

**GCE**

**Chemistry B (Salters)**

Unit **F335**: Chemistry by Design

Advanced GCE

**Mark Scheme for June 2017**

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This mark scheme is published as an aid to teachers and students, to indicate the requirements of the examination. It shows the basis on which marks were awarded by examiners. It does not indicate the details of the discussions which took place at an examiners' meeting before marking commenced.

All examiners are instructed that alternative correct answers and unexpected approaches in candidates' scripts must be given marks that fairly reflect the relevant knowledge and skills demonstrated.

Mark schemes should be read in conjunction with the published question papers and the report on the examination.

OCR will not enter into any discussion or correspondence in connection with this mark scheme.

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## Annotations

| Annotation          | Meaning  |
|---------------------|--|
| <b>DO NOT ALLOW</b> | Answers which are not worthy of credit                     |
| <b>IGNORE</b>       | Statements which are irrelevant                            |
| <b>ALLOW</b>        | Answers that can be accepted                               |
| ( )                 | Words which are not essential to gain credit               |
| —                   | Underlined words must be present in answer to score a mark |
| <b>ECF</b>          | Error carried forward                                      |
| <b>AW</b>           | Alternative wording  |
| <b>ORA</b>          | Or reverse argument  |

| Annotation | Meaning   |
|------------|---|
| ✓          | correct response – there must be one tick for every one mark awarded  |
| ×          | incorrect response – These should not be used for every mark lost; just use them in places where it makes your marking clearer. |
| bod        | benefit of the doubt given. Please give a tick as well  |
| nbod       | benefit of the doubt <b>not</b> given   |
| ECF        | error carried forward   |
| ^          | information omitted   |
| I          | Ignored   |
| SEEN       | to be used on any other page where there is a response but no other annotation  |
| BP         | indicates a blank page that has been checked.   |
| CON        | contradicts a correct response and negates the mark   |
| SF         | to draw attention to the significant figures  |

## Subject-specific Marking Instructions

### INTRODUCTION

Your first task as an Examiner is to become thoroughly familiar with the material on which the examination depends. This material includes:

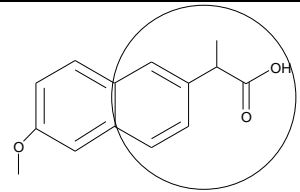
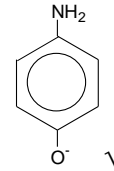
- the specification, especially the assessment objectives
- the question paper
- the mark scheme.

You should ensure that you have copies of these materials.

You should ensure also that you are familiar with the administrative procedures related to the marking process. These are set out in the OCR booklet **Instructions for Examiners**. If you are examining for the first time, please read carefully **Appendix 5 Introduction to Script Marking: Notes for New Examiners**.

Please ask for help or guidance whenever you need it. Your first point of contact is your Team Leader.

| Question | Answer  | Marks | Guidance   |
|----------|---|-------|--|
| 1a       | $C_{13}H_{18}O_2$   | 1     | <b>ALLOW</b> any order of the elements   |
| 1bi      | $CH_3CH(CH_3)CH_2Cl$ ✓<br>1-chloro-2-methylpropane ✓  | 2     | <b>ALLOW</b> any unambiguous structural formula  |
| 1bii     | $AlCl_3$ (catalyst) <b>AND</b> anhydrous ✓  | 1     | <b>IGNORE</b> reaction conditions<br><b>DO NOT ALLOW</b> other reagents  |
| 1ci      | arachidonic acid  | 1     | <b>IGNORE</b> minor spelling errors  |
| 1cii     | binds with/bonds with/fits into ... <u>active site</u> ✓<br>stops (some) substrate binding/fitting/bonding OR<br>competes with substrate/prevents enzyme-substrate<br>complexes forming/blocks active site AW ✓<br>(is reversible/unbinds because) does not bond<br>covalently/forms temporary / weak bonds with active<br>site/can be reversed by increasing substrate<br>concentration (AW) ✓ | 3     |  |
| 1d       | aqueous bromine/bromine solution/ bromine in (named)<br>organic solvent ✓<br><br>turns colourless from yellow/orange /brown <b>OR</b><br>decolourised ✓   | 2     | Not just 'bromine' for mp1 but <b>ALLOW</b> for 2 marks<br>Bromine changing from red/red brown to colourless<br><br>If aqueous bromine/bromine solution <b>ALLOW</b> any<br>colour combinations of orange/yellow/brown<br>If bromine in solvent <b>ALLOW</b> any combination of<br>brown, red or orange.<br><br><b>ALLOW</b> decolourises bromine for 1 mark |
| 1ei      | (isomers that have) same <u>structural formula</u> AND<br>different arrangements in space   | 1     | <b>DO NOT ALLOW</b> atoms bonded in different order<br><b>DO NOT ALLOW</b> 'same molecular formula'  |
| 1eii     | cis <b>AND</b> Z  | 1     |  |
| 1fi      | ether   | 1     | <b>ALLOW</b> alkoxy  |
| 1fii     | part of molecule (AW) ✓<br>where pharmaceutical (AW) action occurs/(that) causes<br>pharmacological effect AW/that causes biological action<br>(AW) ✓   | 2     | <b>DO NOT ALLOW</b> medicinal action   |

|       |   |           |  |
|-------|---|-----------|--|
| 1fiii |    | 1         | Must include all 6 carbons of right hand ring  |
| 1g    |  $\checkmark$ $\text{CH}_3\text{COO}^-$ $\checkmark$ | 2         | <b>ALLOW</b> 1 mark if both structures correct except show OH group<br><b>IGNORE</b> cations |
|       |   | <b>18</b> |  |
|       |   |           |  |

|     |  |   |   |
|-----|--|---|---|
| 2a  | <p>1. (trend because) instantaneous (dipole) induced dipole (bonds/forces) get stronger down the group <math>\checkmark</math></p> <p>2. (trend because) more electrons (down the group)/molecules are larger <b>AW</b> (down the group) <math>\checkmark</math></p> <p>3. water has H-bonds <math>\checkmark</math></p> <p>4. <math>\text{H}_2\text{S}</math> has pd-pd/id-id/ weaker H bonds <math>\checkmark</math></p> <p>5. imb in water (are stronger, so ...) need more energy to break (AW) <b>ORA</b> <math>\checkmark</math></p> | 5 | <p><b>DO NOT ALLOW</b> abbreviations e.g. id-id</p> <p><b>DO NOT ALLOW</b> 'more' id-id down group</p> <p>Mark independently</p> <p><b>QWC</b> only allow marking point 4 if 3 scored</p> <p><b>IGNORE</b> overcome</p> |
| 2b  | ice (molecules are) held in (more) open structure/held further apart (AW) $\checkmark$<br>by hydrogen bonds (in ice) $\checkmark$  | 2 | Air spaces negates mp1  |
| 2ci | <p>O is more electronegative than C (ora) so C=O bonds polarised <math>\checkmark</math></p> <p>Bond polarities cancel / charges balance / the dipoles balance / dipoles cancel out/ centre of +/- charge coincide <math>\checkmark</math></p>   | 2 | <p><b>ALLOW</b> O is more electronegative than C (ora) with bond polarities shown on C=O</p> <p>Must be a comparison (of electronegativity)</p> <p>Mark independently</p>   |

|      |   |           |   |
|------|---|-----------|---|
| 2cii | H-bonds broken in water ✓<br>H-bonds formed between CO <sub>2</sub> and water ✓<br>id-id bonds broken in CO <sub>2</sub> ✓<br>(there is) little <u>energy</u> change when imb are made/broken<br>AW ✓   | 4         | <b>IGNORE</b> pd pd bonds   |
| 2di  | $\underline{\text{Na}^+(\text{g}) + \text{Cl}^-(\text{g})}$ ✓<br>$\underline{\text{NaCl}(\text{s})}$ ✓<br>NaCl(aq)/ <u>Na<sup>+</sup>(aq) + Cl<sup>-</sup>(aq)</u> ✓<br>sum of enthalpy changes of hydration (of ions)<br>= -775 kJ mol <sup>-1</sup> ✓ | 4         | For mp2, energy levels must be in sequence shown/<br><b>IGNORE</b> arrows |
| 2dii | Ion (-)dipole ✓   | 1         |   |
| 2e   | $\text{Mg}_3\text{N}_2 + 6\text{H}_2\text{O} \rightarrow 3\text{Mg}(\text{OH})_2 + 2\text{NH}_3$<br>NH <sub>3</sub> formed ✓ completely correct ✓   | 2         | <b>IGNORE</b> state symbols   |
|      |   | <b>20</b> |   |

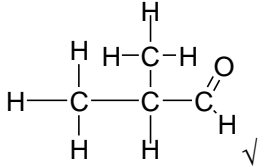
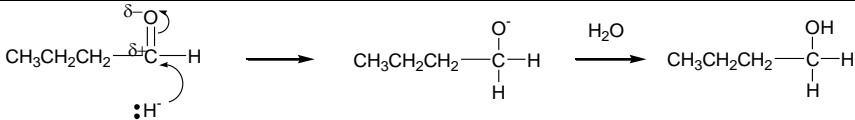
|      |   |   |  |
|------|---|---|--|
| 3ai  | to prevent equilibrium position moving $\checkmark$<br>rates (of back +fwd) slower/stops reaction $\checkmark$  | 2 | <b>ALLOW</b> 'prevents conc of I <sub>2</sub> changing'<br><b>IGNORE</b> quench  |
| 3aii | $[I_2] = 26.3 \times 0.5 \times 0.0687 / 100 \checkmark$<br>$= 9.03 \times 10^{-3} \checkmark$  | 2 | <b>ALLOW</b> 2 or more sf<br>Correct answer on answer line scores 2 without reference to working.<br>1.80...x 10 <sup>-2</sup> (no factor of 2) scores 1 without reference to working.<br>9.03 x 10 <sup>-4</sup> (divides by 1000 not 100) scores 1 without reference to working<br>mp 2 ecf only for a transcription error |
| 3bi  | $2 \times 0.00863 / 0.01726$ (mol dm <sup>-3</sup> ) of HI react (to give 0.00863 mol dm <sup>-3</sup> I <sub>2</sub> ) $\checkmark$<br>(Initial conc of HI) = 0.004 x 10 <b>or</b> 0.04 (mol dm <sup>-3</sup> )<br><b>AND</b> $0.04 - 0.01726 = 0.0227 \checkmark$ | 2 | Method must be shown to score either mark<br><b>ALLOW</b> alternative methods  |
| 3bii | $[H_2] = [I_2]$ <b>OR</b> $[H_2] = 0.00863 \checkmark$<br>$K_c = [H_2] [I_2] / [HI]^2$ <b>OR</b> $[I_2]^2 / [HI]^2$ <b>OR</b> $[H_2]^2 / [HI]^2$ <b>OR</b> numbers substituted $\checkmark$<br>$= 0.14453 \dots \checkmark$<br>$= 0.145$ (3 sf) $\checkmark$        | 4 | First marking point can be inferred from later working<br>0.14453...or 0.144 score 3 without working<br>0.145 scores 4 without working<br><b>ALLOW</b> mp4 for any calculation correctly evaluated and shown to 3sf  |
| 3c   | I/iodine has been oxidised from -1 to 0 $\checkmark$<br>S/sulfur has been reduced from +6 to -2 $\checkmark$  | 2 | Give one mark for correct identification of iodine (oxidised) and sulfur (reduced) without/with wrong oxidation states.<br><b>DO NOT ALLOW</b> I <sub>2</sub> for 'iodine'   |
| 3d   | Endothermic $\checkmark$<br>(because) <u>iodine</u> is formed so equilibrium (position) must have moved to right $\checkmark$   | 2 | <b>IGNORE</b> favouring  |
| 3ei  | $N_2H_4 + 2I_2 \rightarrow N_2 + 4HI$   | 1 | <b>IGNORE</b> state symbols<br><b>ALLOW</b> $N_2H_4 + I_2 \rightarrow N_2 + 2HI + H_2$   |

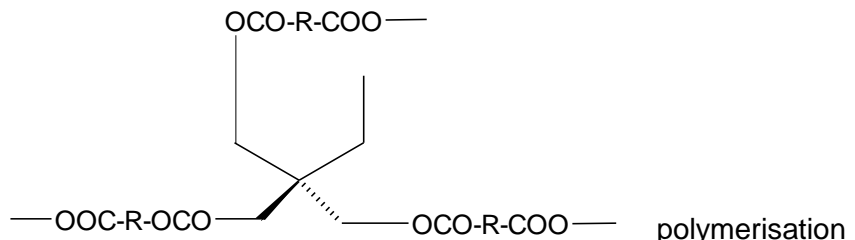


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|-------|--|----|--|
| 3eii  | $\begin{array}{c} \text{H} \cdot \cdot \text{N} \cdot \cdot \text{N} \cdot \cdot \text{H} \\   \quad   \quad   \\ \text{H} \quad \text{H} \end{array}$ <p>5 electrons around each N ✓ completely correct ✓</p>   | 2  |  |
| 3eiii | <p>109 ✓<br/>4 pairs/ groups of electrons/ 4 areas of electron density/<br/>4 regions of <u>negative</u> charge ✓<br/>repel <b>AND</b> get as far away from each other as possible ✓</p> <p>pyramidal ✓</p>  | 4  | <p><b>ALLOW</b> 104 – 110</p> <p>mp3 <b>ALLOW</b> 'minimise repulsion'<br/><b>IGNORE</b> 'repel as much as possible'<br/><b>QWC</b> 'pyramidal' must be spelled correctly to score<br/><b>IGNORE</b> tetrahedral/triangular<br/><b>ALLOW</b> trigonal pyramid spelled correctly<br/>no ecf</p>                       |
| 3eiv  | <p><math>\text{N}_2\text{H}_5^+\text{HSO}_4^- / \text{H}_2\text{NNH}_3^+ \text{HSO}_4^- / \text{H}_3\text{NNH}_3^{2+} \text{SO}_4^{2-}</math></p> <p>Formula ✓<br/>All correct ✓</p>   | 2  | <p><b>ALLOW</b> one mark for <math>\text{N}_2\text{H}_5\text{HSO}_4</math><br/>or <math>\text{N}_2\text{H}_6\text{SO}_4</math> (with or without correct/incorrect charges)</p>   |
| 3f    | <p><math>([\text{HI}] =) 26.25 \times 0.110/25</math> <b>or</b> <math>0.1155</math> ✓<br/><math>V \times 0.1155 = 1000 \times 0.1</math> <b>or</b> <math>V = 1000 \times 0.1/0.1155</math> ✓</p> <p><math>V = 866/ 865.8</math> (<math>\text{cm}^3</math>) ✓</p> | 3  | <p><b>ALLOW</b> 0.116 for 0.1155</p> <p><b>ALLOW</b> 862.068..... if 0.116 used<br/><b>ALLOW</b> answer to 2 or more sf (870)<br/><b>ALLOW</b> ecf from first marking point.<br/><b>ALLOW</b> 865.1 for 3 marks (early rounding mp1)</p> <p>Correct answer on answer line scores 3 without reference to working.</p> |
|       |  | 26 |  |

|       |   |   |   |
|-------|---|---|---|
| 4a    | ester   | 1 |   |
| 4bi   | <u>acid anhydride</u>   | 1 |   |
| 4bii  | mol phenol = $85000/94 = 904(.255\dots)$ ✓<br>mol cpd A = $690000/148 = 466(.216\dots)$ ✓<br>cpd A in excess AND because reacting mols is 2:1 ratio ✓   | 3 | <b>ALLOW</b> calculations in kilomoles but maximum 1 mark for mp1 and mp2 if units omitted and 0.904 AND 0.466<br><br>2:1 ratio can be shown from calculation<br><b>ALLOW</b> ecf   |
| 4biii | $0.81 \times 904/2 \times 318 = 120/116 \text{ kg}$ ✓✓  | 2 | <b>ALLOW</b> ecf from numbers (but not wrong conclusion) in ii<br><b>ALLOW</b> one mark for incorrect evaluation of correct expression<br><b>ALLOW</b> for 1 mark correct evaluation of expression with 0.81 or 2 missing.<br>ie 143.76 scores 1 (missing x 0.81); 232.85 scores 1 (missing /2)<br><b>ALLOW</b> answers to 2 or more sf |
| 4ci   | 1 electron from each carbon ✓<br>in ring(s) over all <u>carbon</u> atoms ✓<br>(rings are) one above and one below ✓   | 3 | <b>IGNORE</b> 'clouds'<br>'rings above and below carbon (atoms)' scores mp2 and 3   |
| 4cii  | 1. <u>electrons</u> raised to higher <u>energy level</u> ✓<br>2. radiation/light/photon/frequencies in electromagnetic spectrum absorbed <b>AND</b> $\Delta E = hv/f$ ✓<br>3. more delocalisation in ion / larger chromophore ✓<br>4. reduces $\Delta E$ so light/visible radiation absorbed/lower frequency radiation absorbed ORA ✓<br>5. (for ion) complementary colour <u>reflected/ transmitted</u><br><b>OR</b> <u>molecule</u> absorbs in uv ✓ | 5 | mp2 <b>ALLOW</b> $E = hv/f$ if 'energy gap' clearly indicated<br><br><b>QWC</b> only award 4. if 3. scored<br><br>mp5 <b>ALLOW</b> 'complimentary' or 'colour(s)/frequency(ies) not absorbed' for 'complementary'.<br><b>DO NOT ALLOW</b> mention of 'dropping back down' or 'emit' in mp5  |

|       |  |    |  |
|-------|--|----|--|
| 4d    | $pK_a = pH/9.3 \checkmark$ $K_a = 5.0 \times 10^{-10} \checkmark$  | 2  | <b>ALLOW</b> 1sf<br>Correct answer on answer line scores 2 without reference to working.   |
| 4ei   | 0.1 mol of HCl and NH <sub>3</sub> react to form 0.1 mol NH <sub>4</sub> Cl $\checkmark$<br>double volume so 0.05 mol dm <sup>3</sup> $\checkmark$       | 2  | 2 <sup>nd</sup> mark dependent on first  |
| 4eii  | $[NH_3] [H^+]/[NH_4^+]$  | 1  | Must be square brackets  |
| 4eiii | $[H^+] = \sqrt{(5.6 \times 10^{-10} \times 0.05)}$ or 5.3 (5.29...) $\times 10^{-6} \checkmark$<br>pH = 5.3 (5.276....) $\checkmark$                     | 2  | <b>ALLOW</b> first mark if 'H <sup>+</sup> =' stated.<br><b>ALLOW</b> ecf for second mark if [H <sup>+</sup> ] or 'H <sup>+</sup> ' quoted incorrectly as 10 <sup>-3</sup> or smaller and correctly evaluated. |
| 4eiv  | pink colour of phth only develops after/around pH 9.3 $\checkmark$<br>indicator needs to change colour around a pH of 5 for this titration. $\checkmark$ | 2  | <b>ALLOW</b> ecf from 4eiii provided an acidic pH<br><b>ALLOW</b> +/- 0.5 pH units   |
| 4fi   | benzene sulfonic acid  | 1  | <b>ALLOW</b> 'sulphonic'   |
| 4fii  | conc/c sulfuric acid/H <sub>2</sub> SO <sub>4</sub> $\checkmark$<br>heat/reflux $\checkmark$   | 2  |  |
| 4fiii | electrophilic substitution $\checkmark$  | 1  |  |
| 4fiv  | no reaction $\checkmark$<br>pink/purple colour $\checkmark$  | 2  | If starting colour given, must be orange/yellow or colourless.   |
|       |  | 30 |  |

|       |   |   |  |
|-------|---|---|--|
| 5ai   | increasing pressure increases yield ( <b>ORA</b> ) ✓<br>fewer/less moles/molecules/particles on right/products <b>ORA</b> ✓   | 2 | Mark independently<br><b>IGNORE</b> 'fewer/less products'  |
| 5aai  | increasing temperature increases rate ✓<br>more frequent (AW) <u>collisions</u> with energy greater than $E_a$ /activation enthalpy /more frequent <u>successful collisions</u> ✓   | 2 |  |
| 5aiii | disorder <b>or</b> number of ways of arranging particles/molecules/quanta of energy ✓<br>-/negative since fewer/less moles/molecules/particles on right/products ✓  | 2 | <b>IGNORE</b> 'fewer/less products'  |
| 5bi   | butanal   | 1 |  |
| 5bii  | <br>(2-)methylpropanal ✓   | 2 |  |
| 5biii | CH <sub>3</sub> CH <sub>2</sub> CH <sub>2</sub> COOH <b>AND</b> butanoic acid   | 1 | <b>ALLOW</b> any unambiguous structure<br><b>ALLOW</b> molecular formula C <sub>4</sub> H <sub>8</sub> O <sub>2</sub>  |
| 5ci   | <br>attack of H <sup>-</sup> and curly arrow on aldehyde ✓<br>intermediate ✓<br>attack by water to give product ✓<br>partial charges correct on C=O <b>AND</b> lone pair on H <sup>-</sup> ✓ | 4 | Mark independently<br>Curly arrow should start either from lone pair or negative charge on H <sup>-</sup> and end pointing at the C atom or the bond about to form between C and H<br><b>IGNORE</b> curly arrows and lone pairs on water |

|      |  |    |  |
|------|--|----|--|
| 5cii | $\text{CH}_3\text{CH}_2\text{CH}_2\text{CH}=\text{NH}$ / $\text{CH}_3\text{CH}_2\text{CH}=\text{CHNH}_2$ ✓✓  | 2  | <b>ALLOW</b> any unambiguous representation.<br><br><i>for one mark, allow:</i><br>any structure that shows attack by $\text{NH}_3$ on the right-hand carbon and the correct molecular formula eg $\text{C}_3\text{H}_7\text{CHNH}$ or $\text{CH}_3\text{CH}=\text{CHCH}_2\text{NH}_2$ |
| 5di  | heat/reflux with conc/c sulfuric acid/ $\text{H}_2\text{SO}_4$   | 1  | <b>ALLOW</b> conc HCl  |
| 5dii | $\text{CH}_3\text{CH}_2\text{CH}_2\text{CH}_2\text{OH} + \text{CH}_3\text{COOH} \rightarrow \text{CH}_3\text{COOCH}_2\text{CH}_2\text{CH}_2\text{CH}_3 + \text{H}_2\text{O}$<br>correct formula of ester ✓<br>completely correct equation ✓  | 2  | <b>ALLOW</b> any unambiguous structures<br><b>ALLOW</b> 1 mark for use of $\text{C}_4\text{H}_9$ for $\text{CH}_3\text{CH}_2\text{CH}_2\text{CH}_2$  |
| 5e   | IR – ester (not acid), as no O–H > 3000 / has C=O ester at 1750 ✓<br><br>ester: $\text{CH}_3\text{COOCH}_2\text{CH}(\text{CH}_3)_2$<br>$(\text{CH}_3)_2\text{CH} -$ ✓<br>Remainder correct ✓<br><br><i>two from</i> ✓✓<br>• Peak at 2.0* indicates CHCO<br>• 6 protons at 0.9* indicates 2 $\text{CH}_3$ <b>or</b> 4 proton environments<br>• 2 proton doublet at 3.9* indicates $\text{CH}_2\text{O}$ next to CH<br>* allow $\pm 0.1$ | 5  | <b>ALLOW</b> range 2500 to 3200 for O-H<br><br><b>DO NOT ALLOW</b> ranges for chemical shift   |
| 5f   | <br>occurs at these bonds<br>3 ester links to TMP ✓<br>indication where polymerisation can continue ✓   | 2  | 2 <sup>nd</sup> mark depends on first being correct<br><br><b>ALLOW</b> indication of further polymerisation from COOH   |
|      |  | 26 |  |

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