

International Symposium on Immunohistochemistry

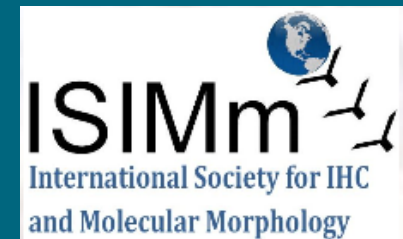
January 4th - 7th, 2018

Hosted by Dept. of Histopathology, Tata Medical Center, Kolkata, India

In collaboration with NordiQC, Alborg, Denmark and ISIMM, California, USA



The impact of proficiency testing on lab immunoassays



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Director of NordiQC
Aalborg University Hospital,
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Nordic Immunohistochemical Quality Control



Denmark



AALBORG



AALBORG UNIVERSITY HOSPITAL



NORTH DENMARK REGION

Budolfi Church



Aalborg Harbour Front

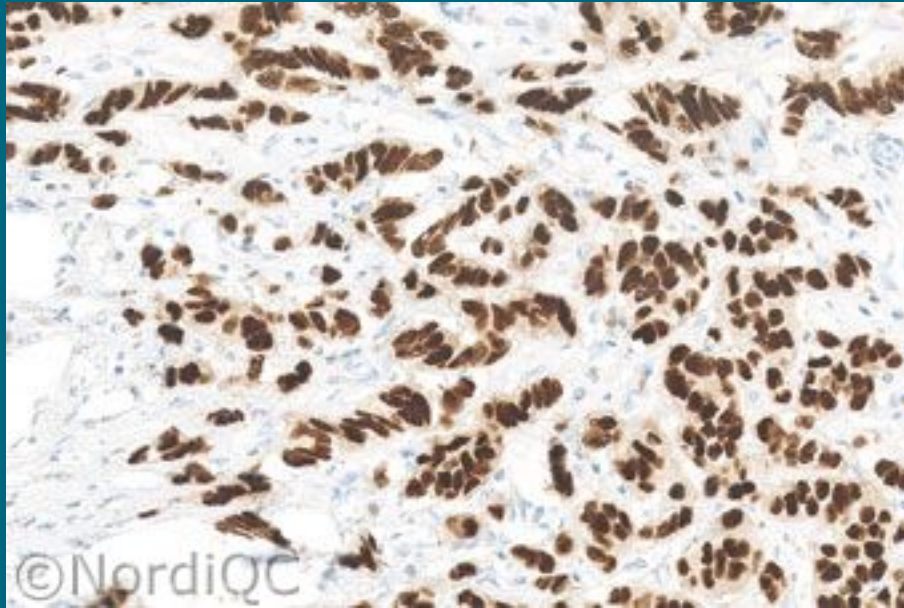


Aalborg House of Music

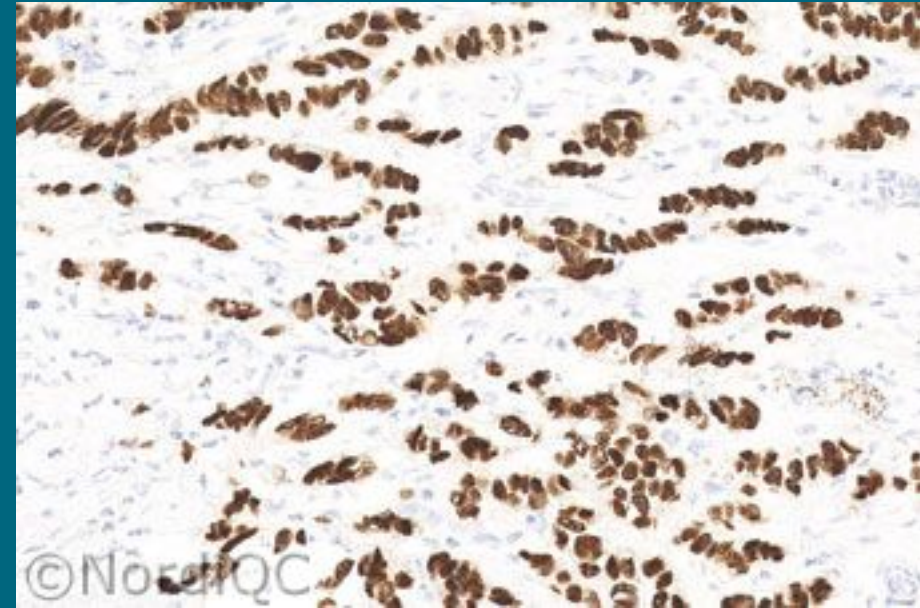




Serial sections stained for Estrogen receptor



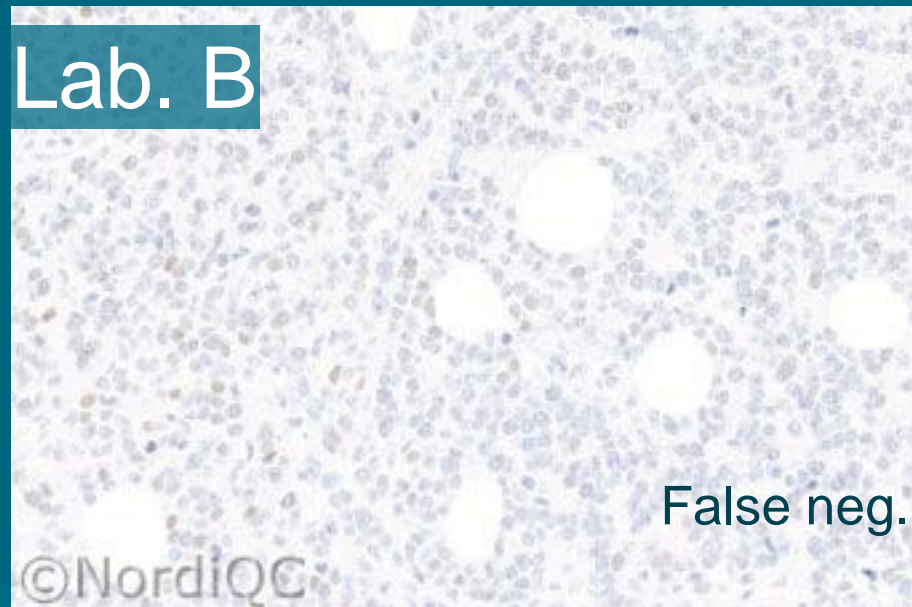
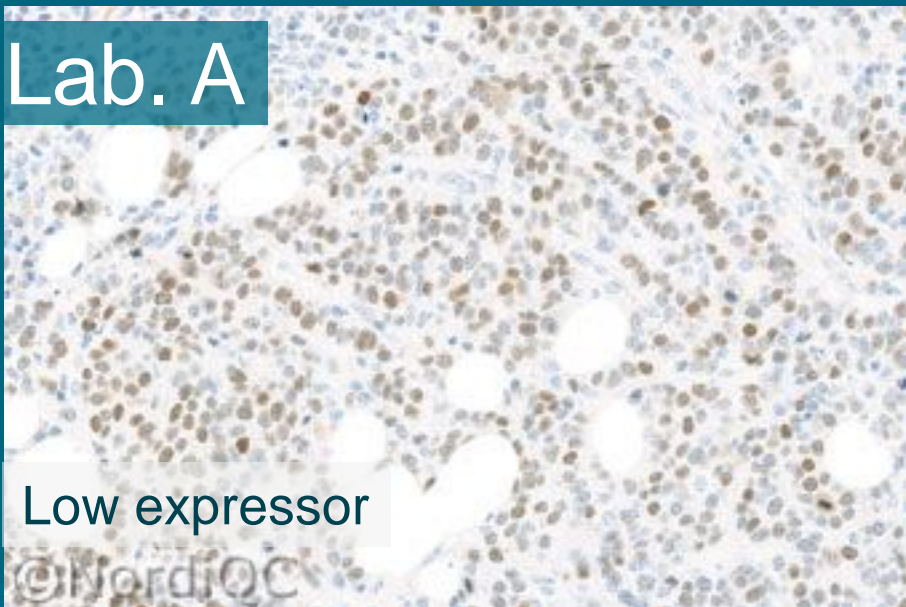
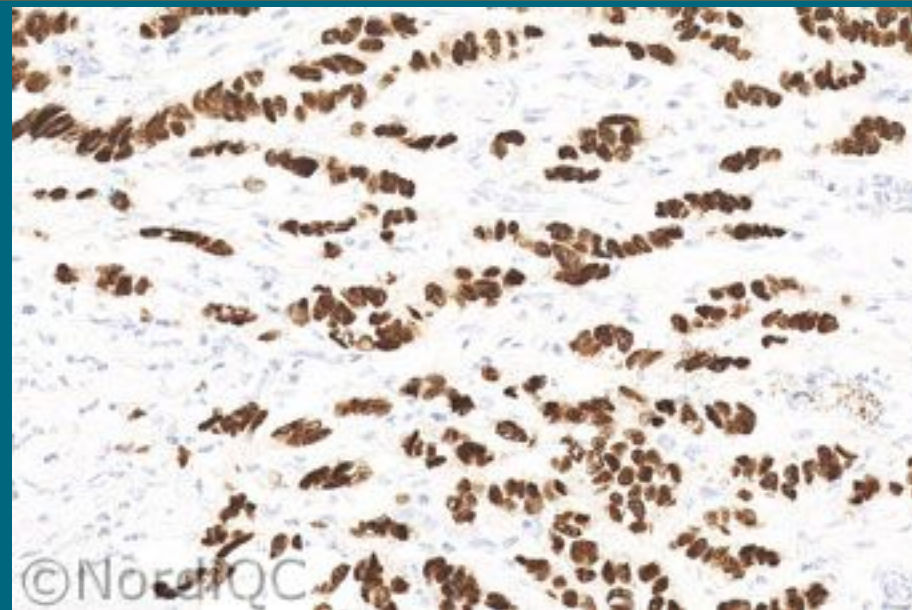
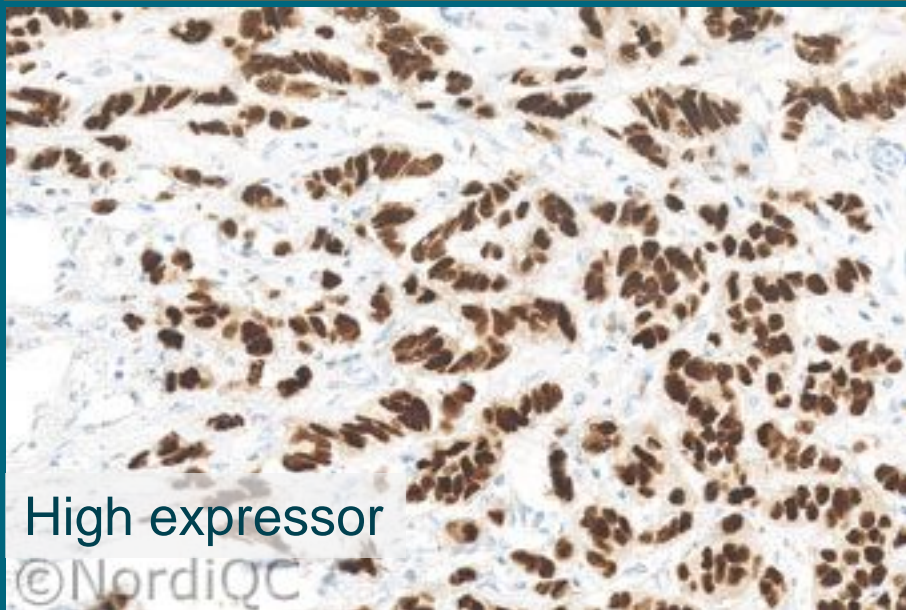
Lab. A



Lab. B

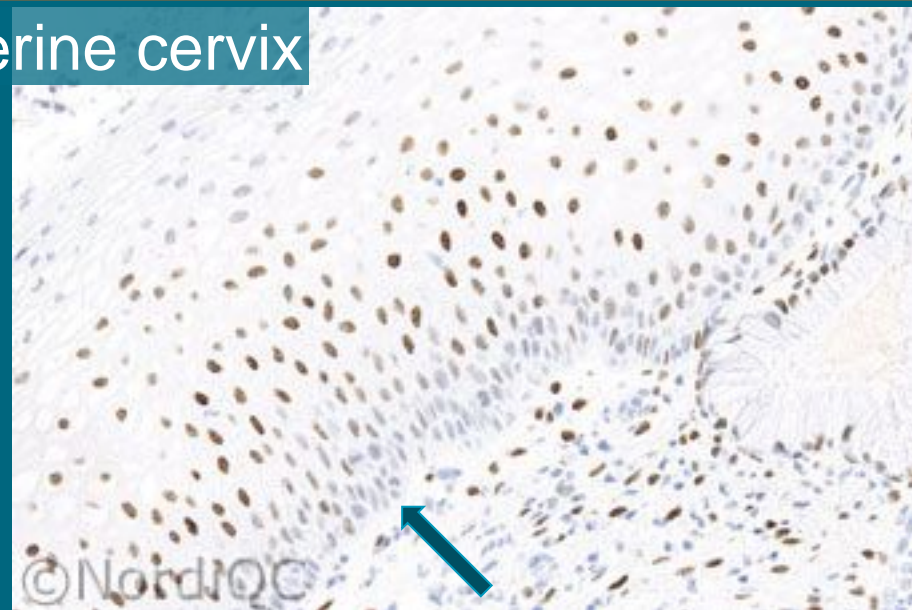
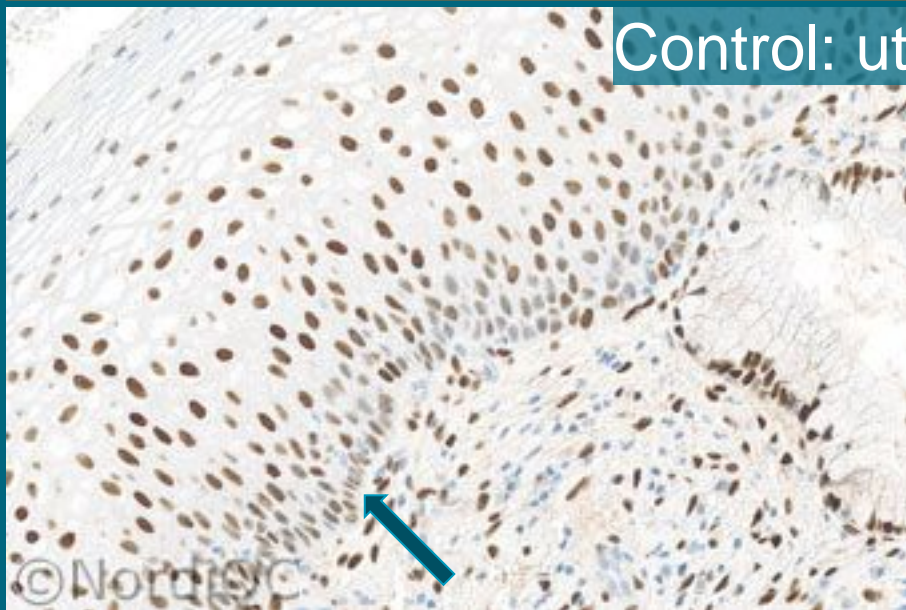
Optimally processed ductal breast carcinoma tissue

Serial sections stained for Estrogen receptor

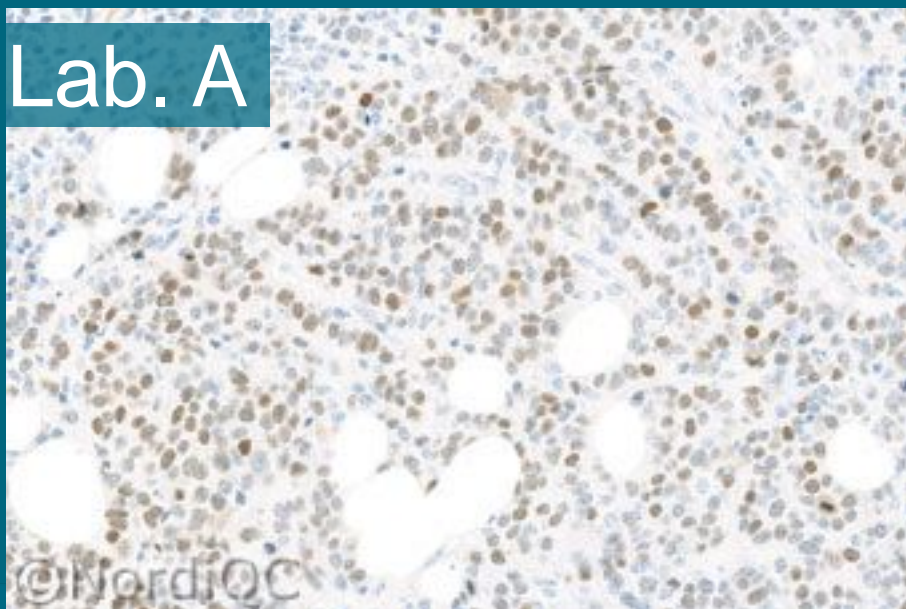


Serial sections stained for Estrogen receptor

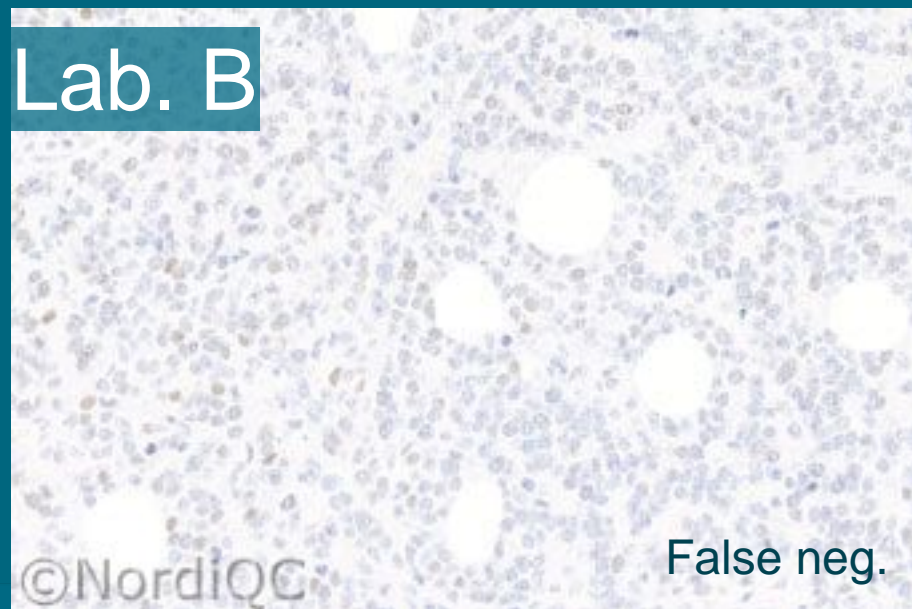
Control: uterine cervix



Lab. A

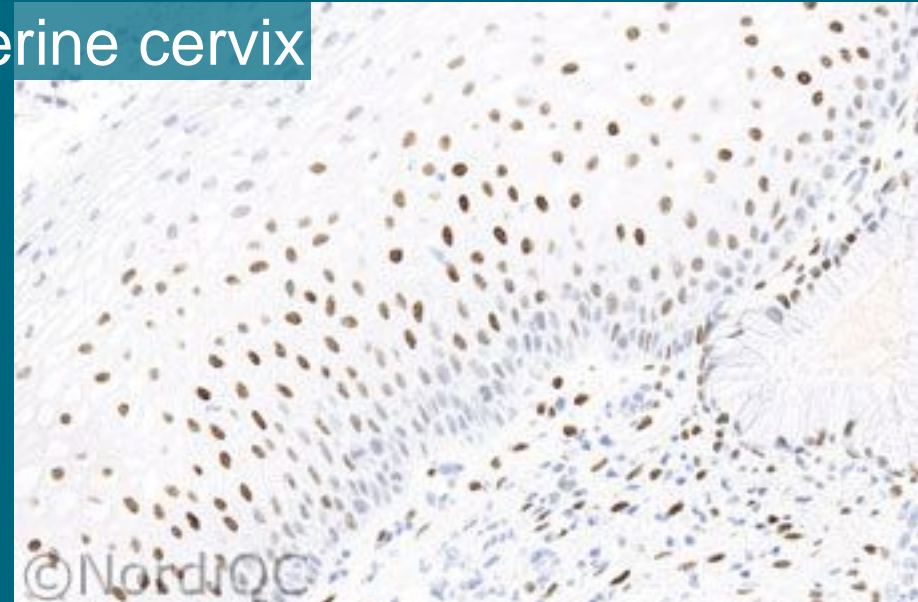
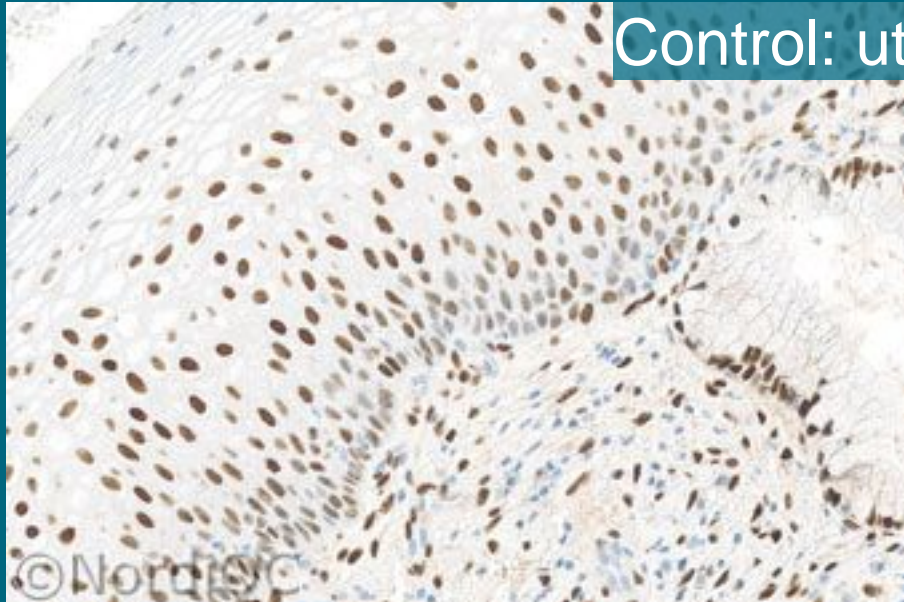


Lab. B

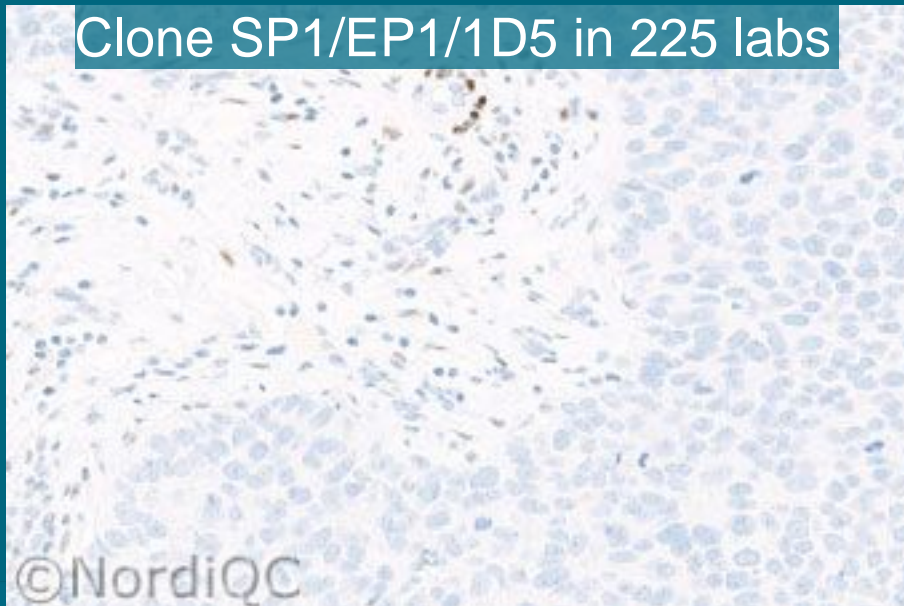


Serial sections stained for Estrogen receptor

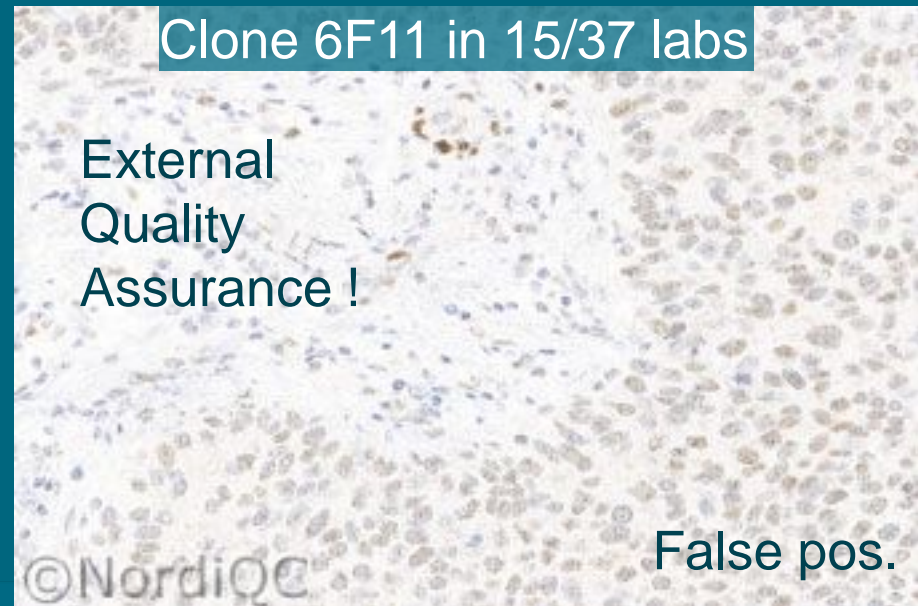
Control: uterine cervix



Clone SP1/EP1/1D5 in 225 labs

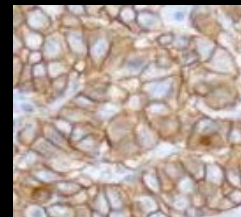
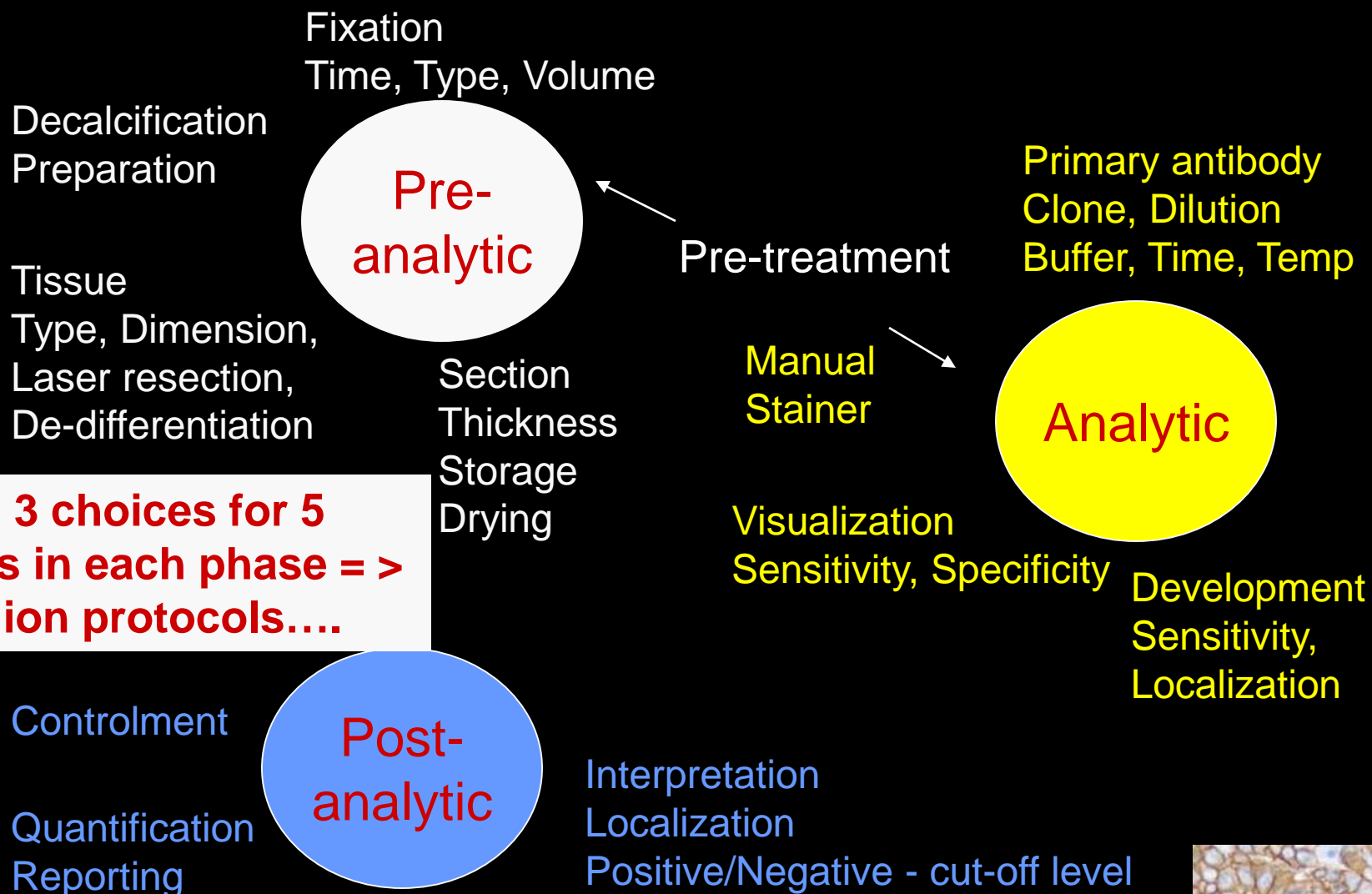


Clone 6F11 in 15/37 labs



External
Quality
Assurance !

False pos.



Suboptimal IHC assays may be due to:

- Preamanalytical issues
 - Fixation too short, too late, decalcification too soon...
- Analytical issues:
 - Less successful or too dilute antibody clones/RTUs
 - Insufficient epitope retrieval
 - Insensitive visualization systems
 - Platform problems
- Post-analytical issues
 - Interpretation criteria, interobserver variation ...

Should be identified
with proper external
on-slide controls



"Cold spots" – a problem with ~30% of all slides

AaS 16-400001



21.01.13 R
VIM



010116-5JM9

AaS 16-400001



21.01.14 R
VIM



010116-5JM9

AaS 16-400001



21.01.15 R
VIM



010116-5JM9

AaS 16-400001



21.01.16 R
VIM



010116-5JM9

AaS 16-400001



21.01.17 R
VIM



010116-5JM9

AaS 16-400001



21.01.18 R
VIM



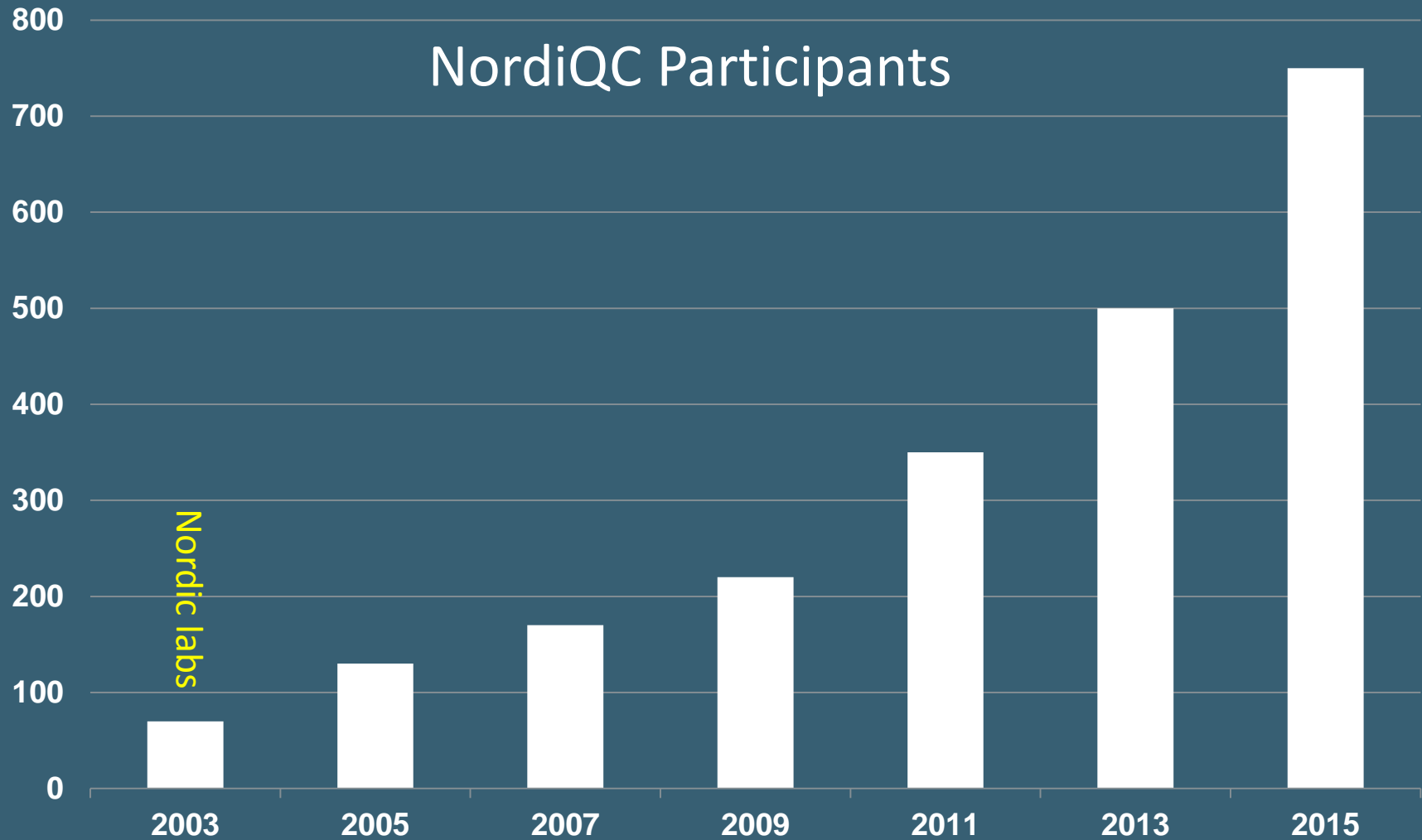
010116-5JM9

- International organization for **proficiency testing** of IHC
- Founded 2003 by Nordic pathologists
- Independent, scientific, not-for-profit organisation
- Institute of Pathology, Aalborg University Hospital, DK
- General module: 3 runs/year
 - 15-18 different marker challenges
- Breast cancer IHC module: 2 runs/y
 - HER-2, ER/PR, Ki67/E-Cad ...
- HER-2 ISH module: 2 runs/year
 - BRISH, FISH
- Companion module: 2 runs/year
 - PD-L1 ...



~ 90 IHC
markers
in
NordiQC
Runs
Tested
1-15
times

Alpha-methylacyl-CoA racemase	CyclinD1	MLH1
Alpha-smooth muscle actin	Cytokeratin 5	MSH2
Anaplastic lymphoma kinase	Cytokeratin 7	MSH6
B-cell specific activator protein	Cytokeratin 19	Multiple myeloma oncogene 1
bcl-2protein	Cytokeratin 20	Myosin, smooth muscle heavy chain
bcl-6protein	Cytokeratin, high molecular weight	Napsin A
Calretinin	Cytokeratin, low molecular weight	Neurofilament protein
Cancer antigen 125	Cytokeratin, pan-	Octamer transcription factor-3/4
Carcinoembryonic antigen	Desmin	p16 ^{ink4a}
CD3	Detected on GIST-1	p40
CD4	E-cadherin	p53
CD5	Epithelial cell adhesion molecule	p57
CD8	Epithelial membrane antigen	p63
CD10	Estrogen receptor alpha	Paired box gene-2 protein
CD14	Factor VIII related antigen	Paired box gene-8 protein
CD15	GATA3	Placental alkaline phosphatase
CD19	Glial fibrillary acidic protein	PMS2
CD20	Glypican 3	Podoplanin
CD23	Gross cystic disease fluid protein-15	Prostate specific acid phosphatase
CD30	HER-2	Prostate specific antigen
CD31	Hepatocyte antigen	Prostein
CD34	Human chorionic gonadotropin	Progesterone receptor
CD45	Immunoglobulin kappa	S-100 protein beta
CD56	Immunoglobulin lambda	Sal-like protein 4
CD68	Immunoglobulin M	SOX10
CD79a	Ki-67	Synaptophysin
CD99	Mammaglobin	Terminal deoxynucl. transferase
CD117	Melan-A	Vimentin
Chromogranin	Melanosoma specific antigen	Wilm's tumour-1 protein



Virchows Arch (2016) 468:19–29
DOI 10.1007/s00428-015-1829-1



ANNUAL REVIEW ISSUE

Proficiency testing in immunohistochemistry—experiences from Nordic Immunohistochemical Quality Control (NordiQc)

Mogens Vyberg^{1,2} · Søren Nielsen¹

[Free PMC Article](#)

Vyberg et al. *BMC Health Services Research* (2015) 15:352
DOI 10.1186/s12913-015-1018-6



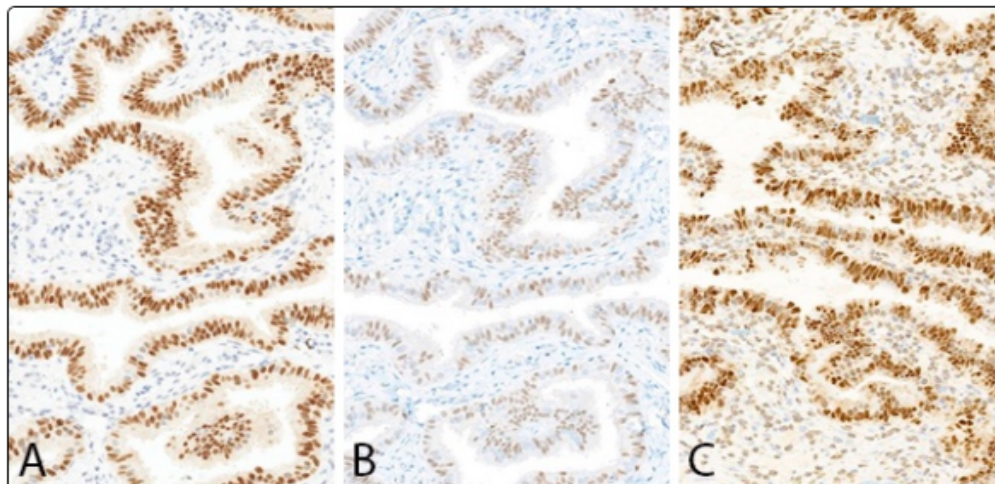
RESEARCH ARTICLE

Open Access

Immunohistochemical expression of HER2 in breast cancer: socioeconomic impact of inaccurate tests



Mogens Vyberg^{1*}, Søren Nielsen¹, Rasmus Røge¹, Beth Sheppard², Jim Ranger-Moore², Eric Walk², Juliane Gartemann³, Ulrich-Peter Rohr³ and Volker Teichgräber³



Serial sections of Fallopian tube stained for PAX8 in three laboratories. Lab A gets an optimal result, lab B a too weak staining with false negative reaction in ciliated cells, and lab C a too strong staining with false positive reaction in stromal cells. See the details in the PAX8 assessment, run 51.

Results - module 51, B24, H12, C2

15-Dec-2017

Individual results for the runs 51, B24 and H12 are now available (after logging in). Results from C2 will be available on the 14th January 2018. Click to see an overview of the results.

WWW.NORDIQC.ORG
FREE ACCESS

Events

[International Symposium on Immunohistochemistry](#)

4-7 Jan 2018: Tata Medical Center, Kolkata, India

[QulP/NordiQC Workshop in Applied Immunohistochemistry](#)

13-15 Jun 2018: Brugge, Belgium

[NordiQC Workshop in Diagnostic Immunohistochemistry](#)

18-20 Sep 2018: Aalborg, DK

Important dates

Run 52, H13, C3, B25

Protocol submission deadline

3 Jan 2018

Slide circulation

9 Jan 2018

Slide return deadline

13 Feb 2018

Publication of results

20 Apr 2018

Questions

Check out our [FAQ](#) (Frequently asked questions) or [contact us](#)

Google Custom Search

Search x

Collaborators



NordiQC assessment scheme 2018

Module	Winter	Spring	Autum
General	Run 52 <u>CR MSH6 SYP TdT</u> <u>VIM</u>	Run 53 <u>BSAP CGA ECAD</u> <u>OCT3/4 PMS2</u>	Run 54 <u>CD8 CEA CK-PAN</u> <u>GATA3 Podop</u>
Breast	Run B25 <u>ER HER2 IHC</u>		Run B26 <u>ER HER2 IHC PR</u>
HER2	Run H13 <u>HER2 ISH</u>		Run H14 <u>HER2 ISH</u>
Companion	Run C3 <u>PD-L1</u>		Run C4 <u>PD-L1</u>

Dates	Winter	Spring	Autum
Protocol submission opens	1 Dec 2017	12 Feb	1 Aug
Protocol submission closes	3 Jan	13 Mar	4 Sep
Shipping of slides	9 Jan	21 Mar	12 Sep
Deadline for slide return	13 Feb	1 May	11 Oct
Assessment General	6 Mar - 8 Mar	23 May - 25 May	24 Oct - 26 Oct
Assessment Breast	15 Mar - 16 Mar		8 Nov - 9 Nov
Assessment HER2	23 Mar		16 Nov
Assessment Companion	5 Apr		20 Nov
Publication of results	20 Apr	9 Jul	7 Dec

CD31

Characteristics

CD31 is a transmembrane glycoprotein, 130-140 kDa, also designated platelet-endothelium cell adhesion molecule = PECAM-1, belonging to the immunoglobulin super family. CD31 is ligand for CD38 and plays a role in thrombosis and angiogenesis. CD31 is strongly expressed in endothelial cells and weakly expressed in megakaryocytes, platelets, occasional plasma cells, lymphocytes (espc. marginal zone B-cells, peripheral T-cells) and neutrophils.

Neoplasms

CD31 is expressed in the vast majority of all types of vascular neoplasms, such as hemangioendothelioma, angiofibroma, hemangioma, and angiosarcoma. CD 31 is also expressed in most cases of Kaposi sarcoma and epithelioid hemangioendothelioma.

CD 31 may be expressed in a haematolymphoid neoplasms like chronic lymphatic leukaemia, plasmacytoma, histiocytosis and juvenile xanthogranuloma. When it comes to carcinomas, very few cases have been reported to express CD31. However, there are reports where singular cases of mucoepidermoid carcinoma, papillary thyroid carcinoma, sweat gland tumours, and metaplastic breast carcinoma with spindle cells have been reported to stain for CD31. There are also reports on CD 31-positivity in a minority of malignant fibrous histiocytoma of giant cell type and malignant mesothelioma. Finally CD31 has been detected in malignant gliomas, not only in the vessels but also the neoplastic cells.

Application

CD31 is most used in the panel for recognizing endothelial cell differentiation in tumours and should be considered a reliable marker for all types of vascular neoplasms.

Controls

Liver and tonsil are recommendable positive tissue controls for CD31. In liver an at least weak to moderate, distinct staining reaction in virtually all hepatic sinusoidal endothelial cells must be seen.

Endothelial cells of the portal tract vessels must show a moderate to strong staining reaction. In tonsil the majority of activated mantle zone B-cells must display an at least weak to moderate, distinct membranous staining reaction, whereas plasma cells and endothelial cells must show a strong staining reaction.

Appendix can be used as negative tissue control, as no staining reaction of the epithelial cells should be seen (only intraepithelial lymphocytes should be demonstrated).

Protocols

[Reommended protocols](#)

Assessments

[Run 46](#)

[Run 38](#)

[Run 32](#)

[Run 26](#)

[Run 11](#)

New protocol - H13 - HER2 ISH

Staining platform and assay

Hybridization target

Select

Select

Staining platform

Select

HER2/CHR17 dual BRISH

Select assay

Select

HER2 single BRISH

HER2/CHR17 FISH

Product lot no.

Heat Induced Epitope Retrieval (HIER)

HIER device

Select

Select

HIER buffer

Select

On Board / On Machine

HIER time (min.)

PT-link / PT-module

HIER temp (°C)

Microwave oven

Pressure Cooker

Water bath

Proteolysis

Enzyme

Select

Other

None

Proteolysis time (min.)

Proteolysis temp (°C)

Denaturation and hybridization of HER2 probe

Denaturation time (min.)

Denaturation temp (°C)

Hybridization time (min.)

Hybridization temp (°C)

Comment

Comments to protocol

Create

Back

Modify protocol ID 635, CDX2, run 48

Staining platform

Staining platform Ventana Benchmark Ultra ▾

Primary antibody

Primary antibody clone Cell Marque (235-Rxx) - EPR2764Y ▾

Lot number 1523802K

Dilution factor : 1:400 400

Diluent buffer Dako - Antibody Diluent (K8006) ▾

Incubation time (minutes) 32

Incubation temperature (Celcius) 36

Epitope Retrieval, HIER

Epitope retrieval, HIER ☒ YES ☐ NO

Device On Board / On Machine ▾

HIER buffer Ventana - Ultra CC1 (950-224) ▾

Efficient Heating Time (minutes) 48

Max. heating temperature (Celcius) 99

Epitope Retrieval, proteolysis

Epitope retrieval, proteolysis ☐ YES ☒ NO

Visualization system

Visualization system OptiView DAB IHC Detection Kit - 760-700 ▾

Amplification None ▾



















Incubation time linker (minutes) 8

Incubation time polymer (minutes) 8

Incubation temperature (Celcius) 36

Participant site

Protocol submission

Run ▲	Module	Epitope ▾	Protocol status ▾	Slide received by NordiQC ▾	Action ▾
49	General Module	CD5	✓	2017-02-13	 
49	General Module	CK-LMW	✓	2017-02-13	 
49	General Module	MLA	✓	2017-02-13	 
49	General Module	MLH1	✓	2017-02-13	 
49	General Module	NKX3.1	✓	2017-02-13	 
49	General Module	PSA	✓	2017-02-13	 
B23	Breast Cancer Module	ER	✓	2017-02-13	 
B23	Breast Cancer Module	HER2 IHC	✓	2017-02-13	 
C1	Companion Diagnostic Module	PD-L1	✓		 

Module status

49	General Module	Slides sent
B23	Breast Cancer Module	Slides sent
H11	HER2-ISH Module	Slides sent
C1	Companion Diagnostic Module	Slides sent

Open Homepage open for protocol submission. New protocols can be created, edited and deleted.

Closed Homepage closed for new protocol submission. Protocols already submitted can be edited. NordiQC are preparing to send slides.

Slides sent Slides for the submitted protocols have been sent to participants. Only protocol corrections are allowed.

Multi-tissue FFPE blocks

10% NBF 24-48 h (ASCO/CAP guidelines ...)

- Normal and clinically relevant tumour tissues
- Different levels of antigen expression
 - high, moderate, low, none

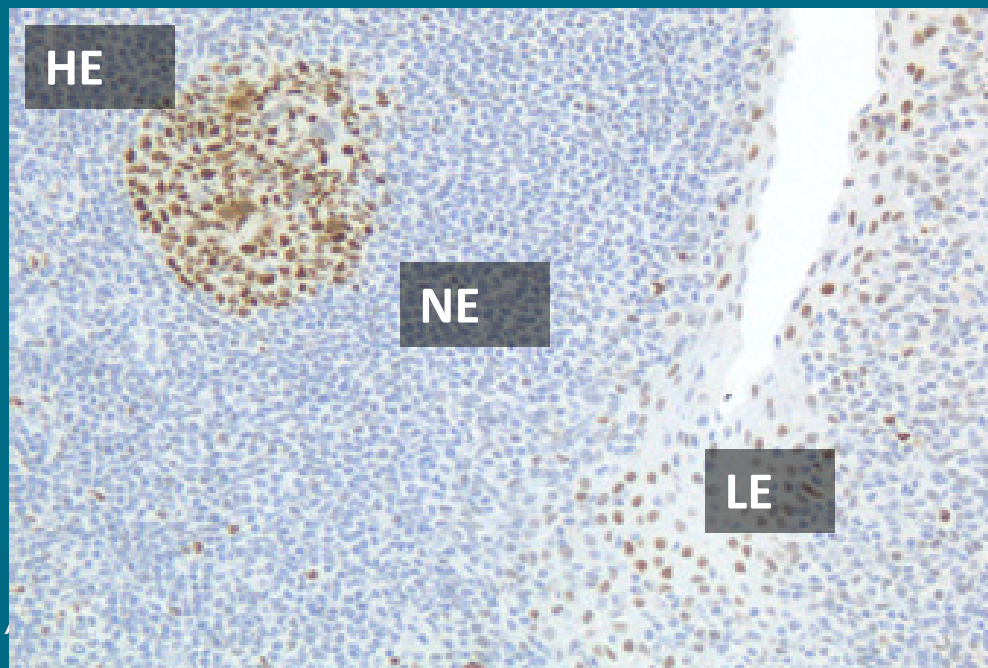


2 unstained slides for each marker send to the participants

1 stained slide returned for central assessment

The slide to be stained for **Bcl-6** comprised:

- 1-2. Tonsils, 24 h/48 h
3. Follicular lymphoma, grade I
4. Follicular lymphoma, grade II
5. Diffuse large B-cell lymphoma



Tissue selection:

- **High Expressor**
 - to confirm antibody
- **Low Expressor**
 - to ensure sensitivity
- **No-Expressor**
 - to ensure specificity

Material

The slide to be stained for [Bcl-6](#) comprised:

1. Tonsil, 24h fixation, 2. Tonsil, 48h fixation*, 3. Follicular lymphoma grade I,
4. Follicular lymphoma grade II, 5. Diffuse large B-cell lymphoma, non-Germinal Centre B-cell type (DLBCL non-GCB), 6. DLBCL, GCB.

All tissues were fixed in 10% neutral buffered formalin.

Criteria for assessing a Bcl-6 staining as optimal included:

- A moderate to strong distinct nuclear staining reaction of virtually all normal germinal centre B-cells in the tonsil fixed 24h.
- An at least weak to moderate distinct nuclear staining reaction of the majority of the squamous epithelial cells in the tonsil fixed 24h.
- A moderate to strong distinct nuclear staining reaction of the neoplastic cells in the two follicular lymphomas.
- An at least weak to moderate nuclear staining reaction of the majority of the neoplastic cells in the DLBCL, GCB subtype, tissue core no. 6.
- No or only a nuclear staining reaction in dispersed neoplastic cells of the DLBCL, non-GCB subtype, tissue core no. 5

* The tonsil fixed for 48h (tissue core no 2) was excluded from the assessment due to an aberrant inconsistent staining reaction in the circulated material.

Participation

Number of laboratories registered for Bcl-6, run 42	244
Number of laboratories returning slides	228 (93%)

Results

228 laboratories participated in this assessment. Of these, 168 (74%) achieved a sufficient mark (optimal or good). Table 1 summarizes the antibodies (Abs) used and assessment marks (see page 2).

The most frequent causes of insufficient staining reactions were:

- too low concentration of the primary antibody
- less successful performance of the mAb clone PG-B6p
- use of low sensitivity detection systems

Performance history

This was the third NordiQC assessment of Bcl-6. An increased pass rate was seen compared to the two previous runs 17, 2006 and 28, 2010 (see table 2).

Table 2. Proportion of sufficient results for Bcl-6 in the three NordiQC runs performed

	Run 17 2006	Run 28 2010	Run 42 2014
Participants, n=	69	132	228
Sufficient results	42%	48%	74%

Conclusion

The mAbs clones GI191E/A8, LN22 and PG-B6p could all be used to produce optimal staining results for Bcl-6. Irrespective of the clone applied, efficient HIER in alkaline buffer, use of a high sensitive detection system and careful calibration of the primary antibody were the most important prerequisites for an optimal staining result. The mAb clone GI191E/A8 and LN22, which both had a detection sensitivity of 69% and 66% respectively, were used by 16 of the 228 participating laboratories. The mAb clone PG-B6p, which had a detection sensitivity of 44%, was used by 10 of the 228 participating laboratories.

Table 1. Antibodies and assessment marks for Bcl-6, run 42

Concentrated antibodies	n	Vendor	Optimal	Good	Borderline	Poor	Suff. ¹	Suff. OPS ²
mAb clone GI191E/A8	13	Cell Marque 1 Immunologic 1 Zytomed	6	8	0	1	93%	100%
mAb clone LN22	38	Leica/Novocastra 2 DBS 1 Biocare 1 BioGenex 1 Zeta Corporation	20	16	4	3	84%	100%
mAb clone PG-B6p	43	Dako 1 DBS 1 Thermo/Neomarkers	9	22	11	3	69%	86%
Ready-To-Use antibodies								
mAb clone GI191E/A8 760-4241	59	Ventana/Cell Marque	24	25	9	1	83%	84%
mAb clone GI191E/A8 227M-9x	1	Cell Marque	0	0	1	0	-	-
mAb clone LN22 PA0204	10	Leica/Novocastra	3	7	0	0	100%	100%
mAb clone LN22 PM410	1	Biocare	1	0	0	0	-	-
mAb clone LN22 MAD-00638QD	1	Master Diagnostica	0	1	0	0	-	-
mAb clone PG-B6p IR/IS625	44	Dako	4	17	21	2	48%	75%
mAb clone PG-B6p GA625	7	Dako	2	2	3	0	57%	75%
mAb PG-B6p MAD-004023QD	2	Master Diagnostica	0	1	1	0	-	-
Total	228		69	99	50	10	-	
Proportion			30%	44%	22%	4%	74%	

1) Proportion of sufficient stains (optimal or good)

2) Proportion of sufficient stains with optimal protocol settings only, see below.

Detailed analysis of Bcl-6, Run 42

The following protocol parameters were central to obtain optimal staining:

Concentrated antibodies

mAb clone GI191E/A8: Protocols with optimal results were all based on HIER using Cell Conditioning Solution 1 (CC1; Ventana) (6/14)* as retrieval buffer. The mAb was typically diluted in the range of 1:50-1:100 depending on the total sensitivity of the protocol employed. Using these protocol settings 9 of 9 (100%) laboratories produced a sufficient staining result (optimal or good).

* (number of optimal results/number of laboratories using this HIER buffer)

mAb clone LN22: Protocols with optimal results were all based on HIER using Target Retrieval Solution 2 (TRS) pH 9 (3-in-1) (Dako) (2/2), CC1 (Ventana) (9/18) or Epitope Retrieval Solution 2 (BERS2; Ventana) (9/11) as retrieval buffer. The mAb was typically diluted in the range of 1:25-1:200 depending on the sensitivity of the protocol employed. Using these protocol settings 27 of 27 (100%) laboratories produced a sufficient staining result.

PDF file e-mailed to participants with assessment marks and – when needed – explanations and recommendations



Assessment of ER, B24 - individual results
Aalborg Sygehus (101)

Epitope ER
Assessment Optimal

NordIQC has assessed the submitted slides. In general, the assessment is based on staining intensity and distribution in cells expected to be demonstrated, background staining, cross-reactivity, quality of counterstaining and preservation of tissue morphology. Specific criteria for each epitope are described on <http://www.nordiqc.org/epitope.php>.

Each slide was marked as optimal, good, borderline or poor based on the following criteria:

Optimal: The staining reaction is considered perfect or close to perfect in all of the included tissues.

Good: The staining reaction is considered acceptable in all of the included tissues. However, the protocol settings may be optimized to ensure improved sensitivity or higher signal-to-noise ratio.

Borderline: The staining reaction is considered insufficient because of a generally too weak staining reaction, false negative or false positive staining reaction of one of the included tissues. The protocol should be optimized.

Poor: The staining reaction is considered insufficient because of, e.g., false negative or false positive staining reactions of several of the included tissues. An optimization of the protocol is urgently needed.

Moderate or strong cross reaction (due to the character of the primary antibody) or other false positive staining reaction (e.g. due to endogenous biotin) is not compatible with an optimal result and will usually cause downgrading.

For stains assessed as borderline or poor, comments and recommendations are given to the protocols. Good stains may also be accompanied by comments if specific problems are identified.

Recommended protocols from each staining platform are available at the NordIQC homepage (<http://www.nordiqc.org/recommended.php>) for comparison. Implementation of NordIQC recommended protocols as well as changes suggested in this letter must be tested carefully in your own laboratory before implementation into diagnostic work. NordIQC do not take any responsibility for consequences of changes in protocols or methods in your laboratory.

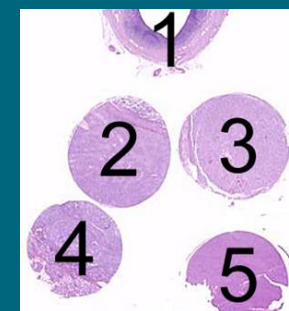
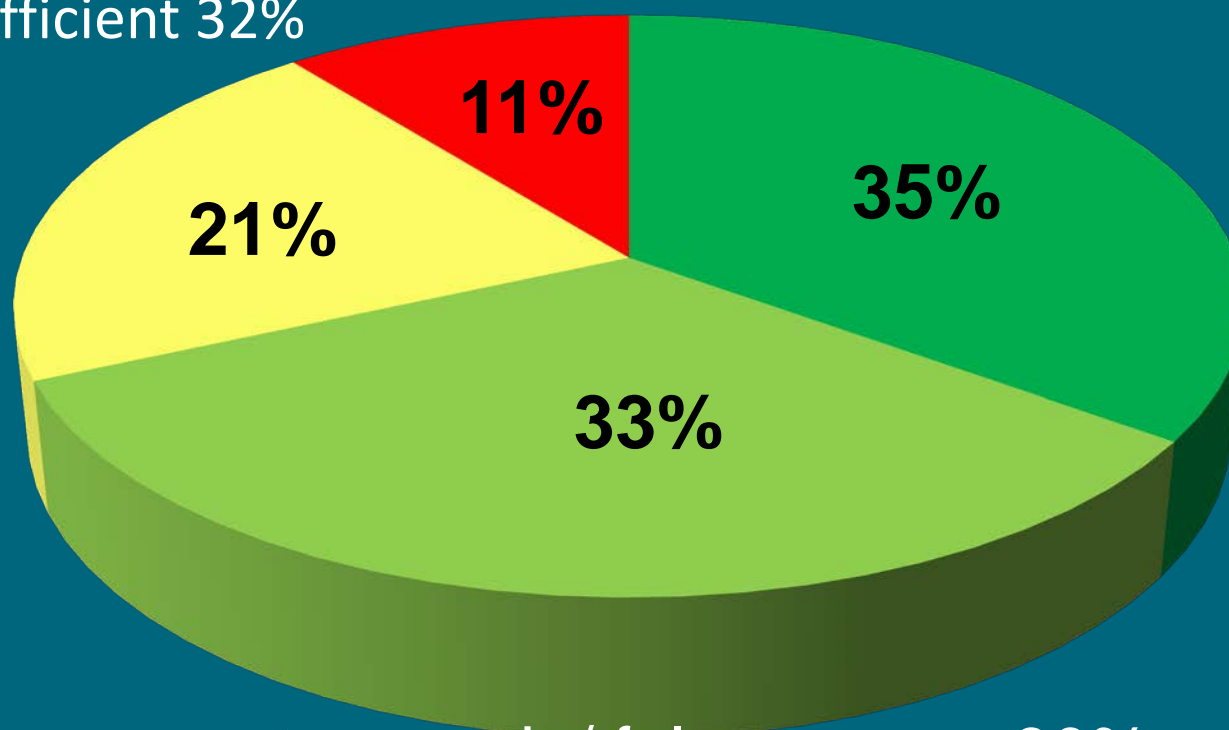
Marker	CD23	CR	CyD1	Ki67	Podop	TTF1
Assessment:	Poor	Optimal	Optimal	Good	Good	Borderline
Comments to the protocol:	False negative	-	-	Excessive counterstain	Weak	Weak
Suggestions for improvement:	Consider change of primary Ab and recalibrate	-	-	-	-	Increase primary Ab conc. and/or prolong HIER

Original stain

* Please read the epitope description and assessment summary carefully, as the choice of the Ab clone will influence the sensitivity and specificity.

General module ~ 20,000 slides (~100.000 core sections)

Insufficient 32%

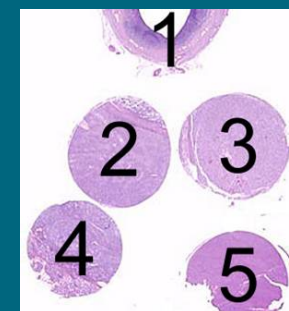
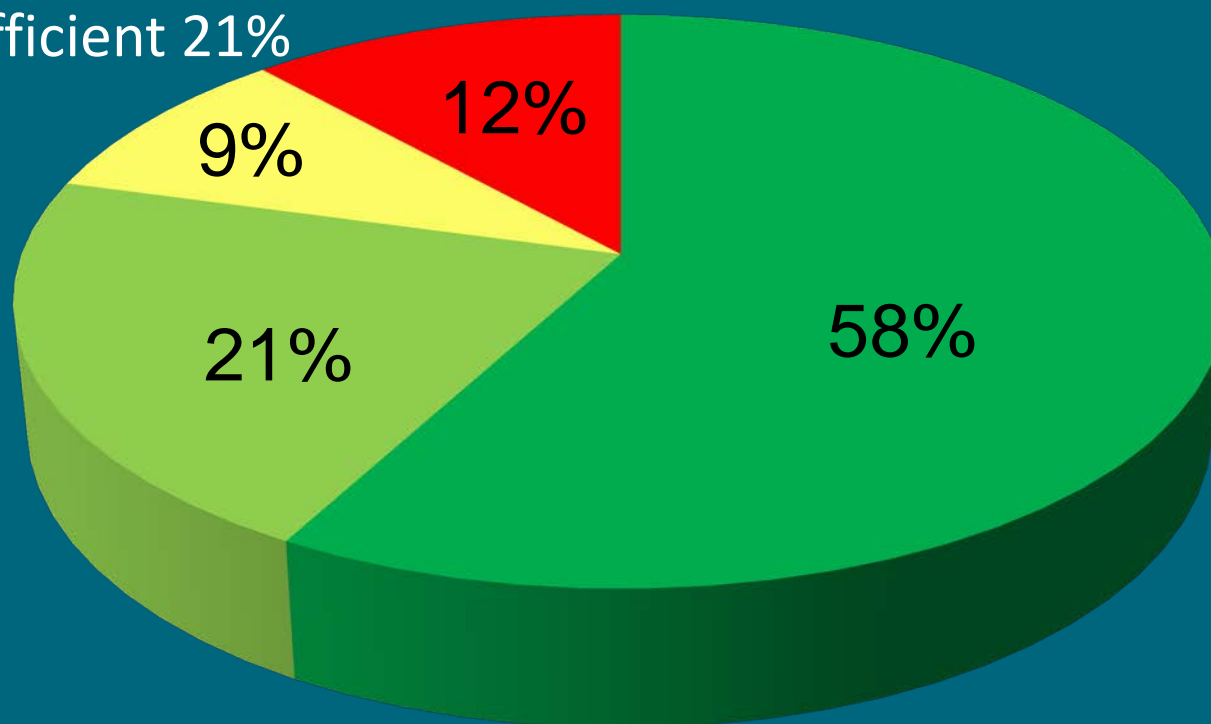


- Optimal
- Good
- Borderline
- Poor

Insuff. { too weak / false neg.: ~ 90%
over-stained / false pos.: ~ 10%

Breast cancer module ~ 9,000 slides (~35,000 core sections)

Insufficient 21%



- Optimal
- Good
- Borderline
- Poor

Insuff. { too weak / false neg.: ~ 90%
over-stained / false pos.: ~ 10%

Major causes of **insufficient** stains in ~ 9,000 slides

Less successful antibodies/RTUs	17 %
---------------------------------	------

Inappropriate antibody dilution	20 %
---------------------------------	------

Inappropriate epitope retrieval	27 %
---------------------------------	------

Inappropriate detection kit	19 %
-----------------------------	------

Other inappropriate lab. performance	17 %
--------------------------------------	------

Endogenous biotin reaction

Section drying-out after HIER

Technical platform error

....

Unexplained

Go for Low antigen expressors ~

Critical Stain Quality Indicators (CSQI)

- essential to evaluate sensitivity
- essential to assure consistency

Normal tissues when ever possible - easier to recognize and ensure the quality

- 90 % of insufficient staining results in EQA are caused by weak/false negative results
- often related to the use of inappropriate positive tissue controls.....

Standardization of Negative Controls in Diagnostic Immunohistochemistry: Recommendations From the International Ad Hoc Expert Panel

Emina E. Torlakovic, MD, PhD,†‡ Glenn Francis, MBBS, FRCPA, MBA, FFSc (RCPA),§||¶
 John Garratt, RT,†‡# Blake Gilks, MD, FRCPC,†‡** Elizabeth Hyjek, MD, PhD,*
 Merdol Ibrahim, PhD,†† Rodney Miller, MD,‡‡ Søren Nielsen, HT, CT,§§||
 Eugen B. Petcu, MD, PhD,§ Paul E. Swanson, MD,¶¶ Clive R. Taylor, MD, PhD,##
 and Mogens Vyberg, MD§§|| AIMM 2014, 22:241*

Standardization of Positive Controls in Diagnostic Immunohistochemistry: Recommendations From the International Ad Hoc Expert Committee

Emina E. Torlakovic, MD, PhD,† Søren Nielsen, HT, CT,‡§ Glenn Francis, MBBS, FRCPA,
 MBA, FFSc (RCPA),||¶# John Garratt, RT,†** Blake Gilks, MD, FRCPC,†††
 Jeffrey D. Goldsmith, MD,‡‡ Jason L. Hornick, MD, PhD,*§§ Elizabeth Hyjek, MD, PhD,*
 Merdol Ibrahim, PhD,|| Keith Miller, FIBMS,|| Eugen Petcu, MD, PhD,||
 Paul E. Swanson, MD,¶¶## Xiaoge Zhou, MD,***††† Clive R. Taylor, MD, PhD,‡‡‡
 and Mogens Vyberg, MD‡§ AIMM 2015, 23:1*

Evolution of Quality Assurance for Clinical Immunohistochemistry in the Era of Precision Medicine:

Part 1: Fit-for-Purpose Approach to Classification of Clinical Immunohistochemistry Biomarkers

AIMM 2016-17

Carol C. Cheung, MD, PhD, JD,† Corrado D'Arrigo, MB, ChB, PhD, FRCPath,‡§||
 Manfred Dietel, MD, PhD,¶ Glenn D. Francis, MBBS, FRCPA, MBA, FFSc (RCPA),#**††
 C. Blake Gilks, MD,‡‡ Jacqueline A. Hall, PhD,§§||| Jason L. Hornick, MD, PhD,¶¶
 Merdol Ibrahim, PhD,### Antonio Marchetti, MD, PhD,*** Keith Miller, FIBMS,##
 J. Han van Krieken, MD, PhD,††† Soren Nielsen, BMS,‡‡‡§§§ Paul E. Swanson, MD,||| ||
 Clive R. Taylor, MD,¶¶¶ Mogens Vyberg, MD,‡‡‡§§§ Xiaoge Zhou, MD,####****
 and Emina E. Torlakovic, MD, PhD,*††††††††*

*From the International Society for Immunohistochemistry and Molecular Morphology (ISIMM)
 and International Quality Network for Pathology (IQN Path)*

Evolution of Quality Assurance for Clinical Immunohistochemistry in the Era of Precision Medicine – Part 2: Immunohistochemistry Test Performance Characteristics

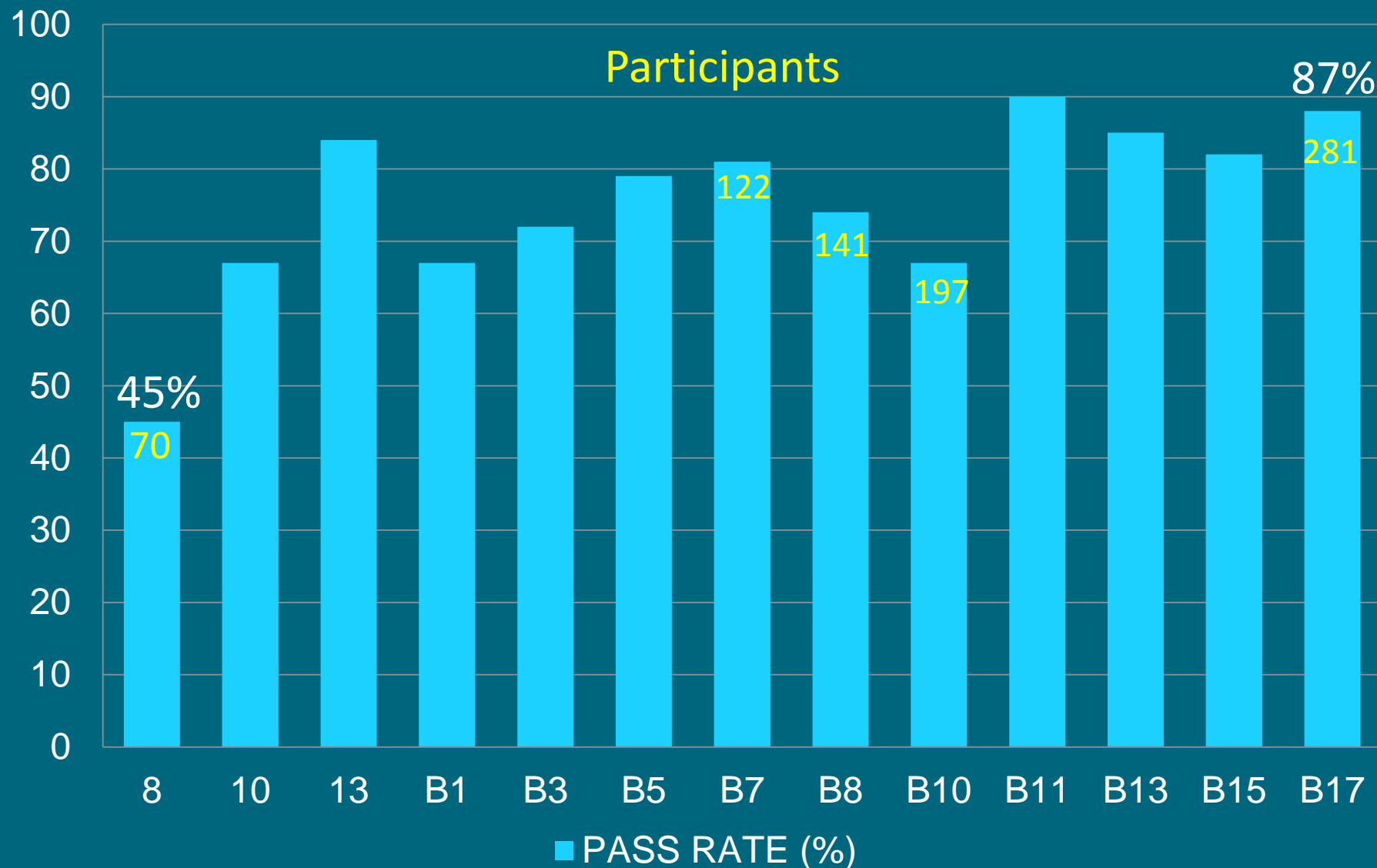
Evolution of Quality Assurance for Clinical Immunohistochemistry in the Era of Precision Medicine.

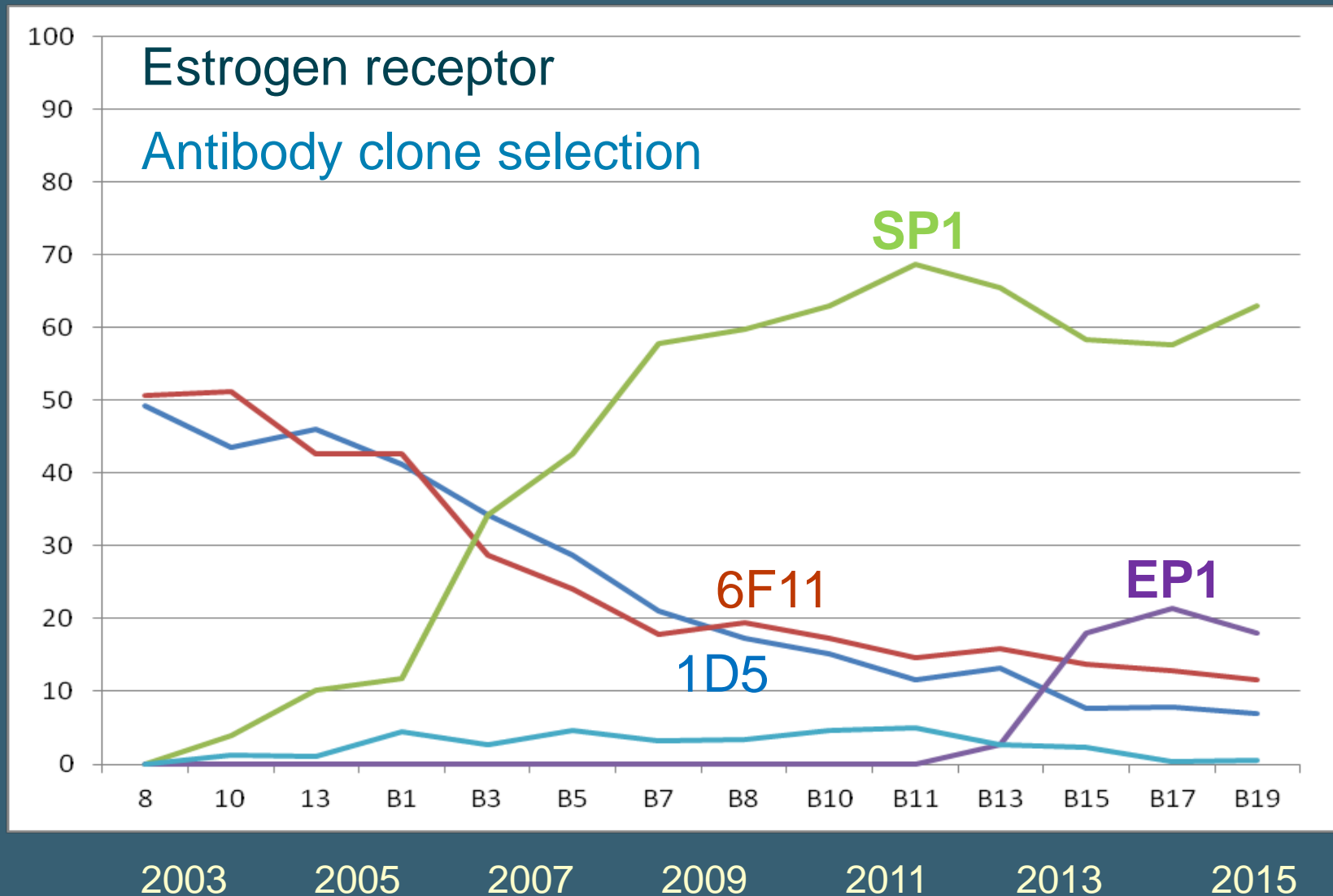
Part 3: Technical Validation of Immunohistochemistry (IHC) Assays in Clinical IHC Laboratories

Evolution of Quality Assurance for Clinical Immunohistochemistry in the Era of Precision Medicine:

Part 4: Tissue Tools for Quality Assurance in Immunohistochemistry

NordiQC EQA: Estrogen Receptor in 13 runs

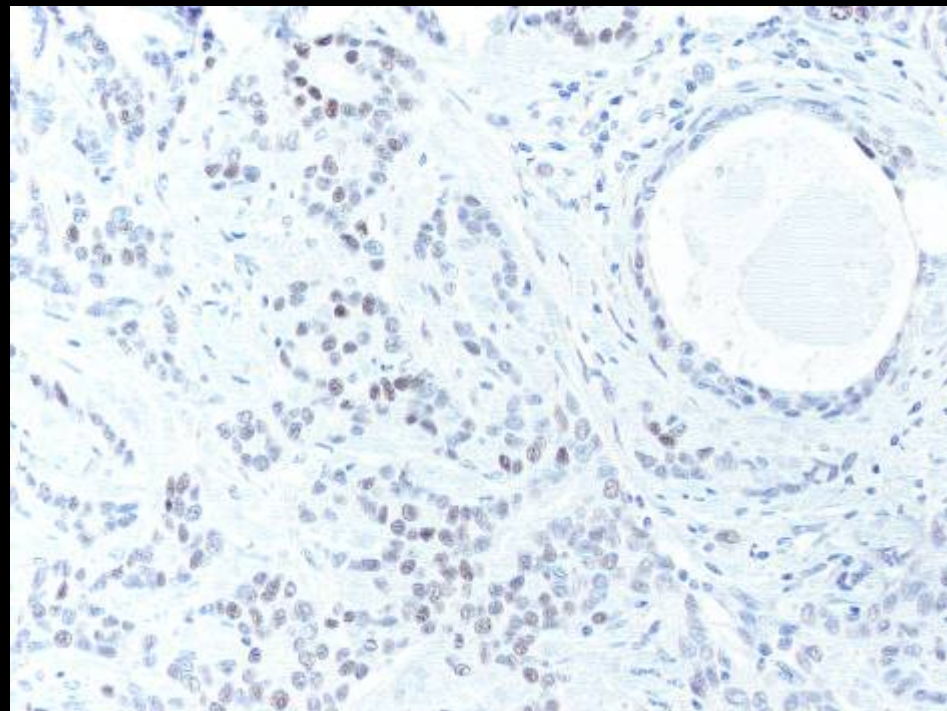




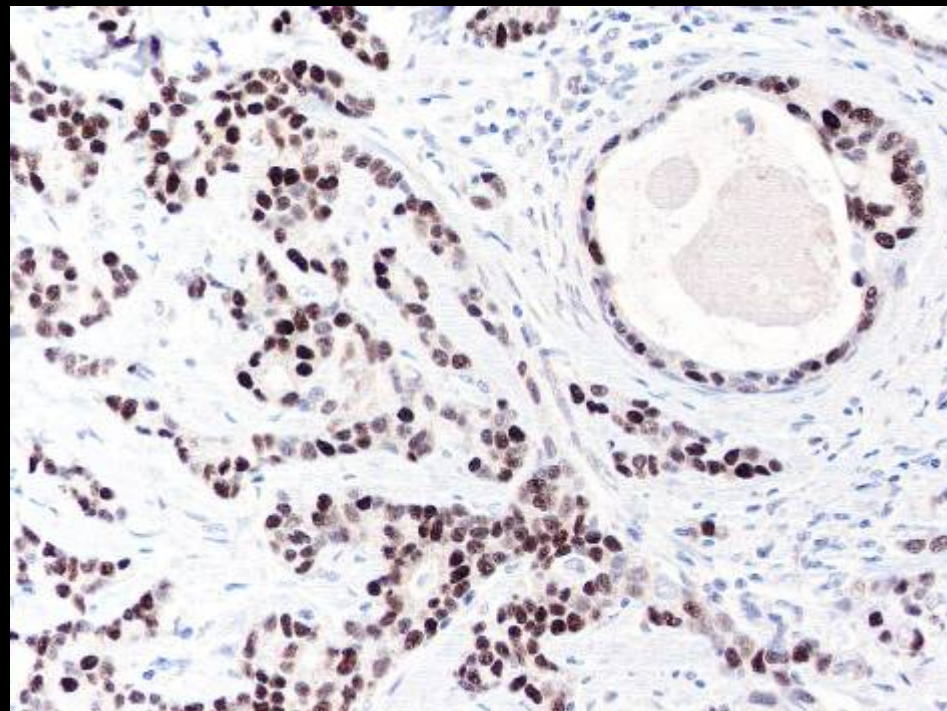
IHC – Optimal performance

ER 1D5 1:100

HIER Ci pH 6



2-step polymer

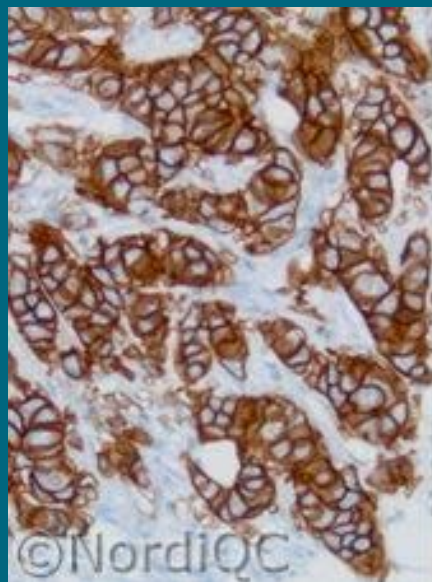


3-step polymer

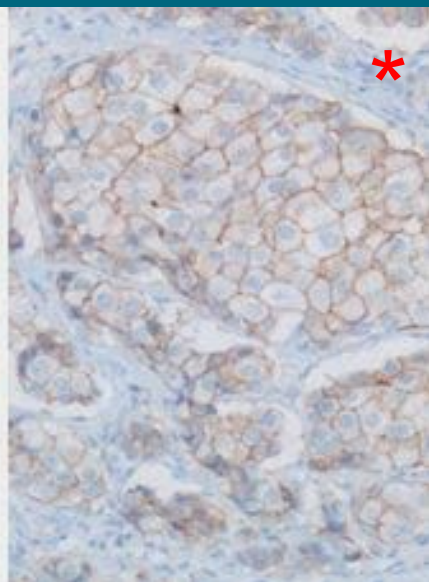
Pass rate (optimal + good) by participant status

Estrogen receptor	New participants	'Old' participants
Run 10, 2004	57%	71%
Run B15, 2010	70%	86%
Run B19, 2015	51%	73%
Average	59%	77%

NordiQC runs for HER2 IHC



Ampl. 3+

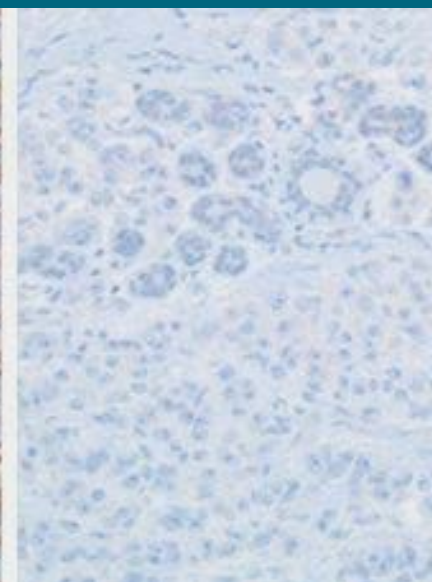


Ampl. 2+

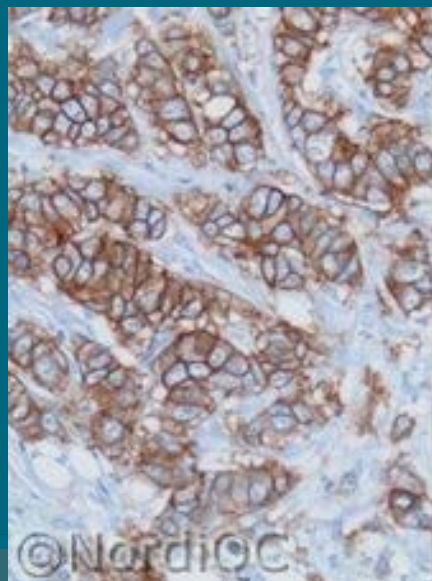
Optimal



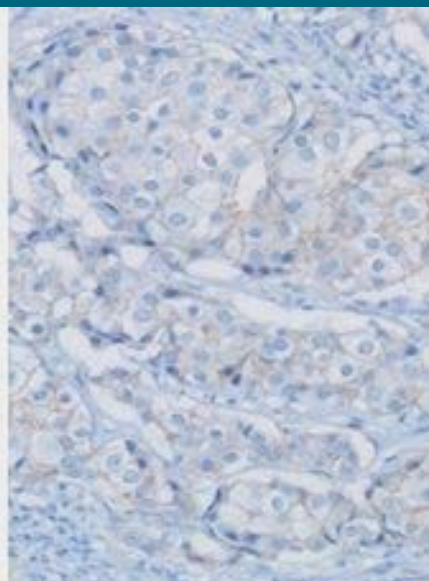
Unampl. 2+



Unampl. 0

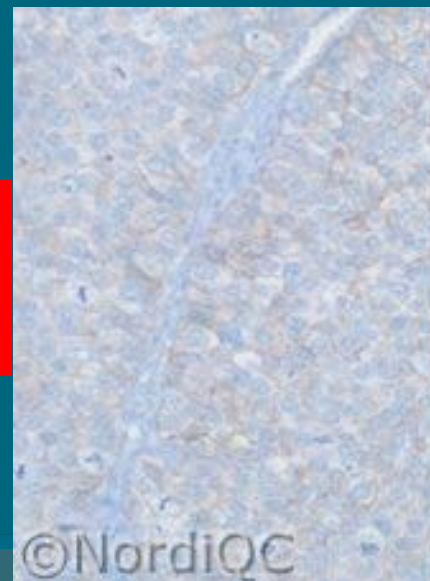


Ampl. 3+

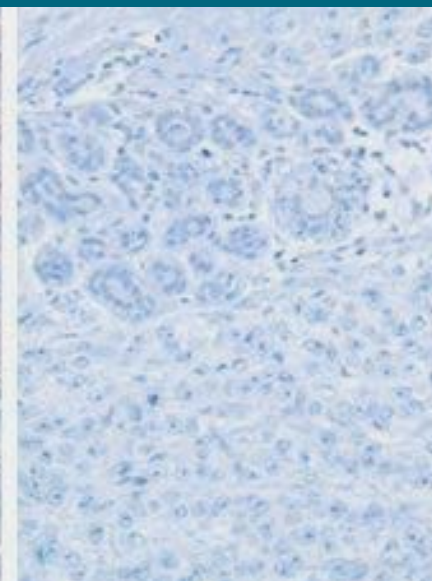


Ampl. 1+

Poor

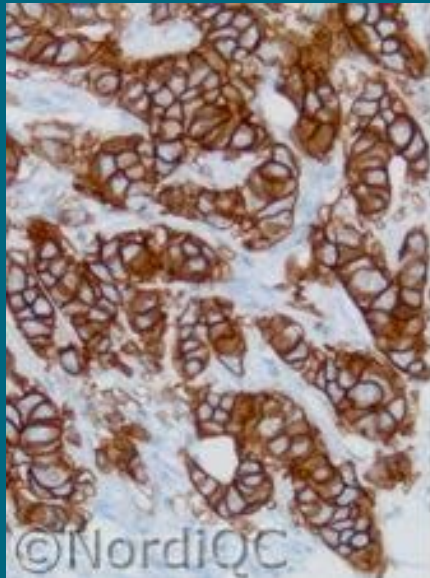


Unampl. 1+

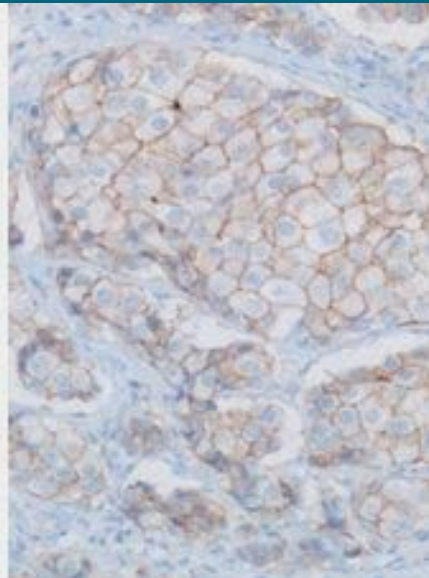


Unampl. 0

NordiQC runs for HER2 IHC



Ampl. 3+

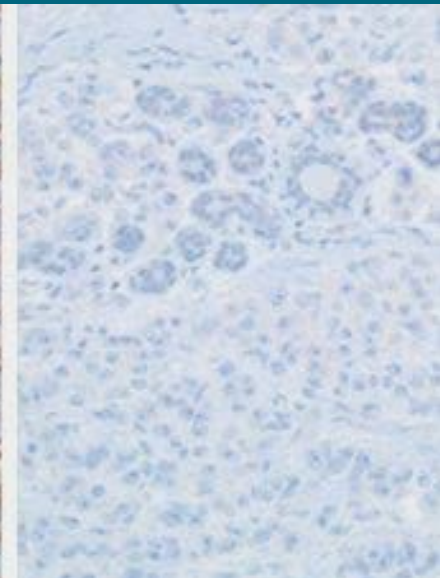


Ampl. 2+

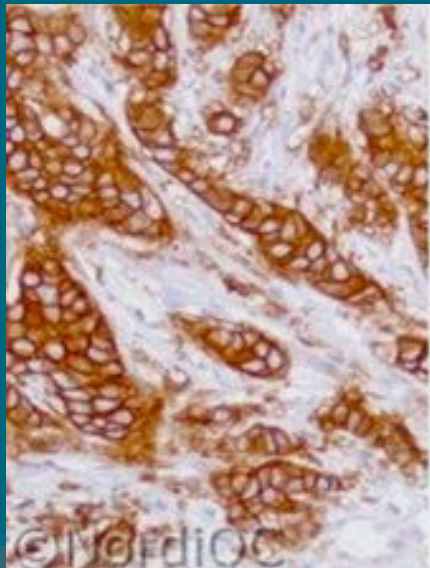
Optimal



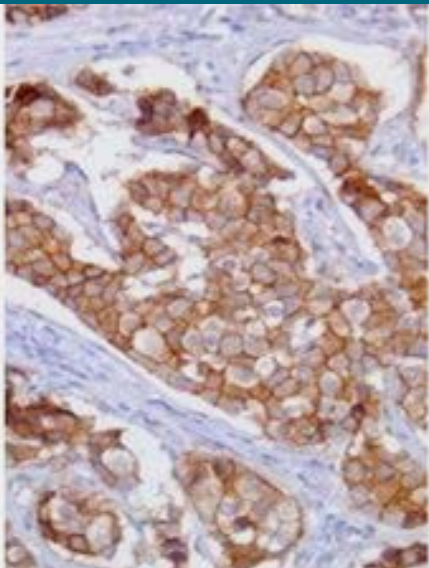
Unampl. 2+



Unampl. 0

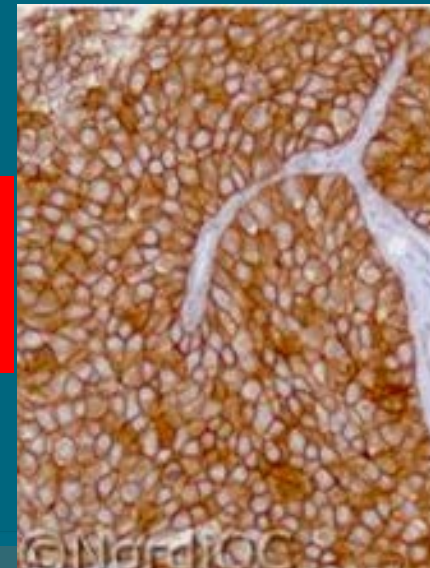


Ampl. 3+

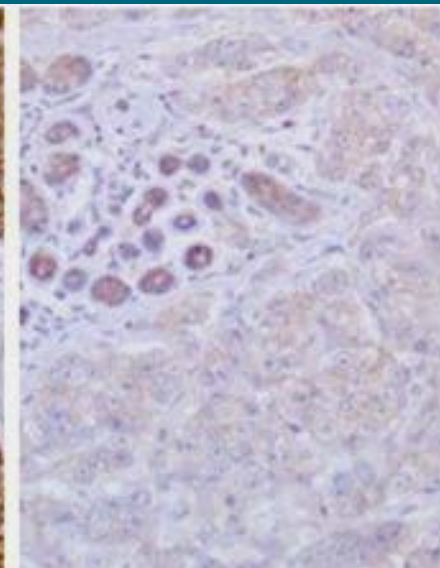


Ampl. 2+

Poor

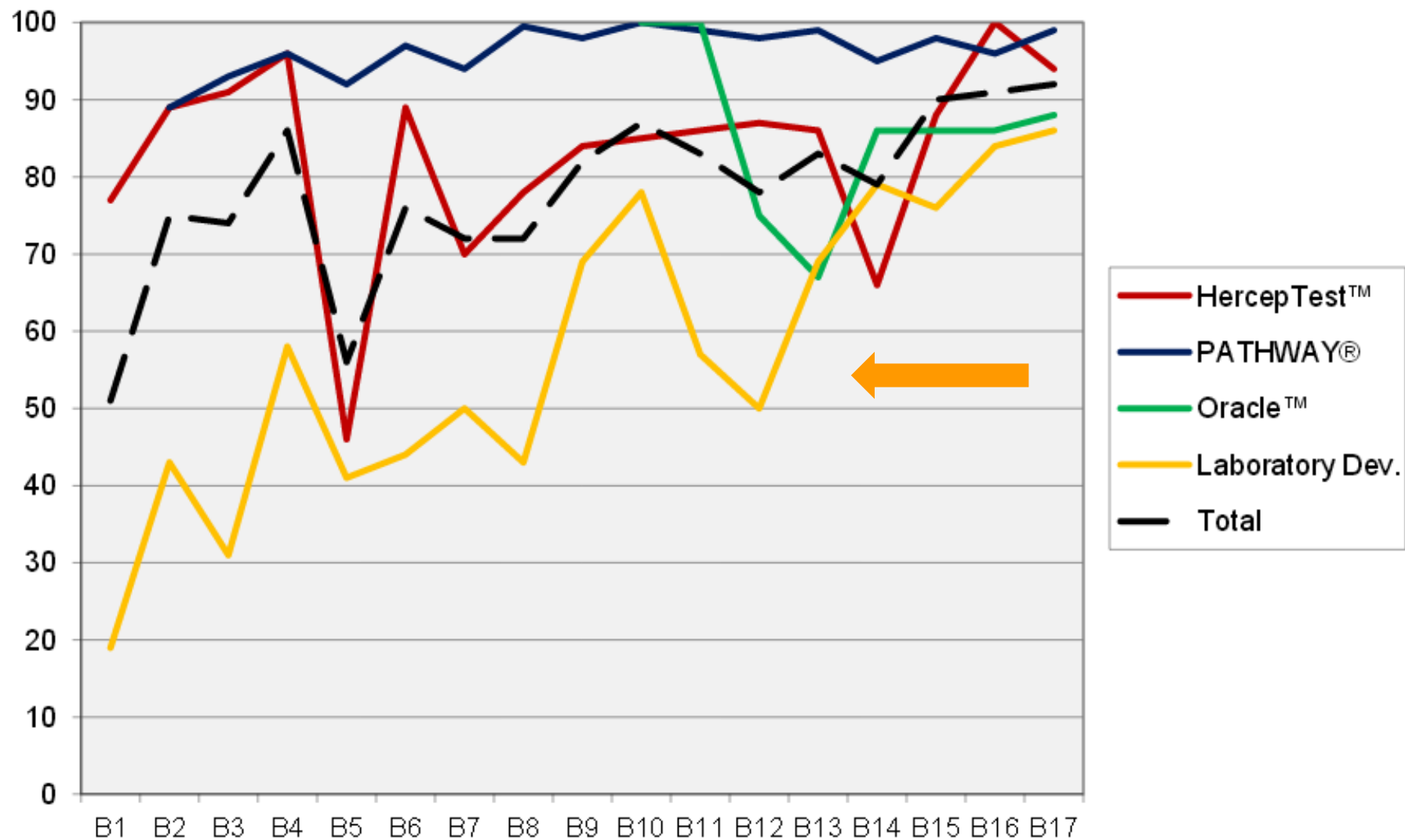


Unampl. 3+



Unampl. 1

HER-2 staining results in 17 runs



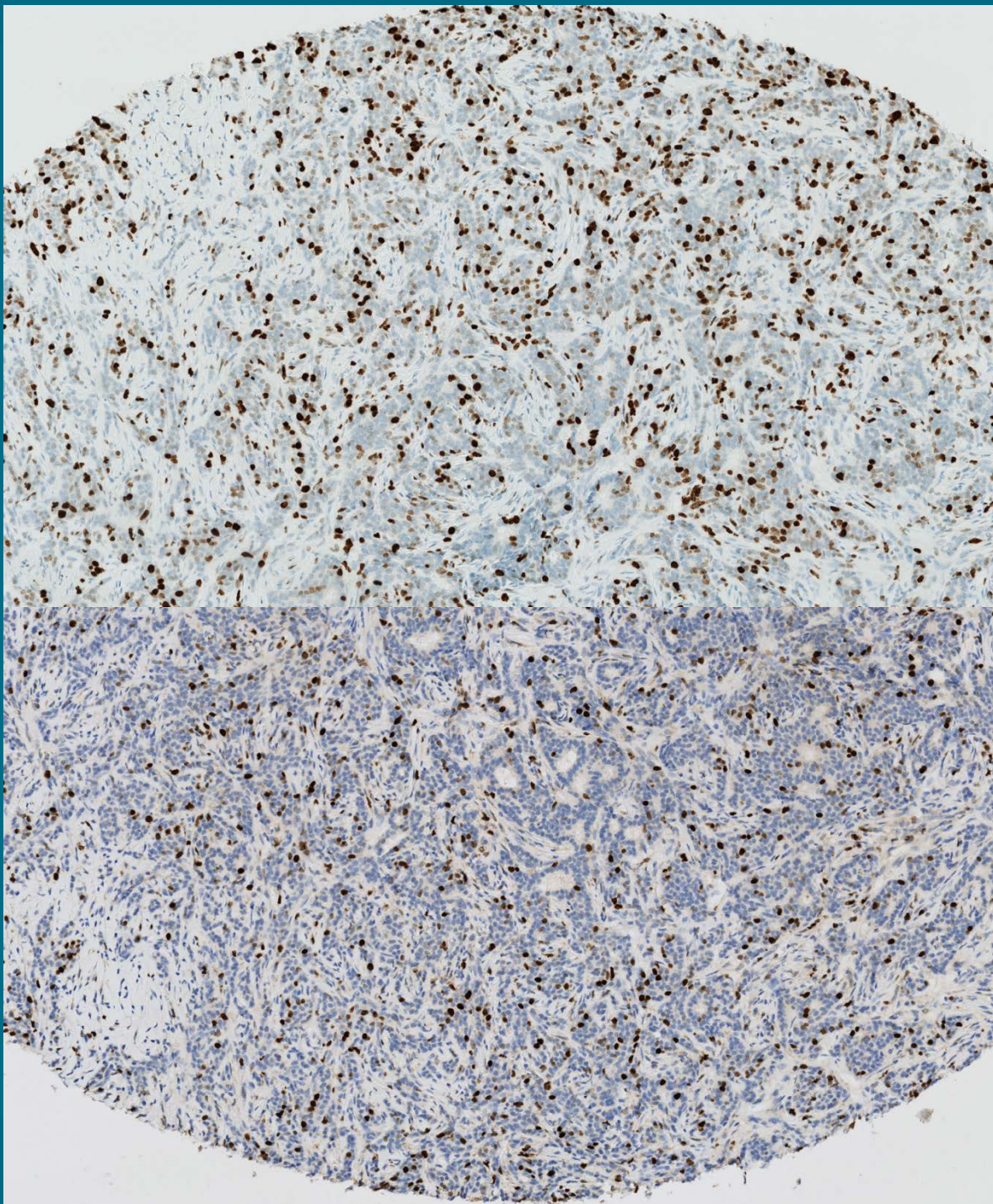
Ki67 Index, HER2 Status, and Prognosis of Patients With Luminal B Breast Cancer

Maggie C. U. Cheang, Stephen K. Chia, David Voduc, Dongxia Gao, Samuel Leung, Jacqueline Snider, Mark Watson, Sherri Davies, Philip S. Bernard, Joel S. Parker, Charles M. Perou, Matthew J. Ellis, Torsten O. Nielsen

J Natl Cancer Inst 2009;101: 736 – 750

Ki67 antibody clone SP6 applied at a 1:200 dilution for 32 minutes, by following the Ventana Benchmark automated immunostainer standard Cell Conditioner 1 (CC1) protocol at 98°C for 30 minutes.

The best Ki67 index cut point to distinguish luminal B from luminal A tumors was **13.25%**.



Ki67 immunoassay

Digital image analysis

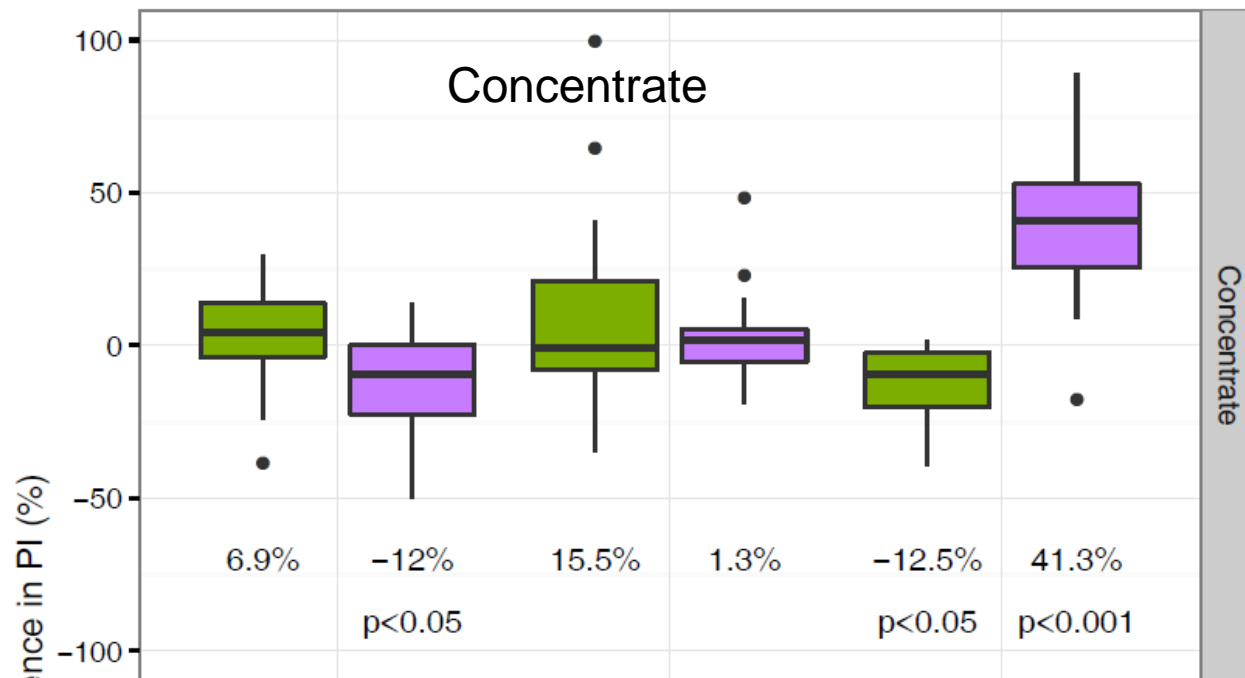
Clone SP6
concentrate
Ventana
platform

Prolif.index
38 %

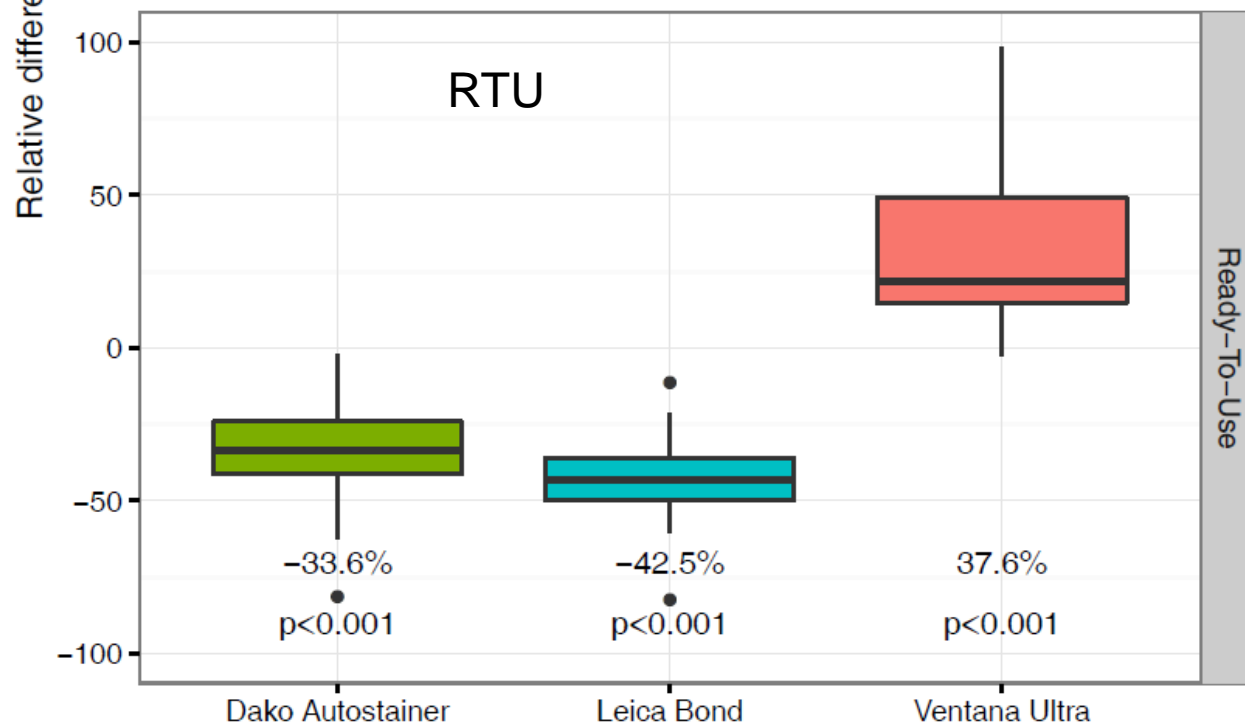
Clone MM1
RTU
Leica platform

Prolif.index
12 %



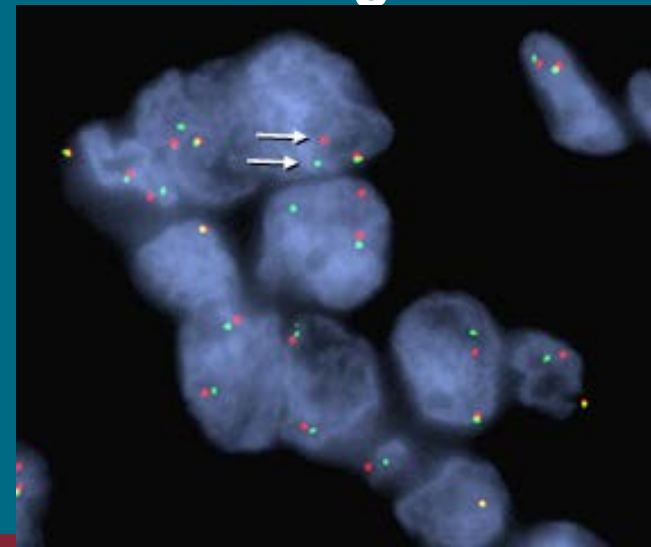


Mib1 as
concentrate
in optimized
protocols on
3 platforms



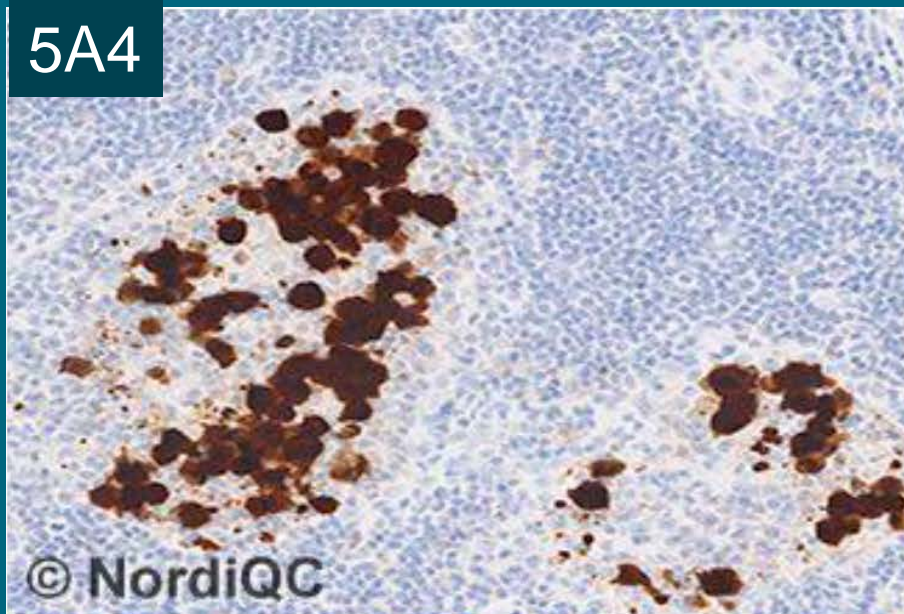
ri Mutations can be identified by mutation-specific proteins:
ALK (anaplastic large cell lymphoma, lung adenocarcinoma).

- Anaplastic lymphoma kinase (ALK): a transmembrane receptor tyrosin kinase
- CD30+ anaplastic large cell lymphomas (ALCLs) may be associated with a balanced (2;5)(p23;5q35) chromosomal translocation
- Anaplastic lymphoma kinase (ALK) rearrangements are present in about **5% of advanced non-small-cell lung cancer**
- Crizotinib was the first ALK tyrosine kinase inhibitor licensed for the treatment of metastatic ALK-positive NSCLC

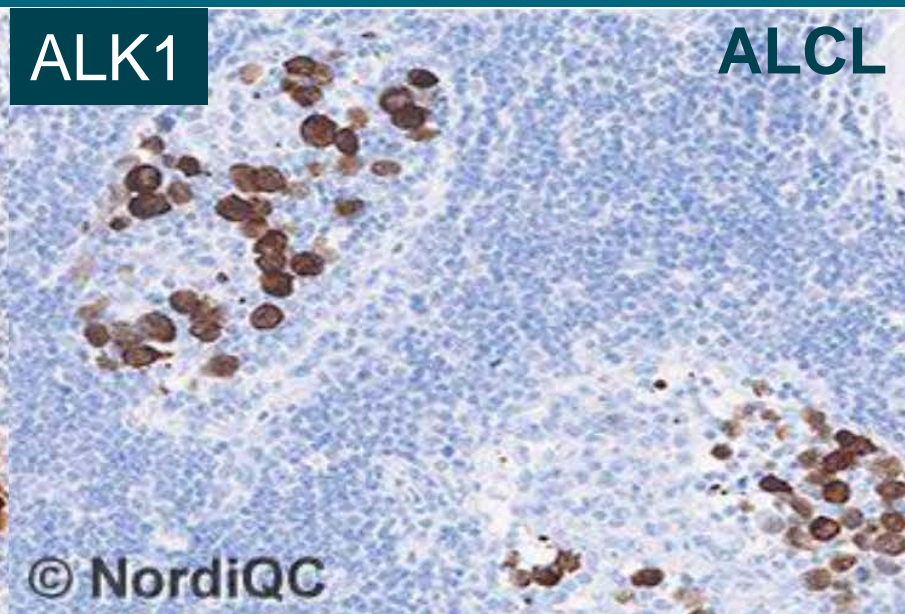


Lung ALK – run 45, 176 labs

5A4

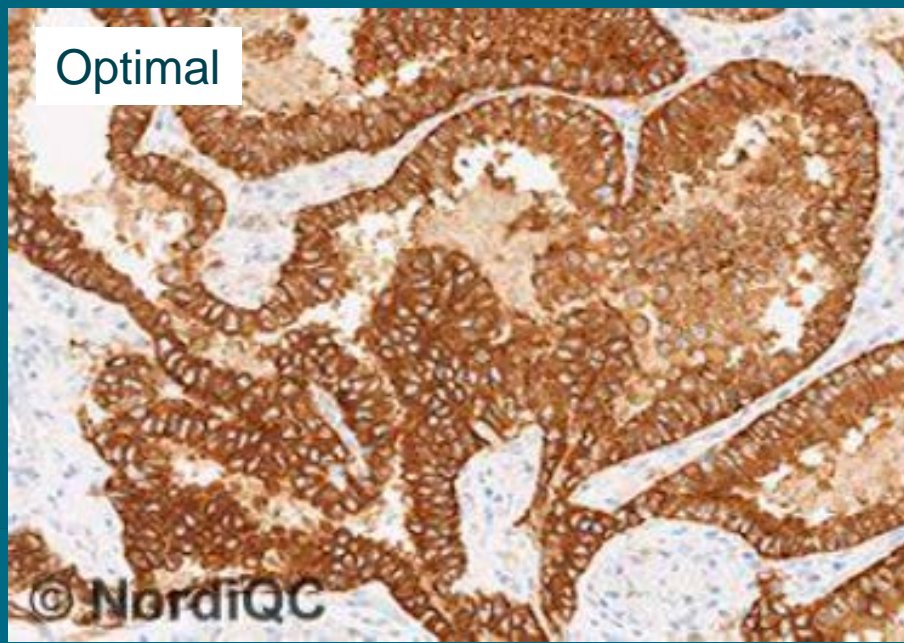


ALK1

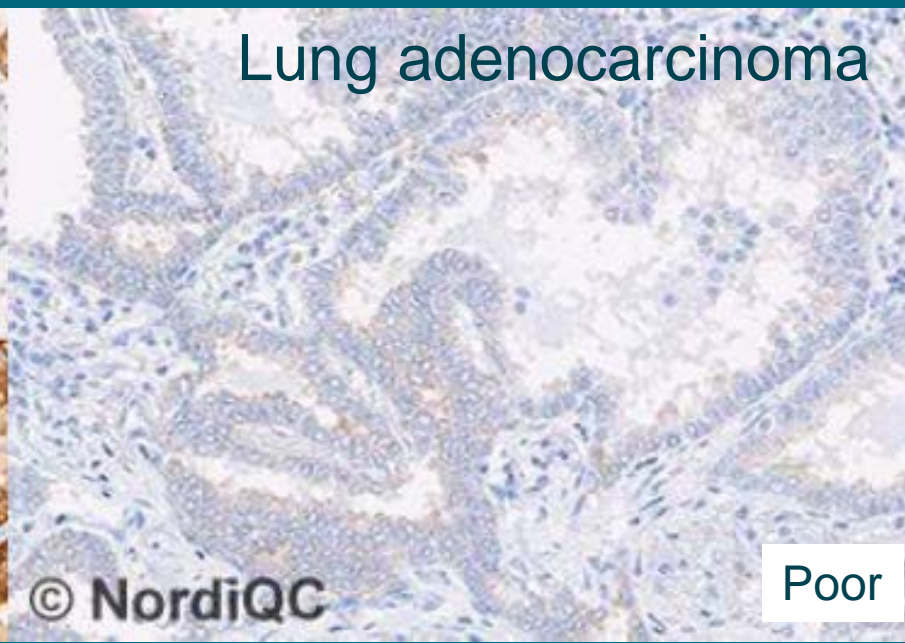


ALCL

Optimal



Lung adenocarcinoma



Poor

Lung ALK – run 45, 176 labs

Table 1. **Antibodies and assessment marks for lu-ALK, run 45**

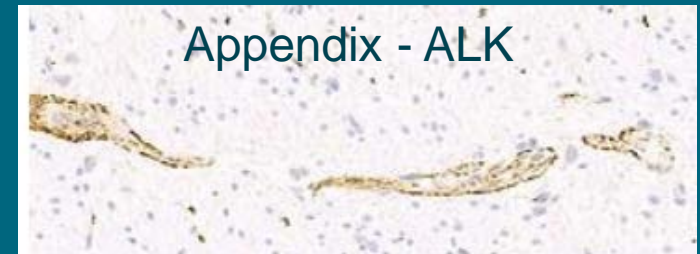
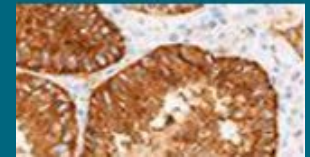
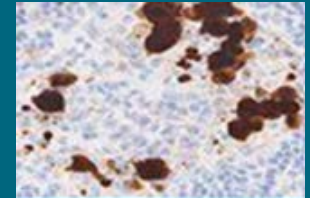
Concentrated antibodies	n	Vendor	Optimal	Good	Borderline	Poor	Suff. ¹	Suff. OPS ²
mAb clone 5A4	46 3 2 1 1 1	Leica/Novocastra Thermo/NeoMarkers Monosan Abcam Biocare Zytomed	24	16	13	1	74%	81%
mAb clone ALK1	8	Dako	0	0	3	5	0%	-
mAb clone OTI1A4	5	ORIGENE	4	1	0	0	100%	100%
rmAb clone D5F3	21 1	Cell Signaling PrimeBioMed	18	2	1	1	91%	95%
rmAb clone SP8	2	Thermo/NeoMarkers	0	0	1	1	-	-
Ready-To-Use antibodies								
mAb ALK1 IR641	15	Dako	0	0	4	11	0%	-
mAb clone ALK1 790/800-2918	10	Ventana	0	1	6	3	10%	-
mAb clone ALK1 204M-18	1	Cell Marque	0	0	0	1	-	-
mAb clone ALK1 GA641	1	Dako	0	0	0	1	-	-
rmAb clone D5F3 790-4794	47	Ventana	41	4	2	0	96%	96%

The immunoassay must fit for the purpose:

- Identify the antibody useful for the specific task

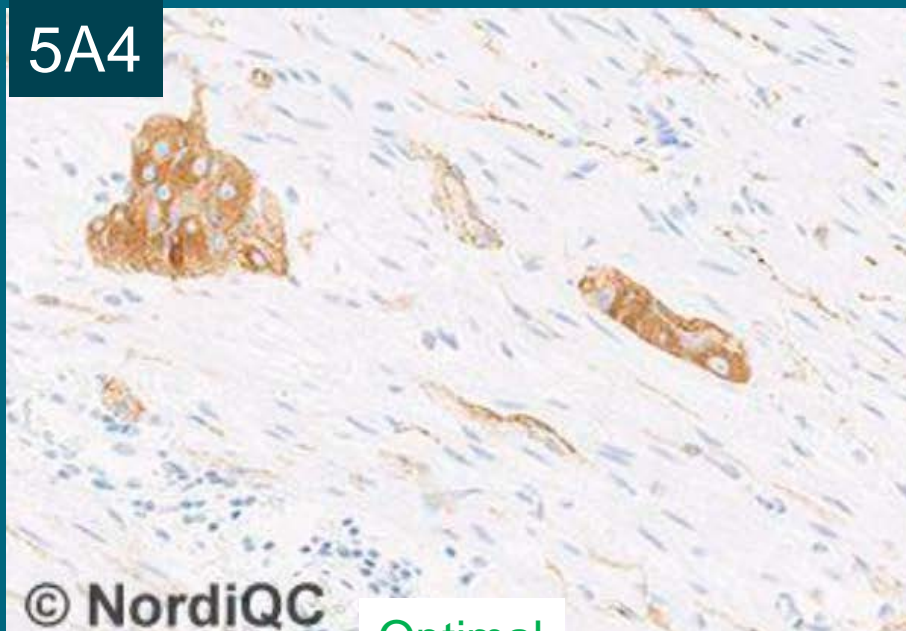
The right external controls must be used:

- Tissue with high epitope expression to identify the right antibody
 - Appendix
- Tissue with low epitope expression to assure the sensitivity:
 - ALK-positive lung adenocarcinoma
- Tissue with no epitope expression to assure the specificity
 - e.g., liver



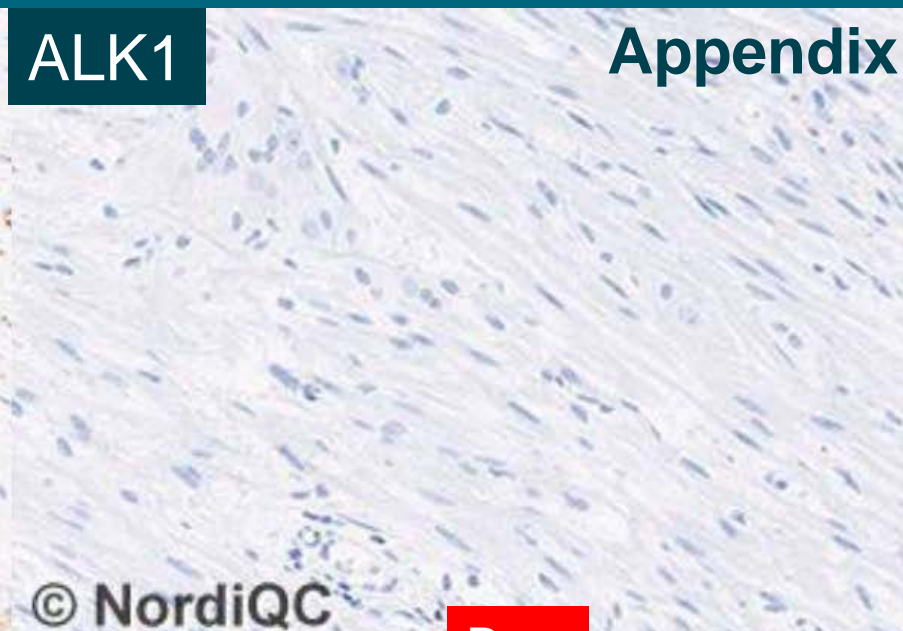
Lung ALK – run 45, 176 labs

5A4

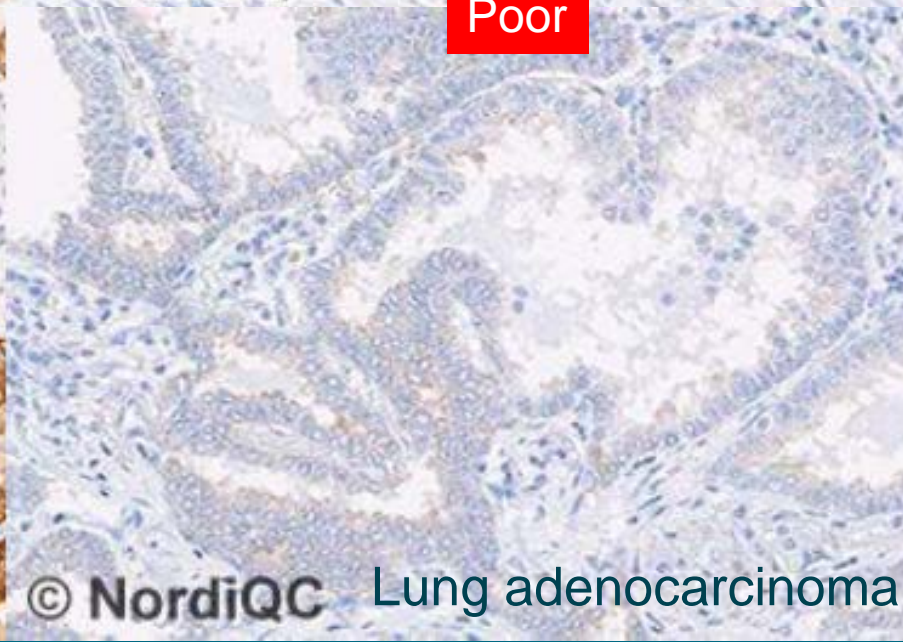
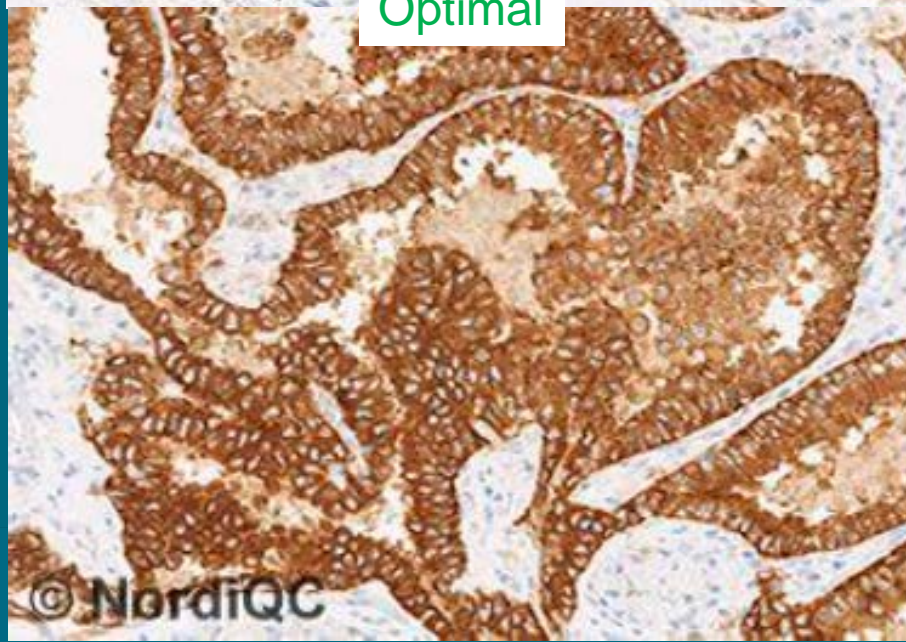


Optimal

ALK1



Poor



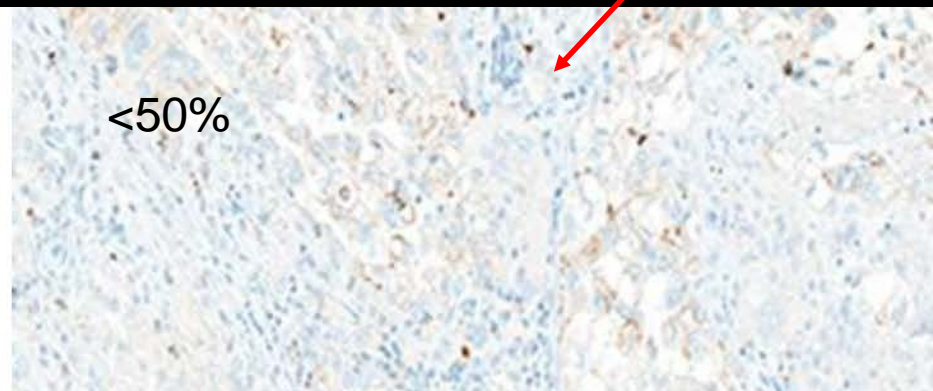
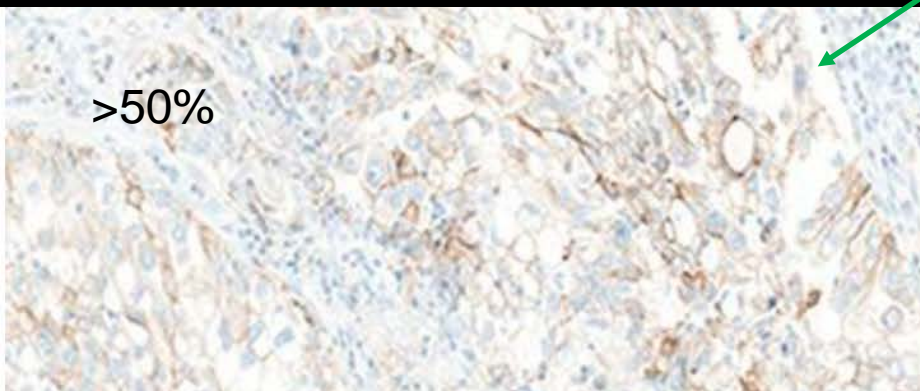
Appendix

Lung adenocarcinoma

External Quality Assurance – PDL1

Table 1. Assessment marks for IHC assays and antibodies run C1, PD-L1 IHC

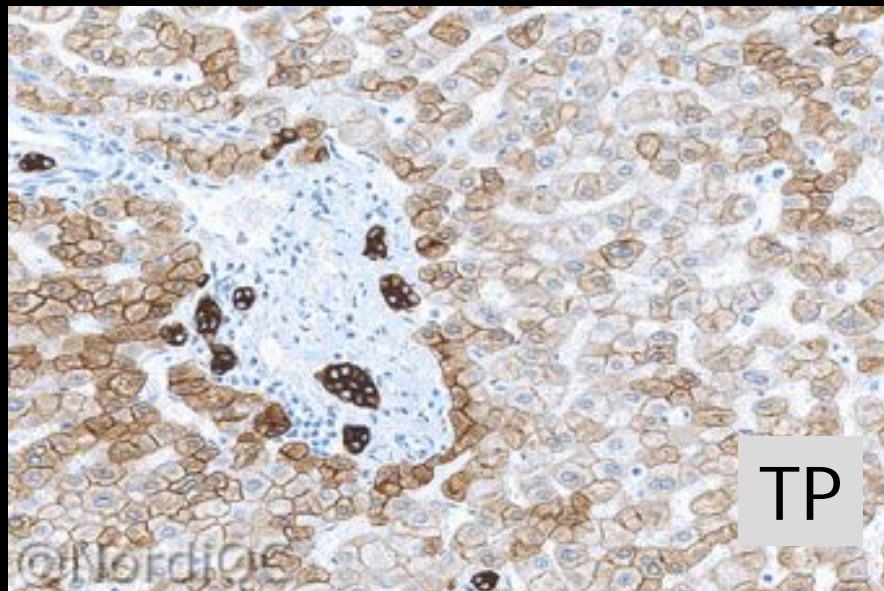
CE-IVD / FDA approved PD-L1 assays			Optimal	Good	Borderline	Poor	Suff. ¹	Suff. OPS ²
22C3 pharmDX, SK006	12	Dako/Agilent	10	1	0	1	92%	92%
22C3 pharmDX, SK006⁴	2	Dako/Agilent	0	0	1	1	-	-
28-8 pharmDX, SK005	7	Dako/Agilent	3	3	1	0	86%	86%
SP263, 790-4905	16	Ventana/Roche	9	2	2	3	69%	77%
SP142, 740-4859	1	Ventana/Roche	0	0	0	1	-	-
Antibodies ³ for laboratory developed PD-L1 assays, conc. antibody			Optimal	Good	Borderline	Poor	Suff. ¹	Suff. OPS ²
mAb clone 22C3	13	Dako/Agilent	1	1	4	7	15%	-
mAb clone E1L3N	8	Cell Signaling	1	1	1	5	25%	-
mAb CAL10	1	Biocare	0	0	1	0	-	-
rmAb clone 28-8	6	Abcam	0	1	1	4	17%	-
rmAb clone ZR3	1	Zeta Corporation	1	0	0	0	-	-



Misleading data sheets

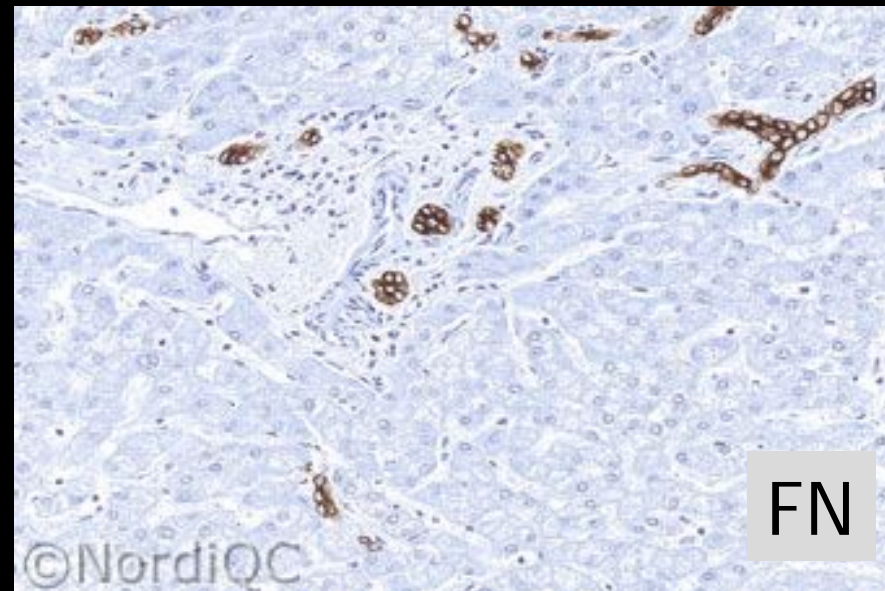
Pan-Cytokeratin

Inappropriate retrieval (31%)



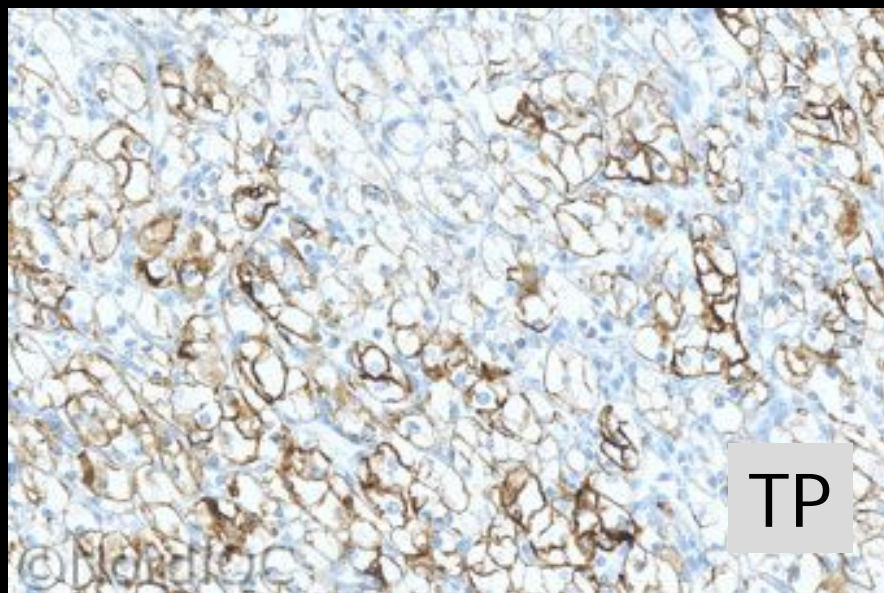
Liver

AE1/AE3 + HIER



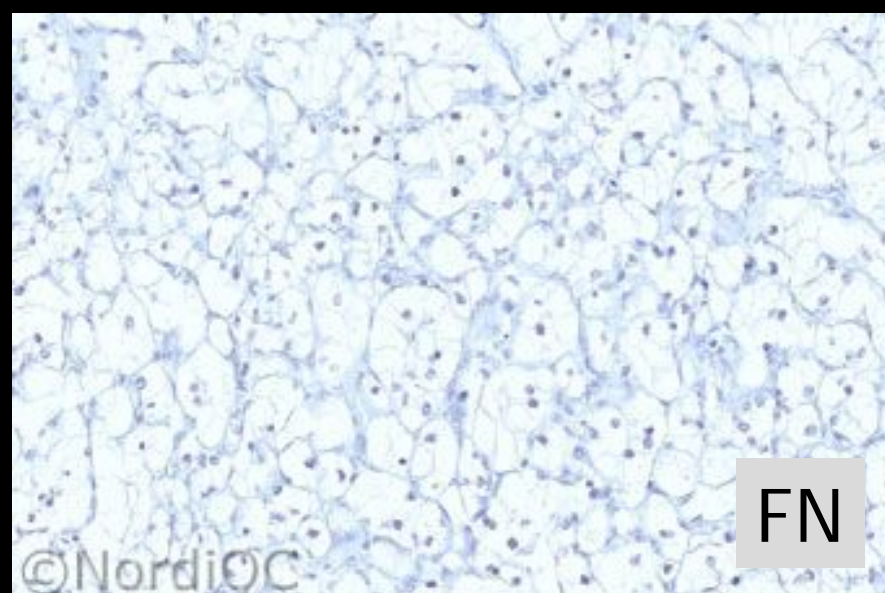
FN

AE1/AE3 + proteolysis



RCC

FN



Performance history

This was the 7th NordiQC assessment of CK-PAN. The overall pass rate has remained almost constant in the last 5 runs performed, as shown in table 2.

Table 2. **Proportion of sufficient results for CK-PAN in the seven NordiQC runs performed**

	<u>Run 8 2003</u>	<u>Run 15 2005</u>	<u>Run 20 2008</u>	<u>Run 24 2008</u>	<u>Run 30 2010</u>	<u>Run 36 2012</u>	<u>Run 41 2014</u>
Participants, n=	72	85	103	123	168	202	233
Sufficient results	53%	58%	62%	60%	65%	65%	67%

AE1/AE3 : Optimal results only obtained by **HIER** in NordiQC runs

Dako: RTU – HIER

Conc: **Proteolysis** or HIER

Leica: RTU – **Proteolysis**

Conc: HIER

Thermo:

Conc: HIER Quanto – **Proteolysis** UltraVision

.....

VMS: RTU - **Proteolysis**

Misleading data sheets + Wrong control material used

CEA – run 47, 2016, 255 labs

Table 1. Antibodies and assessment marks for CEA, run 47

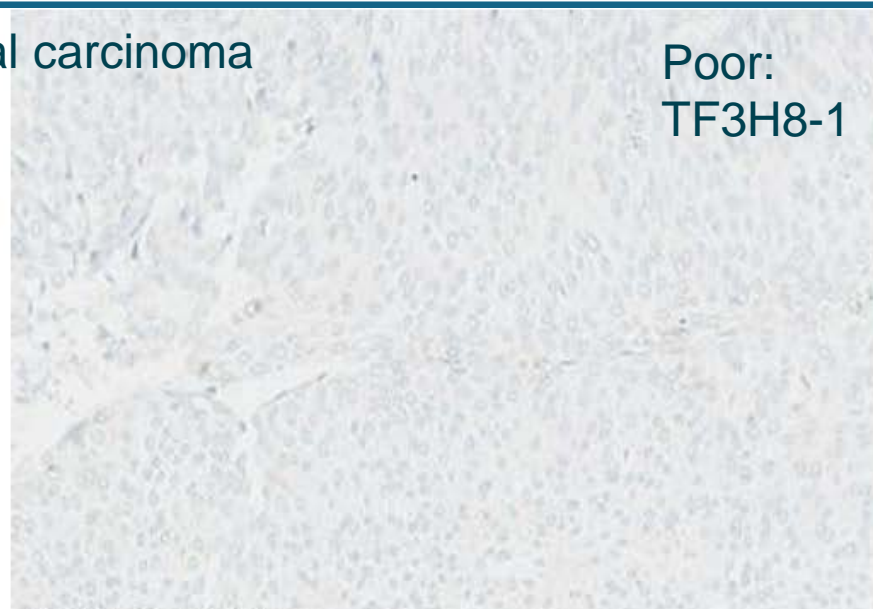
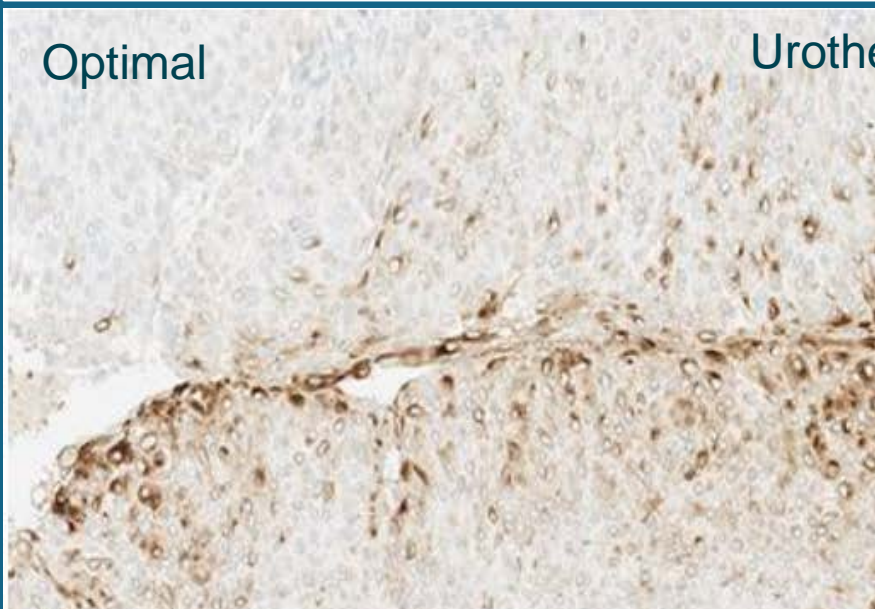
Concentrated Antibodies	n	Vendor	Optimal	Good	Borderline	Poor	Suff. ¹	Suff. OPS ²
mAb clone 12-140-10	3	Leica/Novocastra	0	0	0	3	-	-
mAb clone CEA31	9	Cell Marque	6	0	3	1	67%	75%
mAb COL-1	6	Thermo/Neomarkers	11	7	2	0	90%	94%
	5	Invitrogen/Zymed						
	5	Biocare						
	2	Immunologic						
	1	Zytomed						
	1	GeneTex						
mAb II-7	85	Dako/Agilent	2	19	60	4	25%	58%

Ready-To-Use Antibodies								
mAb clone CEA31 760-4594	53	Ventana/Cell Marque	22	26	5	0	91%	100%
mAb clone II-7 IR/IS622/GA622	47	Dako/Agilent	0	6	40	1	13%	-
mAb clone II-7 PA0004	12	Leica	0	5	6	1	42%	-
mAb clone TF3H8-1 760-2507	13	Ventana/Roche	0	0	0	13	0%	-

Optimal

Urothelial carcinoma

Poor:
TF3H8-1



Optimal

Liver

Poor:
TF3H8-1

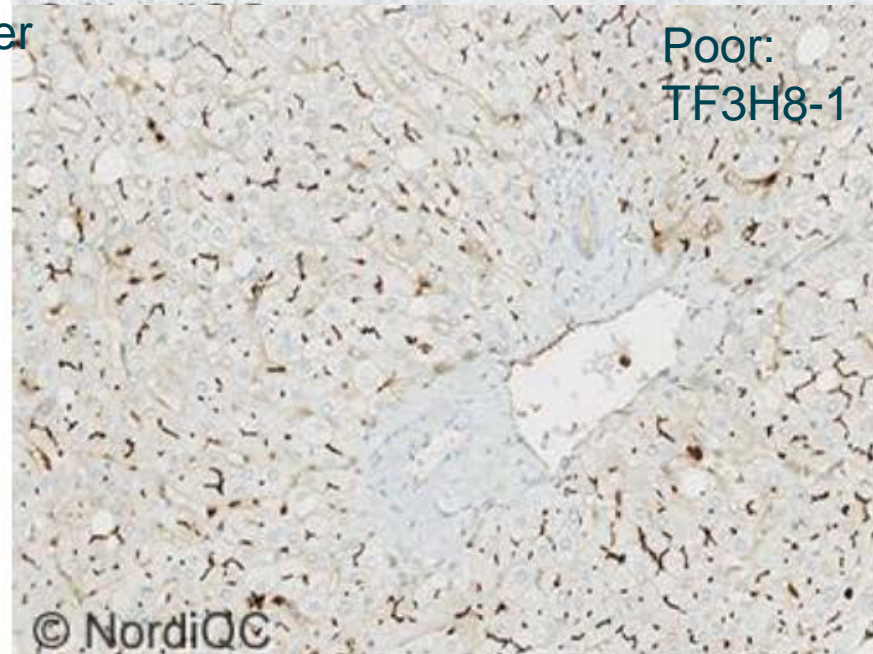
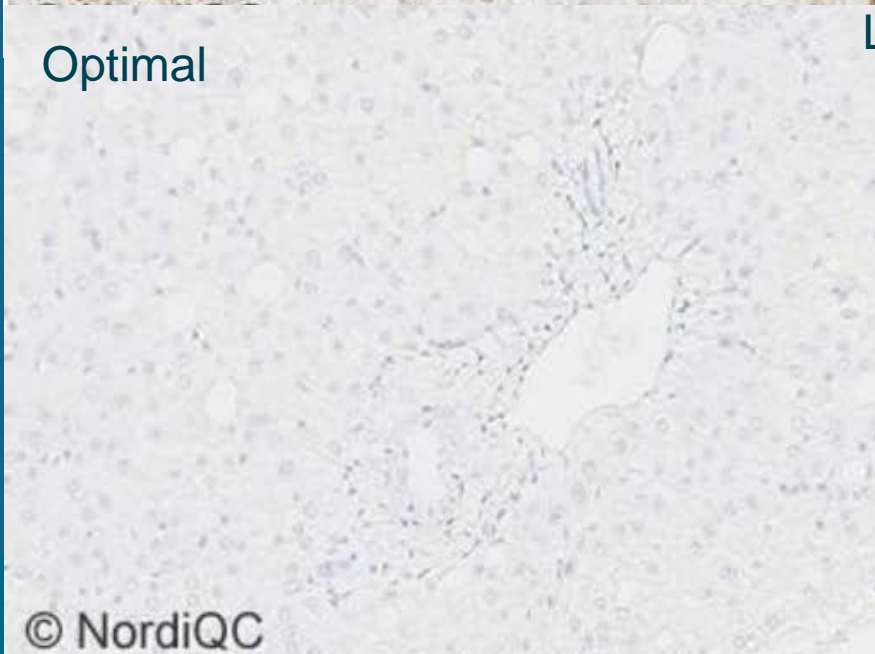


Table 1. Antibodies and assessment marks for CDX2, run 48

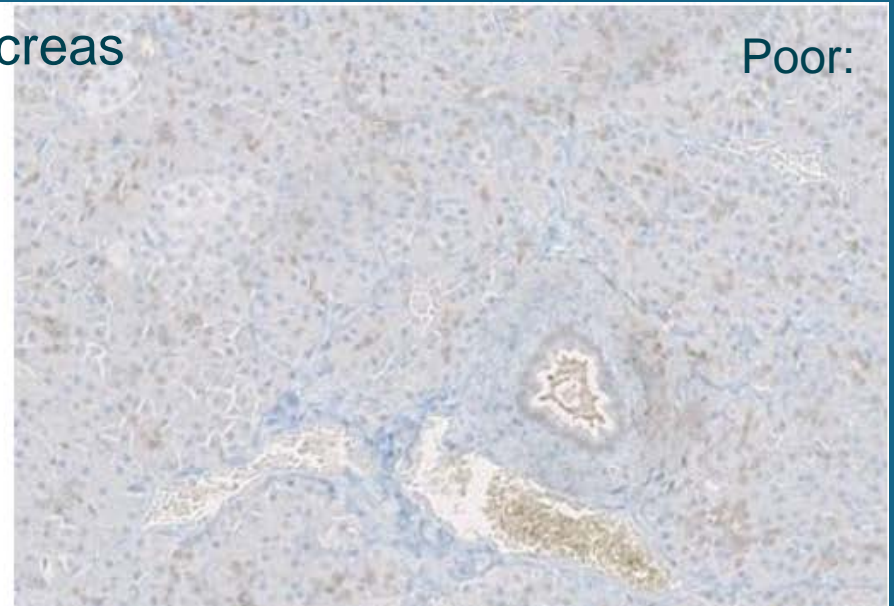
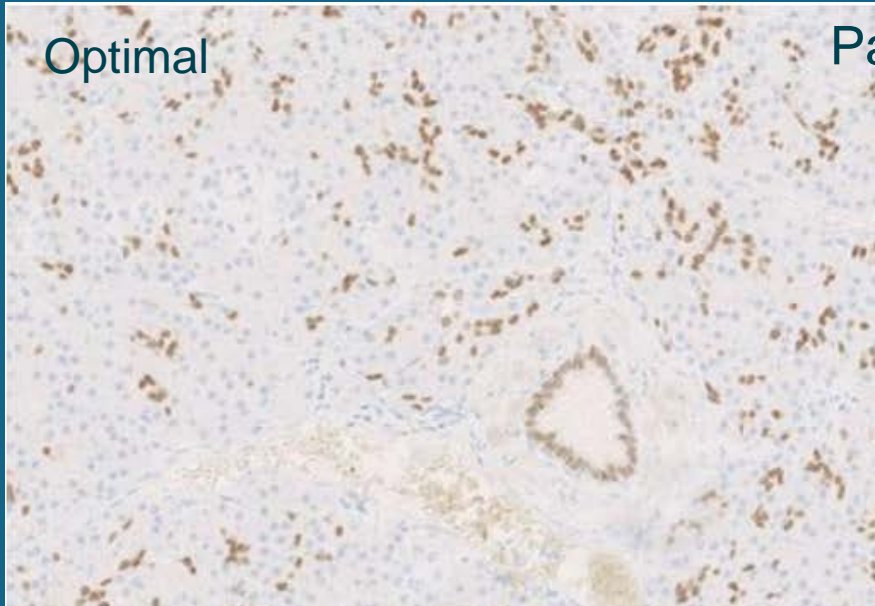
Concentrated antibodies	n	Vendor	Optimal	Good	Borderline	Poor	Suff. ¹	Suff. OPS ²
mAb clone AMT28	2	Leica/Novocastra	0	0	0	2	-	-
mAb clone CDX2-88	2	Biocare	0	0	1	3	-	-
	2	Biogenex						
mAb clone DAK-CDX2	31	Agilent/Dako	6	9	7	9	48%	57%
rmAb clone EPR2764Y	31	Cell Marque						
	5	Thermo/Neomarkers						
	4	Immunologic						
	4	Zytomed						
	2	Monosan						
	2	Zeta Corporation						
	1	Δ Menarini	28	14	7	3	81%	81%
Ready-To-Use antibodies								
mAb DAK-CDX2 IR080/IS080	34	Agilent/Dako	18	10	5	1	82%	93%
mAb DAK-CDX2 GA080	26	Agilent/Dako	16	4	3	3	77%	100%
rmAb clone EPR2764Y 760-4380/ 235R*	103	Ventana/Cell Marque	81	15	5	2	93%	96%

CDX2

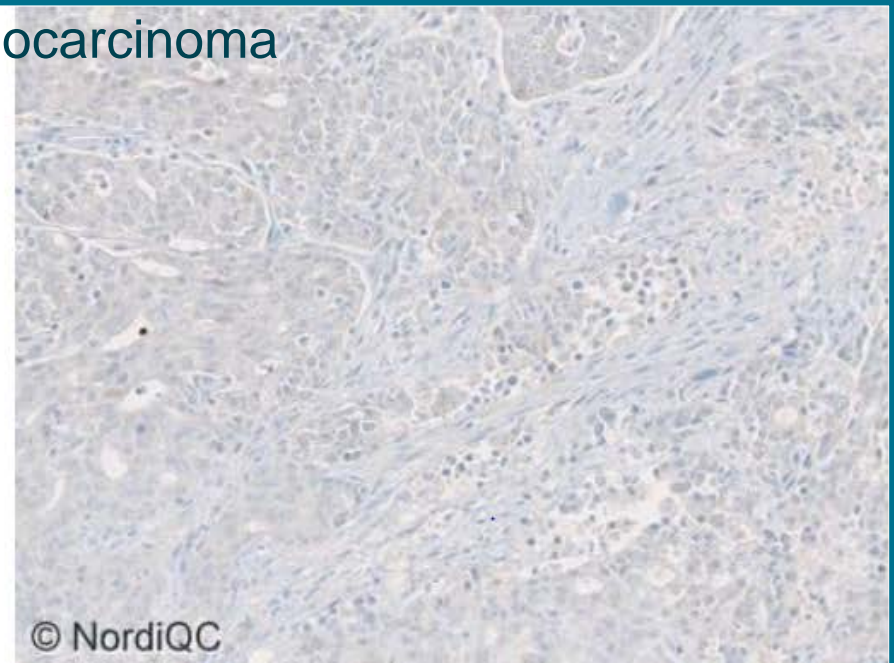
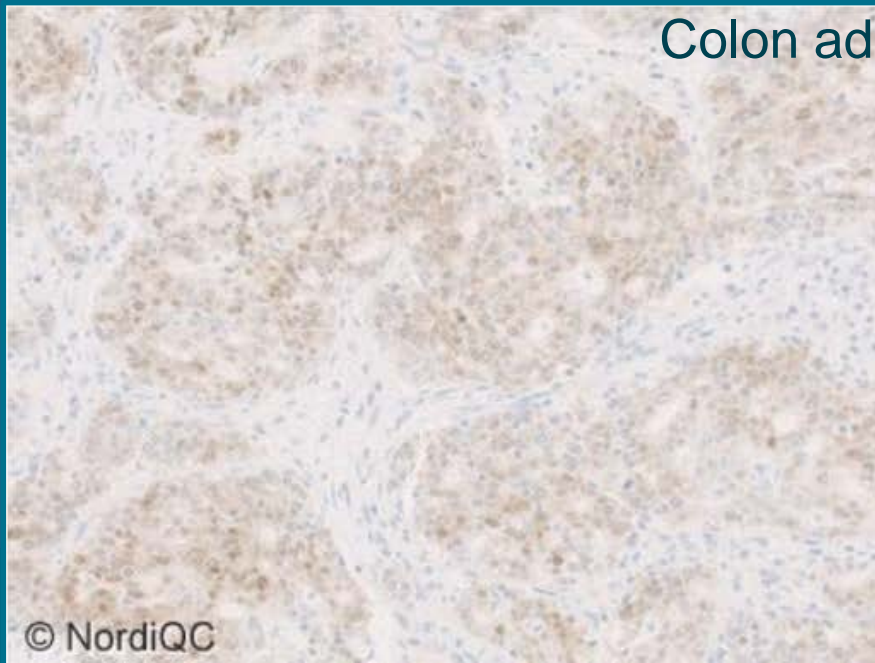
Optimal

Pancreas

Poor:



Colon adenocarcinoma



Calretinin - run 45

Table 1: **Antibodies and assessment marks for CR, run 45**

Concentrates	n	Vendor	Optimal	Good	Borderline	Poor	Suff. ¹	Suff. OPS ²
mmAb clone 2E7	1	Immunologic	1	0	0	0	-	-
mmAb clone 5A5	21	Leica/Novocastra	3	10	8	1	59%	56%
mmAb clone CAL6	1	Immunologic						
mmAb clone CAL6	6	Leica/Novocastra	4	1	0	2	71%	-
mmAb clone CAL6	1	Monosan						
mmAb clone DAK-Calret 1	35	Dako	10	13	9	3	66%	87%
mmAb clone SP13	3	Thermo/Neomarkers						
mmAb clone SP13	1	Spring Bioscience	1	2	2	1	50%	-
mmAb clone SP13	2	Cell Marque						
pAb 18-0211	16	Invitrogen/Zymed	2	8	6	0	63%	-
pAb 232A	5	Cell Marque	0	1	2	2	20%	-
pAb 61-0006	1	Genemed	0	1	0	0		

Optimal

Poor

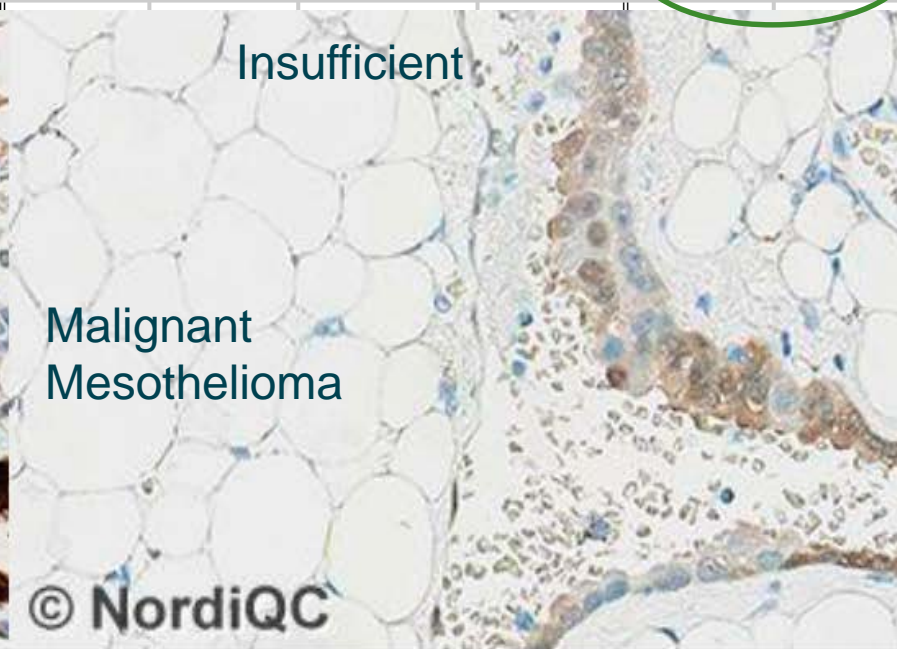
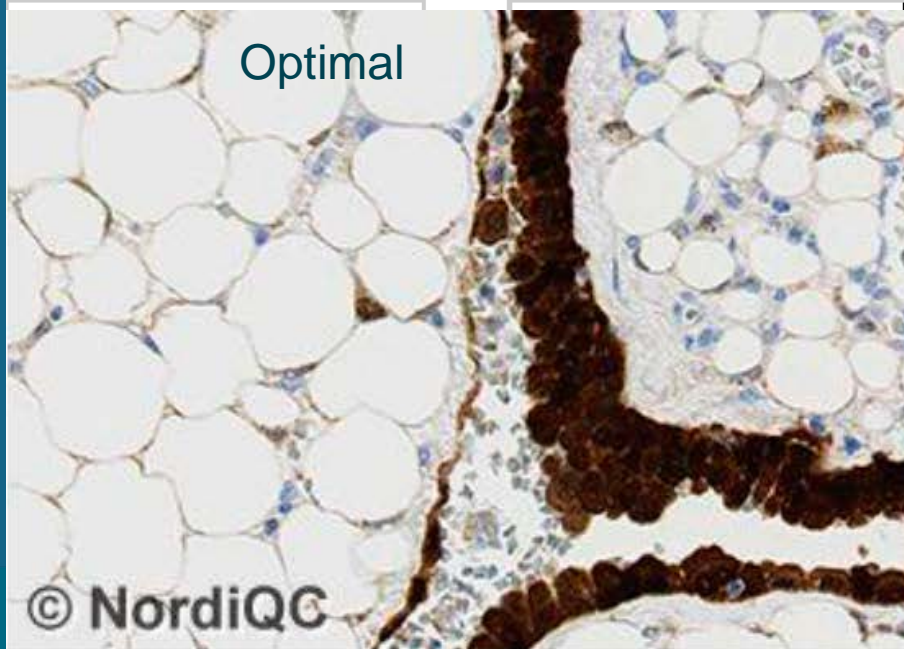
Adrenal gland

Calretinin – 113 labs using RTUs

Ready-To-Use
antibodies

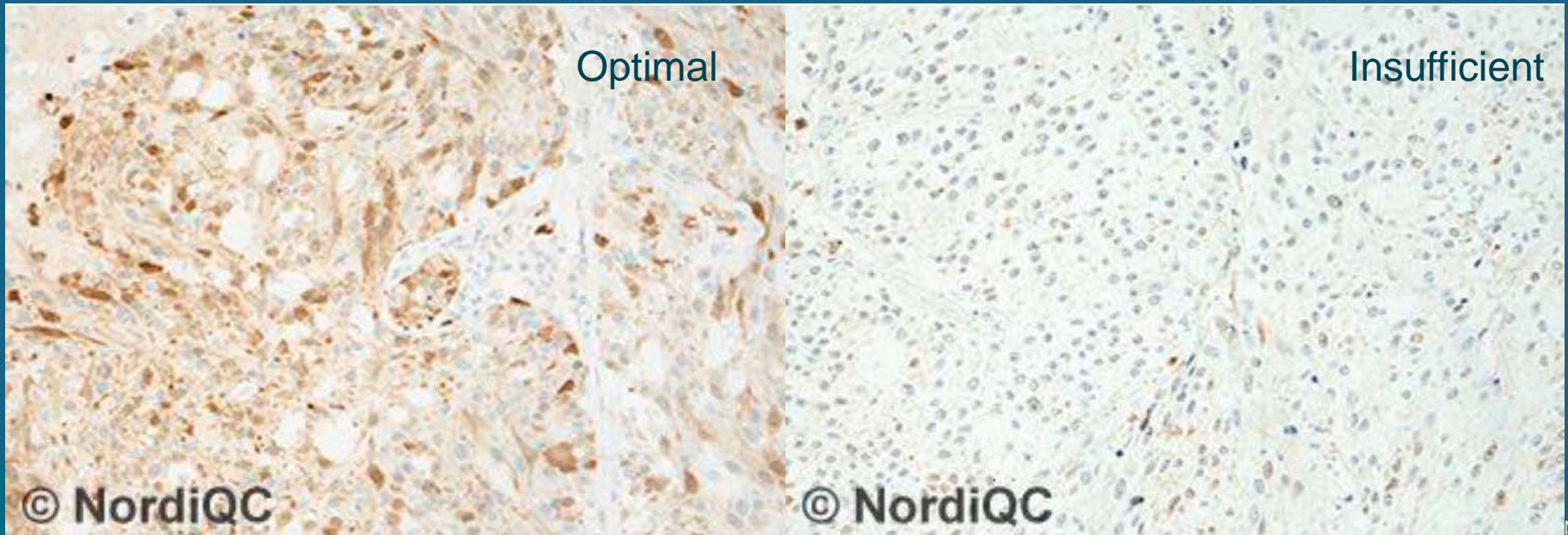


mmAb clone CAL6 PA0346	8	Leica/Novocastra	2	3	2	1	63%	-
mmAb clone DAK-Calret 1 IS/IR627	38	Dako	9	17	10	2	68%	79%
rmAb SP13 RMA-0524	1	Maixin	1	0	0	0	-	-
rmAb SP13 232R-18	1	Cell Marque	0	1	0	0	-	-
rmAb SP13 MAD-000315QD	1	Master Diagnostica	0	1	0	0	-	-
rmAb clone SP65 790-4467	64	Ventana	52	8	2	2	94%	94%



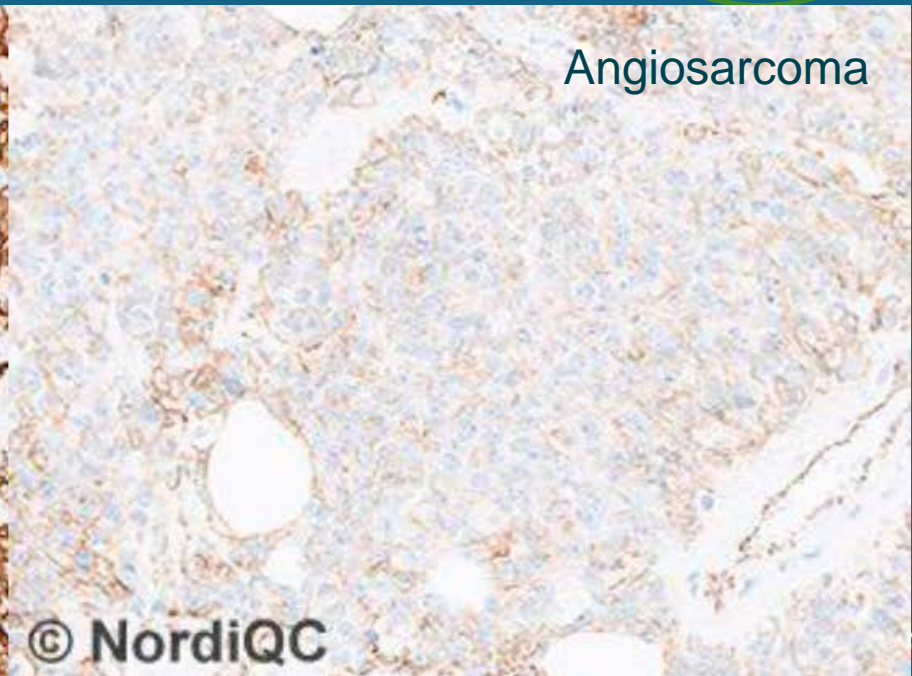
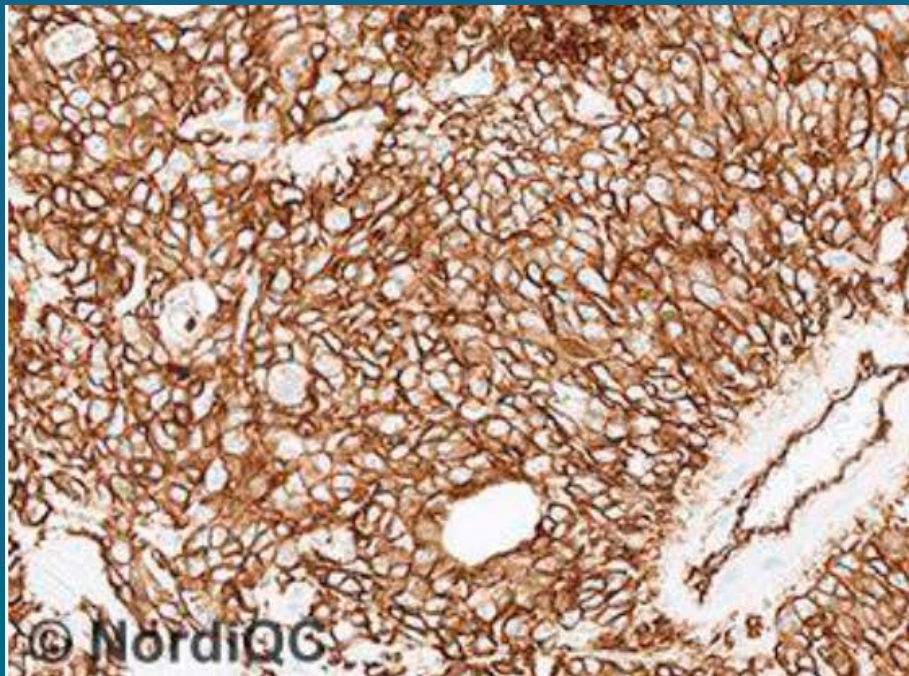
Ready-To-Use systems								
mAb clone 4C4.9 790-2914	24	Ventana	2	8	9	5	42%	100%
pAb IR504	34	Dako	3	27	4	0	88%	95%
pAb GA504	13	Dako	5	6	2	0	85%	90%
pAb 760-2523	26	Ventana	1	11	12	2	46%	100%

Mal. melanoma



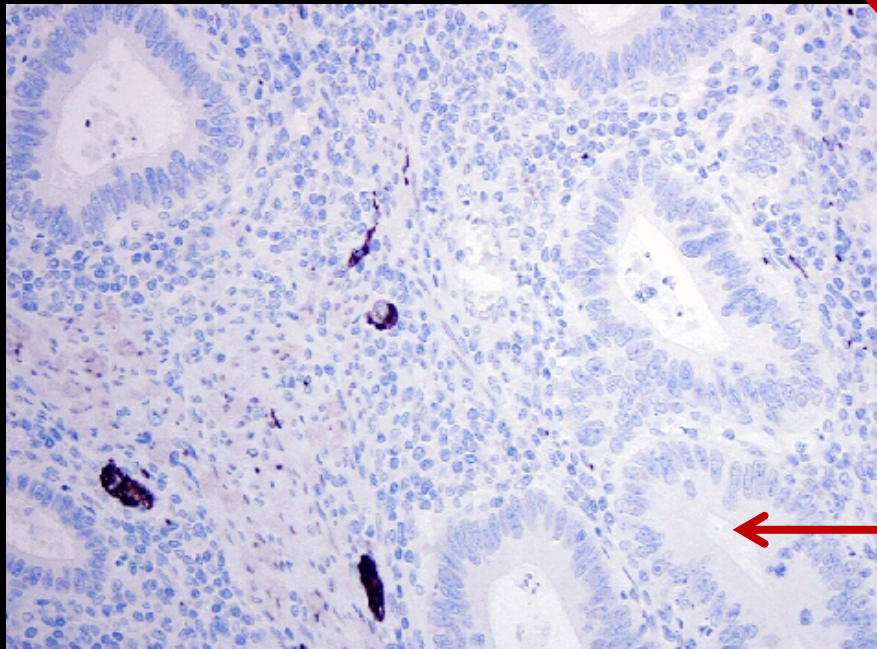
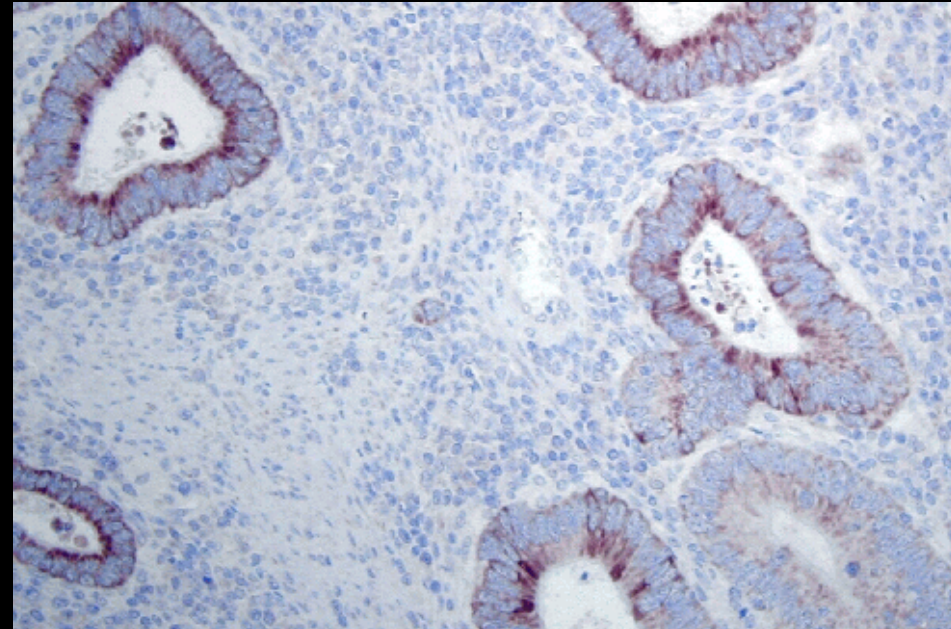
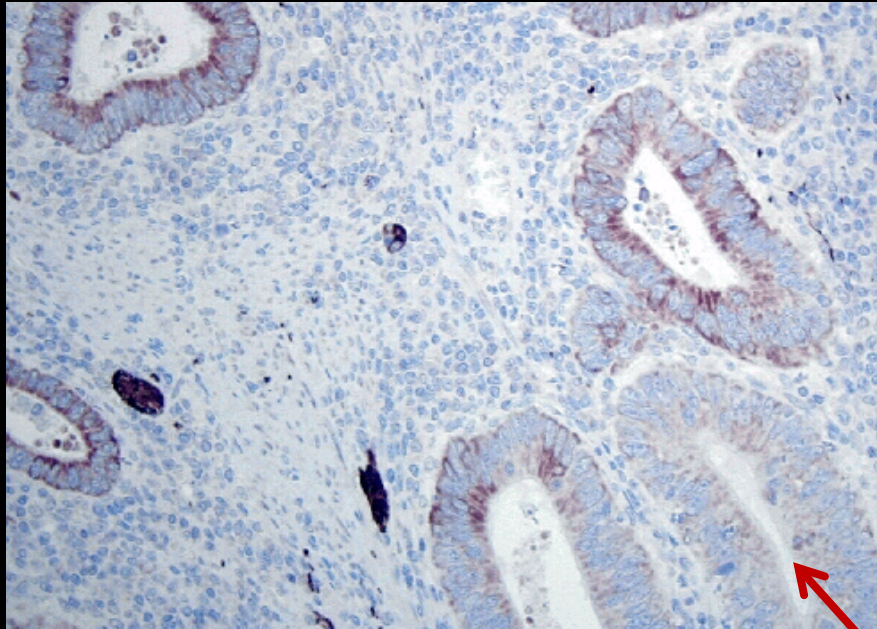
CD31

Ready-To-Use antibodies								
mAb clone 1A10 PA0250	5	Leica/Novocastra	0	0	0	5	0%	-
mAb clone BC2 PM347	1	Biocare	0	1	0	0	-	-
mAb clone JC70A 760-4378	54	Ventana/Roche	23	10	18	3	61%	67%
mAb JC70A IR/IS610	47	Dako/Agilent	27	15	3	2	89%	95%
mAb JC70A GA610	19	Dako/Agilent	15	4	0	0	100%	100%



Angiosarcoma

Biotin based system giving false positives

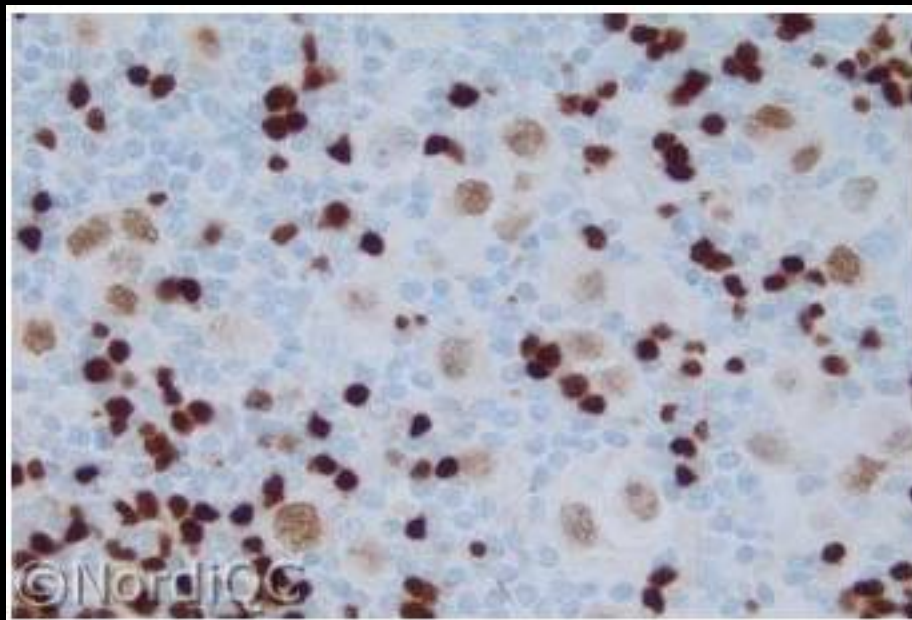


Synaptophysin
Labelled Streptavidin-Biotin system

No antibody
Labelled Streptavidin-Biotin system:
Neg. reagent control mandatory

Synaptophysin
Polymer based system

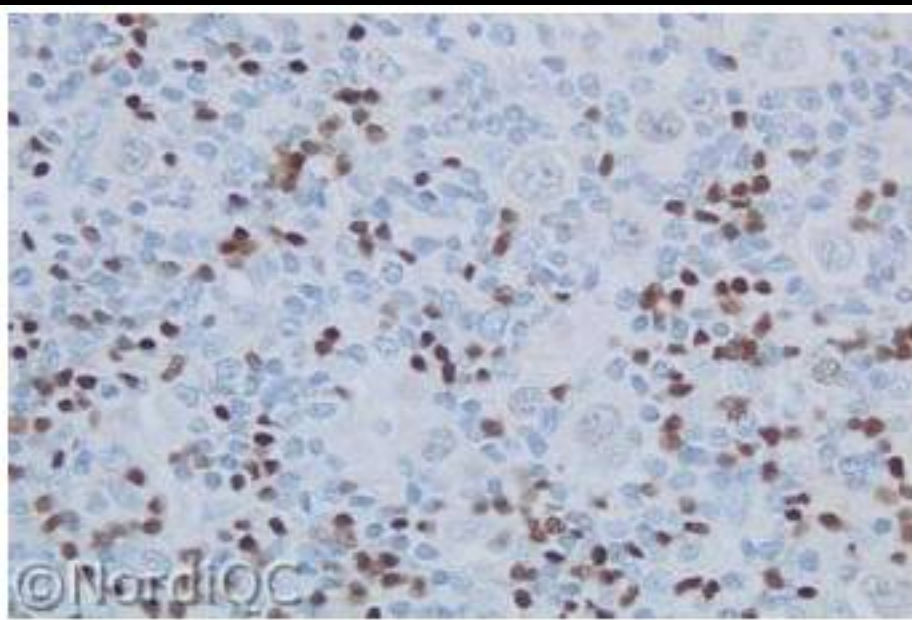
Hodgkin lymphoma NS



clone SP34

RTU VMS/CM

x200



clone 24

RTU VMS/CM

x200

Original article

Call for a European programme in external quality assurance for bone marrow immunohistochemistry; report of a European Bone Marrow Working Group pilot study

J Clin Pathol 2009;**62**:547–551.

E E Torlakovic,¹ K Naresh,² M Kremer,³ J van der Walt,⁴ E Hyjek,⁵ A Porwit⁶

Results: Eight different fixatives and nine different decalcification methods were used. While 93% of participants believed that they produced excellent results in BMTB IHC, only 4/19 (21%) laboratories did not have any poor results. CD117 and Ki-67, with 53% and 50% poor results, respectively, were the most problematic immunostains, while CD20 was the least problematic, with only 11% poor results.

- Replace less successful antibodies (conc./RTU)
- Calibrate the antibody concentration
- Use HIER (instead of proteolysis or no retrieval)
- Increase HIER time / temperature / buffer pH
 - For 95% of epitopes pH 8-9 is preferable to pH 6
- Use a non-biotin based viz. system
- Use FDA approved kits instead of home-brews
-
- Improve the internal QC: Identify the right controls –
Select well defined normal low expressor cells/tissues

419 advices for 11 markers

	<u>No.</u>	<u>Improved</u>	<u>%</u>
Positive	268	195	73
Negative	151	21	14

External Quality Assurance (EQA)

- Provides objective evidence of lab performance
- Identifies methodological errors
- Provides directions for improvements & controls

The results of the NordiQC work indicate that

- Improvement of IHC is strongly needed
- EQA schemes, industry and KOL must align - describing the requirements for optimal IHC performance.

- Almost 1/3 of all IHC stains produced by NordiQC participants are still insufficient !
 - New labs
 - New antibodies, techniques, platforms
 - Increasing demands
- How many IHC stains produced by labs not participating in an EQA scheme are insufficient ?
- How many scientific publications are based on insufficient IHC stains ?
- What are the consequences for the patients ?

JNCI Journal of the National Cancer Institute Advance Access published June 10, 2008

NEWS |

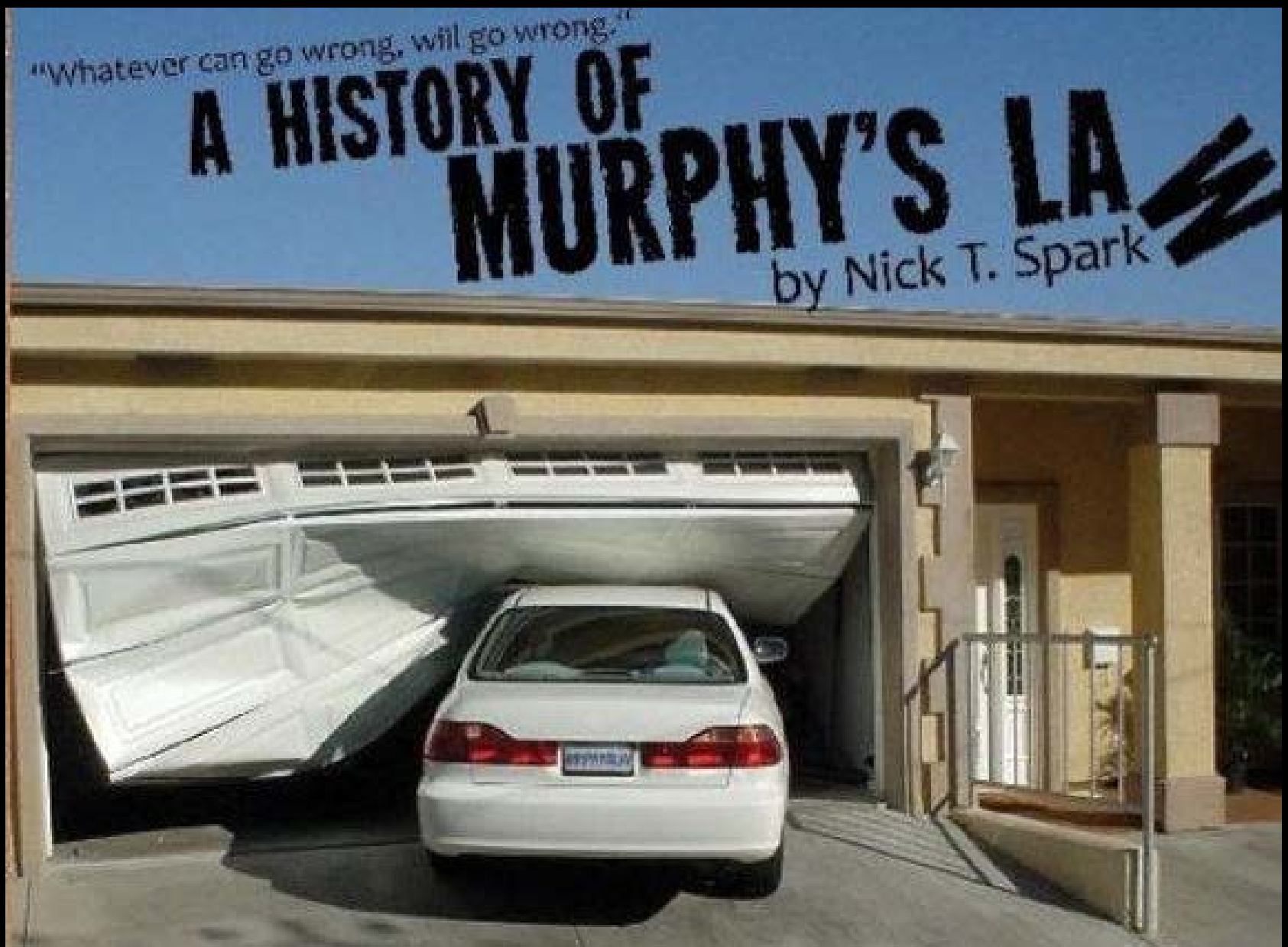
Breast Cancer Testing Scandal Shines Spotlight on Black Box of Clinical Laboratory Testing

By Karyn Hede

“Through the inquiry, the public learned that between 1997 and 2005 nearly 400 of about 1,000 breast cancer patients received incorrect test results of the ER status of their breast tumors.”

“There are no good data on the quality of ER testing in the United States. The scary thing about the debacle in Canada is that we would never have known about this if results hadn’t been checked in a central lab.”

Craig Allred



When you believe in automation and stop thinking

International Symposium on Immunohistochemistry

January 4th - 7th, 2018

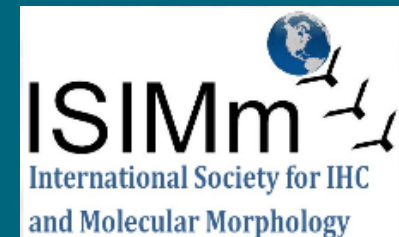
Hosted by Dept. of Histopathology, Tata Medical Center, Kolkata, India

In collaboration with NordiQC, Aalborg, Denmark and ISIMM, California, USA



The impact of proficiency testing on lab immunoassays

Thank you
for your attention



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Director of NordiQC
Aalborg University Hospital,
Aalborg, Denmark