Version 1

**General Certificate of Education (A-level)
January 2011**

Chemistry   **CHEM4**

(Specification 2420)

Unit 4: Kinetics, Equilibria and Organic Chemistry

Post-Standardisation

Mark Scheme

Mark schemes are prepared by the Principal Examiner and considered, together with the relevant questions, by a panel of subject teachers. This mark scheme includes any amendments made at the standardisation events which all examiners participate in and is the scheme which was used by them in this examination. The standardisation process ensures that the mark scheme covers the candidates' responses to questions and that every examiner understands and applies it in the same correct way. As preparation for standardisation each examiner analyses a number of candidates' scripts: alternative answers not already covered by the mark scheme are discussed and legislated for. If, after the standardisation process, examiners encounter unusual answers which have not been raised they are required to refer these to the Principal Examiner.

It must be stressed that a mark scheme is a working document, in many cases further developed and expanded on the basis of candidates' reactions to a particular paper. Assumptions about future mark schemes on the basis of one year's document should be avoided; whilst the guiding principles of assessment remain constant, details will change, depending on the content of a particular examination paper.

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Mark Scheme – General Certificate of Education (A-level) Chemistry – Unit 4: Kinetics, Equilibria and Organic Chemistry – January 2011

Question	Marking Guidance		Mark	Comments
1(a)(i)	propyl methanoate		1	must be correct spelling
1(a)(ii)	rate = k[X][OH⁻]		1	allow $HCOOCH_2CH_2CH_3$ (or close) for X allow ( ) but penalise missing minus
1(a)(iii)	$k = \dfrac{8.5 \times 10^{-5}}{(0.024)(0.035)}$	In (a)(iii), if wrong orders allow	1	mark is for insertion of numbers in correct expression for k If expression for k is upside down, only score units conseq to their expression
	$= 0.10(12)$ 2sf minimum	1 for conseq answer	1	
	$mol^{-1}\,dm^3\,s^{-1}$	1 for conseq units	1	any order
1(a)(iv)	$2.1(3) \times 10^{-5}$		1	or $2.1(2) \times 10^{-5}$ ignore units allow 2 sf **NB If wrong check the orders in part (a)(iii) and allow (a)(iv) if conseq to wrong k** **See * below**

1(a)(v)	1.3×10^{-4} (1.28×10^{-4})	1	allow (1.26×10^{-4}) to (1.3×10^{-4}) ignore units
			allow 2 sf
			NB If wrong check the orders in part (a)(iii) and allow (a)(iv) if conseq to wrong k
			See ** below
	For example, if orders given are 1st in X and second in OH⁻ [The mark in a(ii) and also first mark in a(iii) have already been lost] So allow mark *** in (iv) for rate = their k ×(0.012)(0.0175)² = their k ×(3.7 × 10⁻⁶) (allow answer to 2sf) ** in (v) for rate = their k × (0.012)(0.105)² = their k ×(1.32 × 10⁻⁴) (allow answer to 2sf) **The numbers will of course vary for different orders.**		

| 1(a)(vi) | Lowered | 1 | if wrong, no further mark |
| | fewer particles/collisions have energy >E_a
OR
fewer have sufficient (activation) energy (to react) | 1 | not just fewer successful collisions |

| 1(b) | Step 2 | 1 | if wrong, no further mark |
| | (this step with previous) involves one mol/molecule/particle A and two Bs
or 1:2 ratio or same amounts (of reactants) as in rate equation | 1 | |

Question	Marking Guidance	Mark	Comments
2(a)(i)	- log[H$^+$] or log 1/[H$^+$]	1	penalise missing square brackets here only
2(a)(ii)	0.81	1	2dp required, no other answer allowed
2(a)(iii)	**M1** mol H$^+$ = 1.54 ×10^{-3}	1	if wrong no further mark if 1.5 ×10^{-3} allow M1 but not M2 for 2.82
	M2 pH = 2.81	1	allow more than 2dp but not fewer
2(b)	**M1** [H$^+$] = 3.31 ×10^{-3}	1	
	M2 K_a = $\dfrac{[H^+][X^-]}{[HX]}$ or $\dfrac{[H^+]^2}{[HX]}$ or using numbers	1	do not penalise ( ) or one or more missing [ ]
	M3 [HX] = $\dfrac{[H^+]^2}{K_a}$ = $\dfrac{(3.31\times10^{-3})^2}{4.83\times10^{-5}}$	1	allow conseq on their [H$^+$]^2/(4.83 × 10^{-5}) (AE) if upside down, no further marks after M2
	M4 [HX] = 0.227	1	allow 0.225 – 0.23
2(c)	**M1** extra/added OH$^-$ removed by reaction with H$^+$ or the acid	1	
	M2 correct discussion of equm shift i.e. HX $\rightleftharpoons$ H$^+$ + X$^-$ moves to right **OR** ratio $\dfrac{[HX]}{[X^-]}$ remains almost constant	1	

2(d)(i)	**M1** mol HY = $(50 \times 10^{-3}) \times 0.428 = 0.0214$	**OR** $[Y^-] = .0236 \times \dfrac{1000}{50} = 0.472$	1	mark for answer
	M2 $[H^+] = 1.35 \times 10^{-5} \times \dfrac{0.0214}{0.0236}$ **OR** $1.35 \times 10^{-5} = [H^+] \times \dfrac{0.0236}{0.0214}$	**OR** $[H^+] = 1.35 \times 10^{-5} \times \dfrac{0.428}{0.472}$ **OR** $1.35 \times 10^{-5} = [H^+] \times \dfrac{0.472}{0.428}$	1	must be numbers not just rearrangement of Ka expression If either HY value or Y⁻ value wrong, (apart from AE -1) lose M2 and M3
	M3 $[H^+] = 1.22 \times 10^{-5}$		1	mark for answer
	M4 pH = 4.91		1	allow more than 2dp but not fewer **allow M4 for correct pH calculation using their [H⁺] (this applies in 2(d)(i) only)**
	If Henderson Hasselbalch equation used:			**If Henderson Hasselbalch equation used:**
	M1 mol HY = $(50 \times 10^{-3}) \times 0.428$ = 0.0214	**OR** $[Y^-] = .0236 \times \dfrac{1000}{50} = 0.472$	1	mark for answer
	M2 pKa = 4.87		1	
	M3 $\log(\dfrac{0.0214}{0.0236}) = -0.043$	$\log(\dfrac{0.428}{0.472}) = -0.043$	1	If either HY value or Y⁻ value wrong, (apart from AE-1) lose M3 and M4
	M4 pH = 4.87 − (− 0.043) = 4.91		1	allow more than 2dp but not fewer

2(d)(ii)			Can score full marks for correct consequential use of their HY and Y⁻ values from d(i)
	M1 Mol HY after adding NaOH = $0.0214 - 5.0 \times 10^{-4} = 0.0209$	1	AE in subtraction loses just M1 If wrong initial mol HY (i.e. not conseq to part d(i)) or no subtraction or subtraction of wrong amount, lose M1 and M3
	M2 Mol Y⁻ after adding NaOH = $0.0236 + 5.0 \times 10^{-4} = 0.0241$	1	AE in addition loses just M2 If wrong mol Y⁻ (i.e. not conseq to part d(i)) or no addition or addition of wrong amount lose M2 and next mark gained
	M3 $[H^+] = 1.35 \times 10^{-5} \times \dfrac{0.0209}{0.0241}$ $(= 1.17 \times 10^{-5})$ if convert to concentrations $[H^+] = 1.35 \times 10^{-5} \times \dfrac{0.418}{0.482}$ $(= 1.17 \times 10^{-5})$	1	if HY/Y⁻ upside down, no further marks
	M4 pH = 4.93	1	allow more than 2dp but not fewer NOT allow M4 for correct pH calculation using their $[H^+]$ (this allowance applies in 2(d)(i) only)

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	If Henderson Hasselbalch equation used:		If Henderson Hasselbalch equation used:
			Can score full marks for correct consequential use of their HY and Y^- values from d(i)
M1	Mol HY after adding NaOH = $0.0214 - 5.0 \times 10^{-4} = 0.0209$	1	AE in subtraction loses just M1 If wrong initial mol HY (i.e. not conseq to part d(i)) or no subtraction or subtraction of wrong amount lose M1 and M3
M2	Mol Y^- after adding NaOH = $0.0236 + 5.0 \times 10^{-4} = 0.0241$	1	AE in addition loses just M2 If wrong mol Y^- (i.e. not conseq to part d(i)) or no addition or addition of wrong amount lose M2 and next mark gained
M3	$\log \left(\dfrac{0.0209}{0.0241} \right) = -0.062$	1	if HY/Y^- upside down, no further marks
M4	pH = $4.87 - (-0.062) = 4.93$	1	allow more than 2dp but not fewer

Question	Marking Guidance	Mark	Comments
3(a)	mol CH_3OH $= 0.07(0)$	1	
	mol H_2 $= 0.24(0)$	1	
3(b)(i)	$\dfrac{[CH_3OH]}{[CO][H_2]^2}$ or $\dfrac{(0.082/1.5)}{(0.210/1.5)(0.275/1.5)^2}$	1	allow ( ) but expression using formulae must have brackets alternative expression using numbers must include volumes
3(b)(ii)	**M1** divides by vol	1	Mark independently from (b)(i) any AE is –1 if volume missed, can score only M3 and M4
	M2 $\dfrac{(0.082/1.5)}{(0.210/1.5)(0.275/1.5)^2}$ $\left(= \dfrac{(0.05467)}{(0.14)(0.1833)^2} \right)$	1	mark is for correct insertion of correct numbers in correct Kc expression in b(ii) If Kc expression wrong, can only score M1 & M4 If numbers rounded, allow M2 but check range for M3
	M3 11.6 or 11.7	1	mark for answer above 11.7 up to 12.2 scores 2 for M1 and M2 if vol missed, can score M3 for 5.16 (allow range 4.88 to 5.21)
	M4 $mol^{-2} dm^6$	1	Units conseq to their Kc in (b)(ii)
3(b)(iii)	no effect or no change or none	1	

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3(c)	**M1** T_1		1	if wrong - no further marks
	M2	(forward) reaction is exothermic **OR** gives out heat	<u>backward</u> reaction is endothermic	1	only award M3 if M2 is correct
	M3	shifts to RHS <u>to replace lost heat</u> **OR** <u>to increase the temperature</u> **OR** <u>to oppose fall in temp</u>	<u>backward reaction</u> takes in heat **OR** to lower the temperature	1	not just to oppose the change

3(d)	fossil fuels used **OR** CO_2 H_2O produced/given off/formed which are <u>greenhouse gases</u> **OR** SO_2 produced/given off/formed which causes acid rain **OR** Carbon produced/given off/formed causes global dimming	1	not allow electricity is expensive ignore just global warming ignore energy or hazard discussion

3(e)	$C_{17}H_{35}COOCH_3$ or $C_{17}H_{31}COOCH_3$ or $C_{17}H_{29}COOCH_3$ **OR** $CH_3OOCC_{17}H_{35}$ or $CH_3OOCC_{17}H_{31}$ or $CH_3OOCC_{17}H_{29}$	1	

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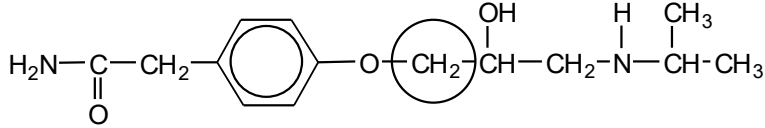
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Question	Marking Guidance	Mark	Comments
4(a)	3-hydroxypropanoic acid	1	allow 3-hydroxypropionic acid must be correct spelling
4(b)(i)	must show trailing bonds or can start at any point in the sequence, e.g. 	1	not allow dimer allow $-O-CH_2CH_2COOCH_2CH_2CO-$ or $-CH_2CH_2COOCH_2CH_2COO-$ ignore ( ) or n NB answer has a total of 6 carbons and 4 oxygens
4(b)(ii)	condensation (polymerisation)	1	Allow close spelling
4(c)(i)	C=C or carbon-carbon <u>double</u> bond	1	
4(c)(ii)		1	must show **ALL** bonds including O–H
4(c)(iii)	must show trailing bonds 	1	allow polyalkene conseq on their c(ii) ignore n

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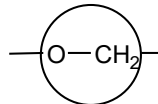
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4(d)		1	allow NH_3^+— allow COO^-
4(e)(i)		1	In 4(e), do not penalise a slip in the number of carbons in the $-CH_2CH_2-$ chain, but all must be bonded correctly NB two carboxylate groups Allow $COONa$ or COO^-Na^+ but not covalent bond to Na allow NH_2-
4(e)(ii)	 **OR** 	1	In 4(e), do not penalise a slip in the number of carbons in the $-CH_2CH_2-$ chain, but all must be bonded correctly NB two ester groups allow NH_2- or $^+NH_3-$

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4(e)(iii)		1	In 4(e), do not penalise a slip in the number of carbons in the -CH_2CH_2- chain, but all must be bonded correctly allow anhydride formation on either or both COOH groups (see below) with or without amide group formation
4(f)	**M1** phase or eluent or solvent (or named solvent) is <u>moving or mobile</u>	1	
	M2 stationary phase or solid or alumina/silica/resin	1	
	M3 separation depends on balance between solubility or affinity (of compounds) in each phase **OR** different adsorption or retention **OR** (amino acids have) different R_f values **OR** (amino acids) travel at different speeds or take different times	1	

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Question	Marking Guidance	Mark	Comments
5(a)	**J** (acid) amide	1	not peptide, not N-substituted amide
	K (secondary) amine or amino	1	penalise primary or tertiary allow N-substituted amine
5(b)	(δ =) 3.1-3.9	1	Not 3.7 – 4.1
	doublet **OR** duplet	1	Not secondary name required not the number 2
5(c)(i)	Solvent must be proton-free **OR** $CHCl_3$ has protons or has H or gives a peak	1	
5(c)(ii)	$CDCl_3$ is polar **OR** CCl_4 is non-polar	1	
5(d)	11 **OR** eleven	1	
5(e)(i)	$Si(CH_3)_4$ **OR** SiC_4H_{12}	1	ignore TMS
5(e)(ii)	a single number or a range within 21-25	1	penalise anything outside this range
5(e)(iii)		1	allow ring around the C only and also allow

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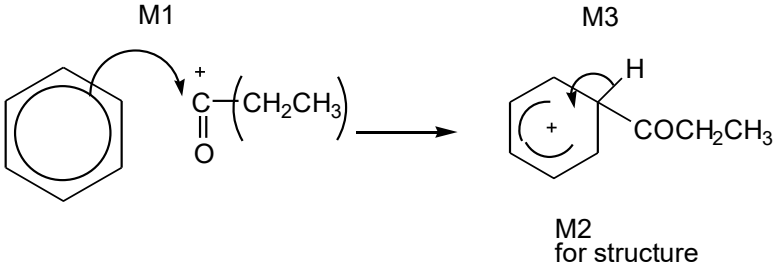
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5(f)(i)	$NaBH_4$	1	ignore name if formula correct ignore solvent allow $LiAlH_4$ Zn/HCl Sn/HCl H_2/Ni H_2/Pt
5(f)(ii)	$H_2N-\underset{\underset{O}{\parallel}}{C}-CH_2-\bigcirc-O-CH_2-\overset{OH}{(CH)}-CH_2-\overset{H}{\underset{\vert}{N}}-\overset{CH_3}{\underset{\vert}{CH}}\text{-}CH_3$	1	allow ring around the C only

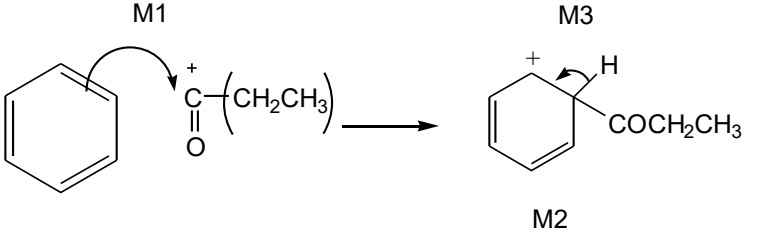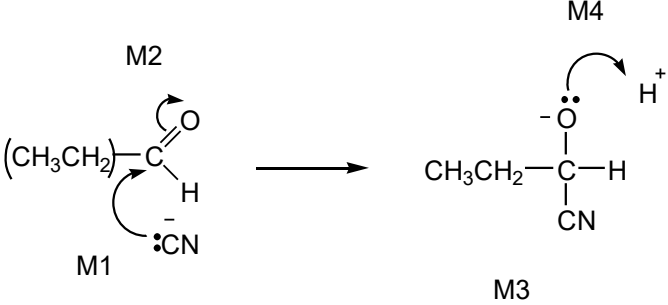| 5(f)(iii)	(plane) polarised light **OR** light in a polarimeter	1	
	polarised light is not rotated or is unaffected	1	penalise bent/diffracted/deflected/reflected Not just solution is optically inactive
5(f)(iv)	**adv** cheaper medicine due to cost or difficulty <u>of separation</u> or both can lower blood pressure 　　**OR** more effective/beneficial <u>with a reason</u> **disadv** may be side effects from one enantiomer in the mixture or only half the product works or one enantiomer may be ineffective or double dose required	1 1	or no need to separate

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Question	Marking Guidance	Mark	Comments
6(a)(i)	C_6H_6 + CH_3CH_2COCl → $C_6H_5COCH_2CH_3$ + HCl **OR** C_6H_6 + $CH_3CH_2CO^+$ → $C_6H_5COCH_2CH_3$ + H^+	1	allow C_2H_5 penalise C_6H_5–CH_3CH_2CO allow + on C or O in equation
	phenylpropanone **OR** ethylphenylketone **OR** phenylethylketone	1	Ignore 1 in formula, but penalise other numbers
	$AlCl_3$	1	can score in equation
	CH_3CH_2COCl + $AlCl_3$ → $CH_3CH_2CO^+$ + $AlCl_4^-$	1	allow C_2H_5 allow + on C or O in equation
	$AlCl_4^-$ + H^+ → $AlCl_3$ + HCl	1	

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| 6(a)(ii) | electrophilic substitution | 1

3 | can allow in (a)(i) if no contradiction

M1 arrow from circle or within it to C or to + on C horseshoe must not extend beyond C2 to C6 but can be smaller
+ not too close to C1

M2 penalise C_6H_5–CH_3CH_2CO (even if already penalized in (a)(i))

M3 arrow into hexagon unless Kekule

allow M3 arrow independent of M2 structure

ignore base removing H in M3 |

| 6(b)(i) | $CH_3CH_2\underline{CHO}$ + HCN   → $CH_3CH_2CH(OH)CN$ **OR**
$C_2H_5CH(OH)CN$

2-hydroxybutan<u>e</u>nitrile **OR** 2-hydroxybutan<u>o</u>nitrile | 1

1 | aldehyde must be -CHO
brackets optional

no others |

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6(b)(ii)	nucleophilic addition	1	
		4	M1 includes lp and arrow to Carbonyl C and minus charge (on either C or N) Not allow M2 before M1, but allow M1 to C^+ after non-scoring carbonyl arrow Ignore $\delta+$, $\delta-$ on carbonyl group, but if wrong way round or full + charge on C lose M2 M3 for correct structure including minus sign. Allow C_2H_5 M4 for lp and curly arrow to H^+
6(b)(iii)	(propanone) slower **OR** propanal faster inductive effects of alkyl groups **OR** C of C=O less $\delta+$ in propanone **OR** alkyl groups in ketone hinder attack **OR** easier to attack at end of chain	1 1	if wrong, no further marks

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Question	Marking Guidance	Mark	Comments
7(a)	diethylamine **OR** ethyl ethanamine **OR** ethyl aminoethane	1	ignore N-

Question	Marking Guidance				Mark
7(b)	For 7(b) and (c) There are three valid routes for this synthesis called Routes **A**, **B** and **C** below • Decide which route fits the answer best (this may not be the best for part b) to give the candidate the best possible overall mark. • Mark part (b) • For this best route mark the mechanism and reagent independently • Migration from one route to another is not allowed • Either name or formula is allowed in every case. • Ignore conditions unless they are incorrect.				
		Route A	**Route B**	**Route C**	
	F	CH_3CH_2Br or CH_3CH_2Cl	C_2H_6	CH_3CH_2OH	1
	G	$CH_3CH_2NH_2$ ethylamine **OR** ethanamine **OR** aminoethane	CH_3CH_2Br **OR** CH_3CH_2Cl	CH_3CH_2Br **OR** CH_3CH_2Cl	1

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7(c)			Route A	Route B	Route C	
	Step 1	Reagent(s)	HBr **OR** HCl	<u>H$_2$ / Ni</u> (Not NaBH$_4$)	H$_2$O & H$_3$PO$_4$ **OR** H$_2$O & H$_2$SO$_4$	1
		Mechanism	Electrophilic addition	addition (allow electrophilic **OR** catalytic but not nucleophilic) ignore hydrogenation	Electrophilic addition	1
	Step 2	Reagent(s)	NH$_3$	Cl$_2$ **OR** Br$_2$	HBr **OR** KBr & H$_2$SO$_4$ **OR** PCl$_3$ **OR** PCl$_5$ **OR** SOCl$_2$	1
		Mechanism	Nucleophilic substitution	(free) radical substitution	Nucleophilic substitution	1
	Step 3	Reagent(s)	CH$_3$CH$_2$Br **OR** CH$_3$CH$_2$Cl	CH$_3$CH$_2$NH$_2$ **OR** NH$_3$ but penalise excess ammonia here	CH$_3$CH$_2$NH$_2$ **OR** NH$_3$ but penalise excess ammonia here	1
		Mechanism	Nucleophilic substitution	Nucleophilic substitution	Nucleophilic substitution	1

7(d)	tertiary amine **OR** triethylamine **OR** (CH$_3$CH$_2$)$_3$N Quaternary ammonium salt **OR** tetraethylammonium bromide **OR** chloride **OR** ion **OR** (CH$_3$CH$_2$)$_4$N$^+$ (Br$^-$ **OR** Cl$^-$)	1	
	further substitution will take place **OR** diethylamine is a better nucleophile than ethylamine	1	

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General principles applied to marking CHEM4 papers by CMI+ (January 2011)

It is important to note that the guidance given here is generic and specific variations may be made at individual standardising meetings in the context of particular questions and papers.

Basic principles
- **Examiners should note that throughout the mark scheme, items that are underlined are <u>required information</u> to gain credit.**
- **Occasionally an answer involves incorrect chemistry and the mark scheme records CE = 0, which means a chemical error has occurred and no credit is given for that section of the clip or for the whole clip.**
-

A. The "List principle" and the use of "ignore" in the mark scheme

If a question requires **one** answer and a candidate gives two answers, no mark is scored if one answer is correct and one answer is incorrect. There is no penalty if both answers are correct.

N.B. Certain answers are designated in the mark scheme as those which the examiner should ─Ignore". These answers are not counted as part of the list and should be ignored and will not be penalised.

B. Incorrect case for element symbol

The use of an incorrect case for the symbol of an element should be penalised **once only** within a clip.
For example, penalise the use of ─h" for hydrogen, ─CL" for chlorine or ─br" for bromine.

C. Spelling

In general
- The names of chemical compounds and functional groups **must be spelled correctly** to gain credit.
- Phonetic spelling may be acceptable for some chemical terminology.

N.B. Some terms may be required to be spelled correctly or an idea needs to be articulated with clarity, as part of the ─Quality of Language" (**QoL**) marking. These will be identified in the mark scheme and marks are awarded only if the QoL criterion is satisfied.

D. <u>Equations</u>

In general
- Equations **must** be balanced.
- When an equation is worth two marks, one of the marks in the mark scheme will be allocated to one or more of the reactants or products. This is independent of the equation balancing.
- State symbols are <u>generally</u> ignored, unless specifically required in the mark scheme.

E. <u>Reagents</u>

The command word ─Identify", allows the candidate to choose to use **either** the name or the formula of a reagent in their answer. In some circumstances, the list principle may apply when both the name and the formula are used. Specific details will be given in mark schemes.

The guiding principle is that a reagent is a chemical which can be taken out of a bottle or container. Failure to identify complete reagents **will be penalised**, but follow-on marks (e.g. for a subsequent equation or observation) can be scored from an incorrect attempt (possibly an incomplete reagent) at the correct reagent. Specific details will be given in mark schemes.
For example, **no credit** would be given for

- the cyanide ion or CN^- when the reagent should be potassium cyanide or KCN;
- the hydroxide ion or OH^- when the reagent should be sodium hydroxide or NaOH;
- the $Ag(NH_3)_2^+$ ion when the reagent should be Tollens' reagent (or ammoniacal silver nitrate). In this example, no credit is given for the ion, but credit could be given for a correct observation following on from the use of the ion. Specific details will be given in mark schemes.

In the event that a candidate provides, for example, **both** KCN and cyanide ion, it would be usual to ignore the reference to the cyanide ion (because this is not contradictory) and credit the KCN. Specific details will be given in mark schemes.

F. <u>Oxidation states</u>

In general, the sign for an oxidation state will be assumed to be positive unless specifically shown to be negative.

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### G. <u>Marking calculations</u>

In general
- A correct answer alone will score **full marks** unless the necessity to show working is specifically required in the question.
- An arithmetic error may result in a one mark penalty if further working is correct.
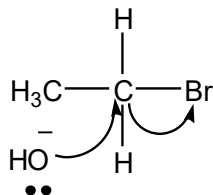- A chemical error will usually result in a two mark penalty.

H. <u>Organic reaction mechanisms</u>

Curly arrows should originate either from a lone pair of electrons or from a bond.
The following representations should not gain credit **and will be penalised each time** within a clip.
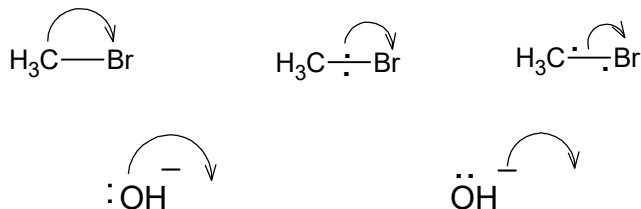
For example, the following would score zero marks

When the curly arrow is showing the formation of a bond to an atom, the arrow can go directly to the relevant atom, alongside the relevant atom or **more than half-way** towards the relevant atom.
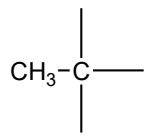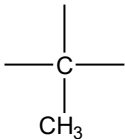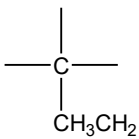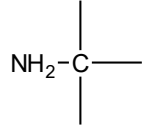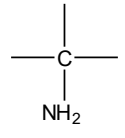
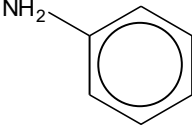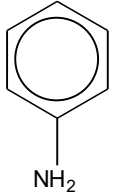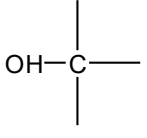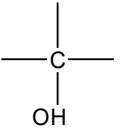In free-radical substitution
- The absence of a radical dot should be penalised **once only** within a clip.
- The use of double-headed arrows or the incorrect use of half-headed arrows in free-radical mechanisms should be penalised **once only** within a clip

In mass spectrometry fragmentation equations, the absence of a radical dot on the molecular ion and on the free-radical fragment would be considered to be two independent errors and both would be penalised if they occurred within the same clip.

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Mark Scheme – General Certificate of Education (A-level) Chemistry – Unit 4: Kinetics, Equilibria and Organic Chemistry – January 2011

I. **Organic structures**

In general
- Displayed formulae must show all of the bonds and all of the atoms in the molecule, but need not show correct bond angles.
- Bonds should be drawn correctly between the relevant atoms. This principle applies in all cases where the attached functional group contains a carbon atom, e.g nitrile, carboxylic acid, aldehyde and acid chloride. The carbon-carbon bond should be clearly shown. Wrongly bonded atoms will be penalised **on every occasion**. (see the examples below)
- The same principle should also be applied to the structure of alcohols.   For example, if candidates show the alcohol functional group as C ─ HO, they should be penalised **on every occasion**.
- Latitude should be given to the representation of C ─ C bonds in alkyl groups, given that CH_3─   is considered to be interchangeable with H_3C─   even though the latter would be preferred.
- Similar latitude should be given to the representation of amines where NH_2─ C will be allowed, although H_2N─ C would be preferred.
- Poor presentation of vertical C ─ CH_3 bonds or vertical C ─ NH_2 bonds should **not** be penalised. For other functional groups, such as ─ OH and ─ CN, the limit of tolerance is the half-way position between the vertical bond and the relevant atoms in the attached group.
    By way of illustration, the following would apply.

allowed	**allowed**	not allowed			
allowed	**allowed**	**allowed**	**allowed**	not allowed	not allowed

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CN—C— (structure)	—C—CN (structure)	COOH—C— (structure)	—C—COOH (structure)	—C—COOH (structure)	
not allowed	not allowed	not allowed	not allowed	not allowed	
CHO—C— (structure)	—C—CHO (structure)	—C—CHO (structure)	COCl—C— (structure)	—C—COCl (structure)	—C—COCl (structure)
not allowed	not allowed	not allowed	not allowed	not allowed	not allowed

- In most cases, the use of "sticks" to represent C ─ H bonds in a structure should **not** be penalised. The exceptions will include structures in mechanisms when the C ─ H bond is essential (e.g. elimination reactions in haloalkanes) and when a displayed formula is required.
- Some examples are given here of **structures** for specific compounds that should **not** gain credit

CH_3COH　　　　　for　　　　　ethanal

CH_3CH_2HO　　　　for　　　　　ethanol
$OHCH_2CH_3$　　　　for　　　　　ethanol
C_2H_6O　　　　　　for　　　　　ethanol

CH_2CH_2　　　　　for　　　　　ethene
$CH_2.CH_2$　　　　　for　　　　　ethene
$CH_2:CH_2$　　　　　for　　　　　ethane

N.B. Exceptions may be made in the context of balancing equations
- Each of the following **should gain credit** as alternatives to correct representations of the structures.

$CH_2 = CH_2$　　　　for　　　　　ethene,　$H_2C=CH_2$
$CH_3CHOHCH_3$　　for　　　　　propan-2-ol,　$CH_3CH(OH)CH_3$

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J. Organic names

As a general principle, non-IUPAC names or incorrect spelling or incomplete names should **not** gain credit.
Some illustrations are given here.

but-2-ol	should be **butan-2-ol**
2-hydroxybutane	should be **butan-2-ol**
butane-2-ol	should be **butan-2-ol**
2-butanol	should be **butan-2-ol**
2-methpropan-2-ol	should be **2-methylpropan-2-ol**
2-methylbutan-3-ol	should be **3-methylbutan-2-ol**
3-methylpentan	should be **3-methylpentane**
3-mythylpentane	should be **3-methylpentane**
3-methypentane	should be **3-methylpentane**
propanitrile	should be **propanenitrile**
aminethane	should be **ethylamine** (although aminoethane can gain credit)
2-methyl-3-bromobutane	should be **2-bromo-3-methylbutane**
3-bromo-2-methylbutane	should be **2-bromo-3-methylbutane**
3-methyl-2-bromobutane	should be **2-bromo-3-methylbutane**
2-methylbut-3-ene	should be **3-methylbut-1-ene**
difluorodichloromethane	should be **dichlorodifluoromethane**