IN THE HIGH COURT OF JUSTICE CHANCERY DIVISION PATENTS COURT

HP-2014-000037

BETWEEN:

HOSPIRA UK LIMITED

Claimant

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CUBIST PHARMACEUTICALS INC

Defendant

AMENDED CLAIMS AS REFERRED TO IN STATEMENT OF REASONS IN SUPPORT OF APPLICATION TO CONDITIONALLY AMEND EUROPEAN PATENT (UK) NO. 1 252 179

1. A method to purify daptomycin, wherein daptomycin is selected from the group consisting of essentially pure daptomycin, daptomycin that is at least 98% pure, daptomycin that is substantially free of anhydro-daptomycin and substantially free of beta-isomer of daptomycin, daptomycin, daptomycin, daptomycin, daptomycin that is free of anhydro-daptomycin and substantially free of beta-isomer of daptomycin, daptomycin, daptomycin, daptomycin that is substantially free of impurities 1 to 14 and daptomycin that is essentially free of impurities 1 to 14, wherein impurities 1 to 14 are as follows:

Impurity	Retention	Molecular	Structure
	<u>time</u>	weight	
	(minutes)		
1	7.96	1638	HO ₂ C NH ₂ CONH ₂ CONH ₂ HO ₂ C NH NH NH ₂ NH NH ₂ NH NH ₂ NH NH ₂ NH NH NH NH NH NH NH NH NH N
2	9.11	1638	HO ₂ C NH ₂ HO ₃ C CONH ₂ HO ₄ C CONH ₂
3	11.54	<u>745</u>	HO CONH ₂ CONH ₂ (CH ₂) ₂ CH ₃ HO

Impurity	Retention time (minutes)	Molecular weight	Structure
4	12.28	1624	HO ₂ C HN O NH CO ₂ M HN O HN HN
<u>5</u>	13.10	<u>1618</u>	= .
<u>6</u>	14.43	587	HO CONH2 (CH2)8CH3
7	14.43	1606	HO ₂ C HN HO ₂ C HN HN HO ₂ C

Impurity	Retention time	Molecular	Structure
	1	<u>weight</u>	
<u>8</u>	(minutes) 15.10	1620	
9	17.92	874	= .
10	19.57	1810	
11	19.57	<u>1635</u>	
11		L	=
12	20.93	859	HO NH ₂ CONH ₂ (CH ₂) ₈ CH ₃ HN NH ₂
13	23.11	1602	HOSC HOSC HOSC HOSC HOSC HOSC HOSC HOSC

1	Impurity	Retention	Molecular	Structure
١		time	weight	
I		(minutes)		
	14	24.53	1634	HO ₂ C HN NH CO ₂ H HN HO ₂ C HN HN HO ₂ C HN HN HN HO ₂ C HN HN HN HO ₂ C HN HN HN HN HN HN HN HN HN H

comprising the steps of:

- a) supplying a daptomycin preparation that contains at least 2.5% of a combined amount of anhydro-daptomycin and β-isomer of daptomycin;
- b) binding the daptomycin preparation to an anion exchange resin in the presence of a modified buffer under conditions in which daptomycin binds to the anion exchange resin in a monomeric and non-micellar state, wherein the modified buffer comprises a buffering agent selected from acetate, phosphate, citrate and Tris-HCI and one or more chaotropic agents selected from ammonia, urea, benzoate and ascorbate;
- c) washing the anion exchange resin in the presence of the modified buffer under conditions that elutes anhydro-daptomycin but retains daptomycin;
- d) eluting daptomycin in the presence of the modified buffer under conditions that separate the purified daptomycin from the β-isomer of daptomycin; and
- e) obtaining purified daptomycin.
- The method according to claim 1, further comprising the step of filtering and concentrating the eluted daptomycin.
- 3. A method to purify daptomycin, wherein daptomycin is selected from the group consisting of essentially pure daptomycin, daptomycin that is at least 98% pure, daptomycin that is substantially free of anhydro-daptomycin and substantially free of beta-isomer of daptomycin, daptomycin that is essentially free of anhydro-daptomycin and substantially free of beta-isomer of daptomycin, daptomycin that is free of anhydro-daptomycin and substantially free of beta-isomer of daptomycin, daptomycin, daptomycin that is

substantially free of impurities 1 to 14 and aptomycin that is essentially free of impurities 1 to 14, wherein impurities 1 to 14 are as follows:

Impurity	Retention	Molecular	Structure
	<u>time</u>	weight	
	(minutes)		
1	7.96	1638	HO ₂ C HO ₂ C HO ₃ C HO ₄ C
<u>2</u>	9.11	1638	HO ₂ C NH ₂ HO HO HO HO HO HO HO HO HO H
<u>3</u>	11.54	745	HO CONH2 HO CON

Impurity	Retention	Molecular	Structure
	<u>time</u>	weight	
	(minutes)		
4	12.28	1624	HO ₂ C HO ₂ C
5	13.10	<u>1618</u>	= .
6	14.43	587	HO CONH2 HO CO2H NO
7	14.43	1606	HO ₂ C HO ₂ C HO ₃ C

Impurity	Retention	Molecular	Structure
	<u>time</u>	weight	
	(minutes)	}	
8	15.10	1620	HO ₂ C H _N CO ₂ H CO ₂ H CO ₂ H HO ₂ C H ₃
9	17.92	874	=
10	19.57	1810	=
11	19.57	1635	=
12	20.93	<u>859</u>	HO NH ₂ NH ₂ NH ₂ NH ₂ NH ₂ CONH ₂ NH ₂
13	23.11	1602	HO ₂ C NH CONNIA CONNIA

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Impurity	Retention time (minutes)	Molecular weight	Structure
14	24.53	1634	HO ₂ C HN O HN HO ₂ C HN HN O HN HO ₂ C HN HN HO ₂ C HN HO ₂ C HN HN HO ₂ C HN HN HO ₂ C HN HO ₂ C HN HN HN HO ₂ C HN HN HO ₂ C HN HN HN HO ₂ C HN HN HN HO ₂ C HN HN HN HN HN HO ₂ C HN HN HN HN HN HN HN HN HN H

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comprising the step of:

- a) fermenting Streptomyces reseosporus with a feed of n-decanoic acid to produce daptomycin in a fermentation broth;
- b) clarifying the fermentation broth;
- subjecting the fermentation broth to anion exchange chromatography to obtain an enriched daptomycin preparation;
- d) subjecting the enriched daptomycin preparation to hydrophobic interaction chromatography to obtain a semi-purified daptomycin preparation; and
- e) subjecting the semi-purified daptomycin preparation to modified buffer enhanced anion chromatography, wherein the modified buffer comprises a buffering agent selected from acetate, phosphate, citrate and Tris-HCI and one or more chaotropic agents selected from ammonia urea, benzoate and ascorbate to obtain purified daptomycin.
- 4. The method according to claim 3, wherein the feed of n-decanoic acid in step a) is regulated to achieve a residual concentration of n-decanoic acid of no more than 50 parts per million (ppm) during fermentation; said clarifying in step b) comprises extracting the fermentation broth with a buffer comprising butanol; the anion exchange chromatography in step c) is performed on FP-DA 13 resin; or either or both steps c) or e) comprises the use of a continuous salt gradient or step salt gradient.
- 5. The method according to claim 3, wherein the modified buffer enhanced anion exchange chromatography in step e) comprises the step of:
 - supplying the semi-purified daptomycin preparation form step d) in a buffer appropriate for modified buffer enhanced anion exchange chromatography;
 - ii. binding the daptomycin preparation to an anion exchange resin in the presence of the modified buffer under conditions in which daptomycin binds to the anion exchange resin in a monomeric and non-micellar state;
 - iii. washing the anion exchange resin in the presence of the modified buffer under conditions that elutes anhydro-daptomycin but retains daptomycin; and
 - iv. eluting daptomycin in the presence of the modified buffer under conditions that permit the separation of daptomycin from β-isomer.
- 6. The method according to claim 3, further comprising: the step of anion exchange chromatography prior to step e); or the step of filtering and/or concentrating daptomycin.

- 7. The method according to claim 3, further comprising the step of depyrogenating daptomycin.
- 8. The method according to claim 7, further comprising the step of lyophilizing daptomycin.