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

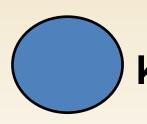


The State of Affairs in Diagnostic **Medicine in the** US

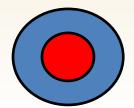


How Much Information is There to Know?

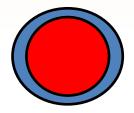




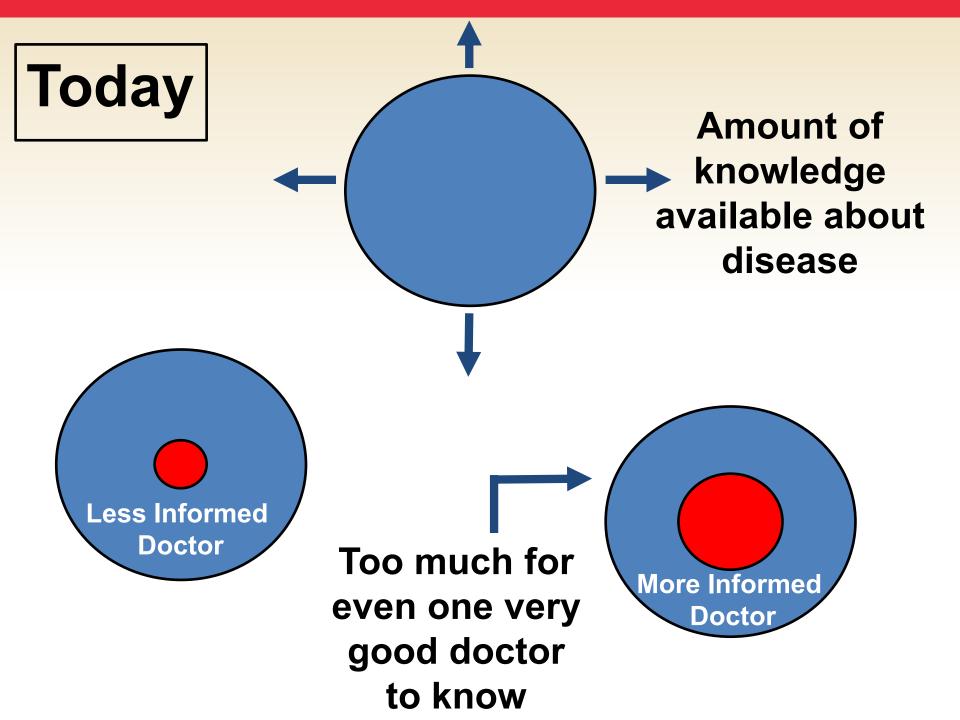
Amount of knowledge available about disease



Doctor does not know much about diagnosis



Doctor knows much about diagnosis



How Can a Clinician Know Enough?

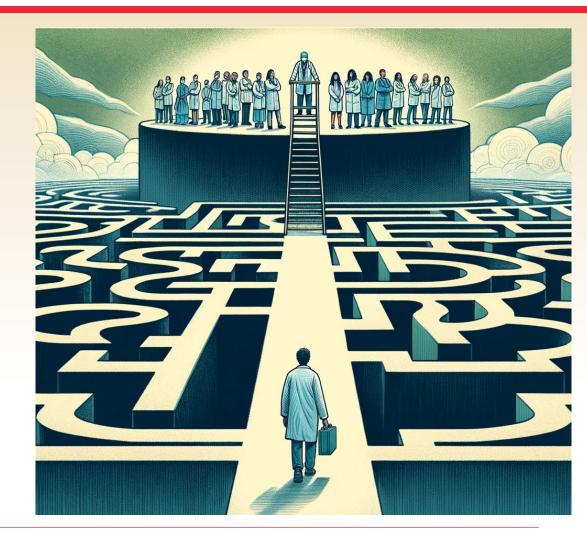
- They are ordering the right nonroutine tests?
- They are making the most up-to-date interpretation of the test results for the patient in their current state?





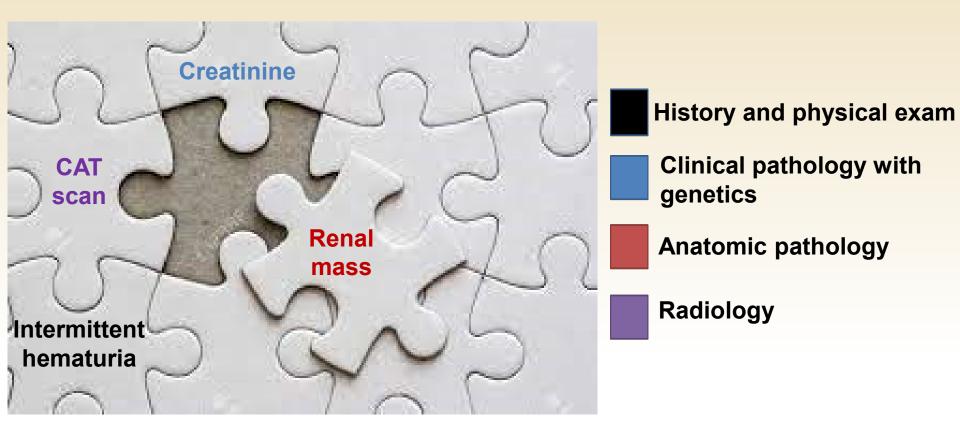
The Diagnostic Center Allows for Easy Access to Experts in All Diagnostic Fields

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



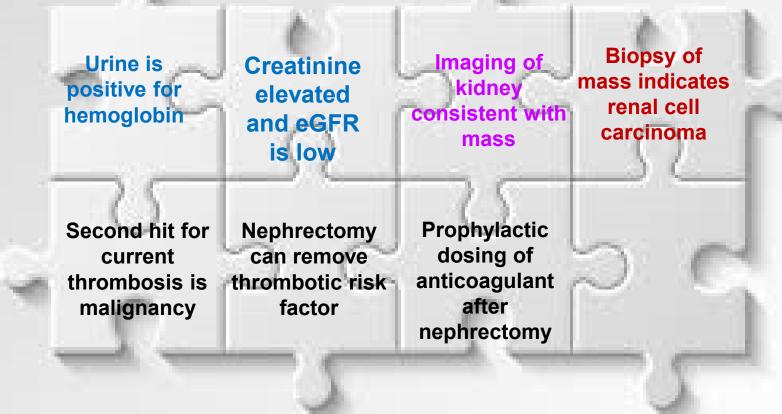


Integrative Diagnostics Must Be Implemented -There are Major Barriers to This Challenge



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





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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 ⁽¹⁾, ^{1,2} Najlla Nassery, ³ Adam C Schaffer, ^{4,5} Chihwen Winnie Yu-Moe, ⁵ Gwendolyn D Clemens, ⁶ Zheyu Wang, ^{6,7} Yuxin Zhu, ^{1,6} Ali S. Saber Tehrani, ¹ Mehdi Fanai, ¹ Ahmed Hassoon, ^{1,2} Dana Siegal^{8,9}

"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**



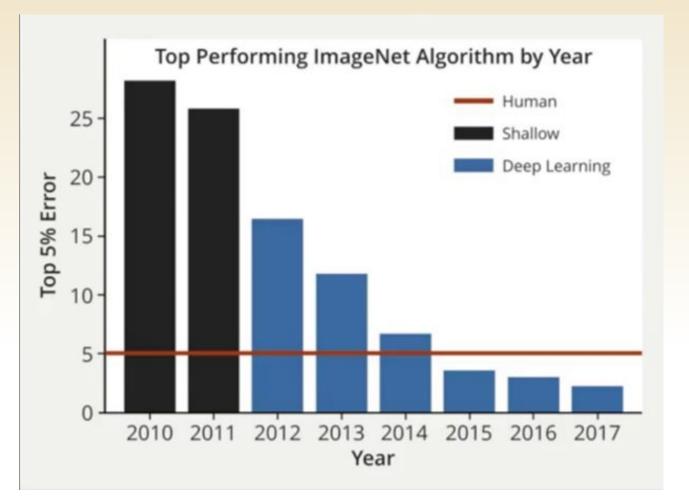
The Fear that Artificial Intelligence will Reduce Pathologist Income

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

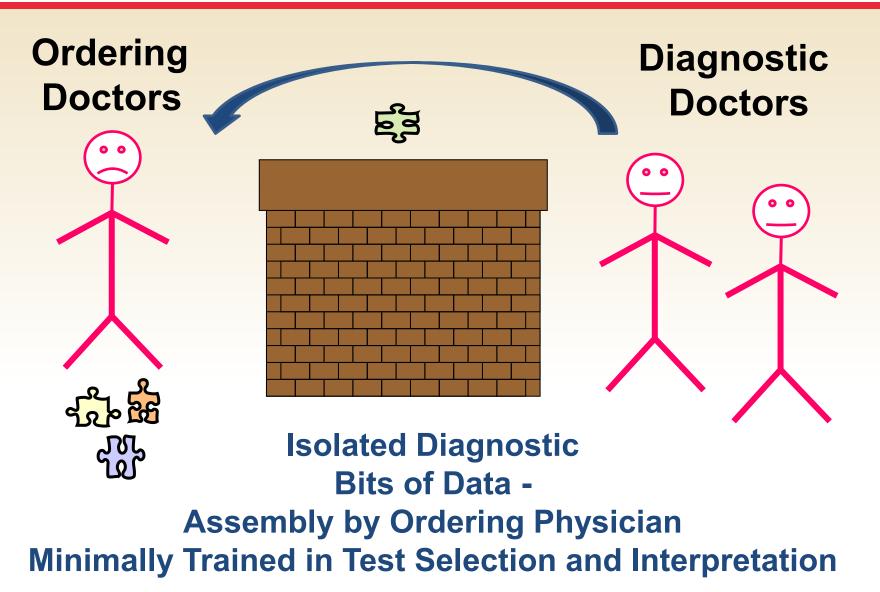
THE CONSEQUENCES OF A MISTAKE ARE TOO ENORMOUS



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



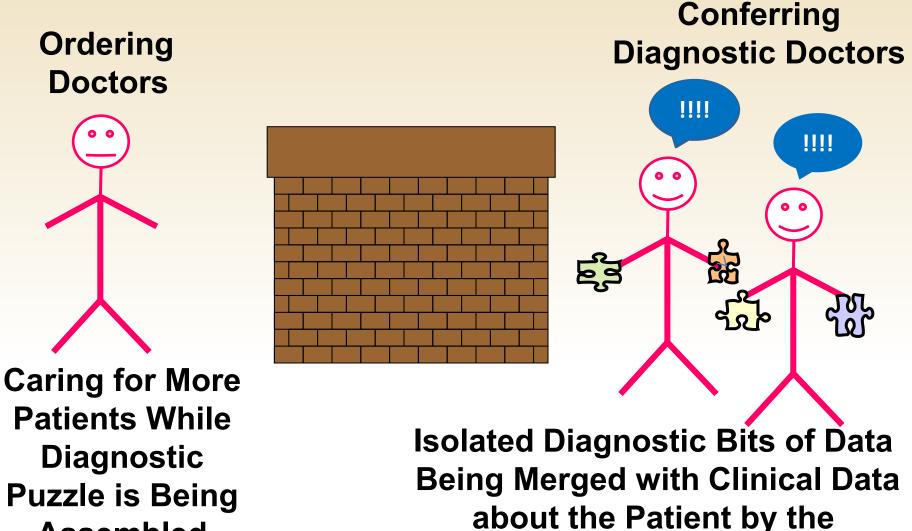
Not choosing pathology – An essay-based survey of first-year clinical residents

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%).



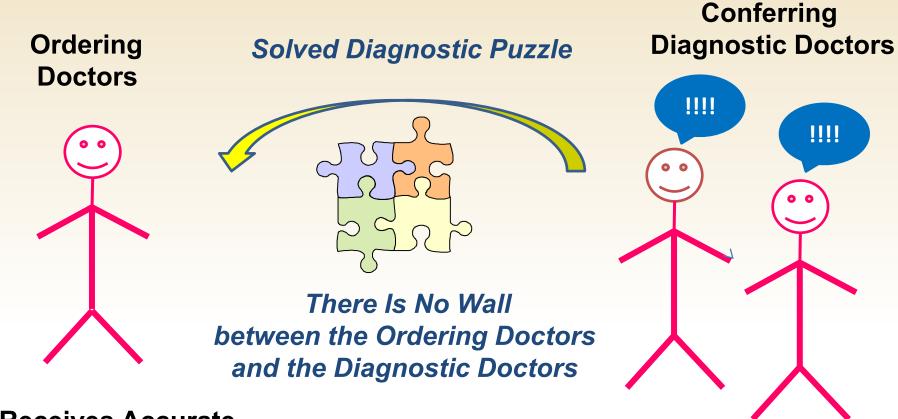
Until recently there was minimal payment for picking up the pieces and establishing a diagnosis



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



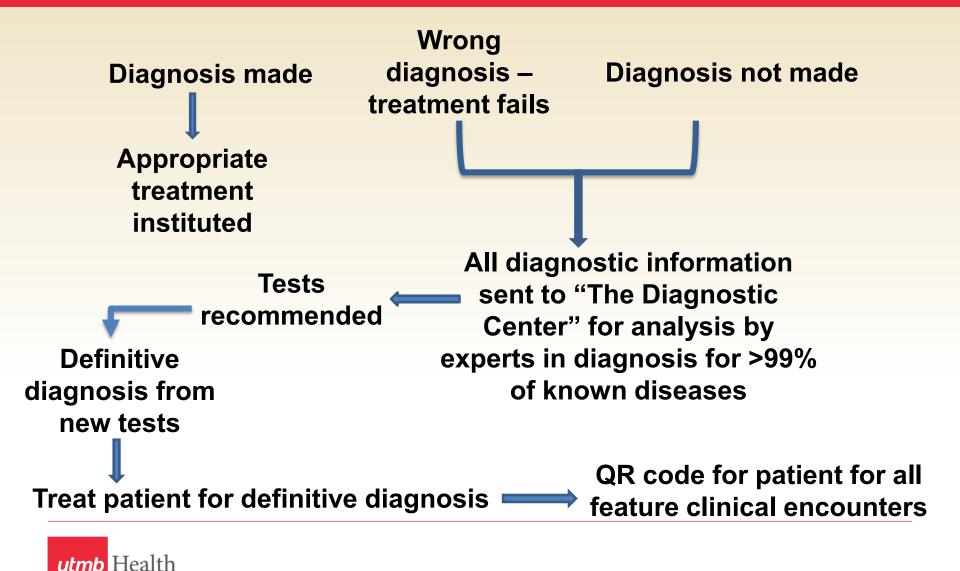
There is no plan to compete for laboratory test performance or the patient

Niche laboratory test performance will be offered as an option

This is a unique service in the practice of medicine



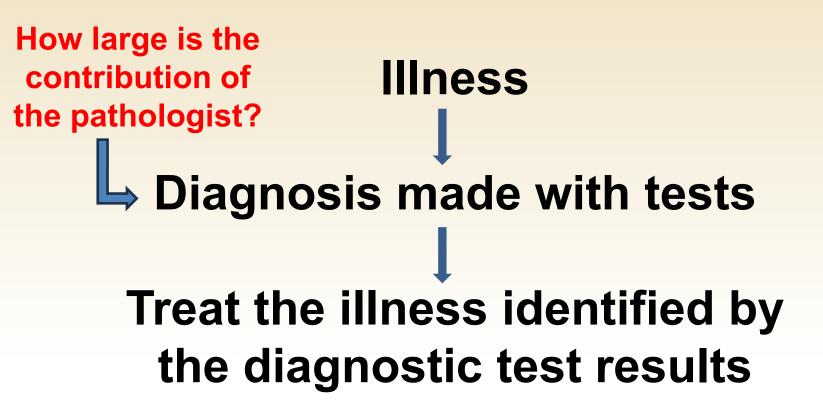
How the Diagnostic Center Works



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

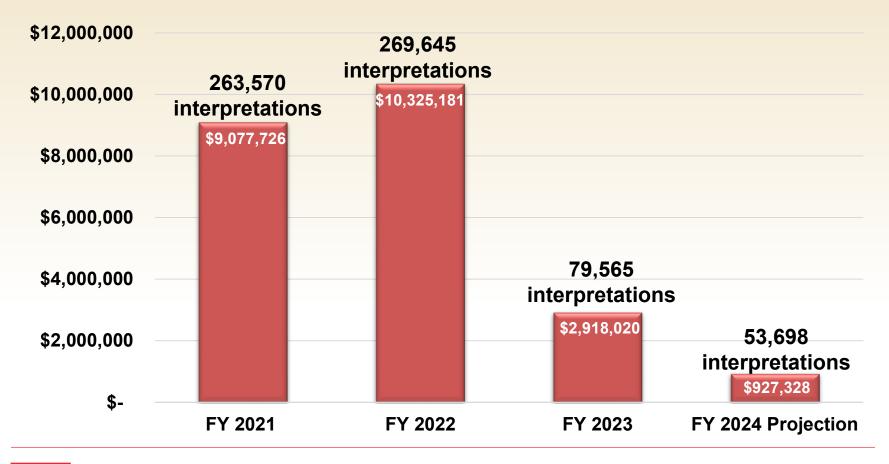
"Literature Scraping" on a Regular Basis

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



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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					
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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



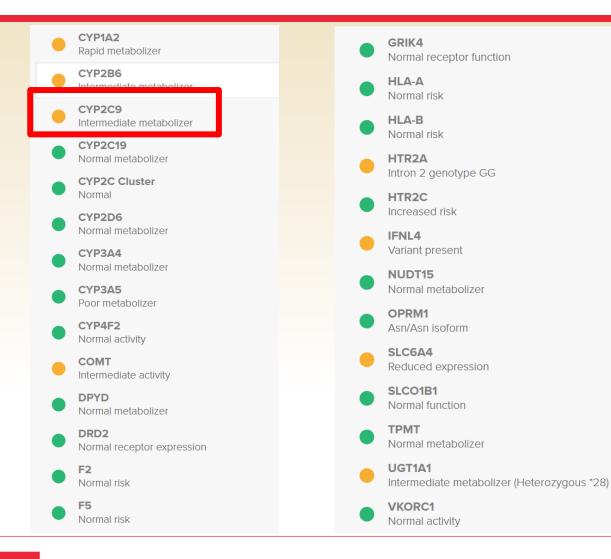
Clinical Benefit from the Pharmacogenomics DMT

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	Recommendations for Prescriber
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

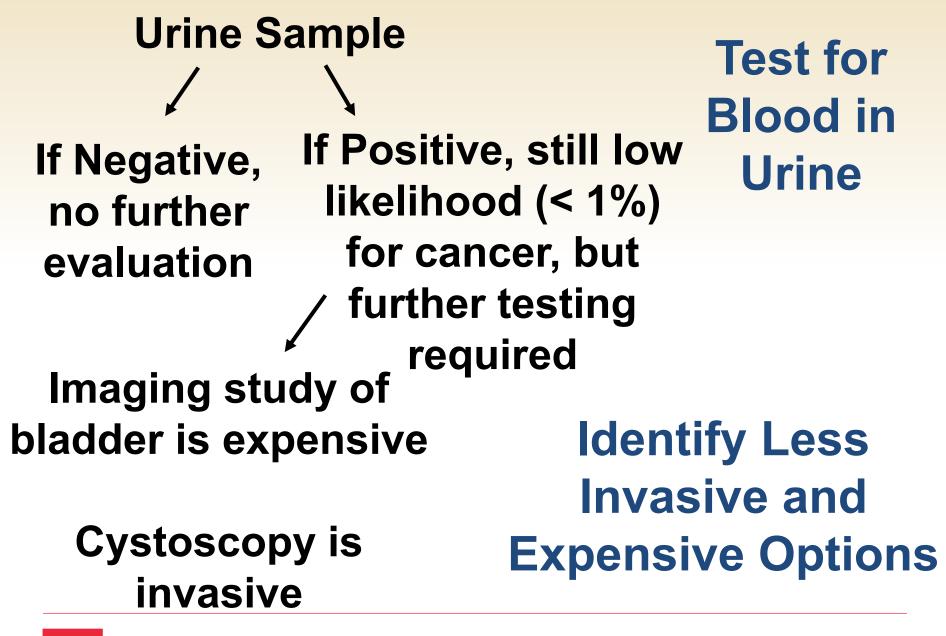
- An algorithm for testing starts with a test that is neither invasive nor expensive
- If the screening test is positive, the challenge is to find a second level test that is only modestly invasive or expensive



Algorithms to Screen for Different Types of Cancer

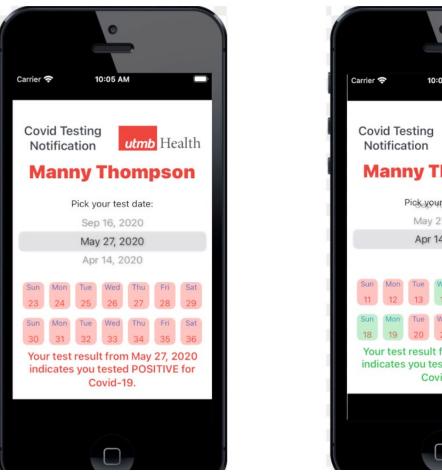
- It is understood that the final confirmatory test for cancer is likely to be invasive, expensive, or both
- The goal is to not take individuals who are suspected of having cancer and move them through an evaluation with invasive or expensive tests which may be harmful and <u>not</u> identify cancer







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

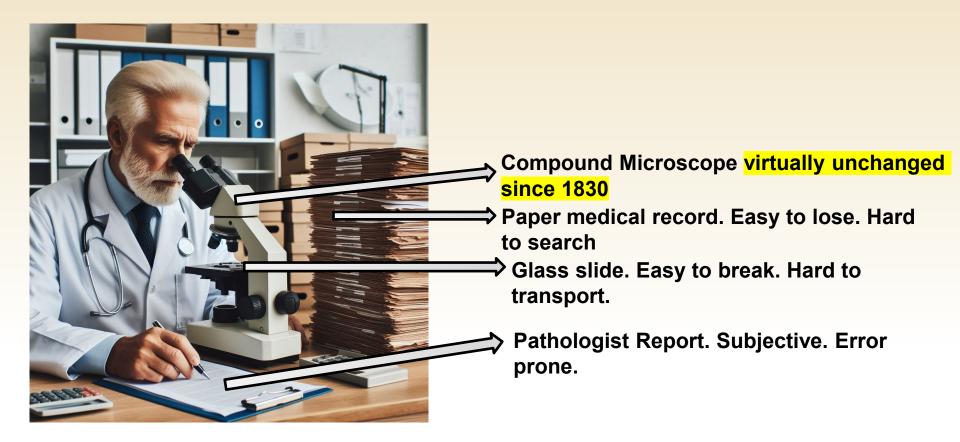
A Comprehensive High Sensitivity Cancer Diagnosis DMT



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

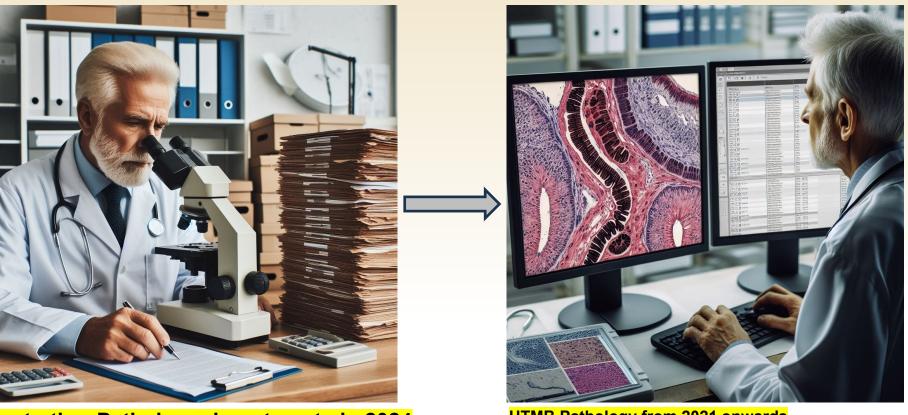


Traditional Anatomic Pathology





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



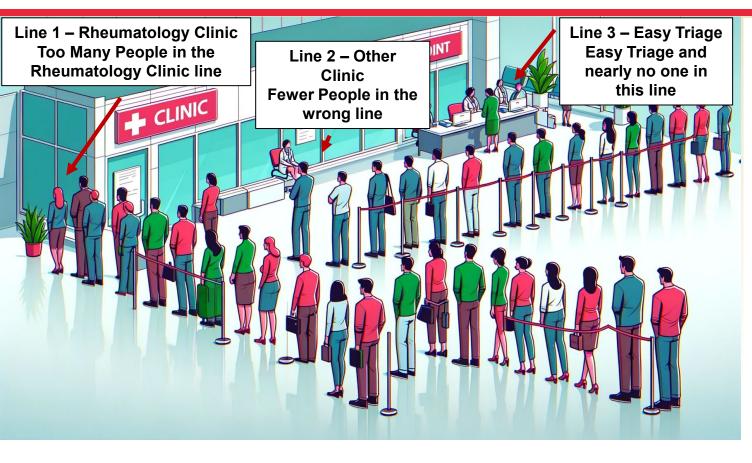
Reduction of Physician "Pajama Time" : Interpretation of Routine Tests with Abnormal Results

- 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...



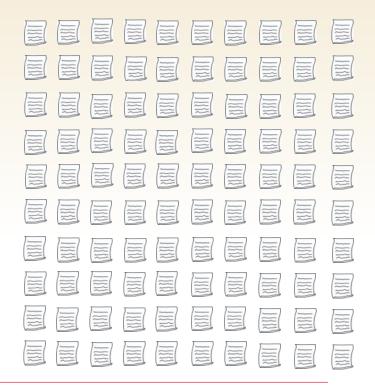
Nearly 30 Years Passed Before the Technical Capabilities Arose to Permit Widespread Interpretation of All Diagnostic Results by Experts in ONE site

- 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

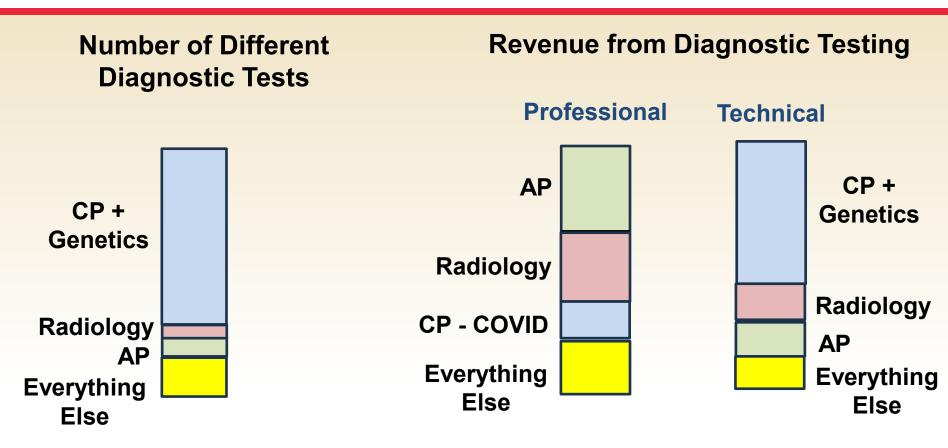


SmartSheets – Toxicology example

																									_	_	_	-
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.

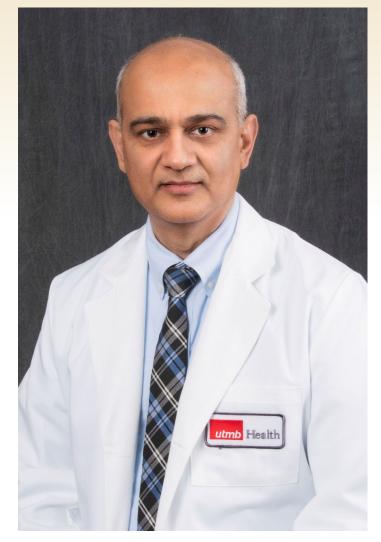
Director, Division of Clinical Pathology

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



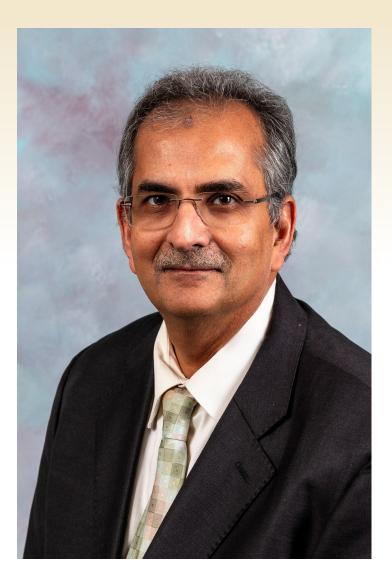


Norma Hernandez, MBA

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

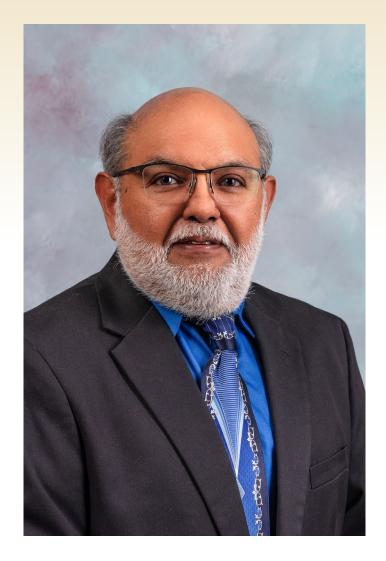




Gladson John

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

