

# Surotomycin

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## 1. DESCRIPTION

Surotomycin (CB-183,315; MK-4261) belongs to the lipopeptides and is a semisynthetic derivative cyclic tail analogue of daptomycin (see [Chapter 45](#), Daptomycin) designed specifically to improve the activity against *Clostridium difficile* (Yin *et al.*, 2015). It has the same peptide sequence as daptomycin but has an aromatic ring containing an unsaturated lipid tail, (E)-3-(4-pentylphenyl) but-2-enoic acid tail. The molecular formula for surotomycin is  $C_{77}H_{10}N_{17}O_{26}$  with a corresponding molecular weight of 1680.7 g/mol (PubChem, 2016). The molecular structure of surotomycin is depicted in [Figure 50.1](#). Its mechanism of action is similar to that of daptomycin, cell membrane depolarization leading ultimately to cell death (Mascio *et al.*, 2012). It was developed by Cubist Pharmaceuticals but is now owned by Merck & Co. (North Wales, PA).

Surotomycin has, similar to daptomycin, good activity against most Gram-positive bacteria, including enterococci and methicillin-resistant *Staphylococcus aureus*. Although it is a promising alternative to vancomycin and metronidazole for the treatment of *C. difficile*-associated diarrhea (CDAD), with activity against both growing and nongrowing *C. difficile* (Alam *et al.*, 2015), further development remains unclear because of disappointing results in clinical trials.

## 2. KEY FEATURES

The antimicrobial spectrum of surotomycin is comparable to daptomycin but is significantly more active against *C. difficile*. Surotomycin has no significant activity against Gram-negatives, and, although concentrations in stool are very high (Chandorkar, 2013b), minimum inhibitory concentration (MICs) are even higher (Citron *et al.*, 2012). It should be

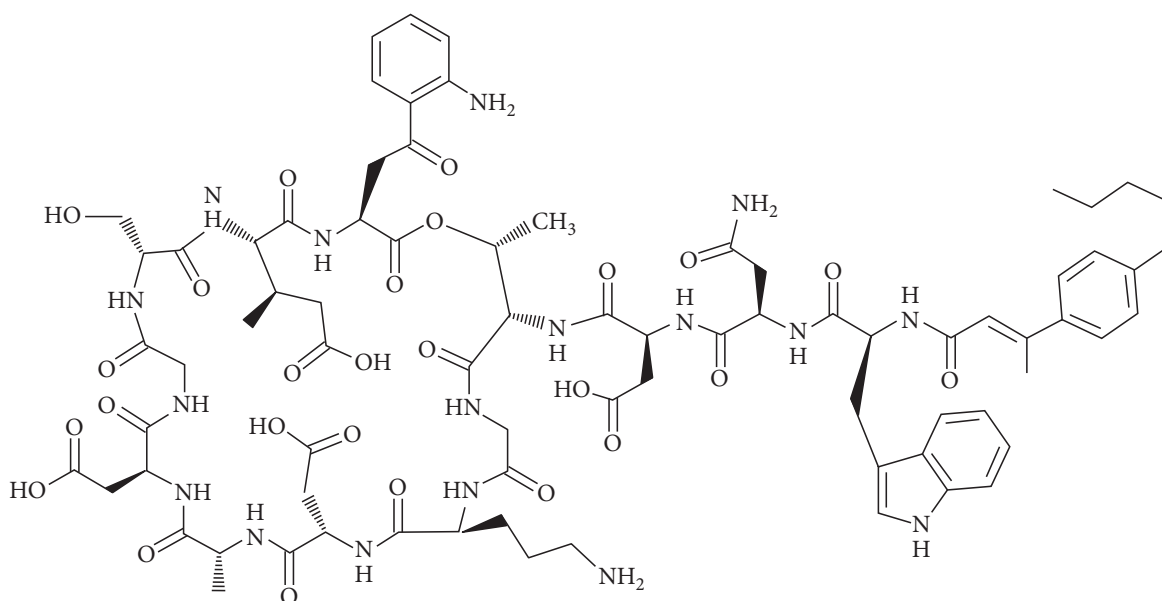


Figure 50.1. Molecular structure of surotomycin.