THE MOLECULAR BASIS OF ANTIBIOTIC ACTION

E. F. Gale E. Cundliffe P. E. Reynolds M. H. Richmond M. J. Waring

SECOND EDITION

The Molecular Basis of Antibiotic Action

SECOND EDITION

E. F. GALE, F.R.S.

Department of Biochemistry, University of Cambridge F. CUNDLIFFE

Department of Biochemistry, University of Leicester

P. E. REYNOLDS

Department of Biochemistry, University of Cambridge M. H. RICHMOND, F.R.S.

Department of Bacteriology, University of Bristol

M. J. WARING

Department of Pharmacology, University of Cambridge

A Wiley-Interscience Publication

JOHN WILEY & SONS

LONDON · NEW YORK · SYDNEY · TORONTO

Copyright © 1972, 1981 by John Wiley & Sons Ltd.

All rights reserved.

No part of this book may be reproduced by any means, nor transmitted, nor translated into a machine language without the written permission of the publisher.

British Library Cataloguing in Publication Data:

The molecular basis of antibiotic action.—2nd ed.

- 1. Antibiotics
- 2. Structure-activity relationship (Pharmacology)
- I. Gale, Ernest Frederick
- 615'.329' RM267 80-41380

ISBN 0 471 27915 3

Printed in Great Britain by J. W. Arrowsmith Ltd, Bristol BS3 2NT.

Preface to the First Edition

In the thirty years which have elapsed since the Second World War there have been many great achievements in medicine and the biological sciences, among which one of the most impressive has been the discovery and development of antibiotics. It is commonplace to remark that antibiotics have revolutionized the treatment of infectious disease, and perhaps equally so to point to their contribution as tools for basic biological research. In both these areas, the remarkable specificity of action of antibiotics has been harnessed to meet a need, for antibiotics have indeed fulfilled the role of 'magic bullets' as originally conceived by Ehrlich. In chemotherapy the magic of the bullets is manifested in selective attack against pathogenic microorganisms. In biological research the magic is no less evident in the use of antibiotics as agents endowed with the capacity to interfere with one or a small number of processes in the complex, integrated reactions of cell metabolism. Small wonder that a knowledge of antibiotic action has become indispensable to the training of medical and biological students.

Most of the antibiotics were discovered before any detailed knowledge was available of the biochemistry underlying their effects. Much of the early work was therefore directed towards uncovering the general sites of action. With the birth and rapid growth of molecular biology it was natural that attempts should be made to elucidate the molecular basis of antibiotic selectivity. Already much has been learned, and it is equally clear that many more important advances lie just around the corner. This book represents an attempt to summarize and take stock of the position to date. There is more to this endeavour than mere curiosity or academic exercise, for it is an axiom of our age that deeper understanding of fundamental biological processes must ultimately be applicable in the service of mankind. In this case a part of the aim must be to improve the efficacy of chemotherapy in medical practice, the philosophy being that a thorough understanding of the nature of such selectivity as is shown by agents known at the present time will help to delineate sites and mechanisms through which selectivity may be achieved, and that, armed with this information, new drugs may be developed to combat disease. In short, the study of the molecular basis of antibiotic action must help to pave the way for a rational approach to chemotherapy. In this light we have allowed ourselves to include in a little reflection and speculation at the beginning and end of the book, including topics such as drug resistance which might not a priori be considered relevant to our title. Similarly, we have not slavishly restricted our attention to antibiotics in the strict sense, i.e. substances of natural (microbial) origin, for in several places we felt that the topic could only be satisfactorily treated in context by including discussion of synthetic antimicrobial drugs, and occasionally even substances which are not chemotherapeutic agents at all.

We make no excuses for choosing to write at the present time: we are all too conscious that, if we were to wait a little longer, there would be many more exciting developments to record and the story would be that much more complete. Such will always be the case to some extent, and at the same time the volume of pertinent literature will increase in its relentlessly exponential fashion, requiring a commensurate increase in the number and/or capacity of authors to deal with it. It could fairly be claimed that we are already too late to hope to deal adequately with the relevant material, and to this contention we would be forced to bow. We cannot know everything, and in the course of writing we have become painfully aware of the enormity of the task and our inadequacy to cope with it as we should like. On the other side of the coin, we have been privileged to consult manuscripts and pre-publication material provided by a number of authors, whose interest and help we gratefully acknowledge, and we have occasionally taken the liberty of including unpublished information obtained in our own and other laboratories. In these instances acknowledgment is given in the text. To cope with the potentially enormous number of references worthy of citation we have, in general, adopted the policy of relying on recent reviews to cover the earlier literature and have tried to concentrate on providing references to the most recent and up-to-date original papers. Very often we have reluctantly decided not to give a direct reference to a classic paper when it is treated in reviews. Any errors, omissions, misinterpretations or other infelicities are of course our own responsibility. In attempting to provide a reasonably comprehensive coverage of the field we have, naturally, each written as individuals on those areas in which we 'specialize'. If, as a result, we have failed to achieve uniformity of style and have afforded a ready means of identifying the writers of particular chapters, we shall not complain if we are each held responsible for our own words.

One of our major problems was to decide how much background biochemistry to include. We have not always been consistent but, in general, have assumed knowledge of basic biochemical matters and, where possible, referred the reader to recent reviews for detailed treatment.

Finally, we should like to acknowledge our debt to colleagues in many laboratories, to students, visitors and correspondents who have discussed, argued, criticized and so helped the development of our ideas over the years.

We also want to thank Mrs B. Gill for her invaluable help in assembling and preparing the manuscript.

Cambridge, August 1971

E.F.G. E.C. P.E.R. M.H.R. M.J.W.

Preface to the Second Edition

In the nine years that have elapsed since the first edition was written there have, inevitably, been many advances in our knowledge of the ways in which antibiotics work and a vast increase in the number of papers published on relevant topics. New antibiotics have been isolated and characterized. We have not attempted to produce a comprehensive guide to all the antibiotics that have been described but have concentrated on those which have been studied in sufficient detail to throw light on their action at a molecular level. In general, advances of recent years have not revealed totally new principles of action but have led to a greater depth of understanding of the interplay between antibiotic structure and the properties of the cellular target. Consequently we have retained the same format as that adopted in the first edition but the new developments have meant that, with the exception of Chapter 1 and parts of Chapter 2, the greater part of the book has been rewritten. As before we have included synthetic drugs where consideration of these has aided our understanding of antibiotic action but, generally, not otherwise.

Again we would like to express our gratitude to many colleagues for the advice, criticisms, and discussions they have provided and especially to those who have allowed us to see and quote papers in preparation for publication. We also want to thank Mrs M. Cumpsty for her invaluable assistance in preparing and collecting the manuscript and particularly for rescuing us from a series of crises in the final stages of that preparation.

Cambridge, May 1980

E.F.G. E.C. P.E.R. M.H.R. M.J.W.

Contents

1.	The	conce	pt 1
	A.	Introd	luction
	В.	Arsen	ical Drugs
		B.1.	The development of arsenical drugs
		B.2.	Mode of action
	C.	Sulph	onamide drugs
		C.1.	Discovery of sulphanilamide
		C.2.	Relationship to p-aminobenzoic acid
		C.3.	
		C.4.	Mode of action
		C.5.	Nature of selectivity
	D.	Antin	netabolites
		D.1.	Early work on antimetabolites
		D.2.	Lethal synthesis
	Ref	erence	-
			· · · · · · · · · · · · · · · · · · ·
ii ii			
2.	Bio	chemic	al targets for drug action 23
	A.	Enzyı	mes
		A.1.	Enzyme structure and activity
		A.2.	Action at the active centre; substrate competition 2
		A.3.	Inhibition at the active centre; cofactor inhibition 33
		A.4.	Allosteric inhibition and 'false feedback' inhibitors 3:
		A.5.	Exo-inhibitors: inhibitors acting outside the active centre 3
		A.6.	'Double blockade'
		A.7.	Transition state inhibitors
	В.	Infor	mation transfer
		B.1.	Synthesis of macromolecules
		B.2.	Nucleic acids
		B.3.	Ribosomes
		B.4.	Repression

xiii

xiv			Contents

C.		sion and protection mechanisms	40
	C.1.	Impairment of membranes	40
	C.2.	1	41
D.	Select	tivity of action	42
	D.1.	Bactericidal and bacteriostatic agents	42
	D.2.	Selective advantage based on the absence of a target from	
		the host	43
	D.3.	Selection based on homologous targets having different	
	2.0.	properties	44
	D.4.	Selective action based on permeability differences	45
	D.5.	Other grounds for selective action	46
Dof	erence:		47
Kei	erence	5	4 /
Inh	bitors	of bacterial and fungal cell wall synthesis	49
Α.	Intro	duction	49
Λ.	A.1.	Composition of bacterial cell walls	49
	A.2.		
		Structure of cell wall polymers	51
-	A.3.	Synthesis of peptidoglycan	56
В.		ition of biosynthetic enzymes	63
	B.1.	Fosfomycin (Phosphonomycin, cis-1,2-epoxypropyl-	
		phosphonic acid)	63
	B.2.	D-cycloserine and O-carbamyl-D-serine	65
		B.2.a. Biochemical investigations	65
		B.2.b. Structure/activity relationships	66
		B.2.c. Molecular basis of action	67
		B.2.d. Resistance to D-cycloserine	73
	B.3.	Alafosfalin (Alaphosphin, L-alanyl-L-1-aminoethyl-	
		phosphonic acid)	74
	B.4.	Tunicamycin	
	B.5.	(i) Diumycin, moenomycin, prasinomycin, macarbomy-	
		cin, 11837 RP; (ii) Janiemycin, enduracidin	78
	B.6.	β -Lactam antibiotics—penicillins and cephalosporins .	
	D .o.	B.6.a. Early biochemical investigations	
		B.6.b. Penicillin-sensitive enzymes and penicillin-	12
		binding proteins	83
		B.6.b.i. Penicillin-sensitive enzymes: trans-	03
		peptidases, D,D-carboxypepti-	
		dases, L,D-carboxypeptidases/	
		transpeptidases and endopep-	212
		tidases	83
		B.6.b.ii. The binding of penicillin	86

		B.o.b.111.	Purincation and activities of peni-	
			cillin-binding proteins	87
		B.6.b.iv.	Function of penicillin-binding pro-	
			teins	92
		B.6.b.v.	Identification of lethal target	100
	B.6.c.	Interaction	n of β -lactam antibiotics and peptide	
		substrat	es with penicillin-sensitive enzymes .	103
		B.6.c.i.	Kinetic studies	103
		B.6.c.ii.	Chemical nature of the penicillin and	
			substrate-binding sites and possi-	
			ble relationship of D,D-carboxy-	
			peptidases/transpeptidases to β -	
			lactamases	106
		B.6.c.iii.	Breakdown of β -lactam-enzyme	
			complexes	111
		B.6.c.iv.	Molecular basis of interaction of β -	
			lactams with proteins	112
	B.6.d.	Structure	activity relationships	116
	5.0.0.	B.6.d.i.	Penicillins	116
		B.6.d.ii.	Cephalosporins	122
		B.6.d.iii.	Non-classical β -lactams	129
	B.6.e.		2	130
	2,5,5,7	B.6.e.i.	Modification of target	130
		B.6.e.ii.	The role of β -lactamases	131
		B.6.e.iii.	Penetrability of the outer mem-	
			brane of Gram-negative bacteria.	133
	B.6.f.	How are	bacteria killed by β -lactam antibi-	
	2.0.1.	otics?		134
C.	Antibiotics wh		e with carrier molecules	137
C.	C.1. Bacitra			137
D.			e with substrates	144
υ.			ocetin A and B, Ristomycin A and B,	
			d B, Avoparcin A and B, and Antibio-	
				144
	D.1.a.		on	144
	D.1.b.		living cells	144
	D.1.c.		cell-free systems	145
	D.1.d.		o bacteria, cell walls, and wall pep-	175
	D.1.d.			147
	D.1.e.		n of vancomycin and ristocetin with	17/
	D.1.C.			153
	D.1.f.	2 (5.25)	Is in vivo	133
	D.1.I.		of vancomycin and other antibiotics of	155

xvi Contents

			D.1.g.	Molecular basis of action: the interaction of aglycones with peptides ending in acyl-D-ala-		
				D-ala	. 15	8
	E.	Antibi	iotics affe	ecting fungal cell wall biosynthesis	. 16	1
		E.1.	Structur	e and biosynthesis of fungal cell walls	. 16	1
		E.2.		on of chitin synthesis by Polyoxin D	. 16	2
		E.3.	Inhibitio	on of mannan biosynthesis by tunicamycin	. 16	5
	Refe	erences			. 16	5
4.	Ant	ibiotics	affecting	g the function of the cytoplasmic membrane	17	5
	A.	The m	nembrane	e	. 17	5
		A.1.	Compos	sition	. 17	5
		A.2.	The cyto	oplasmic membrane: molecular structure	. 17	7
		A.3.	The out	er membrane of Gram-negative organisms .	. 18	(
		A.4.	Artificia	al membranes	. 18	1
	B.	Trans	port acro	oss membranes	. 18	2
	C.	Drugs	producii	ng leakage of small molecules from sensitive cells	, ;	
		earl	ly work		. 18	?
	D.	Drugs	causing	g major disorganization of the cytoplasmi	С	
		mer	mbrane			-
		D.1.	Surface	-active agents		
		D.2.	Tyrocid			
		D.3.	Phenols			
		D.4.	Polymy:	xins and octapeptins		
			D.4.a.	Structure and activity		
			D.4.b.	Site of action		
			D.4.c.	Mode of action		
		D.5.		lar basis of action		-
	E.	Antib	iotics bel	lieved to act by the production of aqueous pores i		
		mei	mbranes			5
		E.1.	Gramic	idins		
			E.1.a.	Structure and activity	. 19	4
			E.1.b.	Mode of action		
			E.1.c.	Molecular basis		
		E.2.	Polyene	es		
			E.2.a.	Structure and activity	. 20	1
			E.2.b.	Action on micro-organisms		
			E.2.c.	Sterol requirement		
			E.2.d.	Selectivity		
			E.2.e.	In vitro studies		
			E.2.f.	Structure of the pore	. 20	19

Contents xvii

			E.2.g. Do polyene-sterol pores explain the action of	
			amphotericin?	214
			E.2.h. Filipin	218
		E.3.	Alamethicin, suzukacillin	219
			E.3.a. Structure and activity	219
			E.3.b. Mode of action	220
		E.4.	Monazomycin	225
		E.5.	Comparison of channels	225
	F.	Drugs	s producing specific changes in cation permeability (iono-	
		pho	res)	226
		F.1.	Dinitrophenol	226
		F.2.	Salicylanilides and other disinfectants	227
		F.3.	Valinomycin, enniatins, macrotetralides	229
			F.3.a. Structure and activity	229
			F.3.b. Molecular basis of ionophore activity	234
		F.4.	Ionophores binding mono- and di-valent cations	241
	G.		piotics which inhibit membrane-bound enzymes involved in	
			rgy transfer	242
		G.1.	Direct inhibition of adenosinetriphosphatase (ATPase) .	242
		G.2.	Indirect inhibition of ATPase	245
	Н.		ition of the synthesis of membrane lipids	245
		H.1.		245
		H.2.		246
	I.		omycins	248
	Rei	erences	s	249
5.	Inhi	bitors (of nucleic acid synthesis	258
	Α.	Introd	duction	258
	B.	Agen	ts which interfere with nucleotide metabolism	260
		B.1.	Inhibitors of nucleotide synthesis	261
		B.2.	Inhibitors of nucleotide interconversion	264
		B.3.	Inhibitors of nucleotide utilization	269
		B.4.	Agents which become incorporated into polynucleo-	
			tides	270
	C.	Agent	ts which impair the template function of DNA	273
		C.1.	The classical intercalating drugs: acridines and ethidium.	274
			C.1.a. Binding to DNA	275
			C.1.b. The intercalation model	280
			C.1.c. Interaction with circular DNA	293
			C.1.d. Selective action on plasmids, kinetoplasts, and	
			mitochondria	
			C.1.e. Frameshift mutagenesis	300

xviii	é.	Contents

	C.2.	Other in		06
		C.2.a.	Anthracycline antibiotics	06
		C.2.b.	Ellipticine, miracil D, chloroquine, and tilo-	
				11
	C.3.	Actinor		14
		C.3.a.		16
		C.3.b.	Molecular models	19
	:*	C.3.c.	Inhibition of RNA polymerase	31
	C.4.	Echino	mycin, triostin, and related antibiotics 3	33
	C.5.	Diacrid	lines and other bis-intercalating drugs 3	37
	C.6.	Non-co	valent interaction with DNA	41
		C.6.a.	Chromomycin, mithramycin, and olivomycin . 3	41
		C.6.b.	Netropsin and distamycin	45
		C.6.c.	Aromatic diamidines	49
		C.6.d.		50
		C.6.e.		51
		C.6.f.		52
	C.7.			53
		C.7.a.		53
		C.7.b.		60
		C.7.c.		62
		C.7.d.	Strand-breaking antibiotics: streptonigrin, bleo-	
				64
D.	Agen	ts which	inhibit enzymic processes in nucleic acid syn-	
				70
	D.1.			70
		D.1.a.	1 2	70
		D.1.b.		75
		D.1.c.		76
	D.2.	Inhibito	ors of DNA replication	77
		D.2.a.		77
		D.2.b.		81
		D.2.c.		82
		D.2.d.		84
		D.2.e.		84
Ref	erence	s		86
				-
i				
Ant	ibiotic	inhibito	rs of ribosome function 4	02
A.	Intro	duction		02
	A.1.			02
				06

6.

Contents	ï					T		xi
							1	- Armon

ntent	S		Ÿ	xix
	A.3.	The med	chanism of protein synthesis	407
		A.3.a.		408
		A.3.b.	Polypeptide chain initiation	408
		A.3.c.	Polypeptide chain termination	409
B.	The n	node of a	ction of puromycin	410
	B.1.		nship of the puromycin reaction to protein	
		synth	esis	412
C.	Ribos	omal spe	cificity of antibiotic action	412
	C.1.	Subunit	localization of antibiotic action	414
		C.1.a.		414
		C.1.b.	Competition with radiolabelled antibiotics of	
			known specificity	414
		C.1.c.	Formation of hybrid ribosomes	415
		C.1.d.	Cross-resistance with antibiotics of known	
			specificity	415
		C.1.e.	Supplementation of inhibited systems	415
		C.1.f.	Inhibition of biochemical processes known to	
			involve a particular subunit	416
		C.1.g.	Examination of components of antibiotic-resis-	
			tant ribosomes	416
	C.2.	Direct of	examination of antibiotic binding sites in ribo-	
		some	s	417
		C.2.a.	Use of antibiotic affinity analogues	417
		C.2.b.	Use of the splitting-reconstitution technique	418
		C.2.c.	Reconstitution with components from resistant	
			ribosomes	418
D.	Inhibi	itors of th	ne smaller ribosomal subunit	418
	D.1.	Strepton	mycin	419
		D.1.a.	, , , , , , , , , , , , , , , ,	419
		D.1.b.	Streptomycin exerts a range of effects in cell-free	
			systems	420
		D.1.c.	Localization of the site of action of streptomycin	421
		D.1.d.	Binding of dihydrostreptomycin to ribosomes .	422
		D.1.e.	The ribosomal binding site for streptomycin	422
		D.1.f.	Misreading of the genetic message caused by	422
		D1 -	aminoglycosides in vitro	423
		D.1.g. D.1.h.	Misreading of the genetic message in vivo	425
		D.1.h. D.1.i.	There are three streptomycin-phenotypes	427
			Effects of streptomycin upon protein synthesis.	428
		D.1.j.	Effects of streptomycin upon partial reactions of protein synthesis	429
		D.1.k.	Properties of reconstituted ribosomes lacking	429
		D.1.K.	protein S12	430
			DIOGHIOLE TELLET	

xx Contents

		D.1.l.	Properties of ribosomes exposed to streptomy-	
			cin in vivo and in vitro	430
		D.1.m.	2	
	•		heterozygotes	431
		D.1.n.	How does streptomycin kill cells?	432
	D.2.		glycosides: Neomycin, kanamycin, gentamicin,	
	_		nygromycin B	433
	D.3.		omycin	438
	D.4.		mycin	439
	D.5.	Aurintr	icarboxylic acid	442
	D.6.		ycin	443
		D.6.a.	Binding of pactamycin to ribosomes	444
		D.6.b.		444
		D.6.c.	Effects of pactamycin in bacterial systems	446
	D.7.	Edeine		446
	D.8.	Tetracy	clines	448
		D.8.a.	Resistance to tetracyclines	448
		D.8.b.	Binding of tetracyclines to ribosomes	448
		D.8.c.	The mode of action of tetracyclines	450
		D.8.d.	Effects of tetracyclines on polyribosome	
			metabolism	451
		D.8.e.	Functions of the ribosomal A site	452
		D.8.f.	Structure and function in tetracycline: β -chelo-	
			cardin	452
	D.9.		e and tubulosine	453
			pleurine, tylophorine, and tylocrebrine	454
E.		itors of t	he larger ribosomal subunit	456
	E.1.	Puromy		456
	E.2.	Chlora	nphenicol	460
		E.2.a.	Structural modifications and analogies	460
		E.2.b.	Mode of action of chloramphenicol	462
		E.2.c.	Binding of chloramphenicol to ribosomes	464
		E.2.d.	The ribosomal binding site(s) for chloram-	
			phenicol	465
		E.2.e.	Resistance to chloramphenicol	46
		E.2.f.	Concluding remarks on chloramphenicol	467
	E.3.	Macrol	ides	
		E.3.a.	Binding of macrolides to ribosomes	469
		E.3.b.	Effects of macrolides in vitro	472
		E.3.c.	Effects of macrolides upon polyribosomes	
		E.3.d.	An anomalous effect of chalcomycin	
	E.4.	Erythro	omycin	
			Binding of erythromycin to ribosomes	

Contents xxi

		E.4.b.	Effects of erythromycin on protein synthesis	475
		E.4.c.	Resistance to erythromycin and other macro-	
			lides	477
	E.5.	Lincom	ycin	478
		E.5.a.	Binding of lincomycin to ribosomes	478
		E.5.b.	Mode of action of lincomycin	479
	E.6.	Streptog		480
		E.6.a.	Binding of streptogramins to ribosomes	483
		E.6.b.	Mode of action of streptogramins	484
		E.6.c.	Resistance to streptogramins	485
	E.7.	Sparson	nycin	485
	E.8.		ıycin	488
	E.9.	Fusidic		489
			epton and related antibiotics	492
		E.10.a.	Inhibition of polypeptide chain elongation	494
		E.10.b.		496
		E.10.c.		496
		E.10.d.		496
		E.10.e.		497
		E.10.f.	Resistance to thiostrepton	498
		E.10.g.	Concluding remarks on thiostrepton	499
	E.11.	Viomyc	in and capreomycin	500
			eximide and related glutarimide antibiotics	502
		E.12.a.		503
		E.12.b.	Effects of cycloheximide on nucleic acid	
			synthesis	506
	E.13.	Anisom		507
			hecenes	508
			ds active on the 60S ribosomal subunit	511
		E.15.a.		511
	· ·	E.15.b.		512
		E.15.c.		514
		E.15.d.		514
F.	Misce	llaneous	Pederine	515
	F.1.		in, gougerotin, blasticidin S., actinobolin, and	
		anthe	elmycin	515
	F.2.	Bottron		516
	F.3.	Tenuaz	onic acid	517
	F.4.		ycin	518
	F.5.	Kirrom	ycin	518
	F.6.	Pulvom		519
G.	Catal	ytic inhib	pitors of protein synthesis	520
		ATTENDED TO THE PARTY OF THE PA	eria toxin and PA toxin	520

(XI	1													COL	lienis
		G.2.	Abrin ar	nd ricin .											522
		G.3.	Modecci	n			•		•				•		524
		G.4.	Crotin a	nd curcin					ž.				•	•	525
		G.5.	Pokewee	ed anti-vira	l peptid	le .								1.0	526
		G.6.	Alpha sa	arcin											526
		G.7.		E3 and cloa											527
	Ack	nowled													529
		erences													529
7.	Bac	terial r	esistance	to antibiot	ics										548
	A.	Introd	duction												548
	B.	Bioch	emical ba	asis of resis	tance				•		÷			*	550
		B.1.	Modifica	ation of the	target										550
		B.2.	Reducti	on in the ph	iysiologi	ical in	mpo	rtai	nce	of t	he 1	targ	et		553
		B.3.	Duplica	tion of the	target				•						554
		B.4.	Prevent	ion of acce	SS .										554
		B.5.	Resistar	nce to inact											556
			B.5.a.												556
			B.5.b.	Enzymes	inactiva	ting	by s	ubs	titu	tior	ì				557
	C.	Gene	tic basis o	of resistance	e .										561
		C.1.	Gene m	odification											562
			C.1.a.	Modificat	ion of th	he ge	ne s	spec	ifyi	ing	the	tar	get		562
			C.1.b.	Modificat	ion of g	enes	oth	er t	har	the	ose	spe	cif	y-	
				ing the	target							,			565
		C.2.	Resistar	nce based o											566
		C.3.		location o											566
			C.3.a.	Plasmid v	ersus ch	rom	osoi	mal	loc	atio	n				566
			C.3.b.	Replicon	interact	ions									568
			C.3.c.	The adva	ntage of	apl	asm	id le	oca	tion	fo	ra	res	is-	
					ene .										574
		*:	C.3.d.	Transmiss	sibility o	of pla	ismi	ds						(in	575
			C.3.e.	Structure											577
	D.	Evolu		ntibiotic re											582
		D.1.	Evoluti	on of resist	ance de	term	ina	nts	٠						582
			D.1.a.	β -Lactam											583
			D.1.b.	Chloramp	henicol	acet	tyl t	rans	fer	ases	•	(*)			583
			D.1.c.	Other pr											
					for resi										585
		D.2.	Evoluti	on of plasn											586
	E.	The o		cture											587
	References											590			