CHEM 496 Topics in Chem & Biochem: Biochemistry, Physiology & Neurochemistry of Beer, Wine & Alcohol

Block III – EtOH metabolism, alcohol use disorder & pathology



Alcohol – who we metabolize EtOH, what are the effects (chronic and acute), and how does it impact our brain?



Alcohol as a Drug

- Alcohol is a psychoactive drug that is a **CNS depressant**.
- Some claim that alcohol is the most widely consumed drug in the world and for some is as much a part of daily life as eating.
- Alcohol is an addictive substance. Of the approximately 2 million receiving treatment for drug abuse, 64% are being treated for alcoholism.

Negative Impact of Alcohol

- 100,000 140,00 US deaths and 3 mil global deaths associated with alcohol each year (2.5 males for each female).
- Nearly 50% of all Americans will be involved in an alcohol-related traffic accident at some time during their lives.
- More than 2% of nighttime drivers have a blood-alcohol content that exceeds 0.08%, the legal amount in most states.



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Negative Impact of Alcohol

- Shortened the lives of those lost by an average of 26 years for a total of nearly 3.6 million – most due to long term usage leading to cancer, liver disease, and heart disease
- Costs over 287 billion in the US each year
- Illness, accidents, violence, and crime related to alcohol use
- Consumption by college students causes approximately 2,000 deaths per year.
- Fetal alcohol syndrome
- Alcohol is the second leading cause of premature death in America
- 2 million people admitted for treatment of alcohol use disorder annually

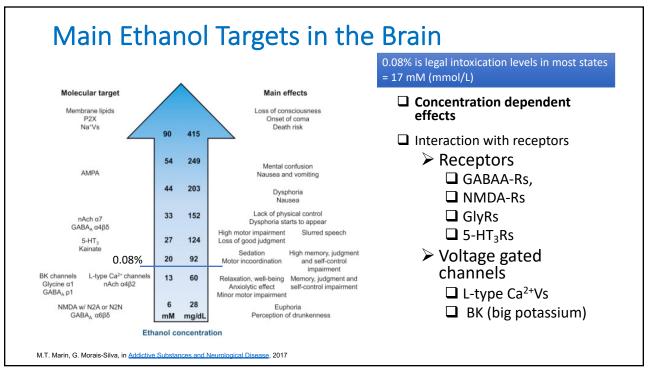
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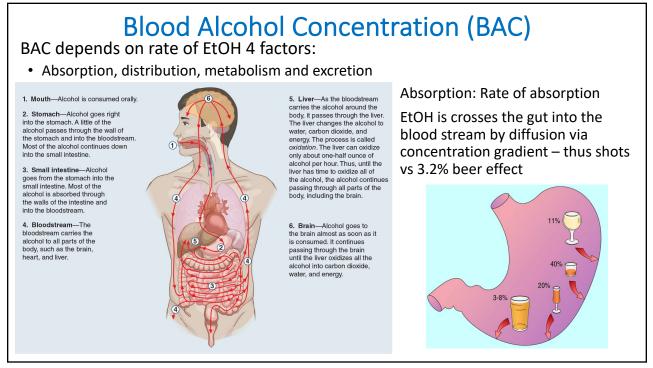
CAN CONTRIBUTE TO **CANCERS** OF THE:

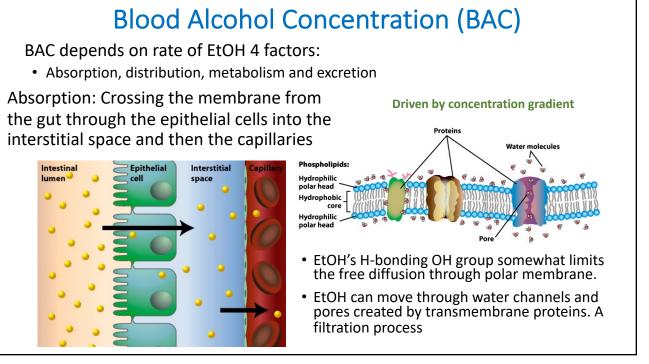
- MOUTH AND THROAT
- LARYNX (voice box)
- ESOPHAGUS
- COLON AND RECTUM
- LIVER

cdc.gov/cance

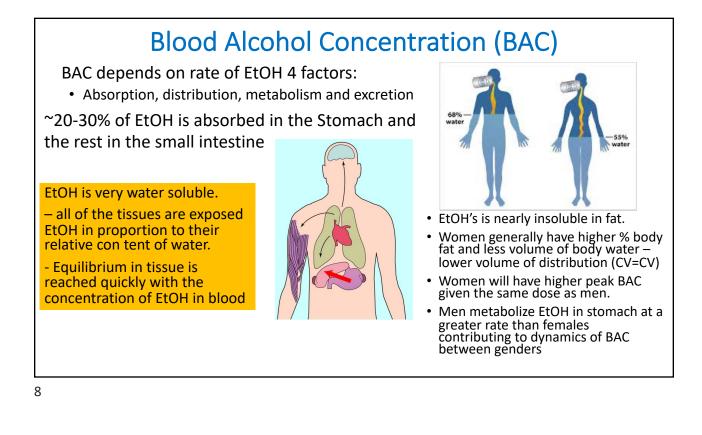
• BREAST (in women)

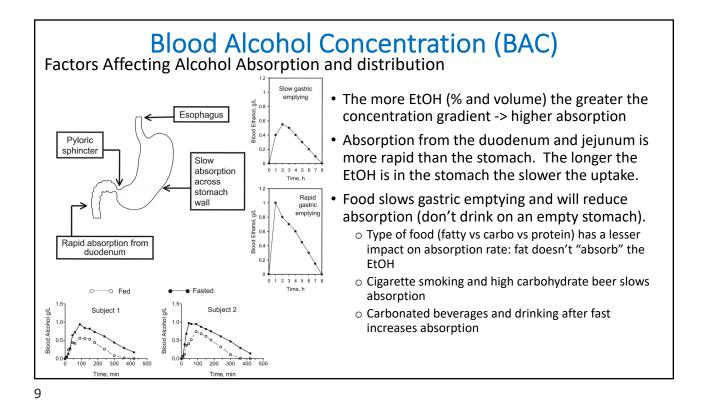








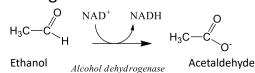




First Pass Metabolism

Some (minor fraction ~10-30%) of the EtOH is metabolized in the stomach (first pass metabolism) rest is metabolized in liver or a small amount is lost as liquid.

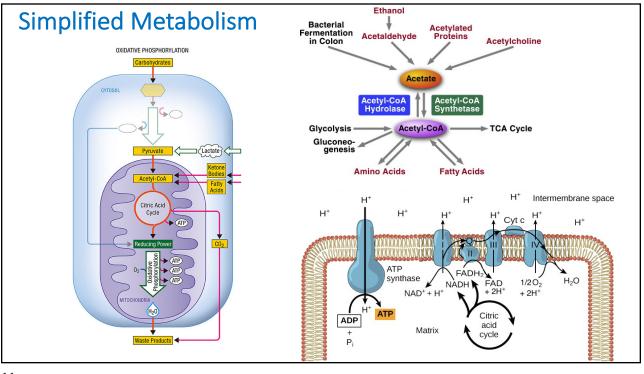
First-pass metabolism is a pharmacokinetic concept where some of a drug is metabolized prior to entering the blood



Alcohol Dehydrogenase (ADH class I and II) expressed in stomach.

- Impact of ADH is diminished in rapid stomach emptying
- Alcoholics (especially women) have decreased ADH activity and may contribute to increased sensitivity in women
- Aspirin and anti-ulcer / acid pump blockers inhibit stomach ADH

2-5% of Ethanol is lost unchanged in urine, sweat or breath

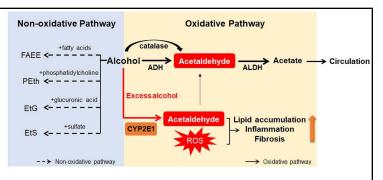


Alcohol Metabolism

Oxidative -Three systems

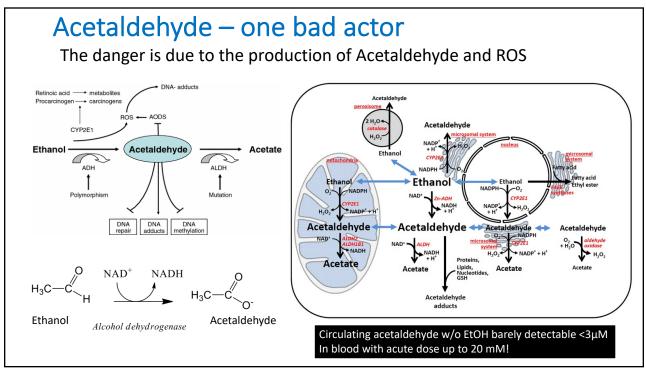
- 1) Alcohol dehydrogenase steady rate based on amount of ADH present
- 2) Cytochrome P450 system (CYP2E1) - induced by heavy use
- 3) Catalase
- Occurs primarily in liver but brain can participate
- Alcohol dehydrogenase is the major component

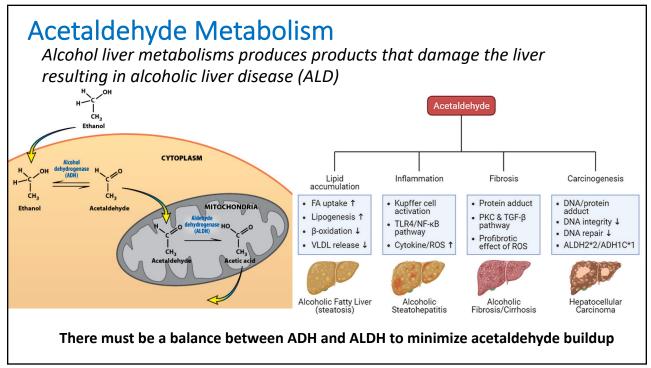
tone bodies and cholestero

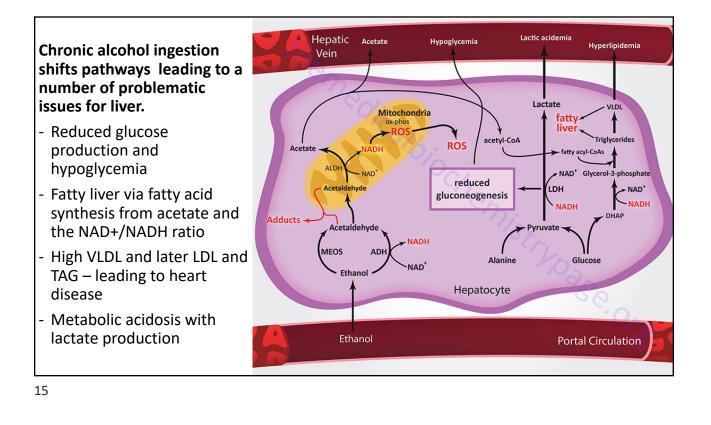


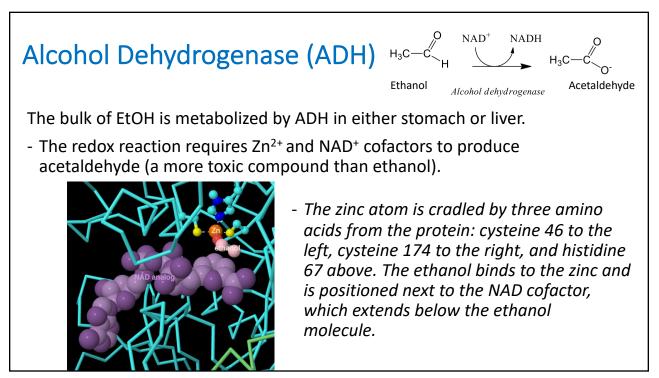
Non-Oxidative -minor pathway

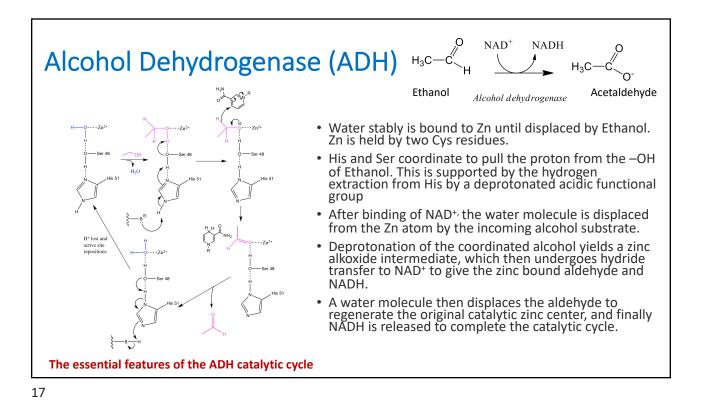
- 1) Ethanol esters conjugation to fatty acids (FAEE)
- PLD generation of PEth
- Conjugation with sugars and sulfates

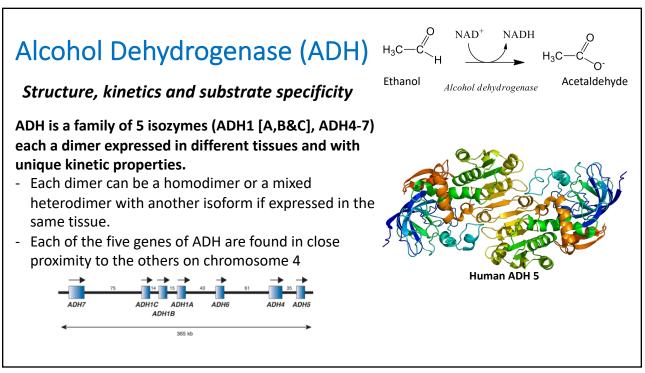


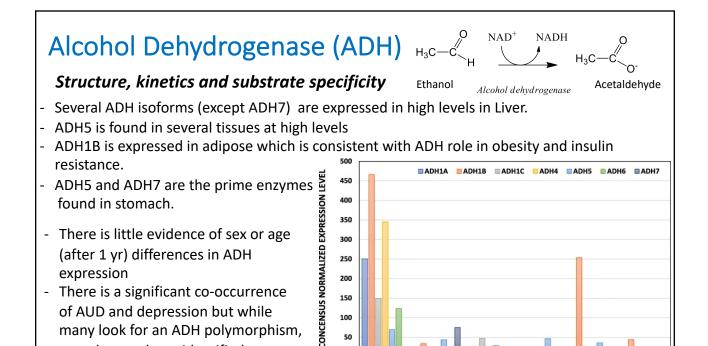












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none has yet been identified.

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Alcohol Dehydrogenase (ADH) Structure, kinetics and substrate specificity

In general at modest to high levels of EtOH most enzymes are saturated and metabolism is zero order

Name/Variant Km (mM) 300 500 150 ADH1A 4.0 ADH1A ADH1B ADH1C 30 Liver 250 400 200 100 ADH1B*1 (Arg/Arg) 0.05 4 Liver, adipose, 300 150 ADH1B*2 (His/Arg) 0.9 350 200 50 100 ADH1B*3 (Arg/Cys) 40.0 300 100 50 ADH1C*1 (Arg/Ile) 1.0 90 Liver, gut n ADH1C*2 (Gln/Val) 0.6 40 Liver issue Liver Colon Gallbladder nall intestine ADH4 30.0 20 Liver ADH5 >1,000 100 Ubiquitous 350 ADH6 --mRNA no protein ADH5 ADH7 300 60 1,800 ADH7 10.0 Throat tongue 250 60 200 40 40 150 BAC (mg/dL) mmol/L* Effect† 100 2(4–11 20-50 Decreased fine motor control 50 11-22 50-100 Decreased coordination 0 100-150 22-33 Difficulty standing Liver Colon mall intestine gland 33-55 Difficulty sitting 150-250 Epididv 66 Unresponsive to voice and/or pain 300 88 400 Respiratory depression

- At low concentration elimination slows as the EtOH falls near Km and maximum velocity is no longer taking place

Alcohol Dehydrogenase (ADH) Structure, kinetics and substrate specificity

Name/Variant	Km (mM)	Vmax	Tissue	
ADH1A	4.0	30	Liver	
ADH1B*1 (Arg/Arg) ADH1B*2 (His/Arg) ADH1B*3 (Arg/Cys)	0.05 0.9 40.0	4 350 300	Liver, adipose,	
ADH1C*1 (Arg/lle) ADH1C*2 (Gln/Val)	1.0 0.6	90 40	Liver, gut	
ADH4	30.0	20	Liver	
ADH5	>1,000	100	Ubiquitous	
ADH6			mRNA no protein	
ADH7	10.0	1,800	Throat tongue	

After 4 cans of beer or 6 shots BAC will reach

~110 mg/dl (23 mM) in about an hour after consumption

Legal intoxication in the US (0.08%) corresponds to 17 mM Levels at 43.2 mM may lose consciousness and above 86 mM is fatal

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First Pass Metabolism

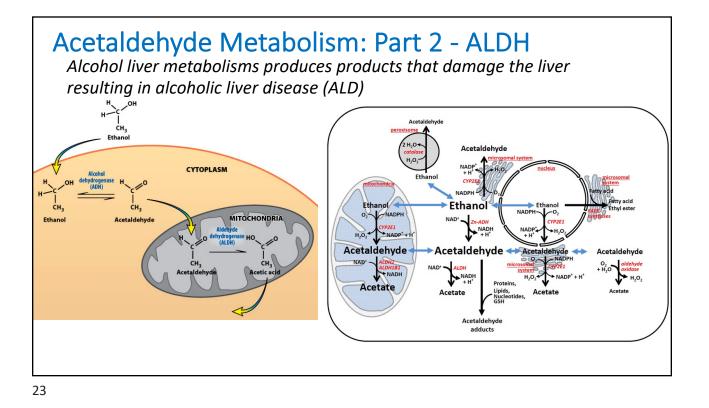
- ADH1C: Low Km and moderate Vmax • Slow and steady EtOH metabolism
- ADH5: High Km (low affinity) med Vmax
- This enzyme is important right after drinking when EtOH conc can reach molar levels
- Helps prevent production of acetaldehyde until EtOH is present
 Those with ADH1B*2 mutations with high Vmax and low Km results
- in high levels of acetaldehyde at even moderate level of drinking
 Those with ADH1B*3 will have higher EtOH clearance w/
- potentially high levels of acetaldehyde but will not metabolize as much EtOH until high levels of BAC

Liver Metabolism

- Major forms ADH1A, ADH1B, ADH4
- Covers a range of Km and moderate clearance rates
- Allows quick and steady metabolism at all levels

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EtOH and FAS: Special Cases Alcohol consumption decreases folate (Vit B9) and B12) limiting DNA methylation and Liver ADH is very low in fetus and elimination of production of important cofactors. Increased acetate from EtOH metabolism EtOH in fetal tissues is slow (FAS). increased histone acetylation in areas of brain - There is some evidence that mothers homozygous that may lead to memory, learning and addiction with ADH1B*1 (low km low Vmax) mutations had a higher risk of children with FAS. - These mothers report higher drinking at and after conception (2.45 vs 1.82 drinking days/week) - Slow metabolism in mother leads to sustained EtOH throughout body including fetus - Mothers and infants who lack ADH1B*3 (high Km high Vmax) and drink are at greater risk of FAS syndromes (compared to ADH1B*3 who didn't drink or ADH1B who did drink) **ADH1B*2** (low Km high Vmax) has a high frequency in China and Japan (~50%). - Heterozygous mutations reduced the risk of alcoholism - Homozygous was even more protective from alcohol abuse - Presumably because of high levels of acetaldehyde which causes several negative outcomes including – hangovers! **ADH1B*3** (high Km high Vmax) is found mostly in African origin people. They metabolize EtOH faster than other but only when high levels of EtOH. This is associated with reduced risk of alcohol dependence





NAD ⁺ NADH	H + H [⁺] NAD [⁺] NAD	H + H⁺
	O CH3 ALDH	
Ethanol	Acetaldehyde	Acetic acid

ALDH catalyzes the NAD(P)+ dependent oxidation of an aldehyde to a carboxylic acid.

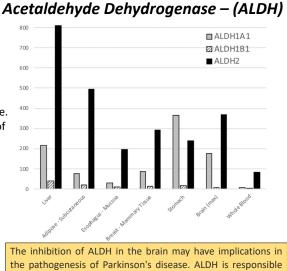
There are 19 ALDH genes grouped into 9 families, each coding for ALDH with a broad specificity of substrate and expression level / tissue.

- ALDH1A1, ALDH1B1 and ALDH2 are most relevant to metabolism of acetaldehyde.

ALDH highest expression in liver with significant amounts found in adipose, stomach and brain.

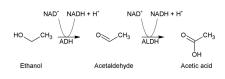
ALDH1A1 is cytosolic. ALDH1B1 & ALDH2 are mitochondrial

Gene Name	Km (mM)	Max (min ⁻¹)
ALDH1B1	55.0	655
ALDH1A1	180.0	380
ALDH2	0.2	280
ALDH2*1. (Glu504)	х	Х
ALDH2*2. (Lys 504)	Х	Х



the pathogenesis of Parkinson's disease. ALDH is responsible for the detoxification of acetaldehyde containing metabolites of dopamine and build up of these metabolites can cause neurotoxicity.

Acetaldehyde Metabolism: Part 2 - ALDH Acetaldehyde Dehydrogenase – (ALDH)



The main isozyme responsible for detoxification is ALDH2

- Low Km (very high affinity) and high reaction velocity (Vmax) rapidly eliminates most of the acetaldehyde as fast as ADH can produce it.
- Any mutation to E504 (results in a large increase in the K_m for NAD⁺, which almost inactivates the enzyme.
- About half of Taiwanese, Han Chinese and Japanese populations have high levels of acetaldehyde after drinking.
- People with both heterozygous and homozygous mutated ALDH2 show virtually no acetaldehyde metabolism and experience negative physiological responses to alcohol
- Alcoholic cirrhosis is reduced by 70% in this population as well as low esophageal and had and neck cancers.

Acute (short term) effect of Acetaldehyde:

Elevated blood acetaldehyde causes facial flushing, severe headache, palpitations, tachycardia, hypertension, respiratory distress, nausea and vomiting. These symptoms commence within 15–30 min of ingesting alcohol and persist for several hrs.

Heterozygote for ALDH2*



The alcohol flush reaction. The primary feature of the alcohol flush reaction is a red face—or flush—but it can also be accompanied by hives, nausea, low blood pressure, the worsening of asthma, or an episode of migraine.

Alcohol flush is due to high acetaldehyde in people with ALDH2 mutations

 Due to the unpleasant effects, these people are protected from drinking

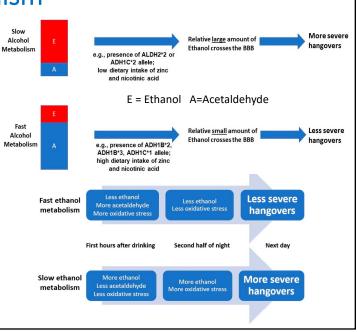
Acetaldehyde Metabolism

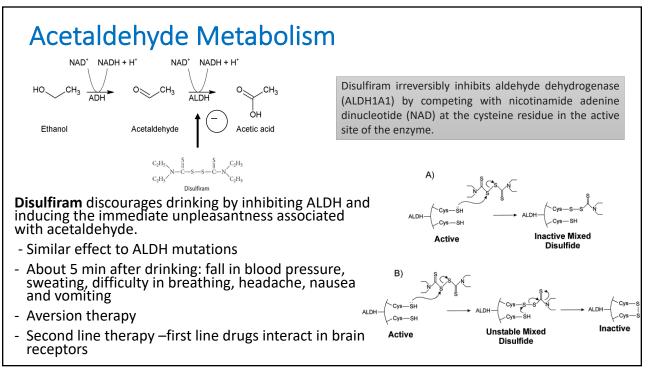
The Hangover – not the movie

- Methanol in some alcohols is slowly metabolized to formaldehyde by ADH.
- Presence of formaldehyde and acetaldehyde along with dehydration and hypoglycemia lead to hangovers

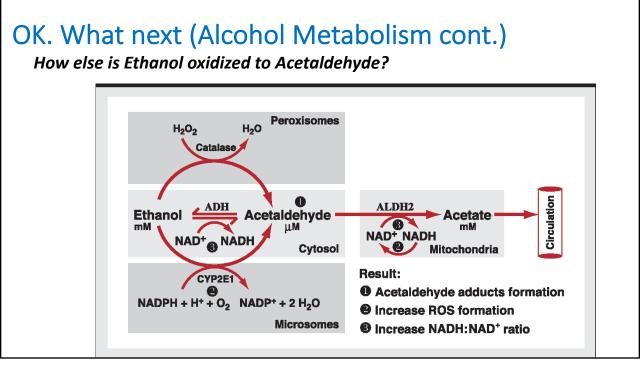
Data suggest that the ethanol elimination rate is a critical determinant of hangover severity

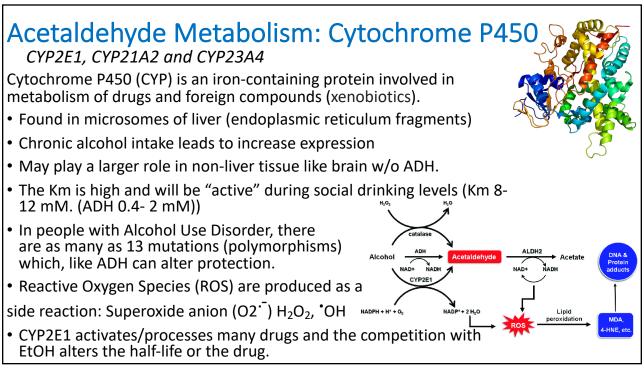
- Significant correlations have been found between ethanol concentration (but not acetaldehyde) and hangover severity.
- ROS metabolism of ethanol leads to oxidative stress and high inflammatory response to alcohol consumption.
- Nutrients, microbiota, and hangover treatments that speed up the conversion of ethanol into acetaldehyde are associated with less severe hangovers.



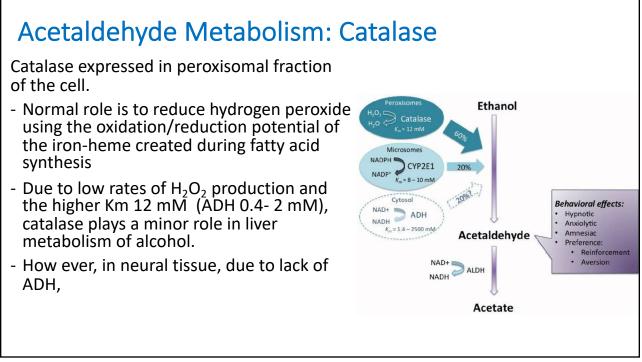


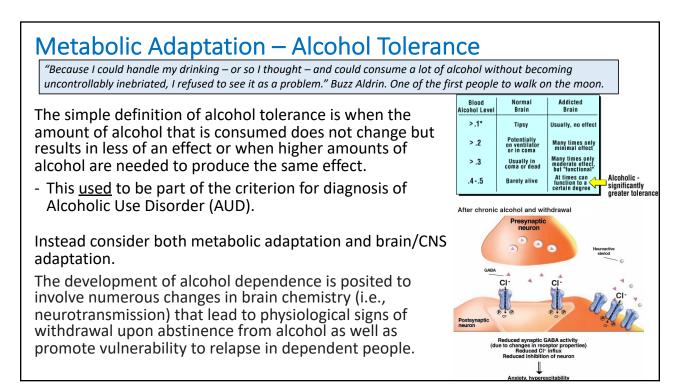


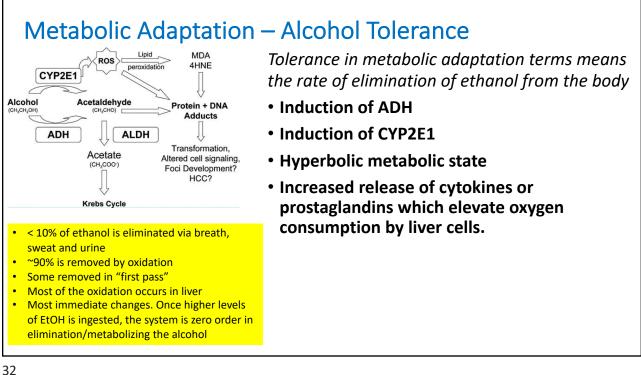




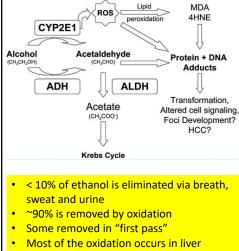








Metabolic Adaptation – Alcohol Tolerance



 Most immediate changes. Once higher levels of EtOH is ingested, the system is zero order in elimination/metabolizing the alcohol

Tolerance in metabolic adaptation terms means the rate of elimination of ethanol from the body.

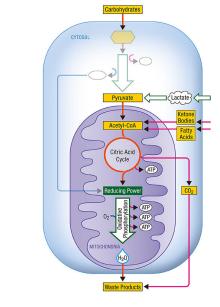
Induction of ADH – chronic alcohol ingestion can decrease thyroid hormones, one of the regulators of ADH expression. Yet some studies show that ADH expression can increase depending on intake.

Induction of CYP2E1

Hyperbolic metabolic state

Increased release of cytokines or prostaglandins which elevate oxygen consumption by liver cells.

Metabolic Adaptation – Alcohol Tolerance



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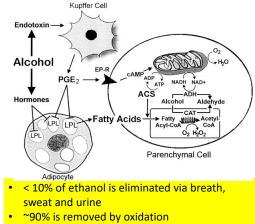
Induction of ADH

Induction of CYP2E1

Hyperbolic metabolic state – Increased activity of ATPase O_2 consumption and NADH oxidation to NAD⁺. This can also lead to hypoxia and liver damage.

Increased release of cytokines or prostaglandins which elevate oxygen consumption by liver cells.





- Some removed in "first pass"
- Most of the oxidation occurs in liver
- Most immediate changes. Once higher levels of EtOH is ingested, the system is zero order in elimination/metabolizing the alcohol

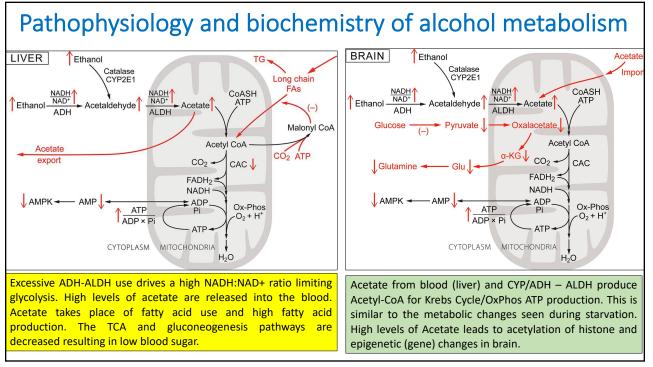
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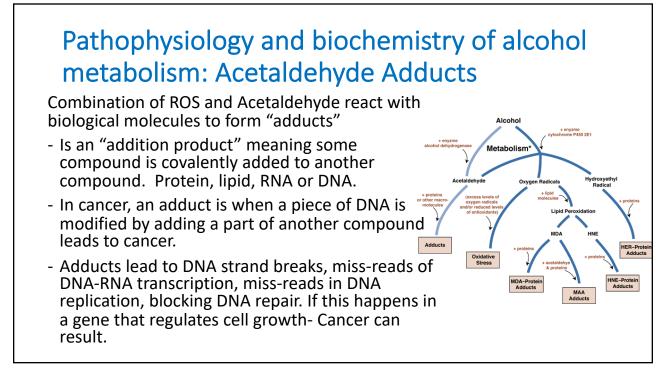
Induction of ADH Induction of CYP2E1

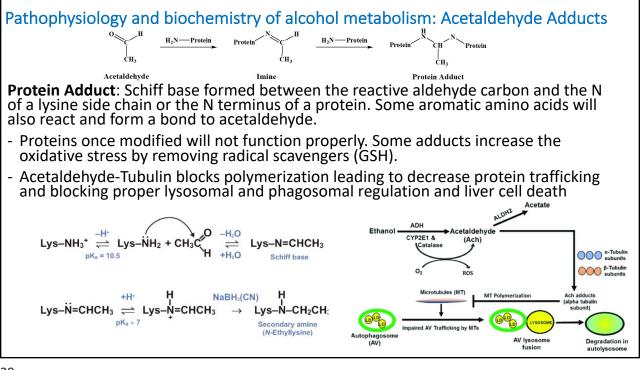
Hyperbolic metabolic

Increased release of cytokines or prostaglandins which elevate oxygen consumption by liver cells- Ethanol increases release of hormones, cytokines and prostaglandins to further increase metabolism and increase oxygen consumption



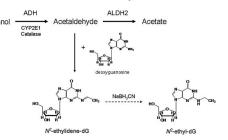


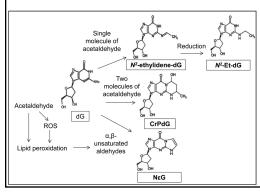




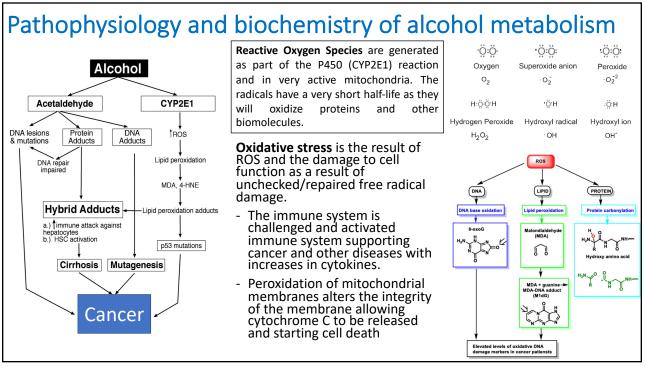
Pathophysiology and biochemistry of alcohol metabolism: Acetaldehyde Adducts

DNA Adduct: The formation of acetaldehyde-derived DNA adducts plays an important role in carcinogenesis. The <u>major</u> DNA adduct in the human body is a Schiff base N^2 -ethylidene-2'-deoxyguanosine (N^2 -ethylidene-dG) adduct. In reactions of acetaldehyde with deoxyribonucleosides, deoxyguanosine (dG) was most reactive, though products are also observed in reactions with deoxyadenosine (dA) and deoxycytidine (dC).





- N^2 -Et-dG blocks DNA synthesis and induces frameshift deletions and G:C > T:A transversions.
- N^2 -Et-dG can rotate around the exocyclic nitrogen and the alpha carbon of acetaldehyde because it has a single bond, whereas N^2 -ethylidene-dG has a double bond, which makes it more hydrophobic than N^2 -Et-dG. These differences may result in significantly different mutagenic potential between N^2 -Et-dG and N^2 -ethylidene-dG.
- CrPdG induces DNA interstrand and intrastrand cross-links. The ring-opened form of CrPdG can react with dG on the opposite strand of the DNA to form DNA interstrand cross-links.
- CrPdG-mediated disruption of the DNA replication process is thought to cause DNA damage



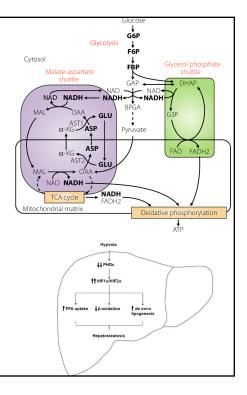
Hypoxia and Liver Damage

Excessive ADH and ALDH reduce the cell's cytosolic and mitochondrial NAD⁺ stores to NADH

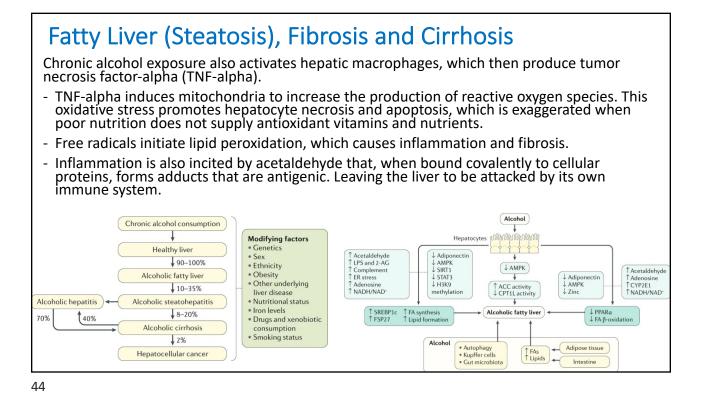
- NADH is oxidized in mitochondria as part of the electron transport system / ATPase (O₂ utilizing)
- The results are high oxygen consumption in liver. Liver cells close to the arterial supply will strip the blood of much of the oxygen leaving areas low of oxygen (hypoxia)
- Hypoxic tissue show damage in chronic alcohol consumption.

Non-alcoholic fatty liver disease is the most common chronic liver disease along with alcohol liver disease (ALD). In both cases, HIF1 α , a transcriptional protein, activates a series of genes leading to cell death in hypoxia.

- ROS activates lipopolysaccharides which stabilize HIF1 α , leading to greater damage.



Fatty Liver (Steatosis), Fibrosis and Cirrhosis The first stage of liver damage following chronic alcohol consumption is the appearance of fatty liver, followed by inflammation, apoptosis, fibrosis and finally cirrhosis. - Lipid oversupply leads to liver increasing uptake of fatty acids and the production of new fatty acids from acetate. Hepatic Steatosis (Fatty liver) - Acetaldehyde induced increases in the NADH/NAD+ ratio and increases in NADPH (from P450) and other changes (AMP) promotes further fatty acid synthesis. - As a part of the process VLDL (lipid vesicles) are produced but not released from liver cells, leaving an enlarged and fatty liver. Fatty live

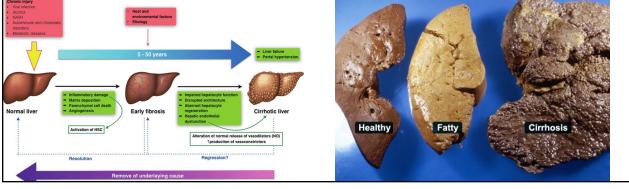


Fatty Liver (Steatosis), Fibrosis and Cirrhosis

Continued chronic alcohol exposure will cause a subset of steatosis livers to develop fibrosis and then cirrhosis.

- Fibrosis is a result of hepatocyte death. This causes the release of tumor necrosis factor and other hormones that cause damaged and normal cells express and secrete extracellular matrix proteins such as collagen and fibronectin. Essentially making the liver a scar tissue.

- Many of the protein adducts generated with acetaldehyde and ROS turn on these profibrotic pathways. Hypoxia supports the further stimulation of fibrosis.



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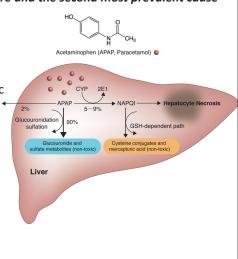
Alcohol and Acetaminophen (Tylenol) Toxicity

Acetaminophen toxicity is one of the most common causes of both intentional and unintentional poisoning in the United States. It has become the most common cause of acute liver failure and the second most prevalent cause of liver failure requiring transplantation.

Upon ingestion, acetaminophen is rapidly absorbed from the gastrointestinal (GI) tract and quickly distributed throughout the body. Peak plasma concentrations are achieved within 30 to 60 min. The half-life of acetaminophen is approximately 2 to 3 hours after therapeutic doses, yet can be increased to more than 4 hours in patients with hepatic injury.

Acetaminophen is extensively metabolized by the liver via three main[®] hepatic pathways: glucuronidation, sulfation, and **CYP2E1** oxidation.

- Approximately 90% of acetaminophen is conjugated to sulfated & glucuronidated metabolites that are eliminated.
- Of the remaining acetaminophen, approximately 2% is excreted unchanged in the urine and the rest undergoes CYP2E1-mediated oxidation to form a reactive imine (NAPQI).



Alcohol and Acetaminophen (Tylenol) Toxicity

With large acute doses or with chronic use, the major metabolic pathways—the glucuronide and sulfate conjugation systems become saturated, and more acetaminophen is metabolized by the CYP2E1 system. When glutathione is approximately 70% depleted, NAPQI begins to accumulate in the hepatocytes, resulting in hepatic damage.

Acetaminophen overdose here is defined as intake of more than 5gm of APAP in one sitting APAP when ingested, is rapidly absorbed in the body leading to the synthesis of toxic intermediate NAPQI within hepatocytes by several P450

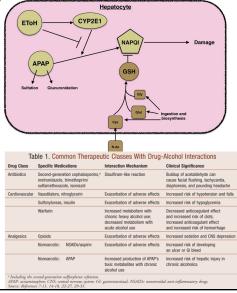
Under normal circumstances, this toxic metabolite reacts with sulfhydryl groups in glutathione, converting it to harmless metabolites before being excreted in the urine.

NAPQI then binds with glutathione (GSH), a well regulated antioxidant found within hepatocytes and is easily eliminated. However when a high dose of acetaminophen is ingested, NAPQI becomes the key pathway and exhausts the GSH(G) reserve and unconjugated NAPQI binds to hepatocellular proteins and other cellular components, damaging hepatocytes.

Patients with AUD or excess alcohol intake induce PY2E1 speeding up the production of NAPQI leading

In 1986, six alcoholics who developed toxicity after taking APAP and reviewed nineteen similar case reports. In 1995, an additional 67 alcoholic patients who developed liver toxicity after taking moderate doses of APAP.

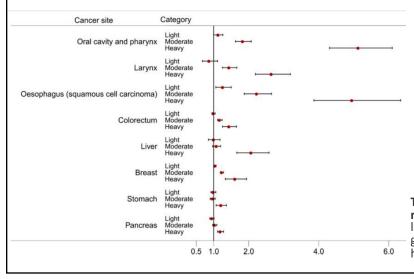
However, the possibility that APAP may be toxic in alcoholics at moderate doses has been disputed The problems in obtaining proper controls makes it difficult to know if alcoholics are more susceptible to APAP hepatotoxicity.

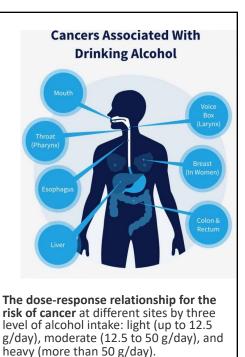


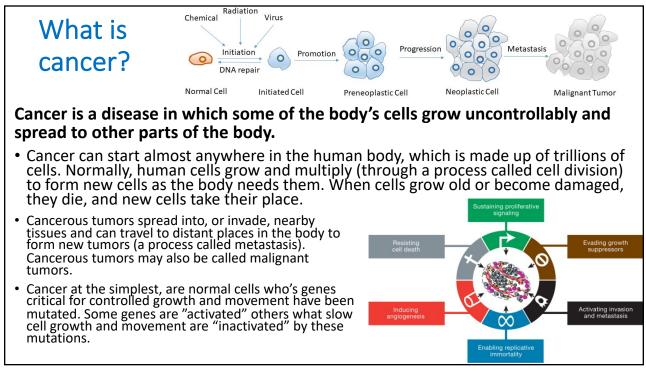
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Alcohol-Related Cancer

Approximately 4% of cancers worldwide are caused by alcohol consumption ~ 740,000 global deaths in 2020.







Alcohol-Related Mechanism of Cancer

- Genetic variations of ALDH2*2 (E487K) with almost no activity, is associated with a high incidence of liver cancer in heavy drinkers.
- ALD-associated cancer is high in populations with the disease who drink.
- Acetaldehyde and ROS combine to produce DNA adducts that lead to DNA double-strand breaks, sister chromatid exchanges and DNA cross-links
- Decreased methylation of DNA can activate some cancer causing genes

