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# Potential of five non-spore-forming bacteria, originated from the European cockchafer, *Melolontha melolontha* (Linnaeus, 1758) (Coleoptera: Scarabaeidae), on three economic insect pests

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## Abstract

Five non-spore-forming bacteria were isolated from the European cockchafer, *Melolontha melolontha* (Linnaeus, 1758) (Coleoptera: Scarabaeidae). Their potential was tested against the three economic insect pests, the great spruce bark beetle, *Dendroctonus micans* Kugelann (Coleoptera: Curculionidae); the pine processionary, *Thaumetopoea pityocampa* (Lepidoptera: Thaumetopoeidae); and the gypsy moth, *Lymantria dispar* (Linn.) (Lepidoptera: Erebidae), to find an effective biological control agent. All isolated bacteria were cultured and identified using VITEK bacterial identification systems and 16S rRNA gene sequence analysis. The bacteria were identified as *Enterobacter cloacae* complex (isolate 1M), *Serratia marcescens* (isolate 3M), *Pseudomonas aeruginosa* (isolate 4M), *Kocuria kristinae* (isolate 5M), and *Serratia liquefaciens* (isolate 8M). Laboratory experiments, carried out to evaluate the virulence of these isolates, showed that all isolated bacteria had a pathogenic effect on the tested pests. *E. cloacae* had 35, 56.7, and 84%; *S. marcescens* 50, 60.9, and 47.8%; *P. aeruginosa* 55, 69.6, and 48%; *K. kristinae* 40, 43.5, and 16%; and *S. liquefaciens* 45, 65.2, and 36% mortality rates on the larvae of *D. micans*, *T. pityocampa*, and *L. dispar*, respectively. The isolated bacteria can be considered in integrated pest control programs.

**Keywords:** *Melolontha melolontha*, *Dendroctonus micans*, *Thaumetopoea pityocampa*, *Lymantria dispar*, Non-spore-forming bacteria, Virulence

## Background

New pest management strategies tend to minimize the impact on the environment and non-target organisms (Ruiu et al. 2013). Entomopathogens have relative specificity and lower environmental impact. The successful use of entomopathogens results in an alternative pest management for insect control. Among them, bacteria and their toxins are the most commercially microbial insecticides used successfully in biological control. Entomopathogenic bacteria such as *Bacillus thuringiensis* are generally known as lower risk pesticides than chemicals. Current goals

include searching for entomopathogenic bacteria and determining their potentials (Egami et al. 2009). Entomopathogenic bacteria are commonly isolated from host insects or soil (Thiery and Frachon 1997). Soil plays an important role as a habitat for spore-forming and non-spore-forming entomopathogenic bacteria. While the entomopathogenic bacteria domain has been well represented by spore-forming bacteria, members of *Bacillaceae* family, non-spore-forming entomopathogenic bacteria have been also documented with insects (Ruiu et al. 2013).

Numerous larvae of Scarabaeidae (Coleoptera) feed on plant materials in the soil, and many of them are known as important plant pests. Scarabs are infected by a number of entomopathogens (Yaman et al. 2016). The European cockchafer, *Melolontha melolontha* (Linnaeus, 1758)

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(Coleoptera: Scarabaeidae) causes significant economic damage. It is a soil-inhabiting insect pest at the larval stage and possible target for entomopathogenic bacteria in soil.

In this study, isolation and characterization of non-spore-forming bacteria from *M. melolontha* larvae were presented and their potentials on three economic insect pests were documented.

## Material and methods

### Insect samples

Samples of *M. melolontha* larvae were collected from Ordu province, Turkey, targeting isolate entomopathogenic bacteria. For bioassay tests, three economic insect pests, namely *Dendroctonus micans* (Yaman et al. 2010), *Thaumetopoea pityocampa*, and *Lymantria dispar*, were used. *D. micans* larvae were obtained from the forest fields near Ordu, Turkey; their cultures were maintained in a laboratory set up by the Turkish Ministry of Forestry in Ordu province. *Lymantria dispar* larvae were collected from İzmir province, Turkey, brought to the laboratory, and fed on apple leaves. *T. pityocampa* larvae were collected from pine trees in Sinop province, Turkey, brought to the laboratory, and fed on fresh pine needles.

### Bacterial isolation

*M. melolontha* larvae suspected of bacterial symptoms after a macroscopic examination were used for bacterial isolation (Thiery and Frachon 1997). Before the isolation process, the dead larvae were individually placed into 70% ethanol and gently shaken for 3 min (Yaman et al. 2010). After surface sterilization, considering aseptic conditions samples were washed by a sterilized water. Sterile 1 ml syringes were used to insert into the hemocoel of the insect. Then, 100 µl hemolymph suspension, taken from the hemocoel, was spread on nutrient agar plates. The plates were incubated at 25–36 °C for 24–48 h. After incubation, the plates were examined and bacterial colonies that were similar in terms of colony and color morphology were selected. Different colony types of bacteria were selected and purified on nutrient agar plates by subculturing. Individual colonies were subcultured twice to ensure purity (Kuzina et al. 2001). One of the subcultured colonies was used for identification of species per isolate.

Bacterial strains were preserved for a long-term storage in nutrient broth with 15% glycerol at –86 °C for further tests at the Department of Molecular Biology and Genetics, Faculty of Arts and Science, Ordu University. The stock culture strains were subcultured onto tryptic soy agar plates to check their purity. All bacterial isolates were initially stained by Gram's dye for the identification of gram-positive or gram-negative bacteria and tested for some biochemical reactions. A standard bacterial suspension was

prepared in 1.8 ml of 0.45% saline, using the VITEK colorimeter for each isolate. The time interval between suspension preparation and card filling was less than 30 min to avoid changes in turbidity. Then, VITEK bacterial identification systems (bioMérieux, Prod. No 21341 and 21342) were used for the identification of the isolated bacteria. Additionally, all isolated bacteria were further identified, using 16S rDNA analysis. For this, pure bacterial suspensions were used for DNA extraction. Equal volumes of bacterial suspension and glass beads were put into an Eppendorf tube and vigorously shaken on the vortex for 1 min at maximum speed. And then, the solutions were incubated by proteinase K at 56 °C for 3 h. After that, nucleic acid extraction was performed by the DNA isolation kit, as according to the manufacturer's guidelines and Hylíš et al. (2005) with some modifications. To amplify and sequence the bacterial 16S rRNA genes and identify the bacterial isolates, the isolated bacterial DNAs were sent to Sentegen Biotech Company (Ankara, Turkey).

### Bioassay tests for potentials of the isolated bacteria

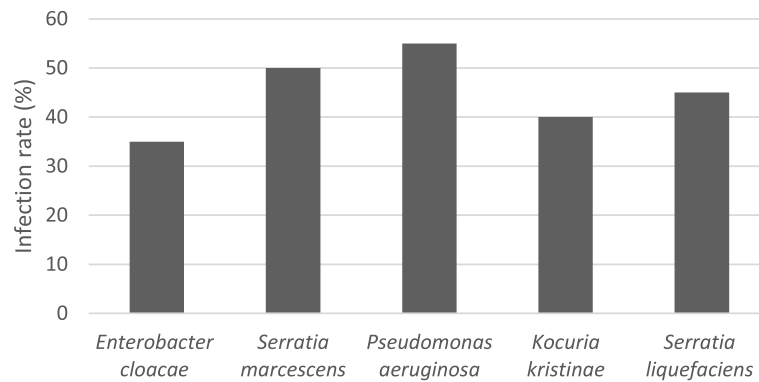
Virulence of the isolated bacteria was tested against the larvae of *D. micans*, larvae of *T. pityocampa*, and *L. dispar*. Bacterial isolates were incubated in nutrient broth medium at 30 °C for 18 h. Bacterial suspensions containing approximately 10<sup>9</sup> bacteria/ml were used in bioassay according to Ben-Dov et al. (1995). *D. micans* larvae were fed on spruce barks dipped into the bacterial suspensions (Yaman et al. 2010), *T. pityocampa* larvae on pine needles, and *L. dispar* on apple leaves. Each bioassay group was performed by 30 insect larvae at the same laboratory conditions. All tested groups were kept at 24–28 °C and 35–45% RH and 18:6 photoperiod in laboratory conditions. Observations were recorded daily and dead larvae were removed immediately. All bioassays were repeated three times on different days and data was corrected, using Abbott's formula (Abbott 1925).

## Results and discussion

### Isolated bacteria

In the present study, five non-spore-forming bacteria were isolated from the larvae of *M. melolontha*. These bacteria were identified as *Enterobacter cloacae* complex (isolate 1M), *Serratia marcescens* (isolate 3M), *Pseudomonas aeruginosa* (isolate 4M), *Kocuria kristinae* (isolate 5M), and *Serratia liquefaciens* (isolate 8M) according to the results from VITEK bacterial identification systems and 16S rRNA gene sequence analysis.

Isolate 1M of *M. melolontha* larvae was identified as *E. cloacae*. It is a rod-shaped, gram-negative, facultatively anaerobic bacterium. Species of the *E. cloacae*



**Fig. 1** Potential of the isolated bacteria on *Dendroctonus micans* larvae

complex are widely encountered in nature, and they can act as pathogens (Mezzatesta et al. 2012). Sezen et al. (2007) also found *Enterobacter* sp. from *M. melolontha*. *E. cloacae* was observed to be the most common species of the poplar pest, *Cryptorhynchus lapathi* (Yaman et al. 2017).

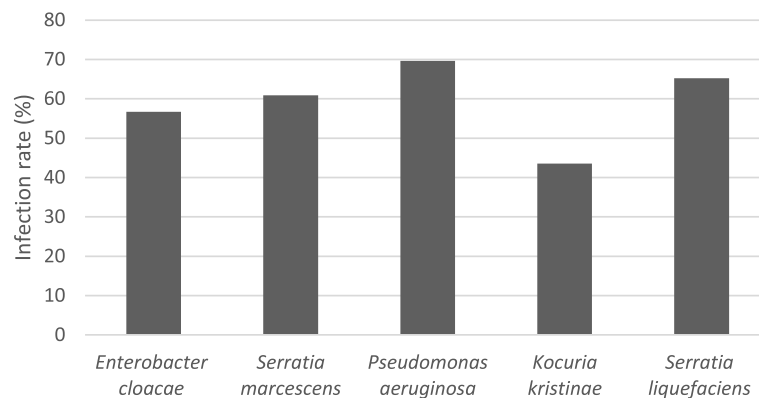
Isolate 3M, isolated from *M. melolontha* larvae, was identified as *Serratia marcescens* (family Enterobacteriaceae). It is a species of rod-shaped, gram-negative and facultative anaerobic bacteria. *S. marcescens* is one of the best-known and mostly isolated pathogenic bacterium from insects (Thiery and Frachon 1997; Lauzon et al. 2003; Pineda-Castellanos et al. 2015). Among the non-spore-forming bacterial genera, the genus *Serratia* includes more effective entomopathogenic species (O'Callaghan et al. 1996) and would possibly be the bio-control agents against some insects (Sezen et al. 2001).

Isolate 4M isolated from *M. melolontha* larvae was identified as *Pseudomonas aeruginosa*. It is a bacil, gram-negative bacterium found throughout the environment. It is known as a potential pathogen for various insects and has frequently been isolated from infected insects, e.g., *Schistocerca gregaria* (Forski) (Orthoptera: Acrididae), *Melanopus* spp.

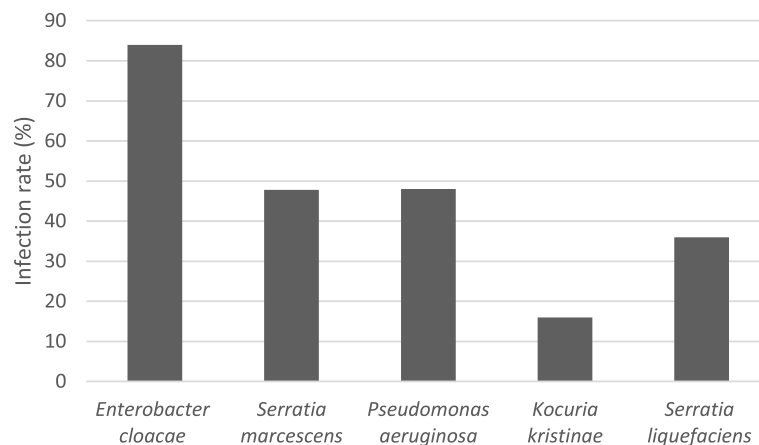
(Orthoptera: Acrididae) (Bucher and Stephens 1959), *Galleria mellonella* (Linnaeus) (Lepidoptera: Pyralidae) (Lysenko 1963), *Stonioxys calcitrans* (Linnaeus) (Diptera: Muscidae), *Amphimallon solstitialis* (Linnaeus) (Coleoptera: Scarabaeidae), and *Eurygaster intergriceps* (Puton) (Hemiptera: Scutelleridae) (Lipa 1975). This bacterium was isolated from dead larvae of *Pieris brassicae* (Yaman and Demirbağ 2000). *P. aeruginosa* had the greatest insecticidal effect (67%) on *P. brassicae* (Bucher and Stephens 1959; Yaman and Demirbağ 2000). Bucher and Stephens (1959) found *P. aeruginosa* as the main cause of disease in grasshoppers, under both laboratory and natural conditions. Obtained results agree with the abovementioned references.

Isolate 5M *Kocuria kristinae* was another bacterium isolated from *M. melolontha* in this study. It is a gram-positive microorganism. Cockburn et al. (2013) isolated and identified this bacterium from the common bed bug, *Cimex lectularius*. Yaman and Ertürk (2016) isolated and identified it from *Crepidodera aurata* (Coleoptera, Chrysomelidae) for the first time, recently.

Isolate 8M *Serratia liquefaciens* was isolated from *M. melolontha* larvae. It is a species of gram-negative bacteria and the most prevalent *Serratia* species in the natural



**Fig. 2** Potential of the isolated bacteria on *Thaumetopoea pityocampae* larvae



**Fig. 3** Potential of the isolated bacteria on *Lymantria dispar* larvae

environment. This bacterium was diagnosed in *Cydia pomonella* (Zimmermann et al. 2013). Muratoğlu et al. (2011) isolated this bacterium from *Ips typographus*. Kati et al. (2017) found both *S. marcescens* and *S. liquefaciens* in *Xyleborinus saxesenii*. Yaman et al. (2017) isolated *S. liquefaciens* from *Cryptorhynchus lapathi*.

#### Bioassay

Bioassay tests showed that all the isolated bacteria had a pathogenic effect against the three economic insect pests tested with different ratios. *E. cloacae* had 35% mortality on *D. micans* larvae, 50% on *S. marcescens*, 55% on *P. aeruginosa*, 40% on *K. kristinae*, and 45% on *S. liquefaciens* (Fig. 1).

Among the tested bacteria, *P. aeruginosa* was the most effective isolate causing 69.6% mortality rate in the larvae of *T. pityocampa*. The other isolates, *E. cloacae*, *S. marcescens*, *K. kristinae*, and *S. liquefaciens* showed different levels of potential such as 56.7, 60.9, 43.5, and 65.2% mortality rate on *T. pityocampa* larvae, respectively (Fig. 2).

On the other hand, the bacterial isolates against the 2nd instar larvae of *L. dispar* was tested. *E. cloacae* was the most effective isolate causing 84% mortality on the larvae of *L. dispar*. The other isolates, *S. marcescens*, *P. aeruginosa*, *K. kristinae*, and *S. liquefaciens* showed different levels of effects such as 47.8, 48, 16, and 36% mortality on *L. dispar* larvae, respectively (Fig. 3).

Sezen et al. (2001) carried out a number of bioassays with *S. marcescens*, using larvae and/or adults of *Agelastica alni*, *Balaninus nucum*, *Curculio elephas*, *Euproctis chrysorrhoea*, *Hyphantria cunea*, *Malocosoma neustria*, *Neodiprion sertifer*, *Pieris brassicae*, and *Y. malinellus*. They determined 70 and 48% effect on the larvae and adults of *A. alni*, respectively; 78% on *B. nucum* adults, 55% on the larvae of *C. elephas*, 58% on *E. chrysorrhoea*, 10% on *H. cunea*, 78% on *M. neustria*, 88% on *N. sertifer*,

100% on *P. brassicae*, and 92% on *Y. malinellus* larvae. On the other hand, Lipa and Wiland (1972) found 100% mortality on *Agrotis* sp., Mathew and Mohamed-Ali (1987) recorded 83.3% on *C. cadambae* larvae, and Onoviran et al. (1985) recorded 65–90% mortality with *S. marcescens* on *Glossiana* spp.

#### Conclusion

Entomopathogenic bacteria are generally known as lower risk pesticides than chemical pesticides. Five entomopathogenic bacteria were isolated from the forest common pests; *D. micans*, *T. pityocampa*, and *L. dispar*. The results confirmed that the five bacteria originated from *M. melolontha* larvae and infected the three tested pests. Therefore, the results of this study would be of great interest to propose some effective entomopathogenic bacteria against these important forest pests.

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#### Authors' contributions

ÖE collected insects and isolated and characterized bacteria and carried out bioassay experiments. MY identified the bacteria, collected insects, and wrote the manuscript. Both authors read and approved the final manuscript.

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#### Availability of data and materials

All datasets are presented in the main manuscript.

#### Ethics approval and consent to participate

Not applicable.

#### Consent for publication

Not applicable.

#### Competing interests

The authors declare that they have no competing interests.

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