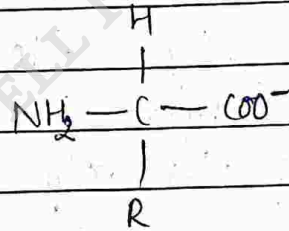
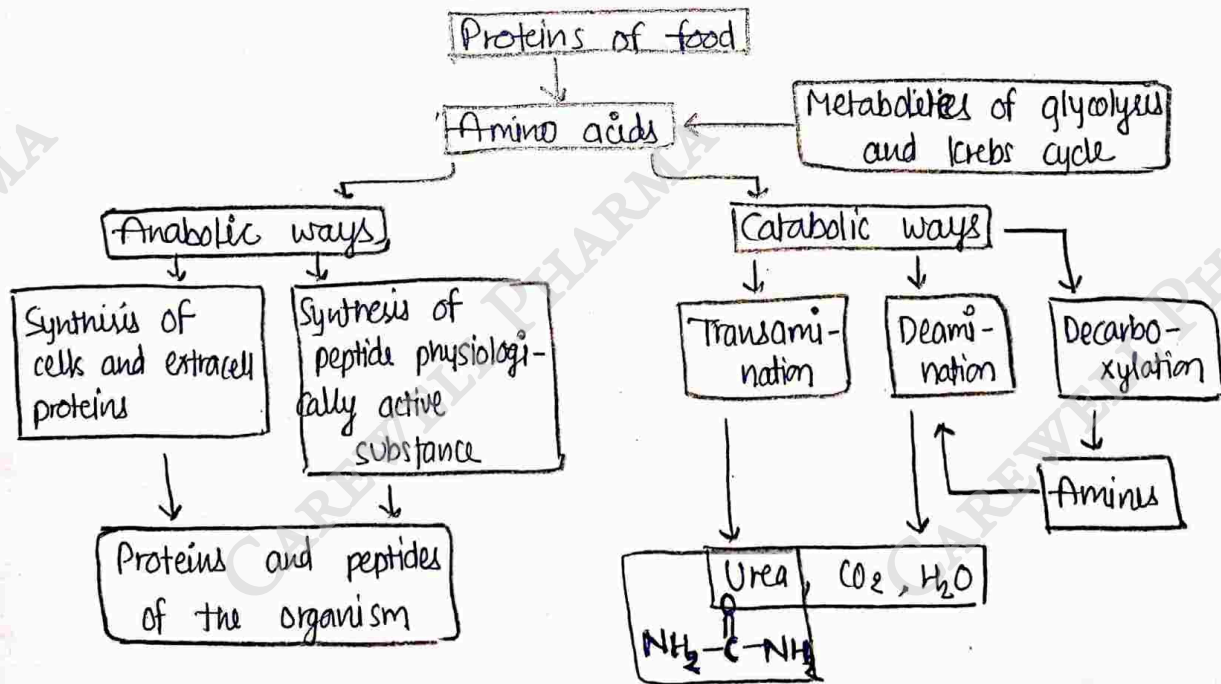


Amino Acid

An amino acid is an organic compound characterized by having a carboxyl group, amino group, and side chain attached to the central carbon atom.



General Pathway of Amino Acid Metabolism



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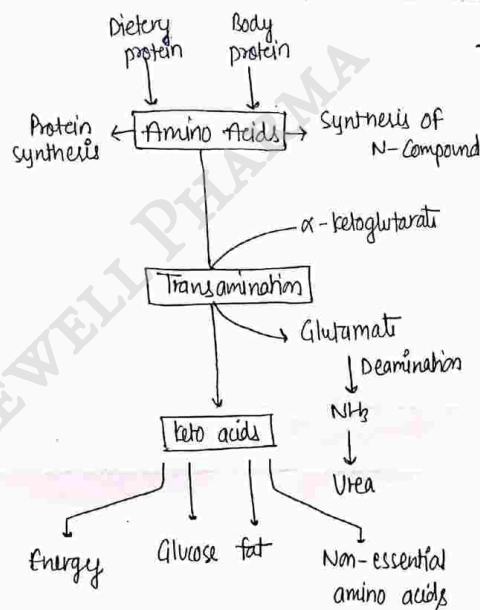
Amino Acid Metabolism

The amino acid undergoes certain common reaction like transamination followed by deamination for the liberation of ammonia. The amino group of the amino acid is utilized for the formation of urea which is an excretory end product of protein metabolism. The carbon skeleton of the amino acids is first converted to keto acids (by transamination) which meet one or more of the following fates.

- (i) Utilized to generate energy.
- (ii) Used for the synthesis of glucose.
- (iii) Diverted for the formation of fat or ketone bodies.
- (iv) Involved in the production of non-essential amino acids.

Transamination

The transfer of an amino ($-NH_2$) group from an amino acid to a keto group is known as transamination. This process involves the interconversion of a pair of amino acids and a pair of keto acid, catalysed by a group of enzyme called transaminases (recently aminotransferase).



•An overview of amino acid metabolism.

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Salient features of transamination

- 1) All transaminase require ~~pyridoxal~~ pyridoxal phosphat (PLP), a Coenzyme derived from Vitamin B₆.
- 2) Specific transaminases exist for each pair of amino acid and keto acids. However, only two - namely, aspartate transaminase and alanine transaminase - make a significant contribution for transamination.
- 3) There is no free NH₃ liberated, only the transfer of amino group occurs.
- 4) Transamination is reversible.

$$\begin{array}{ccc}
 R_1-CH-COO^- & & R_1-C-COO^- \\
 | & & || \\
 NH_3^+ & & O \\
 \text{Amino acid-I} & & \text{keto group acid-I} \\
 & \xleftrightarrow[\text{PLP}]{\text{Transaminase}} & \\
 & & \\
 R_2-C-COO^- & & R_2-C-COO^- \\
 || & & | \\
 O & & NH_3^+ \\
 \text{keto acid-II} & & \text{Amino acid-II}
 \end{array}$$

Transamination reactions

- 5) It is important for production of non-essential amino acids, as per the requirement of the cells. It involves catabolism & anabolism both.
- 6) Transamination diverts the excess amino acid towards energy generation.

• Deamination

The removal of amino group from the amino acids as NH₃ is deamination. Transamination involves only the shuffling of amino groups among the amino acids. On the other hand, deamination result in the liberation of ammonia for urea synthesis. Simultaneously, the carbon skeleton of amino acid is converted of keto

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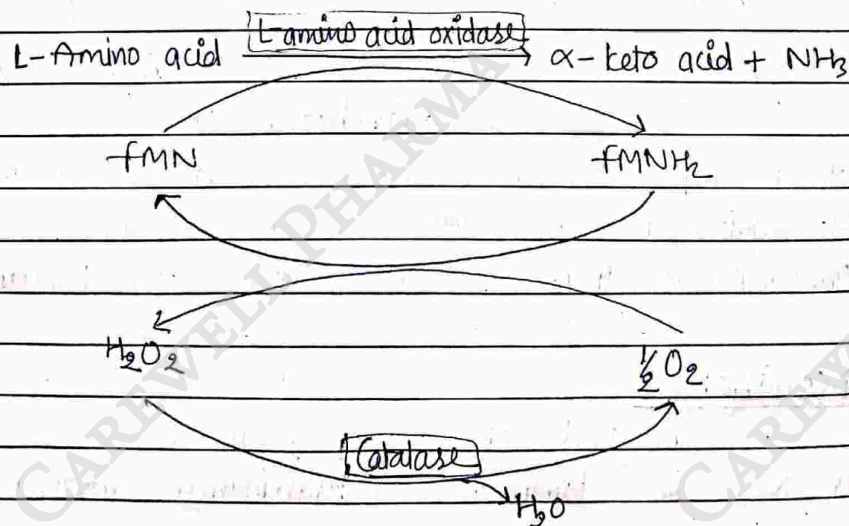
acids. Deamination may be either oxidative or non-oxidative

I. Oxidative deamination

It is the liberation of free ammonia from the amino group of amino acids coupled with oxidation. This takes place mostly in liver & kidney. The purpose of oxidative deamination is to provide NH_3 for urea synthesis and α -keto acids for a variety of reactions, including energy generation.

• Oxidative deamination by amino acid oxidases

L-Amino acid oxidase and D-Amino acid oxidase and flavoproteins, possessing FMN and FAD, respectively. They act on the corresponding amino acid (L or D) to produce α -keto acids and NH_3 . In this reaction, oxygen is reduced to H_2O_2 , which is later decomposed by catalase.



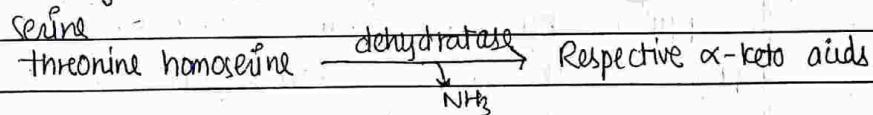
∴ Oxidative deamination of amino acids

The activity of L-amino acid oxidase is much low while that D-amino acid oxidase is high in tissue (mostly liver and kidney). L-Amino acid oxidase does not act on glycine and dicarboxylic acids. This enzyme, due to its very low activity, does not appear to play any significant role in the amino acid metabolism.

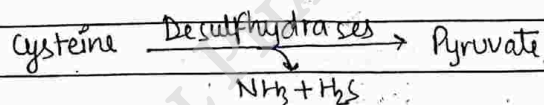
II. Non-oxidative deamination

Some of the amino acids can be deaminated to liberate NH_3 without undergoing oxidation.

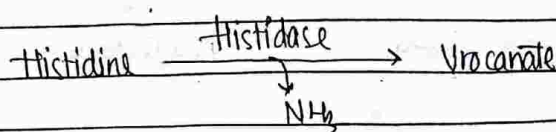
a) Amino acid dehydratases: Serine, threonine and homoserine are the hydroxy amino acids. They undergo non-oxidative deamination catalysed by PLP-dependent dehydratases (dehydratases).



b) Amino acid desulfhydrases: The sulfur amino acids, namely cysteine and homocysteine, undergo deamination coupled with desulfhydration to give keto acids.



c) Deamination of histidine: The enzyme histidase acts on histidine to liberate NH_3 by a non-oxidative deamination process.



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Metabolism of ammonia

- a) formation of ammonia \rightarrow The production of NH_3 occurs from transamination & deamination of amino acids.
- b) Transport & storage of NH_3 \rightarrow The maintenance of NH_3 conc in circulation is done by its transportation between various tissue and the liver mostly occurs in the form of glutamine or alanine & not as free NH_3 .
- c) function of NH_3 \rightarrow In spite of being waste product of nitrogenous metabolism, it is involved directly for the synthesis of many compounds in the body like purines, pyrimidine etc.
- d) Disposal of NH_3 \rightarrow Urea is non-toxic and soluble compound & hence easily excreted in humans.

Toxicity of Ammonia

Ammonia when accumulates in the body results in

↓

Speech slurring & blurring of vision.

↓

which leads to coma, & finally death.

Hyperammonia

Elevation of blood NH_3 level

↓

Cause hepatic coma & mental retardation.

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UREA CYCLE

Urea is the end product of protein metabolism (amino acid metabolism). The nitrogen of amino acids, converted to ammonia, as described is toxic to the body. It is converted to urea and detoxified. As such, urea accounts for 80-90% of the nitrogen containing substance excreted in urine.

Urea is synthesized in liver and transported to kidney for excretion in urine. Urea cycle is the first metabolic cycle that was elucidated by Hans Krebs and Kurt Henseleit (1932), hence it is known as Krebs-Henseleit cycle.

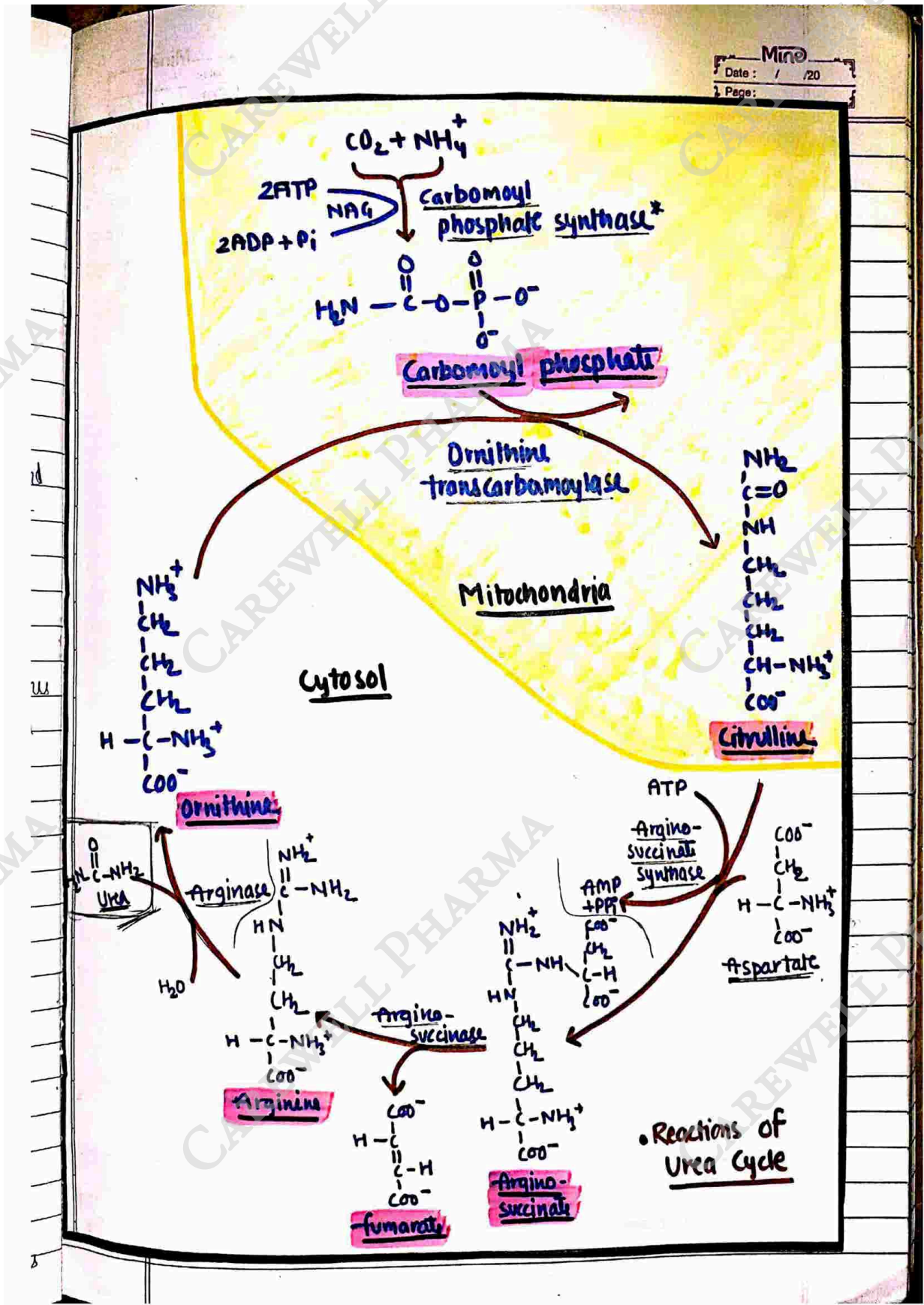
Urea has two amino ($-NH_2$) groups, one derived from NH_3 and the other from aspartate. Carbon atom is supplied by CO_2 . Urea synthesis is a five-step cyclic process, with five distinct enzymes. The first two enzymes are present in mitochondria while the rest are localized in cytosol.

```

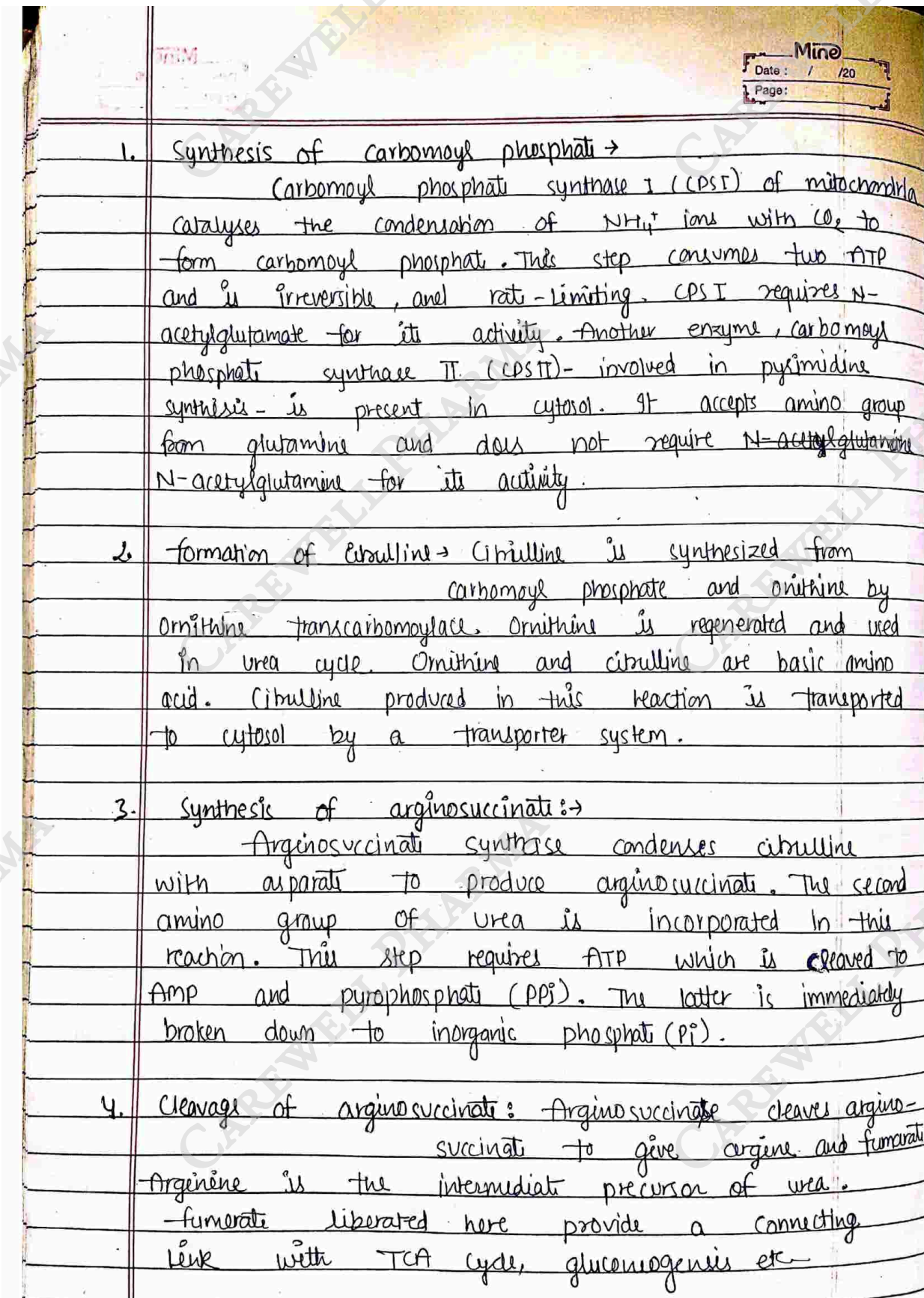
graph TD
    A["CO2 + NH4+"] --> B["Carbamoyl phosphate"]
    B --> C["Citrulline"]
    C --> D["Argininosuccinate"]
    D --> E["Arginine"]
    E --> F["Urea"]
    F --> G["Ornithine"]
    G --> C
    H["Aspartate (R'-NH2)"] --> D
    I["Fumarate"] --> D
  
```

• Outline of urea cycle (In the synthesis of urea one amino group comes from ammonium ion while the other is from aspartate).

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5. formation of urea - Arginase is the fifth and final enzyme that cleaves arginine to yield urea and ornithine. Ornithine, so regenerated, enters mitochondria for its reuse in the urea cycle. Arginase is activated by Co^{2+} and Mn^{2+} . Ornithine and lysine compete with arginine (competitive inhibition). Arginase is mostly found in the liver, while the rest of the enzymes (four) of urea cycle are also present in other tissue for this reason, arginine synthesis may occur to varying degree in many tissue. But only the liver can ultimately produce urea.

- Overall reactions and energetics
The urea cycle is irreversible and consumes 4 ATP. Two ATP are utilized for the synthesis of carbamoyl phosphate. One ATP is converted to AMP and P_i to produce arginosuccinate which equal to 2 ATP. Hence 4 ATP are actually consumed.

$$\text{NH}_4^+ + \text{CO}_2 + \text{Aspartate} + 3\text{ATP} \longrightarrow \text{Urea} + \text{fumarate} + 2\text{ADP} + 2\text{P}_i + \text{AMP} + \text{P}_i$$

- Regulation of urea cycle.
The first reaction catalysed by carbamoyl phosphate synthase I (CPS I) is rate-limiting reaction or committed step in urea synthesis. CPS I is allosterically activated by N-acetylglutamate (NAG). It is synthesized from glutamate and acetyl-CoA by synthase and degraded by a hydrolase.

The rate of urea synthesis in liver is correlated with the concentration of N-acetylglutamate. High concentrations of arginine increase NAG. The consumption

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of a protein-rich meal increases the level of NAG in liver, leading to enhanced urea synthesis.

Carbamoyl phosphati synthase I and glutamati dehydrogenase are localized in the in the mitochondria. They coordinate with each other in the formation of NH_2 , and its utilization for the synthesis of carbamoyl phosphati. The remaining four enzymes of urea cycle are mostly controlled by the concentration of their respective substrati.

```

graph TD
    A[Acetyl CoA] --> B[NAG synthase]
    C[Glutamate] --> B
    B --> D[N-Acetylglutamate]
    B --> E[CoA]
    D --> F[NAG hydrolase]
    F --> G[Acetate]
    F --> H[Glutamate]
  
```

- formation and degradation of N-acetylglutamate.

• Decarboxylation → It is the removal of carbon dioxide from amino acids with formation of amines.

$$\begin{array}{ccc}
 \text{R}-\text{CH}-\text{COOH} & \xrightarrow{\text{CO}_2} & \text{R}-\text{CH}_2-\text{NH}_2 \\
 | & & | \\
 \text{NH}_2 & & \text{amine}
 \end{array}$$

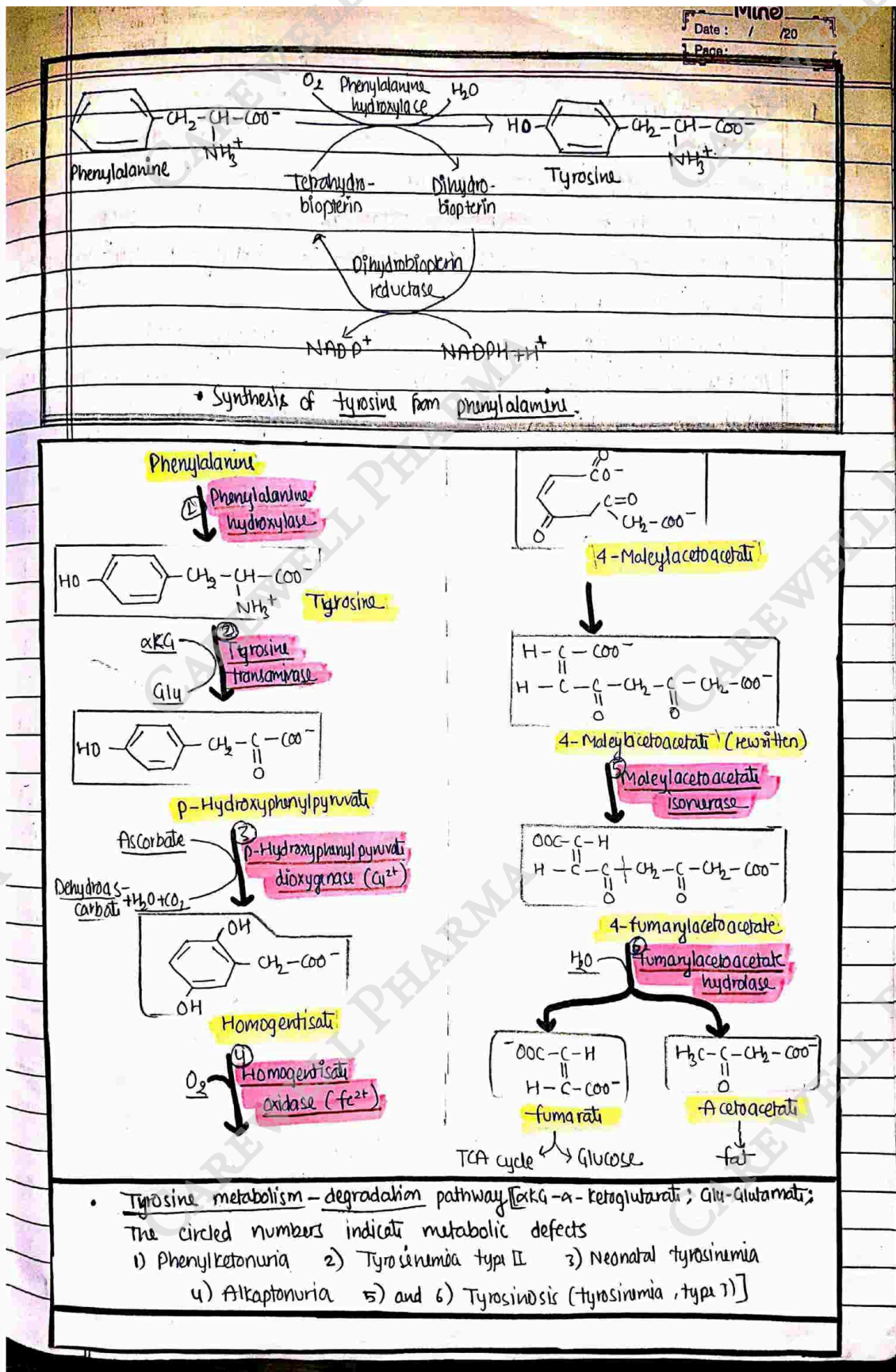
• Degradation (Catabolism) of Tyrosine (Phenylalanine)

The metabolism of phenylalanine and tyrosine is considered together. The sequence of the reaction in the degradation of these amino acids,

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1. As phenylalanine is converted to α -tyrosine, a single pathway is responsible for degradation of both these amino acids, which occurs mostly in liver.
 2. Tyrosine first undergoes transamination to give p -hydroxyphenylpyruvate. This reaction is catalyzed by tyrosine transaminase (PLP dependent).
 3. p -Hydroxyphenylpyruvate hydroxylase (or dioxygenase) is a copper-containing enzyme. It catalyses oxidative decarboxylation as well as hydroxylation of the phenyl ring of p -hydroxyphenylpyruvate to produce homogentisate. This rxn involves a shift in hydroxyl group from para position to meta position, and incorporates a new hydroxyl group at para position. This step in tyrosine metabolism required ascorbic acid.
 4. Homogentisate oxidase (iron metalloprotein) cleaves the benzene ring of homogentisate to form 4-maleylacetoacetate. Molecular oxygen is required for this rxn to break the aromatic ring.
 5. Maleylacetoacetate undergoes isomerization to form 4-fumarylacetoacetate and this rxn is catalyzed by maleylacetoacetate isomerase to form 4-fumarylacetoacetate.
 6. Fumaryl acetoacetase (fumaryl acetoacetate hydrolase) brings about the hydrolysis of 4-fumaryl acetoacetate to liberate fumarate and acetoacetate.
- Fumarate is an intermediate of citric acid cycle and can also serve as precursor for gluconeogenesis. Acetoacetate is a ketone body from which fat can be synthesized. Phenylalanine and tyrosine are therefore, both glucogenic and ketogenic.
- The inborn errors of phenylalanine and tyrosine metabolism are indicated in ~~grey~~ pathway.

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Disorders of Tyrosine & Phenylalanine Metabolism

Disorder	Enzyme defective	Cause symptoms
1) Phenylketonuria	a) Phenylalanine hydroxylase b) Dihydrobiopterin reductase	Growth failure, seizures, mental retardation
2) Tyrosinemia Type II	Tyrosine transaminase	Eyelesion, Dermatitis
3) Neonatal Tyrosinemia	p-Hydroxyphenyl pyruvate dioxygenase	Mental retardation, Growth retardation
4) Alkaptonuria (Black urine disease)	Homogentisate oxidase	Urine resemble colour of coke. Dystonia's Arthritis.
5) Tyrosinosis Tyrosinemia Type-I	a) fumarylacetoacetate hydroxylase b) Maleylacetoacetate isomerase	Liver failure, Diarrhea, Vomiting polyneuropathy
6) Albinism	Tyrosinase responsible for synthesis of melanin	Photophobia

Phenylalanine (Phe, F) and tyrosine (Tyr, R) are structurally related aromatic amino acids. Phenylalanine is an essential amino acid while tyrosine is non-essential. Besides its incorporation into proteins, the only function of phenylalanine is its conversion to tyrosine for this reason, ingestion of tyrosine can reduce the dietary requirement of phenylalanine. This phenomena is referred to as 'sparing action' of tyrosine on phenylalanine. The predominant metabolism of phenylalanine occurs through tyrosine. Tyrosine is incorporated into proteins and is involved in the synthesis of a variety of biological important compounds - epinephrine, norepinephrine, dopamine, thyroid hormones - and the pigment melanin.

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• Disorders of Tyrosine (Phenylalanine) metabolism.

i) Phenylketonuria → (PKU) It is the most common metabolic disorder in amino acid metabolism. The incidence of PKU is 1 in 10,000 births. It is due to deficiency of the hepatic enzymes, phenylalanine hydroxylase, caused by an autosomal recessive gene.

— Phenylalanine metabolism in PKU —

Phenylketonuria primarily causes the accumulation of phenylalanine in tissues and blood, and results in its increased excretion in urine.

Due to disturbance in the routine metabolism, phenylalanine is diverted to alternate pathways, resulting in the excessive production of phenylpyruvate, phenylacetate, phenyllactate and phenyl glutamine. All these metabolites are excreted in urine in high concentration in PKU.

Phenylacetate gives the urine a mousy odour.

ii) Tyrosinemia type II

This disorder also known as Riechner-hanhart syndrome, is due to a defect in the enzyme tyrosine transaminase. The result is a blockade in the routine degradation pathway of tyrosine. Accumulation and excretion of tyrosine and its metabolites— namely p-hydroxyphenyl pyruvate, p-hydroxyphenyllactate, p-hydroxyphenylacetate, N-acetyltyrosine— and tyramine are observed.

Tyrosinemia type II is characterized by skin (dermatitis) and eyes lesions and rarely mental retardation. A disturbed self-coordination is seen in these patients.

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iii) Alkaptonuria (Black Urine disease)

Enzyme defect → The defective enzyme in alkaptonuria is homogentisate oxidase in tyrosine metabolism. Homogentisate accumulates in tissue and blood, and is excreted into urine. Homogentisate, on standing, gets oxidized to the corresponding quinones, which polymerize to give black or brown colour. For this reason, the urine of alkaptonuria patients resembles coke in colour.

iv) Albinism →

Albinism (Greek: albino-white) is an inborn error due to the lack of synthesis of the pigment melanin. It is an autosomal recessive disorder with a frequency of 1 in 20,000.

• Hypopigmentation

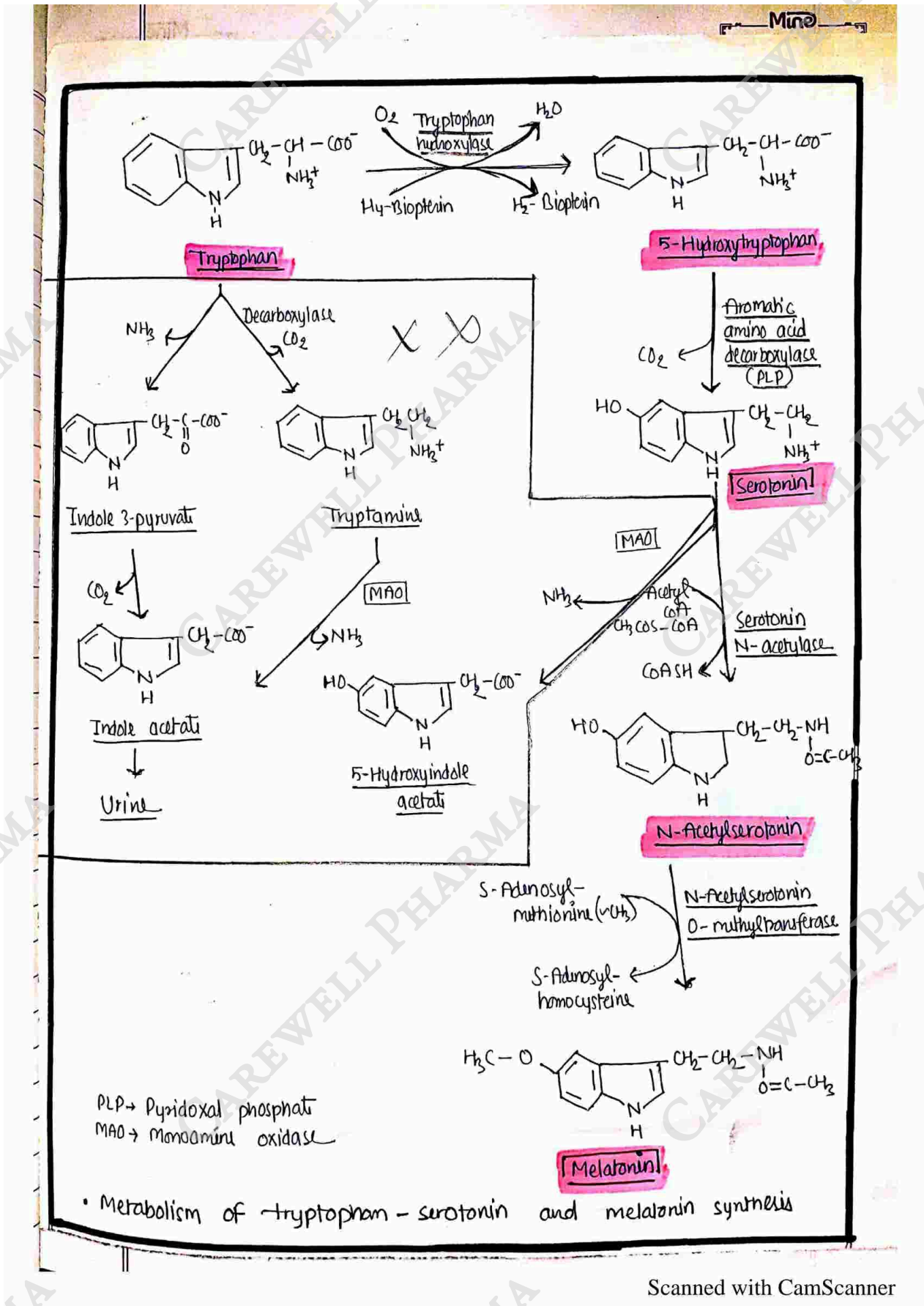
A good example of diffuse hypopigmentation is oculocutaneous albinism which is mostly due to mutations in the tyrosinase gene. The degree of hypopigmentation depends upon the type and severity of mutated genes.

⇒ Synthesis or Significance of biological substance

i) Serotonin pathway

Serotonin or 5-Hydroxytryptamine is a neurotransmitter, synthesized from tryptophan.

— Synthesis of serotonin: In mammals, the largest amount of serotonin is synthesized in the intestinal cells.



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Reactions of serotonin pathway

— Serotonin is a neurotransmitter and performs a variety of functions.

- It is a powerful vasoconstrictor and result in smooth muscle contraction in bronchioles and arterioles.
- It is closely involved in the regulation of cerebral activity (excitation).
- Serotonin controls the behavioural patterns, sleep, blood pressure and body temperature.
- It evokes the release of peptide hormones from gastrointestinal tract.
- It is also necessary for the motility of GIT (peristalsis).

→ When tryptophan hydroxylase enzyme hydrolysed tryptophan to 5-Hydroxytryptophan, further Amino acid decarboxylase enzyme act on it and it produced a Serotonin.

→ Then Serotonin N-acetylase (the rate limiting enzyme) acted on serotonin and give N-Acetylserotonin. The latter undergoes methylation in the presence of enzyme N-Acetylserotonin omethyl transferase, it give melatonin.

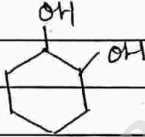
- Melatonin is a hormone, mostly synthesized by the pineal gland.
- It is involved in circadian rhythms or diurnal variations (24 hr cyclic process) of the body. It plays a significant role in sleep and wake process.
- Melatonin inhibits the production of melanocyte stimulating hormone (MSH) and adrenocorticotropic hormone (ACTH).
- It also performs a neurotransmitter function.

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• Biosynthesis of catecholamines (dopamine, norepinephrine, epinephrine).

The name catechol refers to the dihydroxylated phenyl ring. The amine derivatives of catechol are called catecholamines.



Tyrosine is the precursor for the synthesis of catecholamines, namely dopamine, norepinephrine (noradrenalin) and epinephrine (adrenalin).

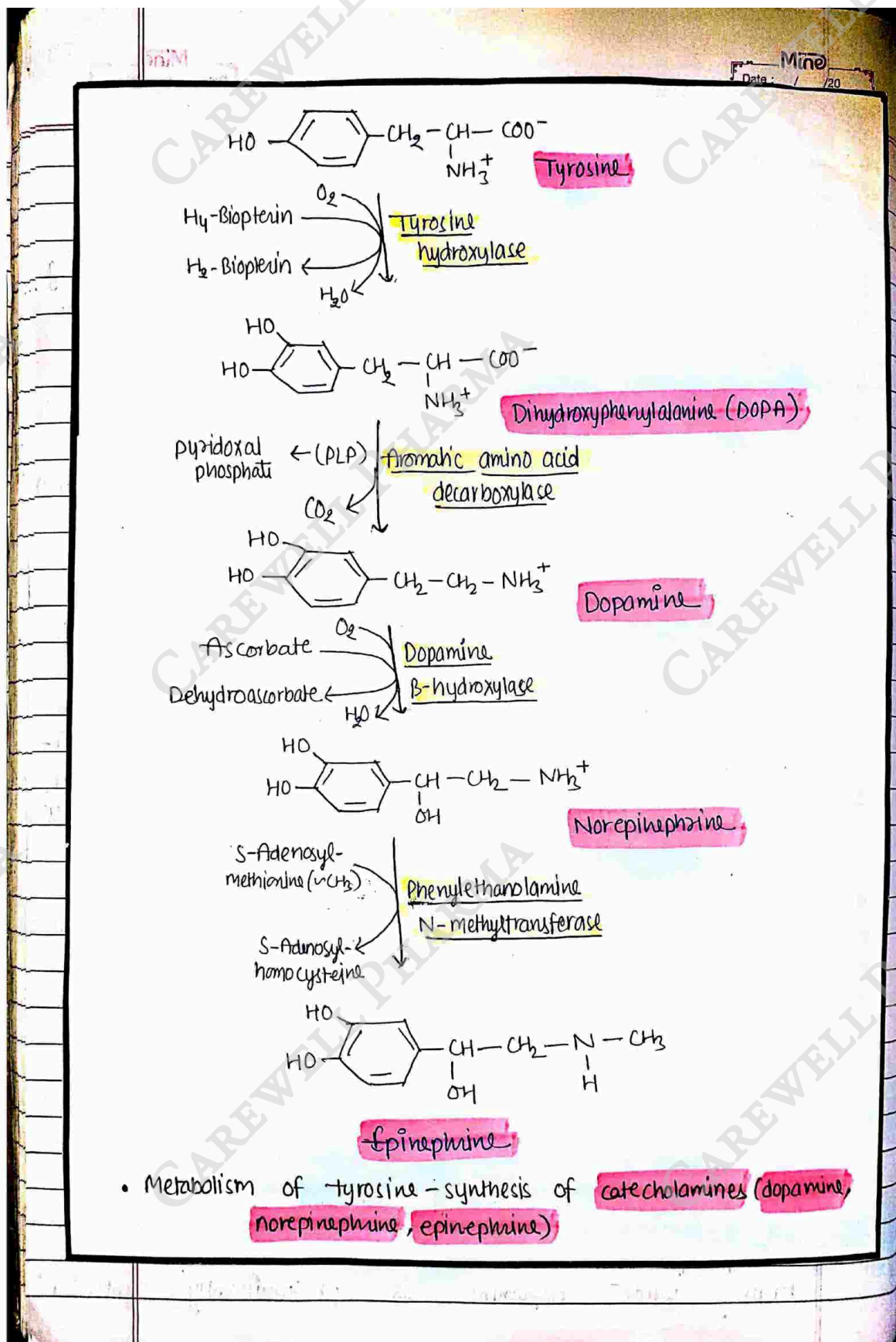
The conversion of tyrosine to catecholamines occurs in adrenal medulla and central nervous system involving the following reactions (in pathway on next page).

→ Tyrosine is hydroxylated to 3,4-dihydroxyphenylalanine (DOPA) by tyrosine hydroxylase. This enzyme catalyzes the rate limiting reaction and requires tetrahydrobiopterin as coenzyme (like phenylalanine hydroxylase). In contrast to this enzyme, tyrosinase present in melanocytes convert tyrosine to DOPA. Hence, two different enzymes system exist to convert tyrosine to DOPA.

→ DOPA undergoes PLP-dependent decarboxylation to give dopamine which, in turn, is hydroxylated to produce norepinephrine. Methylation of norepinephrine by S-adenosyl methionine gives epinephrine. The difference b/w epinephrine is only a methyl group (remember that norepinephrine has no methyl group).

→ There exists tissue specificity in the formation of catecholamines. In adrenal medulla, synthesis of the hormones, norepinephrine and epinephrine is prominent. Norepinephrine is produced in certain areas of the brain while dopamine is predominantly synthesized

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in substantia nigra and coeruleus of brain.

- function of catecholamines → Norepinephrine and epinephrine regulate carbohydrates and lipid metabolisms. They stimulate the degradation of triacylglycerol and glycogen. They cause an increase in the blood pressure. Dopamine and norepinephrine serves as neurotransmitters in the brain and autonomous nervous system.

• CATABOLISM OF HEME

```

    graph TD
      Hemoglobin --> Globin
      Hemoglobin --> Heme
      Heme -- "Heme oxygenase" --> Biliverdin
      Biliverdin -- "Biliverdin reductase" --> Bilirubin
      Bilirubin --> Urobilinogen_in_kidney[Urobilinogen in kidney]
      Bilirubin --> Bilirubin_in_intestine[Bilirubin in intestine]
      Urobilinogen_in_kidney --> Urobilin
      Urobilin --> Urobilin_excreted[Urobilin excreted in urine]
      Bilirubin_in_intestine --> Bilirubin_diglyconide[Bilirubin diglyconide in intestine]
      Bilirubin_diglyconide --> Urobilinogen_in_intestine[Urobilinogen in intestine]
      Urobilinogen_in_intestine --> Stercobilinogen
      Stercobilinogen --> Stercobilin
      Stercobilin --> Stercobilin_excreted[Stercobilin excreted in stools/feces]
  
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•	<p><u>Hyperbilirubinemia</u> → The condition having high bilirubin level is called Hyperbilirubinemia.</p> <p>The cause of hyperbilirubinemia are Intrahepatic cholestasis and extrahepatic obstruction of biliary tract which prevent bilirubin from moving to intestine.</p>	
•	<p><u>Jaundice</u> → It is caused by build up of bilirubin (waste material in blood). An inflamed liver or obstructed bile duct can lead to jaundice. Symptoms</p> <p>Symptoms → Yellowing of eyes, Dark urine itching</p>	
	<u>Types</u>	<u>Symptoms</u>
1)	Pre Hepatic jaundice	Abdominal pain, fever, dark urine
2)	Hepatic Jaundice	Bloody Nose, weakness, fever, appetite
3)	Post hepatic Jaundice	Diarrhea, fever, Abdominal swelling, weightless

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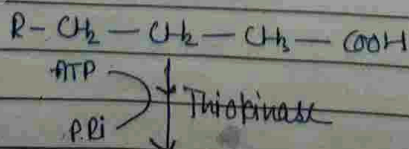
LIPID METABOLISM

A lipid is a chemically defined as a substance that include insoluble in water and soluble in alcohol ether and chloroform. Lipid are an important component are living cells together with carbohydrates, and (protein) lipid are the most abundant lipid comprises 85-90% of body lipids. Most of the triglyceride are stored in the adipose tissue and served as energy receiver wise of the body.

- fatty acid oxidation → The fatty acid in the body are mostly in the oxidation may be defined as the oxidation of fatty acid on the carbon atom. This results in the sequential removal of carbon compound, β -oxidation involves (acetyl CoA).
- i) Activation of fatty acids occurs in the cytosol.
 - ii) Transport of fatty acid into mitochondria.
 - iii) β -oxidation proper in the mitochondrial matrix.

fatty acid activation :-

fatty acid are activated to acyl CoA by thiolase or acyl CoA synthase the rxn occurs in 2 steps and require ATP co-enzyme A and magnesium enzyme fatty acid react with ATP to form acyl adenylate which then combine with co-enzyme A to produce acyl CoA.



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$$\begin{array}{c}
 \downarrow \\
 R-CH_2-CH_2-CH_2-C(=O)-AMP \\
 \text{CoASH} \quad \text{Mg}^{2+} \\
 \text{AMP} \quad \downarrow \\
 R-CH_2-CH_2-CH_2-C(=O)-S-CoA \quad \text{Acetyl CoA}
 \end{array}$$

Transport of acyl CoA into mitochondria -
 The inner mitochondrial membrane is impermeable to fatty acid a specialized carnitine carrier system operates to transport activated fatty acid from cytosol to mitochondria.

Cytoplasm	Mitochondria
$R-C(=O)-S-CoA$ Carnitine Acyl transferase I $R-C(=O)-Carnitine$	$R-C(=O)-S-CoA$ Carnitine Carnitine and transferase II $R-C(=O)-Carnitine$

→ Oxidation proper in mitochondrial matrix
 Each cycle of β -oxidation liberating 2 carbon unit acetyl CoA occurs in sequence of rxn

- oxidation
- hydration
- oxidation
- cleavage

$$\begin{array}{c}
 R-CH_2-CH_2-CH_2-C(=O)-S-CoA \\
 \text{FAD} \quad \text{Acyl CoA dehydrogenase} \quad \text{① - oxidation} \\
 \text{FADH}_2 \quad \downarrow \\
 R-CH_2-CH=CH-C(=O)-S-CoA \quad \text{a}^2 \text{ trans- enoyl CoA}
 \end{array}$$

H_2O } Enoyl CoA hydrolase ② hydration

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$$\begin{array}{c}
 \text{OH} \\
 | \\
 \text{R}-\text{CH}_2-\text{CH}-\text{CH}_2-\overset{\text{O}}{\parallel}{\text{C}}-\text{SCoA} \\
 \downarrow \text{NAD}^+ \\
 \text{NADH} + \text{H}^+ \quad \text{3-hydroxyacyl CoA dehydrogenase} \\
 \text{R}-\text{CH}_2-\overset{\text{O}}{\parallel}{\text{C}}-\text{CH}_2-\overset{\text{O}}{\parallel}{\text{C}}-\text{SCoA} \\
 \beta\text{-ketoacyl CoA}
 \end{array}$$

③ oxidation

$$\begin{array}{c}
 \text{CoASH} \\
 \downarrow \\
 \text{R}-\text{CH}_2-\overset{\text{O}}{\parallel}{\text{C}}-\text{SCoA} + \text{CH}_3-\overset{\text{O}}{\parallel}{\text{C}}-\text{SCoA}
 \end{array}$$

④ cleavage

Polmitoyl CoA undergoes

7 cyclic of β -oxidation to give acetyl CoA

Polmitoyl CoA (16 carbon)

$\text{NAD}^+ \rightarrow \text{NADH}$
 $\text{FAD} \rightarrow \text{FADH}_2$
 $\text{CoA} \rightarrow \text{Acetyl CoA}$

35 ATP

8 Acetyl CoA (2c)

Energy yield from β -oxidation = 7 cycle \times 5 = 35

Energy from 8 acetyl CoA = 8 \times 9 = 72

Energy from Citric acid cycle = 12 ATP \times 8 = 96

Total = 35 + 72 + 96 = 203

Energy utilised for activation = 2

Net yield = 203 - 2 = 201

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* Ketogenesis → Synthesis of ketone bodies occurs in the liver. The enzymes where ketones synthesis are located in mitochondrial matrix.

$$\begin{array}{c}
 \text{CH}_3-\overset{\text{O}}{\parallel}{\text{C}}-\text{S}(\text{CoA}) + \text{CH}_3-\overset{\text{O}}{\parallel}{\text{C}}-\text{S}(\text{CoA}) \\
 \downarrow \text{CoA} \quad \beta\text{-ketothiolase} \\
 \text{CH}_3-\overset{\text{O}}{\parallel}{\text{C}}-\text{CH}_2-\overset{\text{O}}{\parallel}{\text{C}}-\text{S}(\text{CoA}) \\
 \downarrow \text{CoA} \quad \text{Acetoacetyl CoA} \\
 \text{S}(\text{CoA})-\text{CH}_2-\overset{\text{O}}{\parallel}{\text{C}}-\text{CH}_2-\overset{\text{O}}{\parallel}{\text{C}}-\text{S}(\text{CoA}) \\
 \downarrow \text{HMG CoA lyase} \\
 \text{O}=\text{C}-\text{CH}_2-\overset{\text{O}}{\parallel}{\text{C}}-\text{CH}_2-\overset{\text{O}}{\parallel}{\text{C}}-\text{S}(\text{CoA}) \\
 \downarrow \text{Spontaneous} \\
 \text{CH}_3-\overset{\text{O}}{\parallel}{\text{C}}-\text{CH}_3 \quad \text{Acetone}
 \end{array}$$

β-hydroxy β-methyl glutarate CoA

Acetoacetyl CoA

β-hydroxybutyrate

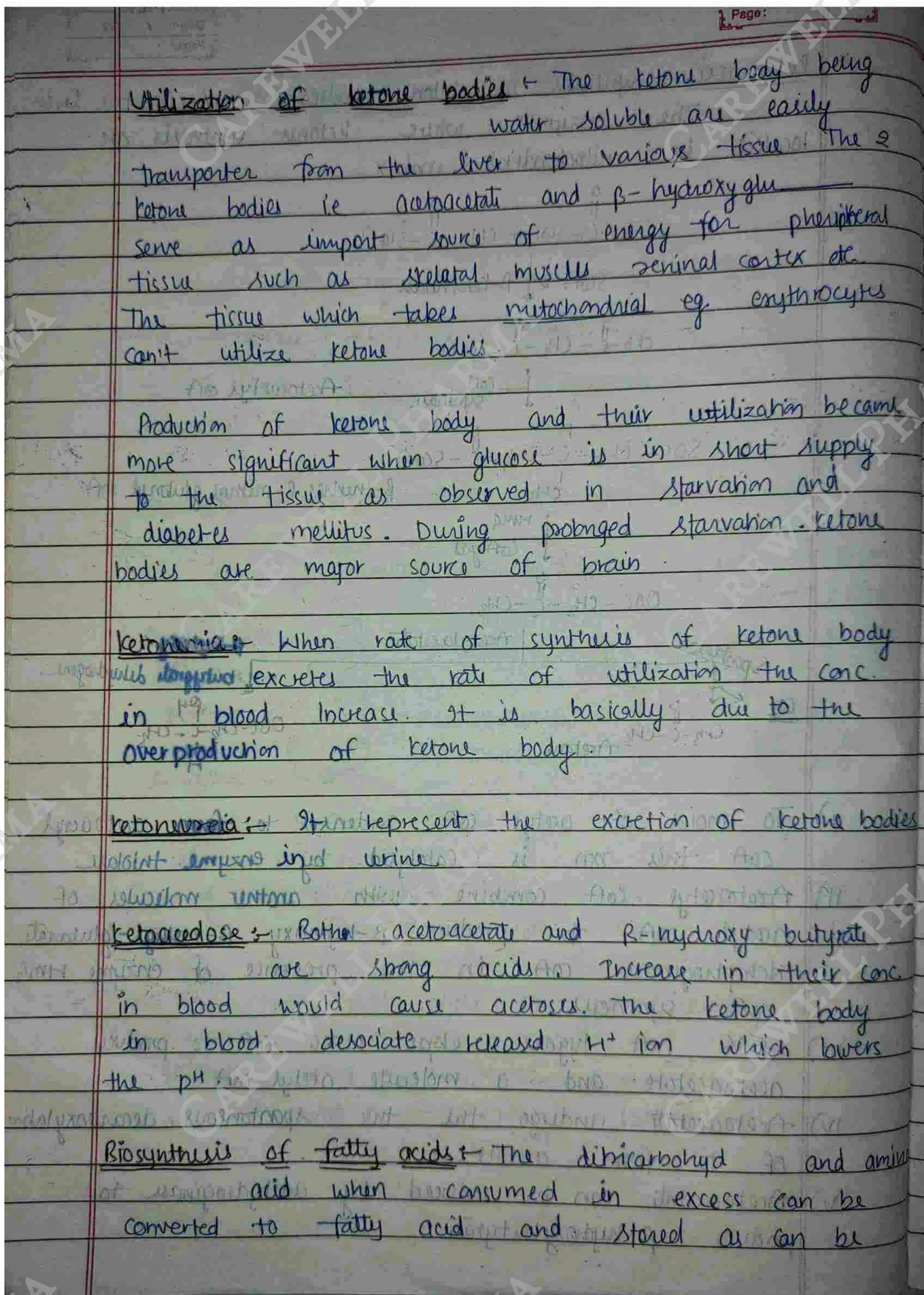
- i) 2 moles of acetyl CoA condenses to form acetoacetyl CoA this rxn is catalysed by enzyme thiolase.
- ii) Acetoacetyl CoA combine with another molecules of acetyl CoA to produce β-hydroxy-β-methyl-glutarate dehydrogenase CoA in the presence of enzyme HMG CoA synthase.
- iii) HMG CoA lyase cleaves HMG CoA to produce acetoacetate and a molecule acetyl CoA.
- iv) Acetoacetate undergoes the spontaneous decarboxylation of to form acetone.
- v) Acetoacetate can be reduced by dehydrogenase to produce β-hydroxybutyrate.

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converted to fatty acid and stored as triglyceride. The de novo biosynthesis of fatty acid occurs predominately in occur in adipose tissue & kidney the enzyme machinery for fatty acid synthesis is located in cytosomal fraction of cells. The fatty acid synthesis takes place in 3 step

- i) Production of acetyl CoA and NADH.
- ii) Conversion of acetyl CoA to malonide CoA.
- iii) Rxn of fatty acid synthesis complex.

i) Production of acetyl CoA and NADPH

Acetyl CoA is the main requirement for fatty acid synthesis. It is produced by oxidation of pyruvate in the mitochondria during TCA cycle, amino acid degradation and also from ketone bodies. In TCA cycle, the first product is citrate that is transported to cytosol. In this reaction step, acetyl CoA and oxaloacetate is synthesized by the cleaved reaction of citrate with citrate lyase enzyme. Oxaloacetate is further converted into pyruvate with the formation of NADPH. Therefore, acetyl CoA and NADPH are both important for the initiative steps for synthesis of fatty acids.

Conversion of acetyl CoA to malonyl CoA $\text{Acetyl CoA} + \text{NO}_2 + 7 \text{ATP}$

Carbohydrate and amino acids

Acetyl CoA and NADPH are prerequisites for fatty acids synthesis. Acetyl CoA is produced in mitochondria.

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Mitochondria power are not permeable to acetyl coA
 An alternate or by pass arrangement is made for
 the transfer of acetyl coA to cytosol.

Cytosol

Pyruvate

Malate

Oxaloacetate

Citrate

→

mitochondrial matrix

Pyruvate

Acetyl coA

Oxaloacetate

Citrate → Citrate synthase

Citrate cycle

ATP

Mitochondrial matrix

Oxaloacetate

Citrate

ATP

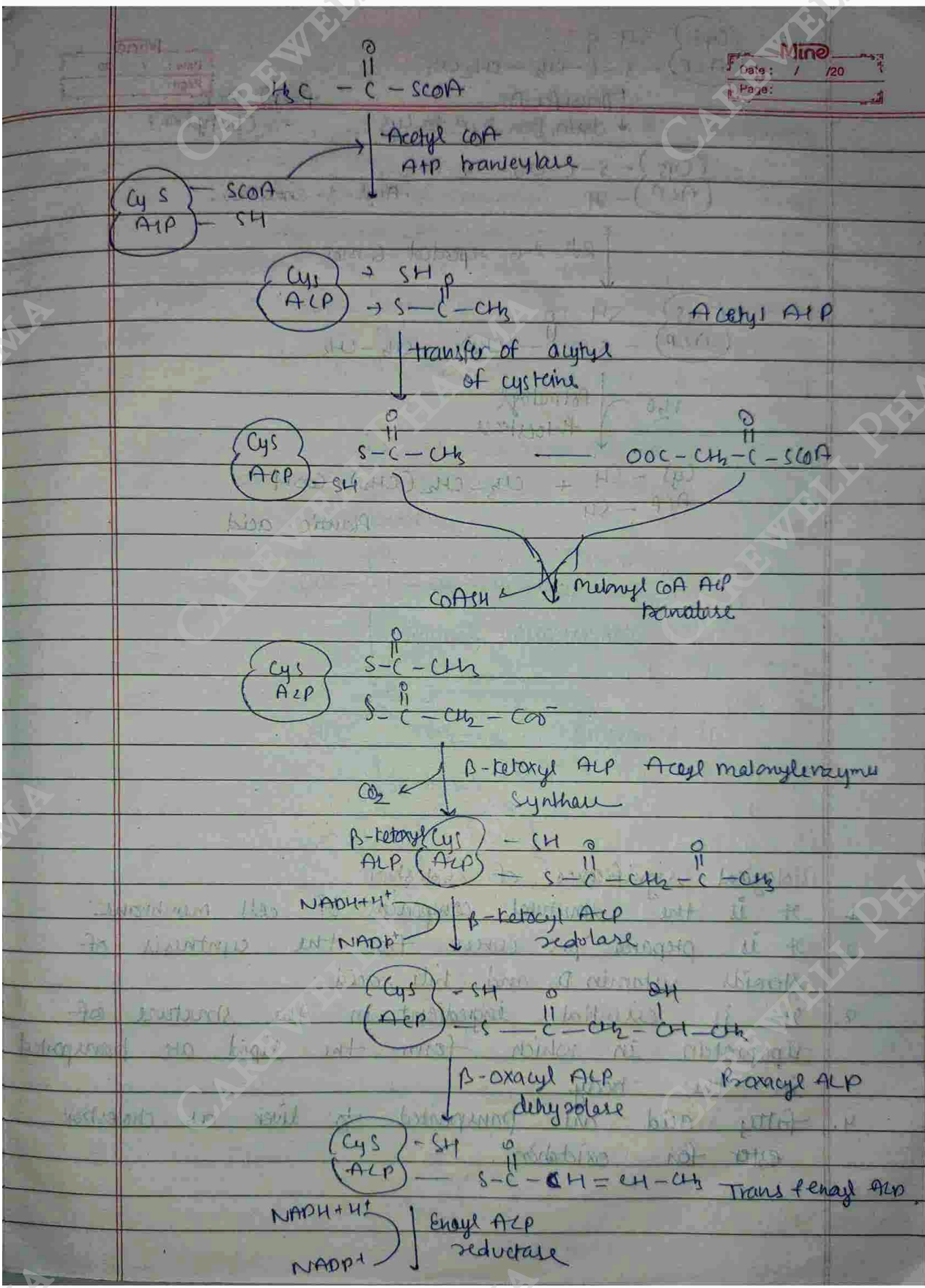
2. Formation of malonyl coA :-
 Acetyl coA is carboxylated to malonyl coA

$$\text{CH}_3 - \text{C}(=\text{O}) - \text{SCoA} + \text{CO}_2 + \text{ATP} \xrightarrow{\text{Carboxylase}} \text{OOC} - \text{CH}_2 - \text{C}(=\text{O}) - \text{SCoA} + \text{ADP}$$

3. Reaction of fatty acid synthase complex :-
 The remaining reaction of fatty acid synthesis
 are catalysed by a multifunction enzyme known
 as fatty acid synthase complex (FAS)

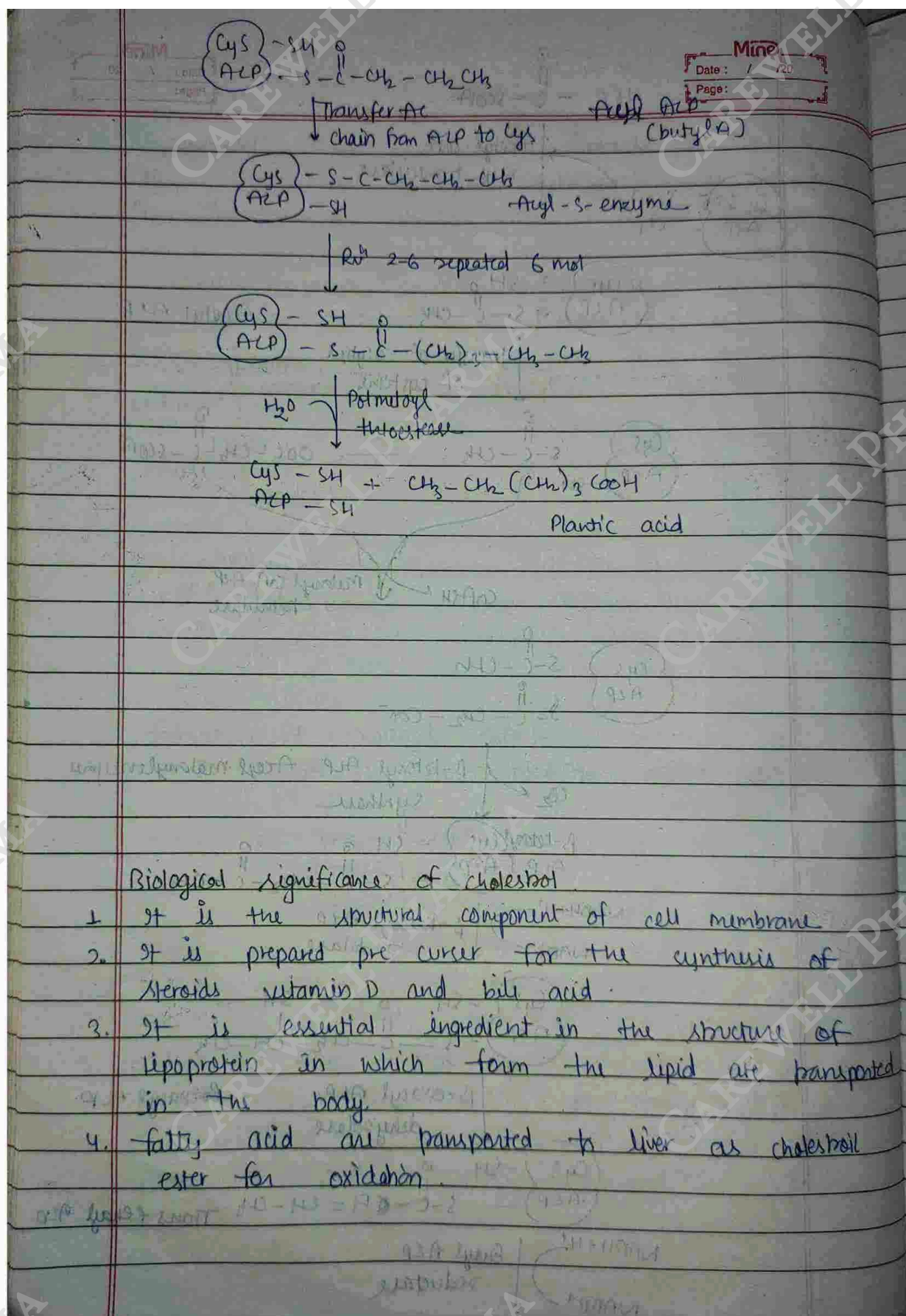
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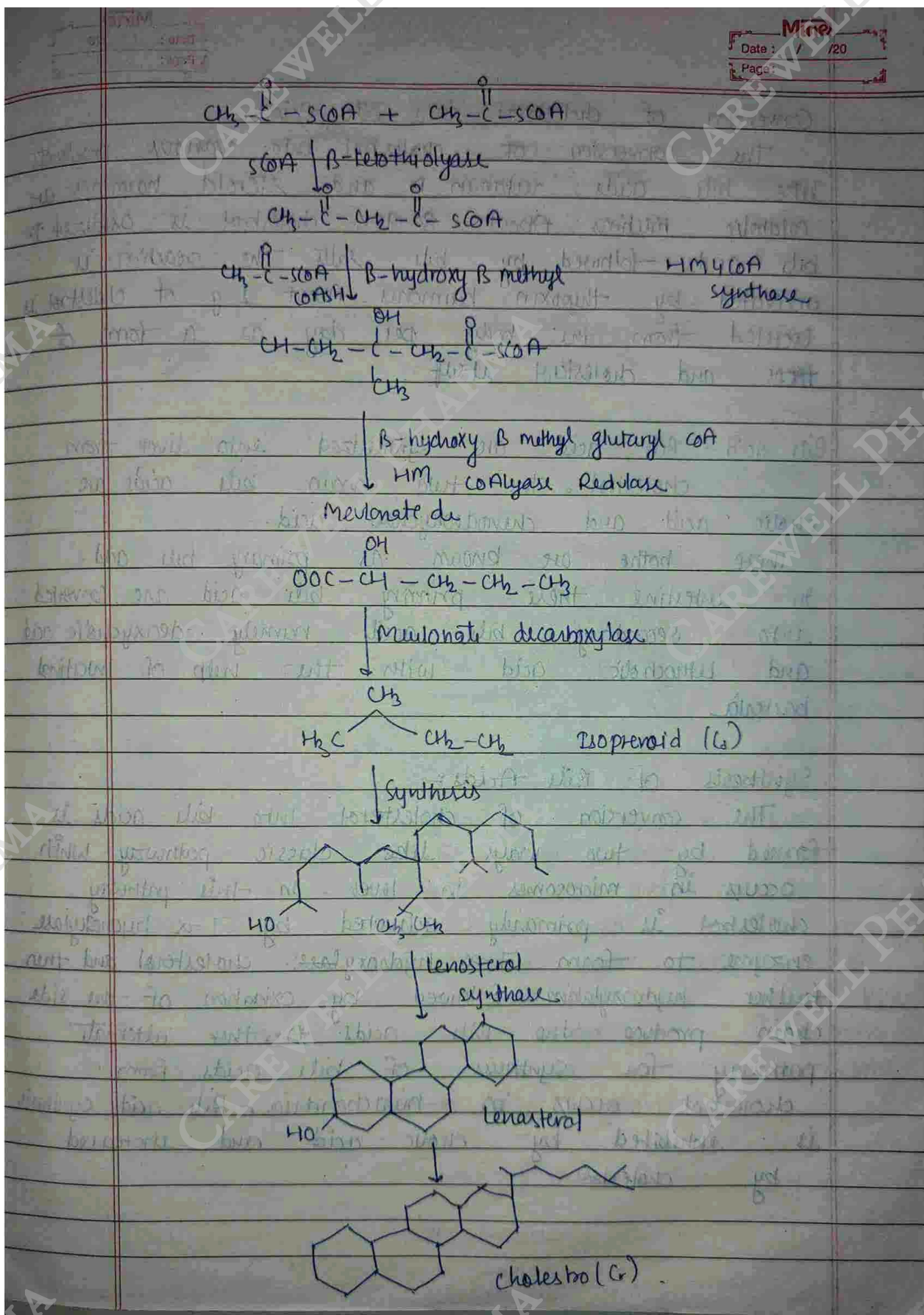
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Conversion of cholesterol into bile Acids

The conversion of cholesterol into various products like bile acids, vitamin D and steroid hormones are catabolic reactions. About 80-90% cholesterol is oxidized to bile acids followed by bile salts. The reaction is activated by thyroxin hormones. About 1g of cholesterol is excreted from the body per day as a form of feces and cholesterol itself.

Bile acid → Bile acid are synthesized into liver from cholesterol. The two main bile acids are cholic acid and chenodeoxycholic acid.

These both are known as primary bile acid.

In intestine, these primary bile acid are converted into secondary bile acids namely deoxycholic acid and lithocholic acid with the help of intestinal bacteria.

Synthesis of Bile Acids:-

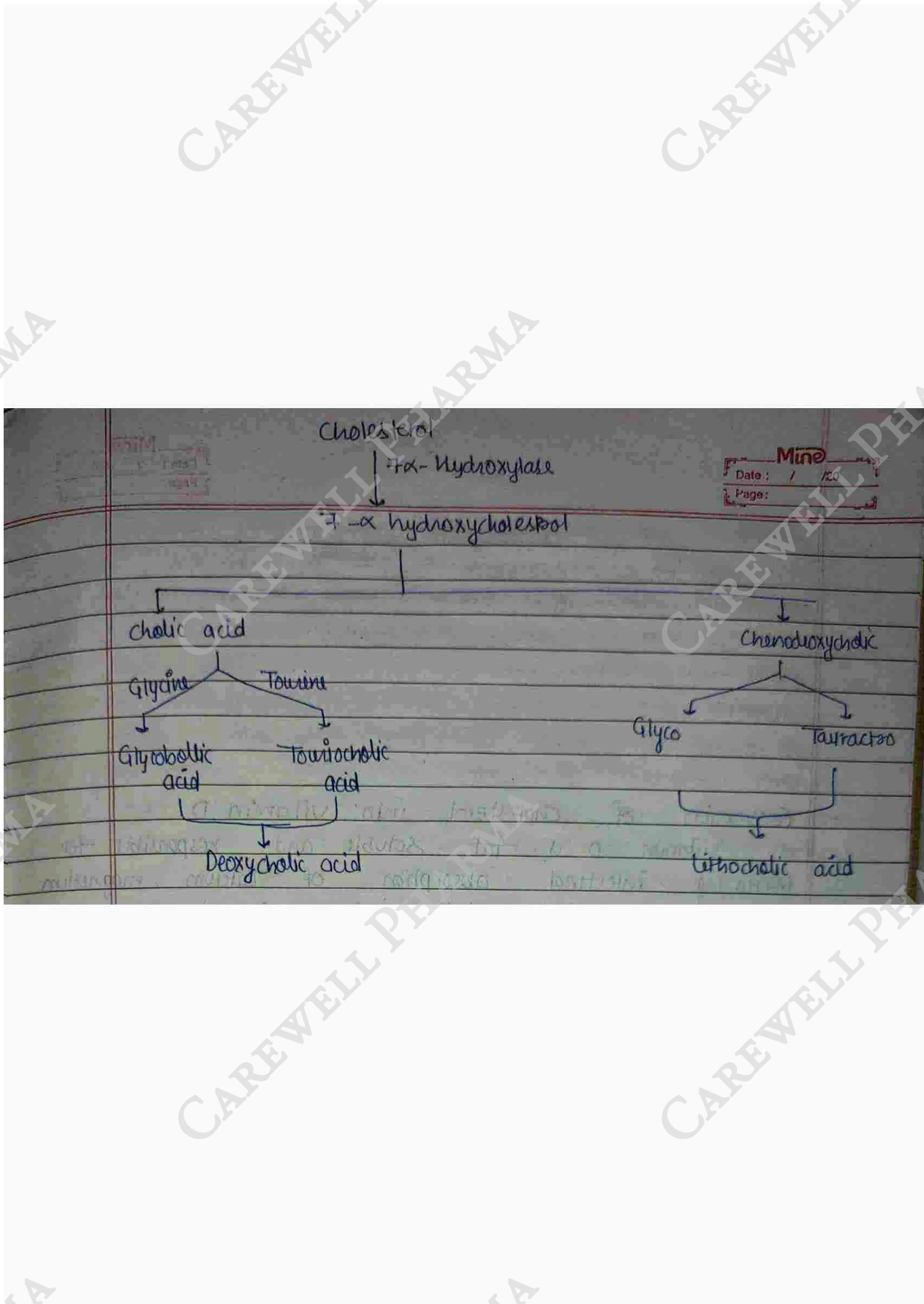
The conversion of cholesterol into bile acids is formed by two ways like classic pathway which occurs in microsomes in liver. In this pathway, cholesterol is primarily activated by 7- α hydroxylase enzyme to form 7- α hydroxylated cholesterol and then further hydroxylation followed by oxidation of the side chain produce the bile acids. Another alternate pathway for synthesis of bile acids from cholesterol occurs in mitochondria. Bile acid synthesis is inhibited by cholic acid and increased by cholesterol.

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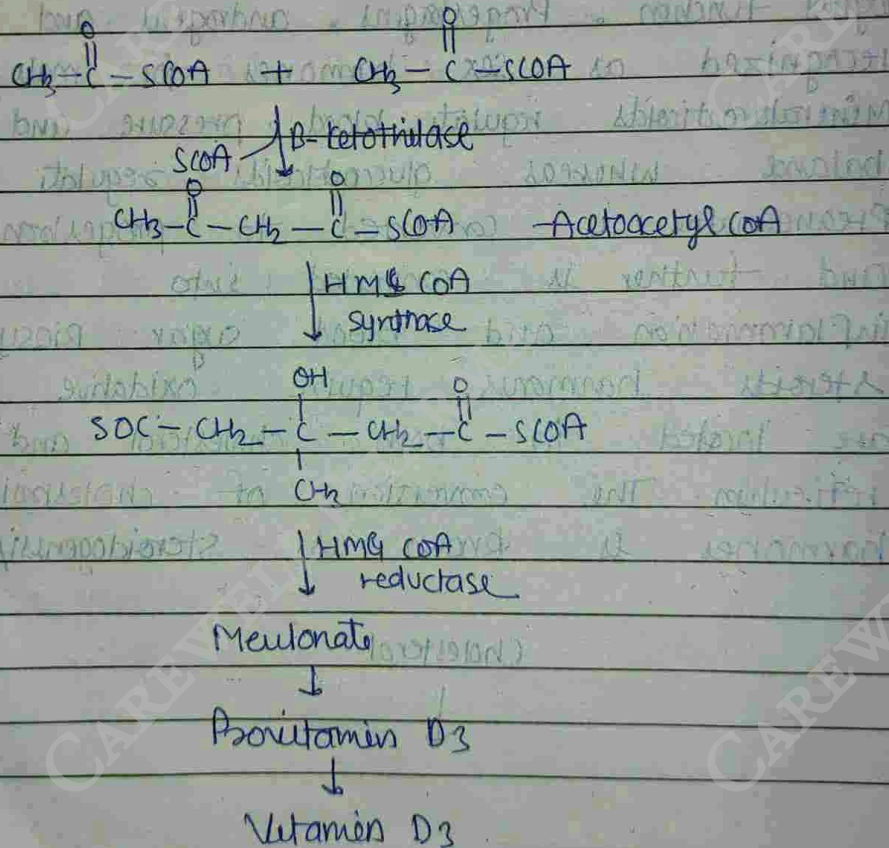


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• Conversion of cholesterol into vitamin D.

Vitamin D is fat soluble and responsible for increasing intestinal absorption of calcium, magnesium and phosphati. Chemically, Vitamin D is a secosteroid. During cholesterol synthesis, 7-dehydrocholesterol is formed as an intermediate product. On exposure to sun light 7-dehydrocholesterol is converted into cholecalciferol (Vit D₃) in the skin and finally converted into calcitriol.



Disorders of lipid metabolism.

Lipids are an important source of energy. These are stored as fat and constantly broken down to release energy with the help of some group of specific enzymes. But sometimes certain abnormalities in these enzymes can lead to the accumulation of fatty substances that are harmful to many organs of the body which is known as Lipidosis. Cholesterol level varies by age, weight and gender. It is measured in three categories viz. total cholesterol, LDL or (Low Density Lipid), HDL or (High Density Lipid), VLDL (Very low Density Lipid) and triglycerides. The normal amount of total cholesterol levels less than 200 milligram per deciliter. LDL cholesterol level should be less than 100 mg/dl. Some of the important ones are discussed in this section.

(I) Hypercholesterolemia → It is the situation of high levels of cholesterol present in the blood. People with hypercholesterolemia have a high risk of coronary artery disease. It is observed in many disorders like diabetes mellitus, in case of hypothyroidism, nephrotic syndrome etc. Plant sterols such as sitosterol esters reduce the plasma cholesterol level. Fibres in vegetables also decrease the cholesterol absorption. Some drugs like statins inhibit HMG CoA reductase and decrease cholesterol synthesis. It is diagnosed by using blood test.

(II) Atherosclerosis → It is a disease in which the inside of an artery narrows due to the build up of plaque which is made up of fat,

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cholesterol and other substance found in the blood. Atherosclerosis begins with damage to the endothelium. It is caused by high blood pressure, smoking or high cholesterol. The development of this condition is directly related with plasma cholesterol and low density lipid. This conditions leads to develops some disease like diabetes mellitus, nephrotic syndrome, hyperlipoproteinemia etc. eating a healthy diet and exercising are often the appropriate treatment for atherosclerosis. Some medicines like calcium channel blocker, B-blocker, antiplatelet agents are used to reduce the risk of atherosclerosis.

(11) Fatty Liver - It is the condition when the excessive fat deposit in the liver and results, deposition of toxic materials in the liver. This condition is also known as hepatic steatosis. In this condition the weight of the liver increases about 10% than that of normal weight and cause inflammation. Some common symptoms are like poor appetite, weight loss, abdominal pain, fatigue, weakness etc.

Liver is having the capability to regenerate new cells when the old cells are damaged but during sometimes liver fails to do that such condition is known as liver fibrosis/cirrhosis.

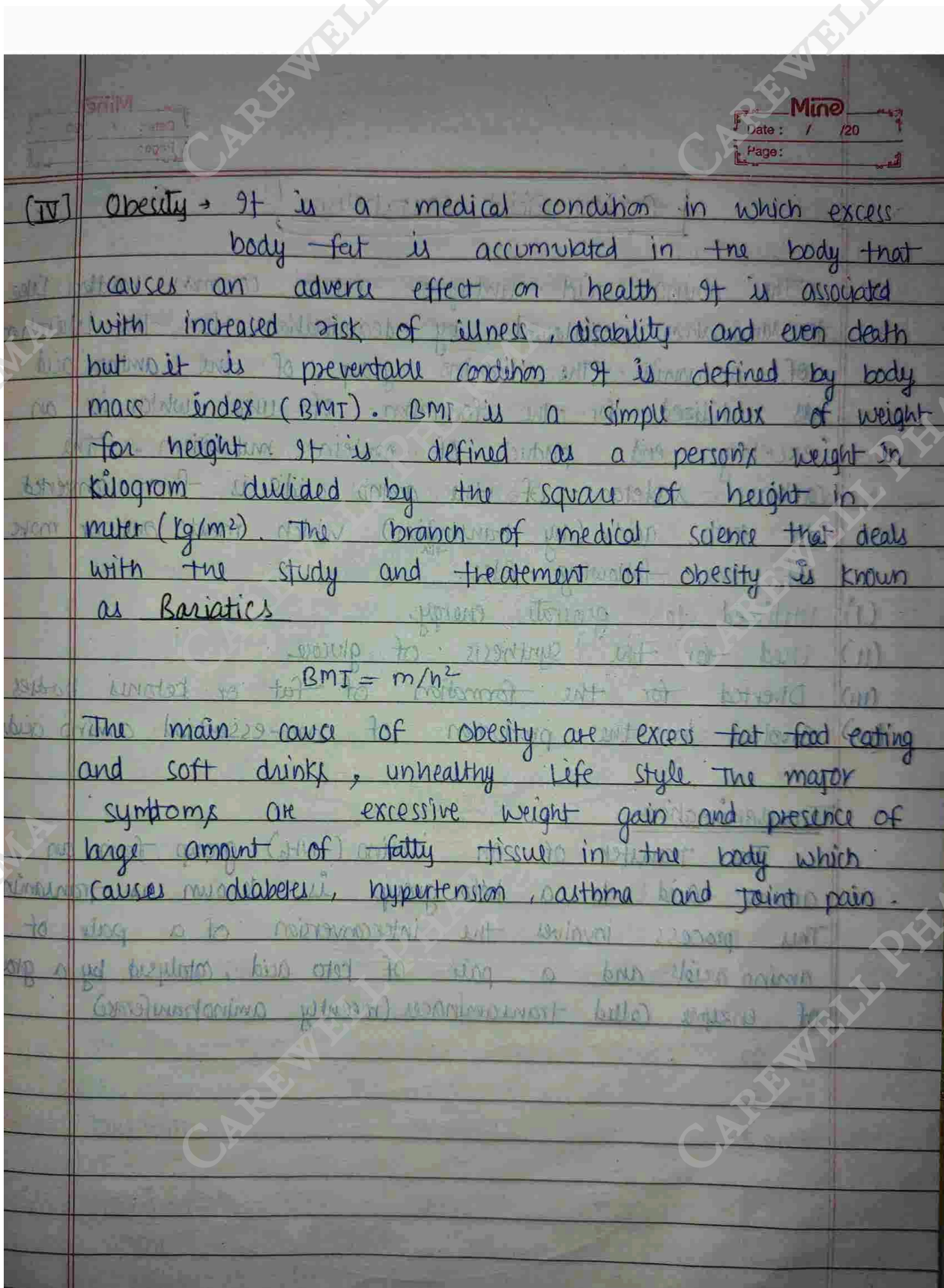
The most common cause of fatty liver is alcoholism and heavy drinking, diabetes, side effects of medicine like aspirin.

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