

BSc/MSci Course Unit Examination

May Assessment Period 2022

CHE206B Pharmaceutical Chemistry Semester B

Examiners: Dr G. Chianello and Dr A. Fornili

	For examiners only:			
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6	AF		GC	
Total				

Answer all the questions on this paper. The total number of marks available is 80.

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Question 1

Answer all parts.

(a) A series of gastro protectors (2-7) have been prepared. The lead compound 1 is shown below. These analogues were classified as being active or inactive in comparison to 1.

Using the information provided, propose a structure for the pharmacophore present in 1.

Clearly explain your reasoning. [10 marks]

Question 1 continued overleaf

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(b) Consider the three lead compounds **8**, **9** and **10**, which are being investigated for optimisation in fragment-based drug design. Predict which fragment has the highest potential for optimisation. Explain your answer. [6 marks]

HO
$$_{OH}$$
 $_{OH}$ $_$

(c) Lead compound 11 below undergoes extensive metabolism. Draw two analogues of 11 in which metabolism is inhibited. [2 marks]

(d) An optimised analogue of compound 11 is being used as Active Pharmaceutical Ingredient (API) in a tablet. It has been found that the density of the tablet is 0.99 g/mL and the density of the particulate is 1.1 g/mL. What is the porosity of the tablet? Would this tablet be prone to cracking or crumbling? Justify your answer. [2 marks]

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Question 2

Answer all parts.

(a) Variation of a substituent **X** in a series of amide derivatives (12) led to the Hansch equation shown below. Using the Craig plot and MR table provided (Appendix 1 and 2), identify the *three* best substituents in order to maximise biological activity. Justify your choices.

[6 marks]

(b) An optimised analogue of compound 12 is being used as the API in a suspension. Assuming the particle radius of this suspension is 650 nm and the particle density variation is 0.5, how many years would it take for these particles to precipitate at 60 °C in water in a container 0.5 dm tall? Show your calculations. (Use Appendix 3 for viscosity values).

[4 *marks*]

Question 3

The following scheme shows a medicinal chemistry route for the synthesis of compound 13.

(a) The synthesis above has a high Environmental factor (E factor). Identify *two* features of the synthesis above which contribute to the high E factor. Explain your answers. [4 *marks*]

Question 3 continued overleaf

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(b) Identify *two* undesirable solvents used in the synthesis and suggest a preferred alternative. Explain your answer. [2 *marks*]

(c) Identify *two* logistical challenges of the above synthesis. Explain your answer. [4 marks]

Question 4

The structures of four potential β -blockers (14-17) are given below:

- (a) Rank the compounds in order of *increasing* activity as β -blockers. [2 marks]
- (b) For *each* of the four compounds, justify the ranking you have provided in part (a). [8 marks]

Question 5

Consider compounds 18-21.

- (a) Comment on the ability of *each* compound to act as a cholinergic agonist or antagonist. [8 marks]
- (b) For each of the compounds identified as having cholinergic activity in part (a), indicate if they are likely to produce CNS effects. Justify your answer. [2 marks]

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Question 6

Consider compounds 22-25.

MeN
$$MeO$$
 MeO
 MeO

(a) Comment on their ability to act as muscle relaxants.

[8 *marks*]

- (b) For each of the compounds identified as muscle relaxants in part (a), indicate if they are depolarising or non-depolarising. Justify your answer. [4 marks]
- (c) In a drug optimisation study, a docking calculation was run to generate models of the complex between one of the active compounds and the target receptor.
 - (i) The docking calculation returned five possible models, which are listed below together with an estimate of the binding affinity between the compound and the target. Which model(s) do you think should be retained for further validation and why?

Model	Binding affinity (kcal/mol)		
Model A	-8.3		
Model B	20.3		
Model C	0.5		
Model D	-9.2		
Model E	-9.0		

[3 *marks*]

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(ii) A virtual screening calculation was then run against the 3D-pharmacophore based on one of the docking models. The screening returned six analogues with the following Root Mean Square Deviation (RMSD) values from the 3D pharmacophore:

Compound	RMSD (Å)		
A	0.3		
В	1.2		
С	1.3		
D	0.2		
Е	2.4		
F	1.9		

Which compound(s) do you think should be retained for further validation and why?

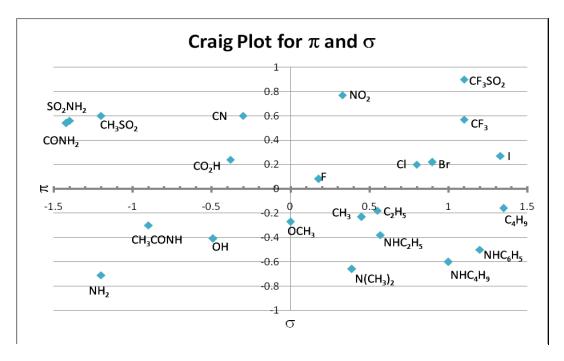
[3 marks]

(iii) In some cases, pharmacophore-guided virtual screening might fail to return any analogue. Provide a possible explanation for this and a possible solution. [2 marks]

End of Paper. An Appendix of 2 pages follows

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Appendix 1



Appendix 2

Subst.	MR	Subst.	MR	Subst.	MR
CF ₃	5.02	F	0.92	CN	6.33
Me	5.65	Н	1.03	CH ₃ SO ₂	14.1
Et	10.3	Cl	6.03	CO ₂ H	6.9
c-Pr	13.5	Br	8.88	CH ₂ OH	7.2
i-Pr	14.9	I	13.94	CONH ₂	9.8
n-Pr	14.9	ОН	2.85	CO ₂ Me	12.9
NHEt	18.6	OMe	7.39	NH ₂	5.4
t-Bu	19.6	OAc	12.47	NO ₂	7.4
Ph	25.4	OEt	12.47	NHMe	10.3
NHPh	26.7	NH-tBu	22.3	NMe ₂	15.6
CH ₃ CONH	15.6	SO ₂ NH ₂	16.9	CF ₃ SO ₂	14.8

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Appendix 3

Temperature	Dynamic viscosity	
[°C]	[Pa s], [N s/m2]	
0.01	0.0017914	
10	0.001306	
20	0.0010016	
25	0.00089	
30	0.0007972	
40	0.0006527	
50	0.0005465	
60	0.000466	
70	0.0004035	
80	0.000354	
90	0.0003142	