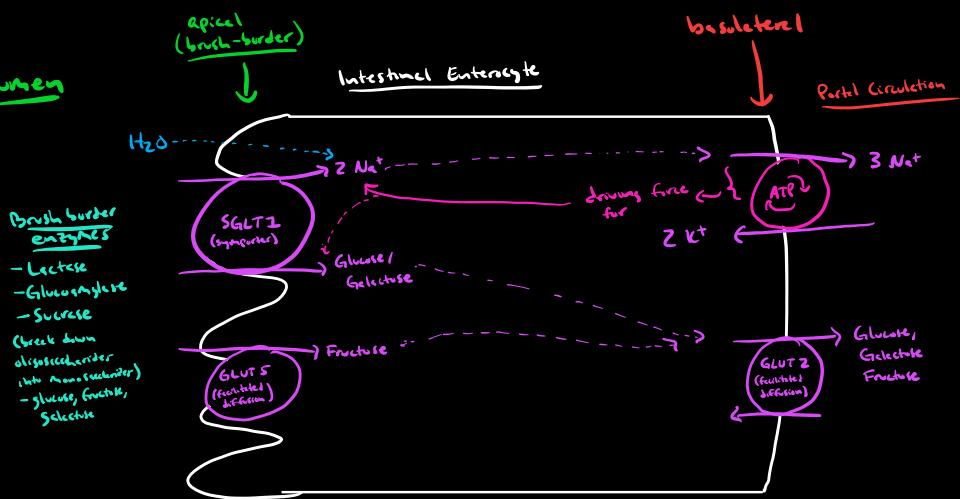
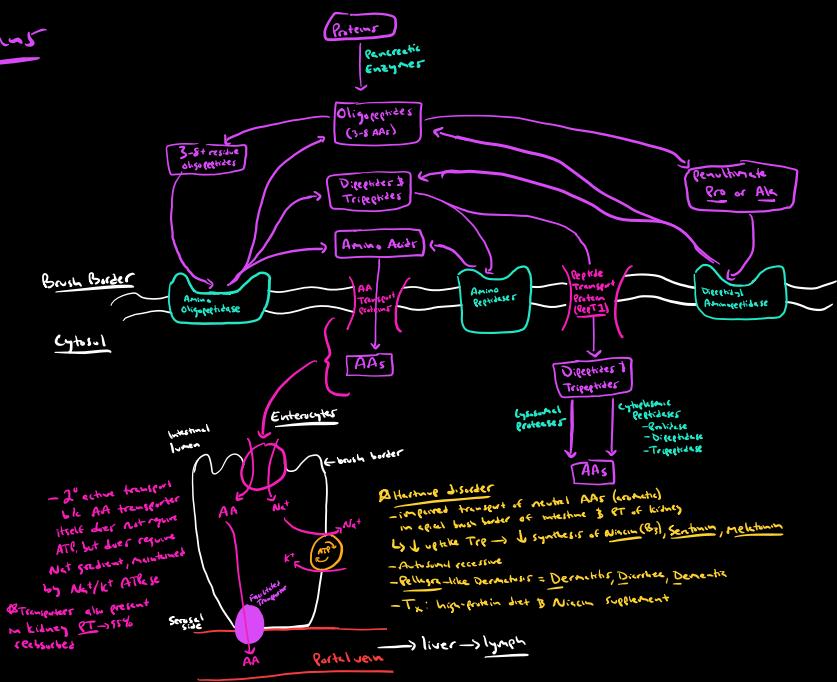


## Carbohydrate



## Proteins



④ Most fat digestion occurs in duodenum  
most fat absorption occurs in jejunum

## Facts



Micelles w/ LCFAs don't use Cerrier-  
Meeker transport to get into enterocyte  
- Fatty Acid Transport Protein 4  
- Caveolin 1

- Enterocytes

- Re-esterification of cholesterol

```

graph LR
    A[Cholesterol] + B[Free FA] --> C["Acyl-CoA cholesterol  
acyl transferase (ACAT)"]
    C --> D[TAGs]
    B --> E[Glycerol]
    E --> D
  
```

The diagram illustrates the re-esterification of cholesterol. It shows a reaction between cholesterol and free fatty acids (FA) to produce triacylglycerols (TAGs). This process is catalyzed by Acyl-CoA cholesterol acyl transferase (ACAT). The free fatty acids are also shown reacting with glycerol to form TAGs.

vitamins A, D, E, K + Cholesterin esters + phospholipid + TAGs

**Apolipoprotein B48**

- Only found on chylomicrons
- Contains 49% of apo-B protein
- Required for **secretion** from enterocytes

**Chylomicron**

**Enterocyte**

**Lymph**

**Blood**

**Lipoproteins**

**Transport Pathway:** Enterocyte → Lymph → Blood → Lipoproteins

**Annotations:**

- Binds to liver receptors
- Used for uptake of TAG
- FAST + EASY to tissues
- extracellular to capillary

The diagram illustrates the role of HDL in remnant lipid metabolism. It shows HDL (orange circle) containing ApoE (green oval) and ApoC-II (green oval). ApoE is labeled as a "co-factor for lipoprotein lipase". The text "to know" points to the top right. A bracket on the right indicates that HDL is "Carried by chylomicrons" and "facilitates binding to hepatic remnant receptors".

```

    graph TD
        FAs[FAs] --> Adipocytes[Adipocytes]
        Adipocytes --> FattyAcidCatabolism[Fatty Acid catabolism]
        FattyAcidCatabolism --> F[FA]
    
```

The diagram illustrates the metabolic pathway of fatty acids (FAs) in adipocytes. FAs enter the cell, where they undergo catabolism to produce energy. The resulting products are then released back into the blood stream.

```

graph TD
    Liver[Liver] --> ChylRem[Chylomicron Remnants]
    ChylRem --> LDL[LDL]
    LDL --> VLDL[VLDL]
    VLDL --> Chol[Cholesterol]
    VLDL --> HDL[HDL]
    Chol --> Tissues[Tissues]
    HDL --> Tissues
    Tissues --> CholTissue[Cholesterol]
    CholTissue --> Tissues
  
```

- extracellular enzyme anchored to capillary walls
- mostly found in adipose tissue, skeletal muscle, breast
  - not in liver (liver has hepatic lipase)
- converts TAG<sub>triglycerides</sub> → FAs → storage or fuel (triacylglycerols)
- requires apo C-II for activation

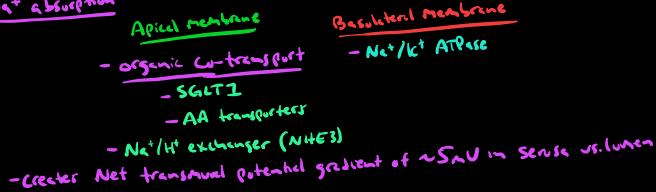
## Intestine vs. Colon Salt & H<sub>2</sub>O Absorption

- Crypts of Lieberkühn - found in mucosa of both small & large intestine
  - Contain different cells depending on location
  - Most concentrated in duodenum
  - In all areas, can secrete isotonic NaCl
    - Apical membrane - CFTR (Cl<sup>-</sup>) channels
    - Basolateral membrane - NKCC cotransporter
    - Na<sup>+</sup> & H<sub>2</sub>O follow paracellularly to via AQP channels
- Fluid can either enter intestinal lumen or be absorbed via osmotic gradients

### Small Intestine

- Under normal conditions, fluid moves rapidly in & out to achieve isotonicity by early jejunum
- Subsequently, fluid absorbed isotonically using standing osmotic gradient in lateral spaces between enterocytes

### Na<sup>+</sup> absorption



- Ca<sup>2+</sup>
  - Apical: normal to P (Ca<sup>2+</sup>)
    - ↳ paracellular diffusion, nonselective
  - Basolateral: (Plasma membrane ATPase (PMCA))
    - ↳ regulated by Vit.D
    - Active transport via TRPV6 (ion channel)
    - Na<sup>+</sup>/Ca<sup>2+</sup> exchanger (NCX1)
    - regulated by Vit.D
    - inhibited by okadaic, phytosterols, dietary fiber, Mg<sup>2+</sup>, PO<sub>4</sub><sup>3-</sup>, tannins
    - inhibited by  $\text{Ca}^{2+}$  /PO<sub>4</sub><sup>3-</sup> combine w/ Ca<sup>2+</sup> → stearate
    - increased by acidic pH, Vit.D, estrogen, lactose

- Phosphate
  - in plant seeds, storage form → phytate, only 25% available to humans b/c don't have phytase
  - meets, Sefat, being, leg, SF, cereal
  - Apical: passive, paracellular
  - Active Type 2B Na<sup>+</sup>-P cotransporter
    - stimulated by Vit.D
    - driven by Na<sup>+</sup> [Ca<sup>2+</sup>] gradient, driven by Na<sup>+</sup>/K<sup>+</sup> ATPase

- Mg<sup>2+</sup>
  - stored in bones
  - reabs in muscle & skin (Ca<sup>2+</sup> in ECs)
  - used as cofactor for ATP-driven reactions (e.g. neuromuscular & muscle contraction)
  - Apical: passive, paracellular
  - Active: (Na<sup>+</sup>-Mg<sup>2+</sup> cotransporter)
    - stimulated by Vit.D
    - driven by Na<sup>+</sup> [Mg<sup>2+</sup>] gradient, driven by Na<sup>+</sup>/K<sup>+</sup> ATPase
  - Basolateral: (Na<sup>+</sup>-Mg<sup>2+</sup> cotransporter)
    - can only absorb Fe<sup>2+</sup> at brush border
    - absorption promoted by: acidic, zinc, cysteine
    - absorption inhibited by: phytate, lactose
    - Ferritin mRNA has competing F<sup>+</sup>-IRE (Fe<sup>2+</sup> free to bind IRE → F<sup>+</sup> free to bind IRE → block translation → ↓ storage)
    - (Ferritin mRNA has 3'-I<sup>+</sup>-IRE (Fe<sup>2+</sup> free to bind IRE) but IRE not protected by endonuclease → ↓ translation → ↓ Fe uptake)
    - Transferrin receptor mRNA has 3'-I<sup>+</sup>-IRE (Fe<sup>2+</sup> free to bind IRE) but IRE not protected by endonuclease → ↓ translation → ↓ receptor translation → ↓ uptake

- Once in circulation, peripheral cells uptake iron from blood w/ transferrin receptor (mediated endocytosis)
  - Apical: Heme-Fe<sup>2+</sup> → Heme-Carrier Protein (HCP)
    - Fe<sup>2+</sup> transporter family
    - then deliver Heme-Fe<sup>2+</sup> to Heme oxygenase inside cell
    - Fe<sup>2+</sup> released & enters
    - O<sub>2</sub> + Fe<sup>2+</sup> → Ferroin (heme deoxygenation)
    - Ferroin → Hemopexin (Fe<sup>2+</sup> decon)
    - transported through circulation via ferritin
  - Non-heme Fe<sup>2+</sup> → Divalent Cytochrome b (CYTB) on brush border
    - reduces Fe<sup>2+</sup> to Fe<sup>3+</sup> → DMIT
    - enhanced by Vit.D

- HFE protein inhibits transferrin receptor @ ↓ [Fe<sup>2+</sup>]
  - $\text{Fe}^{2+}$  → HFE protein dissociates → ↑ Fe<sup>2+</sup> uptake
  - mutations in ferroportin, & molecule that promote hepcidin secretion (HFE, BMPR) → ↑↑ Fe<sup>2+</sup> absorption
  - Uptake/distribution
    - most common cause = HFE mutation
    - HFE = hemochromatosis protein
    - Hereditary HFE-associated cutaneous recessive inheritance
    - all types involve dysfunction in ferroportin pathway → ↑ iron transients from enterocytes → blood

- Cl<sup>-</sup> absorption
  - ~5mV potential gradient due to Na<sup>+</sup> absorption
    - ↳ driver ↑ Cl<sup>-</sup> flux via "leaky" epithelial junctions (paracellular)

- K<sup>+</sup> absorption
  - ~5 (K<sup>+</sup>) rise in lumen or fluid is reabsorbed
    - ↳ creates transmucosal gradient
    - ↳ driver K<sup>+</sup> absorption paracellularly

### Divalent Metal Transporter 1 (DMT1/SLC1)

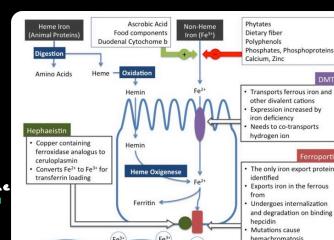
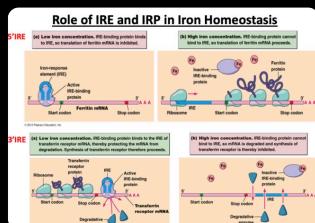
- Intestinal transporter responsible for absorbing:
  - Iron
  - Manganese
  - Nickel
  - Zinc
  - Calcium } Tonic
  - Lead

### Colon

- Colonic epithelium effectively secretes K<sub>2</sub>HCO<sub>3</sub> in exchange for Na<sup>+</sup> absorption
  - ↳ excessive diarrhea → ↑↑ secretion of K<sub>2</sub>HCO<sub>3</sub>
  - ↳ Metabolic acidosis & hypokalemia
- Significantly decreased absorptive capacity for electrolyte & H<sub>2</sub>O due to:
  - ① shorter length
  - ② smaller surface area (no villi)
  - ③ tighter epithelial junctions (no paracellular transport)
- Colonic bacteria produce Short Chain Fatty Acids (SCFA) from Soluble Fiber
  - ↳ ↑ Na<sup>+</sup> & H<sub>2</sub>O absorption via Sodium Monocarboxylate Transporter (SM1)
  - transports SCFAs & Na<sup>+</sup> across apical membrane of colonicocyte
  - ↳ antibiotics → destroy colonic bacteria → ↓ SCFA production → ↓ Na<sup>+</sup> & H<sub>2</sub>O absorption
    - diarrhea

### Pernicious Anemia

- autoimmune disorder where antibodies attack pernicious cells
  - ↳ ↓ intrinsic factor → Vit. B<sub>12</sub> deficiency
  - ↳ impairs RBC growth → megaloblastic anemia (MCV > 120, normal = 100-105)
  - ↓ HCl secretion → Parietal → ↑ Gastrin release by G cell



## Minerals

Iron  
- Iron absorption must be tightly regulated b/c mostly recycled  
and found in [↑] in dietary sources

- only lose iron via:
  - natural shedding of skin & mucosa
  - Blood loss (menstruation/GI bleeding)
  - Sweat

- Iron readily absorbed in Ferric (Fe<sup>3+</sup>) state  
↳ Stomach proton Fe<sup>3+</sup> chelation b/c DMT1 (P1B1) promoter ↑ Fe<sup>3+</sup> (ferric state)

- absorbed on apical surface of enterocyte in Duodenum

2 forms:

- Heme iron

- complexed w/ nonheme heme group
- b) readily absorbed via receptor-mediated endocytosis
- Iron = Fe<sup>2+</sup>
- found in red meat

- Nonheme iron

- Unbound/free

- Fe<sup>2+</sup> or Fe<sup>3+</sup>

- poorly absorbed due to charge

↳ requires divalent metal transporter I (DMT1)

- couples 1st absorption w/ Fe<sup>2+</sup> (NBT Fe<sup>2+</sup>)

- acidic pH of GI lumen keeps iron in Fe<sup>3+</sup> state

↳ Duodenal Cytochrome b (Dcyb) iron-reductase

- reduces Fe<sup>3+</sup> → Fe<sup>2+</sup>

- Vit. C cofactor

- once Fe<sup>2+</sup> absorbed by enterocyte...

① stored as Fe<sup>2+</sup> in Apo ferritin

② shuttled across basolateral membrane into systemic circulation

- Fe<sup>2+</sup> transported across membrane via Ferritinoprotein

↳ oxidized to Fe<sup>3+</sup> via Ferricidase (the 1MEG/Hephaestin Complex)

- Ferricidase = transmembrane protein

- binds Fe<sup>2+</sup> in blood → Fe<sup>3+</sup>

↳ Fe<sup>3+</sup> binds Transferrin → Liver → dispersed to body

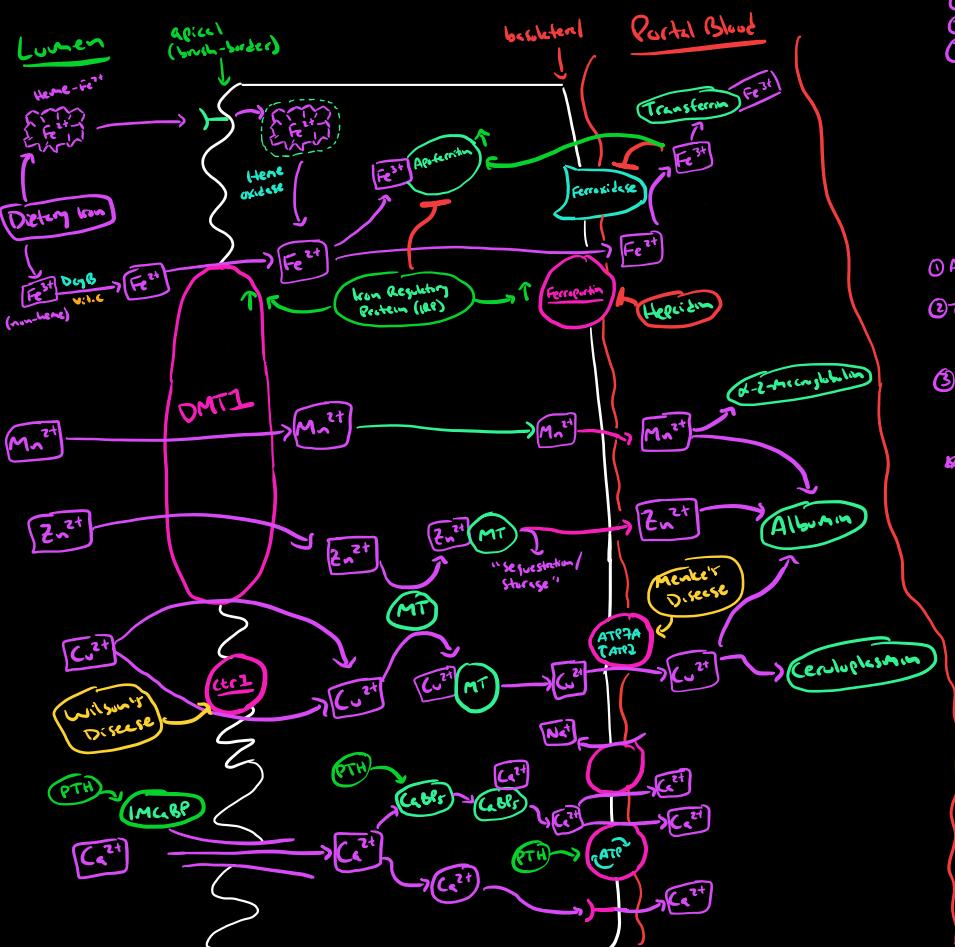
- Transferrin = Fe<sup>3+</sup> binding protein

- Regulation

- ↑ [Fe] in blood → Liver releases Hepcidin

- Hepcidin binds to ferritinoprotein → ferritinoprotein degraded

↳ ↓ [Fe] release



## Copper

- Regulated/controlled by metabolism & excretion in liver

- not big uptake like w/ iron

① Absorbed into enterocyte in Duodenum to Stomach

- High-affinity Copper transport protein I (Ctr1)

- DMT1

↳ MP(Cu) → Ctr1 saturation → ↑ [Cu] uptake inc

DMT1 → ↓ uptake of other divalent ions

like Zn<sup>2+</sup>, Fe<sup>2+</sup>, etc.

- Cu<sup>2+</sup> absorption inhibited by:

- T fiber

- ↑ Zn<sup>2+</sup>

② Once inside enterocyte...

- binds to Metallothionein (MT)

- MT = important regulator of heavy metals in body

- transports iron to basolateral membrane

③ Menkes P-type ATPase (ATP7A)

↳ transports Cu<sup>2+</sup> from cell → blood

④ In blood (central view)

- Cu<sup>2+</sup> binds to albumin → transported to liver

⑤ In liver, Cu<sup>2+</sup> complexes with protein to form Ceruloplasmin

↳ Ceruloplasmin excreted or secreted

- excreted w/ bile via common bile duct

- travels back into blood → dispersed throughout body

- Wilson's disease - defective enzyme in liver responsible for

- metabolizing/excreting Copper → ↑ [Cu<sup>2+</sup>] in blood

- excess toxicity → oxidation → ROS → oxidative damage

- Menke's disease - defective Copper transporter → Copper deficiency

- most commonly due to ATP7A defect, X-linked mitochondrial loss of function

- deficiency of basal myelinating (collagen), ceruloplasmin, dopamine hydroxylase

- "Wimley" / "Steely" hair, stunted growth, abnormal skin pigmentation, mental deterioration

## Zinc

- required for immune function & tissue regeneration

- utilized as cofactor by metalloenzymes that hydrolyze collagen

① absorbed via DMT1

- absorption inhibited by ↑ Zn<sup>2+</sup> or Fe<sup>2+</sup> (Ceruloplasmin)

② Binder to MT → sequestered for storage or transported to blood

→ Zn<sup>2+</sup> → upregulates MT expression

③ In blood, binds to plasma proteins (albumin, α<sub>2</sub>-macroglobulin) → Liver → dispersed

Deficiency → Anendermic Enteropathy (anemia, Zn<sup>2+</sup> transport pattern)

- intestinal absorptive

Tolerance → ↑ MT → that can bind Copper → Copper deficiency → iron deficiency

## Manganese

① Absorbed via DMT1

② Transported to basolateral membrane

③ Transported to blood via iron exporter (ferritinoprotein → transferrin)

- in blood, binds albumin or α<sub>2</sub>-Macroglobulin → Liver → dispersed

- important for metabolism of proteins & fat, supports immune system, regulates glucose homeostasis

- Cofactor for Arginase (urea cycle), pyruvate carboxylase (gluconeogenesis)

- works w/ Vit C to provide collagen

- Cofactor for Superoxide dismutase (front line defense against free radicals)

Toxicity → neurological problems

- Seizures

- Parkinsonian symptoms

## Calcium

① Absorbed via Ca<sup>2+</sup> channel

- Channel activated by Intestinal Membrane Ca<sup>2+</sup> Binding Protein (IMCBP)

② Transcribed through cytoskeleton via:

- vesicles

- Ca<sup>2+</sup> binding protein (CaBP), aka Calbindin (different types exist)

③ Secreted across basolateral membrane via:

- Ca<sup>2+</sup> ATPase

- Ca<sup>2+</sup>/Na<sup>+</sup> exchanger

- vesicle exocytosis

④ Regulation by PTH

↓ [Ca<sup>2+</sup>] → ↑ PTH → ↑ formation of active Vit.D<sub>3</sub> (1,25(OH)<sub>2</sub>D<sub>3</sub>) → ↑ protein

Synthesis of IMCBP, Calbindin, Ca<sup>2+</sup>/Na<sup>+</sup> ATPase

Site of Absorption	Key Proteins
Carbohydrates	Enterocytes (small intestine)
Proteins	Enterocytes (small intestine)
Fats	Enterocytes (small intestine)
Iron	Duodenum
Copper	Duodenum
Zinc	Duodenum
Manganese	Duodenum

⇒ Primary sites of absorption mnemonic

Dude Is Just Feeling Ill Brr

Duodenum - Iron (Fe<sup>2+</sup>)

Jejunum - Zinc

Ileum (terminal) - Brr & Bile Salt