

Kineosporia babensis sp. nov., isolated from plant litter in Vietnam

Sakiyama Y., Thao N.K.N., Giang N.M., Miyadoh S., Hop D.V., Ando K.

NITE Biological Resource Center (NBRC), National Institute of Technology and Evaluation (NITE), Chiba 292-0818, Japan; Institute of Microbiology and Biotechnology (IMBT), Vietnam National University Hanoi (VNUH), Hanoi, Viet Nam

Abstract: Three actinomycetes, designated strains VN05A0342, VN05A0351 and VN05A0415^T, were isolated from plant-litter samples collected in the north of Vietnam and examined in a polyphasic taxonomic study. Phylogenetic analysis based on the 16S rRNA gene sequences showed that these isolates were most closely related to the type strain of *Kineosporia mikuniensis* (98.5% sequence similarity). Morphological properties (the formation of spore domes and motile spores) and chemotaxonomic data supported the assignment of the three isolates to the genus *Kineosporia*. The isolates all contained the following: meso-diaminopimelic acid in the peptidoglycan (with small amounts of the LL isomer); ribose, mannose, galactose and glucose as the whole-cell sugars; MK-9(H₄) as the predominant isoprenoid quinone; C_{18:1} and C_{16:0} as the major cellular fatty acids; and phosphatidylcholine, phosphatidylglycerol, diphosphatidylglycerol and phosphatidylinositol as the phospholipids. The high DNA-DNA relatedness (>71%) among the three isolates showed that they represented a single species. On the other hand, the DNA-DNA relatedness between the novel isolates and all type strains of *Kineosporia* species was less than 46%. The physiological properties of our isolates were distinct from those of all of the *Kineosporia* species with validly published names, e.g. decomposition of L-tyrosine and aesculin and the utilization of raffinose and D-arabitol. Therefore, strains VN05A0342, VN05A0351 and VN05A0415^T represent a novel species of the genus *Kineosporia*, for which the name *Kineosporia babensis* sp. nov. is proposed. The type strain is VN05A0415^T (=VTCC-A-0961^T =NBRC 104154^T). © 2009 IUMS.

Index Keywords: bacterial RNA; cardiolipin; diaminopimelic acid; fatty acid; galactose; glucose; mannose; peptidoglycan; phosphatidylcholine; phosphatidylglycerol; phosphatidylinositol; phospholipid; quinone derivative; ribose; RNA 16S; Actinobacteria; article; bacterium isolate; chemotaxonomy; gene sequence; *Kineosporia babensis*; new species; nonhuman; nucleotide sequence; phylogeny; plant litter; priority journal; sequence homology; type strain; Actinobacteria (class); *Kineosporia*; *Micromonospora echinaurantiaca*

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Molecular Sequence Numbers: GENBANK: AB003931, AB003932, AB003933, AB003935, AB025317, AB377116, AB377117, AB377118, AB377119, AY831385, L41048, X77958, X92357, X93190

Chemicals/CAS: diaminopimelic acid, 583-93-7; galactose, 26566-61-0, 50855-33-9, 59-23-4; glucose, 50-

99-7, 84778-64-3; mannose, 31103-86-3, 3458-28-4; peptidoglycan, 9047-10-3; phosphatidylcholine, 55128-59-1, 8002-43-5; ribose, 34466-20-1, 50-69-1, 93781-19-2

Correspondence Address: Sakiyama, Y.; NITE Biological Resource Center (NBRC), National Institute of Technology and Evaluation (NITE), Chiba 292-0818, Japan; email: sakiyama-yayoi@nite.go.jp

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Authors with affiliations:

- Sakiyama, Y., NITE Biological Resource Center (NBRC), National Institute of Technology and Evaluation (NITE), Chiba 292-0818, Japan
- Thao, N.K.N., Institute of Microbiology and Biotechnology (IMBT), Vietnam National University Hanoi (VNUH), Hanoi, Viet Nam
- Giang, N.M., Institute of Microbiology and Biotechnology (IMBT), Vietnam National University Hanoi (VNUH), Hanoi, Viet Nam
- Miyadoh, S., NITE Biological Resource Center (NBRC), National Institute of Technology and Evaluation (NITE), Chiba 292-0818, Japan
- Hop, D.V., Institute of Microbiology and Biotechnology (IMBT), Vietnam National University Hanoi (VNUH), Hanoi, Viet Nam
- Ando, K., NITE Biological Resource Center (NBRC), National Institute of Technology and Evaluation (NITE), Chiba 292-0818, Japan

References:

- Ezaki, T., Hashimoto, Y., Yabuuchi, E., Fluorometric deoxyribonucleic acid-deoxyribonucleic acid hybridization in microdilution wells as an alternative to membrane filter hybridization in which radioisotopes are used to determine genetic relatedness among bacterial strains (1989) *Int J Syst Bacteriol*, 39, pp. 224-229
- Felsenstein, J., Confidence limits on phylogenies: An approach using the bootstrap (1985) *Evolution*, 39, pp. 783-791
- Goodfellow, M., Numerical taxonomy of some nocardioform bacteria (1971) *J Gen Microbiol*, 69, pp. 33-80
- Gordon, R.E., Mihm, J.M., A comparative study of some strains received as nocardiae (1957) *J Bacteriol*, 73, pp. 15-27
- Gordon, R.E., Barnett, D.A., Handerhan, J.E., Pang, C.H.-N., *Nocardia coeliaca*, *Nocardia autotrophica*, and the nocardin strain (1974) *Int J Syst Bacteriol*, 24, pp. 54-63
- Hayakawa, M., Nonomura, H., Humic acid-vitamin agar, a new medium for selective isolation of soil actinomycetes (1987) *J Ferment Technol*, 65, pp. 501-509
- Hayakawa, M., Otaguro, M., Takeuchi, T., Yarnazaki, T., Iinuma, Y., Application of a method incorporating differential centrifugation for selective isolation of motile actinomycetes in soil and plant litter (2000) *Antonie Van Leeuwenhoek*, 78, pp. 171-185
- Itoh, T., Kudo, T., Parenti, F., Seino, A., Amended description of the genus *Kineosporia*, based on chemotaxonomic and morphological studies (1989) *Int J Syst Bacteriol*, 39, pp. 168-173
- Kluge, A.G., Farris, J.S., Quantitative phyletics and the evolution of anurans (1969) *Syst Zool*, 18, pp. 1-32

- Kudo, T., Matsushima, K., Itoh, T., Sasaki, J., Suzuki, K., Description of four new species of the genus *Kineosporia*: *Kineosporia succinea* sp. nov., *Kineosporia rhizophila* sp. nov., *Kineosporia mikuniensis* sp. nov. and *Kineosporia rhamnosa* sp. nov., isolated from plant samples, and amended description of the genus *Kineosporia* (1998) *Int J Syst Bacteriol*, 48, pp. 1245-1255
- Marmur, J., A procedure for the isolation of deoxyribonucleic acid from microorganisms (1961) *J Mol Biol*, 3, pp. 208-218
- Mesbah, M., Premachandran, U., Whitman, W.B., Precise measurement of the G + C content of deoxyribonucleic acid by high-performance liquid chromatography (1989) *Int J Syst Bacteriol*, 39, pp. 159-167
- Minnikin, D.E., O'Donnell, A.G., Goodfellow, M., Alderson, G., Athalye, M., Schaal, A., Parlett, J.H., An integrated procedure for the extraction of bacterial isoprenoid quinones and polar lipids (1984) *J Microbiol Methods*, 2, pp. 233-241
- Nozawa, Y., Sakai, N., Arai, K., Kawasaki, Y., Harada, K., Reliable and sensitive analysis of amino acids in the peptidoglycan of actinomycetes using the advanced Marfey's method (2007) *J Microbiol Methods*, 70, pp. 306-311
- Pagani, H., Parenti, F., *Kineosporia*, a new genus of the order Actinomycetales (1978) *Int J Syst Bacteriol*, 28, pp. 401-406
- Saito, H., Miura, K., Preparation of transforming deoxyribonucleic acid by phenol treatment (1963) *Biochim Biophys Acta*, 72, pp. 619-629
- Saitou, N., Nei, M., The neighbor-joining method: A new method for reconstructing phylogenetic trees (1987) *Mol Biol Evol*, 4, pp. 406-425
- Shirling, E.B., Gottlieb, D., Methods for characterization of *Streptomyces* species (1966) *Int J Syst Bacteriol*, 16, pp. 313-340
- Staneck, J.L., Roberts, G.D., Simplified approach to identification of aerobic actinomycetes by thin-layer chromatography (1974) *Appl Microbiol*, 28, pp. 226-231
- Tamura, T. & Hatano, K. (2001). Phylogenetic analysis of the genus *Actinoplanes* and transfer of *Actinoplanes minutisporangius* Ruan et al. 1986 and '*Actinoplanes aurantiacus*' to *Cryptosporangium minutisporangium* comb. nov. and *Cryptosporangium aurantiacum* sp. nov. *Int J Syst Evol Microbiol* 51, 2119-2125
Tamura, T., Nakagaito, Y., Nishii, T., Hasegawa, T., Stackebrandt, E., Yokota, A., A new genus of the order Actinomycetales, *Couchioplanes* gen. nov., with descriptions of *Couchioplanes caeruleus* (Horan and Brodsky 1986) comb. nov. and *Couchioplanes caeruleus* subsp. *azureus* subsp. nov (1994) *Int J Syst Bacteriol*, 44, pp. 193-203
- Thompson, J.D., Gibson, T.J., Plewniak, F., Jeanmougin, F., Higgins, D.G., The CLUSTAL_X Windows interface: Flexible strategies for multiple sequence alignment aided by quality analysis tools (1997) *Nucleic Acids Res*, 25, pp. 4876-4882
- Wayne, L.G., Brenner, D.J., Colwell, R.R., Grimont, P.A.D., Kandler, O., Krichevsky, M.I., Moore, L.H., Murray, R.G.E., International Committee on Systematic Bacteriology. Report of the ad hoc committee on reconciliation of approaches to bacterial systematics (1987) *Int J Syst Bacteriol*, 37, pp. 463-464
- Willoughby, L.G., A study on aquatic actinomycetes, the allochthonous leaf component (1969) *Nova Hedwigia*, 18, pp. 45-113

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