

# QA of

# Prognostic / predictive markers in breast pathology

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## Breast panel:

- GCDFP-15
- Mammaglobin
- Gata 3
- Smooth MHCM
- ASMA
- (p63)
- E-cadherin
- p120
- ER
- PR
- HER-2

• Is it primary breast ?

• Is it invasive ?

• Is it lobular or ductal ?

• Which teraphy ?

2000 2002 2004 2006 2008 2010 2012 2014 2016

#### App. 600-700 laboratories in total







#### Assessment Run B23 2017

#### ER

#### Material

The slide to be stained for ER comprised:

No.	Tissue	ER-positivity*	ER-intensity*	
1.	Uterine cervix	80 - 90%	Moderate to strong	
2.	Tonsil	2 - 5%	Weak to strong	2 3
3.	Breast carcinoma	0%	Negative	2
4.	Breast carcinoma	90 - 100%	Moderate to strong	1 5
5.	Breast carcinoma	50 - 70%	Weak to moderate	
6.	Breast carcinoma	60 - 80%	Weak to moderate	

\*ER-status and staining pattern as characterized by NordiQC reference laboratories using the rmAb clones EP1 and SP1.

All tissues were fixed in 10% neutral buffered formalin for 24-48 hours and processed according to Yaziji et al. (1).

### Focus: Appropriate technical quality; signal-to-noise, morphology etc

Appropriate analytical sensitivity and specificity – indicated by concordance of ER status in the included tumours to reference





#### **Performance history**

This was the sixteenth NordiQC assessment of ER. The proportion of sufficient results was similar compared to the latest run (see Figure 1).



#### Fig. 1. Participant numbers and pass rates for ER during 16 runs

2003 2005 2007 2009 2011 2013 2015 2017



#### Performance history

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Estrogen receptor;

#### Fig. 1. Participant numbers and pass rates for ER during 16 runs



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	New participants	"Old" participants
Run B10, 2004	57% (n=61)	71% (n=134)
Run B15, 2010	70% (n=54)	86% (n=208)
Run B19, 2015	51% (n=86)	73% (n=259)

## NordiQC



## Estrogen receptor;

#### Pass rate influenced by protocol harmonization and availability of fully automated IHC systems

#### Performance history

This was the sixteenth NordiQC assessment of ER. The proportion of sufficient results was similar compared to the latest run (see Figure 1).



Fig. 1. Participant numbers and	pass rates for ER during 16 runs
	pace rates for an auting at rans

	2003 B8	2017 B23
Ready-To-Use format	21%	81%
HIER by in-house buffer	88%	5%
HIER by high pH	70%	94%
Polymer/multimer kit	56%	97%
Fully automated system	6%	78%







Table 1. Antibodies an	nd asse	essment marks for ER, r	un B23					
Concentrated antibodies	n	Vendor	Optimal	Good	Borderline	Poor	Suff.1	Suff. OPS <sup>2</sup>
mAb clone <b>1D5</b>	1	Dako/Agilent	0	0	0	1	-	-
mAb clone <b>6F11</b>	22 1	Leica/Novocastra Celnovte	10	11	1	1	91%	86%
rmAb clone <b>EP1</b>	15 1	Dako/Agilent Cell Marque	6	5	5	0	69%	89%
rmAb clone <b>SP1</b>	30 3 3 1	Thermo/Neomarkers Cell Marque Spring Bioscience Immunologic	27	7	2	1	92%	97%
Ready-To-Use antibodies								
mAb clone 1D5 IR/IS657	4	Dako/Agilent	о	2	2	0	-	-
mAb clone 1D5 BMS008	1	Zytomed	0	0	1	0	-	-
mAb clones 1D5 + ER-2-123 SK310	2	Dako/Agilent	o	1	0	1	-	-
mAb clone <b>6F11</b> PA0009/PA0151	14	Leica	7	6	1	0	93%	100%
rmAb <b>EP1</b> IR/IS084	59	Dako/Agilent	34	21	3	1	93%	95%
rmAb EP1 GA084	16	Dako/Agilent	12	3	1	0	94%	92%
rmAb <b>EP1</b> AN710-5M	1	Biogenex	0	1	0	0	-	-
rmAb EP1 249R-2	1	Cell Marque	o	0	1	0	-	-
rmAb clone <b>SP1</b> 790-4324/5	209	Ventana/Roche	122	79	7	1	96%	95%
rmAD clone SP1 249R-1	3	Cell Marque	o	2	1	0	-	-
rmAb clone <b>SP1</b> KIT-0012	2	Maixin	1	1	0	0	-	-
rmAb clone <b>SP1</b> <b>RMPD001</b>	1	Diagnostic Biosystems	o	0	0	1	-	-
rmAb clone <b>SP1</b> ILM30142-R25	1	Immunologic	1	0	0	0	-	-
rmAb clone <b>SP1</b> MAD-000306QD	1	Master Diagnostica	o	1	0	0	-	-
rmAb clone <b>SP1</b> M3011	1	Spring Bioscience	1	0	0	0	-	-
rmAb clone <b>SP1</b> <b>RM-9101-R7</b>	1	Thermo/Neomarkers	1	0	0	0	-	-
Total	394		222	140	25	7	-	
Proportion			56%	36%	6%	2%	92%	

HIER alk. pH 2- & 3-step kits Carefully calib.



Uterine cervix; all epithelial cells Tonsil; scattered T-cells



1) Proportion of sufficient stains (optimal or good).

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



### Estrogen receptor;



Too low titre (EP1, SP1 conc.) Insufficient HIER, Clone 1D5

Clone 6F11 by HIER at high pH, 3-step pol. (not observed on VMS)

Clone 1D5 at high titre, Biotin-based kits, HIER in pressure cooker



### Breast panel: Estrogen Receptor

Basic protocol settings for an optimal staining result (NQC)

	Retrieval	Titre	Detection	RTU	Detection
mAb 6F11*	HIER Ci, High	1:50-200	2- & 3-step	Leica	3-step
rmAb EP1	HIER High	1:25-30	2- & <u>3</u> -step	Dako	2- & <u>3</u> -step
<u>rmAb SP1</u>	HIER High	1:30-100	2- & 3-step	Ventana	<u>2</u> - & 3-step

\* Efficient HIER, high conc., 3-step pol. & low stringent washing can give aberrant nuclear staining Not seen on Ventana stainer, rarely on Autostainer and most commonly on Bond stainer.



Use uterine cervix and tonsil to verify level of sensitivity and specificity;

#### Fig. 1a

Optimal ER staining result of the uterine cervix using the rmAb clone SP1as Ready-To-Use format, Ventana 790-4324 with HIER in CC1 and UltraView as detection system. Virtually all the squamous and columnar epithelial cells show a moderate to strong, distinct nuclear staining reaction. The majority of the stromal cells are demonstrated and only endothelial and lymphoid cells are negative. Also compare with Figs. 2a – 4a, same protocol.



#### Fig. 3a

Optimal ER staining result of the breast ductal carcinoma no. 6 with 80 - 100% cells positive using same protocol as in Figs. 1a and 2a. The vast majority of the neoplastic cells show a



#### Fig 4a

Optimal ER staining result of the breast ductal carcinoma no. 4 with 40 - 60% cells positive using



Fia. 3b

Fig. 3a.

Insufficient ER staining result of the uterine cervix - same field as in Fig. 1a.

The proportion and intensity of the staining reaction in the squamous and especially in columnar epithelial cells is reduced. Also compare with Figs. 2b - 4b, same protocol.

The protocol was based on the rmAb clone EP1 applied with protocol settings giving a too low sensitivity - most likely due to a too dilute titre of the primary Ab and insufficient HIER.

ER staining result of the breast ductal carcinoma

no. 6 with 80 - 100% cells positive using same

protocol as in Figs. 1b and 2b - same field as in



Optimal ER staining result of the tonsil using same protocol as in Fig. 1a

A weak to moderate nuclear staining reaction of dispersed germinal centre lymphocytes is seen. The nuclear staining reaction can be seen at low magnification, x100.

However note that the vast majority of lymphocytes are negative.



#### Fia. 2b

Insufficient ER staining result of the tonsil using same protocol as in Fig. 1b - same field as in Fig. 2a.

Compared to the result obtained in Fig. 2a, only a faint nuclear staining reaction in a significantly reduced proportion of germinal centre lymphocytes is seen.

### Use right performance controls

No tumour material

### What is a high ER pos tumour..



Insufficient ER staining result of the breast ductal carcinoma no. 4 with 40 - 60% cells positive using





Assessment Run B20 2015 Progesterone Receptor (PR)

Recommended PR protocols

**Recommended PR control tissue** 

#### Material

The slide to be stained for PR comprised the following tissues:

No.	Tissue	PR-positivity*	PR-intensity*
1.	Uterine cervix	80 - 90 %	Moderate to strong
2. Tonsil		Negative	Negative
3. Breast carcinoma		Negative	Negative
4. Breast carcinoma		40 - 60%	Weak to strong
5. Breast carcinoma		60 - 80%	Weak to strong
6.	Breast carcinoma	80 - 100%	Moderate to strong



\*PR-positivity and intensity as characterized by NordiQC reference laboratories using the mmAb clone 16 (Leica/Novocastra)

All tissues were fixed in 10% neutral buffered formalin for 24 – 48 hours and processed according to Yaziji et al. (1).

#### Focus:

Appropriate technical quality; signal-to-noise, morphology etc

Appropriate analytical sensitivity and specificity – indicated by concordance of PR status in the included tumours to reference





#### Performance history

This was the ninth NordiQC assessment of PR. A small decrease in the proportion of sufficient results was seen compared to the previous runs, as shown in figure 1:

100% 90% 80% 70% 60% Pass rate 50% 312 Insufficient 224 415 40% 136 60 Sufficient 54 152 30% 20% 10% 0% Run 10 B2 B4 B6 B9 B12 B20 B14 B18

Figure 1 – pass rate in the 9 NordiQC assessments for PR

2004 2006 2007 2008 2010 2011 2012 2014 2015







Comparison of different antibodies for detection of progesterone receptor in breast cancer

Steroids 67 (2002) 799–813 Michael Press<sup>a,b</sup>, Betsy Spaulding<sup>c</sup>, Susan Groshen<sup>b</sup>, David Kaminsky<sup>°</sup>, Margaret Hagerty<sup>°</sup>, Lori Sherman<sup>e</sup>, Kurt Christensen<sup>e</sup>, Dean P. Edwards<sup>e</sup>;\*

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#### Potential for False-Positive Staining With a Rabbit Monoclonal Antibody to Progesterone Receptor (SP2)

Findings of the UK National External Quality Assessment Scheme for Immunocytochemistry and FISH Highlight the Need for Correct Validation of Antibodies on Introdu

Arr J Clin Pathol 2008, 129:398-409 Merdol Ibrahim, PhD,<sup>1</sup> Androw Dodson, MSc,<sup>2</sup> Sarah Barnett, MSc,<sup>1</sup> David Fish, MSc,<sup>2</sup> Bharat Jasani, PhD,<sup>4</sup> and Keith Miller, MSc<sup>1</sup>

Key Words: Breast hormonal receptors; Progesterone receptor; Rabbit monoclonal antibody; SP2; Quality assurance; Antibody validation



#### Table 1: Antibodies and assessment marks for PR, run B20

Concentrated antibodies	n	Vendor	Optimal	Good	Borderline	Poor	Suff.1	Suff. OPS <sup>2</sup>
mAb clone <b>16</b>	48 1 1	Leica/Novocastra Biocare Vector	39	7	3	1	92%	96%
mAb clone cocktail 16 + SAN27	1	Leica/Novocastra	0	1	0	0	-	-
mAb clone <b>1A6</b>	5	Leica/Novocastra	2	1	0	2	-	-
mAb clone <b>PgR 636</b>	68	Dako	48	15	4	1	93%	94%
mAb clone <b>PgR 1294</b>	17	Dako	13	4	0	0	100%	100%
rmAb clone SP2	5	Thermo/NeoMarkers	2	2	0	1	-	-
rmAb clone SP42	1	Zytomed	1	0	0	0	-	-
rmAb clone <b>Y85</b>	1 1	Cell Marque Thermo/NeoMarkers	1	1	0	0	-	-
Ready-To-Use antibodies								
mAb clone <b>16</b> PA0312	13	Leica/Novocastra	11	2	0	0	100%	100%
mAb clone 16 MAD-000670QD	2	Master Diagnostica	2	0	0	0	-	-
mAb <b>PgR 636</b> IR/IS068	78	Dako	60	15	1	2	96%	96%
mAb clone <b>PgR 1294</b> <b>K4071/SK310</b>	4	Dako	1	3	0	0	-	-
mAb clone PR88 AM328-5ME	1	Biogenex	0	0	1	0	-	-
rmAb clone <b>1E2</b> 790-2223/4296	239	Ventana	85	98	53	3	77%	88%
rmAb clone <b>SP2</b> Kit-0013	1	Maixin	1	0	0	0	-	-
rmAb clone SP2 RM-9102	1	Thermo/NeoMarkers	0	0	1	0	-	-
pAb E2071	1	Spring Bioscience	0	0	0	1	-	-
Total	489		266	149	63	11	-	
Proportion			54%	31%	13%	2%	85%	

HIER alk. pH 2- & 3-step kits Carefully calib.



Uterine cervix – all columnar epith. cells, and majority of basal squam. epith. cells (can be neg. in some pts).





1) Proportion of sufficient stains (optimal or good).

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

\*discontinued product



### Progesterone receptor;



Too low titre (16, PgR636) Insufficient HIER Clone SP2 and 1E2. 1E2 mainly by off-label protocol (ext. sensitivity) Clone 1A6, Biotin-based kits, HIER in pressure cooker



#### Fig. 1a

Optimal staining for PR of the uterine cervix using the mmAb clone 16 optimally calibrated at a titre of 1:50, efficient HIER for 48 min. in CC1 pH 8,5 using a 3-step multimer based detection system (OptiView, Ventana). The vast majority of basal squamous epithelial cells show a weak to moderate nuclear staining reaction, whereas the majority of columnar epithelial cells and stromal cells show a moderate to strong nuclear staining reaction.

#### Fig. 1b

Insufficient staining for PR of the uterine cervix, using the mmAb clone 16 with protocol settings giving a too low sensitivity - same field as in Fig. 1a. The stromal cells are demonstrated, but the squamous and columnar epithelial cells are virtually negative. Also compare with Figs. 2b - 3b - same protocol.



#### Fig. 2a

Optimal staining for PR of the breast carcinoma no. 6 with 80 - 100% cells positive using same protocol as in Fig. 1a.

A moderate to strong nuclear staining reaction is seen. A weak cytoplasmic staining reaction in the neoplastic cells is seen, but no background staining.

#### Fig. 2b

Staining for PR of the breast carcinoma no. 5 with 80 -100% cells positive using same protocol as in Fig. 1b. same field as in Fig. 2a.

A weak to strong distinct nuclear staining reaction in virtually all neoplastic cells is seen.

However also compare with Fig. 3b - same protocol.

# © NordiQC Fig. 3a

Optimal staining for PR of the breast carcinoma no. 4 with 40 - 60% cells positive using same protocol as in Figs. 1a - 2a.

The PR positive cells are easily recognized and the appropriate proportion of cells is demonstrated.

#### © NordiQC

Fig. 3b Insufficient staining for PR of the breast carcinoma no. 4 with 40 - 60% cells positive using same protocol as in Figs. 1b - 2b - same field as in Fig. 3a. Only dispersed cells are demonstrated and a significant reduced proportion of cells are identified compared to the level expected.

#### © NordiQC

#### Fig. 4a

Optimal staining for PR of tonsil using same protocol as in Figs. 1a - 3a.

No nuclear staining reaction is seen.

This staining pattern was consistently seen for protocols based on mmAb clone 1A6, 16, PgR 636 and PgR 1294 irrespective of protocol settings applied.

#### © NordiQC

#### Fig. 4b

Insufficient staining for PR of tonsil – same field as in Fig. 4a.

The majority of germinal cells show a weak to moderate and aberrant false positive nuclear staining reaction. This aberrant staining reaction was only seen for rmAb clones 1E2 (RTU, Ventana) and SP2.

For mAb clone 1E2 prolonged antibody incubation time in combination with a reduced HIER time compared to the recommendations provided by Ventana seemed to enhance the aberrant staining pattern. In case of aberrant positive nuclear staining reaction in tonsil and otherwise an expected staining pattern in the other tissues was seen, the result was evaluated as borderline. In case also an aberrant and false positive staining in the PR negative breast carcinoma was observed, the result was assessed as poor – see Fig. 5b.

#### as assessed as poor - see Fig. 5

#### © NordiQC

Fig. 5a Optimal staining for PR of the breast carcinoma no. 3 expected to be negative using same protocol as in Figs. 1a - 4a.

No nuclear staining reaction in the neoplastic cells is seen. The PR status was tested and confirmed by different Abs and protocol settings in the NordiQC reference laboratories. The tumour was ER negative. Normal glands serve as internal positive tissue control.

#### © NordiG Fig. 5b

Insufficient staining for PR of the breast ductal carcinoma no. 3 expected to be negative – same field as in Fig. 5a. Virtually all neoplastic cells show a weak to moderate and aberrant false positive nuclear staining reaction. The protocol was based on the rmAb clone SP2, using HIER in an alkaline buffer and a 2-step polymer based detection system.

### Use right performance controls





### Breast panel: Progesterone Receptor Basic protocol settings for an optimal staining result (NQC)

	Retrieval	Titre	Detection	RTU	Detection
mAb 16	HIER High	1:75-800	2- & 3-step	Leica	3-step
mAb PGR636*	HIER (High)	1:100-800	2- & 3-step	Dako	3-step
mAb PGR1294	HIER (High)	1:250-5.000	2- & 3-step	Dako	2-step
rmAb 1E2**	HIER High	-	-	Ventana	2-step

mAb clone PGR636 has shown to be less successful on Ventana BenchMark Ultra

\*\* rmAb clone 1E2, RTU might provide aberrant false pos. result by 3-step protocols, reduced HIER and prolonged Ab incubation time compared to Ventana guidelines



Use uterine cervix and tonsil to verify level of sensitivity and specificity;





Use metrics

% of ER pos % of PR pos

Some types <u>must</u> be pos (Ductal I, Lobular etc)





#### Assessment Run B23 2017 HER-2 IHC

#### Material

The slide to be stained for HER-2 comprised the following 9 materials:

	IHC: HER-2 Score* (0, 1+, 2+, 3+)	FISH: HER-2/chr17 ratio**
1. Cell line 1, Horizon Discovery***	3+	
2. Cell line 2, Horizon Discovery***	2+	
3. Cell line 3, Horizon Discovery***	1+	
<ol> <li>Cell line 4, Horizon Discovery***</li> </ol>	0	
5. Breast carcinoma, no. 1	3+	> 6.0 (clusters) (amplified)
6. Breast carcinoma, no. 2	2+	2.3 – 2.9 (amplified)
7. Breast carcinoma, no. 3	0-1+	1.1 – 1.4 (unamplified)
8. Breast carcinoma, no. 4	1-2+	1.3 – 1.7 (unamplified)
9. Breast carcinoma, no. 5	0-1+	1.2 – 1.4 (unamplified)

 \* HER-2 immunohistochemical score (see table below) as achieved by using the three FDA approved kits and antibodies, HercepTest™ (Dako), Oracle™ (Leica) and PATHWAY<sup>®</sup> (Ventana), in NordiQC reference laboratories.
 \*\* HER-2/chr17 ratios achieved using Zyto*Light* <sup>®</sup> SPEC HER2/CEN 17 Dual Color FISH (Zytovision)

\*\*\* The cell lines were not included in the assessment. Data will be analyzed subsequently by digital image analysis.

All carcinomas were fixed for 24 - 48 h in 10% neutral buffered formalin.

### Focus: Appropriate technical quality; signal-to-noise, morphology etc

Appropriate analytical sensitivity and specificity – indicated by concordance to FISH status and IHC level established by reference data in all the included tumours.



Table 1. Assessment (	Table 1. Assessment marks for IHC assays and antibodies run B23, HER-2 IHC							
FDA approved HER-2 assays	n	Vendor	Optimal	Good	Borderline	Poor	Suff.1	Suff. OPS <sup>2</sup>
PATHWAY <sup>®</sup> rmAb clone <b>4B5, 790-2991</b>	204	Ventana/Roche	186	13	2	3	98%	98%
CONFIRM™, rmAb clone <b>4B5, 790-4493</b>	15	Ventana/Roche	12	3	0	0	100%	100%
HercepTest™ <b>SK001</b>	33	Dako/Agilent	32	0	0	1	97%	97%
HercepTest™ <b>SK001</b> <sup>4</sup>	8	Dako/Agilent	8	0	0	0	100%	-
HercepTest™ K5207	2	Dako/Agilent	2	0	0	0	-	-
HercepTest™ K5204	2	Dako/Agilent	1	1	0	0	-	-
Oracle™ mAb clone CB11, TA9145	9	Leica	4	4	0	1	89%	88%
Antibodies <sup>3</sup> for laboratory developed HER-2 assays, conc. antibody	n	Vendor	Optimal	Good	Borderline	Poor	Suff.1	Suff. OPS <sup>2</sup>
mAb clone <b>BS24</b>	1	Nordic Biosite	0	1	0	0	-	-
mAb clone <b>CB11</b>	7 1	Leica/Novocastra Biogenex	4	4	0	0	100%	100%
rmAb clone <b>EP3</b>	1 1 1 1	Biocare Cell Marque Celnovte Thermo/NeoMarkers PathnSitu	3	2	0	0	100%	100%
rmAb clone <b>SP3</b>	17 3 1 1 1	Thermo/NeoMarkers Zytomed Cell Marque Spring Bioscience Thermo/Pierce	12	8	0	3	87%	100%
pAb clone A0485	50	Dako/Agilent	35	11	1	3	92%	93%
Antibodies for laboratory developed HER-2 assays, RTU	n	Vendor	Optimal	Good	Borderline	Poor	Suff.1	Suff. OPS²
mAb clone CB11, NCL-L-CB11	4	Leica/Novocastra	0	0	3	1	-	-
mAb clone CB11, BMS014	1	Zytomed	0	0	1	0	-	-
rmAb clone EP3, AN726	1	Biogenex	1	0	0	0	-	-
rmAb clone EP3, RMPD049R	1	Diagnostic Biosystems	1	0	0	0	-	-
rmAb clone SP3, 237R	2	Cell Marque	1	1	0	0	-	-
rmAb clone SP3, MAD-000308QD	1	Master Diagnostics	1	0	0	0	-	-
Ab clone MXR001, RMA-0701	2	Maixin	1	1	0	0	-	-
Total	372		304	49	7	12	-	-
Proportion			82%	13%	2%	3%	95%	-



	IHC: HER-2 Score* (0, 1+, 2+, 3+)
1.Breast carcinoma	2-3+
2.Breast carcinoma	0-1+
3.Breast carcinoma	1-2+
4.Breast carcinoma	3+
5.Breast carcinoma	0-1+***
	FISH: HER-2 gene/chr
	17 ratio**
1.Breast carcinoma	2.3 - 2.8 (a)
2.Breast carcinoma	0.9 - 1.3 (u)
3.Breast carcinoma	1.2 - 1.5 (u)
4.Breast carcinoma	> 6.0 (clusters) (a)
5.Breast carcinoma	1.2 – 1.5 (u)

### Material processed according to ASCO/CAP

JOURNAL OF CLINICAL ONCOLOGY

ASCO SPECIAL ARTICL

Recommendations for Human Epidermal Growth Factor Receptor 2 Testing in Breast Cancer: American Society of Clinical Oncology/College of American Pathologists Clinica Practice Guideline Update

Antonio C. Wolff, M. Elizabeth H. Hammond, "David G. Hicks," Mitch Dovesett, "Lisa M. McShane," Komberly H. Allison, Donald C. Albed, John M.S. Barden, Michael Bilson, Patrick Fitzgöbern, Wolda J. Robert B. Jenkon, Pannela B. Mango, Sconwrynny Path, Edith A. Perez, Michael F. Prens, Patricia A. Spe

1) Proportion of sufficient stains (optimal or good).

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

3) mAb: mouse monoclonal antibody, rmAb: rabbit monoclonal antibody, pAb: polyclonal antibody.

4) RTU system developed for the Agilent/Dako's semi-automated systems (Autostainer Link48) but used by laboratories on different platforms (Leica Bond and Dako Omnis)











#### Figure 1. Pass rates of 23 HER-2 IHC assessments in the NordiQC breast cancer module



App 90 % of insuff. results are FN and seen both by FDA / CE-IVD kits and laboratory developed assays.

FP results have virtually only been seen by laboratory developed assays.



Typical causes for insufficient results in the NordiQC HER2 IHC breast module:

FDA / CE-IVD HER2 IHC kits:

PATHWAY<sup>®</sup>, Ventana: Too short HIER (<24M) and/or too short incubation of primary Ab (<12M)

HercepTest<sup>™</sup>, Dako: Too short HIER (<40M) and/or too short incubation of primary & secondary Ab (<30M)

Oracle<sup>™</sup>, Leica: No single or combination of causes have been identified



Laboratory developed assays:

Inappropriate titre of primary Ab, less successful primary Ab, insufficient HIER, etc.....













Assessment Run H11 2017 HER-2 ISH

#### Material

#### Table 1. Content of the multi-block used for the NordiQC HER-2 ISH assessment, run H11

	HER-2 IHC*	Dual - SISH**	FISH***	FISH***
	IHC score	HER-2/chr17 ratio¤	HER-2/chr17 ratio¤	HER-2 copies
1. Breast carcinoma	0	0.8	0.8 - 1.0	< 4
2. Breast carcinoma	2+	2.3	2.8 - 3.3	> 6
3. Breast carcinoma	3+	8.0	6.5 - 8.5	> 6
4. Breast carcinoma	2+	1.1	1.0 - 1.2	$\geq$ 4 and < 6
5. Breast carcinoma	1+	1.5	1.2 - 1.5	< 4

\* PATHWAY® (Ventana), data from two reference labs.

\*\* Inform HER-2 Dual ISH kit (Ventana), range of data from one reference lab.

\*\*\* HER-2 FISH pharmDX™ Kit (Dako) and HER-2 FISH (Zytovision), range of data from one reference lab. ×HER-2/chr17: HER-2 gene/chromosome 17 ratio

All tissues were fixed for 24 - 48 hours in 10% neutral buffered formalin according to the ASCO/CAP 2013 guidelines for tissue preparation of breast tissue for HER-2 ISH analysis.

#### HER-2 BRISH, Technical assessment

The main criteria for assessing a BRISH HER-2 analysis as technically optimal were the ability to interpret the signals and thus evaluate the HER-2/chr17 ratios in all five tissues.

Staining was assessed as good, if the HER-2/chr17 ratios could be evaluated in all five tissues, but the interpretation was slightly compromised e.g. due to excessive retrieval, weak or excessive counterstaining or focal negative areas.

Staining was assessed as borderline if one of the tissues could not be evaluated properly e.g. due to weak signals, large negative areas with no signals (> 25% of the core) or a low signal-to-noise ratio due to excessive background staining.

Staining was assessed as **poor** if two or more of the tissue cores could not be evaluated properly e.g. due to weak signals, large negative areas with no signals (> 25% of the core) or a low signal-to-noise ratio due to excessive background staining.

#### HER-2 BRISH and FISH interpretation

For both BRISH and FISH, participating laboratories were asked to submit a scoring sheet with their interpretation of the HER-2/chr17 ratio. Results were compared to NordiQC FISH data from reference laboratories to analyse scoring consensus.

Consensus scores from the NordiQC FISH reference laboratories

- Breast ductal carcinomas, no. 1 and 5: non-amplified
- Breast ductal carcinomas, no. 4: non-amplified or equivocal
- Breast ductal carcinoma no. 2 and 3: amplified

The ASCO/CAP 2013 guidelines were applied for the interpretation of the HER-2 status

Unamplified: HER-2/chr17 ratio < 2.0 using a dual probe assay or an average < 4 HER-2 gene copies per cell/nucleus (both dual and single probe assay)

Equivocal: HER-2/chr17 ratio of < 2.0 using a dual probe assay with an average of ≥ 4 and < 6 HER-2 gene copies per cell/nucleus (both dual and single probe assay)

Amplified: HER-2/chr17 ratio  $\geq$  2.0 using a dual probe assay or an average  $\geq$  4 HER-2 copies per cell/nucleus. Using a single probe assay an average of  $\geq$  6 HER-2 copies per cell/nucleus.



#### Participation

Number of laboratories registered for HER-2 BRISH	124
Number of laboratories returning slides	121 (98%)
Number of laboratories returning scoring sheet	112 (90%)
Number of laboratories registered for HER-2 FISH	64
Number of laboratories returning scoring sheet	54 (84%)

#### **Results BRISH**, technical assessment

In total, 121 laboratories participated in this assessment. 73 laboratories (60%) achieved a sufficient mark (optimal or good). Results are summarized in table 2.

#### Table 2. HER-2 BRISH systems and assessment marks for BRISH HER-2 run H11.

Two colour HER-2 systems	n	Vendor	Optimal	Good	Borderline	Poor	Suff.1	Suff. OPS <sup>2</sup>
INFORM™ HER-2 Dual ISH 800-4422	91	Ventana	32	24	15	20	62%	65%
INFORM™ HER-2 Dual ISH + IHC 800-4422 + HER-2 IHC	14	Ventana	3	4	6	1	50%	58%
DuoCISH pharmDx™ SK109	3	Dako	1	1	1	0	-	-
ZytoDot® 2C C-3022 / C-3032	5	ZytoVision	2	2	1	0	80%	100%
One colour HER-2 systems								
INFORM™ HER-2 SISH 780-4332	3	Ventana	0	2	0	1	-	-
ZytoDot® C-3003	5	ZytoVision	1	1	1	2	40%	75%
Total	121		39	34	24	24	60%	-
Proportion			32%	28%	20%	20%		

1) Proportion of sufficient stains.

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



Technically optimal results in the NordiQC HER2 ISH breast module: INFORM<sup>™</sup> HER2 Dual ISH, Ventana Zyto*Dot*<sup>®</sup> 2C, ZytoVision











Typical causes for insufficient results in the NordiQC HER2 ISH breast module:

FDA / CE-IVD HER2 BRISH (CISH/DDISH/etc) kits:

INFORM<sup>™</sup> HER2 Dual ISH, Ventana: Excessive proteolysis (>16M), HIER in CC1.

DuoCISH<sup>™</sup> pharmDx<sup>™</sup>, Dako: Insufficient proteolysis, inappropriate handling of chromogen.

Zyto*Dot*<sup>®</sup> 2C, ZytoVision: Excessive proteolysis.

In 90% of insufficient results, no single or combination of causes could be identified



Technically insufficient results in the NordiQC HER2 ISH breast module: INFORM<sup>™</sup> HER2 Dual ISH, Ventana









#### HER-2 BRISH, Technical assessment

The main criteria for assessing a BRISH HER-2 analysis as technically **optimal** were the ability to interpret the signals and thus evaluate the HER2/chr17 ratios in all five tissues.

Graph 1. Proportion of sufficient results for HER-2 BRISH in the NordiQC assessment



C1 C2 B9 B10 B11 B12 H1 H2 H3 H4 H5 H6 H7 H8 H9 H10 H11 H12



HER2 Gene-Protein-Assay (Roche): HER2 IHC + DDISH – lim. EQA data









#### Participation

Number of laboratories registered for HER-2 BRISH	124
Number of laboratories returning slides	115 (93%)
Number of laboratories returning scoring sheet	105 (92%)
Number of laboratories registered for HER-2 FISH	60
Number of laboratories returning scoring sheet	56 (93%)

### Latest Run

#### **Results BRISH, technical assessment**

In total, 115 laboratories participated in this assessment. 73 laboratories (64%) achieved a sufficient mark (optimal or good). Results are summarized in Table 2.

Two colour HER-2 systems	n	Vendor	Optimal	Good	Borderline	Poor	Suff.1	Suff. OPS <sup>2</sup>
INFORM™ HER-2 Dual ISH <b>800-4422</b>	83	Ventana	28	23	19	13	61%	64%
INFORM™ HER-2 Dual ISH + IHC 800-4422 + HER2 IHC	17	Ventana	11	5	1	0	94%	100%
Zyto <i>Dot®</i> 2C C-3022 / C-3032	6	ZytoVision	2	1	1	2	50%	50%
One colour HER-2 systems								
INFORM™ HER-2 SISH <b>780-4332</b>	4	Ventana	1	0	0	3	25%	-
Zyto <i>Dot®</i> C-3003	5	ZytoVision	1	2	1	1	<mark>60%</mark>	60%
Total	115		43	31	22	19	64%	-
Proportion			37%	27%	19%	17%		

#### Table 2. HER-2 BRISH systems and assessment marks for BRISH HER-2 run H12.

1) Proportion of sufficient stains.

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



Central issues to adress for control material of HER2 IHC test

1. External control – no useful internal HER2 expression

2. Must be processed under similar conditions as patient material – otherwise documented that identical results are generated by other methods (fixative, duration, etc)

3. Must reflect the range of HER2 expression seen for diagnostics

4. Consistent and stable HER2 expression throughout the control material

## HER2 – NordiQC perspective



### Central issues to adress for control material of HER2 IHC test In NordiQC app. 60-70% of laboratories use a 3+ tumour as routine positive control for HER2 IHC



Question: Is this reliable to monitor a consistent level of HER2 assay ?

### HER2 – NordiQC perspective





Ampl. 3+

Ampl. 2+

Unampl. 2+

Unampl. 0

### HER2 – NordiQC perspective





Optimally:

Use small TMA with 1+, <u>2+</u> & 3+ mounted on same slide as pt material for daily control of HER2 IHC assay



Control material for HER2 IHC: performace control / consistency

Histology: 3+ tumour

#### <u>2+ tumour</u>



Cell lines: 3+

<u>2+</u>

Applicable for DIA & ref data comparing run-to-run



### Control material for HER2 IHC: performace control / consistency Histocyte cell lines HER2: PATHWAY IHC



Cell line 3 - 1 +





Control material for HER2 IHC: performace control / consistency Histocyte cell lines HER2: PATHWAY IHC





Connectivity	DIA score
≤ 0.40	0/1+
> 0.40 - ≤ 0.64	2+
> 0.64	3+

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Control material for HER2 IHC: performance control / consistency Histocyte cell lines HER2: Horizon cell lines HER2



Pathway Oracle Herceptest Pathway Oracle HercepTest

### IHC - Protocols and controls for Breas



Connectivity	DIA score
≤ 0.40	0/1+
> 0.40 - ≤ 0.64	2+
> 0.64	3+

Reference values Tolerance criteria If ok in cell line – Tumour can ce scored

PATHWAY Run 1 – Opt PATHWAY Run. 2 – Opt.

### IHC - Protocols and controls for Breas

Connectivity	DIA score
≤ 0.40	0/1+
> 0.40 - ≤ 0.64	2+
> 0.64	3+

Reference values Tolerance criteria If ok in cell line – Tumour can ce scored



### IHC - Protocols and controls for Breas

Connect: 0,5	Connect: 0,7
Breast carc. – 2+ (A)	
Connect: 0,2	Connect: 0,5
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PATHWAY Run 1 – Opt

PATHWAY Run 2 – FP

Connectivity	DIA score
≤ 0.40	0/1+
> 0.40 - ≤ 0.64	2+
> 0.64	3+

Reference values Tolerance criteria If ok in cell line – Tumour can ce scored



The combination of cell lines and DIA might be useful for;

### IHC assay reproducibility / consistency Run-to-Run

Implementation of IHC assay (kits) for verification

Require reference values for sufficient versus insufficient test

Potential for EQA for IHC assays (kits) – LDTS to be confirmed



Conclusions:

1. Pass rates for ER, PR and HER2 IHC are improved. Robust clones, high quality IHC systems.

2. CE-IVD labelled RTU assays / systems have shown superior performance compared to laboratory developed assays.

3. HER2 BRISH (DDISH/SISH/CISH) results have not been improved.



