



Can pharmacogenetic enhance our understanding and management of Opioid Use Disorder?

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Objectives

- Explain the impact of the COVID-19 pandemic on the opioid crisis in Ohio and United States
- Describe the chemical modifications made to fentanyl
- Describe the effects of opioids on the dopamine reward pathways in the brain
- Explain the pharmacogenetics in the context of OUD
- Discuss the Ohio pharmacogenetic study of OUD

National Data: Opioid Overdose Deaths

U.S. Drug Overdose Deaths

- More than 100,000 deaths from June 2020 to May 2021
- ~170 fatalities/ day primarily in those 18 to 45
- Primary driver is illicit forms of fentanyl
- Estimated annual economic cost to the U.S. is \$1 trillion
- Drug traffickers in Mexico produce most of the counterfeit tablet

New Challenges to the United States

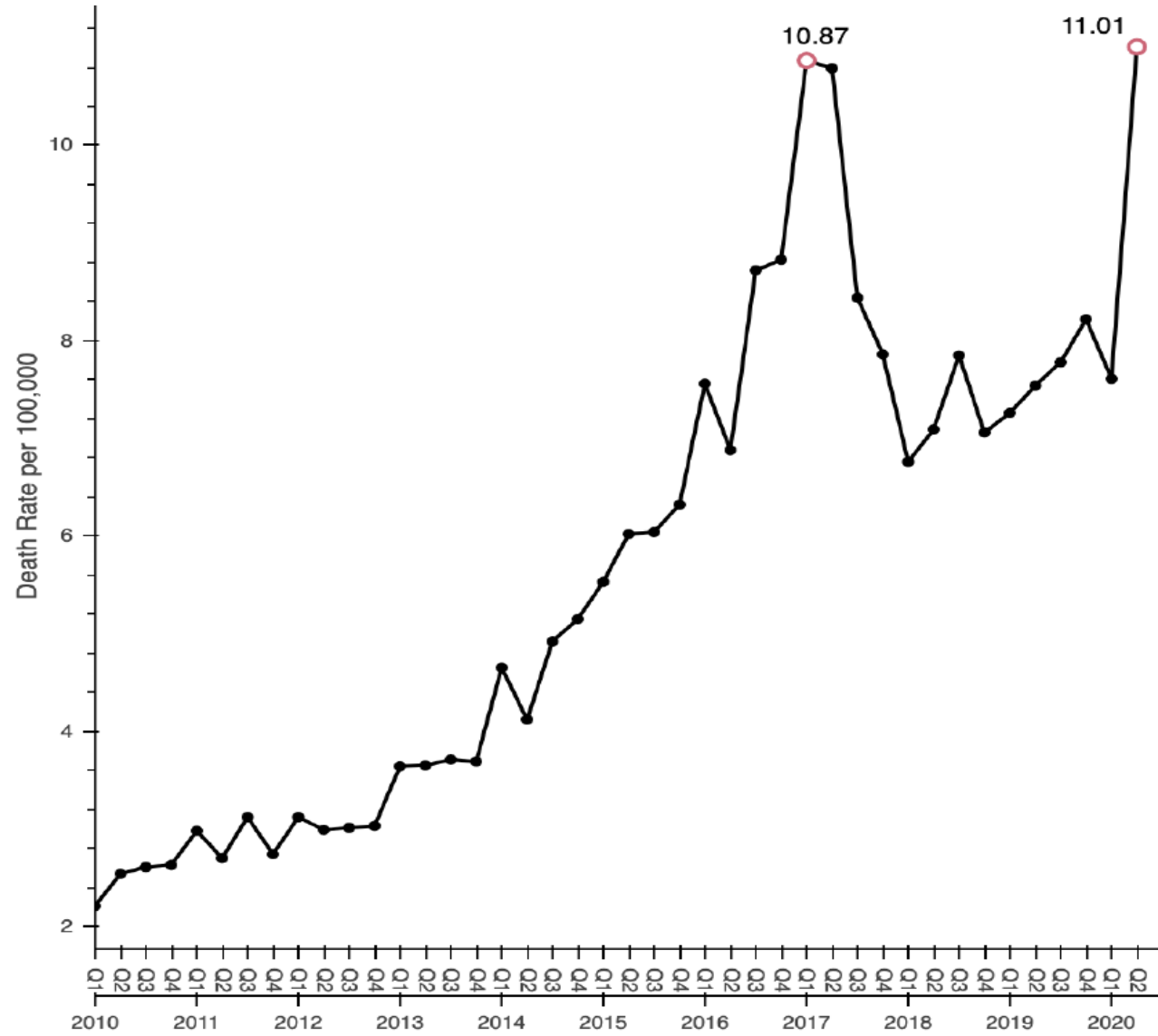
- Illegal drug manufacturing has become easier to conceal by moving from the field to the laboratory.
- Serious geopolitical issues significantly impede actions to disrupt supply.
- Synthetic opioids are highly potent and easy to make, and small amounts can be transported for large profits.
- Social media and encryption platforms, as well as established logistics systems, make distribution difficult to disrupt.
- COVID-19

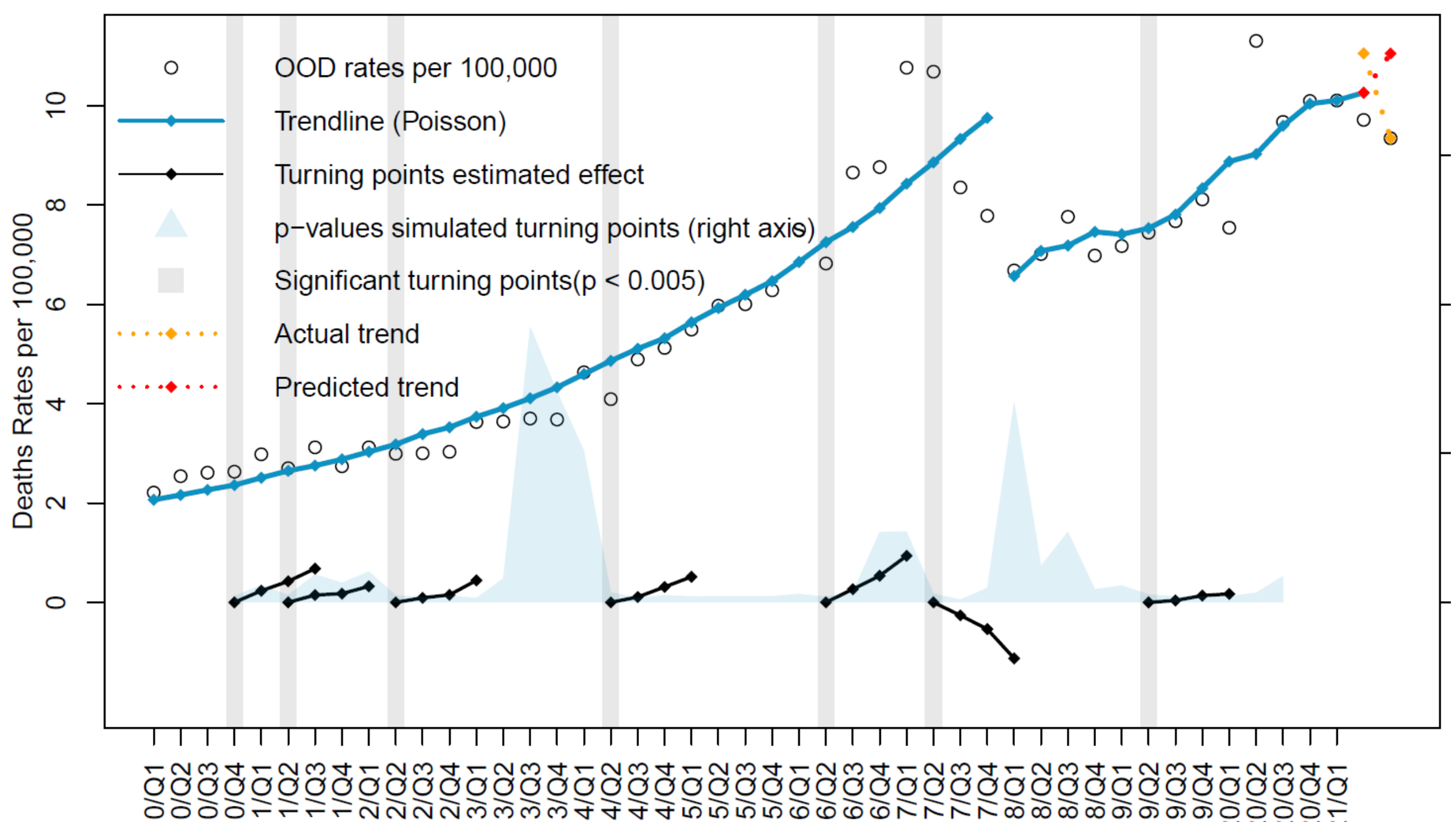
Ohio Data: Opioid Overdose Deaths

Methods

- The ODH database was queried
- Mortality dataset for deaths in Ohio related to opioids
- Temporal trend analyses were utilized to identify significant changes in trends of the quarterly OOD rate per 100,000 from Q1 of 2010 to Q2 of 2020.

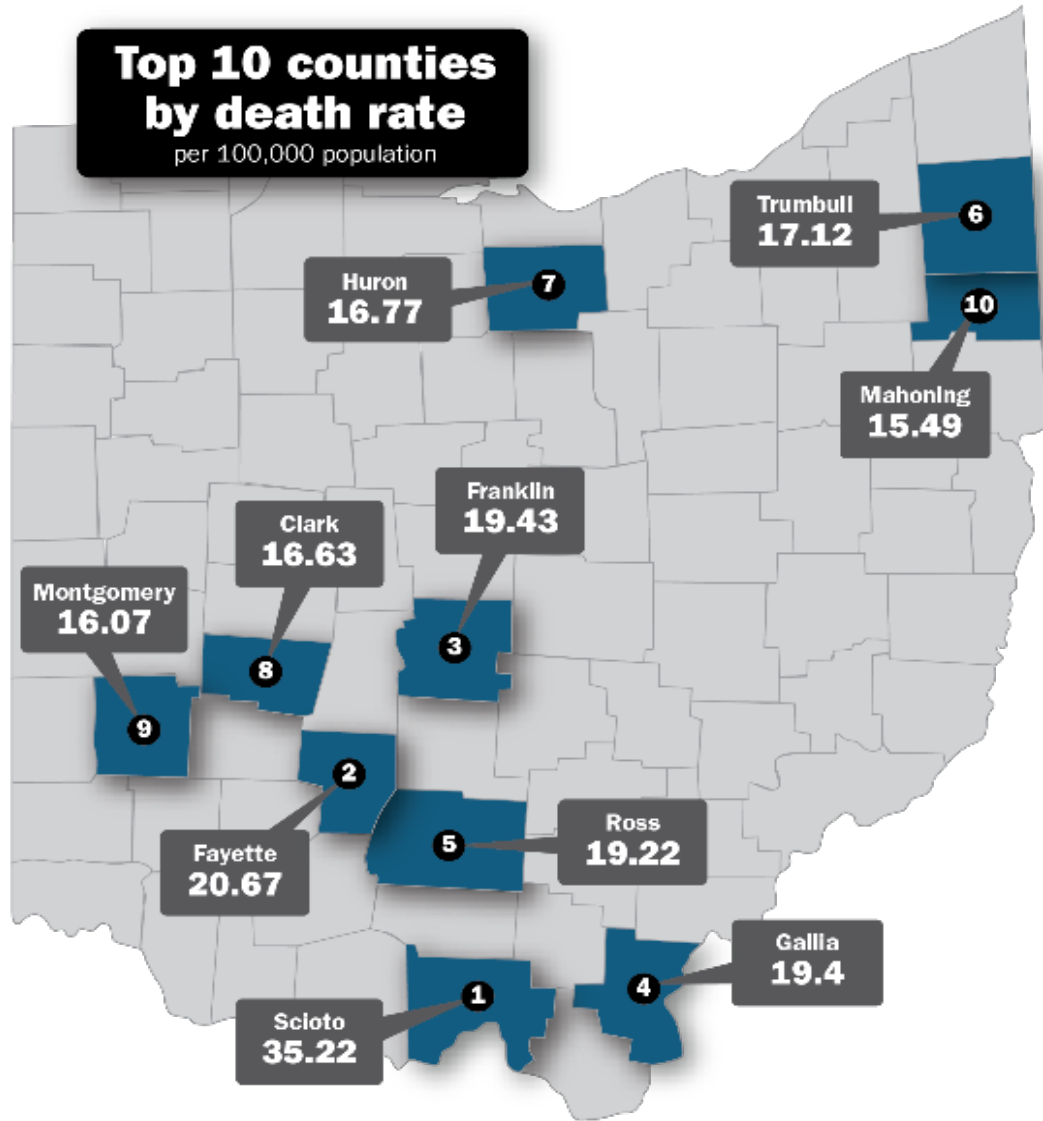
Results





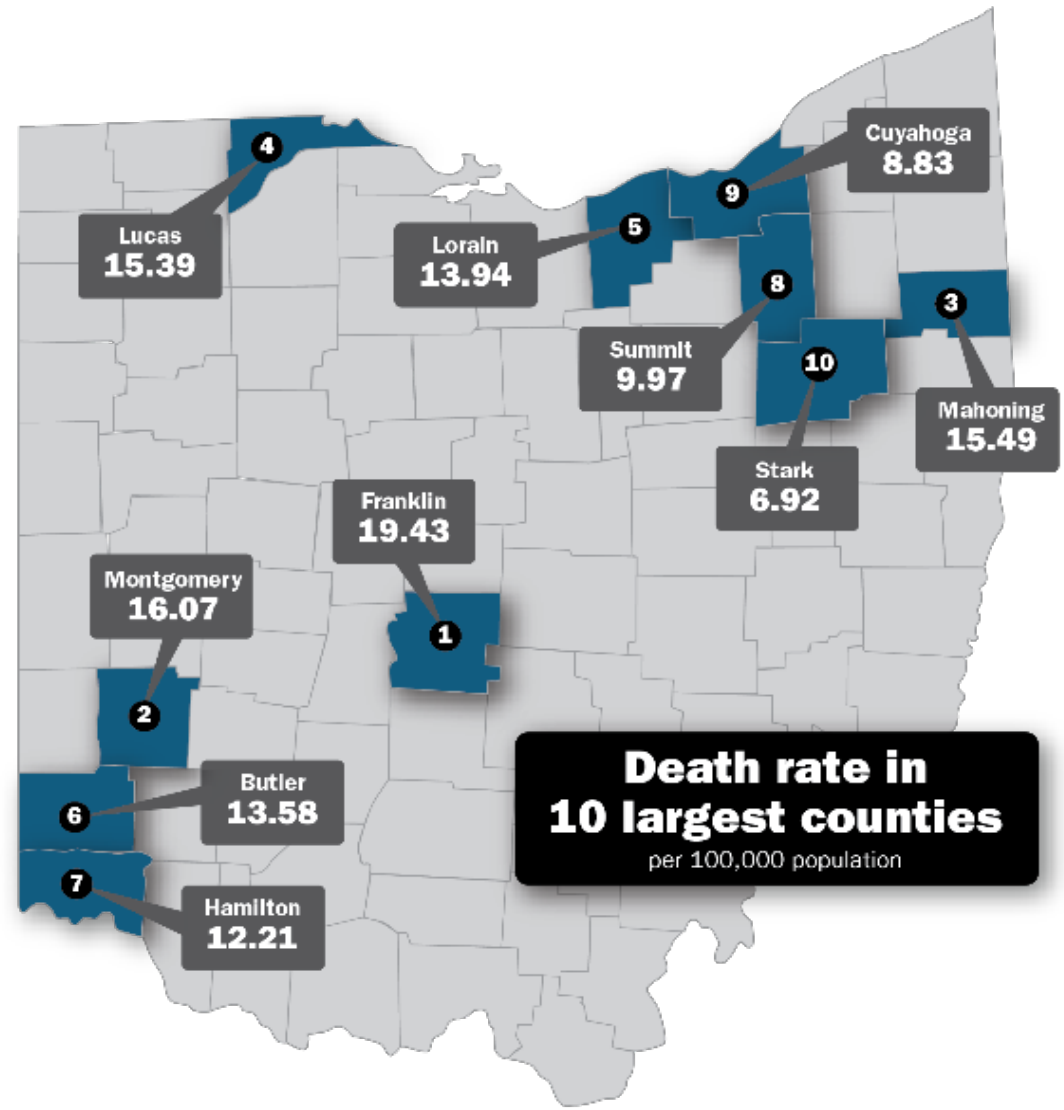
Top 10 counties by death rate

per 100,000 population

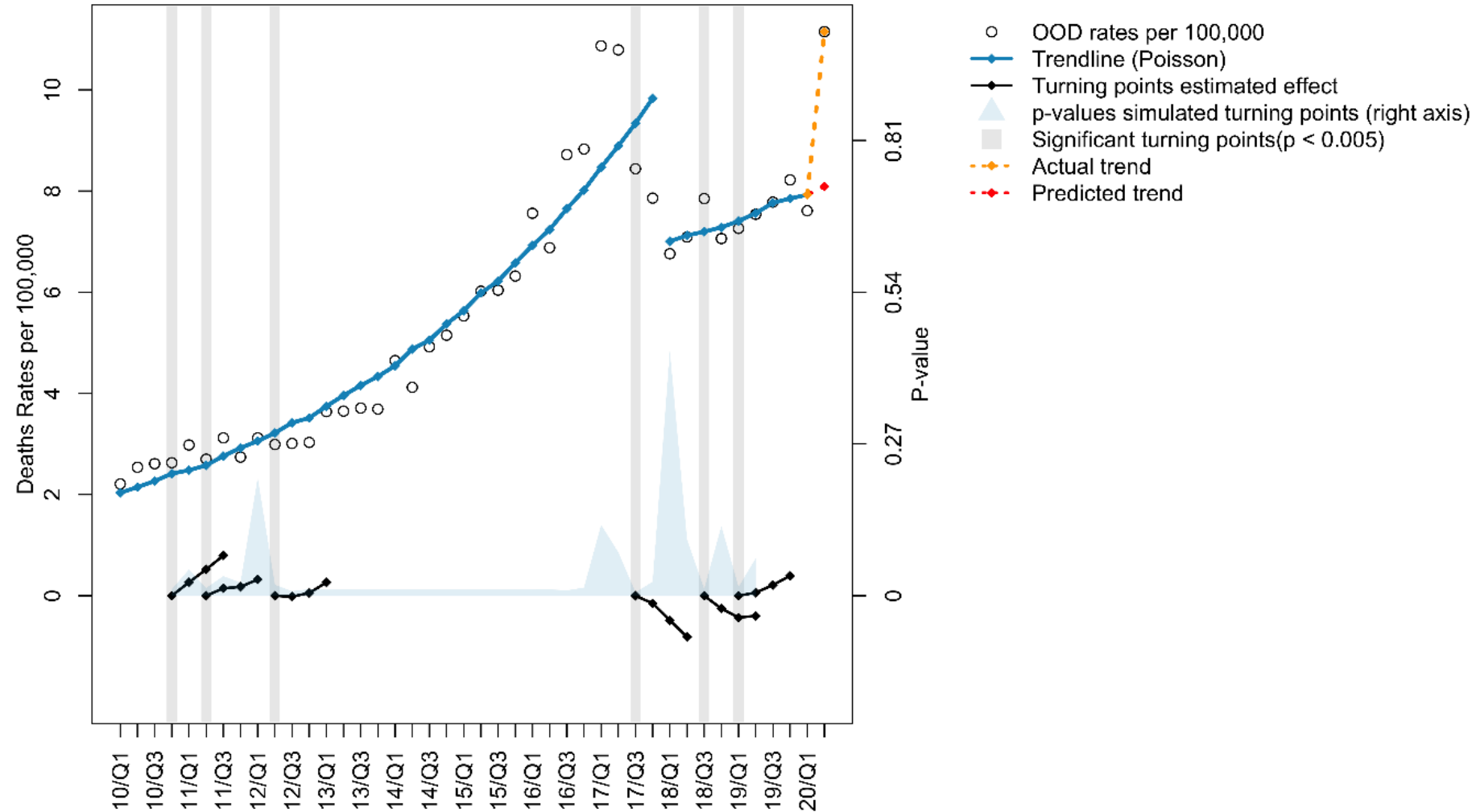


Death rate in 10 largest counties

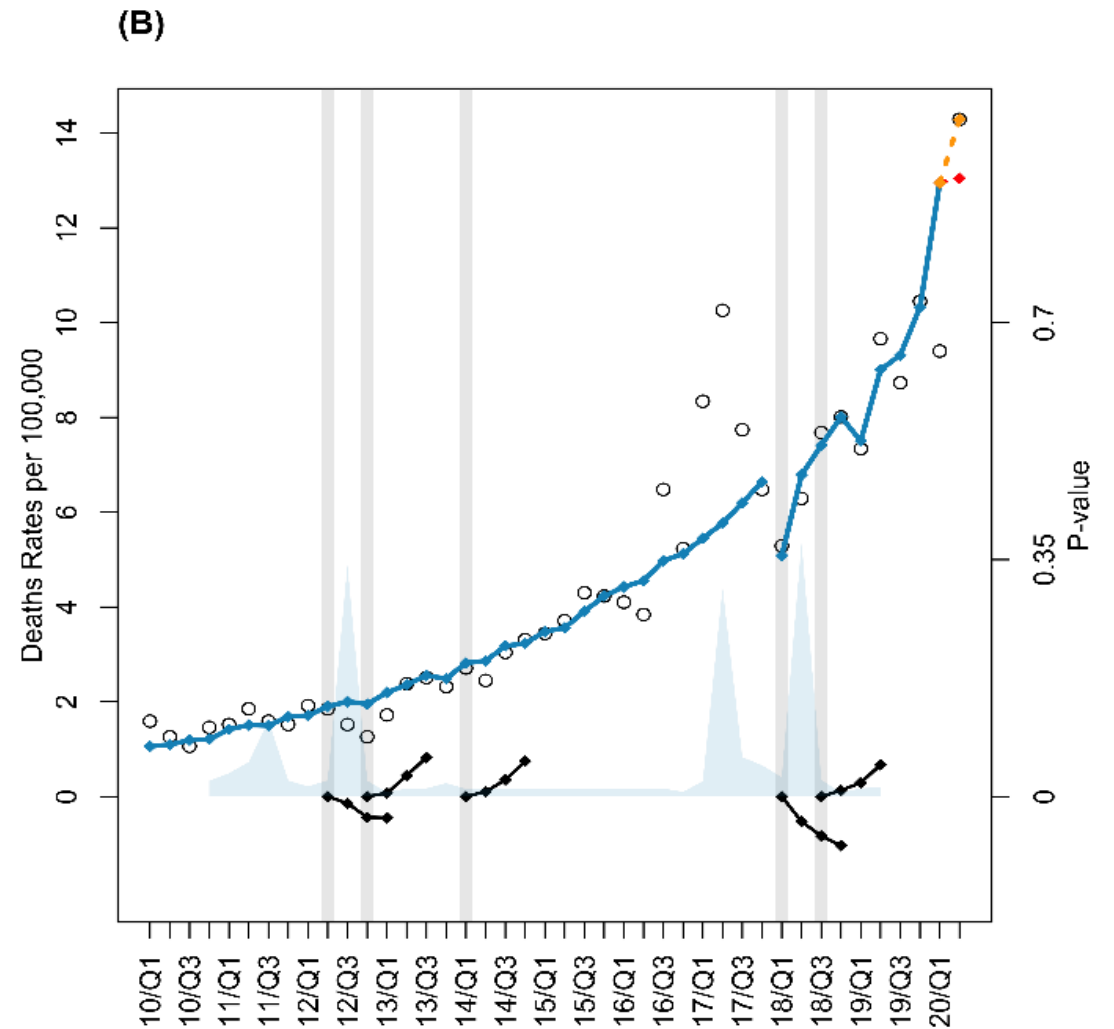
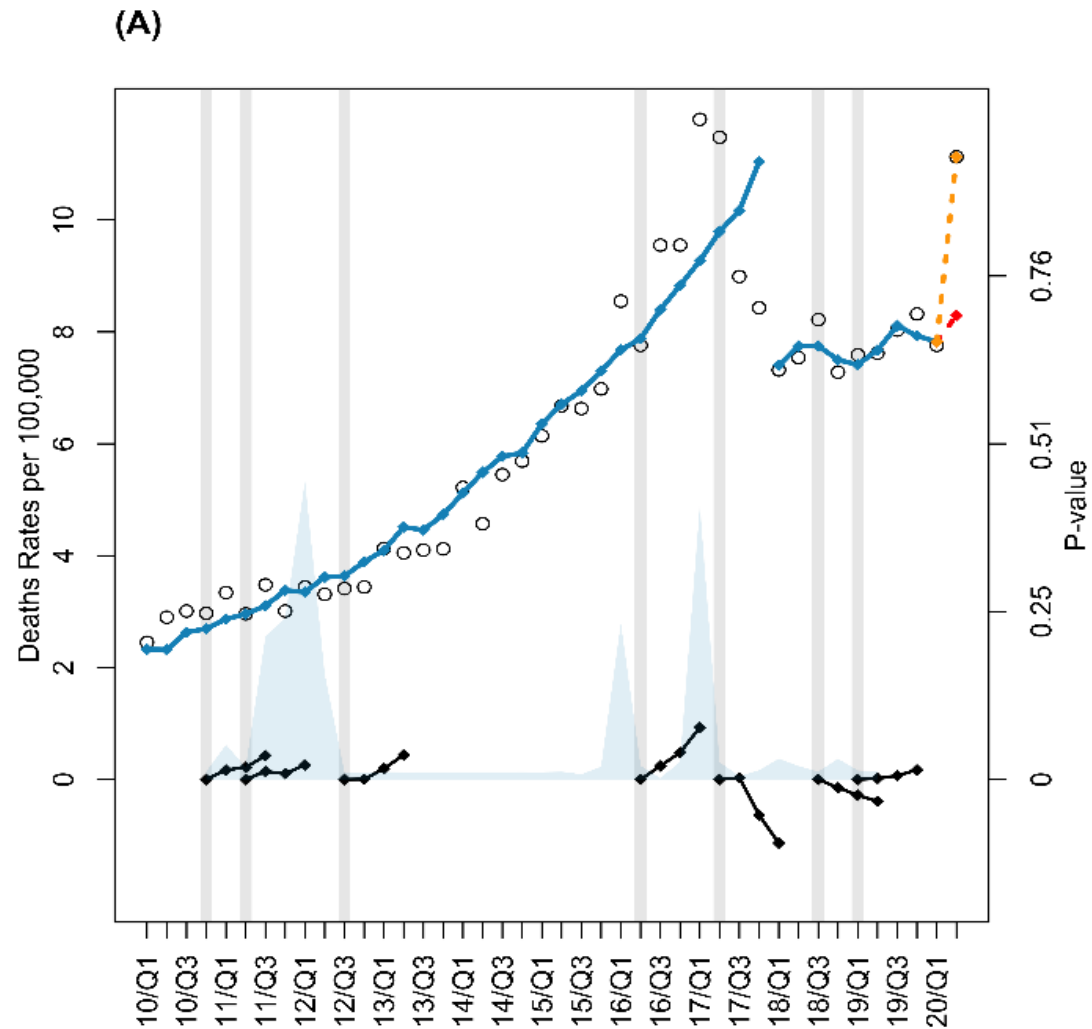
per 100,000 population



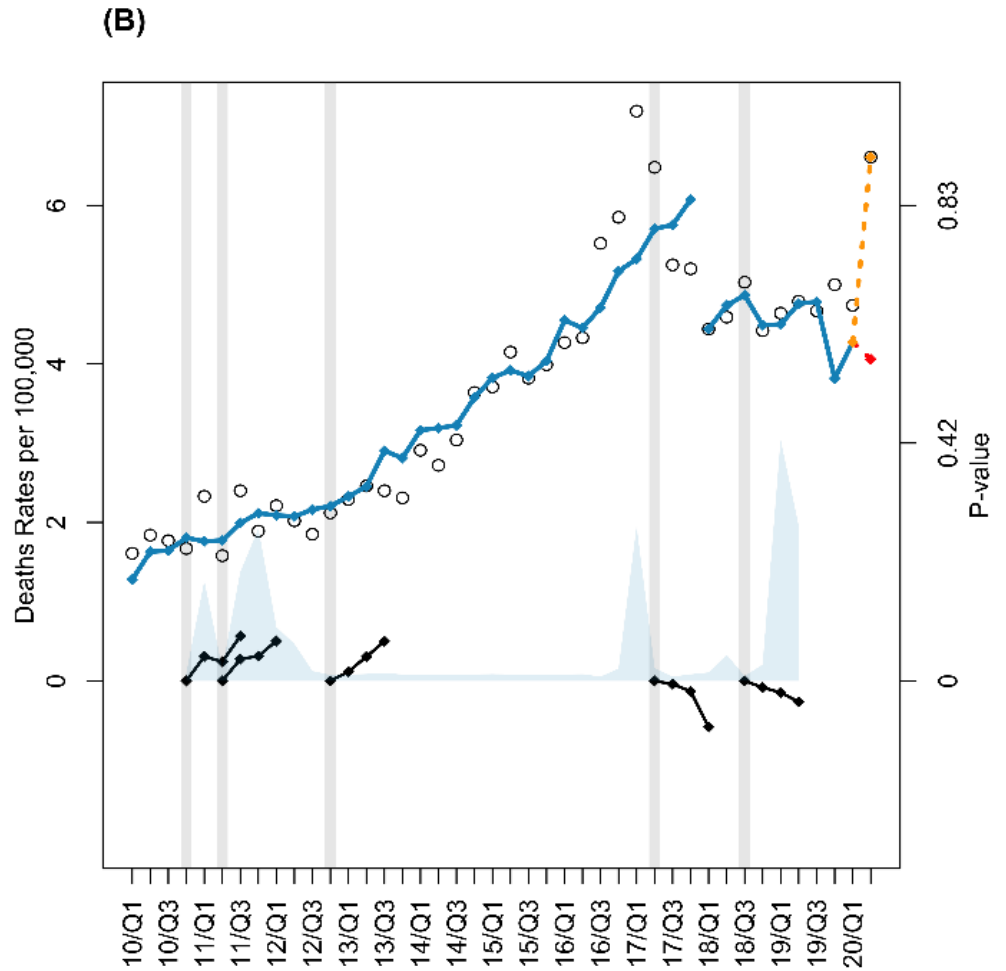
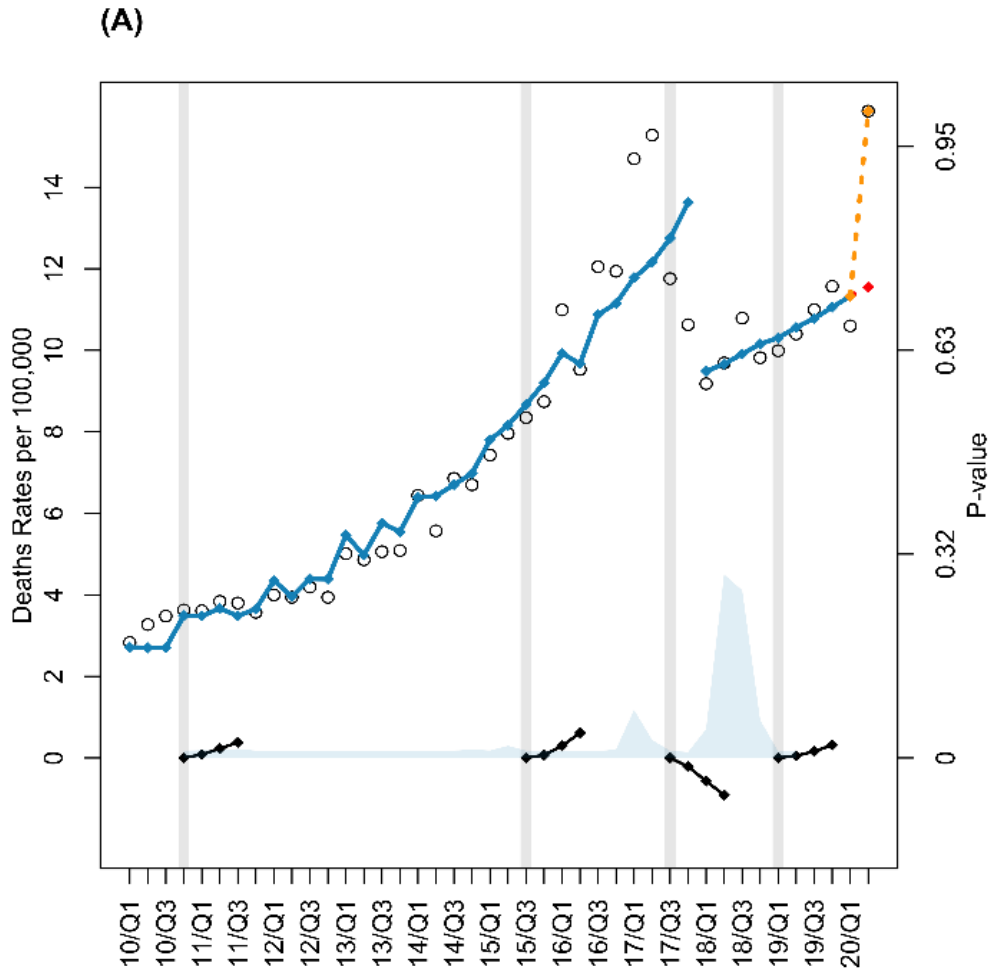
Temporal Trend Analysis of the Overall Quarterly OOD Rate per 100,000



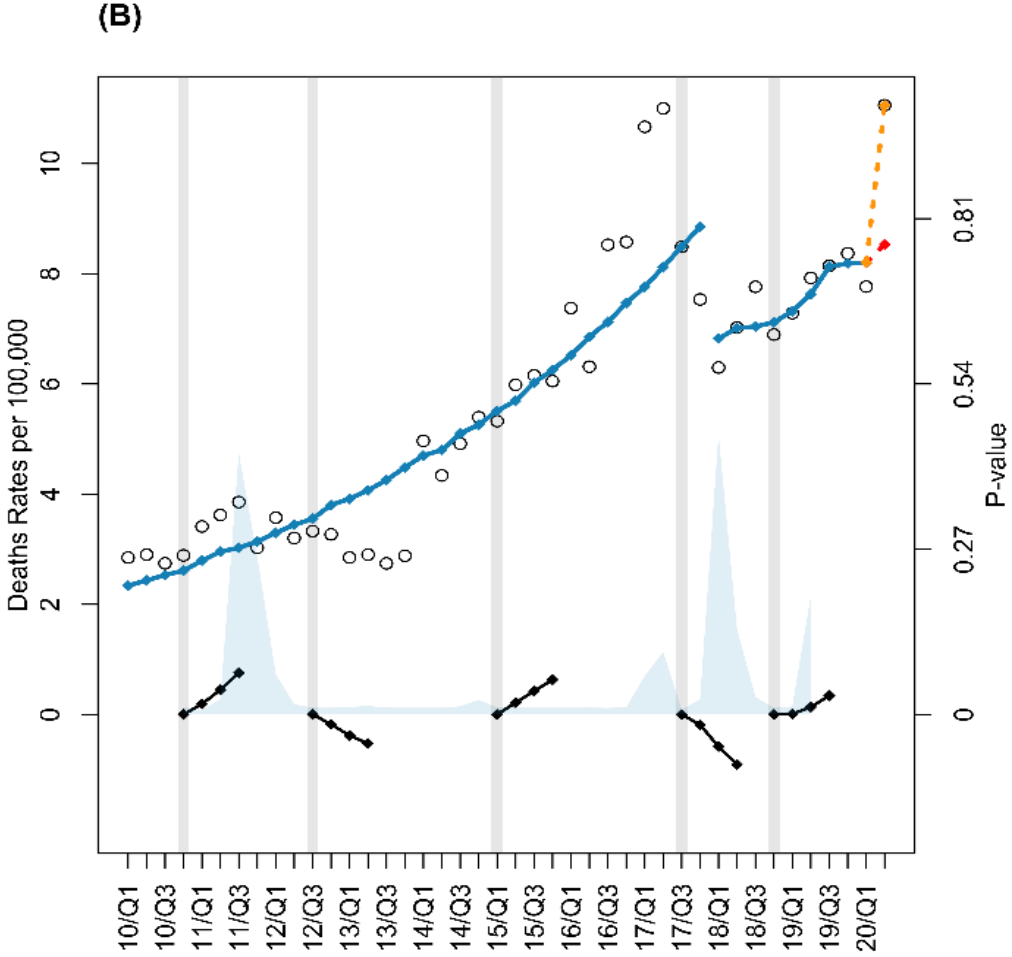
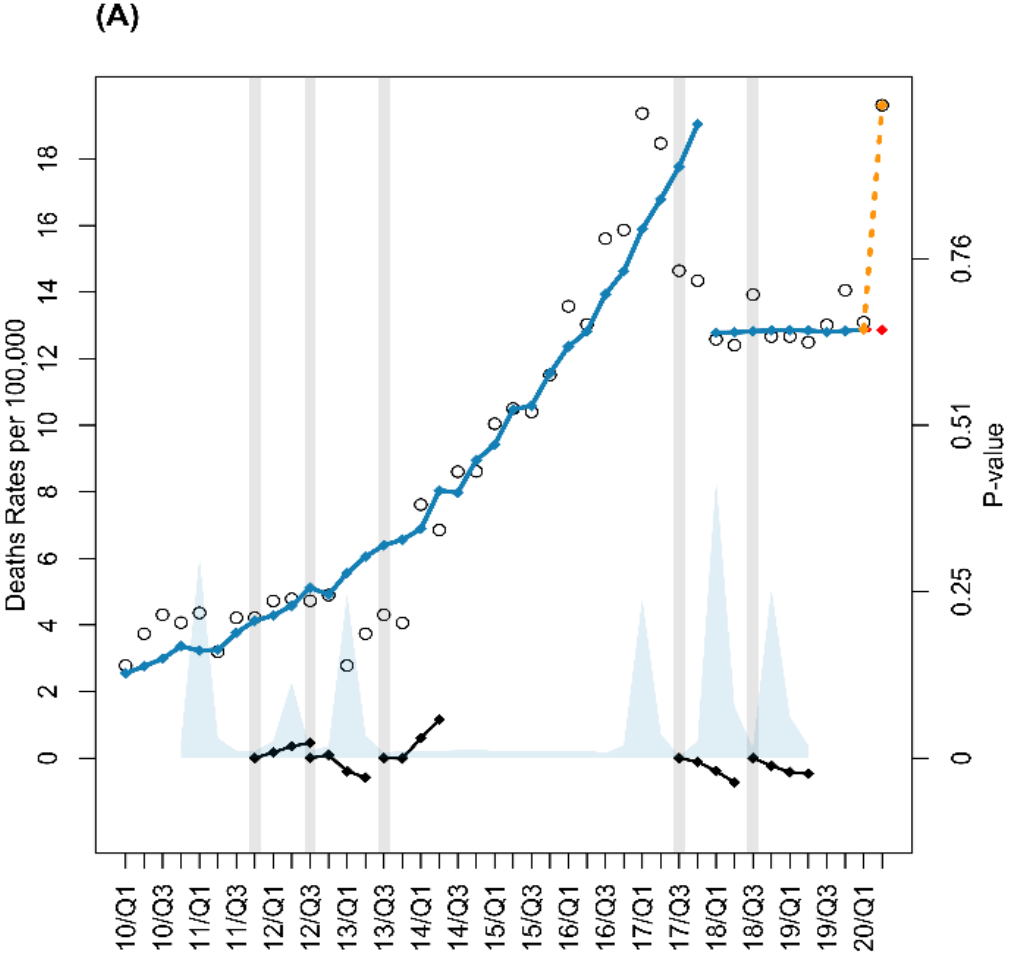
Temporal Trend Analysis of the Quarterly OOD Rate per 100,000 for the White (A) and Black (B) populations



Temporal Trend Analysis of the Quarterly OOD Rate per 100,000 for males (A) and females (B)



Temporal Trend Analysis of the Quarterly OOD Rate per 100,000 for age groups (A) 18-39 and (B) 40+

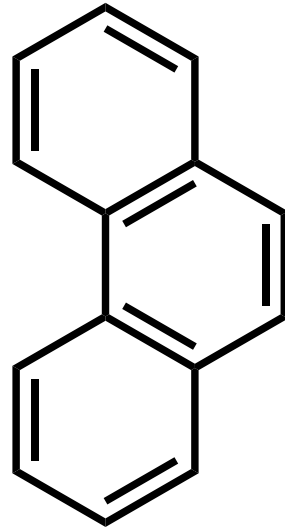


The Pharmacophore Rule

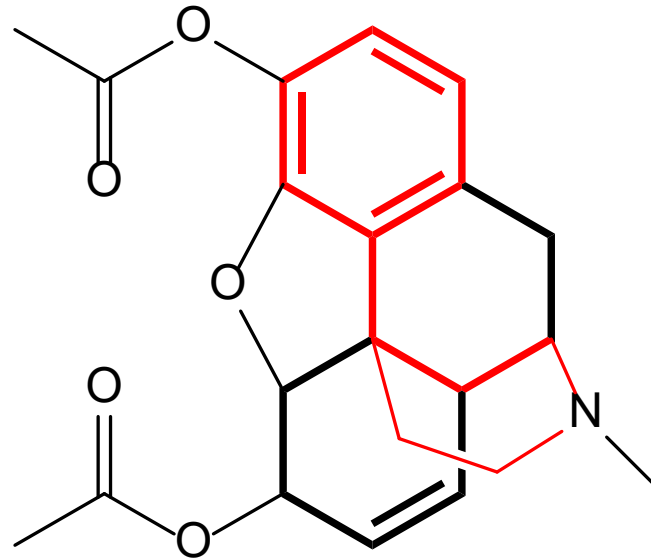
The Pharmacophore Rule was written so chemists would be able to identify the basic structural elements required for a compound to bind to their drug targets.

What is a pharmacophore?

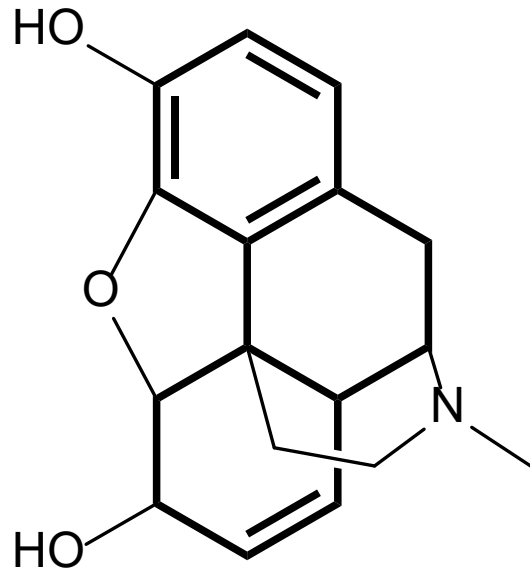
- the portion of drug molecule required for pharmacological activity



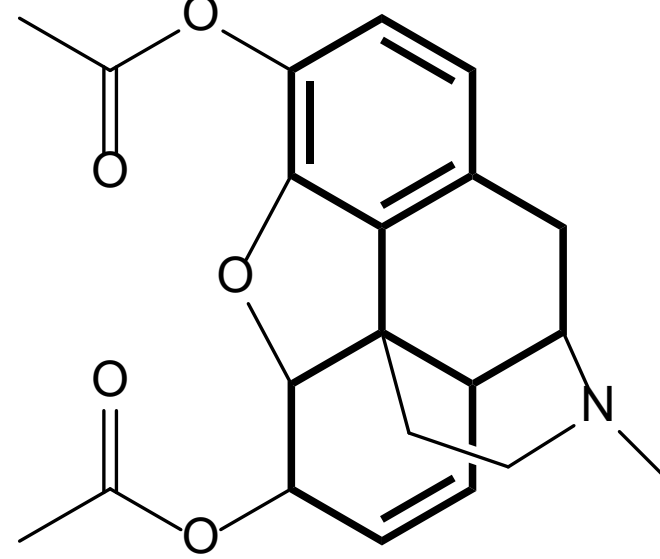
Phenanthrene



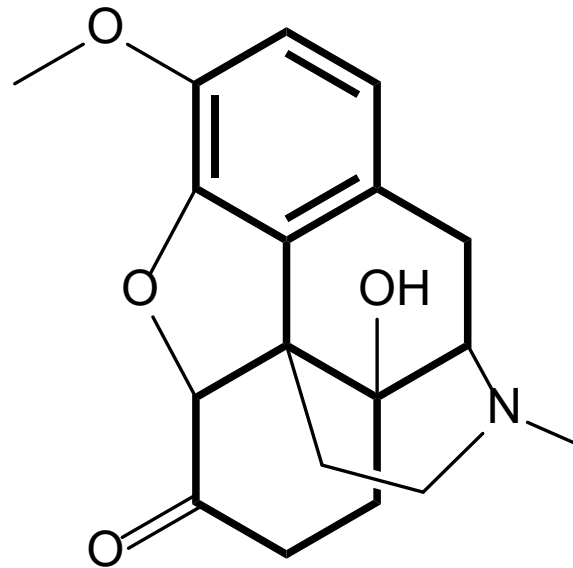
Heroin



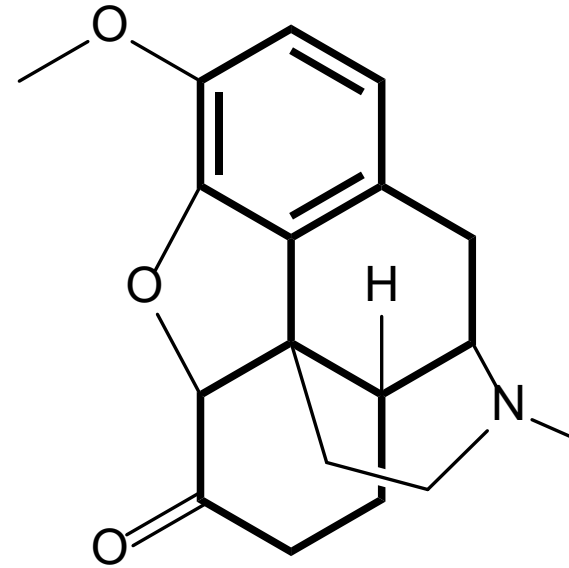
Morphine



Heroin

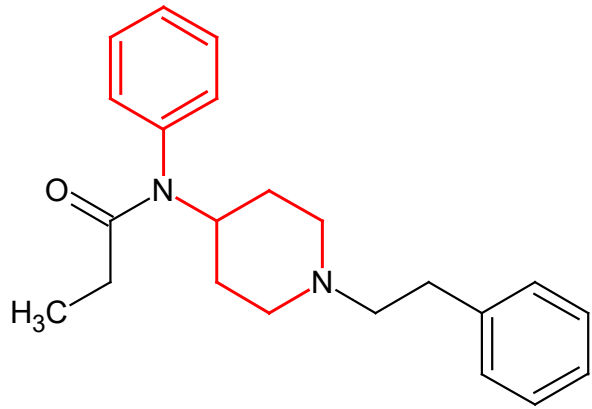


Oxycodone

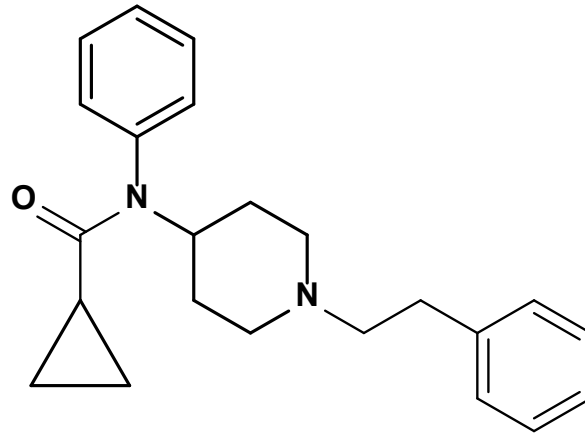


Hydrocodone

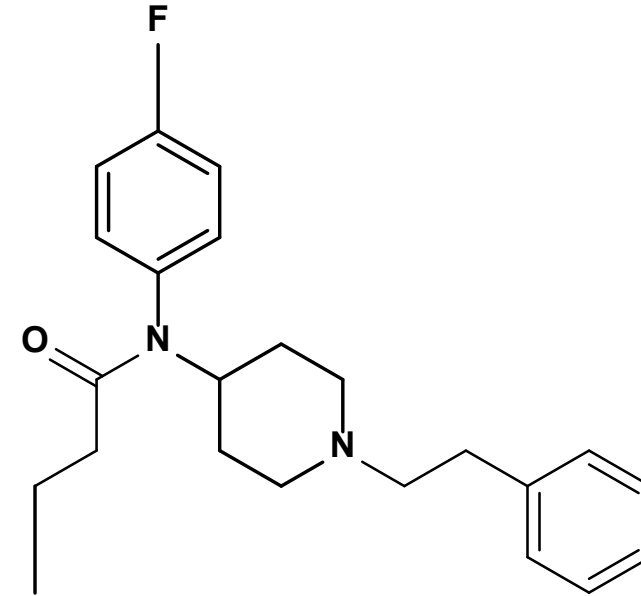
Example Fentanyl Pharmacophores



Fentanyl



Cyclopropyl fentanyl

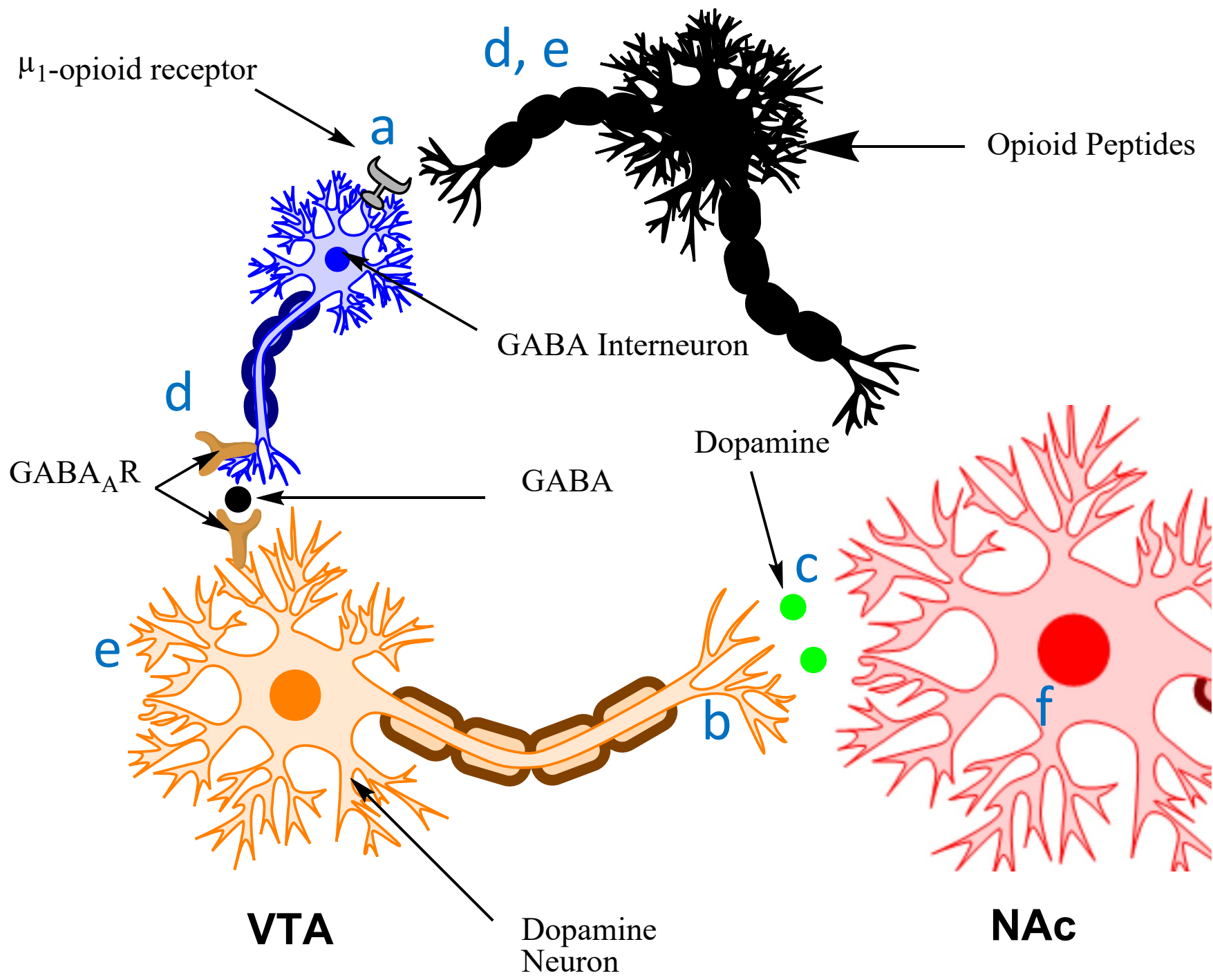


para-fluorobutyryl fentanyl

Dangerous Counterfeit Prescription Tablets



The Dopamine Reward Pathway

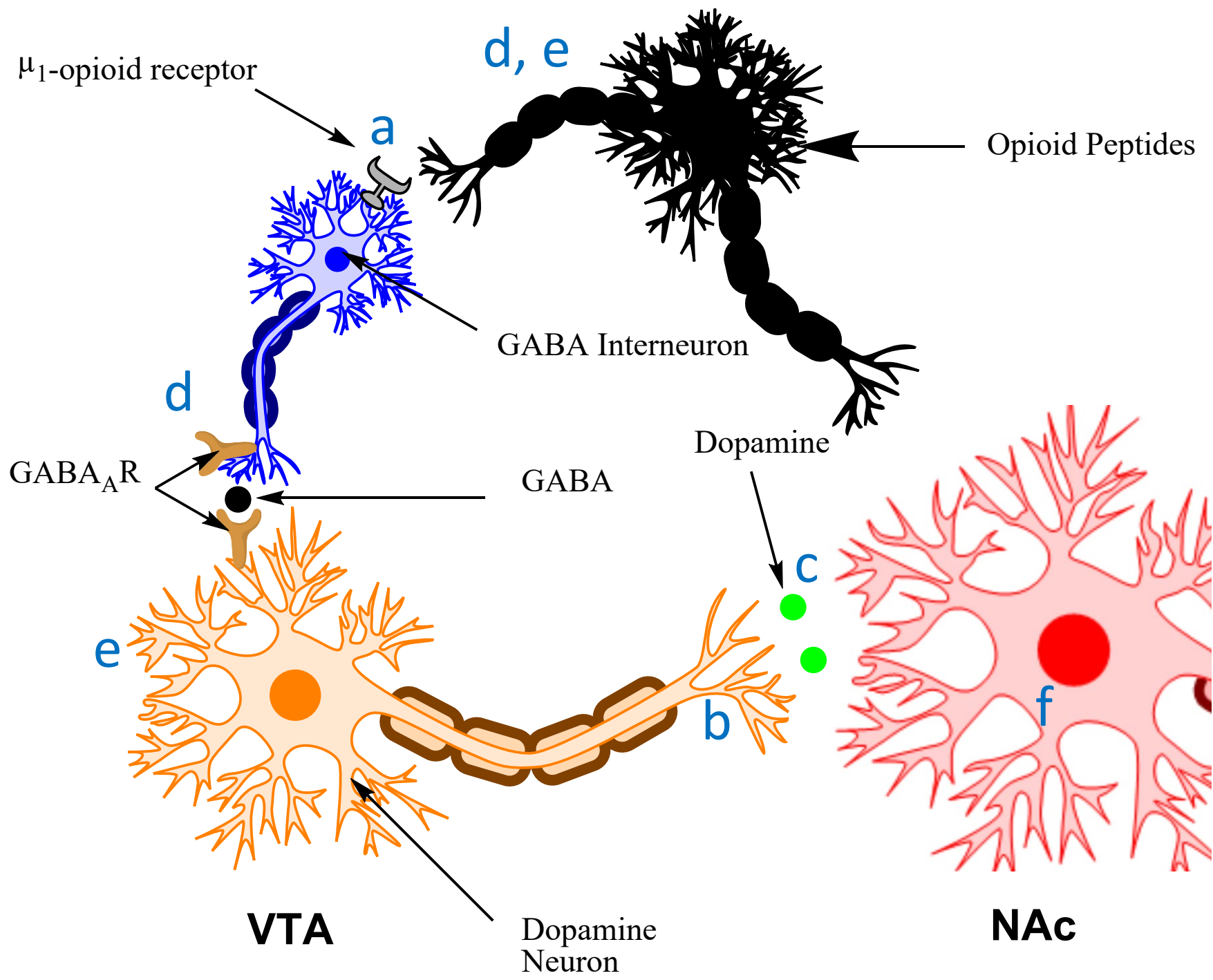


Pharmacological Targets of Drugs of Abuse

Drugs	Target
a. Opioids	Agonists at mainly the μ_1 receptor
b. Cocaine	Increase DA levels by blocking DAT
c. Amphetamine	Stimulate DA release
d. Ethanol	Facilitates GABA _A receptor function
e. Nicotine	Agonist at NACHR
f. Cannabinoids	Agonists at CB ₁ receptors
g. Hallucinogens	Agonists at 5-HT _{2A} receptors

Common Cellular & Molecular Adaptation

Translational Changes



Epigenetic Changes

1. Increases in histone deacetylase in the NAc & VTA
2. Decreases in histone methyltransferase in the NAc

Genetic Changes

1. D2 receptors (D2R)
2. D4 receptors (D4R)
3. Catechol-O-methyltransferase (COMT)

Pharmacogenomic Application

Structure of DNA

- DNA is composed of 4 nucleotide bases
- Matched A-T and C-G

5'-CATGTACCTGGGCCG-3'
3'-GTACATGGACCCGGG-5'

Definitions

Molecular Level

- Pharmacogenomics: The study of variations of DNA and RNA characteristics as related to drug response.¹
- Pharmacogenetics: The study of variations in DNA sequence as related to drug response.¹

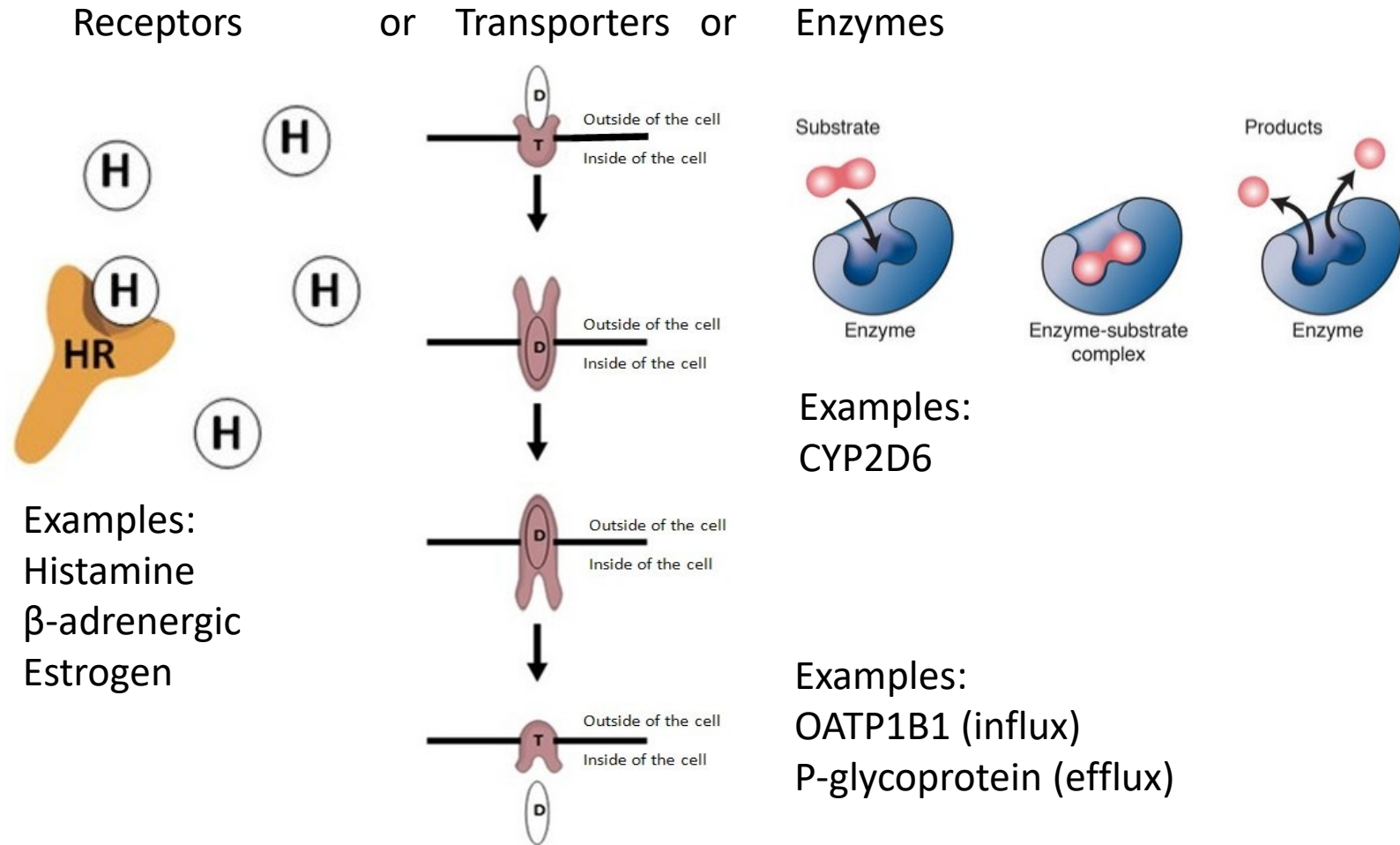
Clinical Level

- Pharmacogenomics: The study of many genes, in some cases the entire genome, involved in response to a drug.²
- Pharmacogenetics: The study of a gene involved in response to a drug.²

¹E15 Definitions for Genomic Biomarkers, Pharmacogenomics, Pharmacogenetics, Genomic Data and Sample Coding Categories. Available at www.fda.gov/downloads/drugs/guidancecomplianceregulatoryinformation/guidances/ucm073162.pdf. Accessed November 4, 2016.

² Kisor DF, Kane MD, Talbot JN, Bright DR, Sprague JE. Pharmacogenes: Scientific Background and Clinical Applications. 2017. Reproduced with permission.

DNA to Protein Drug Targets: Pharmacogenes



Structure of Genes

- A segment of DNA containing all the information needed to encode for one protein is called a gene.

Single Nucleotide Polymorphisms (SNPs)

Influence expression:

5'-CATGTACCTTGGGCCCG-3'

3'-GTACATGGACCCGGGC-5'

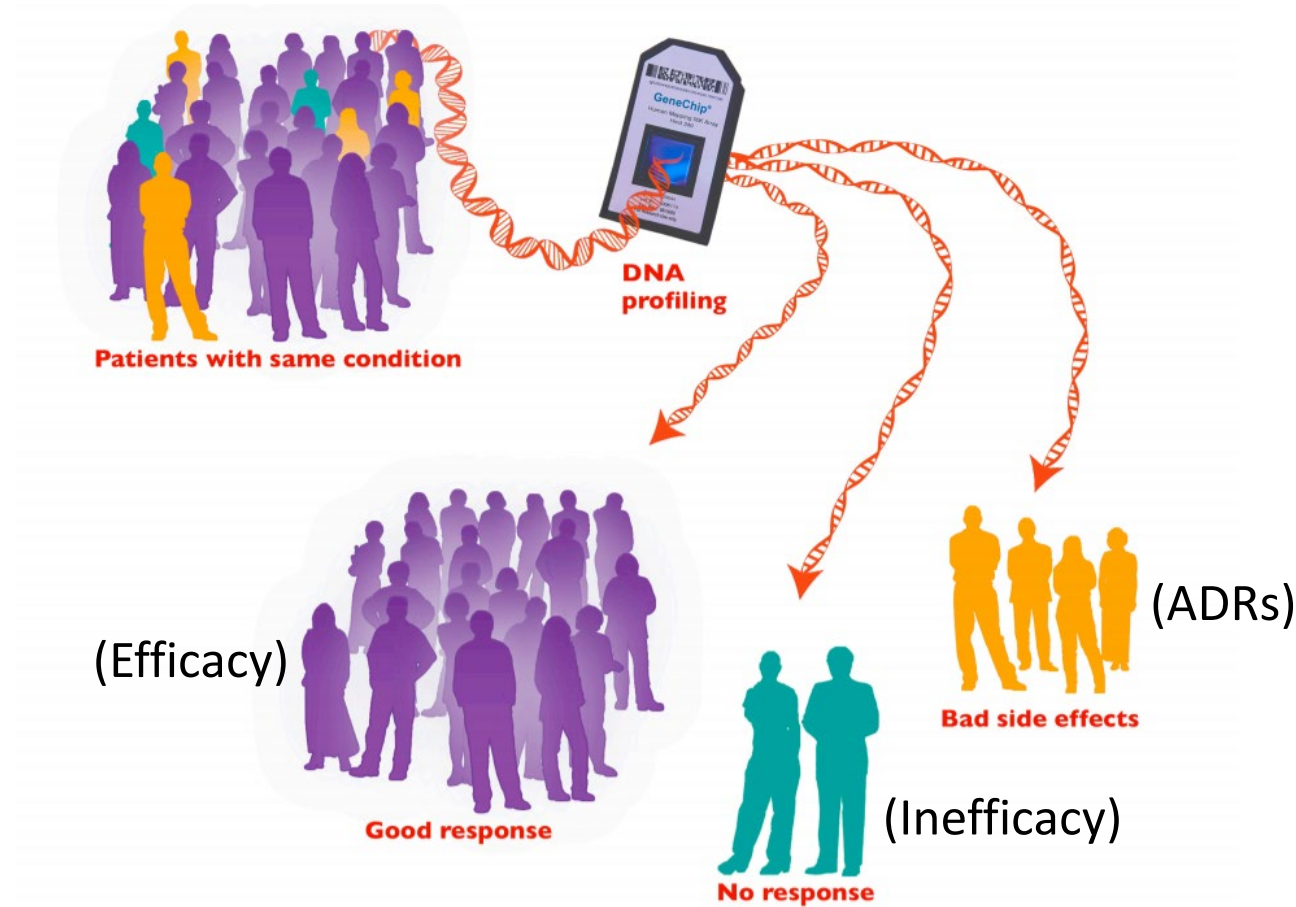
5'-CATGTACCCTGGGCCCG-3'

3'-GTACATGGGCCCCGGGC-5'

Coding Polymorphisms Classifications

- Non-synonymous (missense)
 - ACG codes for threonine
 - CCG codes for proline
- Synonymous (sense)
 - ACG codes for threonine
 - ACA also codes for threonine
- Nonsense

PGx: Drug Efficacy • Adverse Drug Events

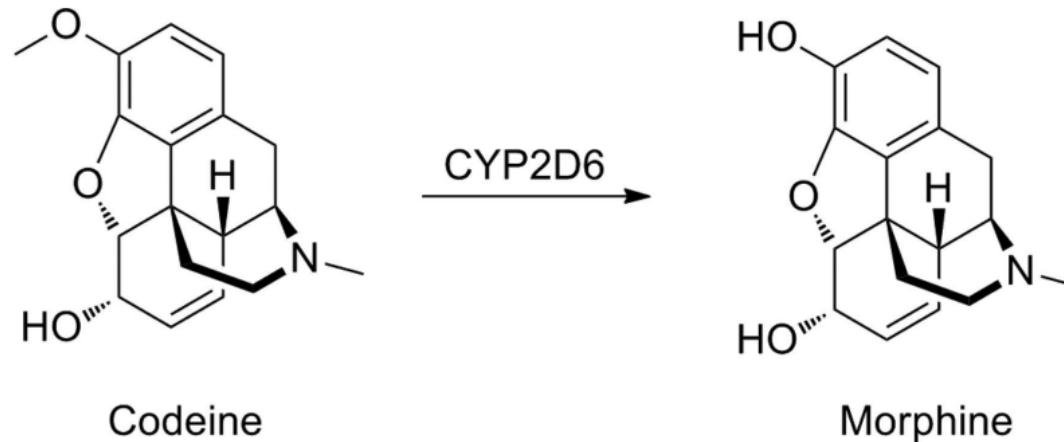


Adverse Drug Events: Example



Rani Jamieson (**Ultra Rapid Metabolizer**)

- Son Tariq was born April 18, 2005;
 - Episiotomy:
 - Received acetaminophen with codeine;
- 12 days later Tariq died.



Adverse Drug Events: Example

- Cause: morphine overdose
- Tariq not receiving morphine
 - Brain/nervous system depression
 - Slow breathing
 - Inactivity/inaction
 - Skin color
 - Poor feeding/failure to thrive

Gene Form	Drug (Std. Dose)	Response	Outcome
<i>CYP2D6</i> *1/*2xN	Codeine	Morphine overdose	Adverse Drug Reaction - Death

CPIC: *CYP2D6*-Codeine

Table 1 Assignment of likely codeine metabolism phenotypes based on cytochrome P450 2D6 (*CYP2D6*) diplotypes

Likely phenotype ^a	Activity score	Genotypes	Examples of diplotypes
Ultrarapid metabolizer (~1–2% of patients)	>2.0	An individual carrying more than two copies of functional alleles	*1/*1xN, *1/*2xN
Extensive metabolizer (~77–92% of patients)	1.0–2.0 ^b	An individual carrying two alleles encoding full or reduced function; or one full-function allele together with either one nonfunctional or one reduced-function allele	*1/*1, *1/*2, *2/*2, *1/*41, *1/*4, *2/*5, *1/*10
Intermediate metabolizer (~2–11% of patients)	0.5 ^b	An individual carrying one reduced-function and one nonfunctional allele	*4/*10, *5/*41
Poor metabolizer (~5–10% of patients)	0	An individual carrying no functional alleles	*4/*4, *4/*5, *5/*5, *4/*6

Table 2 Codeine therapy recommendations based on cytochrome P450 2D6 (*CYP2D6*) phenotype

Phenotype	Implications for codeine metabolism	Recommendations for codeine therapy	Classification of recommendation for codeine therapy ^a	Considerations for alternative opioids
Ultrarapid metabolizer	Increased formation of morphine following codeine administration, leading to higher risk of toxicity	Avoid codeine use due to potential for toxicity.	Strong	Alternatives that are not affected by this <i>CYP2D6</i> phenotype include morphine and nonopioid analgesics. Tramadol and, to a lesser extent, hydrocodone and oxycodone are not good alternatives because their metabolism is affected by <i>CYP2D6</i> activity. ^{b,c}

The Ohio Opioid PGx Study

- Supported by the Ohio Attorney General's Office
- Collaboration with the Emergency Departments at the University of Cincinnati and The Ohio State University
- Sample size: 1300 patients
- PGx screening of 180 genes associated with opioid metabolism and pharmacodynamic response of the reward pathway

The Ohio Opioid PGx Study: Specific Aims

- *Aim 1:* Determine which genes are associated with development of opioid use disorder.
- *Aim 2:* Develop a Genomic Opioid Addiction Risk Score (G-OARs).

Ohio PGx Data Collected

- Genetic samples obtained for a cheek swab
 - Testing for 180 SNPs
- Blood drawn and banked for later testing
- DSM-5 screening for opioid use disorder
- Battery of questions
 - Mental health, prior trauma, prior drug use, environmental influences, risk taking behavior

Planned Analyses

Genetic contribution to opioid use disorder

- Which SNPs are associated with OUD

Machine learning

- Can risk be determined from questions alone?

Highlight prevalence of OUD

- 20% in our participants, despite only 4-5% of all ED visits coded with OUD diagnosis

Report reasons why people participate in research

- Much is known about why people DON'T participate
- Does it differ in the OUD population

Sample Gene Targets

Pharmacokinetic

ABCB1

CYP2D6

CYP2B6

UGT2B7

Pharmacodynamic

TH

COMT

OPRM1

DRD2

General References

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- Walker DM, Cates HM, Heller EA, Nestler EJ. (2015) Regulation of chromatin states by drugs of abuse. *Curr Opin Neurobiol* 112-121.
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