

Glycopeptide Antibiotics

from Conventional Molecules to New Derivatives

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Abstract

Vancomycin and teicoplanin are still the only glycopeptide antibiotics available for use in humans. Emergence of resistance in enterococci and staphylococci has led to restriction of their use to severe infections caused by Gram-positive bacteria for which no other alternative is acceptable (because of resistance or allergy). In parallel, considerable efforts have been made to produce semisynthetic glycopeptides with improved pharmacokinetic and pharmacodynamic properties, and with activity towards resistant strains. Several molecules have now been obtained, helping to better delineate structure-activity relationships. Two are being currently evaluated for skin and soft tissue infections and are in phases II/III. The first, oritavancin (LY333328), is the 4'-chlorobiphenylmethyl derivative of chloroeremomycin, an analogue to vancomycin. It is characterised by: i) a spectrum covering vancomycin-resistant enterococci (VRE), methicillin-resistant *Staphylococcus aureus* (MRSA) and to some extent glycopeptide-intermediate *S. aureus* (GISA); ii) rapid bactericidal activity including against the intracellular forms of enterococci and staphylococci; and iii) a prolonged half-life, allowing for daily administration. The second molecule is dalbavancin (BI397), a derivative of the teicoplanin analogue A40926. Dalbavancin has a spectrum of activity similar to that of oritavancin against vancomycin-sensitive strains, but is not active against VRE. It can be administered once a week, based on its prolonged retention in the organism. Despite these remarkable properties, the use of these potent agents should be restricted to severe infections, as should the older glycopeptides, with an extension towards resistant or poorly sensitive bacteria, to limit the risk of potential selection of resistance.

Glycopeptide antibiotics were introduced in clinical practice 50 years ago, with vancomycin as the only agent for almost 30 years. Teicoplanin was launched in Europe in the mid 1980s and these two molecules remain the only members in this class available for human use; however, related derivatives have been used widely as growth promotants

for livestock. The emergence and spreading of resistance in enterococci and staphylococci towards vancomycin has stimulated active research for new glycopeptides over the last 10 years, leading to the production of a series of semisynthetic derivatives. Starting from vancomycin and moving through a short description of the various compounds obtained

so far, this paper reviews the basis of the selection of oritavancin and dalbavancin as new glycopeptides for clinical development (the chemical structure of these products is presented in figure 1). We also discuss their advantages and potential role in the therapeutic armament.

1. Vancomycin and Teicoplanin

1.1 Mechanism of Action

Biochemical studies indicate that glycopeptides inhibit the late stages of peptidoglycan synthesis.^[1] The biosynthetic pathway of this polymer involves three steps: (i) the synthesis of cytosolic precursors made of pentapeptides fixed on a disaccharide; (ii) the coupling of these precursors with a lipid carrier and the transfer of the resulting amphiphilic molecule to the outer surface of the membrane; and (iii) the reticulation between individual precursors by transpeptidation and transglycosylation reactions, accompanied by the release of the lipid carrier and its recycling to the inner face of the membrane. Bacteria incubated with vancomycin accumulate cytosolic precursors,^[1] suggesting that glycopeptides interfere with the assembly of peptidoglycan and, in particular, with transglycosylation reactions (figure 2). At the molecular level, the primary target of vancomycin was shown to be the D-Ala-D-Ala terminus of the precursors. Molecular modelling and experimental studies^[1-4] indicate that vancomycin forms a stoichiometric complex with the D-Ala-D-Ala dipeptide via the formation of five hydrogen bonds with the peptidic backbone of the glycopeptide. The formation of this complex prevents the transpeptidation reactions by steric hindrance. The tightness of the interaction between the glycopeptide and the D-Ala-D-Ala motif can be enhanced by two mechanisms.

- Formation of homodimers between glycopeptide molecules, which has been demonstrated for glycopeptides of the vancomycin group, such as vancomycin itself,^[6] eremomycin^[7] or chloroeremomycin.^[8] Organisation of molecules in dimers confers a structural rigidity that locks the binding pocket into the correct conformation and

may allow for a cooperative binding to the ligand.^[6,9-12] These dimers are formed by hydrogen bonds between the glycopeptide aglycone, which are maintained in the rigid conformational state favourable for ligand binding by the sugar residues on the glycopeptide molecule.^[13]

- Anchoring of the antibiotic in the membrane, as proposed for glycopeptides carrying a lipophilic tail, such as teicoplanin derivatives.^[11] Lipid moieties are actually a common determinant in the structure of several antibiotics, such as ramoplanin, moenomycin and tunicamycin, which inhibit steps of peptidoglycan synthesis occurring close to the membrane. This suggests that such moieties help to maintain the drug close to their target. The importance of hydrophobic determinants has been further emphasised by the observation that vancomycin derivatives may inhibit transglycosylation without binding to D-Ala-D-Ala.^[14,15]

1.2 Mechanisms and Importance of Resistance

The American National Committee for Clinical Laboratory Standards (NCCLS) has set up breakpoints for susceptibility and resistance, respectively, of ≤ 4 and ≥ 16 $\mu\text{g/mL}$ for vancomycin and ≤ 8 and ≥ 32 $\mu\text{g/mL}$ for teicoplanin (see table I for the main characteristics of the resistant strains). Over the years, two organisms of medical interest, namely the enterococci and *Staphylococcus aureus*, have developed a different, ingenious resistance mechanism.

Resistance of enterococci to vancomycin was first reported in 1988.^[27,28] The situation has evolved quite rapidly in the US, where resistance in blood isolates reached 13% in 1995 and 26% in 2000.^[29] Colonisation by resistant strains is common in critically ill and immunosuppressed patients as well as in patients hospitalised in wards with high antimicrobial use. Hospitalisation is probably the main route for spread of these strains but, currently, outpatients are also often colonised and may, therefore, also represent another important source of contamination.^[30] This alarming observation has stimulated drastic measures in terms of hygiene, restriction in

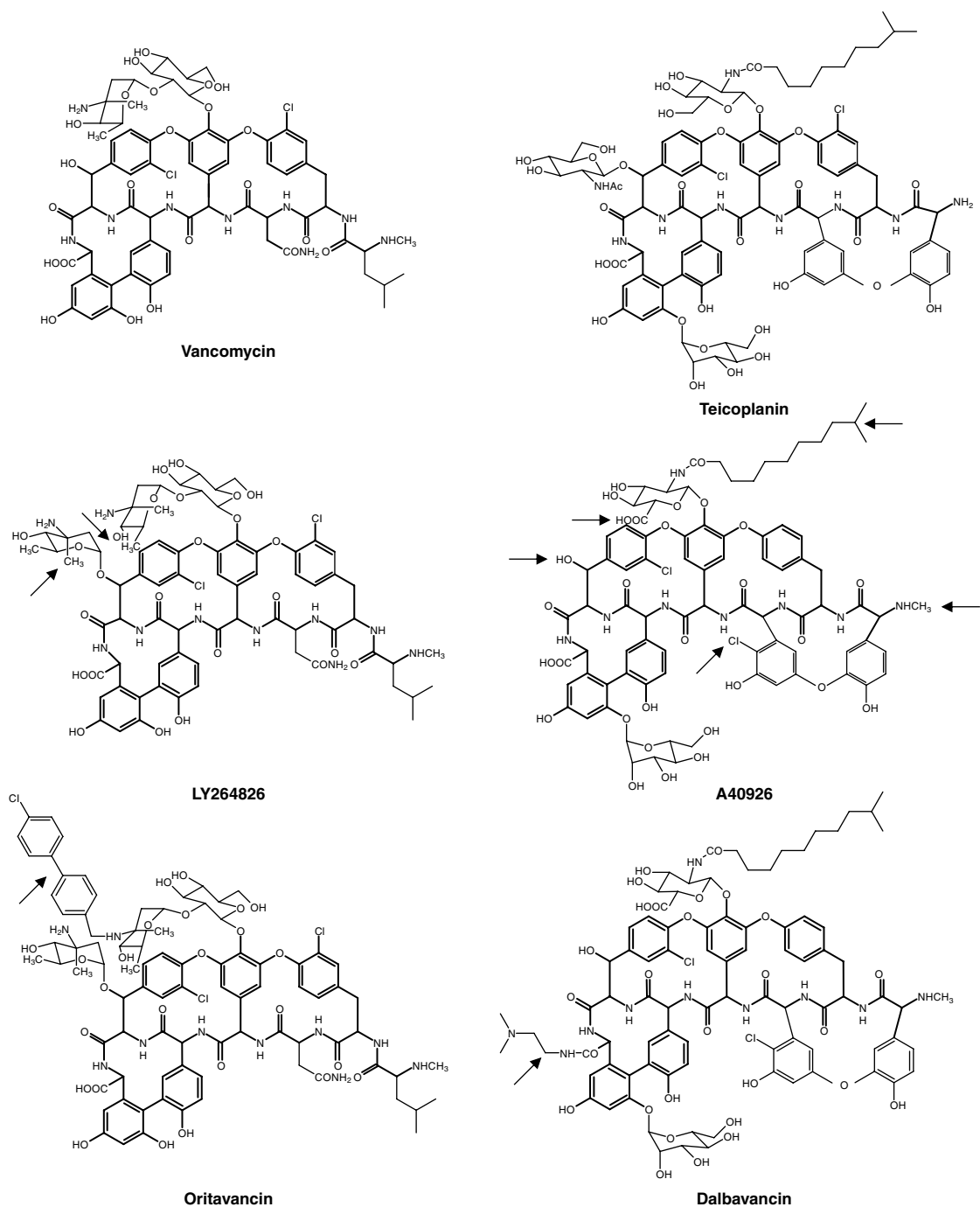


Fig. 1. Chemical structure of glycopeptides. Upper panel: vancomycin and teicoplanin, which are the two glycopeptides currently available in clinical settings; lower panel: oritavancin and dalbavancin, two semisynthetic glycopeptides currently under clinical development. Middle panel: natural derivatives of vancomycin and teicoplanin used for the synthesis of oritavancin and dalbavancin, respectively. The part of the molecule that is common to all glycopeptides appears in bold. The arrows point to the differences between molecules.

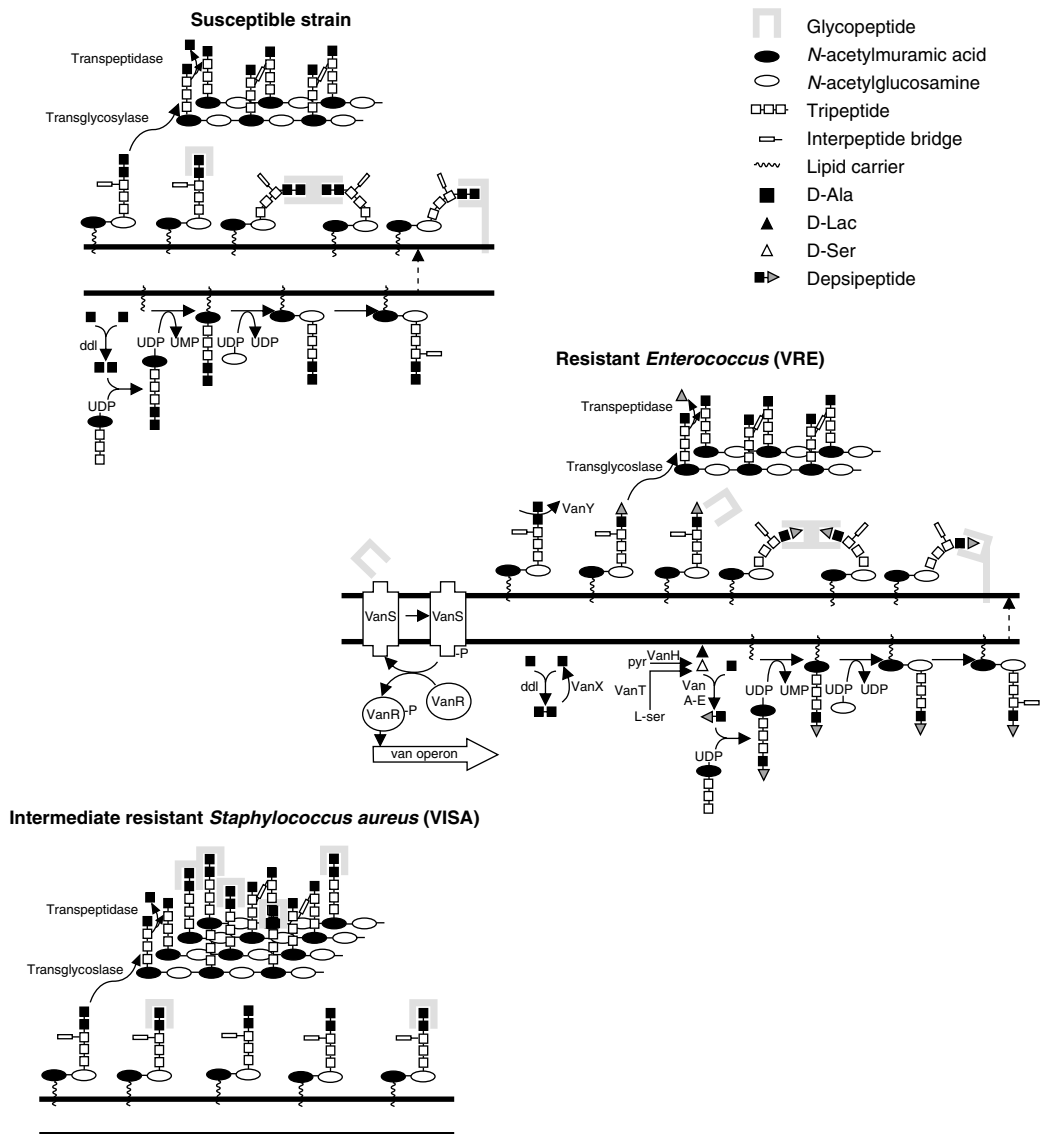


Fig. 2. Peptidoglycan synthesis in glycopeptide-susceptible bacteria (upper panel), in glycopeptide-resistant enterococci (middle panel) and in glycopeptide-intermediate staphylococci (lower panel).^[2,5] In susceptible strains, peptidoglycan precursors are ending by D-Ala-D-Ala termini synthesised by D-Ala:D-Ala ligases (ddl). Glycopeptides interact with D-Ala-D-Ala termini of pentapeptides (through five hydrogen bonds) at the cell surface, preventing transglycosylation reactions and therefore peptidoglycan reticulation. The figure also illustrates how dimerisation of glycopeptide molecules or anchoring in the membrane by a hydrophobic tail may increase the tightness of this interaction. Hydrophobic derivatives may also inhibit the transglycosylation step without binding to peptidoglycan precursors. In glycopeptide-resistant enterococci, the presence of glycopeptides activates the signal-transducing system VanS (sensor) – VanR (regulator), which allows transcription of the resistance genes. These include enzymes involved in the hydrolysis of D-Ala-D-Ala termini (VanX and VanY) or in the synthesis of precursors ending by either D-Ala-D-Lac (VanH and VanA, VanB or VanD) or D-Ala-D-Ser (VanT and VanC, VanE or VanG), characterised by a lower affinity for glycopeptides. In glycopeptide-intermediate staphylococci (so-called VISA [vancomycin-intermediate *Staphylococcus aureus*] or GISA [glycopeptide-intermediate *S. aureus*] strains), more murein monomers are supplied and incorporated in the peptidoglycan, in which more free D-Ala-D-Ala residues remain present because of a decreased level of reticulation. More glycopeptide molecules are therefore trapped in the multiple layers of the thicker cell wall but fewer reach their target at the cell membrane. **UDP** = uridine diphosphate; **UMP** = uridine monophosphate; **VRE** = vancomycin-resistant enterococci.

Table 1. Main characteristics of glycopeptide-resistant bacteria

Resistance type	Bacterial species	Expression	Mechanism of resistance	MIC range of glycopeptides (mg/L)		References
				vancomycin	teicoplanin	
VanA	<i>Enterococcus faecium</i>	Inducible (by vancomycin and teicoplanin)	Modified target (D-Ala-D-Lac)	64–1000	16–512	16
	<i>E. faecalis</i>					
	<i>E. avium</i>					
	<i>E. durans</i>					
	<i>E. hirae</i>					
	<i>E. mundii</i>					
	<i>E. raffinosus</i>					
	<i>E. gallinarum</i>					
	<i>E. casseliflavus</i>					
	<i>Staphylococcus aureus</i>	Inducible (by vancomycin and teicoplanin)	Modified target (D-Ala-D-Lac)			17-20
VanB	<i>E. faecium</i>	Inducible (by vancomycin)	Modified target (D-Ala-D-Lac)	4–1000	0.5–1	16
	<i>E. faecalis</i>					
	<i>Streptococcus bovis</i>					
VanC	<i>E. gallinarum</i>	Constitutive/inducible	Modified target (D-Ala-D-Ser)	2–32	0.5–1	16
	<i>E. casseliflavus</i>					
	<i>E. flavescens</i>					
VanD	<i>E. faecium</i>	Constitutive	Modified target (D-Ala-D-Lac)	64–128	4–64	16,22
VanE	<i>E. faecalis</i>	Inducible (by vancomycin)	Modified target (D-Ala-D-Ser)	16	0.5	16,23
VanG	<i>E. faecalis</i>	Inducible (by vancomycin)	Modified target (D-Ala-D-Ser)	16	0.5	24
VISA	<i>S. aureus</i>		Thickened cell wall with increased proportion of D-Ala-D-Ala termini	8	8–32	5,25
GRSA	<i>S. aureus</i>		Stop codon in the gene of PBP4			26

GRSA = glycopeptide-resistant *Staphylococcus aureus*; **MIC** = minimum inhibitory concentration; **VISA** = vancomycin-intermediate *Staphylococcus aureus*.

antimicrobial use and surveillance to try to control or even reverse the situation.^[29] In sharp contrast, resistance of enterococci in Europe is minimal (3%) and observed only in nosocomial infections;^[31] outbreaks have been reported but remain sporadic. Carriage of resistant enterococci in animals, however, is frequent in Europe and is thought to be due to the massive use of avoparcin, a glycopeptide antibiotic used as growth promoter in animal feed,^[32,33] but which has the same mode of action as vancomycin and, therefore, shares cross-resistance. The ban on avoparcin use instituted by EU countries has caused, in most countries, a fall in the prevalence of resistance in both animals and humans.^[29,34-36]

The mechanism of resistance in enterococci relies on synthesis of peptidoglycan by an alternative pathway, which produces precursors ending in D-Ala-D-Lac or D-Ala-D-Ser instead of D-Ala-D-Ala and concomitantly eliminates precursors ending in D-Ala-D-Ala. The replacement of D-Ala by D-Lac suppresses one of the five hydrogen bonds between the glycopeptides and their target, which leads to a 1000-fold decrease in the binding affinity.^[2,37] The substitution in D-Ser causes a conformational change, which also reduces vancomycin affinity, although not as markedly as with D-Lac. To be phenotypically detectable and significant, resistance requires the coordinated action of several enzymes (see figure 2). Thus, the bacteria need to synthesise

Table II. *In vitro* activity of oritavancin and dalbavancin compared with those of vancomycin and teicoplanin^a

Organism	MIC range (mg/L)				References
	vancomycin	oritavancin	teicoplanin	dalbavancin	
<i>Staphylococcus aureus</i>					
methicillin-susceptible	0.13–1	0.13–1	0.25–8	≤0.03–0.5	49
	0.25–2	0.06–4	0.125–4	0.06–0.5	50-55
methicillin-resistant	0.5–4	0.13–4	0.13–8	0.06–1	49
	0.25–4	0.03–8	0.06–8	0.06–1	50-55
vancomycin-intermediate	8	1–8	8–32	2	56-58
<i>Staphylococcus epidermidis</i>					
methicillin-susceptible	0.13–1	0.25–1	0.25–16	≤0.03–0.25	49
	0.25–2	≤0.03–8	≤0.03–16		50,54
methicillin-resistant	1–4	0.25–4	1–16	≤0.03–1	49
	0.5–4	0.25–16	0.125–16		50,54
<i>Staphylococcus haemolyticus</i>					
methicillin-susceptible	1–4	0.06–1	1–32	≤0.03–0.25	49
	0.25–4	0.12–8	≤0.12–8		50,54
methicillin-resistant	0.5–8	0.13–1	2–128	≤0.03–4	49
	1–4	1–4	0.5–>16		50,54
Other coagulase-negative staphylococci					
methicillin-susceptible	0.5–2	0.06–0.5	0.13–4	≤0.03–0.13	49
	0.25–4	≤0.01–2	≤0.12–32	≤0.03–0.25	51-53
methicillin-resistant	0.5–4	≤0.03–0.5	0.06–32	≤0.03–0.13	49
	0.25–8	0.015–4	0.06–64	0.06–1	51-53
<i>Streptococcus pyogenes</i>	0.5–0.5	0.016–0.13	0.008–0.06	≤0.002–0.06	49
	0.25–0.5	0.06–0.5	≤0.03–0.12		55
<i>Streptococcus pneumoniae</i>					
penicillin-susceptible	0.13–0.5	≤0.002–0.06	0.008–0.06	0.016–0.13	49
	0.06–0.5	≤0.002–0.12	≤0.01–0.25		51,52,54,55
penicillin-resistant	0.25–2	≤0.002–0.06	0.016–0.13	0.008–0.13	49
	0.06–1	0.002–0.25	≤0.01–0.25		51,52,54,55
Viridans streptococci					
	0.25–2		≤0.12–2	≤0.03–0.06	53
penicillin-susceptible	0.12–1	≤0.01	≤0.01–0.5		52
penicillin-resistant	0.25–1	0.01–0.06	≤0.01–0.5		52
β-Haemolytic streptococci	0.25–1		≤0.12–0.25	≤0.03–0.12	53
<i>Enterococcus</i> spp.					
vancomycin-susceptible	0.25–4	0.06–0.25	0.13–0.5	0.06–0.13	49
	≤0.01–4	≤0.03–2	≤0.012–8	≤0.03–1	50-54,59
VanA	>128	0.06–1	64–>128	0.5–>128	49
	64–>256	0.06–1	32–>256		50
VanB	8–128	≤0.03–0.13	0.13–8	0.02–2	49
	8–128	0.25–1	0.125–0.5		60
VanC	4–16	≤0.03–1	0.125–4		50
<i>Bacillus</i> spp.	≤0.12–1	≤0.015–0.5	≤0.12–4	≤0.03–2	53,54
<i>Corynebacterium</i> spp.	0.25–0.5		≤0.12–1	≤0.03–0.12	53

Continued next page

Table II. Contd

Organism	MIC range (mg/L)				References
	vancomycin	oritavancin	teicoplanin	dalbavancin	
<i>Listeria</i> spp.	0.25–2	≤0.03–0.125	0.06–0.25		50
<i>Clostridium difficile</i>	0.5–4	0.016–2	0.064–0.5	0.125–0.5	61,62
<i>Clostridium perfringens</i>	0.025–4	0.016–2	0.064–4	0.03–0.125	61,62
<i>Haemophilus influenzae</i>	≥16	16–32		8–64	53,54

a When two rows of data appear for a single species, the first one refers to studies where the drugs are compared against the same strains; the second one represents the range of MICs reported in other publications.

MIC = minimum inhibitory concentration.

D-Lac (VanH) or D-Ser (VanT) and D-Ala-D-Lac or D-Ala-D-Ser (VanA, B, D or C, E, G), and to degrade D-Ala-D-Ala (VanX) or to remove D-Ala from growing precursors (VanY) or both (VanXY). Moreover, a two-component regulatory system (VanS-VanR) resulting in induction by either vancomycin (VanB, C, E, G phenotype) or vancomycin and teicoplanin (VanA phenotype) plays a critical role.^[2,16,22,23] The resistance proteins are encoded by genes physically grouped in operons that are located on plasmids or in the chromosome, and can be easily transferred, even between different species.^[2,38,39] Six phenotypes of resistance have been described (see table I), which differ by the genetic support, the regulation of expression and the level of resistance conferred. An intriguing phenotype of glycopeptide dependence has also been described, in which synthesis of the D-Ala-D-Ala-ending peptidoglycan precursors is impaired through a mutation in the host ligase. Induction of production of the resistance proteins by glycopeptides restores peptidoglycan synthesis by allowing production of precursors ending in D-Ala-D-Lac.^[40,41]

Resistance in *S. aureus* first emerged in the form of strains with elevated minimum inhibitory concentration (MIC) values towards vancomycin (or both vancomycin and teicoplanin), which have been named vancomycin-intermediate *S. aureus* (VISA) or glycopeptide-intermediate *S. aureus* (GISA). Originally described in Japan in 1996 in methicillin-resistant *S. aureus* (MRSA),^[25] VISA and GISA strains have now been isolated in numerous countries, particularly from patients having received prolonged vancomycin therapy.^[5,42] These bacteria are characterised by a thickened cell wall, production of

an abundant extracellular material of still ill-characterised nature and an impaired ability to divide.^[43] These phenotypic changes can be explained by the production of an altered peptidoglycan with an increased proportion of free D-Ala-D-Ala termini (less reticulation), which can trap vancomycin molecules and prevent their access to the target at the cytosolic membrane (figure 2). The thickened cell wall of a VISA strain may contain up to two to four times more D-Ala-D-Ala residues than that of a susceptible strain and is able to bind up to three to six times more vancomycin molecules before peptidoglycan synthesis becomes impaired.^[5] This mechanism also implies a reduced cross-linkage of peptidoglycan. The latter has been suggested to result from decreased activity of penicillin-binding proteins (PBPs)^[26] or from an alteration of murein precursors.^[44] Yet multiple, additive, but not clearly identified mutations, are probably necessary to obtain resistance. VISA and GISA also need to import a larger amount of precursors than normal strains, which compromises their fitness in an antibiotic-free environment. This explains why VISA and GISA tend to lose their resistance when relieved from vancomycin pressure, giving rise to the so-called hetero-VISA phenotype.^[45] Moreover, a glycopeptide-tolerant phenotype has been observed in clinical isolates of MRSA, in which MICs of glycopeptides are not affected but minimum bactericidal concentrations (MBCs) are considerably increased.^[46] In a still more frightening fashion, two MRSA strains with high levels of resistance to vancomycin and teicoplanin have now been reported in two different hospital institutions in the US.^[17-20] Both of these strains harbour the *vanA* gene cluster,

indicating that they had acquired the corresponding set of genes from enterococci. Should such strains spread in hospitals, treatment options would rapidly become very limited.^[47]

1.3 Pharmacological Properties

The spectrum of activity of glycopeptides covers essentially the Gram-positive organisms and a few anaerobes, and their activity against Gram-negative organisms is most often marginal (table II). Vancomycin and teicoplanin have a similar intrinsic activity, except against streptococci, which are more susceptible to teicoplanin.^[48]

The key pharmacokinetic properties of glycopeptides are summarised in table III. The most striking difference between vancomycin and teicoplanin is their capacity to bind serum proteins, which is much higher for teicoplanin than for vancomycin. This explains the prolonged half-life of teicoplanin in the organism. However, high protein binding also needs to be taken into account for activity predictions (table IV), since only the free serum fraction is directly active. Teicoplanin, perhaps because of its lipophilic character, is characterised by a higher volume of distribution than vancomycin, allowing it to reach therapeutic concentrations in fat, muscles (including pericardium and myocardium) and, to

some extent, bone and cartilage, but not in the CNS.^[63]

In vitro pharmacodynamic models show that vancomycin exhibits time-dependent bactericidal activity against most Gram-positive organisms. However, vancomycin is essentially bacteriostatic against enterococci^[71] but may become bactericidal when combined with an aminoglycoside.^[72,73] Vancomycin activity is adversely affected by a large inoculum but not by acidic pH. The drug also demonstrates faster killing rates against actively growing organisms, which can be easily understood on the basis of its mechanism of action.^[74] Finally, glycopeptides produce persistent effects (postantibiotic effect lasting 1–6 hours^[71,75]). While the dosage recommendations for glycopeptides have insisted on low dosages because of the fear of toxicity, recent developments in our understanding of pharmacodynamics have allowed a more rational basis for recommendations. The first models suggested that the free serum concentration of glycopeptides needs to remain above the MIC for the infecting organism for a prolonged period and, therefore, concluded that activity was primarily driven by the so-called ‘time above MIC’ parameter.^[76,77] However, more recent and thorough studies have indicated that the parameter which best predicts efficacy is the area under the plasma concentration-time curve (AUC)/MIC ratio,^[56] as usually found for most antimicrobials

Table III. Pharmacokinetic properties of glycopeptides as observed after administration of conventional clinical doses^a

Parameter	Vancomycin	Oritavancin	Teicoplanin	Dalbavancin
Peak concentration (mg/L)	20–50	31 ^b	43	300
Trough concentration (mg/L)	5–12 (24h)	1.7 (24h) ^p	<5 (24h)	40 (168h)
Protein binding (albumin) [%]	10–55	90	90	98
Volume of distribution (L/kg)	0.3		0.9–1.6	0.16
Terminal half-life (h)	4–8	360 ^b	83–168	257
Clearance (L/h/kg)	0.058		0.011	0.0006
Renal excretion (%)	80–90	<5 in 14 days ^c	80	42
Calculated AUC (mg • h/L)	260	152 ^{b,d}	550	23 250 ^e

a Vancomycin 1g or 15 mg/kg;^[64] oritavancin 3 mg/kg (in clinical trials);^[65] teicoplanin 6 mg/kg;^[63] dalbavancin 15 mg/kg or 1g (in clinical trials).^[66–68]

b Population pharmacokinetic analysis, data on file, InterMune, Inc.^[69]

c Data on file, InterMune, Inc.^[70]

d AUC 0–24h.

e AUC 0–∞.

AUC = area under the concentration-time curve.

Table IV. Pharmacodynamic parameters predictive of glycopeptide efficiency calculated for conventional dosages (see table III)

PK/PD parameter	Maximum reachable MIC based on PK/PD parameters			
	vancomycin 15 mg/kg bid	teicoplanin 6 mg/kg od	oritavancin 3 mg/kg od	dalbavancin 15 mg/kg od
AUC/MIC = 125^a				
total	4	4	1	185
free	2	0.4	0.1	4
C_{max}/MIC = 10^a				
total	10	4	3	30
free	5	0.4	0.3	0.6

a Target values based on general PD considerations for antibiotics in the absence of actual data for all glycopeptides (lower C_{max}/MIC ratio may be sufficient on the basis of available data for vancomycin and teicoplanin).

AUC = area under the concentration-time curve; **bid** = twice daily; **C_{max}** = peak concentration; **MIC** = minimum inhibitory concentration; **od** = once daily; **PD** = pharmacodynamic; **PK** = pharmacokinetic.

exhibiting time-dependent killing and significant postantibiotic effect.^[78] However, in non-neutropenic animals, the ratio of the free serum peak concentration to the MIC (C_{max-free}/MIC) also plays a determinant role in efficacy.^[79] This C_{max-free}/MIC ratio needs to reach at least 5–6 for vancomycin and 2–3 for teicoplanin (note that the determination of the peak and trough levels for these drugs, which are determined in routine in hospital laboratories, can be used for calculation of both C_{max}/MIC and AUC/MIC ratios). As a consequence, glycopeptides are probably best given by discrete administrations with a total daily dose sufficiently large to match with the MIC of the infecting organism. A recent study comparing vancomycin 30 mg/kg once daily versus two administrations for the treatment of patients with bacteraemia or arthritis did not show a significant difference between the two groups in terms of clinical response (>92% in both cases;^[80] however, no patients with endocarditis were included in this study). On a pharmacodynamic basis, there is therefore little reason to administer vancomycin by continuous infusion^[81,82] even though other parameters such as more sustained concentrations in given tissues, or cost containment and ease of administration, have been advocated.^[83,84] (Note that the stability and compatibility of vancomycin with other drugs in relation to its use by continuous infusion are currently under investigation.^[85])

1.4 Clinical Use: Pros and Cons

Originally introduced into clinical practice as an agent active against β-lactamase-producing *S. aureus*, vancomycin remained largely unused because of development of less toxic alternatives, namely the β-lactamase-resistant penicillins (methicillin, isoxazolyl penicillins) and cephalosporins, as well as the introduction of β-lactamase inhibitors such as clavulanic acid (mainly in Europe) or sulbactam, which have become highly popular when combined with ampicillin or amoxicillin (mostly against Gram-negative bacteria, however). Yet orally administered vancomycin became popular for treatment of *Clostridium difficile*-associated diarrhoea and colitis.^[86] The pandemic of nosocomial MRSA infections which started in the mid 1970s, and the fact that these strains were resistant not only to all β-lactam agents but often also to aminoglycosides, macrolides, lincosamides and fluoroquinolones,^[87] heralded the return of vancomycin for systemic use. Alternatives such as fusidic acid, rifampicin and cotrimoxazole have indeed been poorly studied and did not compete in the face of the extensive data and the everyday experience available with glycopeptides.^[88] Moreover, easy selection of resistant mutants (especially for rifampicin and fusidic acid when used alone) or toxicity problems has further limited their use. As we have seen, however, resistance developed in enterococci almost in parallel with the increase in vancomycin use in the US.^[89-91] This led health authorities and concerned clinicians to try to restrict glycopeptide usage

in order to maintain the activity of these useful agents for situations where alternatives were not usable.^[90,92-95] The introduction of quinupristin/dalfopristin and linezolid in the late 1990s has not really changed this situation, since both drugs are expensive, have rare but potentially worrying adverse effects (haematological toxicity for long-course linezolid treatments, thrombophlebitis, arthralgias and/or myalgias, and drug interactions for quinupristin/dalfopristin), and are already facing emergence of resistance.^[96-101] In particular, linezolid-resistant strains are now increasingly reported, not only in enterococci but also in *S. aureus*, which could become a clinically significant problem in the future.

The recommendations presented in table V were issued in 1995 by the Hospital Infection Control Practices Advisory Committee in the US (HICPAC^[102]) and have been adopted, with adaptations, in most countries (see Gordts et al.^[103] for Belgium). These recommendations have been further extended to special populations (see Nourse et al.^[104] for children). In summary, and outside the field of bacteriologically documented MRSA infections, empirical therapy with glycopeptides should be limited to severe infections in immunocompromised patients (e.g. burn patients, clinical sepsis in the intensive care unit) when local epidemiological data show a high percentage of MRSA, and/or to life-threatening infections associated with the presence of foreign bodies (often infected with methicillin-resistant coagulase-negative staphylococci; e.g. indwelling percutaneous catheters, prosthetic cardiac valves). Indeed, animal models and clinical studies have shown that vancomycin is less bactericidal than isoxazolyl penicillins against methicillin-sensitive staphylococci (causing prolonged periods of fever and persistence of positive blood cultures). This should preclude its use (and, despite fewer data, the use of teicoplanin) when such strains are involved, especially in life-threatening infections (e.g. endocarditis). Therefore, their indiscriminate use in non-documented infections (or infections caused by Gram-positive strains susceptible to other common agents) should be discouraged

Table V. Appropriate clinical use of glycopeptides, based on the recommendations from Hospital Infection Control Practice Advisory Committee (HICPAC)^[102] for vancomycin and on pharmacodynamic considerations (table IV and Gruneberg et al.^[105])

Situations in which the use of glycopeptides is appropriate or acceptable

Treatment

Serious infections caused by β -lactam-resistant Gram-positive microorganisms

Infections caused by Gram-positive microorganisms in patients who have serious allergies to β -lactam agents

Antibiotic-associated colitis failing to respond to metronidazole therapy or potentially life-threatening

Prophylaxis

Endocarditis following certain procedures in patients at high risk for endocarditis

Major surgical procedures involving implantation of prosthetic materials or devices at institutions that have a high rate of infections caused by methicillin-resistant *Staphylococcus aureus* (MRSA) or methicillin-resistant *S. epidermidis*

Situations in which the use of glycopeptides should be discouraged

Routine surgical prophylaxis other than in a patient who has a life-threatening allergy to β -lactam agents

Empirical antimicrobial therapy for a febrile neutropenic patient, unless initial evidence indicates that the patient has an infection caused by Gram-positive microorganisms and the prevalence of infections caused by MRSA in the hospital is substantial

Treatment in response to a single blood culture positive for coagulase-negative staphylococci, if other blood cultures taken during the same time frame are negative

Continued empirical use for presumed infections in patients whose cultures are negative for β -lactam-resistant Gram-positive microorganisms

Appropriate dosages^{a, b}

Vancomycin 15 mg/kg twice daily

Teicoplanin 6–12 mg/kg once daily (after three loading doses of 6 mg/kg every 12 hours)

a Higher intravenous doses are needed for treating peritonitis in patients undergoing peritoneal dialysis.^[106,107]

b Dosage reduction required in patients with renal insufficiency.^[63,64]

with the exception of true β -lactam allergy. Of these severe infections by multiresistant staphylococci, the main indications for vancomycin are, therefore, restricted to: (i) serious diphtheroid infections (when the strain is penicillin resistant or the patient has IgE-mediated allergy to β -lactam agents); (ii) penicillin-resistant *Streptococcus pneumoniae* infections of the CNS in combination with cefotaxime or ceftriaxone; and (iii) antibiotic-associated colitis that is life-threatening or fails to respond to metroni-

dazole (only oral vancomycin; low faecal concentrations by intravenous route and few data with teicoplanin). Routine prophylaxis with glycopeptides should also be prohibited^[102,103] and restricted to prevention of bacterial endocarditis in penicillin/ampicillin-allergic patients at risk, undergoing gastrointestinal or genitourinary procedures (plus gentamicin), or dental procedures (vancomycin alone). In selected surgical wards with a high incidence of methicillin-resistant staphylococci infections, a single dose before surgery may be used in prosthetic surgery, but one should always try to solve the underlying fundamental hygiene problem. Teicoplanin could be used in most of the same indications, with the exception of the CNS infection, because of the poor penetration of the drug in the infected compartment. Its main advantages are the possibility of administering it by the intramuscular route with a once-daily schedule (because of its longer half-life) and a lower incidence of adverse effects. However, several indications are less documented than for vancomycin, and there are no well conducted, large, comparative studies between the two drugs. In severe infections (as in *S. aureus* endocarditis), teicoplanin needs to be given at a dose of at least 12 mg/kg/day once daily, after a minimum of three to four loading doses (each given every 12 hours).

Safety issues need also to be taken into account and may limit glycopeptide use in given populations. Earlier, vancomycin was notorious for toxicity related to impurities and to histamine release (causing the so-called 'red man syndrome'). These adverse effects have been markedly reduced through better purification procedures and by giving the drug as a slow infusion over at least 1 hour.^[108-110] With these precautions, vancomycin can be safely used, causing only relatively mild and self-limiting general toxicity.^[111,112] However, the dose should be corrected in the case of renal insufficiency. The main adverse effects are phlebitis at the site of injection as well as nephrotoxicity and ototoxicity, the latter remaining the most problematic because it can be irreversible.^[64] Both nephro- and ototoxicity are aggravated by the concomitant use of aminog-

lycosides or other drugs toxic for these organs.^[112] Infrequent toxic manifestations include neutropenia, thrombocytopenia, fever, bullous dermatosis, necrotising cutaneous vasculitis and toxic epidermal necrolysis.^[113] Teicoplanin has been claimed to be less toxic than vancomycin.^[114] However, a difficulty relates to the fact that the recommended doses of teicoplanin have been increasing over time and that most comparative studies with vancomycin were done with teicoplanin doses (typically ≤ 6 mg/kg) that are now considered insufficient (table V). Therefore, further studies are needed to evaluate toxicity of teicoplanin in the current conditions of use.

2. New Developments in the Glycopeptide Area

Given the potential of glycopeptides in severe infections, considerable effort has been made to obtain new derivatives with improved pharmacological properties and activity against resistant strains. Because of a highly complex structure, which makes total synthesis of glycopeptides very challenging,^[115] most of the new molecules obtained so far are semisynthetic derivatives of existing, natural glycopeptides. Structure-activity relationships (SARs) are presented in figure 3, based on our current understanding of the molecular mode of interaction between glycopeptides and their pharmacological target. Modifications can be subdivided into three main categories, each of them concerning distinct domains of the molecule and affecting the general mode of action of vancomycin in various ways. Other recent strategies have explored the formation of hybrid or dimerised molecules, or the synthesis of inhibitors of the resistance mechanisms to be administered in combination with unmodified glycopeptides.

2.1 Semisynthetic Modification of Natural Glycopeptides

2.1.1 Modification of the Binding Pocket

Binding studies have shown that five sites on the peptidic backbone of glycopeptides are involved in

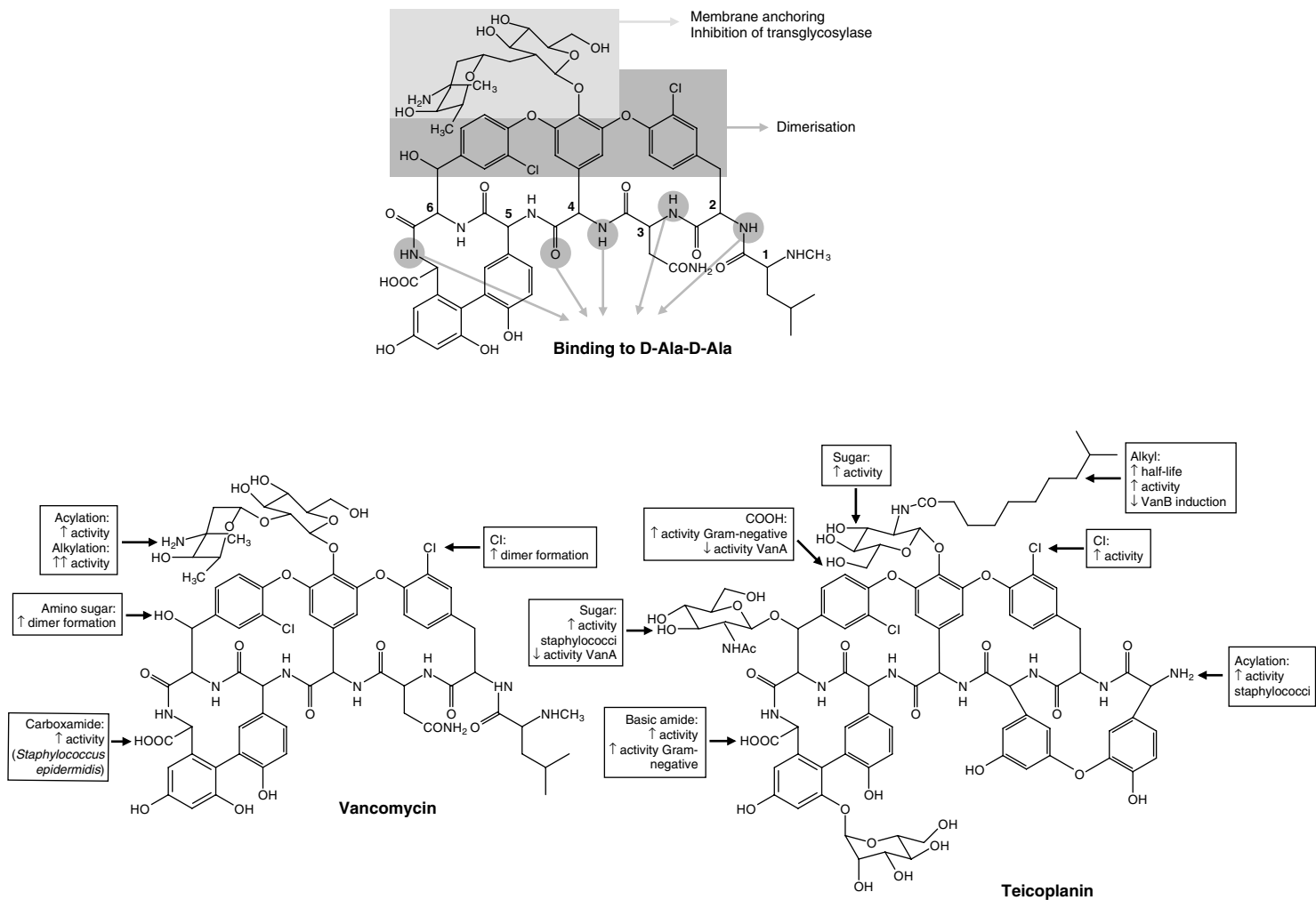


Fig. 3. Structure-activity relationship (SAR) of glycopeptides derived from vancomycin and teicoplanin. Upper panel: current view of the parts of the molecules involved in the various modes of action proposed for glycopeptides. Lower panel: most salient SARs that can be drawn for vancomycin (left) or teicoplanin (right) derivatives. ↑ indicates increased; ↓ indicates decreased.

the interaction with the D-Ala-D-Ala termini (figure 3). Efforts have been made to change these amino acid residues, with the aim to generate compounds with increased affinity for both dipeptide and deipeptide termini. These have included trimming or enlarging the heptapeptide backbone, epimerisation of the C3 chiral centre (in teicoplanin only), or substitution of amino acids 1 and 3.^[116] Unfortunately, most of these modifications have resulted in a reduction of antibacterial activity. Nevertheless, the possibility of obtaining active compounds by this type of approach remains open, based on the observation that substitution of the Asn at position 3 by a hydrophobic amino acid enhances the affinity of vancomycin for D-Ala-D-Lac precursors.^[117]

2.1.2 Modification or Addition of Functional Groups

A few functional groups can be modified out of the binding pocket, essentially causing changes in physicochemical properties. These studies have generated useful knowledge in SAR and have allowed changes in the natural compounds to be proposed.

Modification of Vancomycin

As stated earlier, the affinity of vancomycin for its target is increased by its capacity to dimerise. This process is facilitated by the addition of an amino sugar to residue 6 and by the presence of a chlorine atom in meta on the aromatic substituent of residue 2.^[10,13,116] Other modifications at the extremities of the peptidic backbone have also brought significant pieces of information. Thus, replacement of the free carboxylate terminus by a carboxamide increases the activity on *Staphylococcus epidermidis*, whereas D-amino acids at the N-terminus appear more active than their corresponding L-isomers.^[116]

Modification of Teicoplanin

As for vancomycin, structure-activity studies of teicoplanin have examined the influence of substituting the extremities of the peptidic backbone. At the C-terminus, substitution by a basic, positively charged amide considerably increases activity on staphylococci and also confers moderate activity on Gram-negative bacteria, because of improved ability

to cross their outer membrane.^[118] Esterification with hydrophobic alcohols also leads to marginal activity on Gram-negative bacteria. Similarly, basic amides of the A40,926 derivative of teicoplanin (which lacks the *N*-acetylglucosamine sugar; figure 1) show increased activity against staphylococci (particularly coagulase-negative) and streptococci. The most active compounds also show moderate activity against VanA enterococci.^[119,120] At the N-terminus, acylation or alkylation causes either a slight increase or no modification in activity. Another important modification in the teicoplanin backbone is the introduction of chlorine atoms, which increase both target affinity and antibacterial activity.^[116]

2.1.3 Alteration of Sugars

Despite their location apparently far from the binding pocket, sugars play a determinant role in glycopeptide activity, since aglycones of both vancomycin and teicoplanin are systematically less active than their parent compounds.^[116,121] Substituting these sugars can actually have a considerable impact on glycopeptide pharmacological properties.

Modification of Vancomycin

Besides the beneficial addition of a 4-*epi*-vancosamine on the benzylic function of residue 6,^[122] substitution of the vancosamine sugar by hydrophobic substituents proved mostly useful, probably by somehow mimicking the lipophilic side chain of teicoplanin derivatives. *N*-alkyl derivatives were found to be more active than *N*-acyl products, particularly against enterococci, streptococci and staphylococci.^[116] When combined with the additional 4-*epi*-vancosamine (as in LY264826; see figure 1), an alkyl side chain confers to the molecule an unexpectedly rapid, concentration-dependent bactericidal activity, including against VanA- and VanB-type resistant enterococci.^[123-125] The activity of these hydrophobic derivatives on resistant strains has been proposed to result from the combination of a facilitation of the binding to the peptidoglycan precursors due to their anchoring in the membrane^[9,11] and of a direct inhibition of transglycosylases.^[15,126] Recent studies also suggest that some of these derivatives may act as inhibitors not only of

cell wall synthesis but also of phospholipid synthesis.^[127] Acidic and hydrophilic substituents have also been added on the resorcinol in alpha of the free carboxylate of lipidated analogues of vancomycin, with favourable consequences on distribution and safety profiles, without negative effect on activity against MRSA or vancomycin-resistant enterococci (VRE).^[128]

Modification of Teicoplanin

The impact of sugars on the activity is emphasised by the fact that the removal of *N*-acylglucosamine reduces activity, particularly against streptococci and enterococci, whereas removal of *N*-acetylglucosamine decreases the activity against staphylococci but may confer activity against Gram-negative as well as VanA-type enterococci. This is especially true for derivatives that also have a basic amide substitution of the C-terminus.^[116]

Substituents on sugars also play important roles. As mentioned in the previous section, lipophilic substituents are of major interest for vancomycin activity. Teicoplanin contains naturally a lipophilic side chain that is responsible for its prolonged half-life. This side chain is also important for activity, since deacylated derivatives are only weak antibiotics.^[116] The acyl chain has been proposed not only to favour anchoring in the membrane, and therefore the interaction with the target,^[111] but also to circumvent VanB induction,^[129] explaining teicoplanin activity against resistant strains of the VanB-type. On the other hand, the replacement of the *N*-acetylglucosamine by an *N*-acylglucuronic acid (as in A40926, see figure 1) has a detrimental effect on activity against VanA enterococci but improves anti-Gram-negative activity.^[116]

2.2 Hybrid and Multivalent Glycopeptides

On the basis of current knowledge on the mode of action of glycopeptides, more innovative strategies have been explored to optimise activity against both susceptible and resistant strains. First, considering the potential cooperative binding occurring in multivalent drugs, as well as the enhanced activity of glycopeptides capable of self-association, dimers of vancomycin have been prepared using different

linkers and various linkage positions.^[130,131] Improved activity has been observed against Gram-positive organisms, including GISA and VRE strains, opening promising lines of investigation in this area. Secondly, taking into account the possibility of a dual target for glycopeptides, namely the dipeptide termini of peptidoglycan precursors, and the enzymes involved in the transglycosylation step, vancomycin derivatives have been prepared where the aglycone is separated from the sugars by linkers, with the aim to correctly position each part of the molecule.^[132] This approach could now be extended to other glycopeptides in which both the aglycone and the carbohydrate moieties would be individually optimised for interaction with their own target, so as to generate very active compounds.

2.3 Inhibitors of Resistance Mechanisms

In the same way as clavulanic acid or sulbactam have been designed to restore the activity of β -lactam agents against β -lactamase-producing organisms, inhibitors of glycopeptide resistance mechanisms could be considered to specifically circumvent the mechanisms of resistance developed by enterococci (which, as described in section 1.2, may also apply to the recently described highly resistant *S. aureus* strains). This goal has been achieved by designing small molecules such as ϵ -aminopentanoylated prolinol,^[133] which are able to selectively cleave the ester linkage of the D-Ala-D-Lac dipeptide of resistant bacteria. Concomitant administration of this compound and vancomycin increases the activity of the glycopeptide *in vitro* against VanA-type enterococci, without affecting its activity against susceptible strains. However, activity remains globally low, probably because VanX hydrolyses the D-Ala-D-Ala dipeptide, so that there is no pentapeptide left for binding glycopeptides. These inhibitors will, therefore, need to be combined with inhibitors of the VanX D-D-peptidase.^[134,135] They have been obtained with inhibitory potencies in the micromolar range, but microbiological data are still lacking to demonstrate the interest of combining these inhibitors with vancomycin.

3. Properties of the Ideal Glycopeptide

Based on an examination of the limitations of the current molecules and also on the promising properties of some of the semisynthetic derivatives described in section 2, we delineate in table VI the main desirable characteristics expected from new glycopeptides for future clinical development. Two molecules, oritavancin and dalbavancin, fulfil these conditions in various ways and are currently in phase II/III of clinical development.

4. Glycopeptides Undergoing Clinical Evaluation

4.1 Oritavancin

Oritavancin (LY333328) was discovered at Eli Lilly (Indianapolis, IN, USA) but is now developed by InterMune (Brisbane, CA, USA)^[136] [see figure 1 for structure]. This molecule was obtained by reductive alkylation with 4'-chloro-biphenyl-carboxaldehyde of A82846B (chloroeremomycin; LY264826), a natural glycopeptide that differs from vancomycin by the addition of a 4-*epi*-vancosamine sugar and the replacement of the vancosamine of the disaccharide moiety by an *epi*-vancosamine.^[122] The design of this drug was based on previous observations that chloroeremomycin was more potent than vancomycin because of its stronger tendency to self-associate,^[3] and that substitution of the van-

cosaminy moiety of vancomycin by a lipophilic side chain resulted in enhanced activity and in some cases restored activity towards vancomycin-resistant enterococci.^[121,137]

4.1.1 Microbiological and Pharmacological Properties

Spectrum of Activity and Resistance

The activity of oritavancin (table II) is similar to that of vancomycin against staphylococci, with MICs ranging between 0.03 and 8 mg/L; no differences are seen between methicillin-susceptible and -resistant stains. Against enterococci, oritavancin is more active than vancomycin or teicoplanin, with MICs consistently lower than 1 mg/L. Remarkably, it is as active against glycopeptide-resistant strains as against glycopeptide-susceptible isolates. Oritavancin is also very potent against pneumococci (both penicillin-susceptible and -resistant). It shows low MICs against other Gram-positive species including other streptococci, *Listeria* spp., *Clostridium* spp. and corynebacteria,^[50] but is not active against Gram-negative bacteria, including *Haemophilus influenzae*. Resistance of *S. aureus* to oritavancin has so far not been described in clinical isolates,^[51,52,138] including VISA strains. In contrast, enterococci with reduced susceptibility to oritavancin (MIC 8–16 mg/L) have been obtained *in vitro*, using strains harbouring the *vanA* or the *vanB* gene clusters.^[139] Three mechanisms seem to operate:^[139] (i) the complete elimination of D-Ala-ending precursors by overexpression of the *vanA* gene cluster or by reduced expression of the host D-Ala:D-Ala ligase; (ii) mutations in the VanSB sensor of the *vanB* cluster (which can occur in clinical isolates^[140]); or (iii) the expression of VanZ, a protein encoded by a gene present in the resistance transposon but the precise function of which is still unknown. A clinical strain of oritavancin-dependent *Enterococcus faecalis* has also been isolated but the underlying mechanism should be different from that conferring dependence to vancomycin and teicoplanin, since the growth of this strain is not restored in the presence of D-Ala-D-Ala precursors.^[141]

Table VI. Properties of an ideal glycopeptide

Microbiological properties
High intrinsic activity against Gram-positive organisms, including methicillin-resistant <i>Staphylococcus aureus</i> , vancomycin-resistant enterococci and glycopeptide-resistant <i>S. aureus</i>
Pharmacodynamic properties
Rapid and concentration-dependent bactericidal activity
Pharmacokinetic properties
Area under the concentration/time curve and peak plasma concentration for free fraction adequate to cover minimum inhibitory concentration of target pathogens
Prolonged half-life (once daily administration)
High diffusibility in tissues, including in the CNS
Safety profile
Lower incidence of adverse effects than current molecules

Pharmacodynamic Properties in Relation to the Mode of Action

Oritavancin is bactericidal *in vitro*, with MBCs only 1- to 8-fold higher than the corresponding MICs against most of the covered microorganisms.^[50] Killing curve experiments have revealed a striking difference between vancomycin and oritavancin concerning the rate of killing and its concentration-dependent character. Oritavancin indeed shows very rapid and highly concentration-dependent bactericidal activity (3-log reduction in bacterial counts after 1–8 hours) in conditions where vancomycin requires at least 8–24 hours to reach the same effect.^[50,142] However, oritavancin acts more slowly against VRE.^[143] These properties suggest that oritavancin could have another mode of action than conventional glycopeptides. The current view is that oritavancin combines the advantages of a high capacity to dimerise and to interact with the membrane, and therefore may cooperatively bind to peptidoglycan residues of both susceptible and resistant strains. It may also directly perturb the membrane properties and inhibit transglycosylation reactions.^[125]

Oritavancin also displays a concentration-dependent postantibiotic effect, increasing from ~2 hours at 1 × MIC to 4 hours against VRE and 8 hours against MRSA.^[143] As with vancomycin, the combination of oritavancin with gentamicin is synergistic.^[143-145] The combination with ampicillin enhances the bactericidal activity of oritavancin, without being truly synergistic. It also prolongs its postantibiotic effect against VRE from 18 to 23 hours at 10 × MIC.^[146]

Oritavancin activity is negatively affected by large inocula,^[143] but not by acid pH or by the growth phase of the bacteria.^[147] However, as pointed out by Mercier et al.,^[147] activity might be slightly reduced against VRE in stationary growth phase, as observed in animal model infective endocarditis or in acidic foci of infection.

Pharmacokinetic Profile and Relationship with Pharmacodynamic Properties

The pharmacokinetic properties of oritavancin are presented in table III in comparison with those of

the other glycopeptides. Like teicoplanin, oritavancin is characterised by high protein binding to albumin, which explains its prolonged retention in mammals, but also reduces the rate and extent of its bactericidal activity and shortens the postantibiotic effect.^[146,148] This could be important when serum levels become close to the MIC for the infecting organism. Further studies should address pharmacokinetic-pharmacodynamic relationships for these drugs, but we may anticipate, on the basis of its concentration-dependent bactericidal activity *in vitro*, that high serum levels or AUC relative to MIC will correlate with optimal efficacy. Studies on cultured cells also show that oritavancin accumulates in eukaryotic cells to exceptional levels (the apparent cellular concentration reaching values as high as 400 times the extracellular ones), which may be an advantage for the eradication of intracellular infections.^[149]

Models of Infection

The promising *in vitro* activity of oritavancin has prompted evaluation of its activity in models of difficult-to-treat infections. As anticipated, oritavancin given once daily is as effective as vancomycin given every 8 hours against MRSA rabbit endocarditis^[150] and is bactericidal against *S. pneumoniae* rabbit meningitis,^[151,152] even though the drug penetration in the CSF reaches only 5% of the serum concentration.^[152] Oritavancin activity has also been studied in models of infections by glycopeptide-resistant enterococci. While oritavancin was effective in a model of rat central venous catheter infection,^[153] its activity was more limited in a rabbit endocarditis model, causing a reduction in the number of bacteria in the vegetations but failing to sterilise them.^[154] Increasing serum concentrations to values as high as 80 mg/L did not provide significant improvement and did not prevent the selection of resistant mutants, probably because of the heterogeneous distribution of the drug in the vegetations. Yet combination with gentamicin proved synergistic and bactericidal, and prevented the emergence of resistant mutants.^[155] Intracellular bactericidal activity of oritavancin has been demonstrated in *in vitro* models of polymorphonu-

clear leucocytes infected by vancomycin-resistant enterococci^[156] or by MRSA^[157] as well as in macrophages infected by *S. aureus*.^[158] Taken together, these data suggest a definite clinical interest for this drug, provided it can reach the infected compartment at sufficiently high concentrations. Efficacy against infections occurring in less accessible compartments as well as against intracellular organisms needs, therefore, to be examined in the light of the tissue distribution of the drug *in vivo*.

4.1.2 Clinical Studies

Oritavancin is currently in phase III development, with two studies completed in the treatment of patients with complicated skin and skin structure infection caused by Gram-positive pathogens including MRSA.^[159] The intent-to-treat analysis of one of these, a double-blind, randomised study, has been presented as an abstract^[160] and showed equivalent clinical success with oritavancin at 1.5 or 3 mg/kg for 3–7 days versus vancomycin (15 mg/kg twice daily for 3–7 days followed by oral cephalexin for up to 10–14 days in the vancomycin group).

4.1.3 Safety Profile

The only published data available so far are those of the phase I studies, in which oritavancin was well tolerated.^[161] However, studies on cultured cells indicate that the drug could induce lipid storage, probably in relation to its high level of cellular accumulation,^[162] which may suggest further evaluation in animal models.

4.2 Dalbavancin

Dalbavancin (BI397) is a semisynthetic derivative from the natural glycopeptide A40926 (see figure 1 for structure). It was discovered by Biosearch Italia and outlicensed for North America to Versicor. The two companies have now merged to create Vicuron Pharmaceuticals (King of Prussia, PA, USA)^[163] and continue to develop the product. A40926 differs from teicoplanin by the absence of the acetylglucosamine in the benzylic position, replacement of the acylglucosamine in position 4 by an acylaminoglucuronic acid, the length of the fatty acid chain, the position of one chlorine atom and the

terminal methylamino group.^[164] In dalbavancin, a 3,3-dimethylaminopropylamide replaces the peptide carboxy group of A40926.^[165] SAR studies of teicoplanin indicated an improvement in activity by derivatisation of its carboxy group, and improved activity against VanA-type enterococci is expected to occur when removing the acetylglucosamine.^[48] The most active compound in the series was actually the 6 β -decarboxy-6 β -hydroxymethyl amide of A40926,^[165] but development of this compound was halted because of its poor tolerability in animals (causing adverse effects typical of histamine release, which have been ascribed to its basic character^[48]). Unfortunately, most data on dalbavancin have been published only as abstracts or as short quotations in review papers. This makes an in-depth comparison of dalbavancin with oritavancin difficult.

4.2.1 Microbiological and Pharmacological Properties

Spectrum of Activity and Resistance

In general (table II), dalbavancin is more potent than vancomycin, teicoplanin and, to some extent, oritavancin against staphylococci (with MICs ranging from <0.03 to 1 mg/L for susceptible and methicillin-resistant strains, respectively) and *S. pyogenes* (MIC in the <0.002–0.06 mg/L range).^[49,53,57] It is also very active against *S. pneumoniae*, with MICs 2- to 4-fold higher than those of oritavancin. It is as active as oritavancin or teicoplanin against vancomycin-susceptible enterococci but does not offer the same advantage as oritavancin against glycopeptide-resistant strains. Thus, the MIC distribution range towards VanB- and VanA-type strains includes more susceptible strains but the MIC₅₀ values remain only marginally (4-fold) lower than those of teicoplanin.^[49] Therefore, this drug represents a significant improvement except against glycopeptide-resistant enterococci.^[48]

Subpopulations of staphylococci that have acquired low levels of resistance (2- to 4-fold increase in MIC) have been selected upon serial passage at sub-MIC of dalbavancin.^[166] *In vivo* selection of such mutants may, however, be more difficult, since

trough concentrations of dalbavancin are above the current MIC of the targeted microorganisms.

Pharmacodynamic Properties in Relation to the Mode of Action

Like oritavancin, dalbavancin is bactericidal, with MBC/MIC ratios close to 1, even in the presence of 30% serum.^[66] It is synergic with ampicillin, including against VanA-type enterococci.

Pharmacokinetic Profile and Relationship with Pharmacodynamic Properties

The pharmacokinetic profile of dalbavancin is compared with that of the other glycopeptides in table III. Phase I studies have used higher dosages, providing higher AUC values. However, when corrected for the free fraction, the AUC/MIC and C_{\max} /MIC ratios of dalbavancin are in the same order of magnitude as those of teicoplanin. Plasma dalbavancin concentrations still exceed the MBC values for staphylococci 1 week after administration of a single dose of 1000mg but free serum concentrations are close to the MIC.^[66,67]

Models of Infection

The activity of dalbavancin has been assessed in models of infections by glycopeptide-susceptible Gram-positive organisms and by GISA. Preliminary reports indicate a reduction in bacterial load after a single-dose administration in models of penicillin-resistant *S. pneumoniae* pneumonia or of MRSA rat pouch infection.^[48,167] Its activity is superior to that of conventional glycopeptides in models of endocarditis due to MRSA or GISA, and of acute septicaemia caused by staphylococci, streptococci or enterococci.^[66,168]

4.2.2 Clinical Studies

No clinical data have been published so far but phase II/III clinical trials are ongoing to evaluate the efficacy and safety of dalbavancin for the treatment of complicated skin and soft tissue infections caused by Gram-positive organisms or the use of dalbavancin in a once-a-week mode of administration for other Gram-positive hospital infections.^[66] In a first report of a phase II study,^[169] administration of two doses (1000mg on day 1 and 500mg on day 8) was found to be as effective as comparators (7–21 days

with clindamycin, ceftriaxone, vancomycin or cefazolin) for the treatment of deep skin and soft tissue infections caused by methicillin-susceptible *S. aureus* or MRSA.

4.2.3 Safety Profile

Preclinical studies in rats and dogs show that dalbavancin is well tolerated after intravenous bolus administration at doses several times higher than those expected to be used in humans.^[48] Phase I or phase II clinical trials have not reported major adverse effects in the range of concentrations where dalbavancin is effective.^[66,169] No dosage adjustments are necessary in case of mild renal insufficiency.^[170]

5. New Glycopeptides in the Clinics: for Which Indications?

The improved pharmacological properties of new glycopeptides make them very potent agents, with potentially large clinical indications. However, their use should be as restricted as that of conventional glycopeptides if one wishes to avoid the rapid emergence of resistance.

The advantages of oritavancin that should prove to be useful in clinical practice include: (i) its fast rate of bacterial killing, its spectrum of activity, which covers VRE, methicillin-resistant staphylococci and to some extent the GISA strains; (ii) its activity against intracellular forms of enterococci and staphylococci; and (iii) its prolonged half-life. For dalbavancin, the main improvement seems to be its pharmacokinetic profile, which should allow once-weekly administration. Microbiological properties of dalbavancin are less favourable than those of oritavancin, since VRE are not covered.

On the basis of these properties, and also in view of the results from animal models, oritavancin may become a drug of interest for infections caused by VRE, while both oritavancin and dalbavancin could be considered for severe infections by multiresistant organisms, such as methicillin-resistant staphylococci or *S. pneumoniae*. In the latter case, however, other alternatives such as fluoroquinolones (moxifloxacin or gatifloxacin) should be examined first. In some countries resistance to fluoroquinolones has

already reached alarming levels (15% in Southeast Asia [Hong Kong, South Korea^[171]]). There are also situations such as CNS infections in which fluoroquinolones are poorly efficient.^[172,173] Oritavancin and dalbavancin could, therefore, prove very useful in these 'niche' but important indications. Finally, their activity against GISA strains appears limited (table II), which could be a deterrent to recommending them in these situations. However, synergy has been demonstrated for dalbavancin when combined with β -lactam agents.^[57] Encouraging results have also been obtained in animal models,^[168] which should stimulate further testing of this molecule against GISA infections.

Therefore, reasonable indications could include the following: (i) nosocomial septicaemia or infections of deep organs caused by MRSA or VRE (oritavancin only); (ii) severe and recurrent skin and soft tissue infections caused by MRSA (in relation with activity against both extra- and intracellular bacteria [demonstrated so far only for oritavancin]); and (iii) endocarditis due to resistant organisms (but probably in combination with an aminoglycoside for oritavancin against VRE, based on disappointing animal data^[155]). The usefulness of oritavancin in nosocomial CNS infections caused by MRSA, resistant *S. pneumoniae* or VRE may be suggested on the basis of its activity in animal models. No corresponding data exist for dalbavancin, for which penetration in CSF is low. The potential use of oritavancin in humans in this indication needs to be assessed, as well as the influence of corticosteroids on the CNS penetration of these new glycopeptides (influence of corticosteroids was found to be minor for vancomycin).

Globally speaking, clinical studies or additional animal data are needed to confirm the interest of new glycopeptides in these indications, since the proposals made above rely almost exclusively on an analysis of the pharmacological properties of these drugs and on the results obtained in animal models of infection.

6. Conclusion

Oritavancin, and to a lesser extent dalbavancin, offer considerable possibilities for improvement over vancomycin and teicoplanin, essentially in terms of pharmacokinetic and pharmacodynamic properties, as well as of activity against resistant strains (for oritavancin). Whether these properties can translate into improved clinical efficacy awaits completion of clinical studies. Should these molecules soon become approved for clinical use, it is vital that they are used rationally and prudently to protect against the rapid emergence of resistance and extend their useful life.

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