M1. (a) (i)



Allow -CONH- or - COHN -

Mark two halves separately lose 1 each for missing trailing bonds at one or both ends or error in peptide link or either or both of H or OH on ends

1

1

1

1

1

Not allow $-(C_6H_{12})-$ Ignore n

(ii) **M1** in polyamides - H bonding

M2 in polyalkenes - van der Waals forces Penalise forces between atoms or van der Waals bonds

M3 Stronger forces (of attraction) in polyamides Or H bonding is stronger (must be a comparison of correct forces to score M3) *Do not award if refer to stronger bonds*

(b) (i) (nucleophilic) addition elimination



Not allow N-H₂

Minus sign on NH₂ loses M1

M2 not allowed independent of **M1**, but allow **M1** for correct attack on C+ + rather than δ + on C=O loses **M2** If CI lost with C=O breaking, max 1 for **M1 M3** for correct structure <u>with charges</u> but Ip on O is part of **M4** only allow **M4** after correct/ very close M3 For M4, ignore NH₃ removing H⁺ but lose **M4** for CI removing H⁺ in mechanism, but ignore HCI as a product 1

4

1

1

1

1

(ii) <u>N-methylpropanamide</u> Not N-methylpropaneamide

(c)





(d) (i) <u>2-amino-3-hydroxypropanoic acid</u>

(ii)



Must be salts of aspartic acio allow –CO₂⁻ allow NH₂–

(iii) Penalise use of aspartic acid once in d(iii) and d(iv)



1

1

1

(iv) Penalise use of aspartic acid once in d(iii) and d(iv)



allow $-CO_2^$ must show C-N bond don't penalize position of + on $N(CH_3)_3$

[16]



(ii)



1

(iii) <u>hydrogen</u> bonding (do not allow H-bonding) QWC do not penalise any error twice.

1







1

1

2

1

1

1

(iii) Isomer must be saturated or must not contain a double bond

(c) $\begin{array}{c} ---- (CH_2)_2 - C --- N - (CH_2)_6 - N ---- \\ || & || & | & | \\ 0 & 0 & H & H \end{array}$

(d) (i) heat/reflux with aqu NaOH

poly(alkene) is inert/ no reaction

(ii)	e.g combustion	1	
	heat energy produced	1	
	toxic gases produced	1	[14]

1



[8]

M4. (i) CH₃CH=CHCH₃ (a) 1 Addition or radical (QoL) 1 $CH_{3}CH(OH)CH(OH)CH_{3}$ or with no brackets (ii) 1 butan(e)-2.3-diol or 2.3-butan(e)diol 1 $\begin{array}{cccc} H & H & H & H \\ \downarrow & \downarrow \\ -C & -C & -COCH & allow & ClOC & -C & -COCl \\ \downarrow & \downarrow & \downarrow \\ CH_3 & CH_3 & & CH_3 & CH_3 \end{array}$ HOOC-1 2.3-dimethylbutan(e)dioic acid 2.3-dimethylbutan(e)dioyl chloride ignore -1,4-1 condensation (QoL) 1 (iii) NaOH or HCl etc or Na₂CO₃

Allow conc sulphuric/nitric

$$\begin{array}{ccc} CH_{3} & CH_{2}OH \\ I & I \\ H_{2}N & -C - C - N - C - COOH \\ I & I & I \\ H & O & H \end{array}$$

Allow –CONH– and –COHN– Allow zwitterions **NOT polypeptides/repeating units**

1

1

1

1

1

Structure 2 either of



further substitution/reaction occurs or other products are formed Allow reduction forms only one product

one of (CH₃CH₂CH₂)₂NH (CH₃CH₂CH₂)₃N (CH₃CH₂CH₂)₄N⁺ Br⁻ *Allow salts including NH₄Br Allow HBr*

1