Genomic Predictors of Clinical Outcome in Gastric Cancer: The Singapore Experience

Patrick Tan, MD PhD gmstanp@duke-nus.edu.sg

Global Leaders in Genomic Medicine Conference Washington DC - Jan 2014









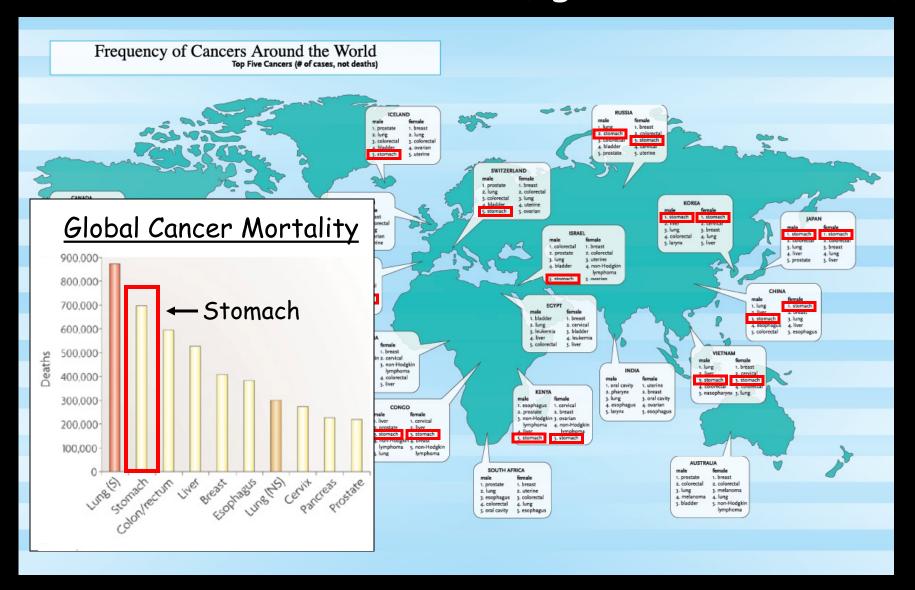
Biomedical Sciences (BMS) in Singapore (2003-2013)

Significant and increasing BMS support from Singapore government

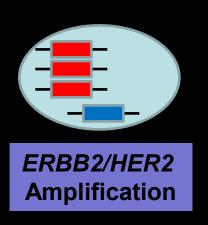
Funding from Three Major Ministries (Trade/Industry, Education, Health)

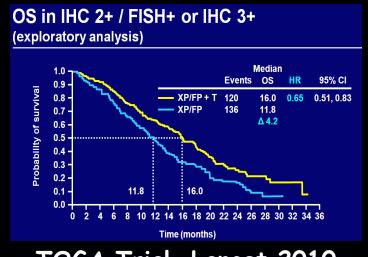
Multiple Research Institutes (eg Biopolis) and Academic Medical Centres (eg Singhealth, National University Hospital)

Focus Area: Asian Cancers (eg Gastric/Stomach)

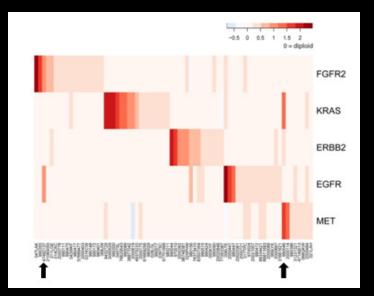


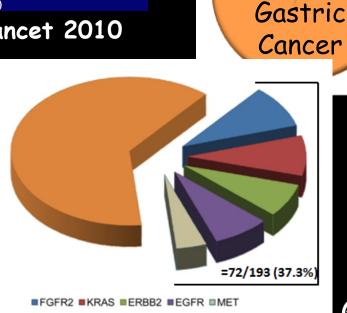
Genomic Amplifications Highlight GC Therapeutic Targets











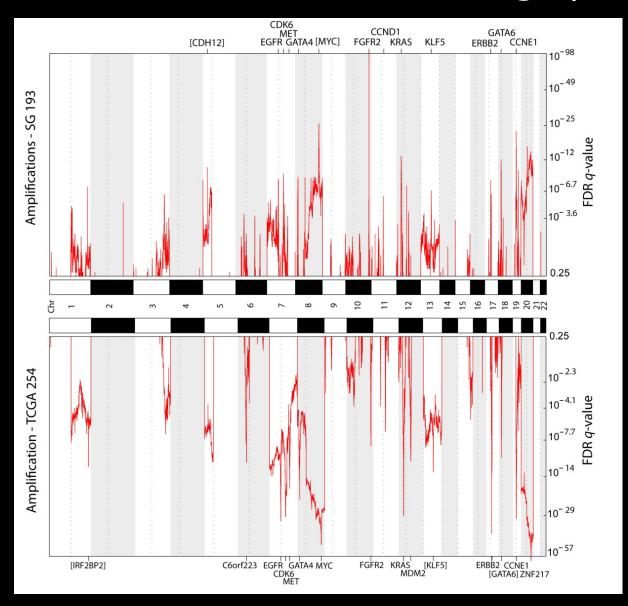
RTK/RAS Absent

ERBB2 Positive

(8-10%)

Gut 2012

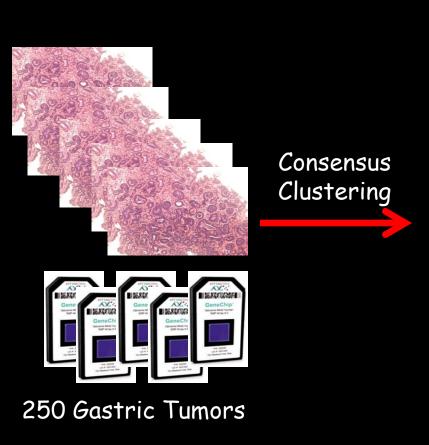
Genomic Amplifications in Asian and Caucasian GCs - Concordant and Largely Similar

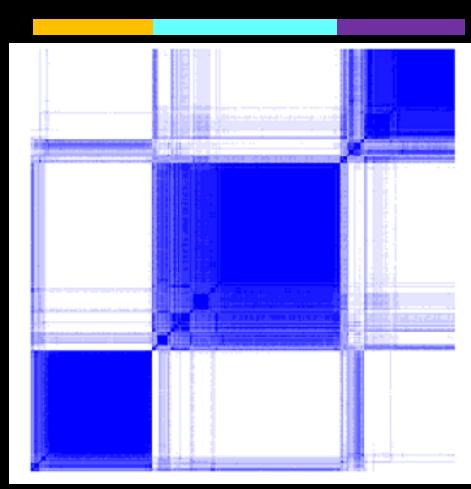


Singapore Cohort

TCGA Cohort (USA)

Transcriptome Clustering Identifies THREE GC Subtypes: Integration with Pathology



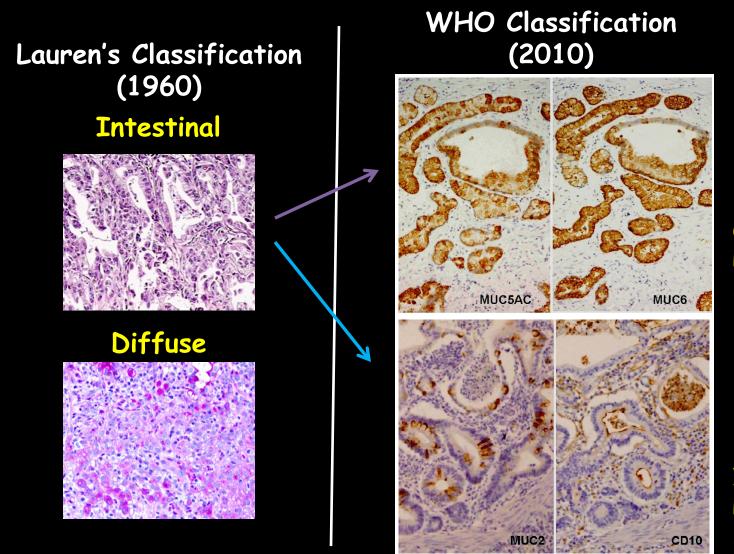


Consensus Subtype Matrix

GC Genomic Subtypes: Mesenchymal, Proliferative, and Metabolic

Pa+	hways				
Pα' β α_	Genomic Subtype	Histological Features	Associated Genes/Pathways	Drug sensitivity (Preclinical)	
alin	Mesenchymal	• Diffuse subtype	 EMT pathways CSC pathways TGFB mTOR signalling 	Sensitive to PI3K/AKT/mTOR inhibitors	
	Proliferative	Intestinal subtype	 Genomic instability TP53 mutations Cell cycle DNA replication Mitosis Copy number alterations (ERBB2/HER2 and KRAS) 		cle eplication s
	Metabolic	Gastric subtype	 Metabolic processes Digestion Secretion SPEM 	 Increased sensitivity to 5- FU 	olic processes ion, Secretion

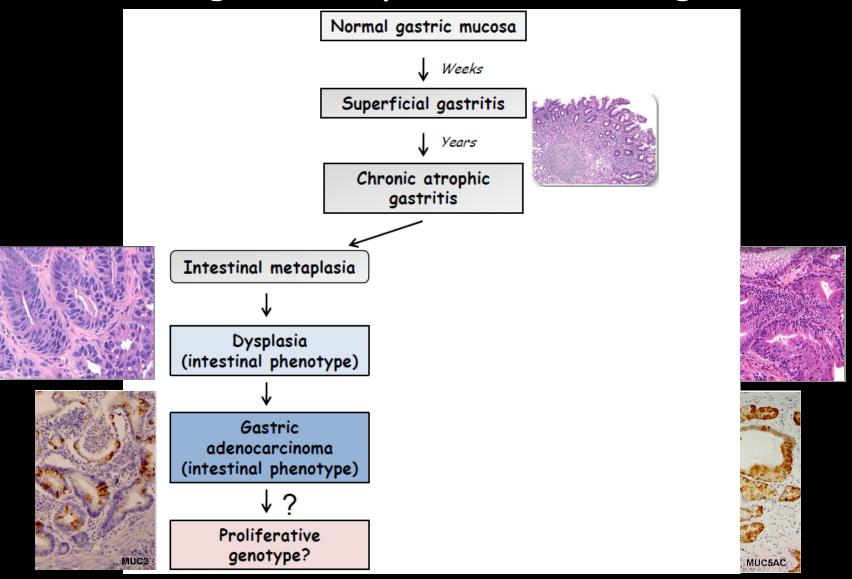
Genomic Subtyping May Drive Improved Pathology



Gastric
Phenotype
Aka Metabolic

Intestinal
Phenotype
Aka Proliferative

Working Roadmap for GC Carcnogenesis



Courtesy Fatima Carneiro, IPATIMUP

Dissecting Asian Cancers - Some Contributions from Singapore

Exome sequencing of gastric adenocarcinoma identifies recurrent somatic mutations in cell adhesion and chromatin remodeling genes

Nature Genetics (2012)

A common *BIM* deletion polymorphism mediates intrinsic resistance and inferior responses to tyrosine kinase inhibitors in cancer

Nature Medicine (2012)

Oncofetal Gene SALL4 in Aggressive Hepatocellular Carcinoma

N Engl J Med (2013)

The POLARIS Program - Enabling Genomic Medicine in a City-State

Funded by A-STAR (Agency for Science, Technology and Research) for 3 years

<u>Pilot</u> clinical use of genomic testing (cancer and genetic diseases)

National network of CAPcertified laboratories at hospitals and research institutes













Some POLARIS Operating Principles

Genomic medicine labs should be deployed WITHIN existing clinical frameworks

Frameworks for GENETIC testing should exist PRIOR to GENOMIC testing

Genomic tests should leverage on EXISTING RESEARCH COMPETENCIES

Tests providing CLINICAL UTILITY will lead to clinician buy-in



POLARIS - Current Status (2013)

First POLARIS Test - TGFBI Eye Test (early 2014)

Genomic labs targeting national certification in mid 2014 (Illumina + Reflex Validation)

Test revenues are distributed among network partners on cost-recovery basis

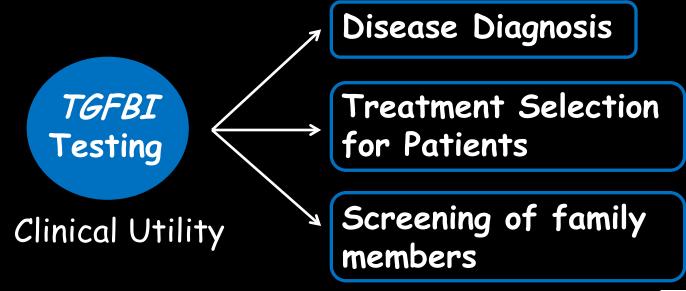
Second POLARIS Test - 90 gene GI Panel (3rd quarter 2014)



Stromal Corneal Dystrophies (SCDs) and TGFBI Testing

- Inherited disorders leading to loss of corneal transparency.
- TGBFI mutations underline the majority of stromal CDs.







PARTIES INVOLVED IN POLARISTM TGFBI TEST





SNEC/SGH

- Patients &Consultation
- Test Ordering
- Blood Collection





GIS/SERI

- Project Management

- Mutation Database



NUHS

- Sequencing
- Mutation Rpt

POLARISTM TGFBI Test



Challenges in Developing a Singapore Framework for Genetic/Genomic Testing

Legal and licensing agreements across institutions and ministries are often complex

Reimbursement options for genetic assays that cross medical centres

General lack of genetic counsellors and advisors

Official polices on patient consent, incidental findings and aggregation of genetic/genomic data

Thanks and Questions

Prof John Wong Prof Wee Joo Chng

Niantao Deng Liang Kee Goh

Zhengdeng Lei Steve Rozen

Khay Guan Yeoh Wei Peng Yong Yoshiaki Ito Christopher Wong Tony Lim Pauline Ng Huck Hui Ng

Prof Jodh Mehta Dianne Poh Evelyn Koay