



The Multidimensional Nature of Cancer Classification: Consequences for Molecular Pathology

International Agency for Research on Cancer
Lyon, France

Ian A Cree FRCPATH
Head, WHO Classification of
Tumours

International Agency for Research on Cancer



creei@iarc.fr

Declaration of Interests

- I am a pathologist, based at the International Agency for Research on Cancer, part of the World Health Organization
- All opinions expressed are personal, and not those of any of the organisations above.

CAROLI LINNÆI
EQUITIS DE STELLA POLARI,
ARCHIATRI REGII, MED. & BOTAN. PROFESS. UPSAL.;
ACAD. UPSAL. HOLMENS. PETROPOL. BEROL. IMPER.
LOND. MONSPEL. TOLOS. FLORENT. SOC.

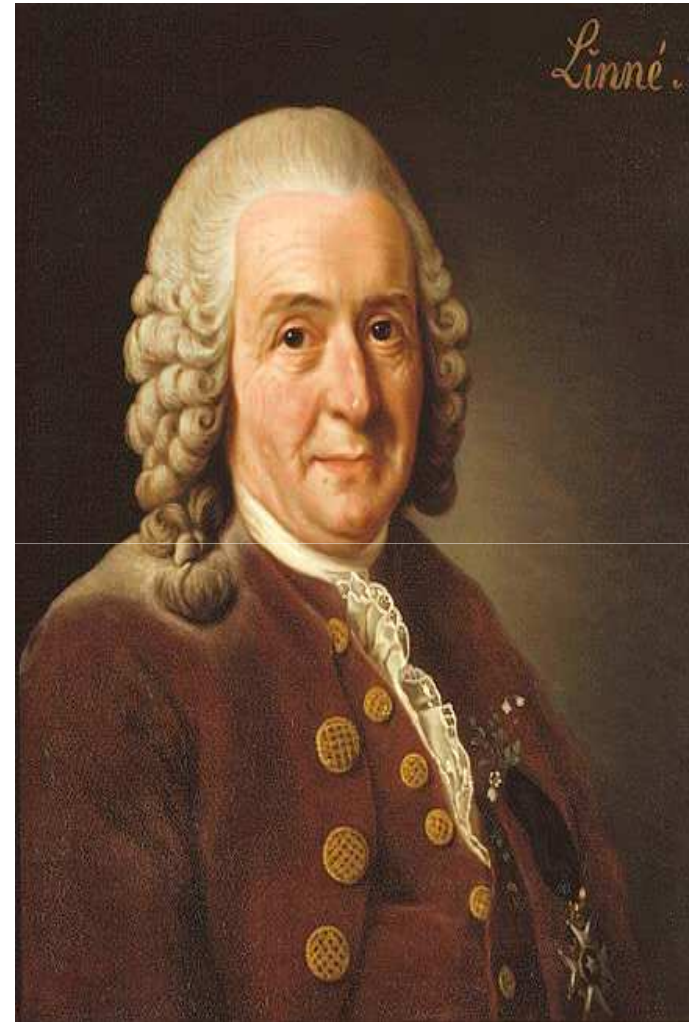
**SYSTEMA
NATURÆ**

PER
REGNA TRIA NATURÆ,
SECUNDUM
CLASSES, ORDINES,
GENERA, SPECIES,
CUM
*CHARACTERIBUS, DIFFERENTIIS.
SYNONYMIS, LOCIS.*

TOMUS I.

EDITIO DECIMA, REFORMATA.
Cum Privilegio Sæ Ræ Mæstis Sveciæ.

HOLMIÆ,
IMPENSIS DIRECT. LAURENTII SALVII,
1758.



Carl Linnaeus (1707 –
1778)

- *Taxonomy* is the science of defining and naming groups of biological organisms on the *basis of shared characteristics*
- *Cancer classification* is based on shared characteristics of cancers – currently mainly histology and genetics.

WHA10.18 The Tenth World Health Assembly resolved, '...to continue work on formulating international definitions of nomenclature and statistical classification...' (May 1957)

WHO Blue Books

Scientific evidence



Illustrative cases

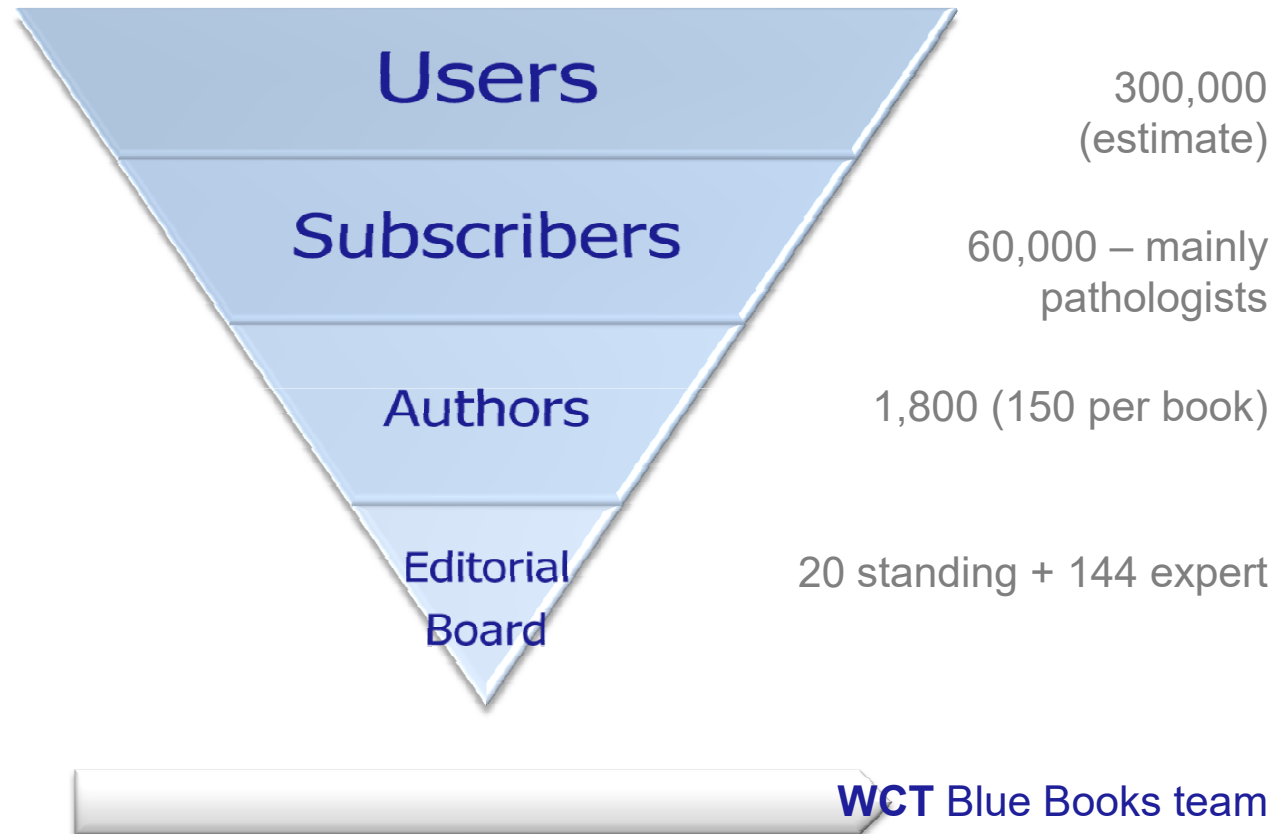


Diagnostic criteria



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WHO Blue Books Faculty



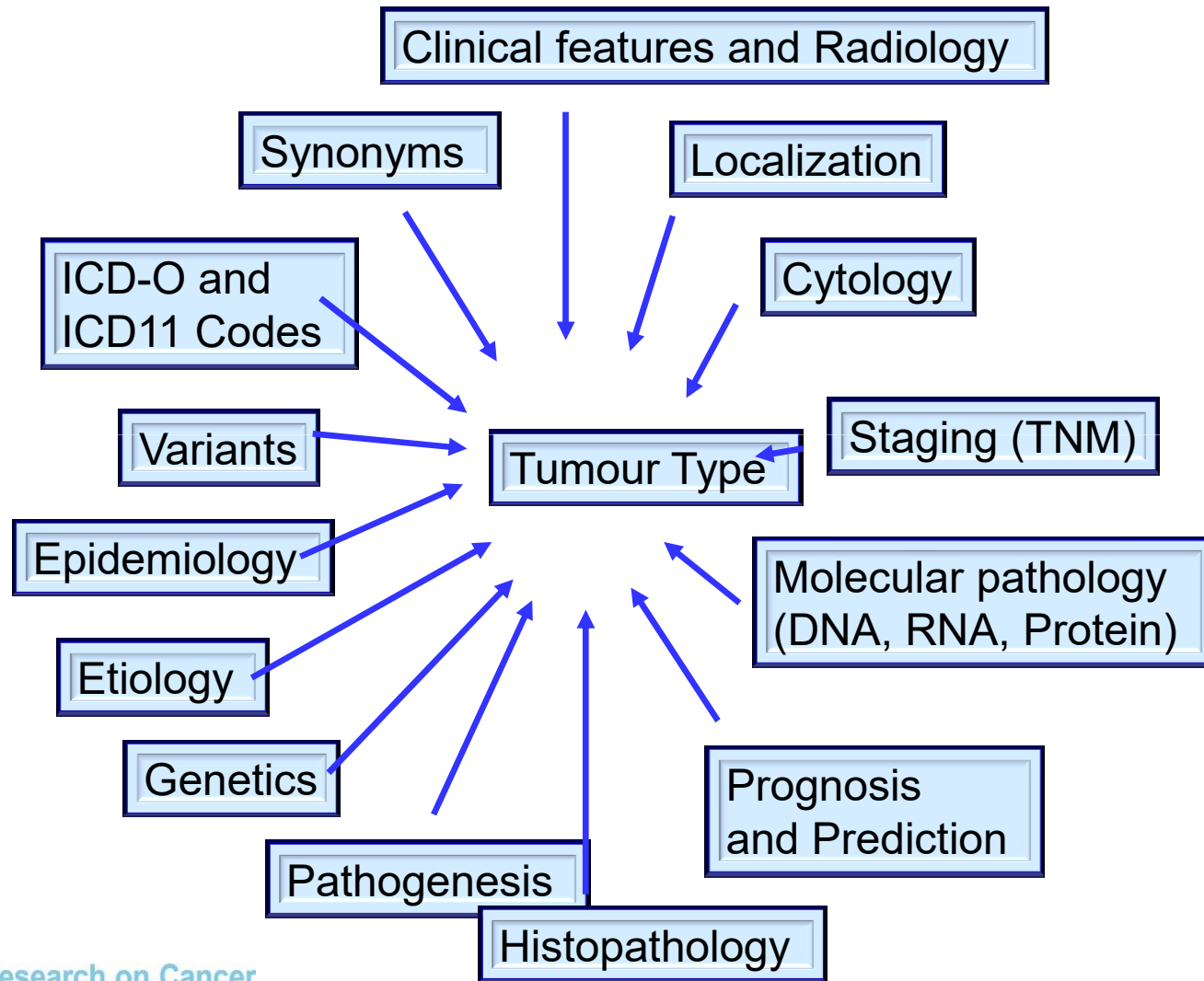
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Classification terms

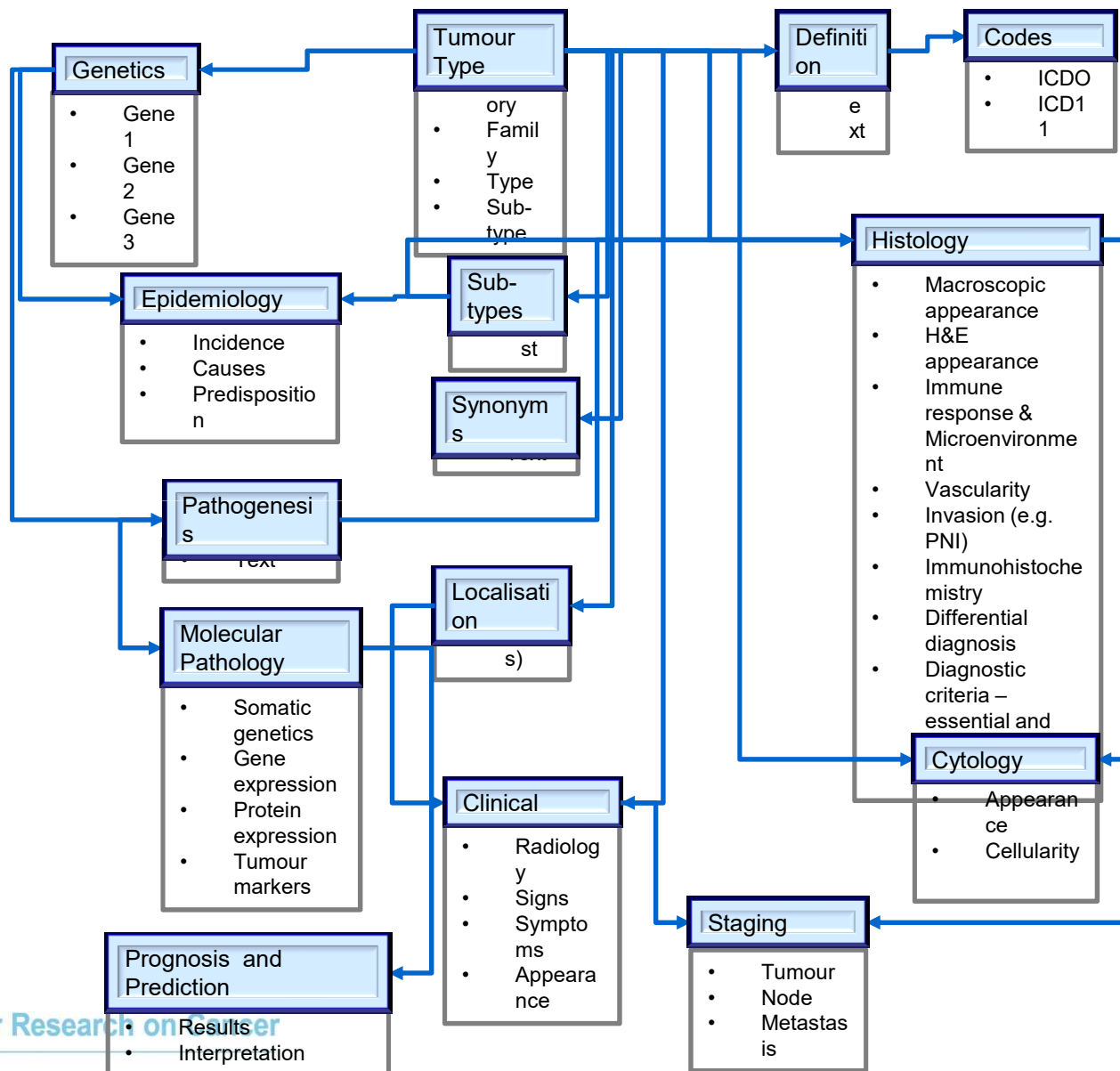
- *Site*, e.g. Stomach
- *Category*, e.g. Epithelial neoplasms
- *Family (Class)*, e.g. Adenomas and other premalignant neoplastic lesions
- *Type*, e.g. Adenoma
- *Variant (Sub-Type)*, e.g. Pyloric-gland type

Stage and Grade are dealt with separately....

The multi-dimensional nature of cancer classification



The multi-dimensional nature of cancer classification



WHO BB Layout (5th Series) DRAFT

- Definition
- ICD-O and ICD11 Codes
- Related Terminology (Synonyms)
- Subtypes
- Localization
- Clinical features and Radiology
- Epidemiology
- Etiology
 - Causes
 - Predisposing factors (Genetics)
- Pathogenesis
- Macroscopic appearance
- Histopathology
 - H&E appearance
 - Immune response & Microenvironment
 - Vascularity
 - Invasion (e.g. PNI)
 - Immunohistochemistry
 - Differential diagnosis
- Cytology
- Molecular pathology
 - Somatic genetics
 - Gene expression
 - Protein expression
 - Tumour markers
- Diagnostic criteria – essential and desirable
- Staging (UICC TNM)
- Prognosis and Prediction
 - Prognostic factors
 - Predictive biomarkers
- Links to other resources
 - ICCR reporting guidance
 - TNM (UICC)

The 5th Series WHO Classification of Tumours

- Digestive System Tumours
- Breast Tumours
- **Soft Tissue and Bone Tumours**
- Female Genital Tumours
- Thoracic Tumours
- Urinary and Male Genital Tumours
- Central Nervous System Tumours
- Head and Neck Tumours
- Endocrine Tumours
- Haematolymphoid Tumours
- Skin and Adnexa Tumours
- Eye and Orbit Tumours
- Paediatric Tumours
- Neuroendocrine Tumours
- Hereditary Tumours

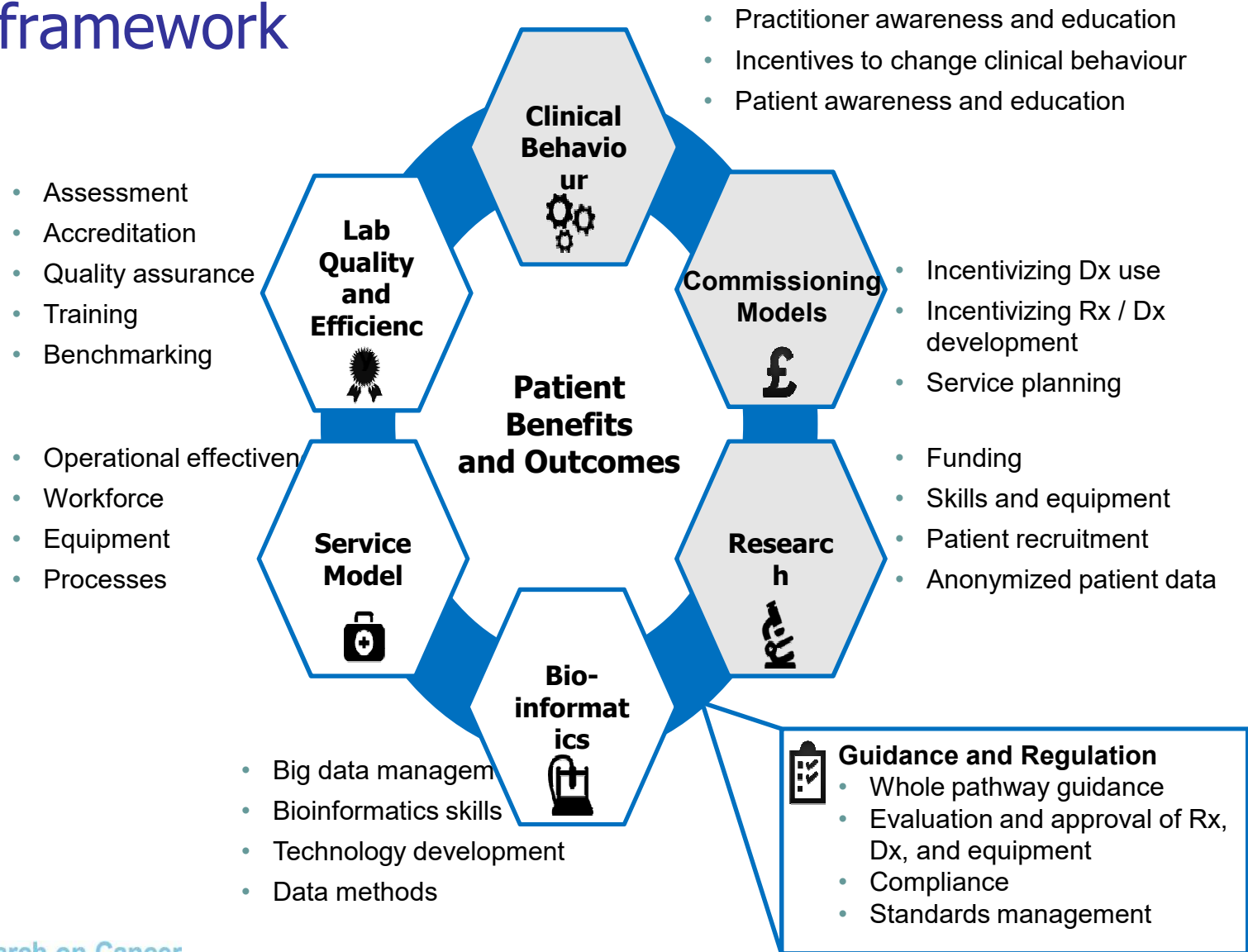
<http://whobluebooks.iarc.fr>

Pathology is changing...

- Cancer classification has previously been based on consensus histopathological opinion with limited molecular input.
- Pathology is undergoing a rapid transformation due to the introduction of new technologies to practice.
- The understanding of cancer at a molecular level is now at a point where it needs to be integrated into its diagnosis.
- Digital pathology and image analysis are now also producing new insights, and providing quantitative justification of many existing diagnostic criteria, while challenging others.

- There is an urgent need to integrate these facets of diagnosis into cancer classification internationally.

Stratified medicine ecosystem framework



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Key Demand side Supply side Guidance and Regulation



Guidance for laboratories performing molecular pathology for cancer patients

Ian A Cree,^{1,2} Zandra Deans,³ Marjolijn J L Ligtenberg,⁴ Nicola Normanno,⁵ Anders Edsjö,⁶ Etienne Rouleau,⁷ Francesc Solé,⁸ Erik Thunnissen,⁹ Wim Timens,¹⁰ Ed Schuurung,¹⁰ Elisabeth Dequeker,¹¹ Samuel Murray,¹² Manfred Dietel,¹³ Patricia Groenen,⁴ J Han Van Krieken,⁴ for the European Society of Pathology Task Force on Quality Assurance in Molecular Pathology and the Royal College of Pathologists

Key guidance for molecular pathology

- Covers every stage of the process
- Produced by EQA providers
- Endorsed by the RCPATH and ESP

Key guidance for clinical use of NGS in cancer

- Covers every stage of the process
- Produced by EQA providers (IQN Path)
- Endorsed by the RCPATH and ESP

Virchows Arch

DOI: 10.1007/s00428-014-2325-7



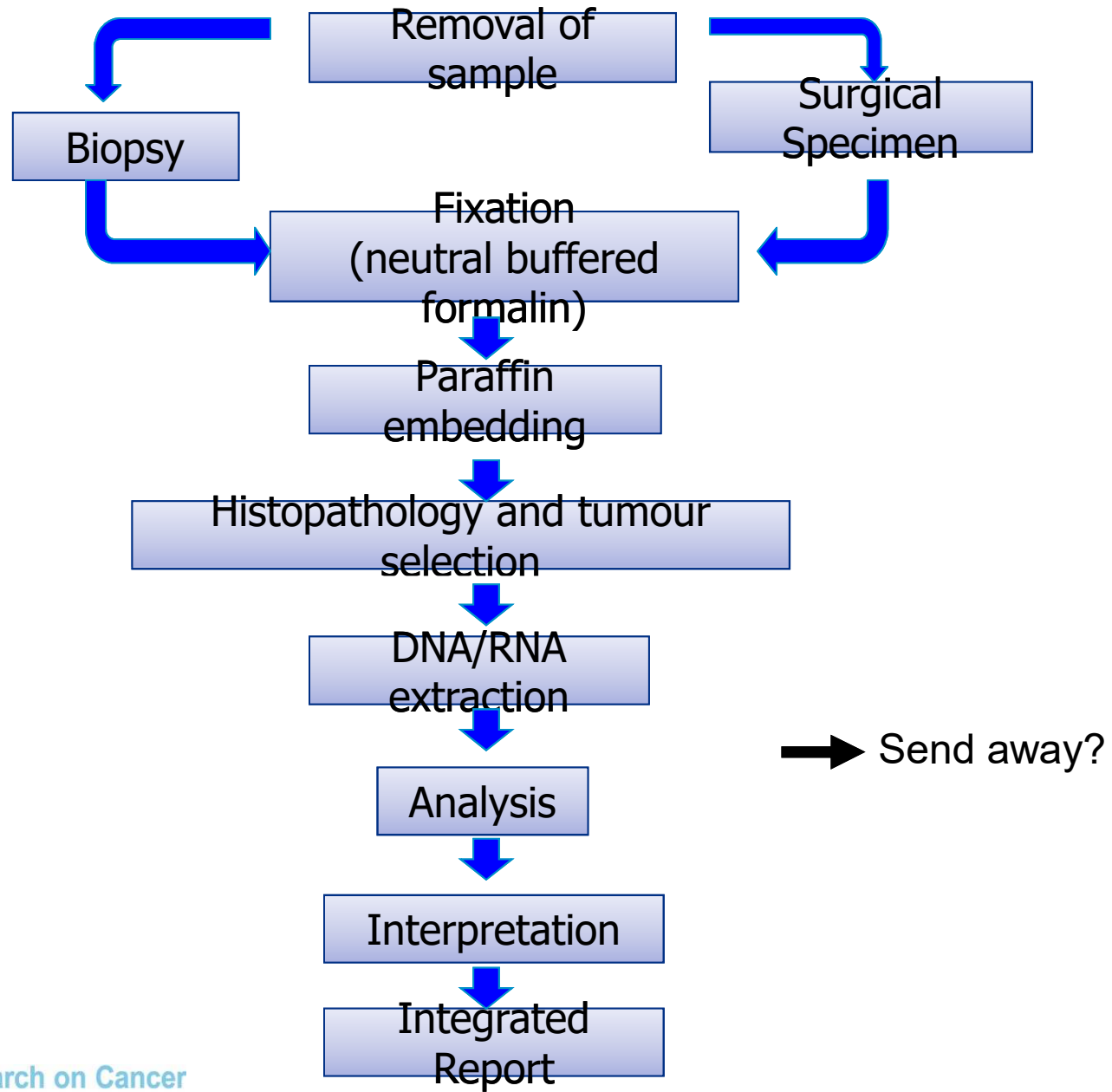
REVIEW AND PERSPECTIVES

Integration of next-generation sequencing in clinical diagnostic molecular pathology laboratories for analysis of solid tumours; an expert opinion on behalf of IQN Path ASBL

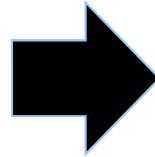
Zandra C Deans¹, Jose Luis Costa², Ian Cree³, Eli Dequeker⁴, Anders Edsjö⁵, Shirley Henderson⁶, Michael Hummel⁷, Marjolijn J L Ligtenberg⁸, Marco Ladda⁹, Jose Carlos Machado¹⁰, Antonio Marchetti¹¹, Katherine Marquis¹², Jianne Mason¹³, Nicola Normanno¹⁴, Etienne Rouleau¹⁵, Ed Schuurung¹⁶, Keesa-Maria Snelson¹⁷, Erik Thunnissen¹⁸, Bastiaan Tops¹⁹, Gareth Williams²⁰, Han van Krieken²¹, Jacqueline A Hall^{13,14}. On behalf of IQN Path ASBL

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The problem....



- Tissue Safe

Sample receipt and handling

- Dedicated SOPs required
- Defined processing – changes can affect DNA and RNA recovery
- The use of fixed tissue that has been previously frozen is not advised, and decalcification is likely to reduce DNA and particularly RNA recovery
- Close collaboration between clinical teams and histopathology essential
- Contamination during cutting – issues exist: mitigate
- Microdissection or punches, not laser capture routinely
- Estimate percentage neoplastic cells present

DNA and RNA extraction

- Methods described – no preference for manual or automated systems
- Storage of DNA and RNA should be controlled carefully. Accession logs or bar-coded vials can be used to prevent sample mis-identification.
- Temperature logs should be maintained - good practice:
 - Store extracted DNA and RNA samples, clearly labelled, at -20 C or -80 C respectively.
 - Store PCR products in a separate freezer at -20 C or -80 C
 - Store sequencing libraries at -20 C or -80 C

Choice of analytical method

- Testing requirements are defined by clinical need
- Panel testing is now recommended for some cancers
- Servicing and training costs a consideration

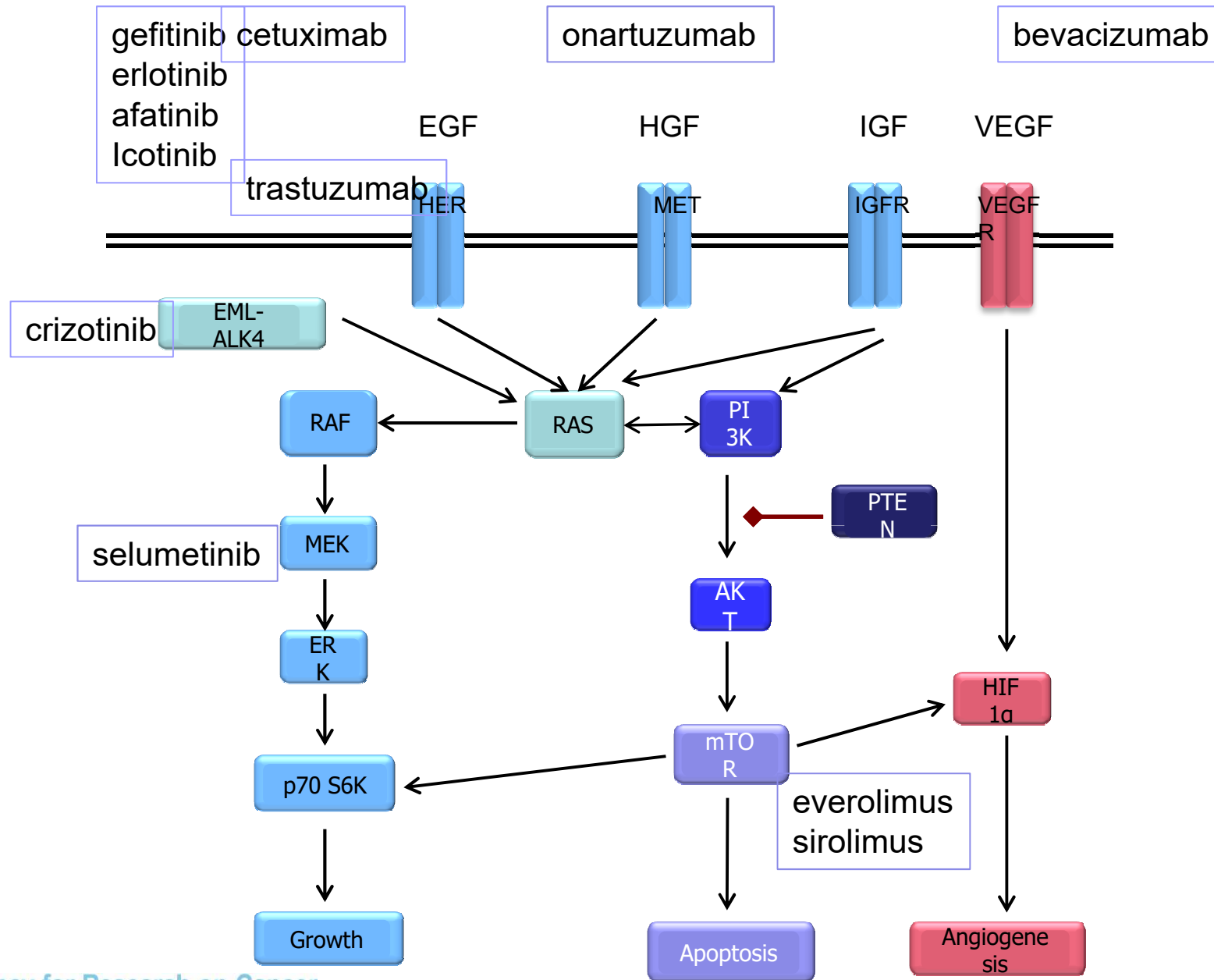
Performance of molecular tests

- Monitor the performance against confidence intervals
- Use externally sourced controls where possible for in house tests
- Turnaround time v. timeliness
- Continuity planning

Mutation testing by Polymerase Chain Reaction (PCR)

- Defined mutations in hotspots
- High sensitivity – use <150 bp primers for FFPE
- Not comprehensive
- Simple set up and performance
- Rapid – results in hours, not days
- Inexpensive
- Can be automated and multiplexed







Contents lists available at ScienceDirect

Pathogenesis

journal homepage: <http://www.pathogenesisjournal.com/>



Original Article

Development and validation of a TaqMan Array for cancer mutation analysis



Hugh Kikuchi ^{a,b}, Anne Reiman ^{a,b}, Jenifer Nyoni ^a, Katherine Lloyd ^c, Richard Savage ^d,
Tina Wotherspoon ^a, Lisa Berry ^a, David Snead ^{a,b}, Ian A. Cree ^{a,e,f,*}

^a Department of Pathology – Coventry and Warwickshire Pathology Services (CWPS), University Hospitals Coventry and Warwickshire, Coventry CV2 2DX, UK

^b Warwick Medical School, University of Warwick, University Hospitals Coventry and Warwickshire, Coventry CV2 2DX, UK

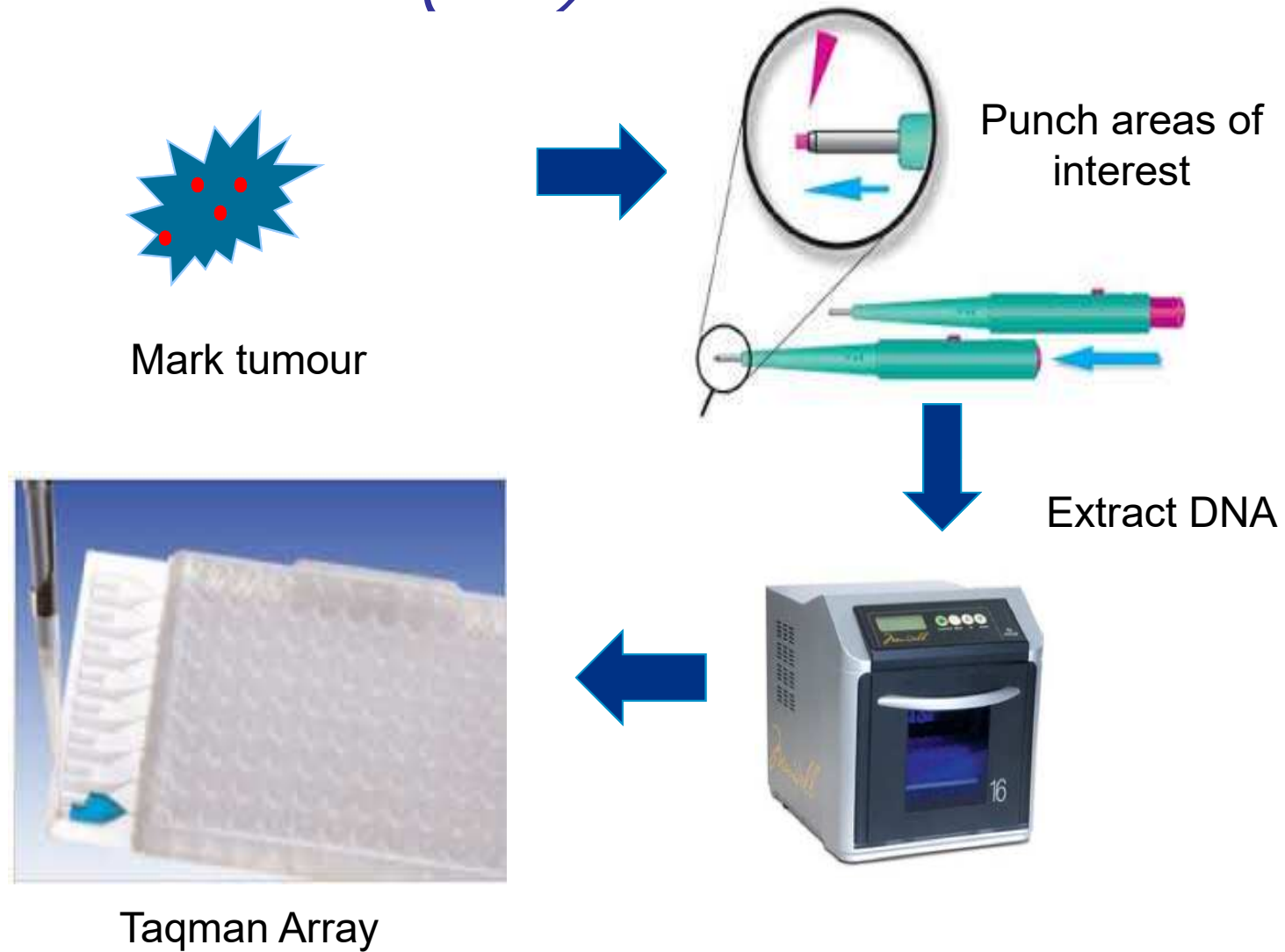
^c MOAC DTC, University of Warwick, Gibbet Hill Road, Coventry CV4 7AL, UK

^d Systems Biology Centre, University of Warwick, Gibbet Hill Road, Coventry CV4 7AL, UK

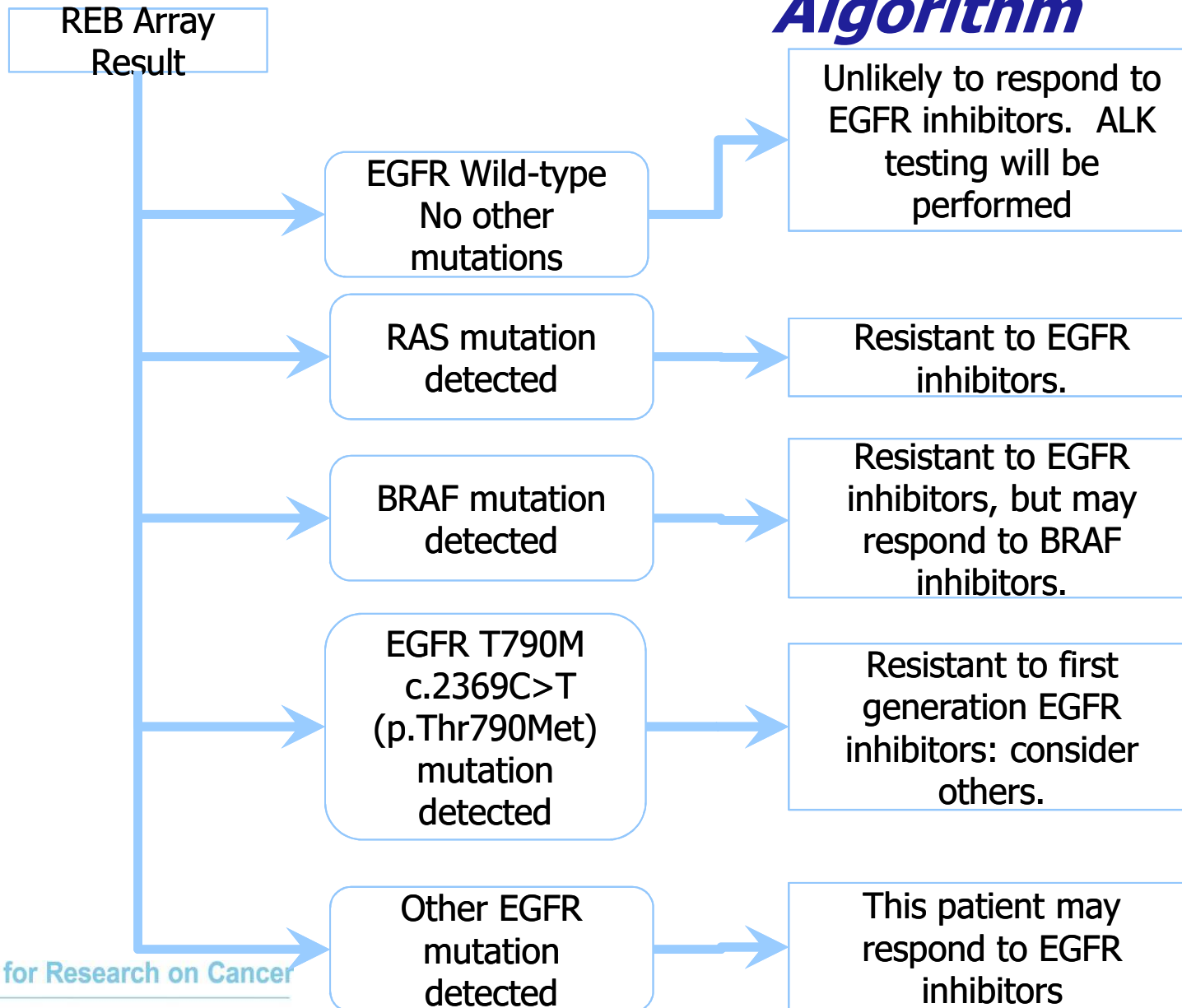
^e Institute of Ophthalmology, University College London, Bath Street, London EC1V 9EL, UK

^f Centre for Technology Enabled Health Research (CTEHR), Faculty of Health & Life Sciences, Coventry University, Coventry CV1 5FB, UK

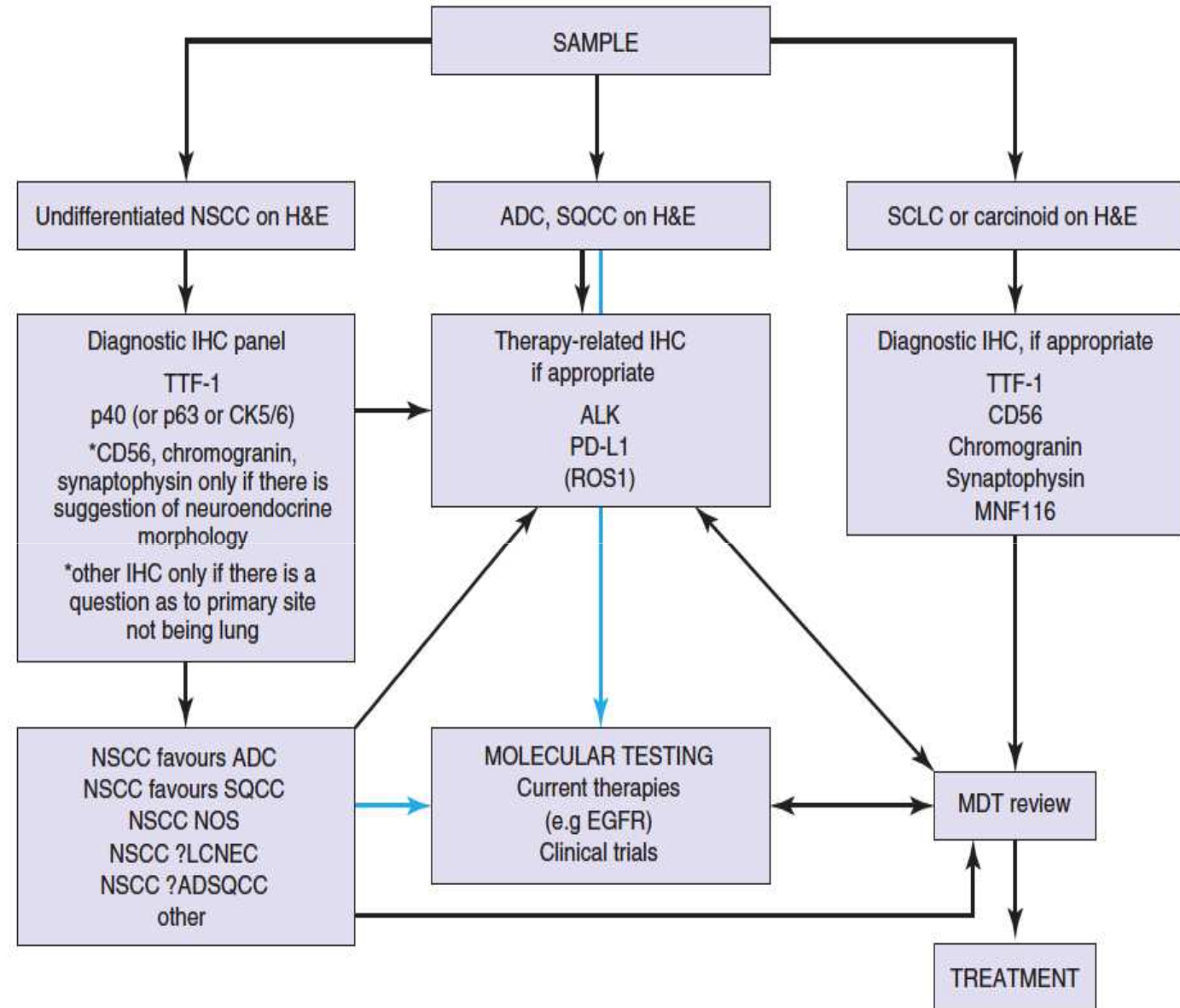
Taqman array PCR panel testing: Ras – EGFR – BRAF (REB) array



NSLSC Algorithm



NSCLC algorithm



IonTorrent next-generation



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Ion Chef™ System

Workflow Automation for Sequencing



- Simple to use
 - Automated template prep AND chip loading – *library in, loaded chips out*
 - Simple reagent and consumables loading – *minutes of hands-on time*
 - Minimizes potential sources of user error and sequencing variability
- High throughput
 - Processes 2 chips and multiple barcoded samples within hours
- Flexible
 - Supports all Ion™ systems, chips, read lengths, and

Melanoma panel

Gene	Reference	Codons included	Number of Mutations
BRAF	NM_004333.4	466, 469, 583, 584, 586, 592, 594, 595, 597, 600, 601, 605, 614, 618	29
GNAQ	NM_002072.3	209, 359	6
GNA11	NM_002067.2	183, 209, 223	3
NRAS	NM_002524.4	12, 13, 18, 50, 59, 61, 68	29
c-KIT	NM_000222.2	553, 557, 559, 560, 566, 569, 576, 642, 655, 816, 820, 822, 823, 829, 853	18
KRAS	NM_033360	12, 61	5
MEK1/MAP2K1	NM_002755.2	111, 124, 203, 264	5

Reiman A, Kikuchi H, Scocchia D, Smith P, Tsang YW, Snead D, Cree IA. Validation of an NGS mutation detection panel for melanoma. BMC Cancer. 2017 Feb 22;17(1):150.

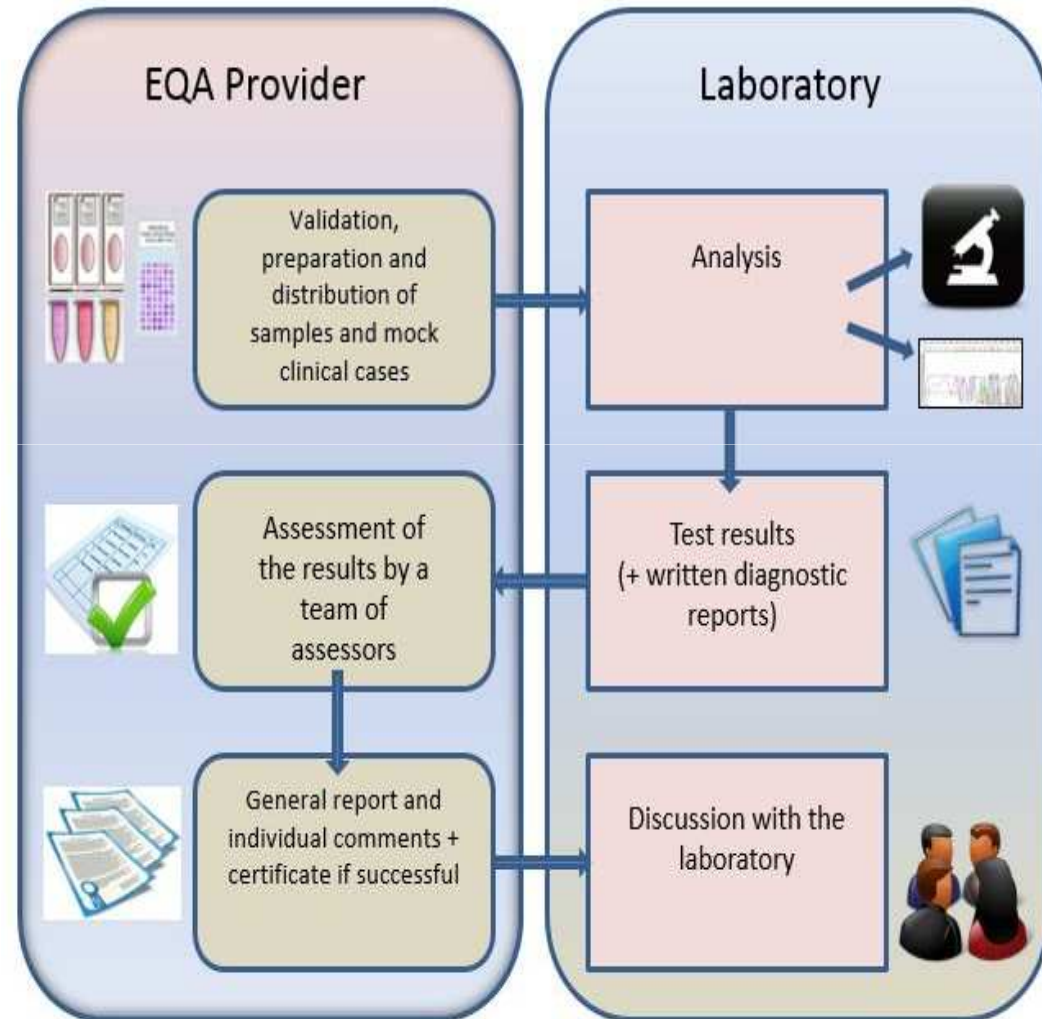
Accreditation and Quality assurance – Internal and External

- All laboratories providing molecular pathology services should have laboratory **accreditation** according to ISO 15189 or equivalent
- All laboratories performing molecular tests for cancer patients should be part of an external quality assessment (**EQA**) scheme
- Internal quality assessment (IQA) - the use of **control** materials within each run is recommended
- **Monitor** to assure end to end performance
- Implement changes when **errors** occur and pre-empt by looking at others' mistakes!

How does EQA help with testing quality?

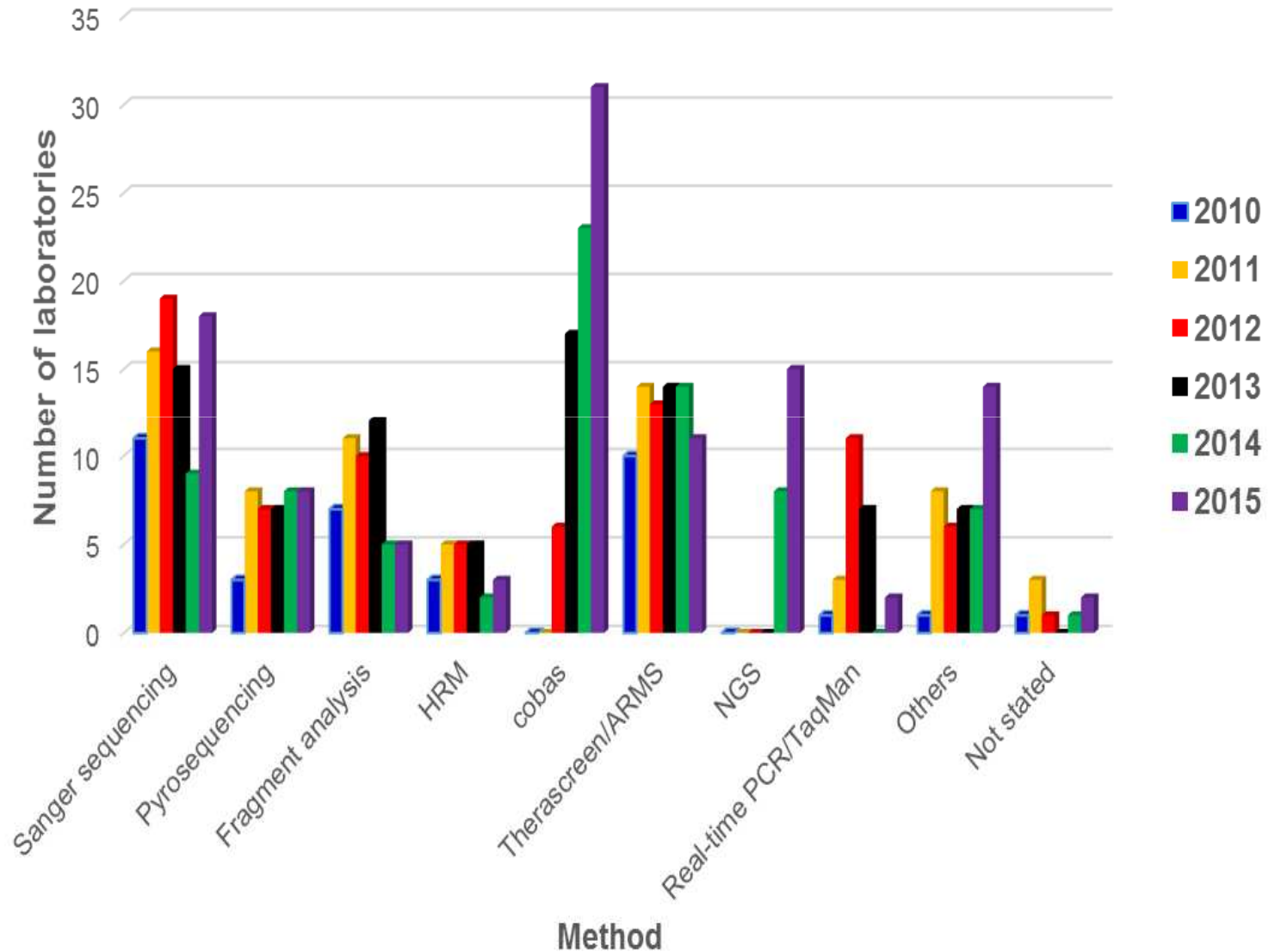
External Quality Assessment (WHO definition)

A system for objectively checking the laboratory's performance using an external agency or facility

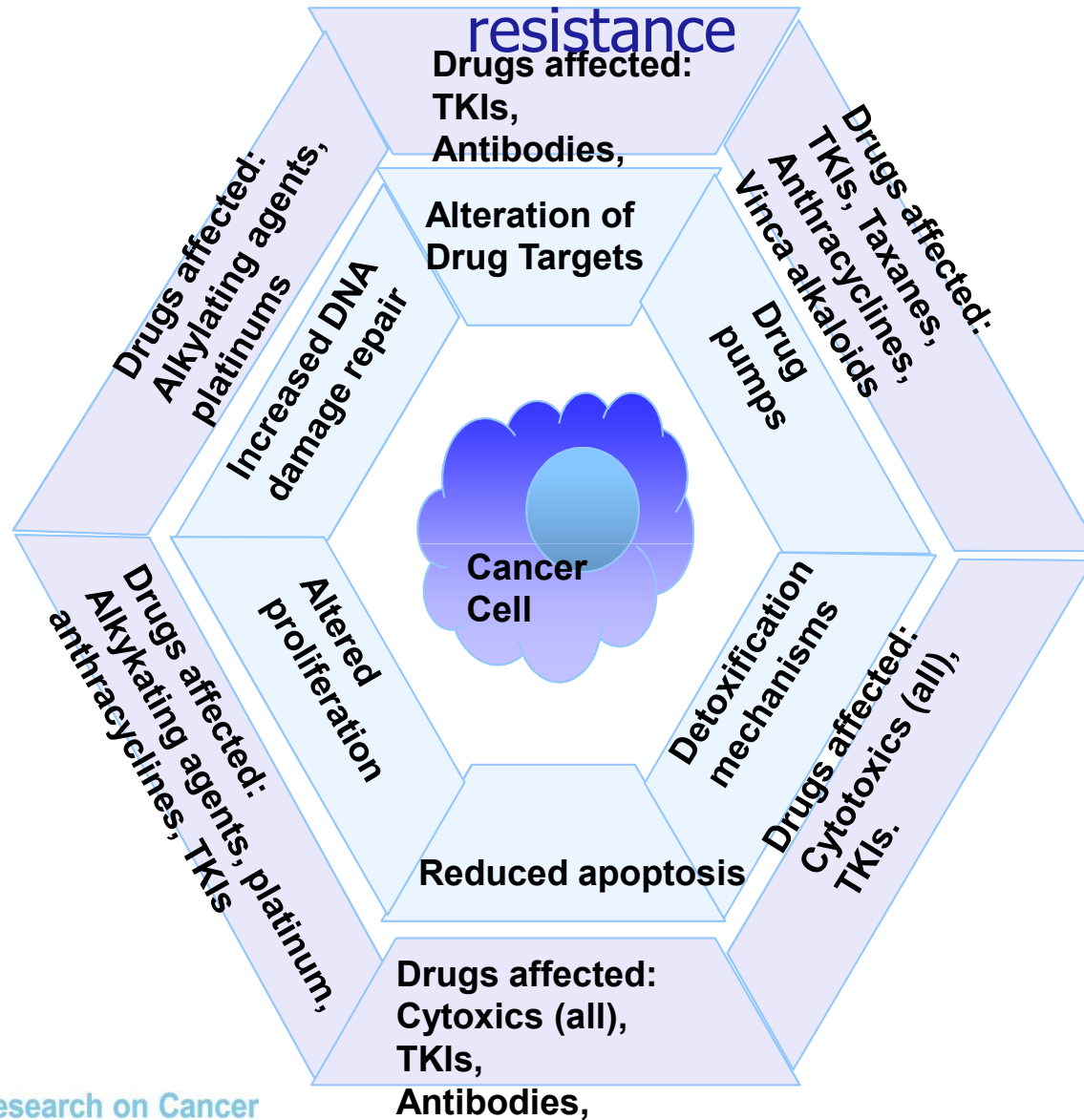


Molecular analysis of lung cancer EQA

**EGFR testing
methodologies**

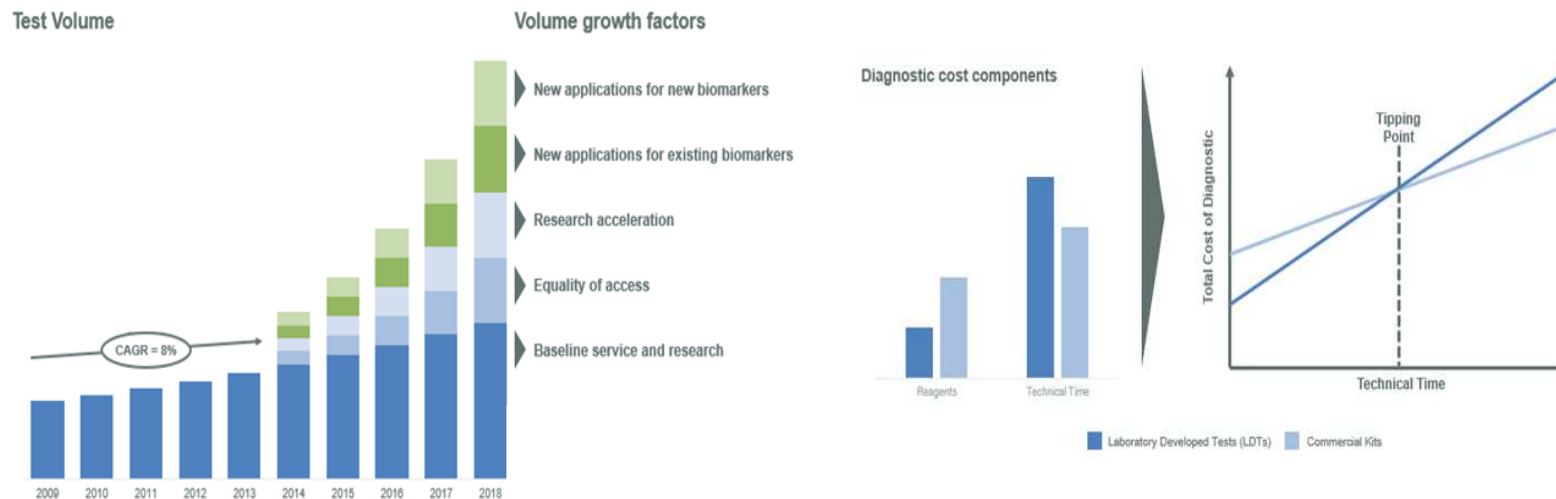


Molecular chess: Hallmarks of anti-cancer drug

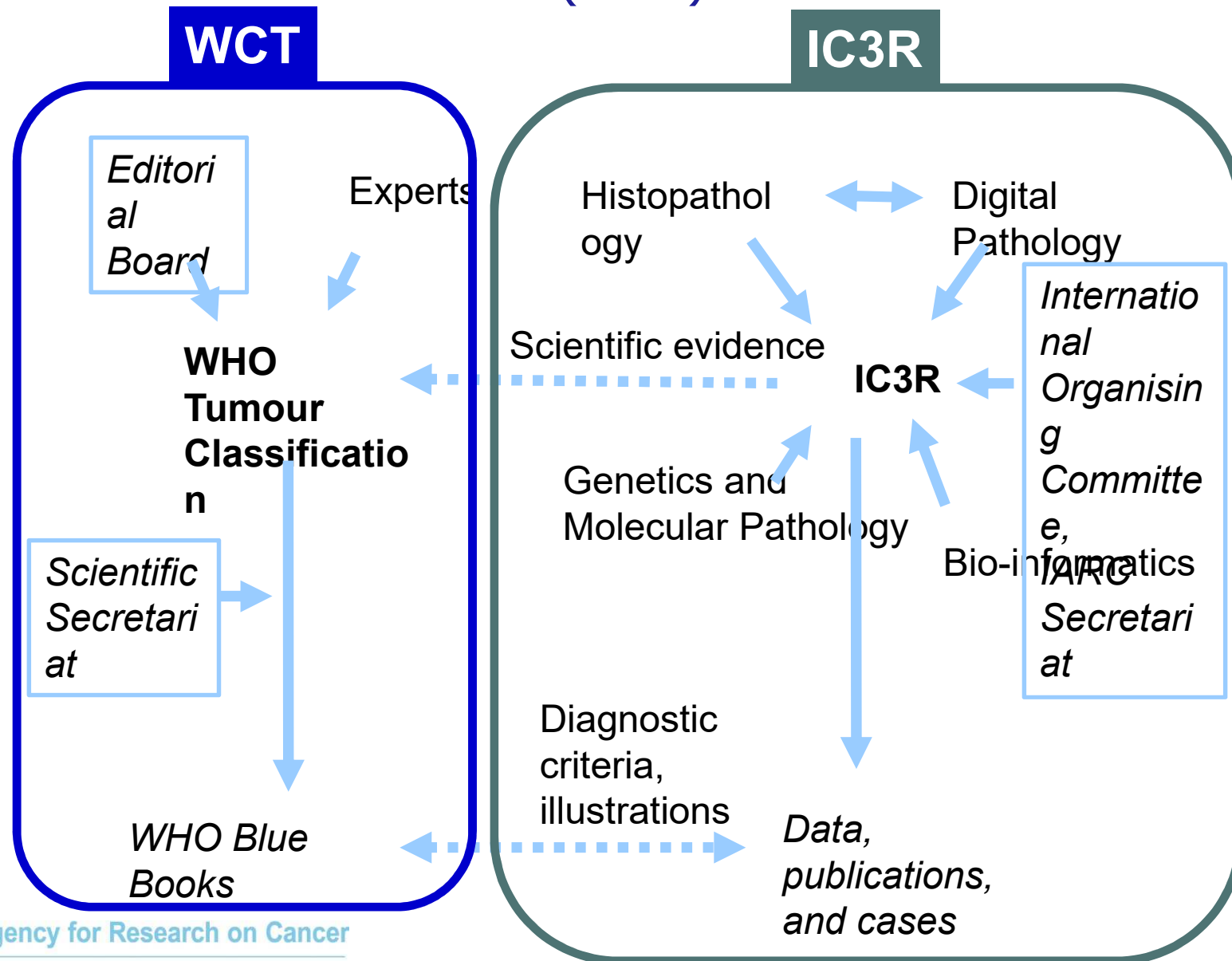


CMD ImPACT business planning tool

The tool is a simple Microsoft Excel programme that allows the costs and income to be modelled against a wide range of funding or demand scenarios, from current requirements to expansion of existing molecular diagnostic services.



Collaboration for Cancer Classification and Research (IC3R)

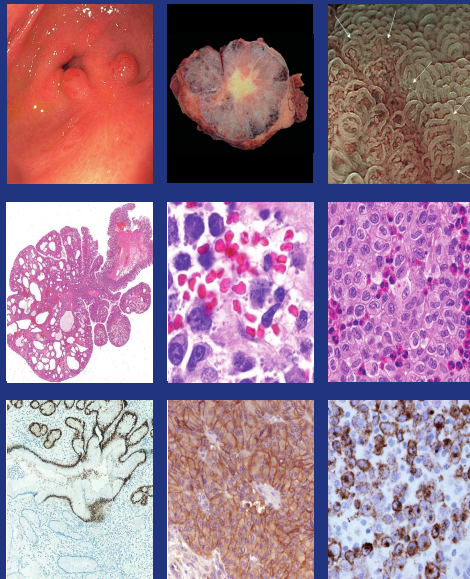


Proofs

WHO Classification of Tumours • 5th Edition

Digestive System Tumours

Edited by the WHO Classification of Tumours Editorial Board



2.0: Tumours of the oesophagus: Introduction

Lam
Cochlin
Otake R

AK
AO

This chapter describes benign and malignant oesophageal tumours of epithelial differentiation and the ICD-O-4 topographical coding for it.

Best.XI ICD-O-4 topographical coding for the anatomical site covered in this chapter

2.1.2.2: Oesophageal squamous dysplasia

Takubo KT
Fuji SF

Chapter 2

There are two main types of precursor lesions: Barrett dysplasia and squamous dysplasia. The latter is typically described in terms of the degree of dysplasia, ranging from low-grade to high-grade. The latter occur in the treatment of low-grade

Definition
Squamous dysplasia of the oesophagus is an unequivocal neoplastic alteration of the oesophageal squamous epithelium, without invasion.

ICD-O coding
807700 Low-grade squamous dysplasia
807702 High-grade squamous dysplasia

ICD-11 coding
2E32.0 & XH43Y37 Benign neoplasm of oesophagus & Oesophageal squamous intraepithelial neoplasia (dysplasia), low-grade
2E50.1 & XH4ND6 Carcinoma in situ of oesophagus & Oesophageal squamous intraepithelial neoplasia (dysplasia), high-grade

Related terminology
None

Subtype(s)
None

Localization
Squamous dysplasia can occur anywhere in the oesophagus, and it is likely to follow the distribution of squamous cell carcinoma.

Clinical features
Patients at high risk of oesophageal squamous cell carcinoma are usually followed using a combination of Lugol's chromoendoscopy and narrow-band imaging (3366). With Lugol's iodine, low-grade dysplasia appears as an unstained or weakly stained area, high-grade dysplasia is completely unstained (3074). Features associated with neoplastic disease include large size, non-flat appearance, positive pink-colour sign, and multiplicity of distinct iodine-unstained lesions (3702). On narrow-band

imaging, dysplastic lesions appear as areas of brownish discoloration (2250,2202). Abnormalities on narrow-band imaging reflect the invasion depth of intramucosal carcinomas and changes of intrapapillary capillary loops (2458).



Fig. 2.1XIX Natural age-standardized incidence rates of oesophageal squamous cell carcinoma (SCC).

4 Tumours of the oesophagus



Fig. 2.1XX Oesophageal squamous dysplasia. A On low-magnification endoscopy with narrow-band imaging, the lesion appears as a reddish-brown area. B On high-magnification endoscopy with narrow-band imaging, the lesion appears as a reddish-brown area. C On white-light endoscopy, the lesion appears as a reddish-brown area. It will disappear and be unstained.

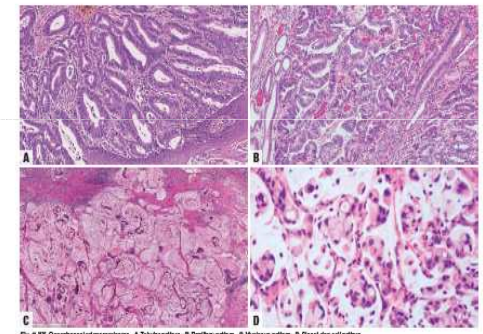


Fig. 2.1XXI Oesophageal adenocarcinoma. A Tubular pattern. B Papillary pattern. C Microcystic pattern. D Signet-ring cell pattern.

In recent years, next-generation sequencing techniques have given rise to global projects involving whole-genome sequencing of oesophageal adenocarcinoma (2568). These projects have revealed key gene pathways and mutations involved in pathogenesis (3327,307). Identified novel genes (818), and shown that the genomic landscapes of prechemotherapy and postchemotherapy samples of oesophageal adenocarcinoma are similar (2867). There are currently no clinical applications for these comprehensive but complex data, but clinically relevant and diagnostically useful prognostic and predictive markers may emerge in the future. Data from The Cancer Genome Atlas (TCGA) also suggest that oesophageal adenocarcinoma strongly resembles gastric carcinoma with chromosomal instability (2832).

Macroscopic appearance

Oesophageal adenocarcinomas often present in advanced stages and appear as stricture, polypoid, fungating, ulcerative, or diffuse infiltrating lesions. In earlier stages, adenocarcinomas may appear as irregular plaques. Early-stage carcinomas may present as small nodules or may not be detected on endoscopy. Adjacent to the carcinoma, there may be irregular tongues of reddish mucosa (resembling a salmon patch) that represent Barrett oesophagus and reflux changes and that contrast with the greyish-white colour of the squamous-lined oesophageal mucosa.

Histopathology

Oesophageal adenocarcinoma shows gastric, intestinal, and mixed (hybrid) lineage, evidenced by a combination of morphological and immunohistochemical features (1548,426).

The mucosa adjacent to the adenocarcinoma may show Barrett dysplasia and intestinal metaplasia (Barrett oesophagus). Oesophageal adenocarcinomas can be classified as having tubular, papillary, mucinous, and signet-ring cell patterns. Only limited evidence of the relevance of these patterns is available; therefore, patterns are described rather than subtypes. A mixture of these patterns is often seen. The tubular pattern is most common. It is characterized by irregular, single or anastomosing tubular glandular structures lined by a layer of single or stratified malignant epithelium; neoplastic glands often show variable amounts of intracellular mucin production and may show dilatation (1756). The papillary pattern is characterized by papillae, with rare cases showing micropapillary architecture (1162). The mucinous pattern generally shows carcinoma cells floating in

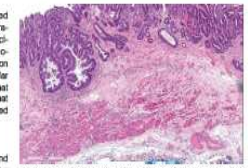


Fig. 2.1XXII Oesophageal adenocarcinoma. An example in Barrett oesophagus with a double layer of mucosal mucosa.

Conclusions

- Change is inevitable, but translation of research findings, and then implementation are difficult.
- In pathology, molecular and digital methods are entering practice.
- This has implications for the way departments are organised, patient pathways, and cost effectiveness.
- Quality control is essential, with clear standard operating procedures.
- The potential gains for patients are considerable!



Thank you!

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