Antimicrobial susceptibility testing

- Bacterial isolate identification was confirmed by standard algorithms and supported by matrix-assisted laser desorption ionization-time of flight mass spectrometry (Staphylococcus epidermidis).
- Multilocus sequence typing (MLST) was performed by extracting the previously defined set of 7 genes (L3, L4, L22).

MATERIALS AND METHODS

INTRODUCTION

- Among 873 isolates included in the 2019 lefamulin surveillance program, 2 (0.2%) isolates had lefamulin MIC results of ≥1 mg/L. These isolates did not have any known resistance determinants associated with pleuromutilin resistance and MIC ≥2 mg/L.