

GAT B Chemistry

Sample Paper – 6

Duration: 23 Minutes

Maximum Marks: 15

Instructions

- This paper contains **15** Multiple Choice Questions (Single Correct Answer), modelled on the Chemistry portion of **GAT B** entrance.
- Each correct answer carries **+1 marks**. There is a **deduction of 0.5 marks for each incorrect answer**.
- Only **one** option is correct. Choose carefully.
- Syllabus level: **NCERT Class 11 & 12 Chemistry — Physical, Organic, and Inorganic Chemistry**.
- Use of mobile phones, calculators, or electronic gadgets is strictly prohibited.

Questions

- Q1.** Which of the following statements **correctly** describes the structural difference between cellulose and starch (amylose)?
- (A) Cellulose has α -1,4 glycosidic bonds, while amylose (starch) has β -1,4 glycosidic bonds.
- (B) Both cellulose and amylose consist of α -D-glucose units and differ only in molecular weight.
- (C) Starch contains β -1,6 branch points while cellulose has only α -1,4 glycosidic bonds.
- (D) Cellulose has β -1,4 glycosidic bonds, while amylose (starch) has α -1,4 glycosidic bonds, leading to very different structural and functional properties.
- Q2.** Which of the following correctly describes the structure and function of transfer RNA (tRNA) in protein synthesis?



- (A) tRNA possesses an **anticodon loop** that base-pairs with the complementary mRNA codon, and carries the cognate amino acid esterified to the hydroxyl group at the **CCA-3'** terminus.
- (B) tRNA carries amino acids at its 5'-phosphate end and recognises codons through a stem-loop structure located at the centre of the molecule.
- (C) The anticodon of tRNA is located at the 3' end, and the amino acid is attached at the CCA sequence at the 5' terminus.
- (D) tRNA functions during transcription in the nucleus to ferry amino acids to the growing pre-mRNA strand.

Q3. Which statement **correctly** distinguishes a cofactor from a coenzyme?

- (A) Cofactors are always organic molecules; coenzymes are always inorganic metal ions.
- (B) *Cofactor* is a broader term encompassing both inorganic metal ions and organic molecules; *coenzyme* refers specifically to the **organic cofactors**, many of which are derived from vitamins.
- (C) Coenzymes are permanently and covalently attached to the enzyme (prosthetic group), while cofactors can freely dissociate.
- (D) Both cofactors and coenzymes are proteins that associate non-covalently with the enzyme active site.

Q4. 50 mL of 0.2 M HCl is titrated with 0.2 M NH_3 ($K_b = 1.0 \times 10^{-5}$ at 298 K). What is the **pH at the equivalence point**?

- (A) 7.00
- (B) 9.00
- (C) 5.00
- (D) 11.00

Q5. An aqueous solution contains 0.1 M Cl^- and 0.1 M I^- ions. When AgNO_3 solution is added dropwise with stirring, which ion is precipitated **first**?



$$[K_{sp}(\text{AgCl}) = 1.8 \times 10^{-10}; \quad K_{sp}(\text{AgI}) = 8.5 \times 10^{-17}]$$

- (A) Cl^- , because it is more commonly encountered and reacts faster with Ag^+ .
- (B) Both Cl^- and I^- simultaneously, since their concentrations are equal.
- (C) Cl^- , because AgCl is less soluble than AgI .
- (D) I^- , because AgI has a much smaller K_{sp} , so a far lower $[\text{Ag}^+]$ is required to initiate its precipitation.

Q6. For a reaction: $\Delta H^\circ = -30 \text{ kJ mol}^{-1}$ and $\Delta S^\circ = -100 \text{ J mol}^{-1}\text{K}^{-1}$. Using $\Delta G = \Delta H - T\Delta S$, at what temperature (in K) is $\Delta G = 0$, and **below** what temperature is the reaction spontaneous?

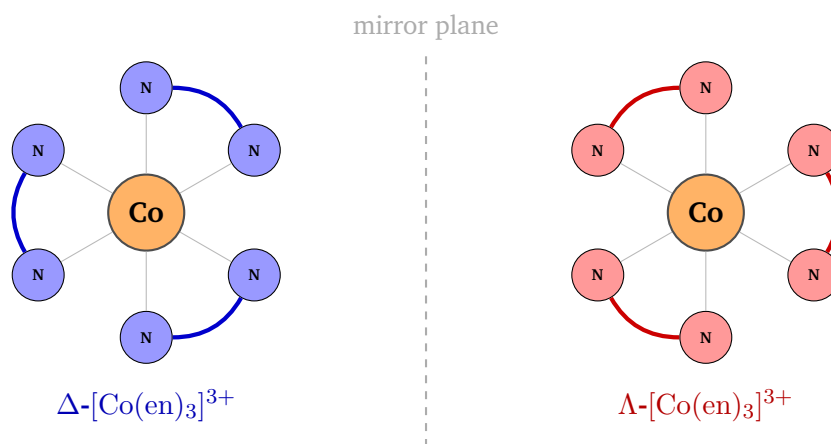
- (A) $T = 300 \text{ K}$; the reaction is spontaneous **below** 300 K.
- (B) $T = 300 \text{ K}$; the reaction is spontaneous **above** 300 K.
- (C) $T = 3000 \text{ K}$; the reaction is spontaneous above 3000 K.
- (D) $T = 3000 \text{ K}$; the reaction is spontaneous at all temperatures since $\Delta H < 0$.

Q7. The standard molar enthalpy of neutralisation of a **strong acid** with a **strong base** at 298 K is approximately:

- (A) $-13.7 \text{ kJ mol}^{-1}$
- (B) $-57.3 \text{ kJ mol}^{-1}$
- (C) $-46.0 \text{ kJ mol}^{-1}$
- (D) $-285.8 \text{ kJ mol}^{-1}$

Q8. The schematic trigonal projections below show the two non-superimposable mirror images (Δ and Λ forms) of a tris-bidentate octahedral cobalt complex. Which of the following complexes **exhibits optical isomerism**?





(Schematic top-view: each thick arc represents one ethylenediamine ligand)

- (A) $[\text{Co}(\text{NH}_3)_4\text{Cl}_2]^+$
- (B) $[\text{PtCl}_4]^{2-}$
- (C) $[\text{Co}(\text{en})_3]^{3+}$
- (D) $[\text{Ni}(\text{CO})_4]$

Q9. According to the **18-electron rule** (effective atomic number rule), which of the following metal carbonyls does **NOT** obey it?

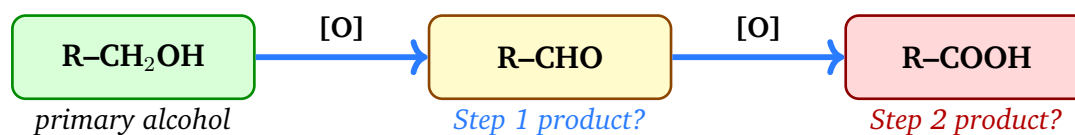
- (A) $\text{Ni}(\text{CO})_4$
- (B) $\text{Fe}(\text{CO})_5$
- (C) $\text{Cr}(\text{CO})_6$
- (D) $\text{V}(\text{CO})_6$

Q10. Ethylmagnesium bromide ($\text{CH}_3\text{CH}_2\text{MgBr}$) is treated first with dry CO_2 gas, then worked up with dilute H_3O^+ . What is the **major organic product**?

- (A) Propanoic acid ($\text{CH}_3\text{CH}_2\text{COOH}$)
- (B) Propanal ($\text{CH}_3\text{CH}_2\text{CHO}$)
- (C) 1-Propanol ($\text{CH}_3\text{CH}_2\text{CH}_2\text{OH}$)
- (D) Ethanol ($\text{CH}_3\text{CH}_2\text{OH}$)

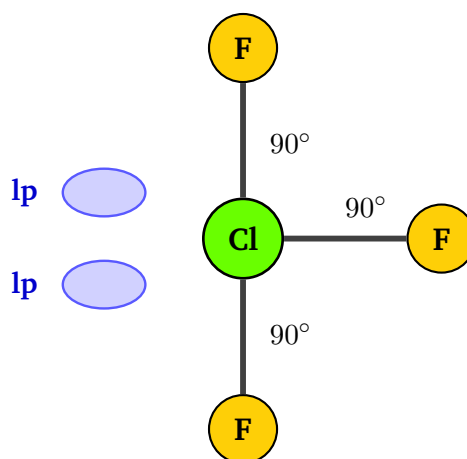


- Q11.** The diagram below illustrates the stepwise oxidation of a primary alcohol. The products formed in the **first** and **second** oxidation steps are, respectively:



- (A) Ketone, then carboxylic acid
(B) Aldehyde, then carboxylic acid
(C) Ether, then ketone
(D) Aldehyde, then ester
- Q12.** Which of the following correctly expresses the standard thermodynamic relationships among ΔG° , the standard cell potential E°_{cell} , and the equilibrium constant K ?
- (A) $\Delta G^\circ = +nFE^\circ = +RT \ln K$
(B) $\Delta G^\circ = -nFE^\circ = +RT \ln K$
(C) $\Delta G^\circ = -nFE^\circ = -RT \ln K$
(D) $\Delta G^\circ = +nFE^\circ = -RT \ln K$
- Q13.** A plot of $\ln k$ versus $1/T$ for a reaction gives a straight line with slope = -9600 K . What is the **activation energy** (E_a) of the reaction?
($R = 8.314 \text{ J mol}^{-1} \text{ K}^{-1}$)
- (A) 9600 J mol^{-1}
(B) 1154 J mol^{-1}
(C) 120 kJ mol^{-1}
(D) 79.8 kJ mol^{-1}
- Q14.** The diagram below depicts the molecular geometry of an interhalogen compound containing one chlorine and three fluorine atoms. Identify the compound and its **correct molecular geometry**:





Molecular geometry of the interhalogen compound

- (A) ClF_3 ; pyramidal geometry due to one lone pair on chlorine
- (B) ClF_3 ; T-shaped geometry arising from two equatorial lone pairs on chlorine
- (C) ClF_3 ; trigonal planar geometry with all bond angles 120°
- (D) Cl_3F ; linear geometry with chlorine at the centre

Q15. Which of the following **correctly** identifies the monomers of **Nylon-6,6** and the type of linkage formed during polymerisation?

- (A) Caprolactam alone; amide linkage by ring-opening polymerisation.
- (B) Adipic acid and ethylene glycol; ester linkage (polyester).
- (C) Adipic acid $[\text{HOOC}(\text{CH}_2)_4\text{COOH}]$ and hexamethylenediamine $[\text{H}_2\text{N}(\text{CH}_2)_6\text{NH}_2]$ **amide** linkage.
- (D) Hexamethylenediamine and sebacic acid $[\text{HOOC}(\text{CH}_2)_8\text{COOH}]$; amide linkage.

Detailed Solutions



Q1.

Solution

Concept — Polysaccharide Stereochemistry: Cellulose and starch are both homopolymers of glucose, but the configuration at the anomeric carbon (C-1) of the glycosidic bond produces completely different structures and biological roles.

Step 1 — Starch (Amylose): Amylose is composed of α -D-glucose units joined by α -1,4 glycosidic bonds. The α configuration bends the chain, causing it to coil into a helix (useful for energy storage; digested by amylase enzymes).

Step 2 — Cellulose: Cellulose is composed of β -D-glucose units joined by β -1,4 glycosidic bonds. Each glucose unit is “flipped” 180° relative to its neighbour, producing a straight, rigid chain that forms inter-chain hydrogen bonds — hence its structural role in plant cell walls. Humans cannot digest cellulose because they lack β -glucosidase.

Why other options are wrong:

- Option A: Reverses the bond types — cellulose is β -1,4 and amylose is α -1,4, not the other way round.
- Option B: Only starch uses α -D-glucose; cellulose uses β -D-glucose. They differ in linkage configuration, not just molecular weight.
- Option C: It is amylopectin (branched starch) that has α -1,6 branch points (not β -1,6); cellulose has β -1,4 bonds, not α -1,4.
- Option D: Correct in every detail.

Final Answer: Cellulose = β -1,4; amylose = α -1,4 $\Rightarrow$ D

Answer: (D) [Go Back to Q 1](#)

Q2.

Solution

Concept — tRNA Architecture: tRNA is the adaptor molecule of translation, possessing two functionally critical sites: the anticodon loop (mRNA recognition) and the 3'-CCA terminus (amino acid attachment).

Step 1 — Anticodon Loop: The anticodon, a triplet of nucleotides, is located at the anticodon loop — roughly at the bottom of the L-shaped 3D structure. It forms Watson–Crick base pairs (antiparallel, $3' \rightarrow 5'$) with the complementary mRNA codon on the ribosome.



Step 2 — CCA-3' Terminus: All mature tRNA molecules share the universal 3' terminal sequence . . .CCA-3'. Aminoacyl-tRNA synthetase attaches the correct amino acid via an ester bond to the 2'- or 3'-OH of the terminal adenosine, forming aminoacyl-tRNA.

Why other options are wrong:

- Option A: Correct — anticodon loop + amino acid at CCA-3' end.
- Option B: Amino acid is attached at the 3' end, not the 5' end; and codon recognition occurs at the anticodon loop, not general stem-loops.
- Option C: The anticodon is at the anticodon loop (central region), not the 3' end.
- Option D: tRNA functions in **translation** (cytoplasm/ribosome), not in transcription in the nucleus.

Final Answer: Anticodon loop reads mRNA; amino acid attached at CCA-3' $\Rightarrow$ A

Answer: (A) [Go Back to Q 2](#)

Q3.

Solution

Concept — Enzyme Auxiliary Molecules: Many enzymes require non-protein chemical species for catalytic activity. The term *cofactor* is the general name; *coenzyme* is a subset.

Step 1 — Cofactor (Broad Category): A *cofactor* is any non-protein component needed for enzyme activity. This includes:

- Inorganic metal ions: Zn^{2+} (carbonic anhydrase), Mg^{2+} (kinases), $\text{Fe}^{2+}/\text{Fe}^{3+}$ (cytochromes).
- Organic molecules (coenzymes): NAD^+ , FAD, Coenzyme A, etc.

Step 2 — Coenzyme (Organic Subset): A *coenzyme* is specifically an **organic cofactor**, usually a vitamin derivative. Coenzymes that bind loosely and transiently are cosubstrates; those bound permanently are *prosthetic groups*. Thus: coenzyme $\subset$ cofactor.

Why other options are wrong:

- Option A: Reverses the definitions; cofactors include inorganic ions, while coenzymes are organic.



- Option B: Correct — cofactor is broader; coenzyme is the organic subset, often vitamin-derived.
- Option C: It is the *prosthetic group* (a specific class of cofactor) that is permanently bound. Coenzymes typically dissociate after each reaction cycle.
- Option D: Cofactors and coenzymes are **not** proteins; they are non-protein helper molecules.

Final Answer: Cofactor (broad) $\supset$ metal ions + coenzymes; coenzyme = organic cofactor $\Rightarrow$ B

Answer: (B) [Go Back to Q 3](#)

Q4.

Solution

Concept — Equivalence Point pH (Strong Acid + Weak Base): At the equivalence point of HCl vs. NH_3 , all base is converted to the salt NH_4Cl . The solution is acidic because NH_4^+ (the conjugate acid) hydrolyses.

Step 1 — Composition at Equivalence Point:

$$n_{\text{HCl}} = 0.050 \text{ L} \times 0.2 \text{ M} = 0.010 \text{ mol}; \quad n_{\text{NH}_3} = 0.050 \text{ L} \times 0.2 \text{ M} = 0.010 \text{ mol}$$

They react completely: $\text{HCl} + \text{NH}_3 \rightarrow \text{NH}_4\text{Cl}$. Total volume = 100 mL.

$$[\text{NH}_4^+] = \frac{0.010 \text{ mol}}{0.100 \text{ L}} = 0.10 \text{ M}$$

Step 2 — Hydrolysis of NH_4^+ :

$$K_a(\text{NH}_4^+) = \frac{K_w}{K_b} = \frac{1.0 \times 10^{-14}}{1.0 \times 10^{-5}} = 1.0 \times 10^{-9}$$

$$[\text{H}^+] = \sqrt{K_a \cdot C} = \sqrt{1.0 \times 10^{-9} \times 0.10} = \sqrt{1.0 \times 10^{-10}} = 1.0 \times 10^{-5} \text{ M}$$

$$\text{pH} = -\log(1.0 \times 10^{-5}) = 5.00$$

Why other options are wrong:

- Option A: pH 7.00 applies only to strong acid–strong base neutralisation (no hydrolysis of salt).
- Option B: pH 9.00 would be expected at the equivalence point of a *weak acid*–strong base titration.



- Option C: pH 5.00 is the correct result.
- Option D: pH 11.00 is strongly alkaline; this titration gives an acidic salt at equivalence.

Final Answer: $\text{pH} = 5.00 \Rightarrow \boxed{\text{C}}$

Answer: (C) [Go Back to Q 4](#)

Q5.

Solution

Concept — Selective Precipitation by Competing Equilibria: When two anions compete for the same cation, the one that requires the *lower* free-cation concentration to exceed its K_{sp} precipitates first.

Step 1 — Minimum $[\text{Ag}^+]$ Needed for Each Precipitate:

$$[\text{Ag}^+]_{\text{AgCl}} = \frac{K_{sp}(\text{AgCl})}{[\text{Cl}^-]} = \frac{1.8 \times 10^{-10}}{0.1} = 1.8 \times 10^{-9} \text{ M}$$

$$[\text{Ag}^+]_{\text{AgI}} = \frac{K_{sp}(\text{AgI})}{[\text{I}^-]} = \frac{8.5 \times 10^{-17}}{0.1} = 8.5 \times 10^{-16} \text{ M}$$

Step 2 — Comparison: Since $8.5 \times 10^{-16} \ll 1.8 \times 10^{-9}$, AgI begins to precipitate when $[\text{Ag}^+]$ is still far below the threshold for AgCl. Therefore I^- precipitates first. Cl^- remains in solution until $[\text{Ag}^+]$ rises to $1.8 \times 10^{-9} \text{ M}$.

Why other options are wrong:

- Option A: Kinetic speed is irrelevant here; selective precipitation is governed by thermodynamics (K_{sp} values).
- Option B: Equal concentrations do not cause simultaneous precipitation when K_{sp} values differ by ~ 7 orders of magnitude.
- Option C: AgCl ($K_{sp} = 1.8 \times 10^{-10}$) is *more* soluble than AgI ($K_{sp} = 8.5 \times 10^{-17}$), so Cl^- precipitates *later*.
- Option D: Correct — AgI has the smaller K_{sp} ; I^- precipitates first.

Final Answer: I^- precipitates first (smaller K_{sp} for AgI) $\Rightarrow \boxed{\text{D}}$

Answer: (D) [Go Back to Q 5](#)



Q6.

Solution

Concept — Gibbs–Helmholtz Equation: When both ΔH and ΔS are negative, $\Delta G = \Delta H - T\Delta S$ changes sign at a specific temperature $T^* = \Delta H/\Delta S$. The reaction is spontaneous only below T^* .

Step 1 — Temperature at $\Delta G = 0$:

$$0 = \Delta H^\circ - T^* \Delta S^\circ \implies T^* = \frac{\Delta H^\circ}{\Delta S^\circ} = \frac{-30,000 \text{ J mol}^{-1}}{-100 \text{ J mol}^{-1} \text{ K}^{-1}} = 300 \text{ K}$$

Step 2 — Spontaneity Range:

$$\Delta G = (-30,000) - T(-100) = -30,000 + 100T$$

- $T < 300 \text{ K}$: $\Delta G = -30,000 + 100T < 0 \Rightarrow$ **spontaneous**.
- $T > 300 \text{ K}$: $\Delta G > 0 \Rightarrow$ non-spontaneous.

Why other options are wrong:

- Option A: Correct — $T^* = 300 \text{ K}$; spontaneous below 300 K.
- Option B: When $\Delta H < 0$ and $\Delta S < 0$, spontaneity holds at *low* temperature, not high.
- Option C and D: Arithmetic error by a factor of 10: $(-30,000)/(-100) = 300$, not 3000.

Final Answer: $T^* = 300 \text{ K}$; spontaneous below 300 K $\Rightarrow$ A

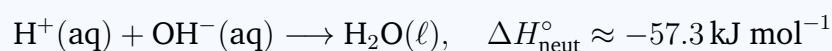
Answer: (A) [Go Back to Q 6](#)

Q7.

Solution

Concept — Standard Enthalpy of Neutralisation: For any strong acid–strong base pair, both are fully dissociated, so the net ionic equation is always the same: $\text{H}^+(\text{aq}) + \text{OH}^-(\text{aq}) \rightarrow \text{H}_2\text{O}(\text{l})$. The enthalpy change is therefore constant and independent of which strong acid or base is used.

Step 1 — Net Ionic Equation and Value:



This value is the standard result quoted in NCERT and all major physical chemistry references.

Step 2 — Identifying the Distractors:

- $-13.7 \text{ kJ mol}^{-1}$: No standard thermodynamic significance in this context.
- $-46.0 \text{ kJ mol}^{-1}$: Approximate value for neutralisation involving a weak acid or weak base (energy is consumed to complete ionisation of the weak species, reducing $|\Delta H|$).
- $-285.8 \text{ kJ mol}^{-1}$: This is $\Delta H_f^\circ(\text{H}_2\text{O}, \ell)$ from elements — an entirely different reaction.

Why other options are wrong:

- Option A: $-13.7 \text{ kJ mol}^{-1}$ is incorrect; no known standard value here.
- Option B: $-57.3 \text{ kJ mol}^{-1}$ is the correct standard enthalpy of neutralisation.
- Option C: $-46.0 \text{ kJ mol}^{-1}$ applies to weak acid/weak base systems, not strong/strong.
- Option D: $-285.8 \text{ kJ mol}^{-1}$ is ΔH_f° of liquid water, not neutralisation enthalpy.

Final Answer: $\Delta H_{\text{neut}}^\circ = -57.3 \text{ kJ mol}^{-1} \Rightarrow \boxed{\text{B}}$

Answer: (B) [Go Back to Q 7](#)

Q8.

Solution

Concept — Optical Isomerism in Coordination Compounds: A complex is optically active (chiral) if it lacks an improper rotation axis (S_n , including mirror planes and inversion centres). Tris-bidentate octahedral complexes $[\text{M}(\text{AA})_3]^{n+}$ with D_3 symmetry are always chiral.

Step 1 — Analysis of Each Complex:

- $[\text{Co}(\text{NH}_3)_4\text{Cl}_2]^+$: Both cis and trans forms have mirror planes (C_{2v} and D_{4h} symmetry, respectively) — optically **inactive**.
- $[\text{PtCl}_4]^{2-}$: Square-planar with D_{4h} symmetry; multiple mirror planes — optically **inactive**.
- $[\text{Co}(\text{en})_3]^{3+}$: Octahedral, three bidentate ethylenediamine ligands, D_3 symmetry. No mirror planes, no S_4 axis. Exists as non-superimposable Δ and Λ enantiomers (shown in the diagram) — optically **active**.



- $[\text{Ni}(\text{CO})_4]$: Tetrahedral with T_d symmetry; has S_4 axis — optically **inactive**.

Step 2 — Confirming $[\text{Co}(\text{en})_3]^{3+}$ via the Diagram: The Δ -form (blue arcs, clockwise) and Λ -form (red arcs, counterclockwise) shown are mirror images that cannot be superimposed by any rotation, confirming chirality.

Why other options are wrong:

- Option A: $[\text{Co}(\text{NH}_3)_4\text{Cl}_2]^+$ possesses mirror planes in both geometric forms; achiral.
- Option B: $[\text{PtCl}_4]^{2-}$ is square-planar with D_{4h} symmetry; achiral.
- Option C: $[\text{Co}(\text{en})_3]^{3+}$ is D_3 -symmetric and chiral — **correct**.
- Option D: $[\text{Ni}(\text{CO})_4]$ is tetrahedral (T_d); achiral.

Final Answer: $[\text{Co}(\text{en})_3]^{3+}$ (D_3 symmetry, no mirror planes) $\Rightarrow$ C

Answer: (C) [Go Back to Q 8](#)

Q9.

Solution

Concept — 18-Electron Rule in Metal Carbonyls: Each CO donates 2 electrons to the metal centre. A stable metal carbonyl satisfies: metal valence electrons (= Group number) + electrons from CO = 18.

Step 1 — Electron Count:

Carbonyl	Metal valence e^-	CO donation	Total
$\text{Ni}(\text{CO})_4$	10 (Group 10)	$4 \times 2 = 8$	18 ✓
$\text{Fe}(\text{CO})_5$	8 (Group 8)	$5 \times 2 = 10$	18 ✓
$\text{Cr}(\text{CO})_6$	6 (Group 6)	$6 \times 2 = 12$	18 ✓
$\text{V}(\text{CO})_6$	5 (Group 5)	$6 \times 2 = 12$	17 ×

Step 2 — Conclusion: $\text{V}(\text{CO})_6$ has only **17 electrons**. It is paramagnetic (one unpaired electron) and is the textbook example of a kinetically stable 17-electron complex that violates the 18-electron rule. It does not dimerize (unlike 17-electron $\text{Mn}_2(\text{CO})_{10}$) due to steric protection by the six CO groups.

Why other options are wrong:



- Option A: $\text{Ni}(\text{CO})_4$ — $10 + 8 = 18$ electrons. Obeys the rule.
- Option B: $\text{Fe}(\text{CO})_5$ — $8 + 10 = 18$ electrons. Obeys the rule.
- Option C: $\text{Cr}(\text{CO})_6$ — $6 + 12 = 18$ electrons. Obeys the rule.
- Option D: $\text{V}(\text{CO})_6$ — $5 + 12 = 17$ electrons. Does **not** obey the rule.

Final Answer: $\text{V}(\text{CO})_6$ has 17 electrons; violates the 18-electron rule $\Rightarrow$ **D**

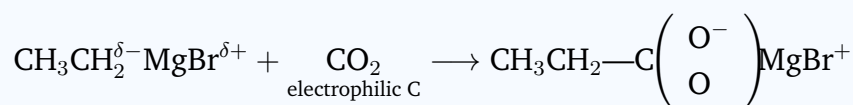
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Q10.

Solution

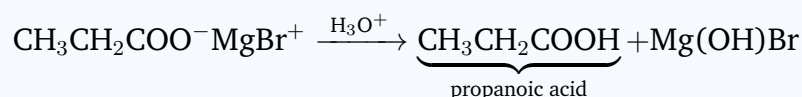
Concept — Grignard Reagent with CO_2 (Carboxylation): A Grignard reagent (RMgX) is a strong carbon nucleophile (carbanion equivalent). Reaction with CO_2 inserts a $-\text{COO}^-$ group; acid workup gives a carboxylic acid one carbon longer than the original alkyl group.

Step 1 — Nucleophilic Addition to CO_2 :



The ethyl carbanion attacks the electrophilic carbon of CO_2 , forming a magnesium carboxylate.

Step 2 — Acid Workup:



Net result: C_2 (ethyl) Grignard + $\text{CO}_2 \rightarrow \text{C}_3$ carboxylic acid (one carbon added).

Why other options are wrong:

- Option A: Propanoic acid ($\text{CH}_3\text{CH}_2\text{COOH}$) — **correct product** of carboxylation.
- Option B: Propanal ($\text{CH}_3\text{CH}_2\text{CHO}$) — requires reaction with DMF or formamide, not CO_2 .
- Option C: 1-Propanol ($\text{CH}_3\text{CH}_2\text{CH}_2\text{OH}$) — would arise from Grignard reaction with *formaldehyde* (HCHO) followed by acid workup.
- Option D: Ethanol ($\text{CH}_3\text{CH}_2\text{OH}$) — formed by protonation of the Grignard with water; no CO_2 involved, no chain extension.

Final Answer: $\text{CH}_3\text{CH}_2\text{MgBr} + \text{CO}_2 \xrightarrow{\text{H}_3\text{O}^+} \text{CH}_3\text{CH}_2\text{COOH} \Rightarrow$ **A**



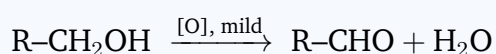
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Q11.

Solution

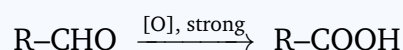
Concept — Stepwise Oxidation of Primary Alcohols: Primary alcohols are oxidised in two stages: first to an aldehyde (by mild oxidants such as PCC, Swern, or pyridinium dichromate), then further to a carboxylic acid (by strong oxidants such as acidic KMnO_4 or $\text{K}_2\text{Cr}_2\text{O}_7/\text{H}^+$).

Step 1 — First Oxidation: Primary Alcohol $\rightarrow$ Aldehyde:



Two hydrogen atoms are removed (one from C-H, one from O-H at the α -carbon). Mild oxidants stop here.

Step 2 — Second Oxidation: Aldehyde $\rightarrow$ Carboxylic Acid:



The aldehyde is oxidised further; the carbonyl carbon's oxidation state increases from +1 to +3.

Why other options are wrong:

- Option A: Ketones arise from oxidation of *secondary* alcohols; a primary alcohol cannot directly give a ketone.
- Option B: Aldehyde then carboxylic acid — **correct sequence**.
- Option C: Ether formation requires acid-catalysed dehydration (H_2SO_4 , 140°C), not oxidation.
- Option D: Ester formation requires the combined action of a carboxylic acid and an alcohol (Fischer esterification); sequential oxidation alone cannot yield an ester.

Final Answer: $\text{R-CH}_2\text{OH} \xrightarrow{[\text{O}]} \text{R-CHO} \xrightarrow{[\text{O}]} \text{R-COOH} \Rightarrow \boxed{\text{B}}$

Answer: (B) [Go Back to Q 11](#)



Q12.

Solution

Concept — Interrelationship of ΔG° , E° , and K : Three pivotal equations of physical chemistry connect these quantities. Knowing the sign conventions and derivations eliminates all distractors.

Step 1 — Derivation of $\Delta G^\circ = -RT \ln K$: At equilibrium, $\Delta G = 0$ and the reaction quotient $Q = K$. Using $\Delta G = \Delta G^\circ + RT \ln Q$:

$$0 = \Delta G^\circ + RT \ln K \implies \Delta G^\circ = -RT \ln K$$

(If $K > 1$, products are favoured and $\Delta G^\circ < 0$; if $K < 1$, $\Delta G^\circ > 0$.)

Step 2 — Derivation of $\Delta G^\circ = -nFE^\circ$: The maximum non-expansion (electrical) work an electrochemical cell can do equals the drop in Gibbs energy:

$$w_{\text{elec,max}} = -\Delta G^\circ = nFE^\circ \implies \Delta G^\circ = -nFE^\circ$$

(A positive E° drives a spontaneous cell, $\Delta G^\circ < 0$.)

Combined: $\Delta G^\circ = -nFE^\circ = -RT \ln K$.

Why other options are wrong:

- Option A: Both terms have the wrong sign; $+nFE^\circ$ and $+RT \ln K$ are both incorrect.
- Option B: $-nFE^\circ$ is correct but $+RT \ln K$ has the wrong sign.
- Option C: $\Delta G^\circ = -nFE^\circ = -RT \ln K$ — **correct**.
- Option D: $\Delta G^\circ = +nFE^\circ$ has the wrong sign for the electrochemical relation.

Final Answer: $\Delta G^\circ = -nFE^\circ = -RT \ln K \Rightarrow \boxed{\text{C}}$

Answer: (C) [Go Back to Q 12](#)

Q13.

Solution

Concept — Arrhenius Plot and Activation Energy: The Arrhenius equation in linear form is $\ln k = \ln A - \frac{E_a}{R} \cdot \frac{1}{T}$. A graph of $\ln k$ vs. $1/T$ yields a straight line with slope $= -E_a/R$, from which E_a is extracted.



Step 1 — Extract E_a from the Slope:

$$\text{slope} = -\frac{E_a}{R} = -9600 \text{ K} \Rightarrow E_a = 9600 \text{ K} \times R = 9600 \text{ K} \times 8.314 \text{ J mol}^{-1}\text{K}^{-1}$$

$$E_a = 79,814 \text{ J mol}^{-1} \approx 79.8 \text{ kJ mol}^{-1}$$

Step 2 — Unit Check: $[\text{slope}] = \text{K}$; $[R] = \text{J mol}^{-1}\text{K}^{-1}$; $[E_a] = \text{K} \times \text{J mol}^{-1}\text{K}^{-1} = \text{J mol}^{-1}$. Units are consistent.

Why other options are wrong:

- Option A: 9600 J mol^{-1} is the numerical value of the slope alone, forgetting to multiply by R .
- Option B: 1154 J mol^{-1} results from dividing by R instead of multiplying ($9600/8.314 \approx 1154$).
- Option C: 120 kJ mol^{-1} is arithmetically wrong; $9600 \times 8.314 = 79,814 \text{ J mol}^{-1} \neq 120,000$.
- Option D: 79.8 kJ mol^{-1} is the correct result ($9600 \times 8.314 = 79,814 \approx 79.8 \text{ kJ mol}^{-1}$).

Final Answer: $E_a = 9600 \times 8.314 \approx 79.8 \text{ kJ mol}^{-1} \Rightarrow \boxed{\text{D}}$

Answer: (D) [Go Back to Q 13](#)

Q14.

Solution

Concept — VSEPR Theory Applied to ClF_3 : Cl has 7 valence electrons; it forms 3 bonds with F and retains 2 lone pairs. The VSEPR notation is AB_3E_2 : electron-pair geometry is *trigonal bipyramidal* (5 pairs), and lone pairs preferentially occupy equatorial positions to minimise repulsion. The resulting molecular geometry is **T-shaped**.

Step 1 — Valence Electron Count for Cl in ClF_3 :

Cl valence electrons = 7; Shared in 3 bonds: 3; Non-bonding: $7-3 = 4 \rightarrow 2$ lone pairs

Total electron pairs on Cl = $3(\text{bonding}) + 2(\text{lone pairs}) = 5 \rightarrow$ trigonal bipyramidal arrangement.

Step 2 — Lone-Pair Position and Molecular Shape: Lone pairs require more angular space than bonding pairs. In a trigonal bipyramid, equatorial positions



(three positions, 120° apart) have less repulsion than axial (two positions, 90° interactions). Both lone pairs occupy equatorial positions. The three F atoms are placed at: top axial, bottom axial, and one equatorial (right). The molecule looks like a "T"; bond angles $\approx 87.5^\circ$ (slightly compressed by lone-pair repulsion).

Why other options are wrong:

- Option A: Pyramidal geometry (AB_3E_1) has 3 bonds + 1 lone pair (e.g., NH_3); ClF_3 has 2 lone pairs.
- Option B: T-shaped with 2 lone pairs at equatorial positions — **correct**.
- Option C: Trigonal planar requires 0 lone pairs (e.g., BF_3 , 120° bond angles); ClF_3 has 2 lone pairs and bond angles $\approx 87.5^\circ$.
- Option D: Cl_3F is not a real chemical formula. ClF_3 is the correct formula and it is not linear.

Final Answer: ClF_3 ; T-shaped geometry (AB_3E_2 , 2 equatorial lone pairs) $\Rightarrow$ B

Answer: (B) [Go Back to Q 14](#)

Q15.

Solution

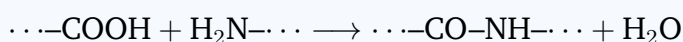
Concept — Condensation Polymers (Nylon-6,6): Nylon-6,6 is a polyamide formed by condensation of a dicarboxylic acid and a diamine. The name "6,6" refers to the six carbon atoms in each monomer unit.

Step 1 — Identifying the Monomers:

- **Adipic acid** (hexanedioic acid): $HOOC-(CH_2)_4-COOH \rightarrow$ **6 carbons**.
- **Hexamethylenediamine** (1,6-diaminohexane): $H_2N-(CH_2)_6-NH_2 \rightarrow$ **6 carbons**.

Hence the name Nylon-6,6 (diamine 6C, diacid 6C).

Step 2 — Linkage Type: Each $-COOH$ reacts with $-NH_2$, eliminating water and forming an **amide bond** ($-CO-NH-$):



The repeating amide units ($-CO-NH-$) allow extensive hydrogen bonding between chains, giving nylon high tensile strength and melting point.

Why other options are wrong:



- Option A: Caprolactam (ring-opening) gives **Nylon-6** (one monomer, 6-membered ring with 6 carbons/N), not Nylon-6,6.
- Option B: Adipic acid + ethylene glycol (2 carbons in diol) gives a **polyester** (–COO– linkage), not a polyamide; and ethylene glycol is not hexamethylenediamine.
- Option C: Adipic acid + hexamethylenediamine with amide linkage — **correct for Nylon-6,6**.
- Option D: Sebacic acid (decanedioic acid, **10** carbons) + hexamethylenediamine (6 carbons) gives **Nylon-6,10**, not Nylon-6,6.

Final Answer: Adipic acid + hexamethylenediamine; amide bond $\Rightarrow$ C

Answer: (C) [Go Back to Q 15](#)



Answer Key

Q	Ans	Q	Ans	Q	Ans	Q	Ans	Q	Ans
1	D	2	A	3	B	4	C	5	D
6	A	7	B	8	C	9	D	10	A
11	B	12	C	13	D	14	B	15	C

