#### Building the Diagnostic Center at UTMB for All of Healthcare

Michael Laposata, M.D., Ph.D. Professor and Chairman Department of Pathology University of Texas Medical Branch Galveston



#### **Disclosures**

#### Director of John Sealy Diagnostic Center at UTMB

#### **Scientific Advisory Board For Werfen**

#### Founding member of Expert Diagnostic Colleague



### **Outline of the Presentation**

- The state of affairs in Diagnostic Medicine in the US
- The recognition of the problem and the assessment of its severity
- The challenges in fixing the problem
- Attempts to provide a solution
- Current activities to create a solution for millions of patients experiencing diagnostic mistakes

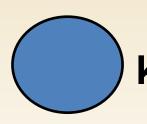


## The State of Affairs in Diagnostic **Medicine in the** US



### How Much Information is There to Know?

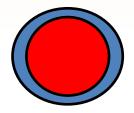




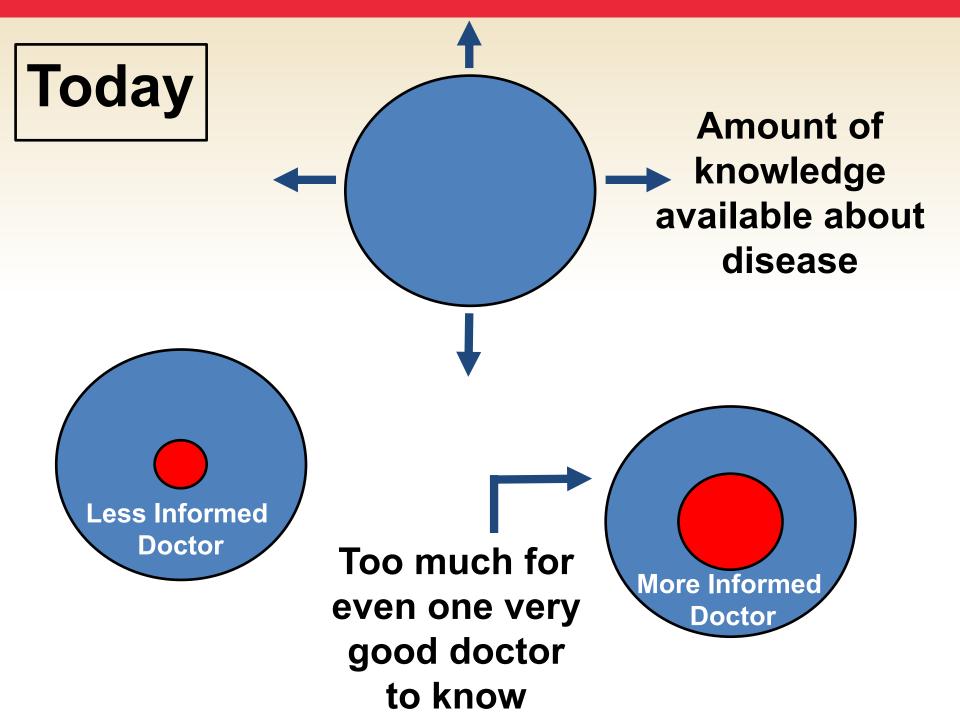
#### Amount of knowledge available about disease



#### Doctor does not know much about diagnosis



Doctor knows much about diagnosis



#### How Can a Clinician Know Enough?

- They are ordering the right nonroutine tests?
- They are making the most up-to-date interpretation of the test results for the patient in their current state?





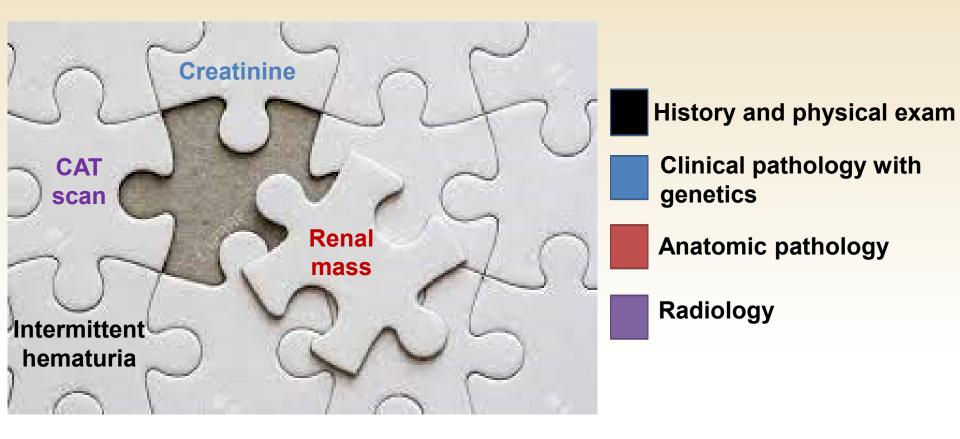
#### The Diagnostic Center Allows for Easy Access to Experts in All Diagnostic Fields

- Up-to-date interpretation
- Patient-specific
- Automated for fast delivery
- Actionable next step recommendations



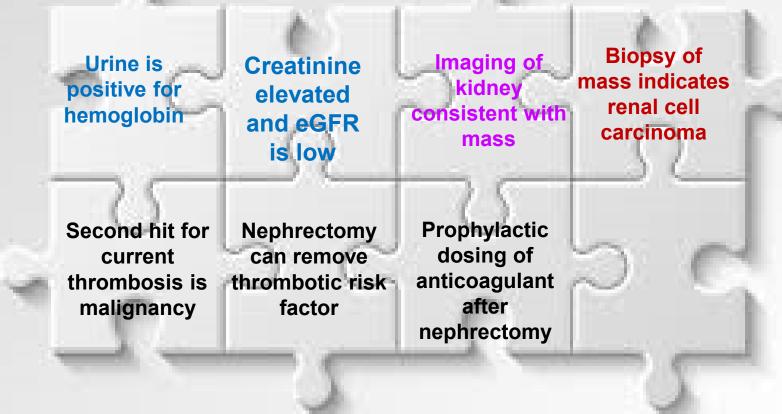


#### Integrative Diagnostics Must Be Implemented -There are Major Barriers to This Challenge



#### Evaluation for Thrombotic Risk to Identify All Contributors to a Thrombotic Event





shutterstock.com · 523973782

## The recognition of diagnostic error and the assessment of its severity



### 2014-2015

# The largest contributor to preventable death from medical error is

## **Diagnostic Error**

as reported by a committee sponsored by the National Academy of Medicine

## Just to show you how little everyone knows about this.....

Why is there no urgency when it is documented by the National Academy of Medicine that there are more than 60,000 preventable deaths in the US annually,

Far more than all the mass shootings combined?



#### Burden of serious harms from diagnostic error in the USA

David E Newman-Toker <sup>(1)</sup>, <sup>1,2</sup> Najlla Nassery, <sup>3</sup> Adam C Schaffer, <sup>4,5</sup> Chihwen Winnie Yu-Moe, <sup>5</sup> Gwendolyn D Clemens, <sup>6</sup> Zheyu Wang, <sup>6,7</sup> Yuxin Zhu, <sup>1,6</sup> Ali S. Saber Tehrani, <sup>1</sup> Mehdi Fanai, <sup>1</sup> Ahmed Hassoon, <sup>1,2</sup> Dana Siegal<sup>8,9</sup>

"Across clinical settings (ambulatory clinics, emergency department and inpatient), we estimate that nearly 800 000 Americans die or are permanently disabled by diagnostic error each year, making it the single largest source of serious harms from medical mistakes."

> BMJ Quality & Safety 2023; 33 82-85 Published Online First: 04 Oct 2023. doi: 10.1136/bmjqs-2023-016496



#### The Concept of Consultation on **Clinical Lab Test Selection and Expert-Driven Result Interpretation** has always been viewed as needed-But Implementation of an effective mechanism to do it has been evasive



The challenges within pathology associated with fixing the problem include: Fear of Obsolescence Fear of Income Loss Fear of Acquisition of Significant **New Clinical Responsibilities** 



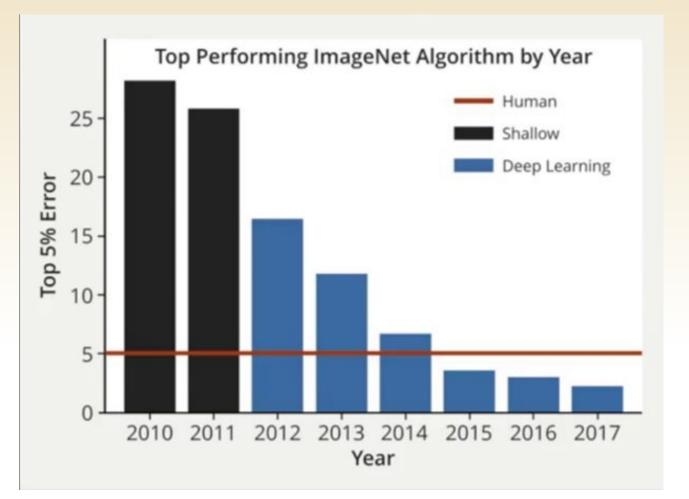
#### The Fear that Artificial Intelligence will Reduce Pathologist Income

#### Developing Quickly in Radiology and Anatomic Pathology



Classification and mutation prediction from non-small cell lung cancer histopathology images using deep learning

#### Diagnostic Algorithms Can Detect Lung Cancer and Classify the Lesions



Deep learning improved the performance of the reading instrument to match and then outperform the pathologist

## An empty cockpit? An instrument deciding if your biopsy is malignant?





Even if there is a computer system to fly an airplane, it will never be without a pilot

The same is true for a pathologist and a biopsy

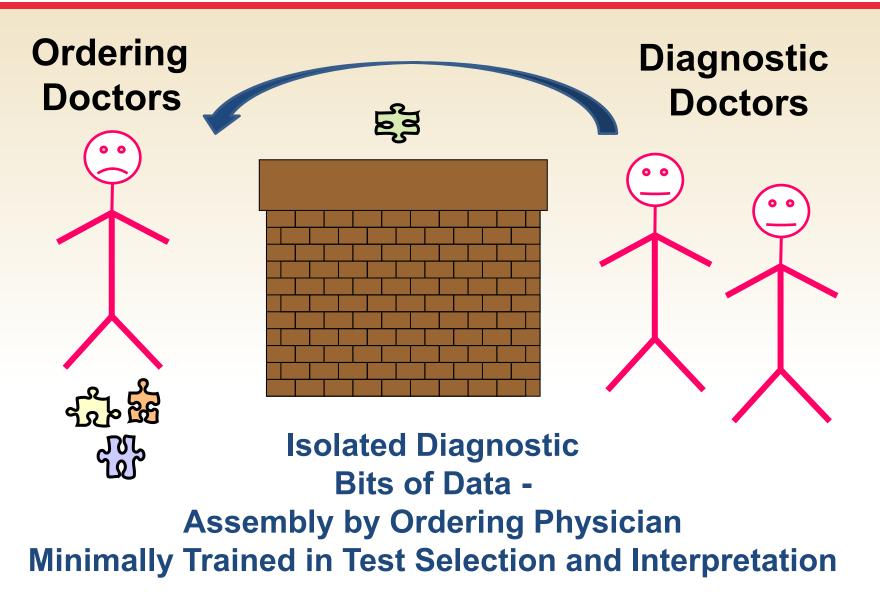
#### THE CONSEQUENCES OF A MISTAKE ARE TOO ENORMOUS



### Expert teams to advise doctors on appropriate test selection and correct result interpretation may reduce the value of the local pathologist



## Passive Laboratory-Virtually every clinical lab in the United States



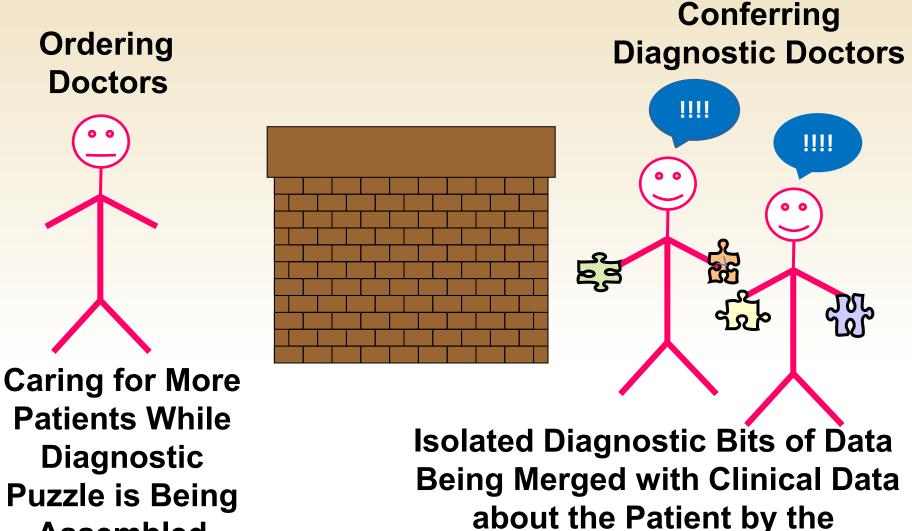
Not choosing pathology – An essay-based survey of first-year clinical residents

The 11 most common reasons why graduates did not choose pathology as a specialty were identified.

The top reason was a perception that pathology lacks practical application to patient care or offers "no real help" (48.2%).



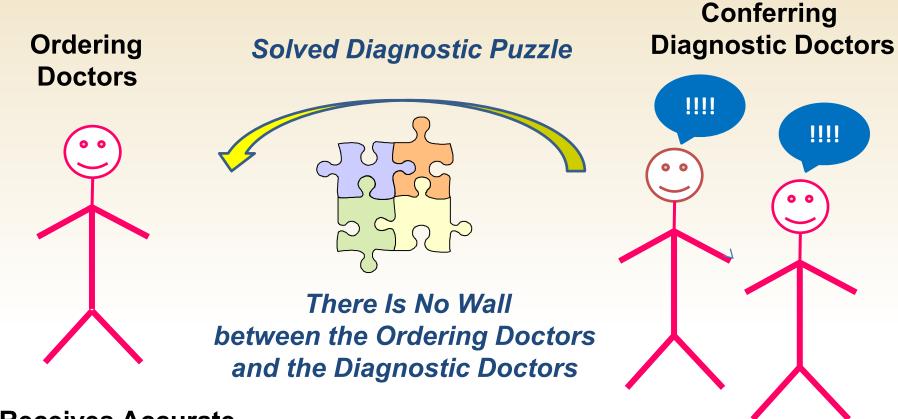
Until recently there was minimal payment for picking up the pieces and establishing a diagnosis



**Diagnostic Doctors** 

Assembled

There was certainly no payment for having a collegial discussion and educating the patient facing physician



Receives Accurate Diagnosis Quickly as a Completed Puzzle Original Idea : DMTs in all major institutions Response : Too much diagnostic medicine to learn with too little financial return

- Diagnostic Center is the source for all diagnostic evaluations
- The Diagnostic Center is paid for its consultation
- If a local pathologist is part of the consultation, diagnostic codes should be able to be used for that activity

utmb Health

### JOHN SEALY DIAGNOSTIC CENTER





### **Diagnostic Center Team**



utmb Health

#### The focus of the Diagnostic Center is almost exclusively on

### **Establishing a Diagnosis**

## with treatment left to the patient facing provider



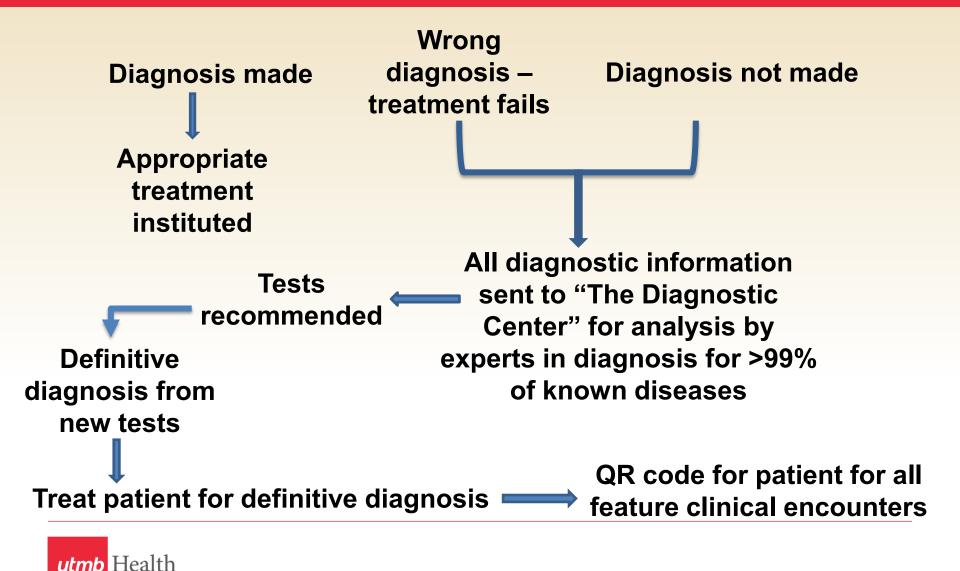
#### There is no plan to compete for laboratory test performance or the patient

## Niche laboratory test performance will be offered as an option

## This is a unique service in the practice of medicine



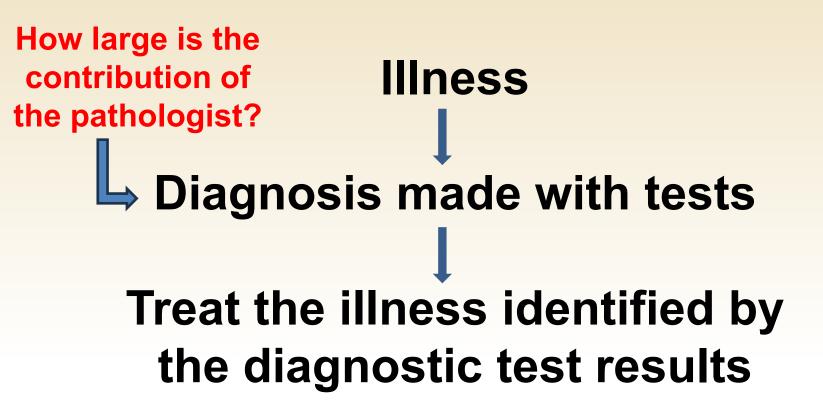
#### How the Diagnostic Center Works



## Attempts to provide a solution



#### The Diagnosis Identifies the Condition and Permits Effective Treatment





## Doing more tests on a small volume of blood does not solve the problem of diagnostic error



#### **Elizabeth Holmes Theranos CEO**





Is performance of battery of diagnostic tests from a fingerstick clinically valuable?

How can someone with only a modicum of engineering experience as a student without the collaboration of clinical chemists or pathologists create a testing device to do >100 tests on a drop of blood

and then convince individuals to invest \$9.2 billion in the company



## The Missing Clinical Service in US Medicine Today :

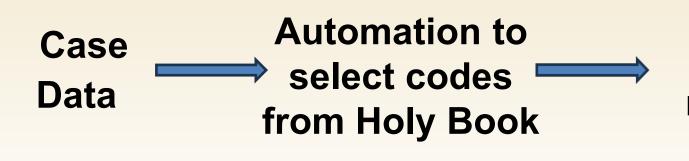
## Leadership in the Diagnostic Evaluation of Patients



## **Current Activities to Create a Solution for** Millions of Patients Experiencing **Diagnostic Mistakes**



### For Each DMT to Completer >100 Cases per Day



Holy Book Interpretive comments reviewed and modified as necessary

## Report sent and case billed



## **Requirements for All DMTs**

Holy Books of hundreds of interpretive comments

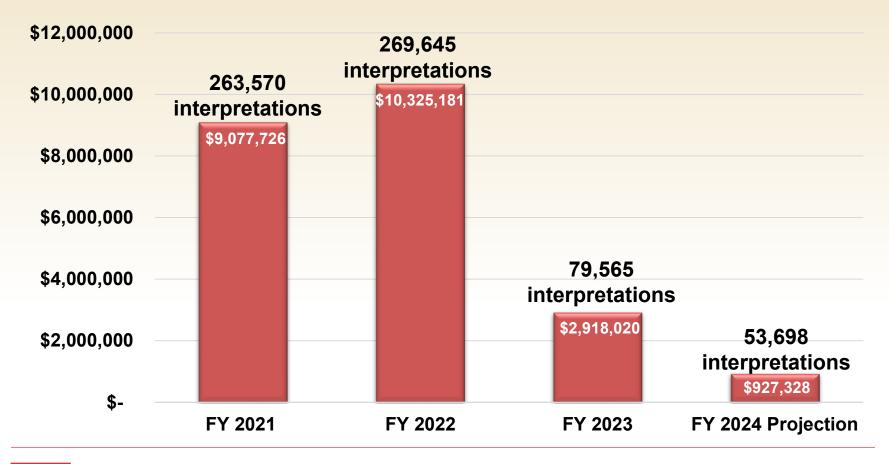
## "Literature Scraping" on a Regular Basis

# Automation from laboratory data to comment selection



#### **COVID Professional Interpretation Revenue**

Total professional revenue of \$22.3M between October 2020 – August 2023 from 612,780 interpretations





This experience taught us:

How to build a Holy Book, and keep it up to date with current findings

How to sign out >1000 cases per day

How to bill for the interpretations

How to automate data extraction from Epic and have the system select an appropriate interpretive comment for acceptance or modification by the pathologist



After 31 meetings with the founding group:

A complete holy book on any topic can be produced in a single day

The automation step which extracts data from the medical record can be done in seconds for any topic – and be linked to references

Scraping the literature now is done with artificial intelligence first reading the published articles and directing findings to the pathologist



## Maintaining Current Content within Holy Book

### "Scraping" the literature

## To obtain the latest findings and incorporate them into the existing ones



Home Insert Draw Page Layo	out Formulas Da	ta Review View Automate Table	💭 Comments 🛛 😥 Share 🗸					
Calibri (Body)	10 A2 A	三三日 砂ィ 池 Wrap Text マ General マ 田子 マ 田子 マ 田子 マ 田子 マ 日本 マ						
Calibri (Body)	✓ 12 ✓ Aˆ A	r ≡ ≡ ≫ v ?2 Wrap Text v General v 🖬 v 🗗 v 👘 v						
Paste Sormat B I U v	⊞ •   <u>∞</u> • <u>A</u> •	E = = → → → Merge & Center → ↓ → % → 60 → 00 Conditional Format Cell Insert	Delete Format Sort & Find & Sensitivity Analyze Data					
A200 $\ddagger$ $\times$ $\checkmark$ $f_{\rm X}$ 10.1001/jar	madermatol.2020.0352							
DOI	PMID v arXiv ID v	Title	▼ Abstract					
78 10.1684/ejd.2020.3880	33021473	Clinical and molecular epidemiology of erythropoietic protoporphyria in Italy	BACKGROUND Erythropoietic protoporphyria (EPP) is a rare inherited disease associated with heme metabolism, charac					
179 10.21037/tgh-2019-rld-05		The acute hepatic porphyrias.	The acute hepatic porphyrias (AHP) are a group of four inherited diseases of heme biosynthesis. They present with simil					
180 10.1101/2020.02.14.949297		Delivery of oligonucleotides to bone marrow to modulate ferrochelatase splicing in a mouse model of Erythropoietic Protoporphyria	Erythropoietic protoporphyria (EPP) is a rare genetic disease in which patients experience acute phototoxic reactions aft					
181 10.1038/s41436-019-0584-0		Targeted resequencing of FECH locus reveals that a novel deep intronic pathogenic variant and eQTLs may cause enythropoietic protoporphyria (EPP)						
182		The Involvement of Anti-Oxidative Response and Mitochondrial Dynamics in the Pathogenesis of Friedreich Aos Ataxia: Relevance to the Development						
183 10.1101/2020.06.14.150904		Human library of cardiac promoters and enhancers	Genome regulatory elements play a critical role during cardiac development and maintenance of normal physiological h					
184 10.5582/irdr.2020.03082 185 10.1080/23808993.2021.1838275		Recent advances in the epidemiology and genetics of acute intermittent porphyria.	Acute intermittent porphyria (AIP) is a dominant inherited disorder with a low penetrance that is caused by mutations in					
185 10.1080/23808993.2021.1838275 186 10.1038/s41467-020-16586-x		Givosiran, a novel treatment for acute hepatic porphyrias Human aminolevulinate synthase structure reveals a eukaryotic-specific autoinhibitory loop regulating substrate binding and product release.	Acute hepatic porphyrias (AHPs) are a group of rare genetic disorders that affect the enzymes of the heme biosynthetic 5,Äs-aminolevulinate synthase (ALAS) catalyzes the first step in heme biosynthesis, generating 5,Äs-aminolevulinate fr					
185 10.1038/s41467-020-16586-x 187 10.1007/s11910-020-01078-8		Human aminolevulinate synthase structure reveals a eukaryotic-specific autoinnibitory loop regulating substrate binding and product release. Porphyric Neuropathy: Pathophysiology, Diagnosis, and Updated Management.	5,AS-aminolevulnate synthase (ALAS) catalyzes the first step in neme biosynthesis, generating 5,AS-aminolevulnate fr PURPOSE OF REVIEW To review the peripheral neurological complications of the acute hepatic porphyrias, as well as the					
188 10.1002/hep4.1503		Porpriet veuropauny: Pathophysiology, Diagnosis, and Opdated Management. Drug-Induced Liver Injury in GI Practice.	Although drug-induced liver injury (DILI) is a rare clinical event, it carries significant morbidity and mortality, leaving it a					
189 10.1016/b978-0-12-819132-3.00018-x		Natural and pharmacological chaperones against accelerated protein degradation: uroporphyrinogen III synthase and congenital erythropoietic porphy						
190 10.1111/bph.15040		Disease pharmacological chaperones against accelerated protein degradation, dioporphymiogen in synchase and ongenital erythoporetic porph Disease pharmacolonetic-pharmacodynamic modelling in acute intermittent porphyria to support the development of mRNA-based therapies.	BACKGROUND AND PURPOSE: Acute intermittent porphysia (AIP) results from haploinsufficiency of the porphobilinogen					
191 10.3324/haematol.2019.232124		Inon metabolism and iron disorders revisited in the hepcidine rra	Iron is biologically essential, but also potentially toxic; as such it is tightly controlled at cell and systemic levels to prevent					
192 10.1136/jclinpath-2020-206647	32605921	Novel frameshift variant (c.409dupG) in SLC25A38 is a common cause of congenital sideroblastic anaemia in the Indian subcontinent.	Aims Congenital sideroblastic anaemias (CSAs) are a group of rare disorders with the presence of ring sideroblasts in the					
193 10.1111/phpp.12501		Acquired erythropoletic protoporphyria: A systematic review of the literature	BACKGROUND: Erythropoietic protoporphyria (EPP) is a semi-dominantly inherited porphyria presenting with photosensi					
94 10.1111/liv.14271		Diagnostic and prognostic assessment of suspected drug-induced liver injury in clinical practice	Idiosyncratic drug-induced liver injury (DILI) is a challenging liver disorder because it can present with a range of phenot					
95 10.1016/b978-0-444-64293-6.00002-6		In silico prediction of drug-induced liver injury: Quo vadis?	Abstract Drug-induced liver injury (DILI) with high incidence and prevalence rates is a potentially severe adverse drug i					
96 10.1007/s40291-019-00438-6		Leading RNA Interference Therapeutics Part 2: Silencing Delta-Aminolevulinic Acid Synthase 1, with a Focus on Givosiran	In November 2019 givosiran became the second small interfering RNA (siRNA)-based drug to receive US Food and Drug					
97 10.1080/13696998.2020.1835306		Cost savings with hemin versus givosiran for the treatment of patients with acute intermittent porphyria (AIP)	BACKGROUND & AIMS Since 1983, hemin has been FDA-approved for acute intermittent porphyria (AIP) attacks. In 201					
98 10.4168/aair.2020.12.3.430		Evaluation of Drug-Induced Liver Injury Developed During Hospitalization Using Electronic Health Record (EHR)-Based Algorithm	PURPOSE: The incidence of drug-induced liver injury (DILI) has been increasing; however, few algorithms are available to					
199 10.1016/j.ymthe.2019.11.010		A Pharmacological Chaperone Therapy for Acute Intermittent Porphyria.	Mutations in hydroxymethylbilane synthase (HMBS) cause acute intermittent porphyria (AIP), an autosomal dominant di					
200 10.1001/jamadermatol.2020.0352	32186677	Association of Afamelanotide With Improved Outcomes in Patients With Erythropoietic Protoporphyria in Clinical Practice.	Importance The effectiveness of afamelanotide treatment in patients with erythropoletic protoporphyria (EPP) in clinica					
201 10.1007/978-3-319-90761-1_53-1		Drug-Induced Liver Injury in Older Adults	The estimated incidence of idiosyncratic drug-induced liver injury (DIU) in the general population is 13.9 to 19.1 per 100,					
02 10.1016/j.cld.2019.09.006	31753253	Drug-Induced Liver Injury in the Setting of Chronic Liver Disease.	Drug-induced liver injury (DILI) is an uncommon but significant cause of liver injury and need for liver transplant. DILI in t					
203 10.1016/j.cld.2019.08.002	31753242	Epidemiology, Predisposing Factors, and Outcomes of Drug-Induced Liver Injury.	Idiosyncratic drug-induced liver injury (DILI) is an underreported and underestimated adverse drug reaction. A recent pop					
204 10.1016/j.ymgme.2020.02.003	32067921	Penetrance and predictive value of genetic screening in acute porphyria.	Abstract Objective Penetrance, predictive value and female patients' perspectives on genetic testing were evaluated a					
205 10.1016/s2468-1253(20)30006-6	32818465	Drug-induced liver injury in older people.	Summary Drug-induced liver injury (DILI) is a rare, unpredictable, and potentially serious adverse reaction. It is induced					
206 10.1056/nejmoa1807838	30726693	Phase 1 Trial of an RNA Interference Therapy for Acute Intermittent Porphyria	Abstract Background Induction of delta aminolevulinic acid synthase 1 (ALAS1) gene expression and accumulation of neu					
207 10.1002/hep.30936	31512765	EXPLORE: A Prospective, Multinational, Natural History Study of Patients with Acute Hepatic Porphyria with Recurrent Attacks	textabstractBackground and Aims: Acute hepatic porphyria comprises a group of rare genetic diseases caused by muta					
208 10.1016/j.ymgme.2018.11.012	30594473	Recent advances on porphyria genetics: Inheritance, penetrance & molecular heterogeneity, including new modifying/causative genes.	Abstract The inborn errors of heme biosynthesis, the Porphyrias, include eight major disorders resulting from loss-of-					
209 10.1111/dth.13014		A first report of porphyria cutanea tarda successfully treated with glycyrrhizin.	Porphyria cutanea tarda (PCT) is a condition that affects liver and skin by reduction of hepatic uroporphyrinogen decar					
210 10.3324/haematol.2018.214320		Dimeric ferrochelatase bridges ABCB7 and ABCB10 homodimers in an architecturally defined molecular complex required for heme biosynthesis	Loss-of-function mutations in the ATP-binding cassette (ABC) transporter of the inner mitochondrial membrane, ABCE					
		Congenital erythropoietic porphyria: Recent advances.	Abstract Congenital erythropoietic porphyria (CEP) is a rare autosomal recessive disorder characterized by photosensiti					
	30391163	Strong correlation of ferrochelatase enzymatic activity with Mitoferrin-1 mRNA in lymphoblasts of patients with protoporphyria.						
12 10.1016/j.ymgme.2018.10.005			Abstract Accumulation of protoporphyrin IX (PPIX) and Zn-PPIX, are the clinical hallmarks of protoporphyria. Phenotypic					
211 10.1016/j.ymgme.2018.12.008 212 10.1016/j.ymgme.2018.10.005 213 10.1016/j.ymgme.2019.01.020		Erythropoletic Protoporphyria and X-Linked Protoporphyria: pathophysiology, genetics, clinical manifestations, and management	Abstract Erythropoletic Protoporphyria (EPP) and X-linked Protoporphyria (XLP) are rare, genetic photodermatoses result					
112 10.1016/j.ymgme.2018.10.005 113 10.1016/j.ymgme.2019.01.020 114 10.1016/j.ymgme.2019.04.013	31076252	Erythropoietic Protoporphyria and X-Linked Protoporphyria: pathophysiology, genetics, clinical manifestations, and management Delta-aminolevulinic acid synthase 2 expression in combination with iron as modifiers of disease severity in erythropoietic protoporphyria.	Abstract Erythropoletic Protoporphyria (EPP) and X-linked Protoporphyria (XLP) are rare, genetic photodermatoses resul Abstract Deficiency in ferrochelatase (FECH), the last enzyme in the heme biosynthetic pathway, leads to an accumula					
112       10.1016/j.ymgme.2018.10.005         113       10.1016/j.ymgme.2019.01.020         114       10.1016/j.ymgme.2019.04.013         115       10.1016/j.ymgme.2018.08.015	31076252 30454868	Erythropoietic Protoporphyria and X-Linked Protoporphyria: pathophysiology, genetics, clinical manifestations, and management Delta-aminolevulinic acid synthase 2 expression in combination with iron as modifiers of disease severity in erythropoietic protoporphyria. Congenital erythropoietic porphyria and erythropoietic protoporphyria: Identification of 7 uroporphyrinogen III synthase and 20 ferrochelatase novel m	Abstract Erythropoletic Protoporphyria (EPP) and X-linked Protoporphyria (XLP) are rare, genetic photodermatoses resu Abstract Deficiency in ferrochelatase (FECH), the last enzyme in the heme biosynthetic pathway, leads to an accumula nu Abstract The erythropoletic porphyrias are inborn errors of heme biosynthesis with prominent cutaneous manifestation					
112       10.1016/j.ymgme.2018.10.005         113       10.1016/j.ymgme.2019.01.020         114       10.1016/j.ymgme.2019.04.013         115       10.1016/j.ymgme.2018.08.015         116       10.1093/nar/gky955	31076252 30454868 30357393	Erythropoietic Protoporphyria and X-Linked Protoporphyria: pathophysiology, genetics, clinical manifestations, and management Delta-aminolevulinic acid synthase 2 expression in combination with iron as modifiers of disease severity in erythropoietic protoporphyria. Congenital erythropoietic porphyria and erythropoietic protoporphyria: Identification of 7 uroporphyrinogen III synthase and 20 ferrochelatase novel m GENCODE reference annotation for the human and mouse genomes.	Abstract Erythropoletic Protoporphyria (EPP) and X-linked Protoporphyria (XLP) are rare, genetic photodermatoses resul Abstract Deficiency in ferrochelatase (FECH), the last enzyme in the heme biosynthetic pathway, leads to an accumula nu Abstract The enythropoletic porphyrias are inborn errors of heme biosynthesis with prominent cutaneous manifestation The accurate identification and description of the genes in the human and mouse genomes is a fundamental requireme					
112 10.1016/j.ymgme.2018.10.005 113 10.1016/j.ymgme.2019.01.020 14 10.1016/j.ymgme.2019.04.013 115 10.1016/j.ymgme.2018.08.015 16 10.1039/nar/gky955 17 10.1182/blood-2018-08.815951	31076252 30454868 30357393 30401706	Erythropoietic Protoporphyria and X-Linked Protoporphyria: pathophysiology, genetics, clinical manifestations, and management Delta-aminolevulinic acid synthase 2 expression in combination with iron as modifiers of disease severity in erythropoietic protoporphyria. Congenital erythropoietic porphyria and erythropoietic protoporphyria: Identification of 7 uroporphyrinogen III synthase and 20 ferrochelatase novel m GENCODE reference annotation for the human and mouse genomes. The molecular genetics of sideroblastic anemia	Abstract Erythropoletic Protoporphyria (EPP) and X-linked Protoporphyria (XLP) are rare, genetic photodermatoses resul Abstract Deficiency in ferrochelatase (FECH), the last enzyme in the heme biosynthetic pathway, leads to an accumula nu Abstract. The erythropoletic porphyrias are inborn errors of heme biosynthesis with prominent cutaneous manifestation. The accurate identification and description of the genes in the human and mouse genomes is a fundamental requireme The sideroblastic anemias (SAs) are a group of inherited and acquired bone marrow diorders defined by pathological in					
112       10.1016/j.ymgme.2018.10.005         13       10.1016/j.ymgme.2019.01.020         14       10.016/j.ymgme.2019.04.013         15       10.1016/j.ymgme.2019.04.013         16       10.0103/nar/gky655         17       10.1182/b/ood-2018-08-815951         18       10.003/nar/gky120	31076252 30454868 30357393 30401706	Erythropoietic Protoporphyria and X-Linked Protoporphyria: pathophysiology, genetics, clinical manifestations, and management Delta-aminolevulinic acid synthase 2 expression in combination with iron as modifiers of disease severity in erythropoietic protoporphyria. Congenital erythropoietic porphyria and erythropoietic protoporphyria: Identification of 7 uroporphyrinogen III synthase and 20 ferrochelatase novel m GENCODE reference annotation for the human and mouse genomes. The molecular genetics of sideroblastic anemia	Abstract Erythropoletic Protoporphyria (EPP) and X-linked Protoporphyria (XLP) are rare, genetic photodermatoses resul Abstract Deficiency in ferrochelatase (FECH), the last enzyme in the heme biosynthetic pathway, leads to an accumula u.bstract. The erythropoletic porphyrias are inborn errors of heme biosynthesis with prominent cutaneous manifestation The accurate identification and description of the genes in the human and mouse genomes is a fundamental requireme The sideroblastic anemias (SAs) are a group of inherited and acquired bone marrow disorders defined by pathological in The GWAS Catalog delivers a high-quality curated collection of all published genome-wide association studies enabling the GWAS Catalog delivers a high-quality curated collection of all published genome-wide association studies enabling the GWAS Catalog delivers a high-quality curated collection of all published genome-wide association studies enabling the GWAS Catalog delivers a high-quality curated collection of all published genome-wide association studies enabling the GWAS Catalog delivers a high-quality curated collection of all published genome-wide association studies enabling the GWAS Catalog delivers a high-quality curated collection of all published genome-wide association studies enabling the GWAS Catalog delivers a high-quality curated collection of all published genome-wide association studies enabling the total studies enabling the					
112       10.1016/j.ymgme.2018.10.005         13       10.1016/j.ymgme.2019.01.020         14       10.1016/j.ymgme.2019.04.013         15       10.1016/j.ymgme.2018.08.015         16       10.1093/nar/gky955         17       10.1182/blood-2018-08-815951         18       10.1093/nar/gky120         19       10.3390/genes10010043	31076252 30454868 30357393 30401706 30445434	Erythropoietic Protoporphyria and X-Linked Protoporphyria: pathophysiology, genetics, clinical manifestations, and management Delta-aminolevulinic acid synthase 2 expression in combination with iron as modifiers of disease severity in erythropoietic protoporphyria. Congenital erythropoietic portophyria and erythropoietic protoporphyria: Identification of 7 uroporphyrinogen III synthase and 20 ferrochelatase novel m GENCODE reference annotation for the human and mouse genomes. The molecular genetics of sideroblastic anemia The NHGRI-EBI	Abstract Erythropoletic Protoporphyria (EPP) and X-linked Protoporphyria (XLP) are rare, genetic photodermatoses resu Abstract Deficiency in ferrochelatase (FECH), the last enzyme in the heme biosynthetic pathway, leads to an accumula nu Abstract The enythropoletic porphyrias are inborn errors of heme biosynthesis with prominent cutaneous manifestatio The accurate identification and description of the genes in the human and mouse genomes is a fundamental requireme The sideroblastic anemias (SAs) are a group of inherited and acquired bone marrow disorders defined by pathological in The GWAS Catalog delivers a high-quality curated collection of all published genome-wide association studies enabling Enhancers are non-coding DNA elements that function in cis to regulate transcription from nearby genes. Through direct					
112       10.1016/j.ymgme.2018.10.005         13       10.1016/j.ymgme.2019.04.013         14       10.016/j.ymgme.2019.04.013         15       10.1016/j.ymgme.2019.04.013         16       10.0103/nar/gky655         17       10.1182/b004-2018.06.815551         18       10.1003/nar/gky120         19       10.330/gene10010043         20       10.016/j.ymgme.2019.01.004	31076252 30454868 30357393 30401706 30445434 30683557	Erythropoietic Protoporphyria and X-Linked Protoporphyria: pathophysiology, genetics, clinical manifestations, and management Delta-aminolevulinic acid synthase 2 expression in combination with iron as modifiers of disease severity in erythropoietic protoporphyria. Congenital erythropoletic porphyria and erythropoietic protoporphyria: Identification of 7 uroporphyrinogen III synthase and 20 ferrochelatase novel m GRNCODE reference annotation for the human and mouse genomes. The molecular genetics of sideroblastic anemia The NHGRI-EBI	Abstract Erythropoletic Protoporphyria (EPP) and X-linked Protoporphyria (XLP) are rare, genetic photodermatoses resu Abstract Deficiency in ferrochelatase (FECH), the last enzyme in the heme biosynthetic pathway, leads to an accumula nu Abstract The erythropoletic porphyrias are inborn errors of heme biosynthesis with prominent cutaneous manifestation The accurate identification and description of the genes in the human and mouse genomes is a fundamental requireme The sideroblastic anemias (SAs) are a group of inherited and acquired bone marrow disorders defined by pathological in The GWAS Catalog delivers a high-quality curated collection of all published genome-wide association studies enabling Enhancers are non-coding DNA elements that function in cis to regulate transcription from nearby genes. Through direc Abstract Porphyria cutanea tarda (PCT) is the most common human porphyria, due to hepatic deficiency of uroporphyri					
112       10.1016/j.ymgme.2018.10.005         13       10.1016/j.ymgme.2019.01.020         14       10.016/j.ymgme.2019.04.013         15       10.1016/j.ymgme.2019.04.013         15       10.1033/nar/gky655         17       10.1182/b004-2018-08-815951         18       10.1033/nar/gky1120         19       10.3390/genes10010043         20       10.1016/j.ymgme.2019.01.004         21       10.106/j.ymgme.2019.01.004	31076252 30454868 30357393 30401706 30445434 30683557 30660387	Erythropoletic Protoporphyria and X-Linked Protoporphyria: pathophysiology, genetics, clinical manifestations, and management Delta-aminolevulinic acid synthase 2 expression in combination with iron as modifiers of disease severity in erythropoletic protoporphyria. Congenital erythropoletic protoporphyria and erythropoletic protoporphyria: Identification of 7 uroporphyringen III synthase and 20 ferrochelatase novel m GENCODE reference annotation for the human and mouse genomes. The molecular genetics of sideroblastic anemia The NIGRI-EBIODE of sideroblastic anemia and Disease Porphyria cutanea tarda: Recent update. GIRXS mutations impair heme biosynthetic enzymes ALA synthase 2 and ferrochelatase in Human congenital sideroblastic anemia	Abstract Erythropoletic Protoporphyria (EPP) and X-linked Protoporphyria (XLP) are rare, genetic photodermatoses resu Abstract Deficiency in ferrochelatase (FECH), the last enzyme in the heme biosynthetic pathway, leads to an accumula u.Abstract The erythropoletic porphyrias are inborn errors of heme biosynthesis with prominent cutaneous manifestatio The accurate identification and description of the genes in the human and mouse genomes is a fundamental requireme The sideroblastic anemias (SAs) are a group of inherited and acquired bone marrow disorders defined by pathological in The GWAS Catalog delivers a high-quality curated collection of all published genome-wide association studies enabling Enhancers are non-coding DNA elements that function in cis to regulate transcription from nearby genes. Through Abstract Porphyria cutanea tarda (PCT) is the most common human porphyria, due to hepatic deficiency of uroporphyri Abstract Non-syndromic microcytic congenital sideroblastic anemia (CAS) is predominantly caused by defective genes					
112       10.1016/j.ymgme.2018.10.005         13       10.1016/j.ymgme.2019.04.013         14       10.1016/j.ymgme.2018.08.015         15       10.1016/j.ymgme.2018.08.015         16       10.1033/nar/gky955         17       10.1182/blood-2018-08.815951         18       10.039/nar/gky9120         19       10.3390/genes10010043         20       10.1016/j.ymgme.2019.01.004         21       10.1016/j.ymgme.2019.01.004	31076252 30454868 30357393 30401706 30445434 30683557 30660387 30733921	Erythropoietic Protoporphyria and X-Linked Protoporphyria: pathophysiology, genetics, clinical manifestations, and management Delta-aminolevulinic acid synthase 2 expression in combination with iron as modifiers of disease severity in erythropoietic protoporphyria. Congenital erythropoietic portoporphyria and erythropoietic portoporphyria: Identification of 7 uroporphyrinogen III synthase and 20 ferrochelatase novel m GENCODE reference annotation for the human and mouse genomes. The molecular genetics of sideroblastic anemia The NHGRIE-BL_S S Catalog of published genome-wide association studies, targeted arrays and summary statistics 2019. Genomic Enhances in Brain Health and Disease Porphyria cutanea tarda: Recent update. GLRXS mutations impair heme biosynthetic enzymes ALA synthase 2 and ferrochelatase in Human congenital sideroblastic anemia Benefits of prophylactic heme therapy in severe acute intermittent porphyria.	Abstract Enthropoletic Protoporphyria (EPP) and X-linked Protoporphyria (XLP) are rare, genetic photodermatoses resu- Abstract Deficiency in ferrochelatase (FECH), the last enzyme in the heme biosynthetic pathway, leads to an accumula u.Abstract The enthropoletic porphyrias are inborn errors of heme biosynthesis with prominent cutaneous manifestatio The accurate identification and description of the genes in the human and mouse genomes is a fundamental requirems the sideroblastic anemias (SAs) are a group of inherited and acquired bone marrow disorders defined by pathological The GWAS Catalog delivers a high-quality curated collection of all published genome-wide association studies enabling Enhancers are non-coding DNA elements that function in cis to regulate transcription from nearby genes. Through direc Abstract Non-syndromic microcytic congenital sideroblastic anemia (cSA) is predominantly caused by defective genes Abstract Acute intermittent porphyria (API), an autosomal dominant inborn error of metabolism, is the most common					
12 10.1016/j.ymgme.2018.10.005 13 10.1016/j.ymgme.2019.01.020 14 10.1016/j.ymgme.2019.04.013 15 10.1016/j.ymgme.2019.04.013 15 10.1033/nar/gky955 17 10.1182/b004-2018-06-815951 18 10.1033/nar/gky1120 19 10.3390/genes10010043 20 10.1016/j.ymgme.2019.01.004 21 10.1016/j.ymgme.2019.01.004 21 10.1016/j.ymgme.2019.01.002 23 10.3390/ph12010017	31076252 30454868 30357393 30401706 30445434 30683557 30660387 30733921 30678075	Erythropoietic Protoporphyria and X-Linked Protoporphyria: pathophysiology, genetics, clinical manifestations, and management Delta-aminolevulinic acid synthase 2 expression in combination with iron as modifiers of disease severity in erythropoietic protoporphyria. Congenital erythropoletic porphyria and erythropoietic protoporphyria: Identification of 7 uroporphyrinogen III synthase and 20 ferrochelatase novel m GENCODE reference annotation for the human and mouse genomes. The molecular genetics of sideroblastic anemia The NHGRI-EBI	Abstract Erythropoletic Protoporphyria (EPP) and X-linked Protoporphyria (XLP) are rare, genetic photodermatoses resu- Abstract Deficiency in ferrochelatase (FECH), the last enzyme in the heme biosynthetic pathway, leads to an accumula nu Abstract The erythropoletic porphyrias are inborn errors of heme biosynthesis with prominent cutaneous manifestatio The accurate identification and description of the genes in the human and mouse genomes is a fundamental requireme The sideroblastic anemias (SAs) are a group of inherited and acquired bone marrow disorders defined by pathological The GWAS Catalog delivers a high-quality curated collection of all published genome-wide association studies enabling Enhancers are non-coding DNA elements that function in cis to regulate transcription from nearby genes. Through direc Abstract Non-syndromic microcytic congenital sideroblastic anemia (SA) is predominantly caused by defective genes Abstract Acute intermittent porphyria (AIP), an autosomal dominant inborn error of metabolism, is the most common Ferritin is a multimeric protein composed of light (L-ferritin) and heavy (H-ferritin) subunits that binds and stores iron in					
121       10.1016/j.ymgme.2018.10.005         131       10.1016/j.ymgme.2019.01.020         141       10.1016/j.ymgme.2019.04.013         125       10.1016/j.ymgme.2019.04.013         125       10.1016/j.ymgme.2019.04.013         126       10.1033/nar/gkyd95         121       10.1182/b004-2018-08-815951         128       10.1033/nar/gkyd120         129       10.1036/j.ymgme.2019.01.004         120       10.1016/j.ymgme.2019.01.002         121       10.1016/j.ymgme.2019.01.002         123       10.3030/gh12010017         124       10.1016/j.ymgme.2019.01.015	31076252 30454868 30357393 30401706 30445434 30683557 30660387 30733921 30678075	Erythropoletic Protoporphyria and X-Linked Protoporphyria: pathophysiology, genetics, clinical manifestations, and management Delta-aminolevulinic acid synthase 2 expression in combination with iron as modifiers of disease severity in erythropoletic protoporphyria. Congenital erythropoletic protoporphyria and erythropoletic protoporphyria: lenntification of 7 uroporphyringen III synthase and 20 ferrochelatase novel m GENCODE reference annotation for the human and mouse genomes. The molecular genetics of sideroblastic anemia The NISRIF.BID_S S Catalog of published genome-wide association studies, targeted arrays and summary statistics 2019. Genomic Enhances in Brain Health and Disease Porphyria cutanea tarda: Recent update. GIXS mutations impair heme biosynthetic enzymes ALA synthase 2 and ferrochelatase in Human congenital sideroblastic anemia Benefits of prophylacit heme therapy in severe acute intermittent porphyria. L-Ferritin: One Gene, Five Diseases; from Hereditary Hyperferritinemia to Hypoferritinemia-Report of New Cases. Regulation and tissue-specific expression of CX-aminolevulinic acid synthases in non-syndromic sideroblastic anemias and porphyrias	Abstract Erythropoletic Protoporphyria (EPP) and X-linked Protoporphyria (XLP) are rare, genetic photodermatoses resul Abstract Deficiency in ferrochelatase (FECH), the last enzyme in the heme biosynthetic pathway, leads to an accumula u.bstract The erythropoletic porphyrias are inborn errors of heme biosynthetic pathway, leads to an accumula The accurate identification and description of the genes in the human and mouse genomes is a fundamental requireme The idenoblastic anemias (SAs) are a group of inherited and acquired born emarwo disorders defined by pathologica. The GWAS Catalog delivers a high-quality curated collection of all published genome-wide association studies enabling Enhancers are non-coding DNA elements that function in cis to regulate transcription from nearby genes. Through direct Abstract Porphyria cutanea tarda (PCT) is the most common human porphyria, due to hepatic deficiency of uroprophyri Abstract. Non-syndromic microcytic congenital sideroblastic anemia (CSA) is predominantly caused by defective genes Abstract. Acute intermittent porphyria (AIP), an autosomal dominant inborn error of metabolism, is the most common Ferritin is a multimeric protein composed of light (L-ferritin) and heavy (H-ferritin) subunits that binds and stores iron in Abstract. Recently, new genes and molecular mechanisms have been identified in patients with porphyrias and siderob					
112       10.1016/j.ymgme.2018.10.005         13       10.1016/j.ymgme.2019.04.013         14       10.1016/j.ymgme.2018.08.015         15       10.1016/j.ymgme.2018.08.015         16       10.1033/nar/gky955         17       10.1182/blood-2018-08.815951         18       10.039/nar/gky9120         10       1182/blood-2018-08.815951         18       10.039/nar/gky120         10       10.1016/j.ymgme.2019.01.004         21       10.0106/j.ymgme.2019.01.002         22       10.1016/j.ymgme.2019.01.002         23       10.3390/gh12010017         24       10.016/j.ymgme.2019.01.015         25       10.1016/j.ajmg.2019.01.015	31076252 30454868 30357393 30401706 30445434 30683557 30660387 30733921 30678075 30712775	Erythropoietic Protoporphyria and X-Linked Protoporphyria: pathophysiology, genetics, clinical manifestations, and management Delta-aminolevulinic acid synthase 2 expression in combination with iron as modifiers of disease severity in erythropoietic protoporphyria. Congenital erythropoietic portoporphyria and erythropoietic protoporphyria: Identification of 7 uroporphyrinogen III synthase and 20 ferrochelatase novel m GENCODE reference annotation for the human and mouse genomes. The molecular genetics of sideroblastic anemia The NHGRI-EBJ_S Catalog of published genome-wide association studies, targeted arrays and summary statistics 2019. Genomic Enhances in Brain Health and Disease Porphyria cutanea tarda: Recent update. GLRXS mutations impair heme biosynthetic enzymes ALA synthase 2 and ferrochelatase in Human congenital sideroblastic anemia Benefits of prophylacic heme therapy in severe acute Intermittent porphyria. L-Ferritin: One Gene, Five Diseases, from Hereditary Hyperferritinemia to Hypoferritinemia-Report of New Cases. Regulation and tissue-specific expression of GE-aminolevulinic acid synthase in non-syndromic sideroblastic anemias and porphyrias Erythroid-Progenitor-Targeted Gene Therapy Using Bifunctional TFR1 Ligand-Peptides in Human Erythropoletic Protoporphyria.	Abstract Erythropoletic Protoporphyria (EPP) and X-linked Protoporphyria (XLP) are rare, genetic photodermatoses resul Abstract Deficiency in ferrochelatase (FECH), the last enzyme in the heme biosynthetic pathway, leads to an accumula Notatact Deficiency in ferrochelatase (FECH), the last enzyme in the heme biosynthetic pathway, leads to an accumula Notatact De enythropoletic porphyria as reinborn errors of heme biosynthesis with prominent cutaneous manifestation The accurate identification and description of the genes in the human and mouse genomes is a fundamental requireme The sideroblastic anemias (SAs) are a group of inherited and acquired bone marrow disorders defined by pathological in The GWAS Catalog delivers a high-quality curated collection of all published genome-wide association studies enabling Enhancers are non-coding DNA elements that function in cis to regulate transcription from nearby genes. Through direct Abstract Non-syndromic microcytic congenital sideroblastic anemia (cSA) is predominantly caused by defective genes Abstract. Acute intermittent porphyria (AIP), an autosomal dominant inborn error of metabolism, is the most common Ferritin is a multimeric protein composed of light (I-ferritin) and heavy (H-ferritin) subunits that binds and stores iron in Abstract. Recently, new genes and molecular mechanisms have been identified in patients with porphyrias and siderob Erythropoletic protoporphyria (EPP) is a hereditary disease characterized by a deficiency is microchelatase (FECH) activity					
212       10.1016/j.ymgme.2018.10.005         213       10.1016/j.ymgme.2019.04.013         214       10.1016/j.ymgme.2019.04.013         215       10.1016/j.ymgme.2019.04.013         216       10.0103/nar/gky955         217       10.1182/biod-2018.06.815551         218       10.1003/nar/gky120         219       10.3390/gene10010043         220       10.1016/j.ymgme.2019.01.004         221       10.1016/j.ymgme.2019.01.002         231       10.330/ghp12010017	31076252 30454868 30357393 30401706 30445434 30683557 30660387 30733921 30678075 30712775	Erythropoletic Protoporphyria and X-Linked Protoporphyria: pathophysiology, genetics, clinical manifestations, and management Delta-aminolevulinic acid synthase 2 expression in combination with iron as modifiers of disease severity in erythropoletic protoporphyria. Congenital erythropoletic protoporphyria and erythropoletic protoporphyria: lenntification of 7 uroporphyringen III synthase and 20 ferrochelatase novel m GENCODE reference annotation for the human and mouse genomes. The molecular genetics of sideroblastic anemia The NISRIF.BID_S S Catalog of published genome-wide association studies, targeted arrays and summary statistics 2019. Genomic Enhances in Brain Health and Disease Porphyria cutanea tarda: Recent update. GIXS mutations impair heme biosynthetic enzymes ALA synthase 2 and ferrochelatase in Human congenital sideroblastic anemia Benefits of prophylacit heme therapy in severe acute intermittent porphyria. L-Ferritin: One Gene, Five Diseases; from Hereditary Hyperferritinemia to Hypoferritinemia-Report of New Cases. Regulation and tissue-specific expression of CX-aminolevulinic acid synthases in non-syndromic sideroblastic anemias and porphyrias	Abstract Erythropoletic Protoporphyria (EPP) and X-linked Protoporphyria (XLP) are rare, genetic photodermatoses resul Abstract Deficiency in ferrochelatase (FECH), the last enzyme in the heme biosynthetic pathway, leads to an accumula u.bstract The erythropoletic porphyrias are inborn errors of heme biosynthetic pathway, leads to an accumula The accurate identification and description of the genes in the human and mouse genomes is a fundamental requireme The idenoblastic anemias (SAs) are a group of inherited and acquired born emarwo disorders defined by pathologica. The GWAS Catalog delivers a high-quality curated collection of all published genome-wide association studies enabling Enhancers are non-coding DNA elements that function in cis to regulate transcription from nearby genes. Through direct Abstract Porphyria cutanea tarda (PCT) is the most common human porphyria, due to hepatic deficiency of uroprhyri Abstract. Non-syndromic microcytic congenital sideroblastic anemia (CSA) is predominantly caused by defective genes Abstract. Acute intermittent porphyria (AIP), an autosomal dominant inborn error of metabolism, is the most common Ferritin is a multimeric protein composed of light (L-ferritin) and heavy (H-ferritin) subunits that binds and stores iron in Abstract. Recently, new genes and molecular mechanisms have been identified in patients with porphyrias and siderob					



## Selected Expert Teams and Benefits to Patient Outcome and Financial Return

## To Be Implemented in the Diagnostic Center



#### Diagnostic Evaluations For People with No Apparent Illness

**Pharmacogenetics** 

**Exome Analysis** 

**Cancer Screening** 



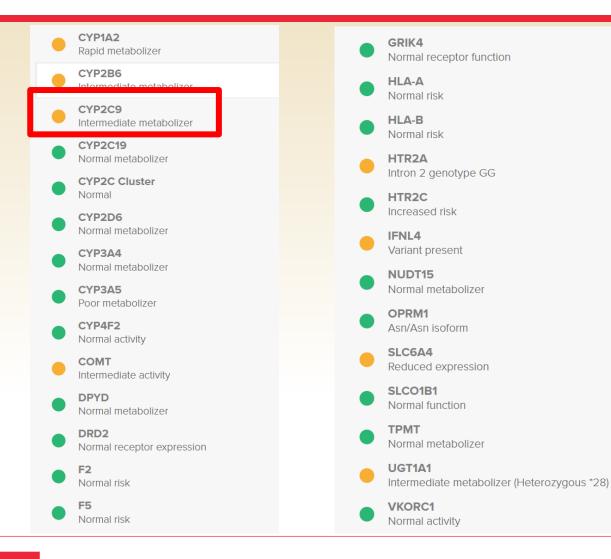
Clinical Benefit from the Pharmacogenomics DMT

### The benefit of taking the most effective drug is obvious, but it requires an initial investment in a lab test to do so

Making an initial investment in people or equipment involves trust in the person recommending the purchase



#### **Genes Relevant to Pharmacogenomics**



Yellow = Altered Metabolism

Green = Normal Metabolism

#### **Focused Report**

Medications	Genes with Variants	Metabolizer Status	<b>Recommendations for Prescriber</b>
Atorvastatin (Lipitor)	SLCO1 B1	Decreased Function	Prescribe ≤ 40 mg as a starting dose and adjust doses of atorvastatin based on disease-specific guidelines. Prescriber should be aware of possible increased risk of myopathy especially for 40 mg dose. If dose > 40 mg is needed for desired efficacy, consider combination therapy (i.e., atorvastatin plus non-statin guideline directed medical therapy.



## Exome Analysis for Single Gene Disorder and Cancer Genetics

 Available upon validation of PacBio

 Can send out testing for special cases at this time



## Exome Analysis for Single Gene Disorder and Cancer Genetics

 If exome analysis shows a high risk for pancreatic cancer, regular abdominal imaging studies to detect an early, surgically curable pancreatic cancer

### **CAN BE LIFESAVING**



## Algorithms to Screen for Different Types of Cancer

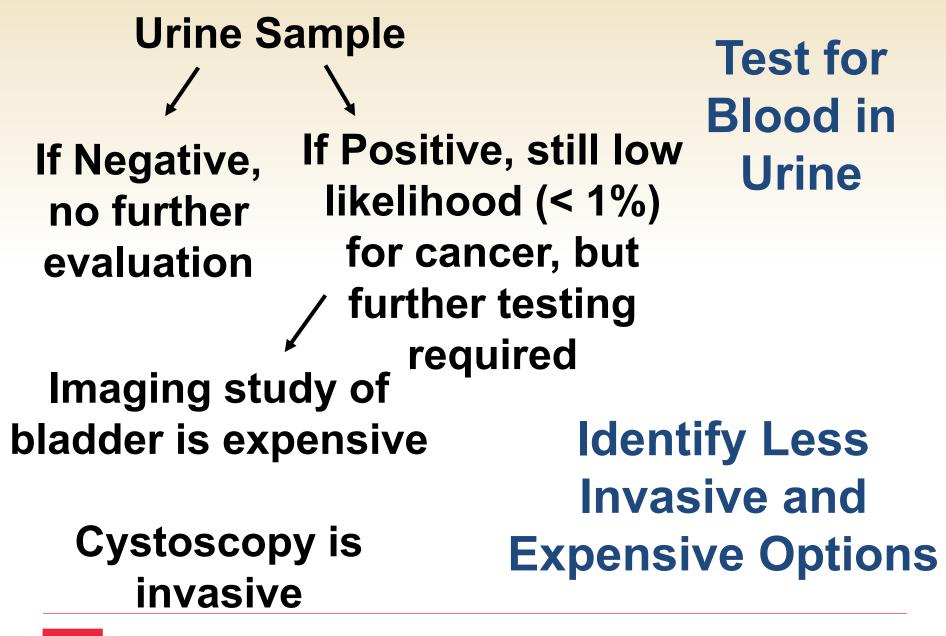
- An algorithm for testing starts with a test that is neither invasive nor expensive
- If the screening test is positive, the challenge is to find a second level test that is only modestly invasive or expensive



## Algorithms to Screen for Different Types of Cancer

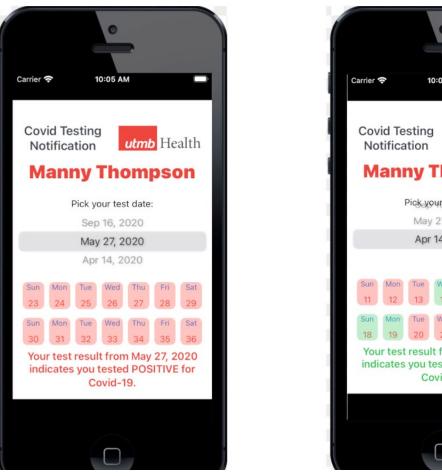
- It is understood that the final confirmatory test for cancer is likely to be invasive, expensive, or both
- The goal is to not take individuals who are suspected of having cancer and move them through an evaluation with invasive or expensive tests which may be harmful and <u>not</u> identify cancer







#### App Sample Concept Similar to COVID-19 Results Notification







#### High Intensity and Low Intensity Screening Will be Available

	Age						
	40	42	44	46	48	50	52
Adrenal Gland Cortex Tumors	X					Χ	
Adrenal Medulla Tumor (pheochromocytoma)		X					X
Bladder Cancer	X	X	X	X	X	X	X



## For the Patient with Cancer

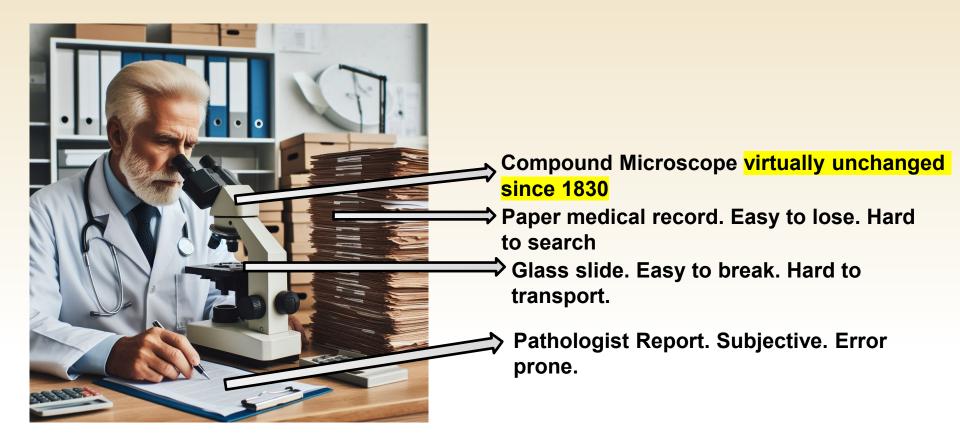
## A Comprehensive High Sensitivity Cancer Diagnosis DMT



## DIGITAL TRANSFORMATION **AND ADOPTION OF** ARTIFICIAL INTELLIGENCE IN ANATOMIC PATHOLOGY AT UTMB

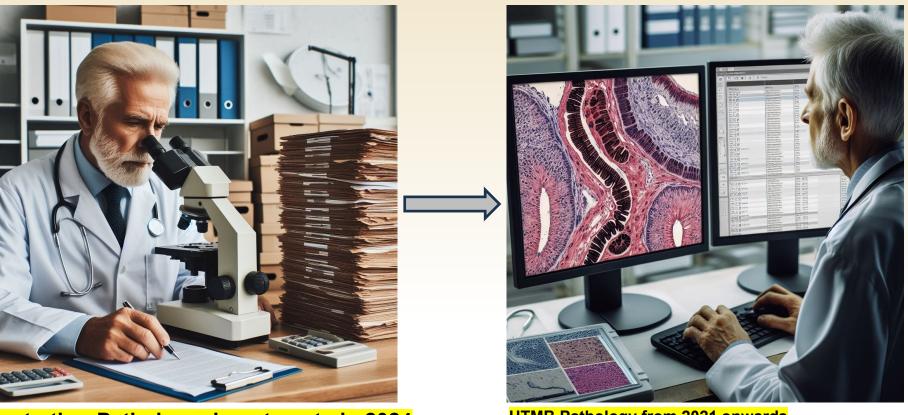


#### **Traditional Anatomic Pathology**





#### **Digital Transformation of Anatomic Pathology**



Most other Pathology departments in 2024

UTMB Pathology from 2021 onwards One of the first all-digital departments in the USA



#### Pathologists Working with Al Algorithm get the Best Results

Metric	Pathologist	Al Algorithm	Pathologist with Al
Sensitivity	93%	93%	100%
Specificity	95%	99%	100%
Positive Predictive Value	89%	99%	100%
Negative Predictive Value	97%	97%	100%

#### Pathologists make mistakes, AI makes mistakes. Together they get it right.



#### **Breast Cancer Genetics: Needed for Prognosis and Selection of Best Therapy**

#### High penetrance genes

BRCA1

BRCA2

#### Medium penetrance genes

- CHK2
- **ATM**
- RAD51C
  - BRIP1
  - PALB2

PTEN

• p53

- **STK11**
- CDH1

#### **Done by PacBio Revio**



### PacBio Revio Instrument



#### Will be used for:

- Pharmacogenomic Sequencing
- Whole Exome and Whole Genome Sequencing
- Single-Cell Genomics



Gene Editing Research

#### Diagnostic Evaluations For People with Clinically Apparent Illness

**Cardiometabolic DMT** 

**Classical Hematology** 

**Cancer Screening** 



#### Diagnostic Evaluations For People with Clinically Apparent Illness

#### **Autoimmunity**

**Opioid toxicology** 

Long Covid and Covid Immunity

Fatty liver of obesity



## **Unexpected benefits of** the test recommendations and result interpretations provided by pathologists in the Diagnostic Center



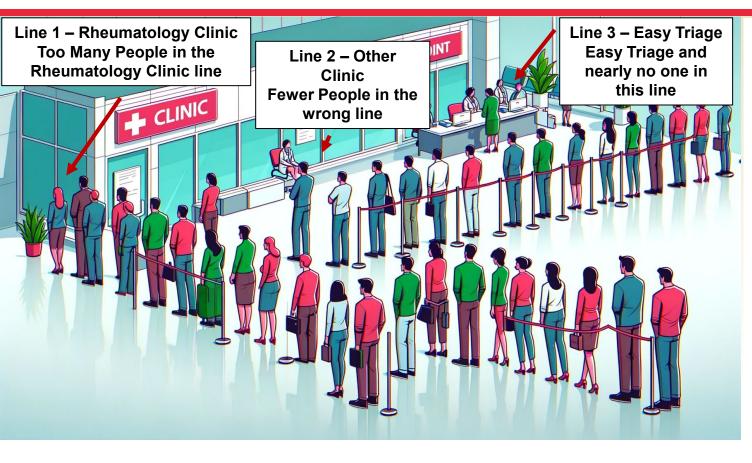
Reduction of Physician "Pajama Time" : Interpretation of Routine Tests with Abnormal Results

- CBC shows MCV with 1 percentage point high
- Chemistry panel shows ALT and AST with minor elevation

# Patient wants an email or phone call with her physician about both



#### **Improve Patient Access By Seeing the Right Ones**



Clothes Key: Red Clothes – High Suspicion for Rheumatologic Disease

Blue and White Clothes – Intermediate Suspicion for Rheumatologic Disease BUT most likely positive for SOME Disease

Green Clothes – Low suspicion of Disease



#### **Survey of Rheumatology**

By our estimation

- <u>33.7%</u> of patients could be Delayed or Triaged completely (Green Clothes)
- <u>17.8%</u> of patients would have additional tests performed prior to visit making it easier for Rheumatology (White/Blue Clothes)



 Leaving <u>48.5%</u> of patients with high suspicion of disease clear to be seen in the Clinic (Red Clothes)



#### There are at least 50 DMTs in line

They can be organized by Disease group: Coagulation Presenting sign/symptom: Bleeding Abnormal laboratory test result: Prolonged PTT

A "Trigger" sign or symptom or lab test result will ideally create an option for a Diagnostic Center consult Potential Clients for the Diagnostic Center

## Health Care Enterprises

## Academic Medical Centers

## Community Hospitals



### Potential Clients for the Diagnostic Center

- Sick patients
- Healthy patients for Pharmacogenomics and Exome Analysis
  - International clients physicians/patients
  - Concierge/VIP patients



# When Can Individual DMTs within the Diagnostic Center Be Fully Operational?

# Only when the potential demand for hundreds of cases per day can be met for every diagnostic area



# Waiting for the New Iphone: Are There Enough?



# Why Did it Take 30 Years?

- Original idea : 1984
- First implementation for coagulation only : 1995
- First major national presentations of concept and service : 1997-98
- Creation of Diagnostic Management Team name : 2010 Three National meetings on the DMT : 2017-2019 Implementation of 5 new DMTs, including one for alleged child abuse cases : 2014-2024
  - And then the pandemic occurred, and a clinical need and a new opportunity appeared that launched the DMT with much greater capacity



"Recent" Technical Advances in Construction of the Eiffel Tower Were Absolutely Required for its Construction

> The tallest structure of its time could only have been built because of technical advances developed within 20 years of its construction – which was just over 2 years from January 28, 1887 to March 15, 1889

With less advanced processes, it might have been limited to a height of a few stories and required much more than 2 years to build.



The "Recent" Technical Advances in Construction of the Eiffel Tower Which Were Absolutely Required for its Construction

- Iron manufacturing techniques enabled the production of high-quality wrought iron
- Newly advanced calculations in structural engineering permitted optimized lattice structure with necessary strength and flexibility
- Pneumatic riveting available to assemble the towers thousands of elements
- And at least 10 more...



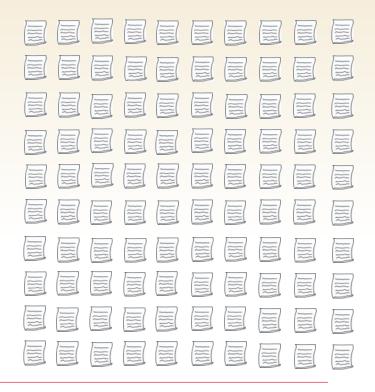
Nearly 30 Years Passed Before the Technical Capabilities Arose to Permit Widespread Interpretation of All Diagnostic Results by Experts in ONE site

- Vectorization to allow searching of Epic and match it with one of many written comments
- Artificial intelligence using generative language models to assemble correct content of interpretive comments

10 laboratory interpretations/day prepared manually



500 interpretations/day prepared with new technology in the same amount of time





# Limitations of the Manual Approach

- Case volume too low to support a full salary
- Content knowledge required of expert is substantial
- No expectation of consultative clinical activity – only technical activity

# **Grant from Sealy & Smith Foundation**

- 9 million dollars over 3 years
- Hiring pathologists and nonpathologists in other roles
- If revenue generation occurs, more pathologists and staff can be hired

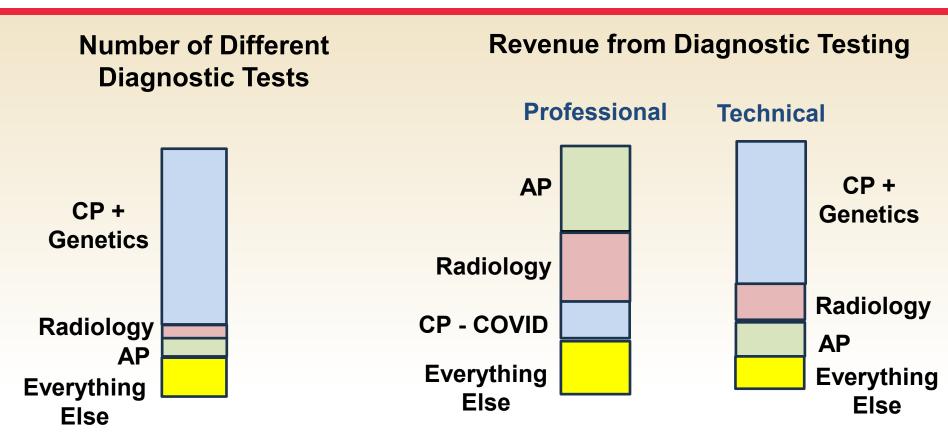


#### **SmartSheets – Toxicology example**

																									_	_	_	-
Jan 22, 2024 Jan 29	9, 2024 Feb	5, 2024	Feb		Feb 19, 2024					Feb 26, 2024					Mar 4, 2024					Mar 11, 2024								
22 23 24 25 26 27 28 29 30 31													_	_								-						_
M T W T F S S M T W	T F S S M T W	T F S S	M T W	T F S	SN	TN	w	TF	S S	M	T	W	F	FS	S	M	Т	w	T	FS	S S	M	T	W	T	FS	i s	
Build case list UI																												
Build holy book vector																												
	Build evaluation UI																											
	Epic clinical data																											
	Build language model pipeline																											
Develop initial promp	ot																											
	Determine RAG parameters																											
	Determine how to represent clinical data																											
													D	oc s	Scor	re In	nterp	os										



# **Approximations for UTMB**



Strategic Effort to Build a Clinical Service Providing Integrated Diagnostics: Pathology Department





Michael Laposata, M.D., Ph.D.

# Professor and Chairman, Department of Pathology

Director, John Sealy Diagnostic Center





#### Peter McCaffrey, M.D.

#### Director of Artificial Intelligence for UTMB

Director, Division of Bioinformatics & Artificial Intelligence

Co-Director, Center for Single Cell Genomics

Medical Director, Laboratory Information Systems

Assistant Professor, Department of Pathology

Assistant Professor, Department of Radiology





Christopher Zahner, M.D.

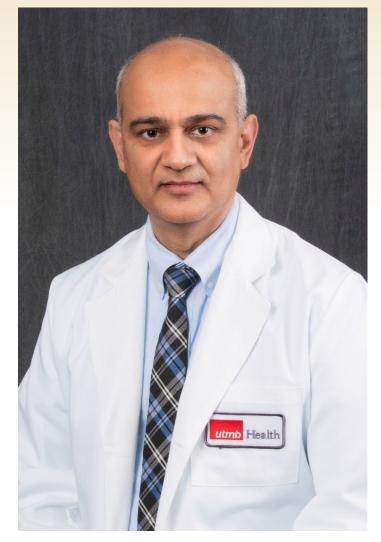
#### Director, Division of Clinical Pathology

Medical Director, Clear Lake, League City, & Angleton Danbury Campus Hospitals

Medical Director, Coagulation, & Point of Care Testing

Assistant Professor, Department of Pathology





# Harsh Thaker, M.D. Ph.D.

# Vice Chair, Anatomic Pathology

Professor, Department of Pathology



#### Juan David Garcia, MBA MLS



### Clinical Enterprise Director, Laboratory Services

### Director of Operations, John Sealy Diagnostic Center





# Bradley Grant, MD, JD

Pharmacogenomics & Molecular Genetics Specialist, John Sealy Diagnostic Center





## Heather Stevenson-Lerner, M.D., Ph.D.

# Director of Transplantation Pathology

Professor, Department of Pathology



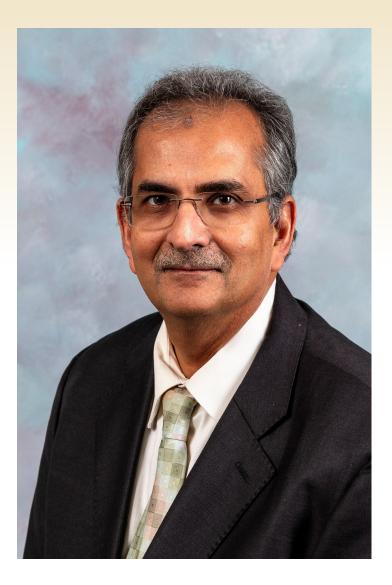


### Norma Hernandez, MBA

### Administrator, UTMB Academic Enterprise

### Director of Finance, John Sealy Diagnostic Center





#### Amin Mohammad, Ph.D.

#### Clinical Consultant for the John Sealy Diagnostic Center from Baylor Scott & White

**Professor, Pathology** 





# Melanie Connolly, MS

Medical Illustrator & Animator, Department of Surgery

Director of Marketing, John Sealy Diagnostic Center





#### Christopher Welch, MBA, CBCS

#### Sr. Finance Manager, John Sealy Diagnostic Center





#### **Carly Goetschius**

### Program Manager, John Sealy Diagnostic Center





Stephanie Kubachka, MHA, MBA, MLS(ASCP)CM

#### Manager, Laboratory Services

#### Project Manager, John Sealy Diagnostic Center

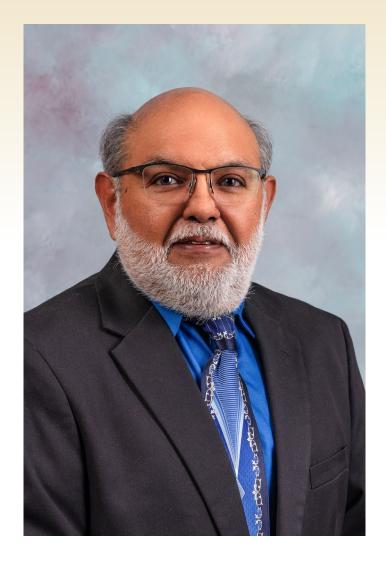




#### **Gladson John**

#### Software Systems Specialist II, John Sealy Diagnostic Center





#### **Victor Luciano**

#### Network Support Specialist, Pathology Administration



# **Links to Promotional Videos**

- **General Overview John Sealy Diagnostic Center.mov** - Dr. Michael Laposata
- **Intergrating Al.mov** Dr. Peter McCaffrey
- **DMT Improves Diagnosis Time.mp4** Dr. Chris Zahner
- Dr. Grant Personalized Medicine Pharmacogenomics.mp4 Dr. Brad Grant
- Digital Transformation and Adoption of Artificial Intelligence in Anatomic Pathology at UTMB.pptx – Dr. Harsh Thaker

