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Thalidomide Metabolism and Metabolites

A thesis submitted to the University of Auckland in fulfilment of the requirements for the degree of Doctor of Philosophy

By

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ABSTRACT

Thalidomide, renowned for causing birth defects in the late 1950s when used for the relief of morning sickness, has attracted new interest for the treatment of inflammatory conditions such as erythema nodosum leprosum and human malignancies such as multiple myeloma. Different species have different sensitivities to thalidomide that could be related to differences in its metabolism. In this study, methodologies using liquid chromatographymass spectrometry were developed to identify thalidomide metabolites formed *in vivo* and *in vitro* in liver microsomes from mice, rabbits and humans, firstly to seek explanations for inter-species differences in sensitivity, and secondly to determine whether thalidomide or its metabolite(s) is the active agent.

Four hydrolysis products were detected in plasma and urine samples from multiple myeloma patients (MMPs) on thalidomide therapy, and mice and rabbits after oral administration of thalidomide. Six hydroxylated metabolites were detected in mice and rabbits, but not in plasma and urine from MMPs. In vitro studies confirmed that murine and rabbit liver microsomes catalysed the hydroxylation of thalidomide efficiently, but significant production of hydroxylation of thalidomide was not observed using human liver microsomes. The degree of hydroxylation both in vivo and in vitro was highest in mice and lowest in humans with rabbits in between. It is unlikely that hydroxylated metabolites are responsible for the effects of thalidomide in the treatment of multiple myeloma, since they were not present in quantifiable amounts in patients who were responding to the treatment. The three major hydrolysis products that were detected in patients were compared with thalidomide for their ability to inhibit tube formation in an *in vitro* angiogenesis assay, to inhibit TNF production induced with LPS in human peripheral blood leucocytes, and to modulate DMXAA-induced TNF production and antitumour activity in mice. One of the three, N-(o-carboxybenzoyl)glutamic acid imide (CG) was found to be as active as thalidomide in all the assays at concentrations (1-2 µg/ml) that are achievable in MMPs. Since CG has been shown by other laboratories to be non-teratogenic, the studies in this

thesis indicate that CG would be a more favourable, non-teratogenic approach to cancer therapy compared with thalidomide.

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TABLE OF CONTENTS

ABSTRACT	I
ACKNOWLEDGMENTS	IV
TABLE OF CONTENTS	VI
LIST OF FIGURES	X
LIST OF TABLES	XVI
LIST OF TABLES	XVI
LIST OF ABBREVIATIONS	XVII
CHAPTER 1. GENERAL INTRODUCTION	1
1.1. THE HISTORY OF THALIDOMIDE AND ITS ROLE IN CANCER THERAPY	1
1.2. THALIDOMIDE AS AN ANTI-CANCER AGENT	4
1.2.1. Pre-clinical Studies	4
1.2.2. Clinical Studies	5
1.2.2.1. Thalidomide in Solid Tumours	5
1.2.2.2. Thalidomide in Haematological Malignancies	6
1.3. MECHANISM OF ACTION IN CANCER TREATMENT	10
1.3.1. Anti-angiogenesis	10
1.3.2. Cytokine Modulation	11
1.3.3. Inhibition of Adhesion Molecule Expression	14
1.3.4. Stimulation of Lymphocytes and Natural Killer Cells	15
1.3.5. Induction of Apoptosis	16
1.4. Pharmacokinetics, Metabolism and Metabolites	18
1.4.1. Pharmacokinetics and Pharmacokinetic Interaction with Other Drugs	18
1.4.1.1. Pharmacokinetics	18
1.4.1.2. Pharmacokinetic Interactions with Other Drugs	20
1.4.2. Metabolism and Metabolites	20
1.5. Summary of the Review	23

1.6. Objectives of This Study	24
CHAPTER 2. DETECTION AND IDENTIFICATION OF THA	LIDOMIDE
METABOLITES IN MICE	26
2.1. Introduction	26
2.2. Methods	
2.2.1. Mice and Tumour	
2.2.2. Drug Administration	27
2.2.3. Metabolite Detection Using LC-MS and HPLC	
2.2.3.1. Preparation of Murine Plasma and Urine Samples	
2.2.3.2. LC-MS Analysis	28
2.2.3.3. Resolution of Phthaloylglutamine (PG) and Phthaloy	rlisoglutamine (PiG)
by HPLC	30
2.2.5.8. Thalidomide Glucuronide Identification	31
2.3. Results	31
2.3.1. Detection of Metabolites	31
2.3.2. Identification of Metabolites	32
2.4. DISCUSSION	44
CHAPTER 3. IDENTIFICATION OF THALIDOMIDE META	ROLITES IN
MULTIPLE MYELOMA PATIENTS	
3.1. Introduction	
3.2. METHODS	
3.2.1. Preparation of Urine and Plasma Samples	
3.2.2. Metabolite Detection and Identification	
3.3. RESULTS	
3.3.1. Detection and Identification of Metabolites in MMPs	
3.3.2. Intra-patient Metabolite Detection Study	
3.4. DISCUSSION	

4.1. Introduction	58
4.2. Methods	59
4.2.1. Murine Studies	59
4.2.2. Rabbit Studies	59
4.2.3. Clinical Studies	60
4.2.4. Metabolite Detection and Identification	60
4.3. Results	61
4.3.1. Thalidomide Metabolite Profile in Mice	61
4.3.2. Thalidomide metabolites in rabbits	66
4.3.3. Thalidomide Metabolites in Patients	66
4.4. DISCUSSION	73
CHAPTER 5. IN VITRO METABOLISM OF THALIDOMIDE IN	MURINE,
RABBIT AND HUMAN LIVER MICROSOMES	
5.1. Introduction	75
5.2. Methods	76
5.2.1. Liver Microsome Preparation	
5.2.2. Bicinchoninic Acid (BCA) Protein Assay	
5.2.3. In Vitro Metabolism	<i>77</i>
5.2.4. Detection of Metabolites Formed in vitro	<i>77</i>
5.2.5. Assay of 5-OH Th	<i>78</i>
5.3. RESULTS	79
5.3.1. Detection of Metabolites	79
5.3.2. Relative Abundance of Metabolites	84
5.3.3. Rate of Metabolism of Thalidomide to 5-OH Th	84
5.4. DISCUSSION	96
CHAPTER 6. BIOLOGICAL ACTIVITY OF THALIDOMIDE'S 1	HYDROLYSIS
METABOLITES	99
6.1. Introduction	99
6.2. Methods	100
6.2.1. Tumour Growth Delay Determinations	100

6.2.2. Modulation of TNF Production in Mice	101
6.2.3. Modulation of TNF Production In vitro	101
6.2.4. TNF Assay	102
6.2.5. Inhibition of Tube Formation In vitro	102
6.2.6. Cytotoxicity Assay	103
6.2.7. Stability and Plasma Concentrations of CG	103
6.2.7.1. Calibration Curves and Quality Controls	103
6.2.7.2. Determination of CG Stability in vitro	104
6.2.7.3. Calculation of Plasma C_{max} , T_{max} , AUC and $t_{1/2}$ of CG in MMPs	104
6.3. Results	105
6.3.1. Potentiation of Anti-tumour Activity of DMXAA in Mice by Thalidomide,	PG,
PiG and CG	105
6.3.2. Effects of Thalidomide, PG, PiG and CG on DMXAA-Induced TNF Prod	luction
in Mice	108
6.3.3. Effects of Thalidomide, PG, PiG and CG on LPS-induced TNF Production	on by
HPBL in Culture	108
6.3.4. Inhibition of Tube Formation in Matrigel	112
6.3.5. Stability of CG at Different pHs	112
6.3.6. Plasma concentrations of CG in MMPs	117
6.4. DISCUSSION	119
CHAPTER 7. GENERAL DISCUSSION	121
7.1. Inter-Species Differences in Thalidomide Metabolism	121
7.2. THE ACTIVE AGENT IN THALIDOMIDE THERAPY	122
7.3. DEVELOPMENT OF CG AS A CLINICAL AGENT	123
APPENDICES	126
APPENDIX 1. CHEMICALS AND REAGENTS	126
APPENDIX 2. PUBLICATIONS DERIVED FROM THIS THESIS	127
REFERENCES	128

LIST OF FIGURES

Figure 1.1 Chemical structures of racemic thalidomide and its stereoisomers
Figure 1.2 A model for the role of TNF in pathophysiology of multiple myeloma
(MM). TNF secreted from MM cells induces modest proliferation, as well as
MEK/MAPK and NF-κB activation, in MM cells. It also augments IL-6 secretion, as
well as activates MEK/MAPK and NF-κB, in BMSCs. Importantly, TNF upregulates
expression of CD49d (VLA-4), CD11a (LFA-1), and Muc-1 on MM·1S cells, as well
as CD54 (ICAM-1) and CD106 (VCAM-1) on BMSCs, which is mediated via NF-κB
activation. Adapted from Hideshima et al., 2001b.
Figure 1.3 Possible role of thalidomide on multiple myeloma (MM) cells' and
BMSCs' microenvironment in vivo. (A) Thalidomide directly inhibits myeloma cell
growth. (B) Thalidomide inhibits MM cell adhesion to BMSCs. (C) thalidomide
blocks IL-6, TNF and IL1β secretion from BMSCs. (D) Thalidomide blocks the
ability of VEGF and bFGF to stimulate neovascularisation of bone marrow. (E)
Thalidomide induces IL-2 and IFN-γ secretion from T-cells. Adapted from
Richardson et al., 2002.
Figure 1.4 Hydrolysis pathway of thalidomide (Adapted from Schumacher et al.,
1965b)
Figure 2.1 UV-detected chromatograms of urine samples from mice without treatment
(dotted lines) and up to 4 h following oral administration of thalidomide (Thal) (50
mg/kg, solid lines).
Figure 2.2 (A) Total ion MS-detected (Signal 1) chromatogram of urine from mice
without treatment (dotted line) and up to 4 h following oral administration of Thal (50
mg/kg, solid line). (B) Mass spectrum of Peak 6 using negative ion-scan mode
showing an [M-H]- mass of 273 amu. (C) Mass spectrum at the retention time
corresponding to Peak 6 in untreated mouse urine
Figure 2.3 (A) Negative SIM mode (Signal 3) MS-detected chromatogram of urine
from mice without treatment (dotted line) and up to 4 h following no. of Thal (50)

mg/kg, solid line). (B) Mass spectrum using negative single-ion monitoring mode
Peak 6 showing a [M-H] response of 273 amu. Note: Peaks 5 & 7 also corresponde
to [M-H] of 273 amu, while Peaks 1 & 4 corresponded to [M-H] of 275 amu, Peak
corresponded to [M-H] of 291 amu and Peak 3 corresponded to [M-H] of 449 am
(spectrum not shown).
Figure 2.4 LC-MS chromatograms of urine samples from Colon 38 tumour-bearing
mice up to 4 h following oral administration of Thal (50 mg/kg). (A) UV-detected
chromatogram, and (B) SIM mode (Signal 3) MS-detected chromatogram 3
Figure 2.5 UV spectra of metabolite peaks (dotted lines) compared with UV spectra of
corresponding authentic standards (solid lines). (A) Peak 1 and CG. (B) Peak 5 ar
cis-5'-OH Th. (C) Peak 6 and trans-5'-OH Th. (D) Peak 7 and 5-OH Th
Figure 2.6 (A) UV-detected chromatogram of Peak 3 following treatment with 1
glucuronidase (solid line) and without treatment (dotted line). (B) Mass spectrum
the Peak II formed following β-glucuronidase treatment showing an [M-H] mass of
273 amu corresponding to 5-OH Th. (C) Mass spectrum at the retention time
corresponding to Peak II in the untreated control.
Figure 2.7 HPLC chromatograms using mobile phase containing cetyltrimethy
ammonium bromide and 1-octanesulfonic acid showing complete separation of: (A
PG and PiG authentic standards; and (B) separation of the Peak 4 fraction from mous
urine into two peaks showing the presence of both PG and PiG.
Figure 2.8 Comparison of UV-detected chromatograms of urine from mid
administered Thal p.o. (solid line) or i.p. (dotted line).
Figure 2.9 Comparison of UV-detected chromatograms of urine (solid line) or plasm
(dotted line) from mice given Thal (50 mg/kg) p.o
Figure 2.10 Proposed pathways of biotransformation of thalidomide in mic
Unconfirmed steps or metabolites are shown in dashed lines
Figure 3.1 LC-MS chromatograms of urine from MMP1 on Thal therapy (100 mg/da
solid lines) and from a healthy volunteer (dotted lines) recorded by: (A) UV at 23
nm, (B) MS at negative TIC mode (Signal 1), (C) MS at positive SIM mode (Sign
2), (D) MS at negative SIM mode (Signal 3).

Figure 3.2 UV chromatograms of urine samples of MMPs on thalidomide therapy
(solid lines) and before treatment (dotted line). A-F correspond to Patients 2-7
respectively51
Figure 3.3 HPLC chromatograms using mobile phase containing cetyltrimethyl-
ammonium bromide and 1-octanesulfonic acid showing complete separation of: (A)
PG and PiG authentic standards; and (B) separation of the Peak 4 fraction from
MMPs' urine into two peaks showing the presence of both PG and PiG
Figure 3.4 UV chromatograms of urine samples of Patient 1 collected on three
occasions after Thal therapy. (A) one month, (B) two months, (C) three months 53
Figure 3.5 Comparison of UV-detected chromatograms of urine (solid lines) or plasma
(dotted lines) from (A) mice given Thal (50 mg/kg, p.o.); and (B) patient 1
approximately 15 h after a prior dose of Thal (100 mg/day p.o.).
Figure 4.1 Chromatograms of urine from mice without treatment (dotted lines) and up
to 4 h following oral administration of Thal (2 mg/kg, solid lines) recorded by: (A
UV at 230 nm, (B) MS at negative TIC mode (Signal 1), (C) MS at positive SIM
mode (Signal 2), (D) MS at negative SIM mode (Signal 3)
Figure 4.2 Negative SIM mode (Signal 3) MS-detected chromatograms of mouse
plasma samples collected before (dotted lines) and after (solid lines) p.o. treatment o
Thal (2 mg/kg). (A) 5 min, (B) 30 min, (C) 4 h
Figure 4.3 Negative SIM mode (Signal 3) MS-detected chromatograms of mouse
plasma samples collected before (dotted lines) and after (solid lines) i.v. treatment o
Thal (2 mg/kg). (A) 1 h, (B) 2 h, (C) 4 h
Figure 4.4 Negative SIM mode (Signal 3) MS-detected chromatograms of rabbi
plasma samples collected before (dotted lines) and after p.o. treatment (solid lines)
(A) 30 min, (B) 2h, (C) 6 h
Figure 4.5 Negative SIM mode (Signal 3) MS-detected chromatograms of rabbit urine
samples collected before (dotted lines) and after treatment (solid lines). (A) 3 h after
p.o. administration, (B) 3 h after i.v. injection.
Figure 4.6 Negative SIM mode (Signal 3) MS-detected chromatograms of rabbi
plasma samples collected before (dotted lines) and after i.v. treatment (solid lines)
(A) 30 min, (B) 2h, (C) 4 h

Figure 4.7	Negative SIM mode (Signal 3) MS-detected chromatograms of MMF
plasma	samples collected before (dotted lines) and after treatment (solid lines). (A) 1
h, from	MMP 8, (B) 4 h, from MMP 10, (C) 24 h, from MMP 11
Figure 4.8	Negative SIM mode (Signal 3) MS-detected chromatograms of MMP 12
urine sa	mples collected before (dotted lines) and after treatment (solid lines). (A) 4 h
(B) 8 h,	(C) 24 h
Figure 4.9	Thalidomide metabolism by hydrolysis (arrows with dashed lines) and CYF
hydroxy	lation and UDPG transferase-mediated glucuronidation (arrows with solic
lines) in	mice, rabbits and MMPs. Unconfirmed metabolites are shown in dotted lines
Structur	es in bold are products formed via hydrolysis only. Numbers in brackets
represen	t metabolite peak number in chromatograms
Figure 5.1	LC-MS chromatograms of Thal metabolites following incubation (60 min
37°C) o	f Thal (400 μM) with liver microsomes (solid lines) of (A) human HL18, (B)
rabbits a	and (C) mice, or with boiled liver microsomes (dotted lines). Metabolites were
detected	by SIM mode (Signal 3) of MS as described in methods
Figure 5.2	HPLC chromatograms with UV detection of Thal metabolites following
incubati	on (60 min; 37°C) of Thal (400 µM) with (A) human HL18, (B) rabbits and
(C) mic	e liver microsomes (solid lines), or with boiled liver microsomes (dotted
lines).	
Figure 5.3	LC-MS chromatograms with UV (A) or MS negative SIM (B) detection of
Thal me	tabolites following incubation (60 min; 37° C) of Thal (400 μ M) with 2 mg/m
human I	HL5 liver micrsomes. 83
Figure 5.4	Enzymatic hydrolysis of PiG by rabbit liver microsomal protein in the
presence	e of NADPH (4mM)
Figure 5.5	Concentration of 5-OH Th with different microsomal protein concentrations
after in	cubating 400 µM of thalidomide with mouse, rabbit and human liver
microso	mes for 60 min
Figure 5.6	Concentration of 5-OH Th at different times after incubating 400 μM or
thalidon	nide with 2 mg/ml of mouse, rabbit and human liver microsomes91

Figure 5.7	HPLC chromatograms showing complete separation of cis-5'-OH Th, trans-
5'-OH T	h, 5-OH Th, Phecacetin and Thal using UV detection at (A) 220 nm or (B)
248 nm.	92
Figure 5.8	Formation of 5-OH Th in rabbit and mouse liver microsomes following
incubatio	on with Thal 93
Figure 5.9	Lineweaver-Burk plots of thalidomide 5-hydroxylation by rabbit and mouse
liver mic	rosomes
Figure 5.10	Eadie-Hoftsee plots of thalidomide 5-hydroxylation by rabbit and mouse
liver mic	rosomes
Figure 6.1	Tumour growth delay in mice untreated, or treated with DMXAA or
DMXAA	combined with Thal or hydrolysis products/metabolites of Thal 106
Figure 6.2	Colon 38 tumour volumes 21 days after treatment in mice. "*" represents
significa	nt difference ($p < 0.05$, student's t-test) compared with DMXAA alone
treatment	107
Figure 6.3	TNF levels in (A) serum and (B) tumour tissue of mice untreated, or treated
with D	MXAA alone or DMXAA combined with Thal or hydrolysis
products/	metabolites of Thal. "*" represents significant difference ($p < 0.05$, student's
t-test) con	mpared with DMXAA alone treatment
Figure 6.4	TNF production by HPBL from seven healthy volunteers at different
concentra	ations of LPS
Figure 6.5	The effect of Thal, CG, PG or PiG on LPS-induced TNF production by
HPBL fi	rom healthy human volunteers. (A) average TNF activity, (B) average
percentag	ge of inhibition
Figure 6.6	Effects of Thal, CG, PG and PiG on tube formation of ECV 304 cells in
Matrigel.	Cells were treated with medium only, medium with vehicle only and
indicated	concentrations of drugs. 113
Figure 6.7	Inhibition of the tube formation of ECV 304 cells in Matrigel by Thal and
CG at dif	ferent concentrations. 114
Figure 6.8	HPLC chromatograms showing complete separation of CG, phenacetin and
Thal at w	ravelength of (A) 220 nm or (B) 248 nm

Figure 6.9 Co	G and Thal concentrations in PBS solutions at different pl	H during 24 h of
incubation	n at 37 °C.	116
Figure 6.10	Plasma concentration-time profiles of CG (from samples	of three MMPs)
compare	d with that of thalidomide (redrawn from Chung et al.,	2004a) after the
treatmen	of 200 mg oral dose fo thalidomide	118
Figure 7.1	Proposed analogues of CG.	125

LIST OF TABLES

Table 1.1 List of diseases in which thalidomide has been trialed
Table 1.2 Pharmacokinetic parameters of orally administered (R-, S-)-racemic
thalidomide (unless stated otherwise).
Table 2.1 Metabolite peaks in UV profiles from murine urine following thalidomid
treatment. 3
Cable 3.1 Comparison of metabolites in mouse and MMP urine samples
Table 4.1 Metabolite peaks in mouse urine LC-MS profiles after thalidomide ora
treatment
Cable 5.1 Metabolite formed after incubating thalidomide with mouse and rabbit live
microsomes
Cable 5.2 Comparison of relative levels of hydroxylated metabolites formed following
a 60 min incubation of Thal with liver microsomal protein (2 mg/ml). Metabolite
were determined by mass spectral detection using single ion monitoring. The respons
of each metabolite peak produced by mouse liver microsomes was normalized to 1.8
Cable 5.3 Comparison of relative levels of hydrolysis products formed following a 6
min incubation of Thal with liver microsomal protein (2 mg/ml). Hydrolysis produc
were determined by mass spectral detection using single-ion monitoring. The
response of each peak produced by mouse liver microsomes was normalized to 18
Cable 5.4 Comparison of the total products of hydrolysis and hydroxylation forme
following a 60 min incubation of Thal with liver microsomal protein (2 mg/ml) 8
Sable 6.1 Comparison of PK parameters of CG and Thal in MMPs

LIST OF ABBREVIATIONS

5-OH Th 5-hydroxythalidomide

5'-OH CG 5'-hydroxy-*N*-(*o*-carboxybenzoyl)glutamic acid imide

5'-OH Th 5'-hydroxythalidomide

 α -MEM α -minimal essential medium

 $\begin{array}{ll} \mu g & microgram \\ \mu l & microlitre \\ \mu M & micromolar \\ ACN & acetonitrile \end{array}$

amu atomic molecular unit

APCI atmospheric pressure chemical ionisation

AU arbitrary unit(s)

AUC area under the concentration-time curve

bFGF basic fibroblast growth factor

BMSC bone marrow stromal cell

C_{max} maximal drug concentration following administration

CG N-(o-carboxybenzoyl)glutamic acid imide

CL_{int} intrinsic clearance
Cox cyclooxygenase

Cl/F apparent clearance rate
CV coefficient of variation

CYP cytochrome P450 enzymes

DMSO dimethylsulphoxide

DMXAA 5,6-dimethylxanthenone-4-acetic acid enzyme linked immunosorbent assay

FBS fetal bovine serum

g gramg gravityh hour

HCl hydrochloric acid

HIV human immunodeficiency virus

HPBL human peripheral blood leucocytes

HPCD 2-hydroxypropyl-β-cyclodextrin

HPLC high performance liquid chromatography
HUVEC human umbilical vein endothelial cell
ICAM intercellular cell adhesion molecule

IFN interferon

IGF insulin-like growth factor

IL interleukin

i.p. intraperitoneali.v. intravenous

KCl potassium chloride

K_M Michaelis-Menten constant

kg kilogram

LC-MS liquid chromatography-mass spectrometry

LPS lipopolysaccharide

mAU milli arbitrary unit(s)

mg milligram
min minute
ml milliliter
mM millimolar

MMP multiple myeloma patient

MRI magnetic resonance imaging

MS mass spectrometry

MSD mass spectral detection

MTT 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyl tetrazolium bromide

NADPH β -nicotinamide adenine dinucleotide phosphate reduced form

NF- κB nuclear factor- κB

NMR nuclear magnetic resonance

PBS phosphate buffered saline

PG phthaloylglutamine

PiG phthaloylisoglutamine

p.o. oral

SEM standard error of mean SIM single ion monitoring

 $t_{1/2}$ drug half-life

 T_{max} time when C_{max} is achieved

TCA trichloroacetic acid

Thal thalidomide

TIC total ion current

TNF tumour necrosis factor-α

UDPG uridine diphosphate glucuronide

UDPG-transferase uridine diphosphate glucuronosyl transferase

UV ultraviolet

V velocity of the reaction

V_{max} maximum velocity of the reaction

V/F volume of distribution

v/v volume/volume

VCAM vascular cell adhesion molecule

VEGF vascular endothelial growth factor