampicillin (10 µg)	+	+	+	+	+	_	+	+	+	+
Ciprofloxacin (5 µg)	+	+	+	+	V	+	_	+	+	+
Kanamycin (30 µg)	_	_	_	+	+	+	_	_	+	+
Methicillin (5 µg)	+	+	+	_	ND	_	_	+	ND	+
Nalidixic acid (30 µg)	_	_	+	+	_	_	_	_	_	+
Norfloxacin (10 µg)	_	_	_	_	_	_	_	+	_	+
Novobiocin (5 µg)	+	+	+	+	+	+	_	+	_	+
Penicillin G (10 IU)	+	_	+	_	_	_	_	+	+	+
Polymyxin B (300 IU)	_	+	_	+	V	_	_	+	+	+
Streptomycin sulfate (10 µg)	_	+	_	+	+	+	_	_	+	+
Sulfonamide (200 µg)	_	_	_	_	_	_	_	_	_	+

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were diaminobutyric acid, glycine, glutamic acid and alanine. Acyl type for these three strains was acetyl. The predominant fatty acids of strains CD1^{T} , CD5^{T} and CD55^{T} were anteiso- $\text{C}_{15:0}$ (50·9, 41·5 and 48·1%, respectively), anteiso- $\text{C}_{17:0}$ (15·3, 34·0 and 13·0%), iso- $\text{C}_{16:0}$ (14·7, 15·5 and 16·5%), iso- $\text{C}_{15:0}$ (14·2, 5·4 and 10·5%) and $\text{C}_{16:0}$ (1·7, 1·3 and 5·9%).

Sequence comparisons of 16S rRNA genes from strains $CD1^{T}$, $CD5^{T}$ and $CD55^{T}$ showed a close phylogenetic relationship to *A. ramosus*, *A. cerinus* subsp. *cerinus*, *A. salentinus*, *A. neolithicus*, *A. albus*, *A. mediolanus* and *A. fucosus* with percentages of similarity ranging from 94 to 98%. Strains $CD1^{T}$ and $CD55^{T}$ showed high similarity

(97%). The recently described species *A. salentinus* (Jurado *et al.*, 2005) was highly related to strains $CD1^{T}$ and $CD55^{T}$ with 97% similarity. *A. neolithicus* was the closest relative to strain $CD5^{T}$ (98%). A phylogenetic tree generated by the neighbour-joining method showing the relationships between members of the genus *Agromyces* and the three novel isolates, $CD1^{T}$, $CD5^{T}$ and $CD55^{T}$, is shown in Fig. 1.

The DNA G + C contents of strains CD1^{T} , CD5^{T} and CD55^{T} were 70.8, 70.6 and 70.4 mol%, respectively. DNA–DNA relatedness studies showed significant differences between the three isolates as well as with their closest phylogenetic neighbours. In all cases, differences in melting temperatures were >7.7 °C (roughly <51% DNA–DNA relatedness;

Table 2. Chemotaxonomic characteristics of strains CD1^T, CD5^T and CD55^T

Components are listed in decreasing order of abundance.

Characteristic	Strain CD1 ^T	Strain CD5 ^T	Strain CD55 ^T		
Whole-cell sugars*	Gal, Rib, Glc, Man	Glc, Gal, Rha, Man	Glc, Gal, Man, Rib		
Major menaquinones	12, 13	13, 12	12, 13		
Polar lipids†	DPG, PG, 2 PL, GL	DPG, PG, PL, 2 GL	DPG, PG, PL, 2 GL		

*Gal, Galactose; Glc, glucose; Man, mannose; Rha, rhamnose; Rib, ribose.

†DPG, Diphosphatidylglycerol; GL, unknown glycolipid; PG, phosphatidylglycerol; PL, unknown phospholipid.

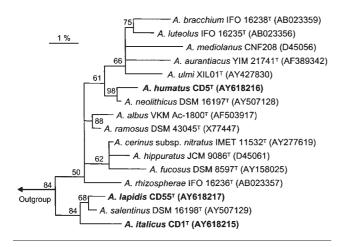


Fig. 1. Neighbour-joining phylogenetic tree showing the relationships between species of the genus *Agromyces* and isolates $CD1^{T}$, $CD5^{T}$ and $CD55^{T}$ based on 16S rRNA gene sequences. Bar, 1% sequence divergence. The outgroup (not shown) was *Corynebacterium* sp. QSSC3-5 (AF170740).

Rosselló-Mora & Amann, 2001), which is well above the 5 °C cut-off point recommended for the delineation of species (Stackebrandt & Goebel, 1994).

The genotypic and phenotypic characteristics of strains $CD1^{T}$, $CD5^{T}$ and $CD55^{T}$ are consistent with their classification in the genus Agromyces (Gledhill & Casida, 1969; Zgurskaya et al., 1992). The isolates can be readily distinguished from representatives of Agromyces species with validly published names, and from one another, by a broad range of phenotypic properties (Table 1) and composition of whole-cell sugars (Table 2). DNA-DNA relatedness data support the above distinctions between strains CD1^T, CD5^T and CD55^T and their closest relatives within the genus Agromyces. Based on the results of this polyphasic approach, we propose that the three studied isolates are classified within novel species of the genus Agromyces, Agromyces italicus sp. nov. (strain $CD1^{T}$), Agromyces humatus sp. nov. (strain $CD5^{T}$) and Agromyces lapidis sp. nov. (strain $CD55^{T}$).

Description of Agromyces italicus sp. nov.

Agromyces italicus (i.ta'li.cus. L. masc. adj. italicus from Italy, the origin of the type strain).

Gram-positive, aerobic to microaerophilic actinomycete that forms branching hyphae (width $0.4-0.6 \ \mu$ m) which break up into irregular diphtheroid and rod-like, non-motile fragments. Colonies are circular, convex, smooth and cream. Colony diameter is about 1 mm. Growth occurs between 10 and 37 °C (optimal growth at 28 °C) and at pH 5–9.5. NaCl is tolerated up to 4 % (w/v), but not at 6 % (w/v). Phenotypic characteristics including antibiotic susceptibility and enzymic activities are reported in Table 1. Chemotaxonomic characteristics are reported in Table 2. Cell-wall amino acids are diaminobutyric acid, glycine,

glutamic acid and alanine. Acyl type is acetyl. Predominant fatty acids are anteiso- $C_{15:0}$ and anteiso- $C_{17:0}$; mycolic acids are absent. G+C content is 70.8 mol%.

The type strain is $CD1^{T}$ (=HKI 0325^{T} =DSM 16388^{T} = NCIMB 14011^{T}), isolated from the wall of a tomb located in the Little Apostle cubicle, Domitilla Catacombs, Rome, Italy.

Description of Agromyces humatus sp. nov.

Agromyces humatus (hu.ma'tus. L. masc. part. adj. *humatus* buried).

Gram-positive, aerobic actinomycete that forms branching hyphae (width $0.3-0.5 \ \mu$ m) which break up into irregular diphtheroid and rod-like, non-motile fragments. Colonies are circular, convex, smooth and yellow. Colony diameter is about 1 mm. Growth occurs between 15 and 37 °C (optimal growth at 28 °C) and at pH 5–9.5. NaCl is tolerated up to 2 % (w/v), but not at 4 % (w/v). Phenotypic characteristics including antibiotic susceptibility and enzymic activities are reported in Table 1. Chemotaxonomic characteristics are reported in Table 2. Cell-wall amino acids are diaminobutyric acid, glycine, glutamic acid and alanine. Acyl type is acetyl. Predominant fatty acids are anteiso-C_{15:0} and anteiso-C_{17:0}; mycolic acids are absent. G+C content is 70.6 mol%.

The type strain is $CD5^{T}$ (=HKI 0327^{T} =DSM 16389^{T} = NCIMB 14012^{T}), isolated from the wall of a tomb located in the Little Apostle cubicle, Domitilla Catacombs, Rome, Italy.

Description of Agromyces lapidis sp. nov.

Agromyces lapidis (la.pi'dis. L. gen. n. lapidis of a stone).

Gram-positive, aerobic to microaerophilic actinomycete that forms branching hyphae (width $0.4-0.6 \mu$ m) which break up into irregular diphtheroid and rod-like, nonmotile fragments. Colonies are circular, convex, smooth and yellow. Colony diameter is about 1 mm. Growth occurs between 10 and 37 °C (optimal growth at 28 °C) and at pH 5–9.5. NaCl is tolerated up to 4 % (w/v), but not at 6 % (w/v). Phenotypic characteristics including antibiotic susceptibility and enzymic activities are reported in Table 1. Chemotaxonomic characteristics are reported in Table 2. Cell-wall amino acids are diaminobutyric acid, glycine, glutamic acid and alanine. Acyl type is acetyl. Predominant fatty acids are anteiso-C_{15:0} and iso-C_{16:0}; mycolic acids are absent. G+C content is 70.4 mol%.

The type strain is $CD55^{T}$ (=HKI 0324^T=DSM 16390^T= NCIMB 14013^T), isolated from a carved stone wall of the Domitilla Catacombs, Rome, Italy.

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