

# *Developing RNA-based molecular diagnostics in the post-genomic era*

National Cancer Policy Forum

Policy Issues in the Development and Adoption of  
Molecularly Targeted Therapies for Cancer

11/10/2014

D. Neil Hayes, MD, MS, MPH

# Introduction

- Computational biologist / translational researcher (The Cancer Genome Atlas 10,000 transcriptomes)
- Physician and clinical trialist
- Not pathologist
- Co-Founder of diagnostics company (GeneCentric)

# Diagnostic Test Adoption

- Science
  - Evidence (hypothesis generation, validation)
  - Platform (tissue and technology)
- Regulatory (federal, state, accrediting bodies)
- Payment
- Practice (adoption in clinical practice)

# Science

## Evidence for RNA

# Mutation Detection

Published online 26 June 2014

Nucleic Acids Research, 2014 • e107  
doi: 10.1093/nar/gku489

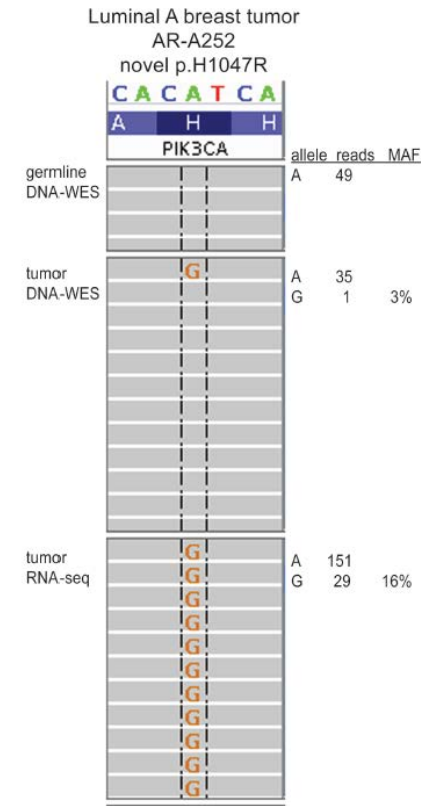
## Integrated RNA and DNA sequencing improves mutation detection in low purity tumors

Matthew D. Wilkerson<sup>1,2,\*</sup>, Christopher R. Cabanski<sup>1,3</sup>, Wei Sun<sup>2,4</sup>, Katherine A. Hoadley<sup>1,2</sup>, Vonn Walter<sup>1</sup>, Lisle E. Mose<sup>1</sup>, Melissa A. Troester<sup>1,5</sup>, Peter S. Hammerman<sup>6,7</sup>, Joel S. Parker<sup>1,2</sup>, Charles M. Perou<sup>1,2</sup> and D. Neil Hayes<sup>1,8,\*</sup>

<sup>1</sup>Lineberger Comprehensive Cancer Center, University of North Carolina at Chapel Hill, Chapel Hill, NC 27599, USA, <sup>2</sup>Department of Genetics, University of North Carolina at Chapel Hill, Chapel Hill, NC 27599, USA, <sup>3</sup>The Genome Institute at Washington University, St. Louis, MO 63108, USA, <sup>4</sup>Department of Biostatistics, University of North Carolina at Chapel Hill, Chapel Hill, NC 27599, USA, <sup>5</sup>Department of Epidemiology, University of North Carolina at Chapel Hill, Chapel Hill, NC 27599, USA, <sup>6</sup>Department of Medical Oncology, Dana-Farber Cancer Institute, Boston, MA 02215, USA, <sup>7</sup>Broad Institute of Harvard and MIT, Cambridge, MA 02142, USA and <sup>8</sup>Department of Internal Medicine, Division of Medical Oncology, Multidisciplinary Thoracic Oncology Program, University of North Carolina at Chapel Hill, Chapel Hill, NC 27599, USA

Received October 14, 2013; Revised April 22, 2014; Accepted May 15, 2014

- Somatic (cancer causing) alterations
- Driver versus passenger
- “Mutant expression”

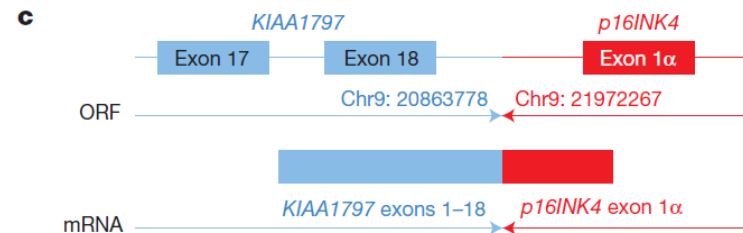
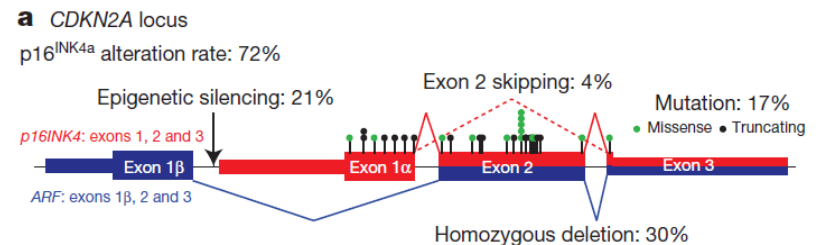


# Structural Alteration

## Detection of Structural Alterations

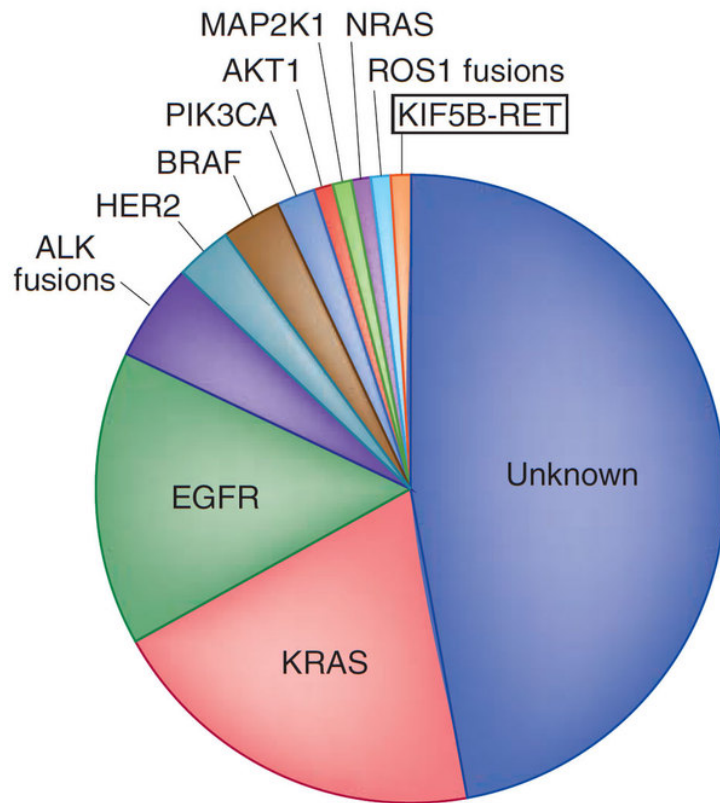
- Whole genome sequencing (expensive)
- Whole exome (limited)
- In situ hybridization (clinical assay, expensive and specific, "one at a time")
- RNA – cheap and all inclusive

## Structural Alterations of CDKN2A by RNA

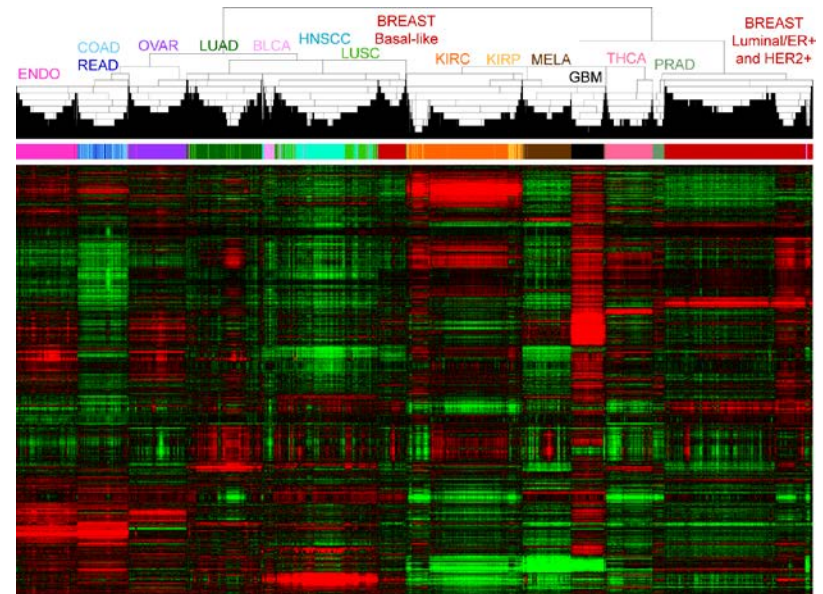


Comprehensive genomic characterization of squamous cell lung cancers. Nature. 2012

# Integrated classification beyond mutation RNA



DNA alterations



Immune System?

# Clinical Validation

- Hypotheses generated (validated) in convenience datasets
- Clinical validation needs to happen in clinical trials datasets
  - Largely absent or unavailable (\$)
  - Generation of new data prohibitive (\$\$\$)



# Science

## Platform



UNC  
LINEBERGER



UNC  
CANCER CARE

# Tissue Requirements and Quality: Lots of opinions, lots of experience, few published data

- % tumor
  - min and max
- Enrichment
  - macro, micro dissection
  - other
- Total amounts
  - Amplification
- Frozen vs paraffin
- Lower amounts and % tumor are useful for finding known variants (1% tumor) and signatures
- More tumor helps find new variants and signatures

# Research Frozen vs Clinical Paraffin

Zhao *et al.* *BMC Genomics* 2014, **15**:419  
http://www.biomedcentral.com/1471-2164/15/419



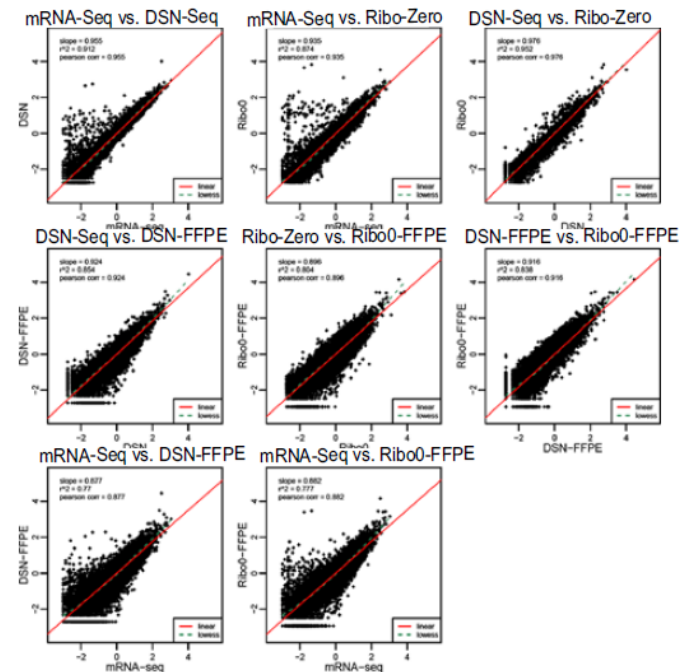
## RESEARCH ARTICLE

## Open Access

Comparison of RNA-Seq by poly (A) capture, ribosomal RNA depletion, and DNA microarray for expression profiling

Wei Zhao<sup>1,2</sup>, Xiaping He<sup>2,3</sup>, Katherine A Hoadley<sup>2,3</sup>, Joel S Parker<sup>2,3</sup>, David Neil Hayes<sup>3,5</sup> and Charles M Perou<sup>1,2,3,4\*</sup>

- Classical teaching
  - RNA degrades quickly
  - Assays on frozen tissue
- Recent experience
  - Intact 200-300 bp RNA fragments remain
  - Technologies targeting 300 bp robust

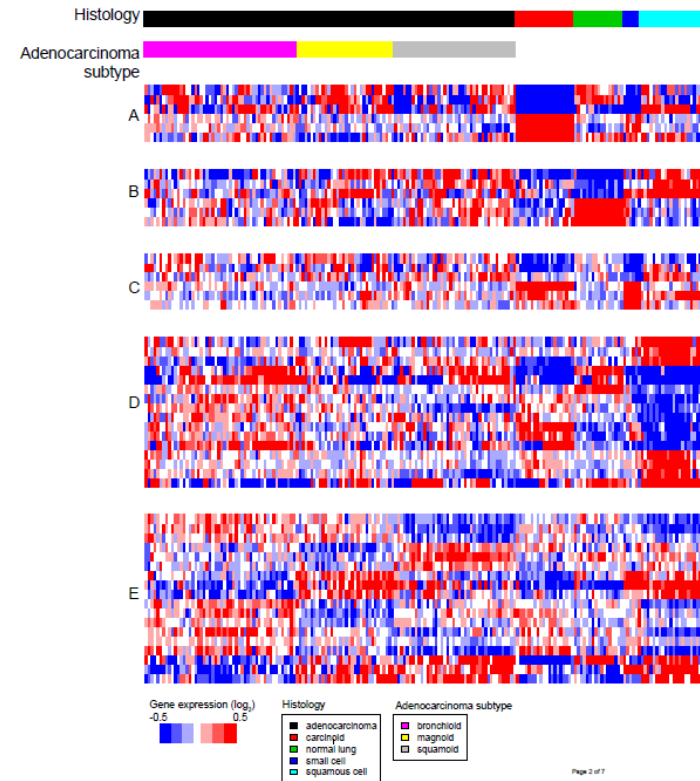
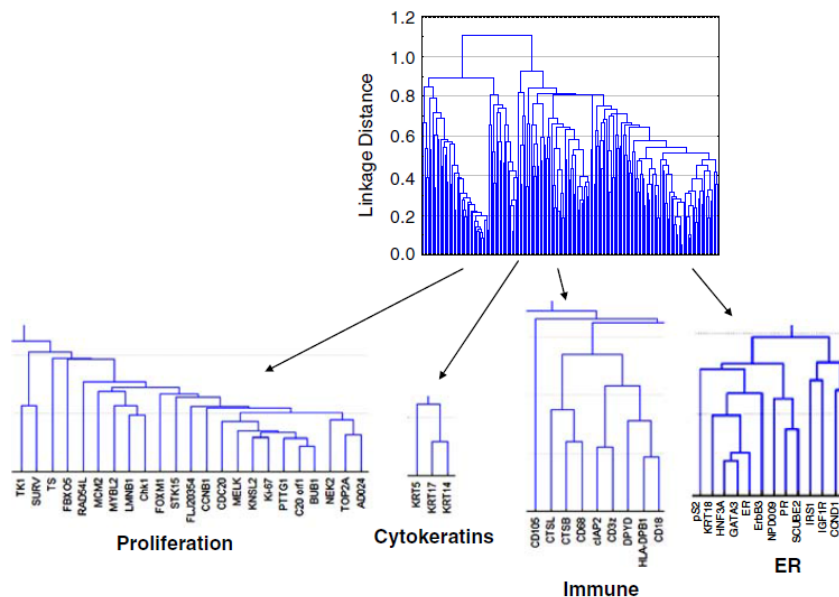


# Paraffin Diagnostics

Genomic Health: Oncotype DX,  
breast Cancer

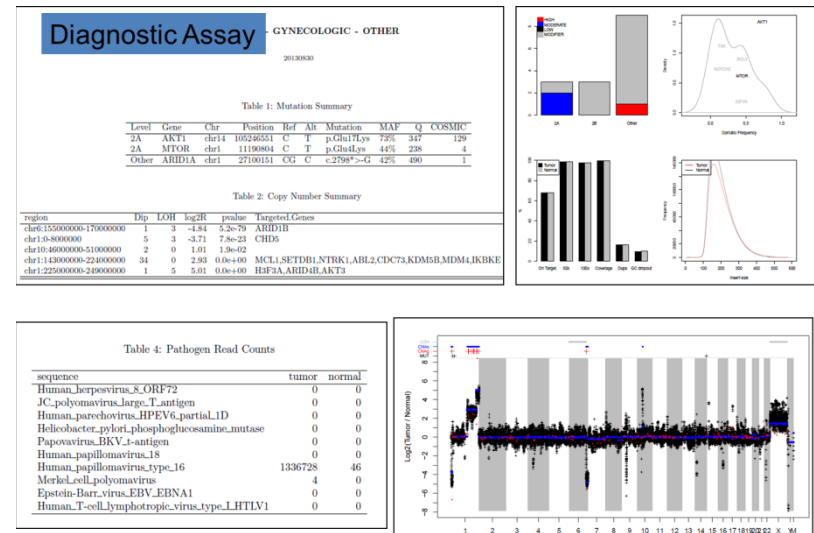
LabCorps: HistoPlus, lung  
Cancer

Figure S1. Lung marker genes (Bhattacharjee et al. cohort  $n=254$ )



# UNC Experience = UNCSeq: LCCC1108

- DNA and RNA assays (capture)
- 1400 patients
- 10 microns tissue (500-1000 ug)
- Variety sources
  - Biopsy (core)
  - Gross resection
- FNA (no quantification)



# Platforms (predicate instrument)

## Issues

- Regulatory clearance for RNA? Mostly no.
- Cost
- Throughput
- Availability to small and large diagnostics labs
- Bridging of “evidence” to commercial assay

## Examples

- Roche LightCycler or similar
- Roche Life Technologies sequencers
- Illumina sequencers
  - miSeq, HiSeq (multiple formats), NextSeq 500
- Nanostring (FDA)

# Regulatory

Tests need to be compliant  
Compliance is complicated and  
expensive

Diagnostic tests may often require  
private sector development



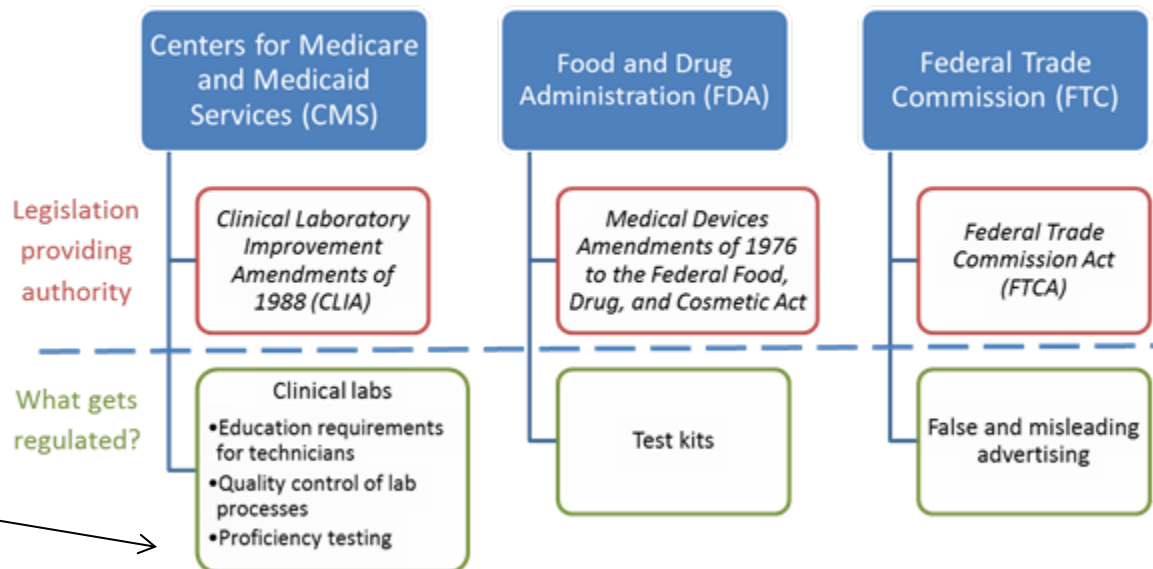
UNC  
LINEBERGER



UNC  
CANCER CARE

# Federal Regulation of Genetic Tests

LDT = Largely unregulated, although the lab itself is regulated



FDA regulates a test is determined by how it comes to market. A test may be marketed as a commercial test "kit," a group of reagents used in the processing of genetic samples that are packaged together and sold to multiple labs. More commonly, a test comes to market as a laboratory-developed test (LDT), where the test is developed and performed by a single laboratory, and where specimen samples are sent to that laboratory to be tested. The FDA regulates only tests sold as kits and, to date, has practiced "enforcement discretion" for LDTs.



# Take Home: 2 strategies

- LDT
  - Potentially cheap
    - IHC
    - Foundation One, Genomic Health
  - Regulatory status is unclear
- FDA
  - 510k – <\$10 million (but >any R01)
  - PMA - >\$10 million

# Laboratory-developed test (LDT)

## Intended

- Diagnostics (IVDs) manufactured. Developed, validated, and offered, within a single laboratory.
- Simple, well-understood pathology tests or
- Diagnosed rare diseases
- Used in a single institution as part of patient care
- Testing outside the institution would be prohibitive to patient care (due to timing between test need and result delivery)

## Actual

- Delivery often is by large corporations
- Test is not simple or well understood
- Disease are common (breast cancer)
- Use in patient care not always clear
- Test is not intended for a single institution but rather reference lab strategy where entire country sends test to the lab
- Common use of an LDT in place of an FDA approved assay
- Examples
  - Genomic health
  - Foundation medicine
  - Labcorp (many LDT's) including GeneCentric

# Payment

1. Cost of assay
2. Investment of development for private sector partners

# Intellectual Property Uncertainty

- “Mayo vs Prometheus”
- “Association for Molecular Pathology v. Myriad Genetics”
- Patent office struggling in light of these decisions, and by extension those wish to develop novel tests

# Test Reimbursement

- Medicare
  - “Medicare Has Stopped Paying Bills For Medical Diagnostic Tests. Patients Will Feel The Effects” *Forbes*. 3/27/2013
- State by State
- Insurer by insurer
- Self pay

# Practice

Adoption in clinical practice  
Impact on clinical workflow



UNC  
LINEBERGER



UNC  
CANCER CARE

# Changing Provider Behavior

- Difficult even when evidence suggests a superior test
- Cancer - multiple physicians involved
  - Subspecialists (biopsy)
  - Surgeons (biopsy and definitive surgery)
  - Med onc
    - User of diagnostic
    - Involved after biopsy / tissue processed

# Pathology Workflow

- Anatomic pathologist diagnosis of cancer have short timeline
- Special tests outside workflow
  - Send out LDT
  - Molecular tests in molecular path lab
  - Default IHC (even if test is inferior)
- Lack of coordination in information management



# Diagnostic Test Adoption

- Science
  - Evidence (hypothesis generation, validation)
  - Platform (tissue and technology)
- Regulatory (federal, state, accrediting bodies)
- Payment
- Practice (adoption in clinical practice)

# Carcinoma of Unknown Primary (CUP)

## Personal experience

- Challenging diagnosis
- Extensive IHC evaluation
- Multiple LDT RNA assays
  - bioTheranostics, Rosetta, others
  - Send out
- Frequently desired by physicians
- Never sent voluntarily by our path department
  - Lack of knowledge about the CUP assays
  - Discussed largely in negative

### Blinded Comparator Study of Immunohistochemical Analysis versus a 92-Gene Cancer Classifier in the Diagnosis of the Primary Site in Metastatic Tumors

Lawrence M. Weiss,<sup>\*</sup> Peiguo Chu,<sup>†</sup> Brock E. Schroeder,<sup>‡</sup> Veena Singh,<sup>§</sup> Yi Zhang,<sup>§</sup> Mark G. Erlander,<sup>§</sup> and Catherine A. Schnabel<sup>§</sup>

**Table 2** Sensitivity and Specificity of the 92-Gene Assay and IHC/Morphology Analysis at the Main Type Level and for the Colon/Appendix Subtype

Main type	No.	IHC/morphology analysis		92-gene assay	
		Sensitivity (%)	Specificity (%)	Sensitivity (%)	Specificity (%)
Melanoma	2	0 (0–66)	100 (97–100)	0 (0–66)	99 (95–100)
Sarcoma	7	86 (49–97)	98 (94–100)	100 (65–100)	98 (94–100)
Mesothelioma	1	0 (0–79)	100 (97–100)	100 (21–100)	99 (95–100)
Lung	24	67 (47–82)	94 (87–97)	75 (55–88)	95 (89–98)
Gynecologic	8	88 (53–98)	98 (94–100)	88 (53–98)	96 (91–99)
Gastrointestinal	26	92 (76–98)	93 (86–96)	92 (76–98)	97 (91–99)
Colon/appendix subtype	17	94 (73–99)	97 (92–99)	94 (73–99)	99 (95–100)
Urinary bladder	11	45 (21–72)	99 (95–100)	82 (52–95)	99 (95–100)
Kidney	13	77 (50–92)	100 (97–100)	77 (50–92)	99 (95–100)
Endocrine	9	56 (27–81)	99 (95–100)	56 (27–81)	99 (95–100)
Hepatocellular	1	100 (21–100)	100 (97–100)	100 (21–100)	100 (97–100)
Head and neck/esophageal squamous	3	67 (21–94)	98 (94–100)	67 (21–94)	97 (92–99)
Salivary gland	1	0 (0–79)	100 (97–100)	0 (0–79)	98 (93–99)
Prostate	4	50 (15–85)	100 (97–100)	100 (51–100)	100 (97–100)
Breast	11	55 (28–79)	97 (92–99)	73 (43–90)	100 (97–100)
Skin basal cell	1	0 (0–79)	100 (97–100)	0 (0–79)	100 (97–100)
Total main type	122	69 (60–76)	99 (98–99)	79 (71–85)	99 (98–99)

The 95% CIs are provided in parentheses.

The Journal of Molecular Diagnostics, Vol. 15, No. 2, March 2013