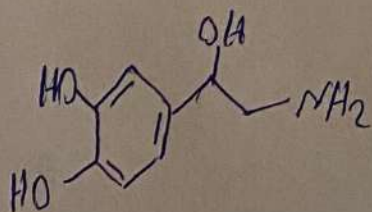
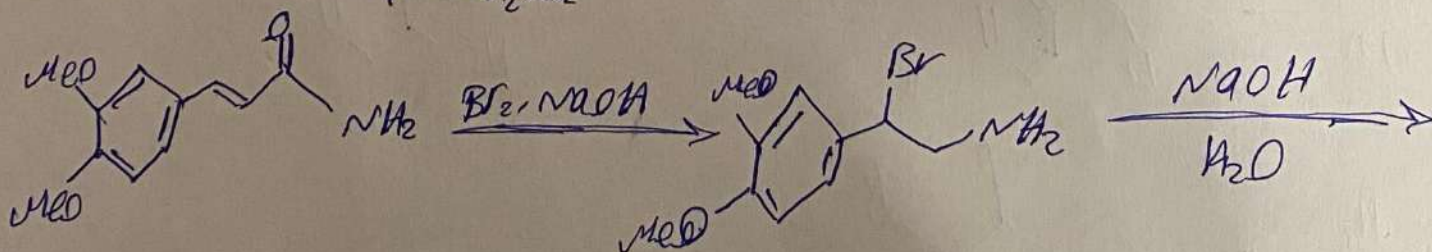
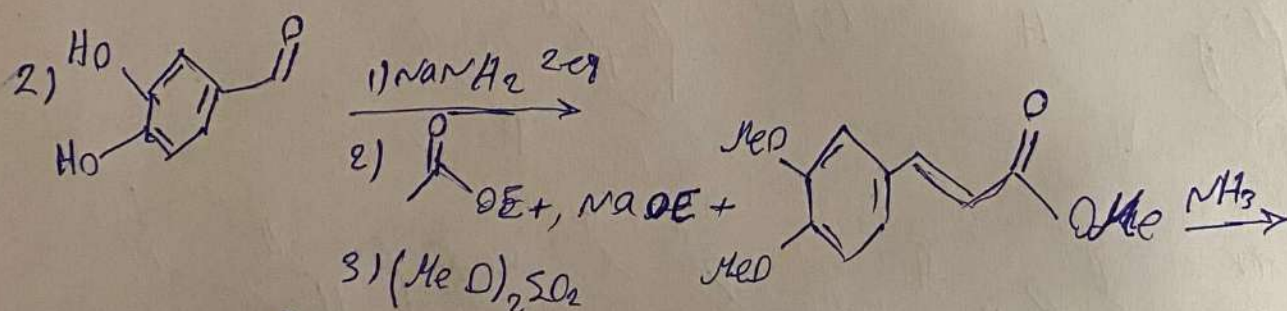
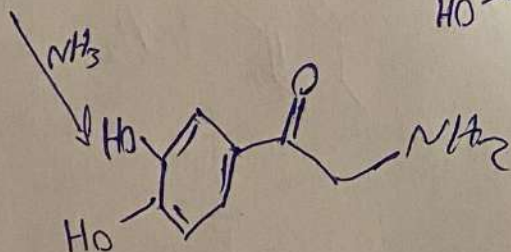
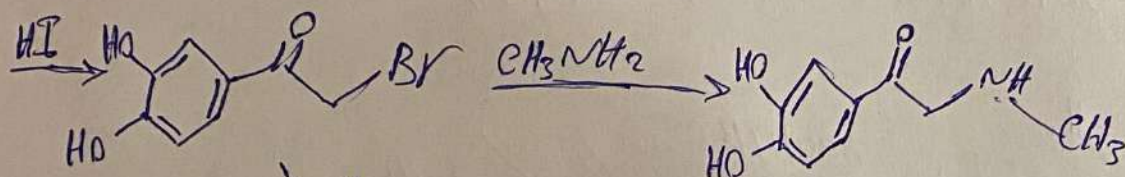
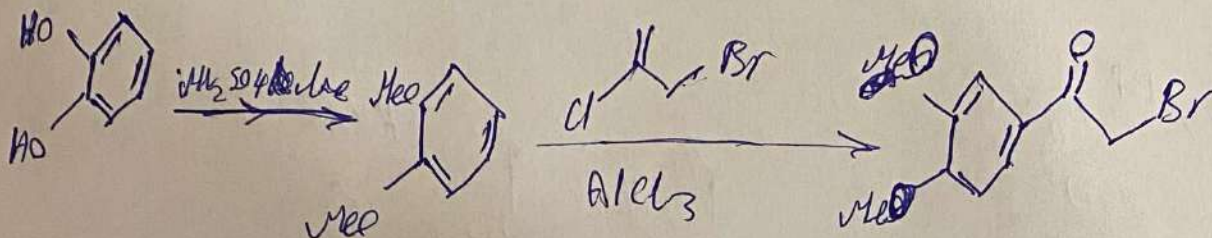
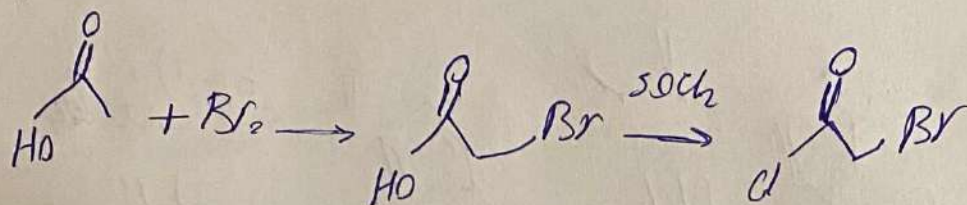
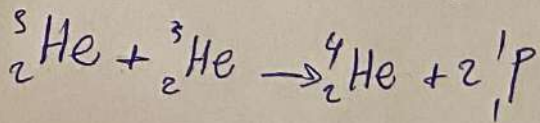
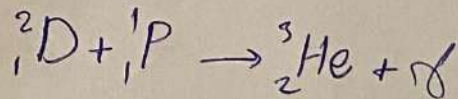
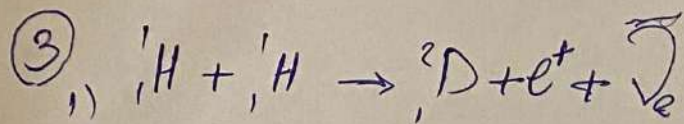


(4)

1)





2) $A = A_0 e^{-kt}$ $k = \frac{\ln 2}{T_{1/2}}$

$\frac{A}{A_0} = e^{-kt}$ $k = 1.216 \cdot 10^{-4} \frac{1}{\text{Year}}$

$0.175 = e^{-kt}$ Object A = 14333 Year

$0.25 = e^{-kt}$ Object B = 11400 Year

$Y = 14$

3) ${}_{40}^{130}\text{K} \xrightarrow{k_1} {}_{40}^{130}\text{Ar}$

${}_{40}^{130}\text{K} \xrightarrow{k_2} {}_{40}^{130}\text{Ca}$

$C = {}_{40}^{130}\text{K} e^{-k_1 t}$ $k_1 = 5.33 \cdot 10^{-6}$

$D = {}_{40}^{130}\text{K} e^{-k_2 t}$ $k_2 = 4.8 \cdot 10^{-9}$

$\frac{C}{D} = \frac{k_1}{k_2} = \frac{1}{9}$ $C = 3$
 $D = 27$
 $k = 8$

${}_{40}^{130}\text{K} = C + D = 3 + 27 = 30$

$\frac{{}_{40}^{130}\text{K}}{{}_{40}^{130}\text{K}_0} = e^{-(k_1 + k_2)t}$

$\ln \frac{3}{30} = -(k_1 + k_2)t$

$t = 292170376 \text{ Years}$

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(2)

1- 120°C - adsorbed water

1100°C - silanol groups

$$V_{H_2O} = \frac{0.8975 \text{ mg}}{18 \text{ mg/mmol}} = 0.02208 \text{ mmol}$$

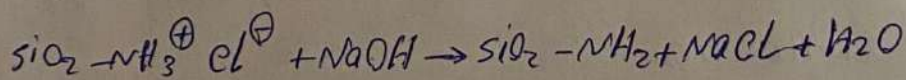
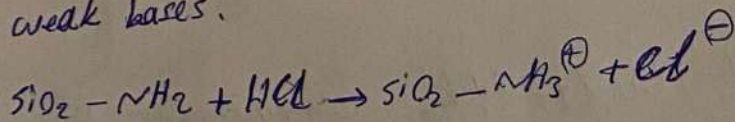
$$C_{H_2O} = \frac{0.02208}{14.2000 \cdot 10^{-3}} = 1.555 \frac{\text{mmol}}{\text{g}}$$

$$V_{\text{silanol}} = \frac{0.1561 \text{ mg}}{18 \text{ mg/mmol}} \cdot 2 = 0.01846 \text{ mmol}$$

$$S = 312.1 \frac{\text{m}^2}{\text{g}} \cdot (14.2000 \cdot 10^{-3} \text{ g}) = 4.439 \text{ m}^2$$

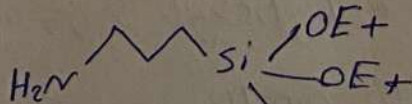
$$C_{\text{silanol}} = 18.46 \frac{\text{mmol}}{\text{m}^2} / 4.439 \text{ m}^2 = 4.159 \text{ mmol/m}^2$$

2- The reverse titration method is used to determine the concentration of weak bases.



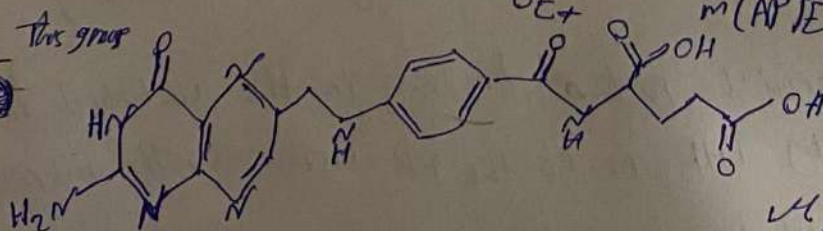
$$C_{\text{NH}_2} = \frac{C_{\text{NaOH}} V_{\text{NaOH}}}{m_{\text{sample}}} = \frac{0.01 \cdot 0.01}{0.1245 \text{ g}} = 1.526 \text{ mmol/g}$$

3- APTES

5.0g SiO₂

$$m(\text{APTES}) = 221 \frac{\text{mg}}{\text{mmol}} \cdot 7.183 = 1.586 \text{ g}$$

4- This group



hydrolysis reaction

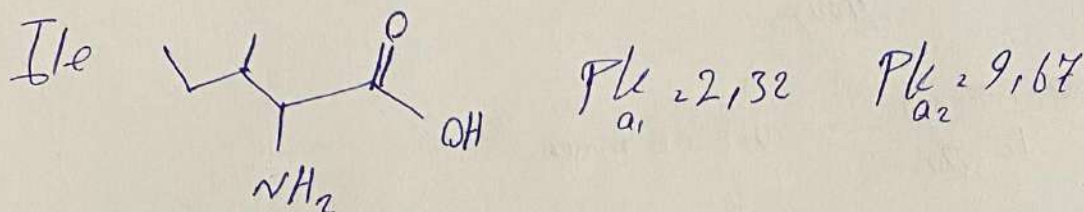
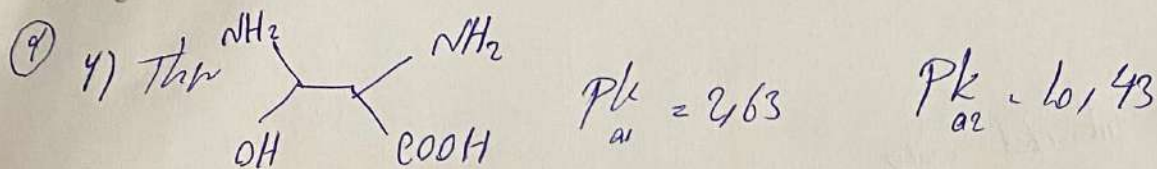
5-

$$\frac{m_{\text{con}}}{m_{\text{g}}} = \frac{0.5009}{1 \text{ g}}$$

$$M_{\text{FA}} = 441 \frac{\text{mg}}{\text{mmol}}$$

$$m_{\text{FA}} = 7.63 \cdot 0.003441 = 0.024 \text{ mg}$$

4



$$pK_{a1} < pH < pK_{a2}$$

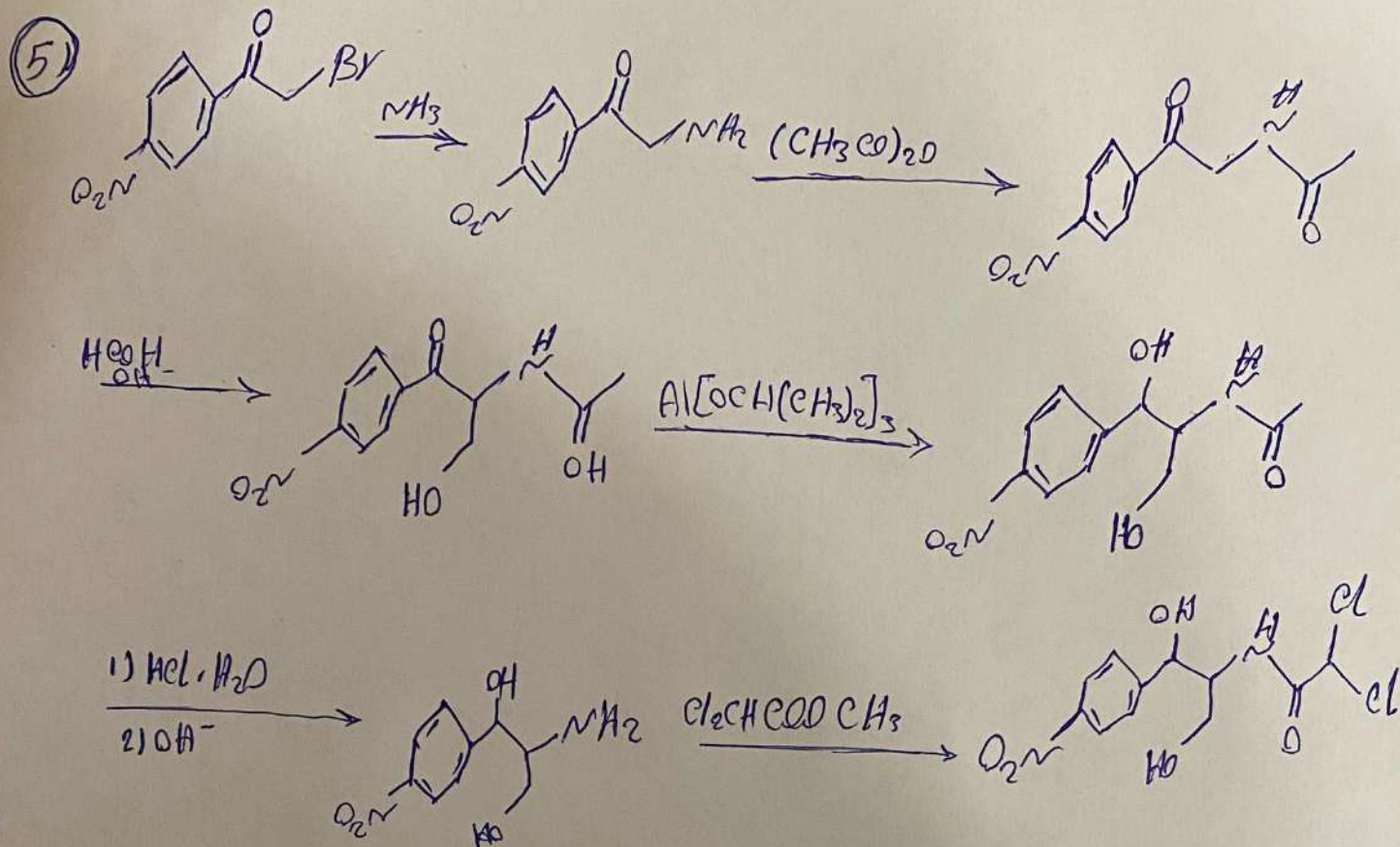
pH	7.0	4.0	9.6	12.0
Thr	<chem>NC(C(O)C(=O)O)N</chem>	<chem>[NH3+](C(C(=O)[O-])O)N</chem>	<chem>[NH3+](C(C(O)C(=O)[O-])O)N</chem>	<chem>[NH3+](C(C(O)C(=O)[O-])O)N</chem>
Ile	<chem>NC(C(C)C)C(=O)O</chem>	<chem>[NH3+](C(C(C)C)C(=O)[O-])</chem>	<chem>[NH3+](C(C(C)C)C(=O)[O-])</chem>	<chem>[NH3+](C(C(C)C)C(=O)[O-])</chem>

If $pH = 9.6$ every aminoacid $pK_{a1} < pH < pK_{a2}$ consequently the Predominant form is zwitter ion

2) The threonine residue in the protein binds to the drug by hydrogen bonds due to presence of a hydroxyl group, and the isoleucine residue with an iso butyl substituent is capable only of interacting with London forces

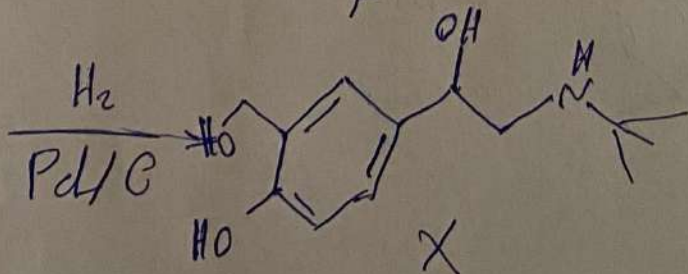
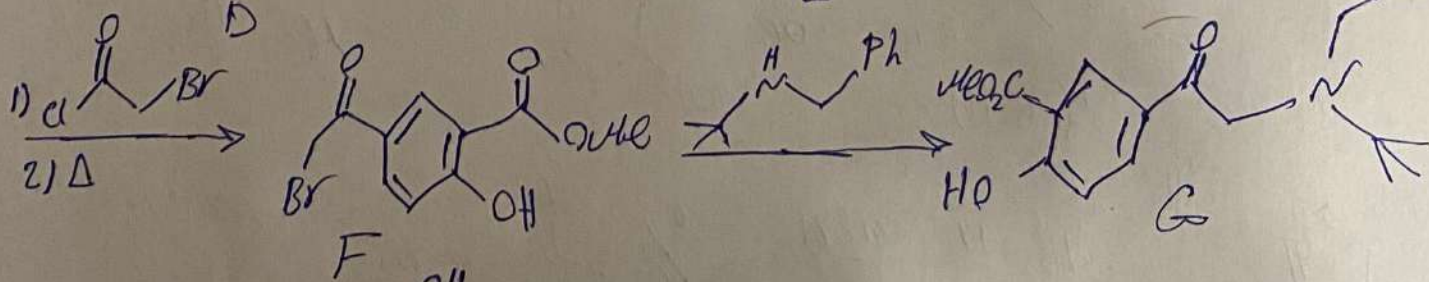
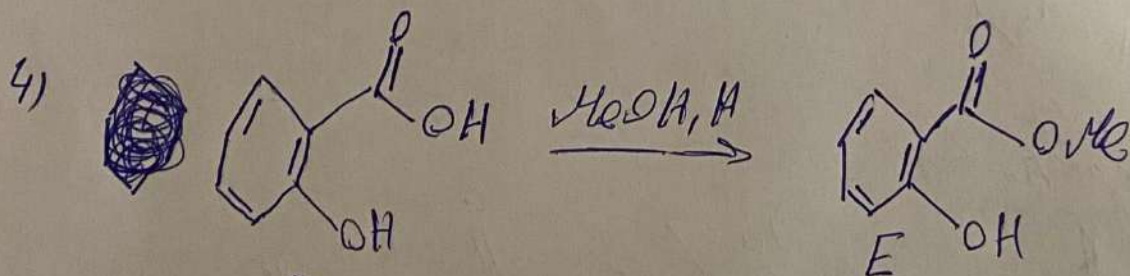
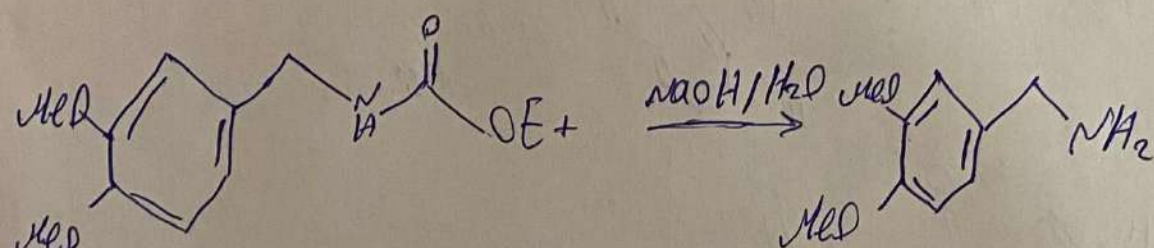
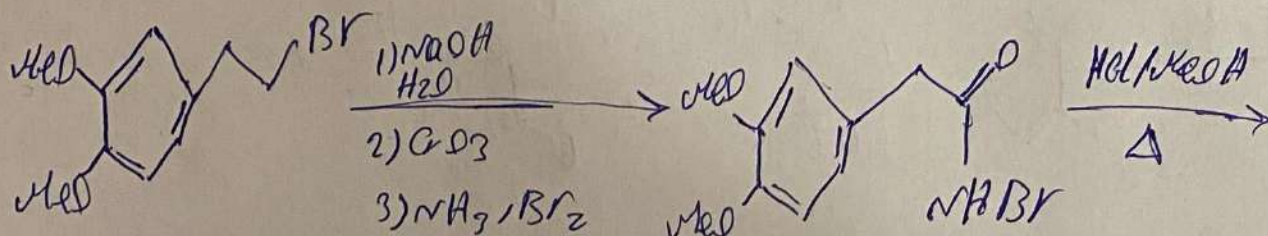
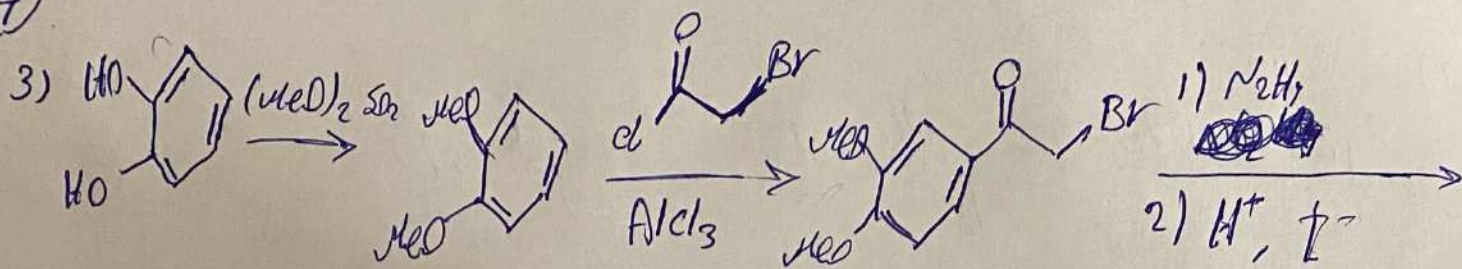
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3) in Fig 3 we see that an important part of binding in the mutated fragment is provided by THR 319. As the pH increases, the nitrogen on the preparation is deprotonated and the binding disappears.



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(4)



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