A new species and four new records of *Bacidia* (Lecanorales, Ramalinaceae) from South Korea, with a key to Korean species

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**Abstract**

A new species, *Bacidia fuscopallida* Lee & Heo and four new records, *B. ekmaniana* R. C. Harris, Ladd & Lendemer, *B. friesiana* (Hepp) Körb., *B. heterochroa* (Müll. Arg.) Zahlbr. and *B. suffusa* (Fr.) A. Schneid., are described from South Korea. *Bacidia fuscopallida* differs from *B. diffracta* S. Ekman, the most similar species, by warted but non-granular thallus, paler and smaller apothecia without pruina, proper exciple without crystals, over 11-septate ascospores and smaller pycnidia and pycnoconidia. *Bacidia ekmaniana* is recorded new to Asia, *B. heterochroa* is reported new to northeastern Asia and *B. friesiana* and *B. suffusa* are new to Korea. Molecular analyses employing internal transcribed spacer (ITS) sequences strongly support the classification of the five species of *Bacidia*. A surrogate key is provided to assist in the identification of all 19 taxa in *Bacidia* of Korea.

**Keywords**

biodiversity, corticolous, lichen, phylogeny, taxonomy

**Introduction**

*Bacidia* has become a species-rich genus since De Notaris’ (1846) introduction. *Bacidia* (230 spp. including *Bacidiopsora*) is one of the largest genera in Ramalinaceae, with *Ramalina* (230 spp.) (Wijayawardene et al. 2020). The genus *Bacidia* was defined in a wide sense by the characteristics of crustose lichens with a chlorococcoid photobiont, biatorine or lecideine apothecia, 8-spored asci with colourless and transversely 3- or
more septate ascospores (Zahlbruckner 1905, 1921–1940). However, the traditional characterisation of the genus has been considered coarse and unnatural. The genus has been split (e.g. Santesson 1952; Vězda 1978) and particularly new taxonomic applications, based on ascus structures (Hafellner 1984), excipulum structures (Vězda 1990) and molecular results (Ekman and Wedin 2000; Ekman 2001) have reclassified the large genus into tens of different genera (e.g. Vězda 1986; Sérusiaux 1986, 1993, 1995; Lücking 1992, 1995; Aptroot and Sipman 1993; Lücking et al. 1994; Ekman 1996; Kistenich et al. 2018). Ekman (2001) represented that Bacidia might be delimited to the B. rosella (Pers.) De Not., the type species, group in a strict sense (Brand et al. 2009) and most Bacidia species with blue-green pigment in ephymenium are closer to Toninia than the type species group, based on molecular phylogeny although B. schweinitzii (Fr. ex E. Michener) A. Schneid.) can be an exception.

Bacidia is one of the least explored genera in Korea and the genus has just been reported since the 2010s. Since Joshi et al. (2011) introduced B. arceutina (Ach.) Th. Fr., B. schweinitzii and B. subincompta (Nyl.) Arnold (syn. Toniniopsis subincompta (Nyl.) Kistenich, Timdal, Bendiksby & S. Ekman), overall 18 species have been recorded in Korea (Zhang et al. 2012; Aptroot and Moon 2014, 2015; Kondratyuk et al. 2016, 2017, 2019a, b; Liu 2018; Yakovchenko et al. 2018). Although detected on diverse substrates (e.g. bark, moss, rock or artificial wood fence), they are mainly corticolous and were collected on deciduous, wide-leaved tree barks in humid forests.

This study describes a new species and four new records of the lichen genus Bacidia. Field surveys for the lichen biodiversity in the main mountains of Korea, i.e. Baekdu-daegan, and several forested wetlands of South Korea were carried out during the spring to summer of 2019–2021 and 54 specimens of Bacidia were collected from barks of deciduous wide-leaved trees and shrubs (Fig. 1). The specimens were comprehensively analysed and identified as a new species, B. fuscopallida, and four new records, B. ekmaniana, B. friesiana, B. heterochroa and B. suffusa. All the collected specimens are deposited in the Herbarium of the Baekdu-daegan National Arboretum (KBA), South Korea.

**Materials and methods**

**Morphological and chemical analyses**

Hand sections were prepared manually with a razor blade under a stereomicroscope (Olympus optical SZ51; Olympus, Tokyo, Japan), examined under a compound microscope (Nikon Eclipse E400; Nikon, Tokyo, Japan) and pictured using a software programme (NIS-Elements D; Nikon, Tokyo, Japan) and a DS-Fi3 camera (Nikon, Tokyo, Japan) mounted on a Nikon Eclipse Ni-U microscope (Nikon, Tokyo, Japan). The ascospores were examined at 1000× magnification in water. The length and width of the ascospores were measured and the range of spore sizes was shown with average, standard deviation (SD), length-to-width ratio and the number of measured spores. Thin-layer chromatography (TLC) was performed using solvent system C according to standard methods (Orange et al. 2001).
Isolation, DNA extraction, amplification and sequencing

Hand-cut sections of ten to twenty ascomata per collected specimen were prepared for DNA isolation (Table 1) and DNA was extracted with a NucleoSpin Plant II Kit in line with the manufacturer’s instructions (Macherey-Nagel, Düren, Germany). PCR amplifications for the internal transcribed spacer region (ITS1-5.8S-ITS2 rDNA) RNA genes were achieved using Bioneer’s AccuPower PCR Premix (Bioneer, Daejeon, Korea) in 20-μl tubes with 16 μl of distilled water, 2 μl of DNA extracts and 2 μl of the primers ITS5 and ITS4 (White et al. 1990). The PCR thermal cycling parameters used were 95 °C (15 sec), followed by 35 cycles of 95 °C (45 sec), 54 °C (45 sec) and 72 °C (1 min) and a final extension at 72 °C (7 min), based on Ekman (2001). The annealing temperature was occasionally altered by ±1 degree in order to obtain a better result. PCR purification and DNA sequencing were accomplished by the genomic research company Macrogen (Seoul, Korea).

Figure 1. Specific collection sites (black symbols) for the new species Bacidia fuscopallida (black star) and four new records, B. ekmaniana (black club), B. friesiana (black diamond), B. heterochroa (black hearth) and B. suffusa (black spade).
Phylogenetic analyses

An independent phylogenetic tree for the genus *Bacidia* was produced from 84 sequences from GenBank and 12 newly-generated sequences for the new species and the new records (Table 2). All ITS sequences were aligned and edited manually using ClustalW in Bioedit v.7.2.6.1 (Hall 1999). All missing and ambiguously aligned data and phylogenetically uninformative positions were removed and phylogenetically informative regions were finally analysed in MEGA X (Stecher et al. 2020). The final alignment comprised 930 bp, in which 102 variable regions were detected. The phylogenetically informative regions were 585. Phylogenetic trees with bootstrap values were obtained in RAxML GUI 2.0 beta (Edler et al. 2019) using the Maximum Likelihood method with a rapid bootstrap with 1,000 bootstrap replications and GTR GAMMA (GTR + G4) for the substitution matrix. The posterior probabilities were obtained in BEAST 2.6.4 (Bouckaert et al. 2019) using the GTR 123454 model, as the appropriate model of nucleotide substitution produced by the Bayesian model averaging methods with bModelTest (Bouckaert and Drummond 2017), empirical base frequencies, gamma for the site heterogeneity model, four categories for gamma and a 10,000,000 Markov Chain Monte Carlo chain length with a 10,000-echo state screening and 1,000 log parameters. Then, a consensus tree was constructed in TreeAnnotator 2.6.4 (Bouckaert et al. 2019) with the first 25% discard as a burn-in, no posterior probability limit, a maximum clade credibility tree for the target tree type and median node heights. All trees were displayed in FigTree 1.4.2 (Rambaut 2014) and edited in Microsoft Paint. Overall analyses in the materials and methods were undertaken based on Lee and Hur (2020).

**Table 1.** Hand-cut section information for DNA isolation.

| Species       | *Bacidia fuscopallida* | *Bacidia ekmaniana* | *Bacidia friesiana* | *Bacidia heterochroa* | *Bacidia suffusa* |
|---------------|------------------------|---------------------|---------------------|-----------------------|------------------|
| Specimens     | KBA-L-0001010 (isotype), KBA-L-0001037 (paratype), KBA-L-0001049 (paratype) | KBA-L-0000072, KBA-L-0002037 | KBA-L-0001910, KBA-L-0001913, KBA-L-0001914, KBA-L-0001917 | KBA-L-0000386, KBA-L-0000391, KBA-L-0002714, KBA-L-0002727, KBA-L-0002734 | KBA-L-0000358, KBA-L-0000359, KBA-L-0000368, KBA-L-0002776, KBA-L-0002778, KBA-L-0002835 |
| Ascomata sections per specimen | 20 | 10 | 20 | 10 | 10 |
| Ascomata sections per species | 60 | 20 | 80 | 40 | 60 |
### Table 2. Species list and DNA sequence information employed for phylogenetic analysis.

| No. | Species          | ITS            | Voucher                      |
|-----|------------------|----------------|------------------------------|
| 1   | Bacidia absistens| AF282085       | Ekman 3223 (BG)              |
| 2   | Bacidia albogranulosa| MK158340    | J. Malicek 9622              |
| 3   | Bacidia albogranulosa| MK158342    | J. Vondrak 11888 (PRA)       |
| 4   | Bacidia arceutina| AF282083       | Ekman 3110 (BG)              |
| 5   | Bacidia arceutina| JQ796851       | LG DNA 579                    |
| 6   | Bacidia areolata  | MH048614       | M-0182592                    |
| 7   | Bacidia aureswaldii| AF282122    | Johansson 20 (UPS)           |
| 8   | Bacidia bagliettiana| AF282123    | Ekman 3137 (BG)              |
| 9   | Bacidia bagliettiana| MG838190    | O-L-175215                   |
| 10  | Bacidia beckhausii| AF282071       | Holien 6744 (TRH)            |
| 11  | Bacidia beckhausii| JF714252       | MSSRF Lichen Herbarium       |
| 12  | Bacidia biatorina | AF282079       | Knutsson 94–148              |
| 13  | Bacidia caligans  | AF282096       | Johansson 21 (UPS)           |
| 14  | Bacidia circumspecta| MH539764    | L-13006                      |
| 15  | Bacidia circumspecta| AF282124    | Ekman L1330 (LD)             |
| 16  | Bacidia cylindrophora| MG926005    | Kurokawa 1692                |
| 17  | Bacidia cylindrophora| MG926006    | Ohmura 7091 (GZU)            |
| 18  | Bacidia diffraecta | AF282090       | Wetmore 26401 (MIN)          |
| 19  | Bacidia diffraecta | MH048620       | Harris 4655-A                |
| 20  | Bacidia ekmaniana | ON352611       | KBA-L-0002037                |
| 21  | Bacidia elongata  | MH048626       | M-0182571                    |
| 22  | Bacidia elongata  | MH048629       | M-0182627                    |
| 23  | Bacidia fraxinea  | AF282088       | Johansson 1620 (BG)          |
| 24  | Bacidia fiesiana  | ON352609       | KBA-L-0001910                |
| 25  | Bacidia fiesiana  | ON352610       | KBA-L-0001913                |
| 26  | Bacidia fiesiana  | MH539765       | L-13159                      |
| 27  | Bacidia fuscopallida| ON352607      | KBA-L-0001010                |
| 28  | Bacidia fuscopallida| ON352608      | KBA-L-0001049                |
| 29  | Bacidia fuscoviridis| AM292665     | Nordin 5058 (UPS)            |
| 30  | Bacidia gigantensis| MT425200      | MCM242                       |
| 31  | Bacidia hemipolia | AF282072       | Toensberg 25091 (BG)         |
| 32  | Bacidia heterochroa| ON352606       | KBA-L-0000386                |
| 33  | Bacidia heterochroa| ON352612       | KBA-L-0002727                |
| 34  | Bacidia heterochroa| ON352613       | KBA-L-0002734                |
| 35  | Bacidia hostheleoides| AF282081    | Seaward 108121               |
| 36  | Bacidia incompta  | AF282092       | Ekman 3144 (BG)              |
| 37  | Bacidia incompta  | MG461697       | KoLRI Udo-32                  |
| 38  | Bacidia kurilensis| MH048612       | M-0182622                    |
| 39  | Bacidia kurilensis| MH048610       | M-0182620                    |
| 40  | Bacidia kurilensis| MH048611       | M-0182621                    |
| 41  | Bacidia laurocerasi| MH048609       | Galanina 424                 |
| 42  | Bacidia laurocerasi subsp. laurocerasi| MN483106       | Spribille 26334 (KLGO)       |
| 43  | Bacidia laurocerasi subsp. laurocerasi| AF282078    | Wetmore 74318 (MIN)          |
| 44  | Bacidia lutescens | MG925952       | Ekman 3655 (BG)              |
| 45  | Bacidia lutescens | AF282082       | Ekman L1161 (LD)             |
| 46  | Bacidia medialis  | AF282102       | Ekman L1193 (LD)             |
| 47  | Bacidia polychoa  | AF282089       | Knutsson 91–215              |
| No. | Species            | ITS         | Voucher            |
|-----|--------------------|-------------|--------------------|
| 48  | Bacidia roSELLa    | AF282086    | Ekman 3117 (BG)    |
| 49  | Bacidia rubella    | AF282087    | Ekman 3021 (BG)    |
| 50  | Bacidia rubella    | HQ650644    | AFTOL-ID 1793      |
| 51  | Bacidia rubella    | JQ796852    | LG DNA 578         |
| 52  | Bacidia rubella    | KX132984    | LIFU076–16         |
| 53  | Bacidia rubella    | MG461695    | AFTOL-ID 1793      |
| 54  | Bacidia rubella    | EU266078    | Hur H06122         |
| 55  | Bacidia rubella    | MH048630    | M-0182581          |
| 56  | Bacidia rubella    | MK158343    | J. Vondrak 12200 (PRA) |
| 57  | Bacidia sabuleTORum| AF282069    | Ekman 3091 (BG)    |
| 58  | Bacidia sachalinENSis | MH048621 | M-0182619          |
| 59  | Bacidia sachalinENSis | MH048625 | M-0182624          |
| 60  | Bacidia schwEinizii| AF282080    | Wetmore 72619 (MIN)|
| 61  | Bacidia schwEinizii| KX151766    | Lendemer 31230A (NY)|
| 62  | Bacidia scopulicola| AF282084    | Ekman 3106 (BG)    |
| 63  | Bacidia sigmoSporae| MW622004   | P.v.d. Boom 55090  |
| 64  | Bacidia sipmani    | JQ796853    | LG DNA 361         |
| 65  | Bacidia sorediata  | KX151772    | Lendemer 33787 (NY)|
| 66  | Bacidia sorediata  | KX151775    | Barton 658 (NY)    |
| 67  | Bacidia squamulosula | MG925955  | Kalb & Kalb in Kalb, Lich. neotrop. 405 |
| 68  | Bacidia subarvolata | MK499342   | MFLU 16-0573       |
| 69  | Bacidia subincompta| AF282125    | Ekman 3413 (BG)    |
| 70  | Bacidia subincompta| KX098342    | WSL DF231          |
| 71  | Bacidia suffusa    | ON352605    | KBA-L-0000359      |
| 72  | Bacidia suffusa    | ON352614    | KBA-L-0002776      |
| 73  | Bacidia suffusa    | ON352615    | KBA-L-0002778      |
| 74  | Bacidia suffusa    | ON352616    | KBA-L-0002835      |
| 75  | Bacidia suffusa    | AF282091    | Wetmore 74771 (MIN)|
| 76  | Bacidia suffusa    | AY756456    | Andersen 99 (BG)   |
| 77  | Bacidia suffusa    | MH048615    | M-0182601          |
| 78  | Bacidia suffusa    | MH048616    | M-0182593          |
| 79  | Bacidia suffusa    | MH048617    | M-0182594          |
| 80  | Bacidia suffusa    | MH048618    | M-0289887          |
| 81  | Bacidia suffusa    | MH048619    | M-0289888          |
| 82  | Bacidia suffusa    | MW728313    | LAH 36839          |
| 83  | Bacidia suffusa    | MW788561    | LAH 36838          |
| 84  | Bacidia vermifera  | AF282109    | Johansson 1619 (BG)|
| 85  | Bacidia vermifera  | KX132992    | LIFU084-16 (verA)  |
| 86  | Bacidia wellingtonii| MG925953  | Ziviagina s.n.     |
| 87  | Bacidia sp.        | AY756133    | KolRI Udo-32       |
| 88  | Bacidia sp.        | KX098339    | WSL DF223          |
| 89  | Bacidia sp.        | KX098340    | WSL DF72           |
| 90  | Bacidia sp.        | KX098341    | WSL DF80           |
| 91  | Bacidia sp.        | MG773660    | Palice 19352       |
| 92  | Biatora bacidioides| MG773663    | Palice 19221       |
| 93  | Biatora bacidioides| MG773664    | Palice 19685       |
| 94  | Biatora pontica    | KF650977    | C. Printzen 6114 (BG)|
| 95  | Biatora pontica    | MK778588    | J. Malicek 10212   |
| 96  | Biatora printzenii | KF650978   | C. Printzen 6837 (BG)|

Overall 96

DNA sequences which were generated for the new species and the new records of *Bacidia* in this study, are presented in bold. All others were obtained from GenBank. The species names are followed by GenBank accession numbers and voucher information. ITS, internal transcribed spacer; Voucher, voucher information.
Figure 2. Phylogenetic relationships amongst available species in the genus *Bacidia*, based on a Maximum Likelihood analysis of the dataset of ITS sequences. The tree was rooted with the sequences of the genus *Biatora*, based on Gerasimova et al. (2018). Maximum Likelihood bootstrap values ≥ 70% and posterior probabilities ≥ 95% are shown above internal branches. Branches with bootstrap values ≥ 90% are shown as thick lines. New sequences produced in this study are presented in bold. All species names are followed by the GenBank accession numbers.
Results and discussion
Phylogenetic analyses

The new species is positioned in the genus *Bacidia* in the ITS tree (Fig. 2). The ITS tree describes *B. fuscopallida*, the new species, being nested with *B. hostheleoides* (Nyl.) Zahlbr., supported by a bootstrap value of 98 and a posterior probability of 1.00 for the branch. *Bacidia fuscopallida* is located in its own clade without any sequences close to it, although *B. fuscopallida* is sister to *B. hostheleoides*.

Taxonomy

*Bacidia fuscopallida* B.G. Lee & T.I. Heo, sp. nov.
MycoBank No: 843830

Fig. 3

**Diagnosis.** *Bacidia fuscopallida* differs from *B. diffracta* by generally non-granular, olivish-green thallus, pale yellow-orange apothecia without pruina, the absence of crystals in proper exciple, slightly narrower ascospores with up to 15-septation and smaller pycnidia and pycnoconidia.

**Type.** South Korea, Gangwon Province, Gangneung, Okgye-myeon, Mt. Seokbyung, 37°34.45'N, 128°55.00'E, 271 m alt., on bark of *Acer pictum* var. *mono* (Maxim.) Maxim. ex Franch., 17 June 2020, B.G. Lee & H.J. Lee 2020-000811, with *Porina hirsuta* Aptroot & K.H. Moon (holotype: KBA-L-0001011!); same locality, on bark of *Acer pictum* var. *mono*, 17 June 2020, B.G. Lee & H.J. Lee 2020-000801 (isotype: KBA-L-0001001); same locality, on bark of *Acer pictum* var. *mono*, 17 June 2020, B.G. Lee & H.J. Lee 2020-000806, with *Mikhtomia gordejevii* (Tomin) S.Y. Kondr., Kärnefelt, Elix, A. Thell, Jung Kim, A.S. Kondr. & Hur, *Straminella varia* (Hoffm.) S.Y. Kondr., Lökös & Farkas, *Phaeophyscia limbata* (Poelt) Kashiw., *Porina hirsuta* (isotype: KBA-L-0001006); same locality, on bark of *Acer pictum* var. *mono*, 17 June 2020, B.G. Lee & H.J. Lee 2020-000810 (isotype: KBA-L-0001010; GenBank ON352607 for ITS); South Korea, Gangwon Province, Gangneung, Okgye-myeon, Mt. Seokbyung, 37°34.39'N, 128°55.01'E, 349 m alt., on bark of *Quercus mongolica* Fisch. ex Ledeb., 17 June 2020, B.G. Lee & H.J. Lee 2020-000837, with *Opeltia flavorubescens* (Huds.) S.Y. Kondr. & Hur (paratype: KBA-L-0001037); South Korea, Gangwon Province, Gangneung, Okgye-myeon, Mt. Seokbyung, 37°34.28'N, 128°54.88'E, 438 m alt., on bark of *Acer triflorum* Kom., 17 June 2020, B.G. Lee & H.J. Lee 2020-000849, with *Biatora pacifica* Printzen, Tønsberg & G. Thor (paratype: KBA-L-0001049; GenBank ON352608 for ITS).

Thallus corticolous, crustose, areoles in young stage and soon coarsely continuous or warty on aging, often overlapping for each other, rarely granular, thin when not overlapping, olivish-green, margin indeterminate, 40–90 μm thick; cortex indistinct, hyaline, up to 5 μm thick; medulla a little shown as mycelia below algal layer; photobiont chlorococcoid, cells globose to subglobose, 5–15 μm thick, algal layer
composing most part of thallus, 35–80 μm thick. Prothallus indistinct or whitish-grey and endosubstratal when present.

Apothecia numerous, solitary, marginate and flat in young stage and seeming im-marginate and convex on aging (consistently marginate and flat on bark of Acer triflorum), 0.1–0.7 mm diam. (mean = 0.33; SD = 0.14; n = 105). Pruina absent. Disc biatorine, thalline exciple absent, pale yellow to pale orange in young stage and slightly more blackish generally around margin when mature (much more blackish on bark of A. triflorum and Q. mongolica from young stage). Proper exciple 65–80 μm wide laterally (SD = 5.7; n = 15), with radiating hyphae of 1–2.5 μm wide (SD = 0.5; n = 10) and outermost cell 2.5–4 μm wide (SD = 0.6; n = 10), hyaline to pale yellow around rim, but darker downwards (pale yellow to pale brown) and the dark colour extending to hypothecium. Epihymenium hyaline, with a little pigment of pale yellow to pale olive-brown locally, smooth and not granular, ca. 5 μm high. Hymenium hyaline, 70–100 μm high (SD = 8.9; n = 10). Hypothecium clearly pigmented, pale orange-brown to brown, prosoplectenchymatous (irregularly arranged), 70–130 μm high (SD = 18.9; n = 10). Crystals absent or a little present in upper hypothecium. Oil droplets absent. Paraphyses simple, rarely branched at tips, 1–1.5 μm wide, tips not or little swollen, not pigmented, 1.5–2 μm wide. Asci cylindrical to narrowly clavate, 8-spored, 49–72 × 11–14 μm (SD = 7.3 (L), 0.9 (W); n = 11). Ascospores 3- to 15-septate, acicular to filiform, 24–69 × 2–3.5 μm (mean = 52.8 × 2.6 μm; SD = 8.7 (L), 0.6 (W); L/W ratio = 3.8–30.5, ratio mean = 17.6, ratio SD = 5.0; n = 104). Pycnidia black, immersed and upper half only shown, globose, 60–65 μm high and 55–75 μm wide (SD = 2.4 (H), 8.2 (W); n = 5), with brownish wall, K–. Pycnoconidia hyaline, filiform, curved or almost straight, 6–17 × 0.3–0.5 μm (mean = 10.4 × 0.5 μm; SD = 2.9 (L), 0.1 (W); n = 53).

Chemistry. Thallus K– or K+ slightly yellow, KC–, C–, Pd–, UV–. Epihymenium K+ purple extending to outermost layers of proper exciple, C–. No lichen substance was detected by TLC.

Distribution and ecology. The species occurs on barks of Acer pictum var. mono, A. triflorum and Quercus mongolica. The species is currently known from the type collections.

Etymology. The species epithet indicates the pale brown colour of the lichen’s apothecia.

Notes. The new species is similar to B. diffracta and B. polychroa (Th. Fr.) Körb. in having colourless epihymenium with pale orange-brown pigment and K+ purple reaction, distinctly pigmented hypothecium with yellow, orange or brown, long ascospores generally with L/W ratio over 10 amongst corticolous species. However, B. diffracta differs from the new species by granular thallus, darker and larger apothecia with pruina, proper exciple with radiating clusters of minute crystals, slightly wider ascospores with up to 11-septation and larger pycnidia and pynoconidia (Ekman 1996) (Table 3). The new species is more similar to B. polychroa in having coarsely continuous or warded thallus. However, B. polychroa differs from the new species by greyish thallus, darker and larger apothecia often with pruina, proper exciple often with radiating clusters of minute crystals, wider ascospores and larger pycnidia and pynoconidia (Ekman 1996; Smith et al. 2009) (Table 3).
### Table 3. Comparison of the new species with close species in the genus Bacidia.

| Species                        | Bacidia fuscopallida | Bacidia diffracta | Bacidia hosteleoides | Bacidia polychroa | Bacidia purpurans |
|--------------------------------|----------------------|-------------------|----------------------|-------------------|------------------|
| Thallus growth form            | warty, rarely granular| finely granular   | wrinkled or granular to subsquamulose | finely wrinkled to warted, sometimes areolate | areolate         |
| Thallus colour                 | olivish-green        | pale grey-green-grey-grey-yellow-grey to grey yellow-grey to pale green-grey | pale grey to pale green-grey | white to grey or yellow-grey | pale grey-green to dark green |
| Prothallus                     | white-grey around margin, endosubstratal | white-pale grey between granules, endosubstratal | absent | – | white, arachnoid |
| Apothecia (mm in diam.)        | 0.1–0.7              | 0.5–1.1           | 0.5–0.8              | 0.4–1.2          | –                |
| Disc colour                    | pale yellow to pale orange (young); more blackish (old) | brown-orange to dark brown | brown-orange | brown-orange to dark brown | dark purple-brown to brown |
| Pruina                         | absent               | white             | absent               | white            | absent           |
| Crystals in proper exciple     | absent               | radiating clusters of minute crystals | absent | with or without radiating clusters of minute crystals | absent           |
| Crystals in hymenium           | small crystals at bottom | –                 | –                   | –                | –                |
| Epiphyemenium colour           | colourless with pale yellow-brown pigment | colourless with pale orange-brown pigment | very pale orange | colourless with brown-orange pigment | greyish          |
| Hymenium height (μm)           | 70–100               | 70–100            | ca. 60               | 55–100           | 100              |
| Hypothecium colour             | pale orange-brown to brown | pale brown to orange-brown | very pale orange | brown-orange to dark brown | orange-brown |
| Hypothecium height (μm)        | 70–130               | –                 | –                   | –                | ca. 60           |
| Ascospore (μm)                 | 24–69 × 2–3.5        | 32–69 × 1.9–4.1   | 16–25 × 2.9–5       | 31–74 × 1.9–5   | 50–75 × 2–4     |
| Ascospore L/W ratio             | 4–31                 | 9–27              | 4–9                 | 7–30             | –                |
| Ascospore septation            | 3–15                 | 3–11              | 3–5                 | 2–15             | 3–15            |
| Pycnidia (μm)                  | 55–75                | 150               | 50–100              | 100–170          | 150–200         |
| Pycnoconidia                   | 6–17 × 0.3–0.5       | 10–15 × 0.5–0.6   | 10–17 × 0.6–0.8     | 20–25 × 0.8     |                  |
| Substance                      | absent               | atranorin, (trace of zeorin) | absent | (trace of atranorin) | atranorin |
| Reference                      | this study           | Ekman (1996)      | Ekman (1996)         | Ekman (1996); Smith et al. (2009) | Lendemer et al. (2016) |

The morphological and chemical characteristics of several species close to the new species are referenced from the previous literature. All information on the new species is produced from type specimens (KBA-L-0001010, KBA-L-0001011 and KBA-L-0001049) in this study.

**Figure 3.** *Bacidia fuscopallida* (KBA-L-0001011, holotype for **A–D, G–O** KBA-L-0001049 for **E, F** KBA-L-0001010 for **P, Q**) in morphology **A, B** habitus and apothecia on bark of *Acer pictum* var. *mono*. Olive-green thallus and pale yellow-orange apothecia **C** vertical section of apothecia **D** prothallus present around margin of habitus (red arrows) **E, F** habitus and apothecia growing on bark of *Acer triflorum* **G** apothecial section **H** epiphyemenium colourless or a little pigmented **I** epiphyemenium K+ purple **J** small crystals (red arrows) present in upper epiphyemenium **K** proper exciple pigmented with pale or colourless margin. Radiating hyphae wider to margin **L** photobiont composing most part of thallus **M, N** asci cylindrical to narrowly clavate. Ascospores not twisted in ascus **O** ascospores aciculat to filiform up to 15-septate **P** pycnidia globose with brown wall **Q** pycnoconidia curved or almost straight. Scale bars: 1 mm (**A, E**); 500 μm (**B, C, F**); 2 mm (**D**); 200 μm (**G**); 50 μm (**H–J, P**); 20 μm (**K, L**); 10 μm (**M–O, Q**).
The new species is quite similar to *B. purpurans* R. C. Harris, Ladd & Lendemer in having greenish thallus with areoles and K+ purple reaction in epihymenium. However, *B. purpurans* differs from the new species by arachnoid prothallus, darker apothecia, green excipular rim adjacent to epihymenium, greyish epihymenium, shorter hypothecium, absence of crystals, larger ascospores and larger pycnidia and pycnoconidia (Lendemer et al. 2016) (Table 3).

The new species can be compared with *B. hostheleoides* in sharing non-pruinose apothecia and proper exciple without crystals. However, *B. hostheleoides* differs from the new species by greyish thallus, absence of prothallus, shorter hymenium, paler hypothecium and shorter ascospores with a few septa (Ekman 1996) (Table 3).

**Bacidia ekmaniana** R. C. Harris, Ladd & Lendemer, The Bryologist 119 (2): 154 (2016)

**Fig. 4**

**Description.** Thallus corticolous, crustose, somewhat granular when young and smoother when mature, grey, greenish-grey to pale grey, margin indeterminate. Prothallus generally not detected or whitish-grey when present.

Apothecia consistently flat or slightly convex when mature, marginate, without pruina, 0.4–1.4 mm diam. (mean = 0.75, SD = 0.23, n = 104). Disc bitorine, without thalline exciple, pale straw, light brown to brown, with a distinct proper margin which is smooth to rugose and becoming thinner but still distinct when mature. Proper exciple pale brown to red-brown, paler or colourless around rim and thicker downwards, 80–120 μm wide laterally. Epithymenium hyaline, smooth but not granular, ca. 5 μm high. Hymenium hyaline, 80–140 μm high. Hypothecium red-brown, prosoplectenchymatous (irregularly arranged), 120–250 μm high. Small crystals present a little in hypothecium, dissolving in K. Oil droplets absent. Asci narrowly clavate, 8-spored, 70–105 × 8–12 μm (n = 5). Ascospores acicular to filiform, cells near head sometimes irregularly swollen, 3- to 9-septate, 52–71 × 2–4.5 μm (n = 15). Pycnidia not detected.

**Chemistry.** Thallus K–, C–. Apothecial section K–, C–. No lichen substance was detected by TLC.

**Notes.** *Bacidia ekmaniana* is easily confused with *B. schweinitzii* under the microscope, as well as in the field because both species often share their habitat and the habit of both species look similar particularly when the ascomata of the latter are paler. Both species are often detected from one specimen under the microscope and those were frequently regarded as one species, i.e. *B. schweinitzii*. Generally, however, *B. ekmaniana* differs from the latter by paler ascomata. *Bacidia ekmaniana* has brown but not black apothecia when mature (Lendemer et al. 2016). *Bacidia ekmaniana* differs from the latter by colourless epihymenium and paler hypothecium as well.

*Bacidia ekmaniana* is more similar to *B. arceutina* than *B. schweinitzii* in morphology in having pale ascomata. However, *B. ekmaniana* differs from *B. arceutina* by the colourless to pale excipular rim, colourless epihymenium and wider ascospores with
Korean Bacidia species

more septation (Ekman 1996; also see the key couplet 23). Bacidia ekmaniana is new to Asia and this is the second record after North America (Lendemer et al. 2016). Bacidia ekmaniana is supposed to occur widespread throughout the world as the species was assumed to be B. schweinitzii in the past. Phylogenetic analysis resulted in B. ekmaniana being located in its own clade in the genus Bacidia (Fig. 2).
Specimens examined. South Korea, North Gyeongsang Province, Bonghwa, Chunyang-myeon, Mt. Munsu, 36°59.28’N, 128°48.17’E, 1,058 m alt., on bark of Quercus mongolica, 29 August 2019, B.G. Lee 2019-000072 (KBA-L-0000072); South Korea, South Jeolla Province, Gokseong, Jukgok-myeon, Taensa Temple, 35°08.06’N, 127°23.26’E, 297 m alt., on bark of Salix pierotii Miq., 25 May 2020, B.G. Lee 2020-000212, with Bacidia schweinitzii (KBA-L-0000412); same locality, on bark of Salix pierotii, 25 May 2020, B.G. Lee 2020-000227, with Bacidia schweinitzii, Coenogonium pineti (Ach.) Lücking & Lumbsch, Phaeophyscia rubropulchra (Degel.) Moberg, Porina melanops Malme (KBA-L-0000427); same locality, on bark of Idesia polycarpa Maxim., 25 May 2020, B.G. Lee 2020-000231, with Bacidia schweinitzii, Porina aff. melanops (KBA-L-0000431); same locality, on bark of Taxicodendron vernicifluum (Stokes) F. A. Barkley, 25 May 2020, B.G. Lee 2020-000233, with Biatora aff. pacifica, Lecidea sp., Phaeophyscia rubropulchra, Rinodina sp., Traponora varians (Ach.) J. Kalb & Kalb (KBA-L-0000433); South Korea, North Gyeongsang Province, Bonghwa, Chunyang-myeon, Mt. Okseok, 37°00.91’N, 128°46.65’E, 1,085 m alt., on bark of Quercus mongolica, 15 September 2020, B.G. Lee & H.J. Lee 2020-001159, with Anisomeridium polypori (Ellis & Everh.) M.E. Barr, Bacidia schweinitzii, Rinodina sp. (KBA-L-0001359); same locality, on bark of Quercus mongolica, 15 September 2020, B.G. Lee & H.J. Lee 2020-001162, with Rinodina sp. (KBA-L-0001362); South Korea, North Jeolla Province, Jangsu, Mt. Youngchui, 35°38.59’N, 127°37.00’E, 907 m alt., on bark of Carpinus tschonoskii Maxim., 08 June 2021, B.G. Lee & H.J. Lee 2021-000563, with Lecanora megalocheila (Hue) H. Miyaw., Rinodina orientalis Sheard (KBA-L-0002035); same locality, on bark of Carpinus tschonoskii, 08 June 2021, B.G. Lee & H.J. Lee 2021-000565, with Arthonia apatetica (A. Massal.) Th. Fr., Lecidella euphora (Flörke) Kremp. (KBA-L-0002037; GenBank ON352611 for ITS); same locality, on bark of Carpinus tschonoskii, 08 June 2021, B.G. Lee & H.J. Lee 2021-000569, with Anisomeridium polypori, Lecidella euphora, Rinodina orientalis, Scoliciosporum sp. (KBA-L-0002041); same locality, on bark of Carpinus tschonoskii, 08 June 2021, B.G. Lee & H.J. Lee 2021-000573, with Arthonia apatetica, Lecanora aff. imshaugii Brodo, Lecidella euphora, Porina hirsuta (KBA-L-0002045); South Korea, North Jeolla Province, Jangsu, Mt. Jangan, 35°38.58’N, 127°36.96’E, 925 m alt., on bark of Carpinus tschonoskii, 09 June 2021, B.G. Lee & H.J. Lee 2021-000759 (KBA-L-0002231); same locality, on bark of Carpinus tschonoskii, 09 June 2021, B.G. Lee & H.J. Lee 2021-000760 (KBA-L-0002232); same locality, on bark of Carpinus tschonoskii, 09 June 2021, B.G. Lee & H.J. Lee 2021-000766, with Lecania sp., Phaeophyscia adiastola (Essl.) Essl., Porina hirsuta, Rinodina orientalis, Scoliciosporum chlorococcum (Graewe ex Stenh.) Vězda (KBA-L-0002232); same locality, on bark of Carpinus tschonoskii, 09 June 2021, B.G. Lee & H.J. Lee 2021-000766, with Lecanora sp., Phaeophyscia sp., Rinodina orientalis (KBA-L-0002238); South Korea, North Jeolla Province, Jangsu, Mt. Baegun, 35°36.76’N, 127°36.85’E, 661 m alt., on bark of Cornus walteri Wangetin, 10 June 2021, B.G. Lee & H.J. Lee 2021-000926 (KBA-L-0002398); same locality, on bark of Cornus walteri, 10 June 2021, B.G. Lee & H.J. Lee 2021-000927 (KBA-L-0002399); same locality, on bark of Cornus walteri, 10 June 2021, B.G. Lee & H.J. Lee 2021-000928 (KBA-L-0002400); same locality, on bark of Cornus wal-
Korean Bacidia species

Bacidia friesiana (Hepp) Körb., Parerga lichenol. (Breslau) 2: 133 (1860) [1865]

Fig. 4

Description. Thallus corticolous, crustose, thin, little developed or indistinct, generally not continuous, minutely granular with contiguous granules when developed, pale grey with slightly brownish colour, margin indeterminate. Prothallus not detected.

Apothecia consistently flat or convex when mature, marginate, without pruina, 0.1–0.5 mm diam. (mean = 0.23, SD = 0.07, n = 107). Disc bitorine, without thalline exciple, pale pink to pale yellow when young and darker (particularly around margin) when mature. Proper exciple hyaline with or without pale brown pigment, the pigment slightly thicker close to hymenium or excipular rim, 40–50 μm wide laterally. Epiphymenium bluish-green, ca. 5 μm high. Hymenium hyaline, 40–45 μm high. Hypothecium hyaline, 50–60 μm high; upper hypothecium paraplectenchymatous (globular to angular), lower hypothecium prosoplectenchymatous (periclinaly or irregularly arranged). Crystals or oil droplets absent. Ascii narrowly clavate, 8-spored, 39–41 × 10–12 μm (n = 3). Ascospores acicular to filiform, 3- or 7-septate, 28–38 × 1.5–2.5 μm (n = 14). Pycnidia not detected.

Chemistry. Epiphymenium K–, C–. Hymenium K– or a few undeveloped asci K+ purplish. No lichen substance was detected by TLC.

Notes. Bacidia friesiana is similar to B. circumspecta (Norrl. & Nyl.) Malme and B. ignarii (Nyl.) Oxner (syn. Scutula ignarii (Nyl.) S. Ekman) in having epiphymenium with green pigments, proper exciple without crystals and dark hypothecium amongst corticolous species. However, B. friesiana differs from the latter two by the excluded margin of apothecia and acicular ascospores. The latter species have a permanent margin of apothecia and bacilliform or clavate ascospores (Ekman 1996).

Phylogenetic analysis resulted in B. friesiana of Korea (ON352609 and ON352610) being nested with the sequences of Russia (MH539765), supported by a bootstrap value of 100 and a posterior probability of 1.00 for the branch (Fig. 2). Bacidia friesiana was previously reported from Europe, North America and Russian Far East (Smith et al. 2009; Gerasimova et al. 2018). This is a new record to Korea.

Specimens examined. SOUTH KOREA, Gangwon Province, Yanggu, Nam-myeon, Dumu-ri, nearby a forested wetland, 38°02.12’N, 128°05.14’E, 421 m alt., on bark of Salix piorotii, 28 April 2020, B.G. Lee 2020-0000164, with Mikhtomia gordejevii, Canodelaria concolor (Dicks.) Arnold, Phaeophyscia adiastola, Porina cf. melanops, Rinodina cf. subminuta (KBA-L-0000364); SOUTH KOREA, Gyeonggi Province, Yangpyeong, Cheonggun-myeon, Dowon-ri, a forested wetland, 37°32.55’N, 127°48.60’E, 443 m alt., on bark teri, 10 June 2021, B.G. Lee & H.J. Lee 2021-000929, with Phaeophyscia adiastola (KBA-L-0002401); same locality, on bark of Cornus walteri, 10 June 2021, B.G. Lee & H.J. Lee 2021-000930, with Phaeophyscia rubropulchra (KBA-L-0002402); same locality, on bark of Cornus walteri, 10 June 2021, B.G. Lee & H.J. Lee 2021-000931, with Lecanora sp., Phaeophyscia adiastola (KBA-L-0002403); same locality, on bark of Cornus walteri, 10 June 2021, B.G. Lee & H.J. Lee 2021-000932 (KBA-L-0002404).
of *Salix pierotii*, 31 May 2021, B.G. Lee & H.J. Lee 2021-000438, with *Lecidella euphora*, *Phaeophyscia adiastola*, *Rinodina orientalis* (KBA-L-0001910; GenBank ON352609 for ITS); same locality, on bark of *Aralia elata* (Miq.) Seem., 31 May 2021, B.G. Lee & H.J. Lee 2021-000440, with *Lecidella euphora*, *Phaeophyscia adiastola*, *Traponora varians* (KBA-L-0001912); same locality, on bark of *Aralia elata*, 31 May 2021, B.G. Lee & H.J. Lee 2021-000441, with *Hyperphyscia adglutinata* (Flörke) H. Mayrhofer & Poelt, *Rinodina orientalis* (KBA-L-0001913; GenBank ON352610 for ITS); same locality, on bark of *Aralia elata*, 31 May 2021, B.G. Lee & H.J. Lee 2021-000442, with *Hyperphyscia adglutinata*, *Rinodina orientalis* (KBA-L-0001914); same locality, on bark of *Aralia elata*, 31 May 2021, B.G. Lee & H.J. Lee 2021-000443, with *Rinodina orientalis*, *Traponora varians* (KBA-L-0001915); same locality, on bark of *Aralia elata*, 31 May 2021, B.G. Lee & H.J. Lee 2021-000444, with *Phaeophyscia adiastola*, *P. rubropulchra*, *Rinodina orientalis* (KBA-L-0001916); same locality, on bark of *Aralia elata*, 31 May 2021, B.G. Lee & H.J. Lee 2021-000445 (KBA-L-0001917).

*Bacidia heterochroa* (Müll. Arg.) Zahlbr., *Cat. Lich. Univers.* 4: 204 (1926) [1927]

**Fig. 4**

**Description.** Thallus corticolous, crustose, continuous, wrinkled, or warted, pale yellowish-grey, margin indeterminate or determinate. Prothallus generally not present or locally present as blackish bordering a different lichen.

Apothecia flat, marginate, without pruina, 0.2–0.6 mm diam. (mean = 0.33, SD = 0.11, n = 72). Disc lecideine, without thalline exciple, blackish or reddish-black. Proper exciple hyaline with pale brown pigment dispersed, pigment slightly thicker close to hymenium, 80–100 μm wide laterally. Epihymenium brown to dark brown, ca. 10 μm high. Hymenium hyaline, 80–95 μm high. Hypothecium hyaline, 80–120 μm high, with a little pale yellow pigment. Crystals or oil droplets absent. Asci narrowly clavate to cylindrical, 8-spored, 42–48 × 12–13 μm (n = 3). Ascospores acicular to filiform, 9- or 10-septate, 36–67 × 2.5–4 μm (n = 11). Pycnidia not detected.

**Chemistry.** Epihymenium K+ purple or intensifying, extending to excipular rim. No lichen substance was detected by TLC.

**Notes.** *Bacidia heterochroa* is the most similar to *B. laurocerasi* (Delise ex Duby) Zahlbr. in having smooth thallus without granules, absence of crystals in exciple, epihymenium without green pigments, pale to colourless hypothecium, K+ purple in apothecial section and narrow ascospores less than 4 μm wide amongst corticolous species. However, *B. heterochroa* differs from *B. laurocerasi* by distinctly brown-pigmented paraphysial tips, less than 16-septate ascospores which are shorter but wider (less than 80 μm long but over 3.5 μm wide) and substrate preference to deciduous trees or shrubs (Ekman 1996; Brodo 2016; also see the key couplet 21).

Phylogenetic analysis resulted in *B. heterochroa* of Korea (ON352606, ON352612 and ON352613) being nested in a sister clade to *B. laurocerasi*, supported by a bootstrap value of 75 without a posterior probability as the Maximum Likelihood analysis did not match with the Bayesian Inference for the clade. The sequences of *B. het-
Bacidia were not compared with previous records due to the lack of data (Fig. 2). *Bacidia heterochroa* was previously reported from Thailand in Asia (Aptroot et al. 2007) and this is a new record to northeastern Asia.

**Specimens examined.** SOUTH KOREA, Gangwon Province, Yanggu, Nam-myeon, Dumu-ri, a forested wetland, 38°02.12’N, 128°05.14’E, 421 m alt., on bark of *Salix koriyana-gr* Kimura ex Goerz, 28 April 2020, B.G. Lee 2020-000186 (KBA-L-0000386; GenBank ON352606 for ITS); SOUTH KOREA, South Jeolla Province, Damyang, Changpyeong-myeon, Oedong-ri, a forested wetland, 35°12.00’N, 127°00.88’E, 338 m alt., on bark of *Fraxinus rhynchophylla* Hance, 12 May 2021, B.G. Lee & D.Y. Kim 2021-000214 (KBA-L-0001686); SOUTH KOREA, Gangwon Province, Jeongseon, Imgye-myeon, Gamok-ri, a forested wetland, 37°32.47’N, 128°57.72’E, 760 m alt., on bark of *Acer tartaricum* subsp. *ginnala* (Maxim.) Wesm., 17 June 2021, B.G. Lee & H.J. Lee 2021-001241, with *Lecanora chionocarpa* Hue (KBA-L-0002713); same locality, on bark of *Acer tartaricum* subsp. *ginnala*, 17 June 2021, B.G. Lee & H.J. Lee 2021-001242, with *Phaeophyscia adiastola* (KBA-L-0002714); same locality, on bark of *Acer tartaricum* subsp. *ginnala*, 17 June 2021, B.G. Lee & H.J. Lee 2021-001255, with *Opeltia flavorubescens*, *Phaeophyscia adiastola* (KBA-L-0002727; GenBank ON352612 for ITS); same locality, on bark of *Acer tartaricum* subsp. *ginnala*, 17 June 2021, B.G. Lee & H.J. Lee 2021-001257, with *Hyperphyscia adglutinata*, *Lecidella euphorea* (KBA-L-0002729); same locality, on bark of *Acer tartaricum* subsp. *ginnala*, 17 June 2021, B.G. Lee & H.J. Lee 2021-001262, with *Lecidella euphorea*, *Phaeophyscia adiastola*, *Rinodina orientalis* (KBA-L-0002734; GenBank ON352613 for ITS); same locality, on bark of *Acer tartaricum* subsp. *ginnala*, 17 June 2021, B.G. Lee & H.J. Lee 2021-001263, with *Opeltia flavorubescens*, *Phaeophyscia adiastola*, *Rinodina orientalis* (KBA-L-0002735); same locality, on bark of *Acer tartaricum* subsp. *ginnala*, 17 June 2021, B.G. Lee & H.J. Lee 2021-001267, with *Lecidella euphorea*, *Porina hirsuta*, *Rinodina orientalis*, *Straminella varia* (KBA-L-0002739); same locality, on bark of *Acer tartaricum* subsp. *ginnala*, 17 June 2021, B.G. Lee & H.J. Lee 2021-001269, with *Lecidella euphorea*, *Opeltia flavorubescens*, *Phaeophyscia rubropulchra*, *Rinodina orientalis* (KBA-L-0002741).

*Bacidia suffusa* (Fr.) A. Schneid., *Guide Study Lich.*: 110 (1898)

**Fig. 4**

**Description.** Thallus corticolous, crustose, continuous, wrinkled, warted or subquamulose, often granular locally, whitish pale grey. Prothallus generally not present or present as dark brown to black between different colonies.

Apothecia flat, marginate, with a little or heavy white pruina, generally more pruinose at margin, 0.3–1.7 mm diam. (mean = 0.75, SD = 0.28, n = 116). Disc lécidéine, without thalline exciple, brown to dark brown. Proper exciple with radiating clusters of crystals produced around hypothecium and expanding to excipular rim and finally shown as pruina on surface, hyaline downwards but brown around rim, the brown concolorous or slightly paler to ephymenium, 80–100 μm wide laterally. Ephymenium brown to dark brown, ca. 10 μm high, with pruina (ca. 10 μm high) on surface. Hymenium hyaline, 70–80 μm high. Hypothecium hyaline, 80–100 μm
high. Other small crystals present a few in upper hypothecium. Oil droplets absent. Asci cylindrical, 8-spored, 65–75 × 10–16 μm (n = 7). Ascospores acicular to filiform, up to 13-septate, 45–70 × 2.5–4.5 μm (n = 10). Pycnidia not detected.

**Chemistry.** Thallus K+ yellow, KC–, C–, Pd–, UV–. Epihymenium K–. Atranorin was detected by TLC.

**Notes.** *Bacidia suffusa* is the most similar to *B. russeola* (Kremp.) Zahlbr. in having dark apothecia, generally colourless epihymenium without green pigment, long ascospores with the L/W ratio over 11, pale or colourless hypothecium and K+ purple reaction on epihymenium and nearby excipular rim amongst corticolous species. However, *B. suffusa* differs from *B. russeola* by the presence of pruina on the disc and in proper exciple as radiating clusters of crystals and more than 10-septate ascospores (Ekman 1996).

Phylogenetic analysis resulted in *B. suffusa* of Korea (ON352605, ON352614, ON352615 and ON352616) being nested in a sister clade of the sequences of Pakistan (MW728313 and MW788561), Russia (MH048615, MH048616 and MH048617) or U.S.A. (MH048618 and MH048619). The molecular data of Korea converged into the previous data of *B. suffusa*, supported by a bootstrap value of 100 and a posterior probability of 1.00 for the branch (Fig. 2). *Bacidia suffusa* was previously detected from North America, North Caucasus, Russian Far East and Pakistan, but rare or absent in Europe (Otte 2007; Gerasimova et al. 2018, 2021; Adrees et al. 2022). This is a new record to Korea.

**Specimens examined.** South Korea, Gangwon Province, Yanggu, Nam-myeon, Dumu-ri, a forested wetland, 38°02.12’N, 128°05.14’E, 421 m alt., on bark of *Salix pierotii* Miq., 28 April 2020, B.G. Lee 2020-000158 (KBA-L-0000358); same locality, on bark of *Salix pierotii*, 28 April 2020, B.G. Lee 2020-000159 (KBA-L-0000359); GenBank ON352605 for ITS); same locality, on bark of *Salix pierotii*, 28 April 2020, B.G. Lee 2020-000168, with *Candelaria concolor*, *Phaeophyscia adiastola*, *Phaeophyscia hirtuosa* (Kremp.) Essl. (KBA-L-0000368); South Korea, Gangwon Province, Gangneung, Okgye-myeon, Mt. Seokbyung, 37°34.45’N, 128°55.01’E, 271 m alt., on bark of *Acer pictum* var. mono, 17 June 2020, B.G. Lee & H.J. Lee 2020-000799 (KBA-L-0000999); South Korea, Gangwon Province, Jeongseon, Imgye-myeon, Gamok-ri, a forested wetland, 37°32.47’N, 128°57.72’E, 760 m alt., on bark of *Fraxinus chiisanensis* Nakai, 17 June 2021, B.G. Lee & H.J. Lee 2021-001304, with *Normandina pulchella* (Borrer) Nyl., *Phaeophyscia* sp. (KBA-L-0002776; GenBank ON352614 for ITS); same locality, on bark of *Fraxinus chiisanensis*, 17 June 2021, B.G. Lee & H.J. Lee 2021-001305, with *Anisomeridium polypori*, *Normandina pulchella*, *Phaeophyscia* sp., *Porina hirsuta* (KBA-L-0002777); same locality, on bark of *Fraxinus chiisanensis*, 17 June 2021, B.G. Lee & H.J. Lee 2021-001306, with *Normandina pulchella*, *Opeltia flavorubescens*, *Phaeophyscia adiastola* (Essl.) Essl. (KBA-L-0002778; GenBank ON352615 for ITS); same locality, on bark of *Fraxinus chiisanensis*, 17 June 2021, B.G. Lee & H.J. Lee 2021-001308, with *Phaeophyscia adiastola* (KBA-L-0002780); same locality, on bark of *Fraxinus chiisanensis*, 17 June 2021, B.G. Lee & H.J. Lee 2021-001320, with *Opeltia flavorubescens* (KBA-L-0002792); same locality, on bark of *Acer tartaricum* subsp. ginatala, 17 June 2021, B.G. Lee & H.J. Lee 2021-001363 (KBA-L-0002835; GenBank ON352616 for ITS).
## Key to the species of *Bacidia* in Korea (19 taxa)

The key is composed of all 19 species in the genus *Bacidia* of Korea, including synonyms in *Bacidina* and *Toniniopsis* species.

| 1 | Epiphymenium with green pigment | 2 |
|---|---------------------------------|---|
| 2 | Epiphymenium colourless, yellow-brown, brown to dark brown, but without green pigment | 5 |
| 3 | Proper exciple with radiating clusters of coarse crystals (up to 7 μm wide); hymenium ca. 100 μm high; ascospores 40–68 × 2.5–3 μm; atranorin present | *B. schweinitzii* |
| 4 | Proper exciple without crystals; hymenium less than 70 μm high; ascospores less than 50 μm long; without substance | 3 |
| 5 | Hypothecium colourless to pale blue-green; thallus pale grey to pale brown-grey without green colour | *B. friesiana* |
| 6 | Proper exciple colourless to brown, dark red-brown; thallus grey-green to green-brown | 4 |
| 7 | Proper exciple with green pigment at rim, pale to colourless downwards; hypothecium K– or K+ green-brown; generally on rock or occasionally on bark or moss | *B. egenula* (*Bacidina egenula*) |
| 8 | Proper exciple colourless at rim, red-brown to black-brown downwards; hypothecium K+ purple; on bark | *B. subincompta* (*Toniniopsis subincompta*) |
| 9 | On rock | 6 |
| 10 | On bark or wood | 12 |
| 11 | Apothecia pruinose | 7 |
| 12 | Apothecia not pruinose | 8 |
| 13 | Thallus coarsely granular without forming soredia; apothecia 0.7–1.2 mm diam.; hymenium 70–100 μm high; hypothecium colourless to pale yellow or pale orange; ascospores 40–70 × 2.5–3 μm, 3- to 7-septate | *B. rubella* |
| 14 | Thallus granular with soredia; apothecia 0.3–0.7 mm diam.; hymenium 40–50 μm high; hypothecium orange-brown to dark red-brown; ascospores 24–46 × 1–2 μm, 1- to 3-septate | *B. arnoldiana* (*Bacidina arnoldiana*) |
| 15 | Disc brown, red-brown to black; hypothecium pale brown to dark brown | 9 |
| 16 | Disc pale yellow, pale orange to dark brown; hypothecium colourless to pale yellow or pale orange | 10 |
| 17 | Proper exciple dark coloured; ascospores 25–35 × 6–10 μm, with L/W ratio less than 10 | *B. hakonensis* |
| 18 | Proper exciple colourless to pale brown; ascospores 24–46 × 1–2 μm, with L/W ratio over 10 | *B. arnoldiana* (*Bacidina arnoldiana*) |
| 19 | Thallus rimose, wrinkled or warted, but not granular; disc pale yellow or pale grey; epiphymenium K– | *B. chloroticula* (*Bacidina chloroticula*) |
| 20 | Thallus granular; disc pale to dark brown; epiphymenium K+ purple | 11 |
11 Thallus granular forming isidia- or coral-like structures; prothallus absent; apothecia flat; ascospores 25–34 × 1.1–1.9 μm; occasionally on old wood.................\textit{B. egenuloidea (Bacidina egenuloidea)}

12 On wood. Thallus granular forming isidia- or coral-like structures; disc pale orange to dark purple-brown; proper exciple orange-brown to brown at rim; on old wood, but generally on rock.................\textit{B. egenuloidea (Bacidina egenuloidea)}

13 Proper exciple with radiating clusters of crystals; white pruina present; atranorin present as a major compound or a trace .........................................................\textit{B. inundata (Bacidina inundata)}

14 Hypothecium brown-orange to dark brown; apothecial section K+ purple-red.........................\textit{B. polychroa}

15 Thallus generally coarsely granular, pale grey to green-grey; prothallus white to pale grey when present; ascospores up to 9-septate .................\textit{B. rubella}

16 Thallus grey; disc not pruinose generally, but sometimes white-pruinose; proper exciple with radiating clusters of minute crystals (ca. 0.5 μm wide); epihymenium without distinct colour; ascospores 50–85 × 2.6–3.4 μm ..................\textit{B. fraxinea}

17 Thallus granular with soredia-like goniocysts.........................................................\textit{B. suffusa}

18 Hypothecium colourless; conidia curved without hook ..............................................\textit{B. delicata (Bacidina delicata)}

19 Disc purple-brown to black or slightly blackish when mature; epihymenium K+ purple ..................\textit{B. sulphurella (Bacidina sulphurella)}

20 Proper exciple colourless to pale yellow at rim; thallus olive-green; apothecia generally pale yellow to pale orange with slightly blackish pigment; epihymenium colourless with a little pale yellow-brown pigment .............\textit{B. fuscopallida}

21 Brown pigment of epihymenium deposited in caps of paraphysial tips; thallus wrinkled or warty, but not squamulose; prothallus blackish on border when present; ascospores 32–67 × 2.5–4.5 μm, 3- to 15-septate.........\textit{B. heterochroa}
white between areoles; ascospores 45–80 × 2–3.5 μm, 7- to 28-septate ...........
.................................................................................................................. B. laurocerasi
22  Thallus rimose, wrinkled or warted; apothecia ca. 0.2 mm diam.; hypothecium colourless; ascospores 24–28 × 1–1.2 μm, 0- to 3-septate; occasionally on rock ..
.................................................................................................................. B. chloroticula (Bacidina chloroticula)
   –  Thallus granular to smooth; apothecia 0.4–1.4 mm diam.; hypothecium straw, yellow-brown to red-brown; ascospores 45–70 × 1.5–4 μm, 3- to 15-septate ..23
23  Proper exciple yellow-brown to brown at rim; epihymenium yellow-brown; ascospores 1.5–2.5 μm wide, 3- to 7-septate ...................... B. arceutina
   –  Proper exciple colourless to pale yellow at rim; epihymenium colourless; ascospores 2–4.5 μm wide, 3- to 15-septate ...................... B. ekmaniana

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