

Antibiotic-Resistant Pathogens in Ear, Nose, and Throat Infections

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Introduction

The management of ear, nose, and throat (ENT) infections requires an accurate clinical and bacteriological diagnosis, followed by an initial empiric antimicrobial therapy that may be adjusted once the identification of the causative organism(s) is available. The increasing antimicrobial resistance of many respiratory tract bacterial pathogens has made the treatment of these infections more challenging [1, 2].

The microflora of the upper airways, including the oral cavity, nasopharynx, and oropharynx, is complex and contains many types of aerobic, facultative, and obligate anaerobic bacteria [3]. The ratio of anaerobic to aerobic bacteria in saliva is approximately 10:1. The total count of anaerobes in the saliva and elsewhere in the oral cavity reaches 10^7-10^8 bacteria/ml.

Table 2.1 lists the major pathogens that cause various ENT infections. *Streptococcus pneumoniae*, *Haemophilus influenzae*, and *Moraxella catarrhalis* are the predominant aerobic pathogens recovered in acute respiratory tract infections. Their resistance to antimicrobials has

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significantly increased in the past 30 years. Staphylococcus aureus, Pseudomonas aeruginosa, and endogenous oropharyngeal anaerobes are commonly recovered in chronic head and neck infections, some of which can be lifethreatening [4]. Because anaerobes are difficult to isolate, they are often overlooked. Furthermore, their exact role is difficult to ascertain from many past reports because of the inconsistent methodologies used for their isolation and identification in many of these studies [5, 6]. Isolation and identification of anaerobes require appropriate methods of collection, transportation, and cultivation of specimens. Treatment of anaerobic infections is complicated by their polymicrobial nature and the growing antimicrobial resistance and slow growth of these bacteria [5, 6].

Antibiotic Resistance Mechanisms

Antibiotics are naturally produced by many bacteria and fungi, and antibiotic-producing microbes are resistant to the antibiotics they produce. Antibiotic resistance therefore preceded the advent of antibiotics by many millennia. Antibiotic resistance genes have been found within bacteria contained in samples of 30,000-year-old permafrost. Selective pressure by human use of antibiotics over the past 80 years has led to rapid expansion in antibiotic resistance in clinically important pathogens. Multidrug-resistant organisms,

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	Aerobic and facultative		
Type of infection	organisms	Anaerobic organism	
Otitis media: acute	Streptococcus pneumoniae	Peptostreptococcus spp.	
	Haemophilus influenzae ^a		
	Moraxella catarrhalis ^a		
Otitis media: chronic, and Mastoiditis	Staphylococcus aureus ^a	Pigmented Prevotella and Porphyromonas sp	
	Escherichia coli ^a	Bacteroides spp. ^a	
	Klebsiella pneumoniae ^a	Fusobacterium spp. ^a	
	Pseudomonas aeruginosa ^a	Peptostreptococcus spp.	
Peritonsillar and retropharyngeal abscess	Streptococcus pyogenes	Fusobacterium spp. ^a	
	S. aureus ^a	Pigmented Prevotella and Porphyromonas spp. ^a	
	S. pneumoniae		
Recurrent tonsillitis	S. pyogenes	Fusobacterium spp. ^a	
	H. influenzae ^a		
	S. aureus ^a		
Suppurative thyroiditis	S. pyogenes	Pigmented <i>Prevotella</i> and <i>Porphyromonas</i> spp. ^a	
	S. aureus ^a		
Sinusitis: acute	H. influenzae ^a	Peptostreptococcus spp.	
	S. pneumoniae		
	M. catarrhalis ^a		
Sinusitis: chronic	<i>S. aureus</i> ^a	Fusobacterium spp. ^a	
	S. pneumoniae	Pigmented <i>Prevotella</i> and <i>Porphyromonas</i> spp. ^a	
	H. influenzae		
Cervical lymphadenitis	S. aureus ^a	Pigmented <i>Prevotella</i> and <i>Porphyromonas</i> spp. ^a	
	Mycobacterium spp.	Peptostreptococcus spp.	
Postoperative infection disrupting oral mucosa	Staphylococcus spp. ^a	Fusobacterium spp. ^a	
	Streptococcus spp. ^a	Bacteroides spp. ^a	
	Enterobacteriaceae ^a	Pigmented <i>Prevotella</i> and <i>Porphyromonas</i> spp. ^a	
	<i>Pseudomonas</i> ^a	Peptostreptococcus spp.	
Deep neck space	Streptococcus spp. ^a	Bacteroides spp. ^a	
	Staphylococcus spp. ^a	Fusobacterium spp. ^a	
		Peptostreptococcus spp.	
Odontogenic complications	Streptococcus spp. ^a	Pigmented <i>Prevotella</i> and <i>Porphyromonas</i> spp. ^a	
	Staphylococcus spp. ^a	Peptostreptococcus spp.	
Oropharyngeal: Vincent's angina	Streptococcus spp. ^a	Fusobacterium necrophorum ^a	
Necrotizing ulcerative gingivitis	Staphylococcus spp. ^a	Spirochetes, Prevotella Intermedia, Fusobacterium sppª	

Table 2.1 Some of the aerobic and anaerobic bacteria isolated in upper respiratory tract and head and neck infections

^aOrganisms that have the potential of producing beta-lactamase

defined as those organisms with resistance to one or more classes of antibiotics, are now prevalent.

Bacteria can be genetically resistant to an antibiotic or acquire resistance through mutation or acquisition of foreign DNA (e.g., uptake of naked DNA left by dying bacteria, or acquisition of a plasmid carrying resistance genes). Plasmids, small circular strands of DNA that replicate independently of chromosomes, are commonly found in bacteria. Plasmids can be transferred from one bacterium to another in several ways, including during bacterial conjugation and via a bacterial virus (bacteriophage). Resistance genes may be continuously expressed ("constitutive"), or expressed only when needed ("inducible"). Resistance usually costs the bacterium energy so inducible resistance is more common.

Bacteria have several mechanisms of resistance (Table 2.2). These include permeability barriers, inactivating enyzmes, target site alteration, overproduction of the target, and efflux mechanisms. An example of a permeability barrier is that of Gram-negative bacilli to penicillin. Gram-negative bacilli have a lipopolysaccharide outer membrane that envelops the cell wall. This outer membrane is absent in Gram-positive bacteria. The outer membrane is hydrophobic, and hydrophilic antibiotics such as nafcillin do not penetrate. Hydrophilic antibiotics may penetrate the outer membrane through their porins (permeability channels), but loss of favorable porins will lead to resistance. This may occur during imipenem treatment of Pseudomonas, for example. Another common mechanism is alteration of the target site of the antibiotic. Penicillin acts by attaching to penicillin binding protein (PBP), a

 Table 2.2
 Some common mechanisms of bacterial resistance and examples of antibiotics affected

Mechanism	Example
Permeability barrier to antibiotic	Outer membrane of Gram- negative bacteria serves as a barrier to nafcillin
Enzymatic inactivation of antibiotic	Beta-lactamases (e.g., <i>Staphylococcus aureus</i> inactivation of penicillin by a beta-lactamase)
Alteration of target site for the antibiotic	 Alteration of the bacterial enzyme, penicillin binding protein, in MRSA so that penicillin cannot bind Alteration of the ribosomal target site by methylation so erythromycin or clindamycin cannot bind
Overproduction of the target	Overproduction of the target bacterial enzyme (dihydropteroate synthase) involved in folate production
Efflux pumps to pump antibiotic out of cell	Efflux of tetracycline by some Gram-negative bacilli, resulting in low intracellular concentrations

MRSA methicillin-resistant Staphylococcus aureus

bacterial enzyme that is used in cell wall synthesis. *Staphylococcus aureus* can acquire a gene (*mecA*) which encodes for an altered PBP (PBP2a) that does not bind penicillin. Acquisition of the *mecA* gene by *S. aureus* results in MRSA (methicillin-resistant *S. aureus*), a bacterial species resistant to all beta-lactams except fifth generation cephalosporins.

Beta-Lactamase Production

A major resistance mechanism is inactivation of the antibiotic by a bacterial enzyme. Betalactamases are the most important examples of such enzymes, and these include penicillinases, cephalosporinases, carbapenemases. Some are produced by the bacterial chromosome and some by a plasmid within the bacterium. Beta-lactam antibiotics have a four-member beta-lactam ring, and beta-lactamases hydrolyze this ring, rendering the antibiotic ineffective (Fig. 2.1).

Beta-lactamase production is an important mechanism of antimicrobial resistance of both aerobic bacteria (e.g., *Staphylococcus aureus*, *H. influenzae*, and *M. catarrhalis*), and anaerobic Gram-negative bacilli (e.g., pigmented *Prevotella* and *Porphyromonas*). Beta-lactamase-producing bacteria can play an important role in respiratory infections [7]. They can cause the infection as well as have an indirect effect through their ability to produce the beta-lactamase [8]. These bacteria may not only survive penicillin therapy but can also, as was demonstrated in vitro [9], in vivo [10, 11], and in clinical [12] studies, protect other penicillin-susceptible bacteria from penicillin by releasing the free enzyme into their environment [8].

Aerobic Bacteria

Haemophilus influenzae

About 40% of *H. influenzae* resist beta-lactam antimicrobials through production of beta-lactamases. Increased prevalence of non-typeable *H. influenzae* strains that resist ampicillin and/or other beta-lactams was noted in the

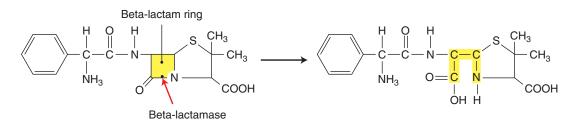


Fig. 2.1 Inactivation of ampicillin by beta-lactamase. The red arrow points to the chemical bond that is hydrolyzed by beta-lactamase

past decade [13]. Ampicillin resistance is usually due to plasmid-mediated production of beta-lactamase so it can be overcome by beta-lactam, beta-lactamase inhibitor combination antibiotics (e.g., amoxicillin-clavulinate). However, H. influenzae resistance to beta-lactams has expanded to include production of an altered penicillin binding protein (PBP3) [14]. This type of resistance cannot be overcome by a beta-lactamase inhibitor, so amoxicillin-clavulinate and similar antibiotics will be ineffective. The frequency of non-beta-lactamase resistance in H. influenzae has increased. In a retrospective study that evaluated 465 H. influenzae isolates from the blood or cerebrospinal fluid from patients in Sweden between 1997 and 2010, a significant increase in beta-lactam-resistant isolates was observed over the course of the study period. Ninety-one isolates (20%) were resistant to one or more betalactam antibiotics (including penicillin, ampicillin, a cephalosporin, or a carbapenem), and nearly half of the resistant bacteria were beta-lactamase-negative [15].

Beta-lactamase–negative, ampicillin-resistant *H. influenzae* strains are being recovered in greater frequency worldwide. The prevalence of such strains has increased in Japan (by 34%) [16], Spain (by 56%) [17], and in other parts of Europe and Canada [18]. Prevalence in the U.S. has remained low (3%) [19]. Possible explanations for this discrepancy include inadequate vaccination against *H. influenzae* type b in some regions, increased use of cephalosporins, and underdosing of ampicillin [16, 17]. These types of ampicillin-resistant, beta-lactamase-negative *H. influenzae* strains are still susceptible to ceftriaxone [20], which may be a good choice for treatment of clinical infections due to these organisms.

Moraxella catarrhalis

Over 90% of *M. catarrhalis* produce a betalactamase and are therefore resistant to ampicillin. Nearly all strains express beta-lactamase from a chromosomal locus. Three types of betalactamases, BRO-1, BRO-2, and BRO-3, that are inducible and intracellular were identified and characterized [21]. *Moraxella catarrhalis* acquired beta-lactamase in the 1970s and the 1980s, and its antimicrobial susceptibility has remained relatively stable. However, recent macrolide and tetracycline-resistant strains were recovered from the Asia Pacific region and China [22].

The oral antibiotics that are active against *M. catarrhalis* as well as *H. influenzae* are amoxicillin-clavulanate, fluoroquinolones, extended-spectrum cephalosporins, newer macrolides, trimethoprim-sulfamethoxazole (TMP-SMX), and tetracyclines. Parenteral antimicrobials effective against these organisms include second and third-generation cephalosporins, aminoglycocides, ticarcillin, and piperacillin. The *M. catarrhalis* strains are resistant to penicillin, ampicillin, and clindamycin [23].

Streptococcus pneumoniae

Resistance of pneumococci to many antimicrobials has increased in the past two decades [23]. Pneumococcal resistance has increased to beta-lactams (penicillins, cephalosporins, and carbapenems), macrolides (erythromycin, azithromycin, clarithromycin), lincosamides (clindamycin), tetracyclines, folate inhibitors (TMP-SMX), and fluoroquinolones (ciprofloxacin, levofloxacin, gemifloxacin, and moxifloxacin). Most strains of penicillin-resistant S. pneumoniae are also resistant to other antimicrobials. Resistance to antimicrobials is determined genetically. The resistance to beta-lactam antimicrobials is through changes in penicillin binding proteins, to chloramphenicol through inactivating enzymes, and to fluoroquinolones through decreased drug permeability [24]. Macrolide resistance is due to efflux pump, and binding blockage. The latter mechanism also blocks clindamycin. There is no resistance to vancomycin or linezolid. Although vancomycin resistance is not known in S. pneumoniae, the phenomenon of vancomycin tolerance has been observed in a few strains [25]. Risk factors for the acquisition of antibiotic-resistant pneumococcal strains include recent antibiotic use; previous time spent in daycare (for children), in an institutional setting, or a shelter for the homeless (for adults); and recent respiratory infections [26-28].

The affinity of beta-lactams for one or more of the penicillin binding proteins is lowered in pneumococcal strains that have reduced susceptibility to penicillins [29]. Decreased susceptibility of pneumococci to beta-lactams can frequently be overcome with higher doses of penicillins, cephalosporins, and carbapenems. Whether invitro resistance to macrolides [30] or the fluoroquinolones [31] can be overcome by increased doses is controversial. Resistance to folate inhibitors or tetracyclines cannot be overcome by increasing the antibiotic dose [32]. Nonsusceptible isolates are divided into intermediate and resistant strains. The penicillin breakpoints for non-meningitis pneumococcal infections are: susceptible minimum inhibitory concentration (MIC) $\leq 2 \text{ mcg/mL}$, intermediate (MIC = 4 mcg/ mL), and resistant (MIC $\geq 8 \text{ mcg/mL}$) [33]. For meningitis, the penicillin breakpoints are much lower and there is no intermediate category: susceptible MIC \leq 0.06 mcg/mL, resistant MIC $\geq 0.12 \text{ mcg/mL}$.

There has been a recent decrease in penicillinresistant pneumococcal strains. This is probably due to both the change in definition of resistance and the widespread use of pneumococcal conjugate vaccine, which has greatly reduced the prevalence of resistant strains in the population. Among isolates obtained in the U.S. from normally sterile sites such as blood culture, pleural fluid, and cerebrospinal fluid (CSF), 95.5% were found to be susceptible, 2.5% intermediate, and 2.2% resistant [34].

Staphylococcus aureus

Staphylococcus aureus can resist beta-lactam antimicrobials through the production of beta-lactamase. It can also resist methicillin which is defined as an oxacillin MIC \geq 4 mcg/mL Isolates resistant to oxacillin or methicillin also resist all beta-lactam agents, including cephalosporins (with the exception of the fifth-generation cephalosporins, ceftobiprole and ceftaroline).

The prevalence of infection and colonization with MRSA is increasing [35] in all infections including head and neck. A 16.3% increase in the rate of pediatric S. aureus head and neck infections occurred between 2001 and 2006 in a study of 21,009 patients [36]. The highest rate of MRSA infections was in otological (34%), followed by sinonasal (28.3%), and oropharynx/ neck (14.2%) infections. The association between previous antimicrobial use and increased isolation of MRSA was noticed in various infections [37, 38], including sinusitis [39, 40]. Brook et al. [39] and Gerencer [40] found that most patients with chronic sinusitis due to MRSA, who were previously treated with antimicrobials, had been treated with either a fluoroquinolone or macrolides.

Methicillin resistance is mediated by the mecA gene that encodes for low-affinity penicillin binding protein, PBP2a. This gene is located on a mobile genetic element called staphylococcal cassette chromosome (SCCmec). Most MRSA strains isolated during the 1960s originated most likely from a single clone; by 2002, five major MRSA clones emerged throughout the globe [41].

Oral Antibiotics Active Against MRSA. Oral antibiotics that can be used for the treatment of MRSA infections include clindamycin, TMX-SMT, tetracyclines (such as doxycycline or minocycline), and linezolid. Because resistance to these agents is rising, their use should be supported by susceptibility testing whenever possible and by clinical response. Clindamycin inhibits bacterial production of toxins, including Panton-Valentine leukocidin and other virulence factors, and has excellent tissue, bone, and abscess penetration [42]. The agent should not be administered empirically when local MRSA resistance rates to clindamycin are >15% [43]. Clindamycin-susceptible isolates that are resistant to erythromycin may become resistant to clindamycin in its presence [44]. Inducible clindamycin resistance can be detected with D testing in the microbiology laboratory [45]. Trimethoprim-sulfamethoxazole and tetracyclines are not advisable for empiric management of infections that may be due to group A streptococci. Resistance of MRSA to fluoroquinolones may emerge during therapy [46]. Oxazolidinones (linezolid or tedizolid) are effective for the treatment of MRSA-related head and neck infections [47]. Their use is limited by cost and toxicity.

Parenteral Agents Active Against MRSA. Parental agents for treating MRSA infections include vancomycin, daptomycin, linezolid, ceftaroline, telavancin, dalbavancin, oritavancin, tedizolid, tigecycline, teicoplanin, and quinupristin-dalfopristin. Some of these are limited by toxicity concerns, as discussed in Chap. 1. The greatest cumulative clinical experience for the treatment of MRSA infections is with the glycopeptide vancomycin. It is still an important agent for treating these infections despite the overall decrease in the in-vitro susceptibility. Its tissue penetration is variable and increases with inflammation. Daptomycin, a cyclic lipopeptide, is inhibited by pulmonary surfactant and should not be used for the treatment of MRSA pneumonia [48]. Previous exposure to vancomycin can increase resistance to daptomycin [49]. Linezolid, a synthetic oxazolidinone, has excellent tissue distribution, and inhibits toxin production [50]. Linezolid resistance has emerged among MRSA isolates, mostly in healthcare associated strains. The mechanism of resistance is via the bacterial cfr gene located in a potentially mobile genetic element [51]. Linezolid use is limited because of safety concerns, including thrombocytopenia, anemia, lactic acidosis, peripheral neuropathy, serotonin toxicity, and ocular toxicity (rare cases of optic neuropathy with treatment beyond 2 weeks).

Ceftaroline, a fifth-generation cephalosporin, is active against Gram-positive organisms (including MRSA, vancomycin-intermediate S. aureus) as well as Gram-negative pathogens (including Enterobacteriaceae but not Pseudomonas species or extended-spectrum beta-lactamase producers) [52]. Telavancin, a semisynthetic lipoglycopeptide, has a half-life of 7-9 h, allowing once-daily dosing. Oritavancin, a semisynthetic glycopeptide, has a half-life of 100 h. Dalbavancin, a semisynthetic lipoglycopeptide, has a half-life of 6-12 days, permitting once-weekly dosing. Teicoplanin, a glycopeptide, can be administered once daily. Quinupristin-dalfopristin, a streptogramin, use is limited by adverse effects (e.g., hyperbilirubinemia, myalgias, arthralgias, and nausea). Tigecycline, a glycylcycline, is active invitro against many Gram-positive cocci (including MRSA, vancomycin-resistant enterococci, and penicillin-resistant S. pneumoniae), aerobic and facultative Gram-negative bacilli (except Pseudomonas and Proteus spp.), anaerobes, and atypical bacteria. However, the U.S. Food and Drug Administration (FDA) issued "boxed warnings" in 2011 and 2013 because of increased risk of death in patients treated with tigecycline compared with other antibiotics.

Pseudomonas aeruginosa

Pseudomonas aeruginosa is commonly found in chronic otitis media and external otitis [53, 54]. *Pseudomonas* possesses intrinsic resistance to several antimicrobials and can attain resistance during therapy. Some strains are highly drugresistant, resisting three or more classes of antibiotics [55]. Only a small number of antimicrobials possess reliable efficacy against *P. aeruginosa*. These include some penicillins (ticarcillinclavulanate, piperacillin-tazobactam), cephalosporins (ceftazidime, cefepime, cefoperazone), monobactams (aztreonam), fluoroquinolones (ciprofloxacin, levofloxacin), carbapemens (imipenem, meropenem, doripenem), aminoglycosides (gentamicin, tobramycin, amikacin), and polymixins (colistin, polymyxin B). All of these antimicrobials are administered parentally except for the fluoroquinolones that can be given also orally [56]. Monobactams require higher dosing. Aminoglycosides are generally not used as single agents because of inadequate clinical efficacy. Polymixins are administered only in the setting of resistance to other antimicrobials because of their toxicity. A combination of anti-*Pseudomonas* antimicrobials can be administered for serious infections due to *P. aeruginosa* [57].

Anaerobic Bacteria

Anaerobic bacteria predominate in the oropharyngeal mucous membranes, and are therefore a common cause of bacterial infections of endogenous origin of upper respiratory tract and head and neck [5, 6]. These infections include chronic otitis media, mastoiditis and sinusitis, pharyngotonsillitis, peritonsillar, retropharyngeal and parapharyngeal abscesses, suppurative thyroiditis, cervical lymphadenitis, parotitis, siliadenitis, and deep neck infections including Lemierre's Syndrome. The recovery from these infections depends on prompt and proper medical and when indicated also surgical management. Because anaerobes generally are isolated mixed with aerobic bacteria, the antimicrobial(s) used should cover these organisms.

The most effective antimicrobials against anaerobic organisms are: metronidazole, the carbapenems (imipenem, meropenem, dorapenem, ertapenem), chloramphenicol, the combinations of a penicillin and a beta-lactamase inhibitor (e.g., amoxicillin plus clavulinate, ampicillin plus sulbactam, ticarcillin plus clavulanate, piperacillin plus tazobactam), tigecycline, cefoxitin and clindamycin. Table 2.3 lists the susceptibility of various anaerobes to antimicrobial agents.

Beta-Lactams and Anaerobes

Penicillins. Penicillin is used when the infecting strains are susceptible. Most *Clostridium* strains and *Peptostreptococcus* spp. are susceptible to penicillin. *Bacillus fragilis* group anaerobes are resistant to penicillin. Other strains that may show penicillin resistance are growing numbers of anaerobic Gram-negative bacilli commonly found in head and neck infections (e.g., pigmented *Prevotella* and *Porphyromonas* spp.,

Table 2.3Susceptibility of common anaerobes to various antibiotics (includes intermediate resistant strains) [58, 74,80, 108]

Anaerobe	Ampicillin- sulbactam (%)	Amoxicililin- clavulinate (%)	Piperacillin- tazobactam (%)	Clindamycin (%)	Moxifloxacin (%)	Imipenem (%)
Anaerobic Gram-positive cocci ^a	100	94–100	97–100	73–95	64–97	100
Clostridium species	100	95-100	100	75-84	47–93	85
Fusobacterium species		89–100	100	69–82	75–90	96
Prevotella species	100	81-100	≥99	67–87	58-89	94–100
Bacteroides fragilis ^b	89–97	63–96	95-100	58-90	59–90	93–99.7
Bacteroides thetaiotaomicron ^b	85–95	63–88	88–100	40–60	25-87	93–100
Bacteroides fragilis group ^b		80–90	92–100	48–68	43-86	≥99

Susceptibility breakpoints (MIC µg/ml), S = susceptible, R = resistant: ampicillin-sublactam ($S \le 8/4$, $R \ge 32/16$); amoxicillin-clavulinate ($S \le 4/2$, $R \ge 16/8$); piperacillin-tazobactam ($S \le 32/4$, $R \ge 128/4$); clindamycin ($S \le 2$, $R \ge 8$); moxifloxacin ($S \le 2$, $R \ge 8$), imipenem ($S \le 4$, $R \ge 16$)

Metronidazole is not listed but >99% of anaerobic Gram-negative bacilli are susceptible

^aIncludes *Peptostreptococcus* species and others

^bThese comprise the majority of *Bacteroides* isolates found in infections above the neck [108]

Prevotella oralis, Prevotella bivia), Bacteroides disiens, strains of clostridia, Fusobacterium spp. (Fusobacterium varium and Fusobacterium mor*tiferum*), and microaerophilic streptococci. Some of these strains show MIC of 8-32 units/mL of penicillin G. In these instances, administration of very high dosages of penicillin G (for non-betalactamase producers) may be effective [58]. Ampicillin and amoxicillin have activity equal to penicillin G, but nafcillin or oxacillin are either not active or have unpredictable activity [59]. Penicillin and ampicillin/amoxicillin are of limited utility because of the production of betalactamases by many oral anaerobes [59–61], but beta-lactam/beta-lactamase inhibitor combinations are effective. Carboxy-penicillins (carbenicillin. ticarcillin) and ureidopencillins (piperacillin, azlocillin, mezlocillin) generally are administered in large quantities to achieve high serum concentrations [62].

Cephalosporins. Cephalosporins have limited utility because many anaerobes produce cephalosporinases [63]. The activity of cephalosporins against the beta-lactamase-producing anaerobic Gram-negative bacilli varies. The antimicrobial spectrum of the first-generation cephalosporins against anaerobes is similar to penicillin G, although on a weight basis, they are less active. Most strains of the B. fragilis group and many Prevotella, Porphyromonas, and Fusobacterium spp. are resistant to these agents [64]. Cephalosporinases have little or no hydrolytic activity for the second-generation cefoxitin (a cephamycin), making it the most effective cephalosporin against the *B. fragilis* group. However, susceptibility to cefoxitin may vary by geographic location and is generally directly related to its clinical use. Cefoxitin is relatively inactive against most species of Clostridium, including Clostridium difficile, with the exception of *Clostridium perfringens* [64–66]. With the exception of moxalactam (not available in the U.S.), the third-generation cephalosporins are not as active against B. fragilis group.

Carbapenems. The carbapenems (imipenem, meropenem, ertapenem, doripenem) have excellent activity against anaerobes [67]. Imipenem is

effective against a wide variety of aerobic and anaerobic Gram-positive and Gram-negative organisms including B. fragilis group [68, 69] It is also effective against most Enterobacteriaceae and about 5-15% of Pseudomonas spp. are resistant [70]. To overcome the problem of renal metabolism of imipenem, it is combined at a 1:1 ratio with an inhibitor of the renal dipeptidase, cilastatin. Imipenem is an effective single agent for the therapy of mixed aerobic-anaerobic infections. Meropenem possesses antibacterial activity similar to imipenem. However, it is less active against staphylococci and enterococci, and provides better coverage of aerobic and facultative Gram-negative bacteria [71, 72]. Ertapenem also has a broad antibacterial spectrum [73] but it is not active against *Pseudomonas*, Enterococcus spp., and Acinetobacter spp. Doripenem has a similar antimicrobial spectrum to meropenem and imipenem [69]. Resistant P. aeruginosa mutants appear to be harder to select in vitro with doripenem than with other carbapenems. Doripenem is not FDA-approved to treat pneumonia. Recent reports have noted the emergence of some carbapenem resistance among anaerobes [74] ranging from 1.1% to 2.5% in a multicenter U.S. survey. Higher resistance was noted in a small number of isolates from Taiwan [75].

Resistance of Anaerobes to Beta-Lactam Antibiotics. Anaerobes exhibit three major resistance mechanisms to beta-lactam antibiotics: inactivating enzymes, mainly beta-lactamases, which include penicilliniases and cephalosporinases; low affinity penicillin binding proteins (PBPs); and decreased permeability through alterations in the porin channel [76]. The production of beta-lactamases is the commonest mechanism, especially among the B. fragilis group and Prevotella spp. [77]. The cephalosporinases are most often of the 2e class type and can be inhibited by three beta-lactamase inhibitors, clavulanic acid, sulbactam, and tazobactam. Each individual cephalosporin may have either a class or specific inhibitor enzyme capable of inactivating it. Carbapenemases are active against the carbapenems as well as all beta-lactam antibiotics.

Carbapenem resistance was found in <1% of U.S. isolates, and up to 3% of *Bacteroides* strains harbor one of the genes that is expressed at a very low level.

With exceptions some among some Clostridium spp., strains of *Clostridium*, Porphyromonas, and Fusobacterium can express resistance through one or more beta-lactamases. Beta-lactamase-producing Fusobacterium and Clostridium spp. express enzymes that are usually inhibited by clavulanic acid [78]. Resistance to beta-lactam antibiotics through changes in the outer membrane porin channels, decreased PBP affinity, and efflux pumps [79] have not been well studied. Bacteroides fragilis group species are generally resistant to penicillins (average 90%), and less often to piperacillin (25%) cefoxitin (25%), cefotetan (30-85%), and third-generation cephalosporins (14–57%) [80, 81].

Beta-lactam/beta-lactamase inhibitor antibiotics and carbapenems have maintained their excellent antibacterial activity against anaerobes, including against members of the B. fragilis group [80]. However, species-to-species variation in susceptibility occurs [40]. Bacteroides fragilis group resistance rates for piperacillintazobactam are generally <1% [82], although one member of the group (Parabacteroides distasonis) has relatively high (20%) resistance. The carbapenems are very effective against all the members of the *B. fragilis* group, and resistance is <0.1% [79, 82, 83]. Some members of the *B*. fragilis group have lower MICs for imipenem and meropenem than for ertapenem [80]. Half of Prevotella spp. may produce beta-lactamases, causing penicillin resistance, and a multicenter survey [68] also detected penicillin resistance in Fusobacterium spp. (9%), Porphyromonas spp. (21%), and Peptostreptococcus spp. (6%). No resistance was found to cefoxitin, cefotetan, betalactam/beta-lactamase combinations, and carbapenems in that survey, with the exception of Peptostreptococcus spp. (4%) and Porphyromonas spp. (5%). Beta-lactamases were identified in several Prevotella and Porphyromonas spp. recovered from pediatric intra-abdominal infections [62].

Chloramphenicol and Anaerobes

Chloramphenicol, a bacteriostatic agent, is active against most anaerobic bacteria but is rarely used in the U.S. [6] due to potentially significant toxicity. The risk of fatal aplastic anemia with chloramphenicol is approximately one per 25,000-40,000 patients treated. This complication is unrelated to the reversible, dosagedependent leukopenia. Other side effects include the production of the potentially fatal "gray baby syndrome" when given to neonates, hemolytic anemia in patients with glucose-6-phosphate dehydrogenase (G6PD) deficiency, and optic neuritis in those who take the agent for a prolonged time. Chloramphenicol has a unique property of lipid solubility that permits penetration across lipid barriers. Levels in the cerebrospinal fluid (CSF), with or without meningitis, usually are one-third to three-fourths the serum concentrations. Levels in brain tissue can be substantially higher than serum levels [83].

Macrolides (Erythromycin, Azithromycin, Clarithromycin) and Anaerobes

The macrolides have moderate to good in vitro activity against anaerobic bacteria other than B. fragilis group and fusobacteria [58, 64]. They are active against microaerophilic streptococci, Gram-positive non-spore-forming anaerobic bacilli, and certain clostridia. They are less effective against *Peptostreptococcus* spp. [84]. Macrolides have relatively good activity against C. perfringens and poor or inconsistent activity anaerobic Gram-negative against bacilli. Clarithromycin is the most active of the macrolides against Gram-positive oral cavity anaerobes. including Actinomyces spp., Propionibacterium spp., Lactobacillus spp., and Bifidobacterium dentium. Azithromycin is slightly less active than erythromycin against these species [84]. Azithromycin is the most active macrolide against Aggragatibacter actinomycetemcomitans, including those isolates resistant to erythromycin. Clarithromycin possess similar activity to erythromycin against most anaerobic Gram-negative bacilli [85]. Emergence of erythromycin-resistant organisms during therapy has been documented [86, 87].

Clindamycin and Anaerobes

Clindamycin has a broad activity against anaerobes, is well absorbed from the gastrointestinal tract [88–90], and rapidly penetrates into most body tissues and fluids [52] although not the central nervous system (CNS). Clindamycin should not be administered in CNS infections. The side effect of most concern is C. difficile associated colitis [91]. Because B. fragilis resistance to clindamycin is increasing worldwide (over 33%) it is no longer recommended as empiric therapy for intra-abdominal infections [65, 74, 80, 92]. Resistance to clindamycin has also increased for other anaerobes. Up to 10% resistance was noted for Prevotella, Fusobacterium, Porphyromonas, and Peptostreptococcus spp., with higher rates for some *Clostridium* spp. (especially *C. difficile*) [68]. Clindamycin has lost some of its activity against anaerobic Gram-positive cocci (i.e., Finegoldia magna-30% resistant), and Prevotella spp. (P. bivia, 70% resistant, P. oralis and Prevotella melaninogenica both 40% resistant), although its activity against Fusobacterium and Porphyromonas spp. remains good. Among the other resistant anaerobes are various species of clostridia especially C. difficile. About 20% of Clostridium ramosum are resistant to clindamycin, as are a smaller number of C. perfringens.

Metronidazole and Anaerobes

Metronidazole and tinidazole are nitroimidazoles with similar in vitro activity against anaerobic bacteria. Metronidazole has excellent in vitro efficacy against most obligate anaerobic bacteria, such as *B. fragilis* group, other species of *Bacteroides*, fusobacteria (including *F. necrophorum*, the etiology of Lemierre's Syndrome), other anaerobic Gram-negative bacilli, and clostridia [93]. These agents have excellent penetration into

the CNS. Resistance to metronidazole among *B*. fragilis group is uncommon [65, 94]. Resistance of anaerobic Gram-positive cocci is rare and resistance of nonsporulating bacilli is common. Most microaerophilic streptococci, P. acnes, and Actinomyces spp. are resistant [94]. Aerobic and facultative anaerobes are usually highly resistant. Because of its lack of activity against aerobic bacteria, an antimicrobial effective against these organisms (e.g., a cephalosporin, a fluoroquinolone) needs to be added when treating a polymi-Adverse crobial infection. reactions to metronidazole include gastrointestinal side effects, central nervous system toxicity, and peripheral neuropathy. Possible mutagenic activity found in mice given large doses of metronidazole [95] was not confirmed by experiments in rats and hamsters [96], and no evidence of mutagenicity was ever found in humans [97].

Tetracyclines and Anaerobes

The tetracycline analogues, doxycycline and minocycline, are more active than the parent compound [58]. However, because of the significant resistance to these drugs, they are useful only when susceptibility tests show efficacy or in less severe infections in which a therapeutic trial is feasible. The use of tetracyclines is not recommended before 8 years of age because of the adverse effect on teeth; tetracyclines are also contraindicated in pregnancy. Tigecycline is a direct analog of minocycline with broad-spectrum activity including anaerobes and some drugresistant pathogens [98, 99]. Resistance of members of the *B. fragilis* group varies from 3.3% to 7.2% [100]. As noted above, tigecycline carries an FDA boxed warning about increased mortality rates compared with other treatments for various infections.

Fluoroquinolones and Anaerobes

Of the systemic quinolones available in the U.S. (ciprofloxacin, levofloxacin, ofloxacin, moxifloxacin, gemifloxacin), moxifloxacin is the most effective against anaerobes [101]. Quinolones with the greatest in vitro activity against anaerobes include clinafloxacin and sitafloxacin [102], but these are not available in the U.S. Quinolones' use is restricted in growing children because of possible adverse effects on the cartilage. In addition, in July 2016, the FDA issued a boxed warning on the use of quinolones for less serious infections, such as acute bacterial sinusitis, due to concern for serious and potentially irreversible side effects on "tendons, muscles, joints, nerves, and the central nervous system" [103]. Increasing resistance to quinolones in B. fragilis group as well as anaerobic Gram-positive cocci has been reported. Bacteroides spp. resistance to fluoroquinolone has been attributed to either an alteration in efflux of the antibiotic or a mutation in gyrase A gene (gyrA) [104]; high-level resistance can be caused by both the mechanisms.

Other Agents Active Against Anaerobes

Bacitracin is active in vitro against pigmented Prevotella and Porphyromonas spp. but is inactive against B. fragilis and Fusobacterium nucleatum [58]. Vancomycin and daptomycin are effective against all Gram-positive anaerobes, but are not active against anaerobic Gram-negative bacilli [105]. Quinupristin/dalfopristin exhibits antibacterial activity against C. perfringens, Lactobacillus spp., and Peptostreptococcus spp. [106]. Linezolid is effective against Fusobacterium spp. (including Fusobacterium nucleatum) and Porphyromonas, Prevotella, and Peptostreptococcus spp. [84, 85]. However, there is little clinical experience in the treatment of anaerobic infections using these agents.

Treating Infections in Otolaryngology

Infections in otolaryngology are often polymicrobial, so antimicrobials effective against both the aerobic and anaerobic components of the infection should be administrated. When such therapy is not given, the infection may persist, and serious complications may occur [5, 6, 107]. A number of factors should be considered when choosing appropriate antimicrobial agents: They should be effective against all target organism(s), induce little or no resistance, achieve sufficient levels in the infected site, cause minimal toxicity, and possess maximum stability and longevity.

When selecting antimicrobials for the therapy of mixed infections, their aerobic and anaerobic antibacterial spectrum and their availability in oral or parenteral form should be considered (Table 2.1). Selection of antimicrobial agents is simplified when a reliable culture result is available. However, this may be particularly difficult in anaerobic infections because of the difficulties in obtaining appropriate specimens. For this reason, many patients are treated empirically based on suspected, rather than established pathogens. Fortunately, the types of anaerobes involved in many infections and their antimicrobial susceptibility patterns tend to be predictable [6, 7]. However, some anaerobes have become resistant to antimicrobials, and many can develop resistance while a patient is receiving treatment [91]. Resistance among some anaerobes has increased significantly over the past three decades. The potential for growing resistance of anaerobes to antimicrobials is especially noted with penicillins, cephalosporins, clindamycin, and fluoroquinolones.

Aside from susceptibility patterns, other factors influencing the choice of antimicrobial therapy include the pharmacologic characteristics of the various drugs, their toxicity, their effect on the normal flora, and bactericidal activity [2, 3]. Although identification of the infecting organisms and their antimicrobial susceptibility may be needed for the selection of optimal therapy, the clinical setting and Gram stain preparation of the specimen may suggest the types of bacteria present in the infection as well as the nature of the infectious process.

Conclusion

Many microbes naturally produce antibiotics and are resistant to the antibiotics they produce. Antibiotic resistant microbes have been present in the environment for millennia. However, the discovery of antibiotics in the twentieth century has led to increasing antibiotic resistance in clinically important microbes. Antibiotics must be chosen carefully and used wisely to prevent further selection and widespread dissemination of multidrug-resistant pathogens.

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