

# NORDIQC DATA FOR BREAST MARKERS

Antibody selection, protocols and controls

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# NORDIQC EQA DATA FOR IHC BREAST MARKERS

Type I IHC tests

Type II IHC tests

		Purpose	Last run	Pass rate	No of labs
Type I IHC tests	GATA3	<u>Breast</u> vs non-breast	Run 63, 2021	68%	320
	Mammaglobin	<u>Breast</u> vs non-breast	Run 25, 2009	83%	23
	GCDFP15	<u>Breast</u> vs non-breast	Run 36, 2012	86%	131
	CK5	CIS vs <u>invasive</u>	Run 65, 2022	71%	311
	SMH	<u>CIS</u> vs invasive	Run 66, 2022	81%	152
	p63	CIS vs <u>invasive</u>	Run 61, 2021	79%	324
	E-Cadherin	<u>Ductal</u> vs lobular	Run 53, 2018	89%	298
Type II IHC tests	KI67	PI index	Run B22, 2016	93%	409
	ER	Predictive for Tamoxifen	Run B35, 2023	91%	422
	PR	Predictive for Tamoxifen	Run B35, 2023	92%	414
	HER2 IHC	Predictive for Herceptin	Run B35, 2023	90%	403
	HER2 BRISH	Predictive for Herceptin	Run H23, 2023	59%	163
	PD-L1 IC	Predictive for Tecentriq	Run C13, 2023	68%	139
	PD-L1 TPS/CPS	Predictive for Keytruda	Run C13, 2023	92%	243



# KEY-POINTS FOR BEST PROTOCOLS

- Clone selection
- RTUs – “Plug and Play” or “Play and Plug”?
- Efficient HIER, preferable in an alkaline buffer
- Use of right detection system
- Use of iCAPS



# CLONE PERFORMANCE FOR SELECTED BREAST MARKERS

Marker	Successful clones	Less successful clones
GATA3	mAb L50-823, rmAb SP368	mAb HG3-31
CK5*	mAb XM26, rmAb SP27	mAb D5/16 B4
SMH	mAb SMMS1	-
p63	mAbs 4A4 & DAK-p63	mAb 7JUL
E-Cadherin	mAbs NCH-38, 36 & 36B5	rmAb EP700Y
KI67	mAb MIB-1, rmAb 30.9	-
ER	rmAbs SP1 & EP1, mAb 6F11	-
PR	mAbs 16 & PgR1294, rmAbs 1E2 & Y85	-
HER2 IHC	rmAbs 4B5 & DG44, Dako pAb	mAb CB11
PD-L1 IC	rmAb SP142	Non-SP142
PD-L1 TPS/CPS*	mAb 22C3, rmAb SP263	rmAb SP142



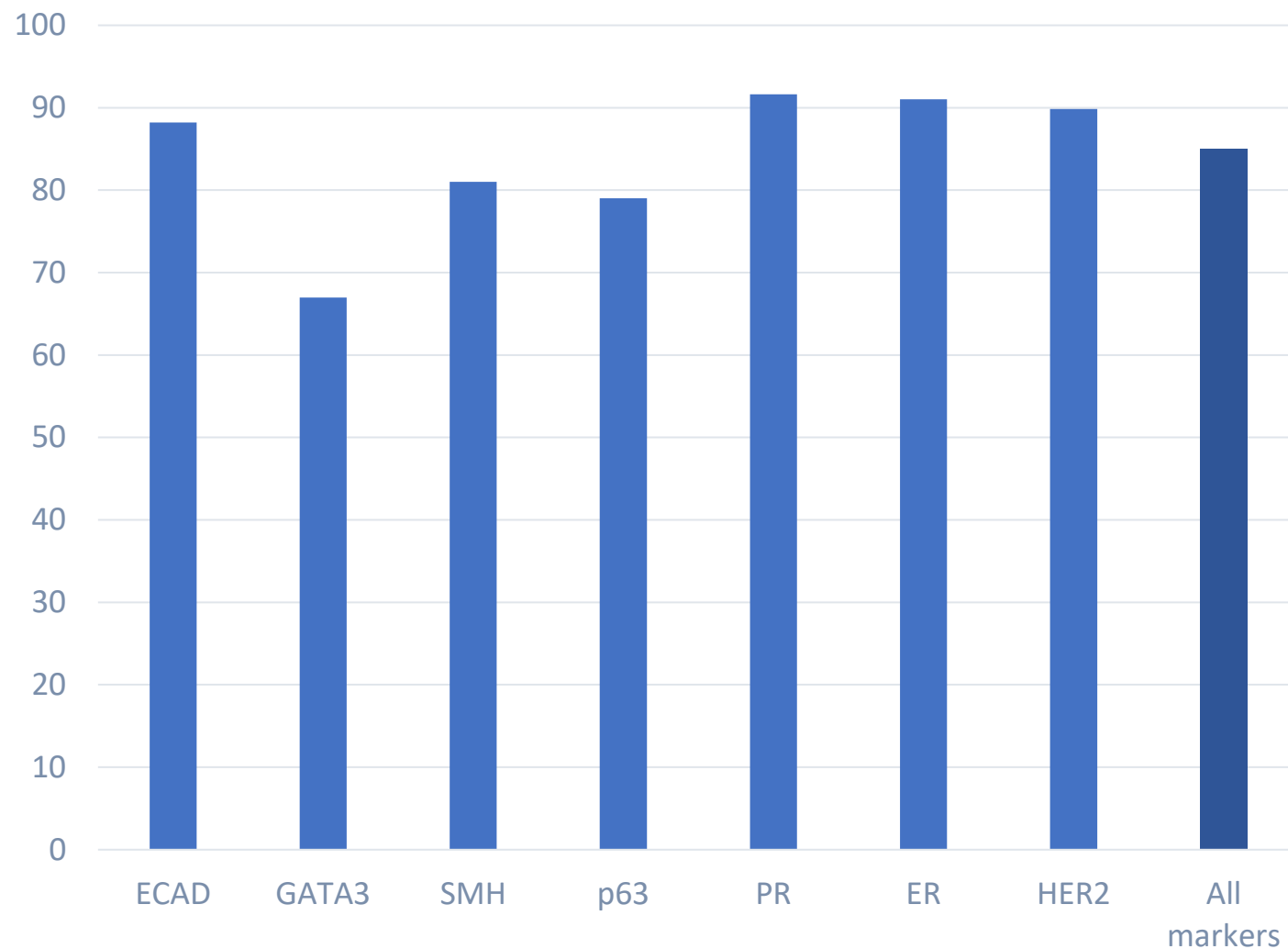
\*see ppt for lung-markers

# ICAPS FOR SELECTED BREAST MARKERS

Marker	IHC critical assay performance controls Low expression	Negative tissue controls No expression	
GATA3	Tonsil: T-helper-cells in the T-zones and germinal centers.	Tonsil: B-cells, squamous epithelial cells, endothelial cells.	<a href="#">Link</a>
Mammaglobin	Skin: Epithelial cells of eccrine sweat glands.	Tonsil: All cell types.	<a href="#">Link</a>
GCDFP15	Skin: Epithelial cells of eccrine sweat glands.	Tonsil: All cell types.	<a href="#">Link</a>
CK5	Pancreas: Scattered epithelial cells of intercalated ducts.	Liver: All cell types.	<a href="#">Link</a>
Smooth MHCM	Tonsil: Follicular dendritic cells in germinal centers.	Tonsil: Epithelial cells.	<a href="#">Link</a>
p63	Placenta: Cytotrophoblastic cells.	Appendix: Epithelial- and smooth muscle cells.	<a href="#">Link</a>
E-Cadherin	Liver: Hepatocytes.	Appendix: Stromal cells, smooth muscle cells, endothelial cells.	<a href="#">Link</a>
KI67	Tonsil: B-cells in the light zones of the germinal centers.	Liver: Hepatocytes	<a href="#">Link</a>
ER	Tonsil: Squamous epithelial cells, T-cells in germinal centres.	Tonsil: B-cells in mantle zone and germinal centres.	<a href="#">Link</a>
PR	Cervix: Basal squamous epithelial cells.	Tonsil: All cells types (especially focus on lymphocytes in germinal centres).	<a href="#">Link</a>
PD-L1 IC	Tonsil: T-cells and macrophages in germinal centres.	Tonsil: Normal squamous epithelial cells, lymphocytes.	<a href="#">Link</a>
PD-L1 TPS/CPS	Tonsil: Germinal center macrophages and T-cells.	Tonsil: Stratified normal squamous epithelial cells and vast majority of lymphocytes.	<a href="#">Link</a>



Pass rates for selected breast markers



## KEY-POINTS FOR BEST PROTOCOLS

### Selected breast markers:

Overall pass rate: **85%**  
(2.078/2.450), ranging from 67%  
for GATA3 till 92% for PR.

# KEY-POINTS FOR BEST PROTOCOLS

## Breast markers:

Overall pass rate: **85%**

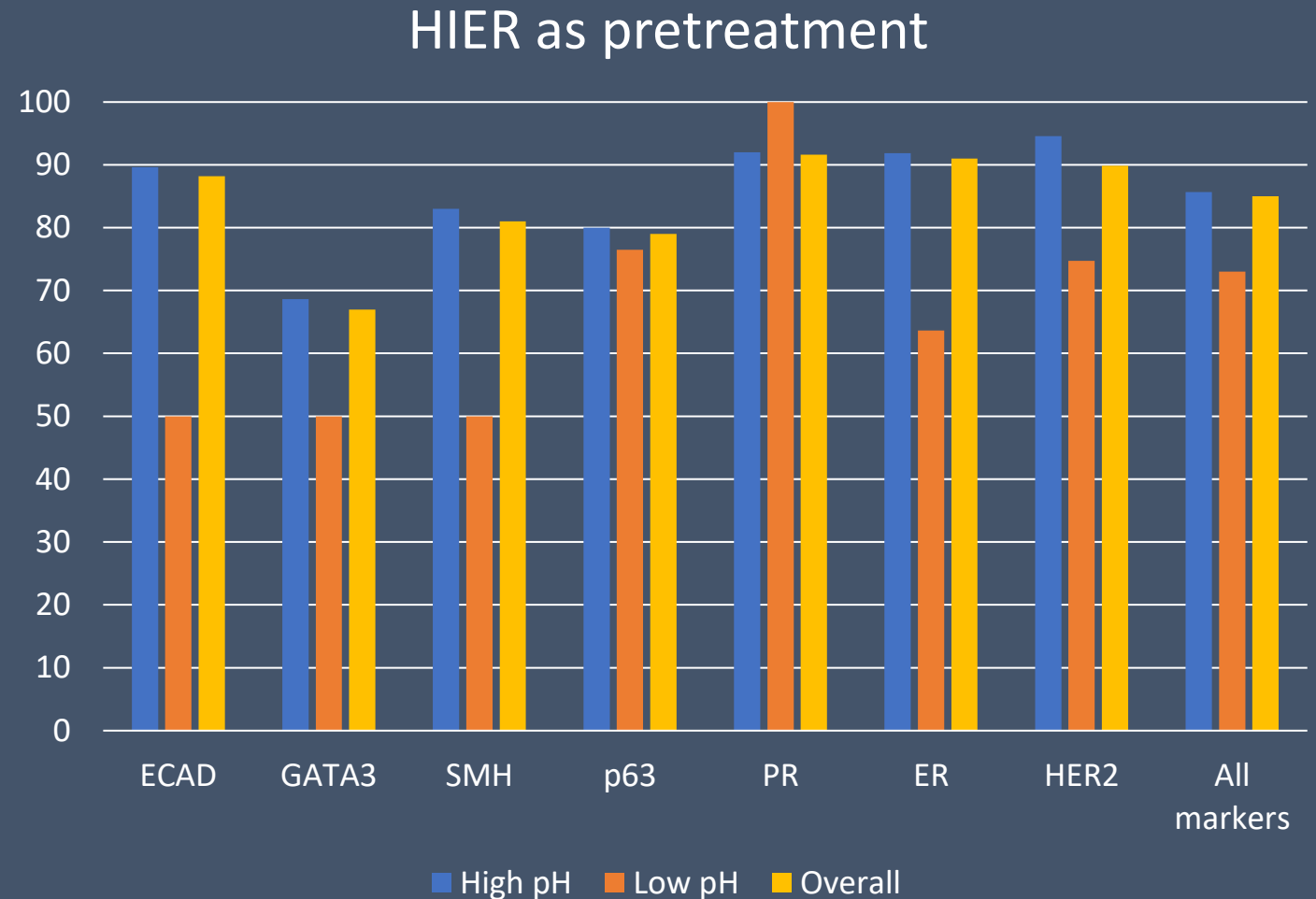
HIER in High pH: 86%

*Ranging from 69% for GATA3 till 95% for HER2*

HIER in Low pH: 73%

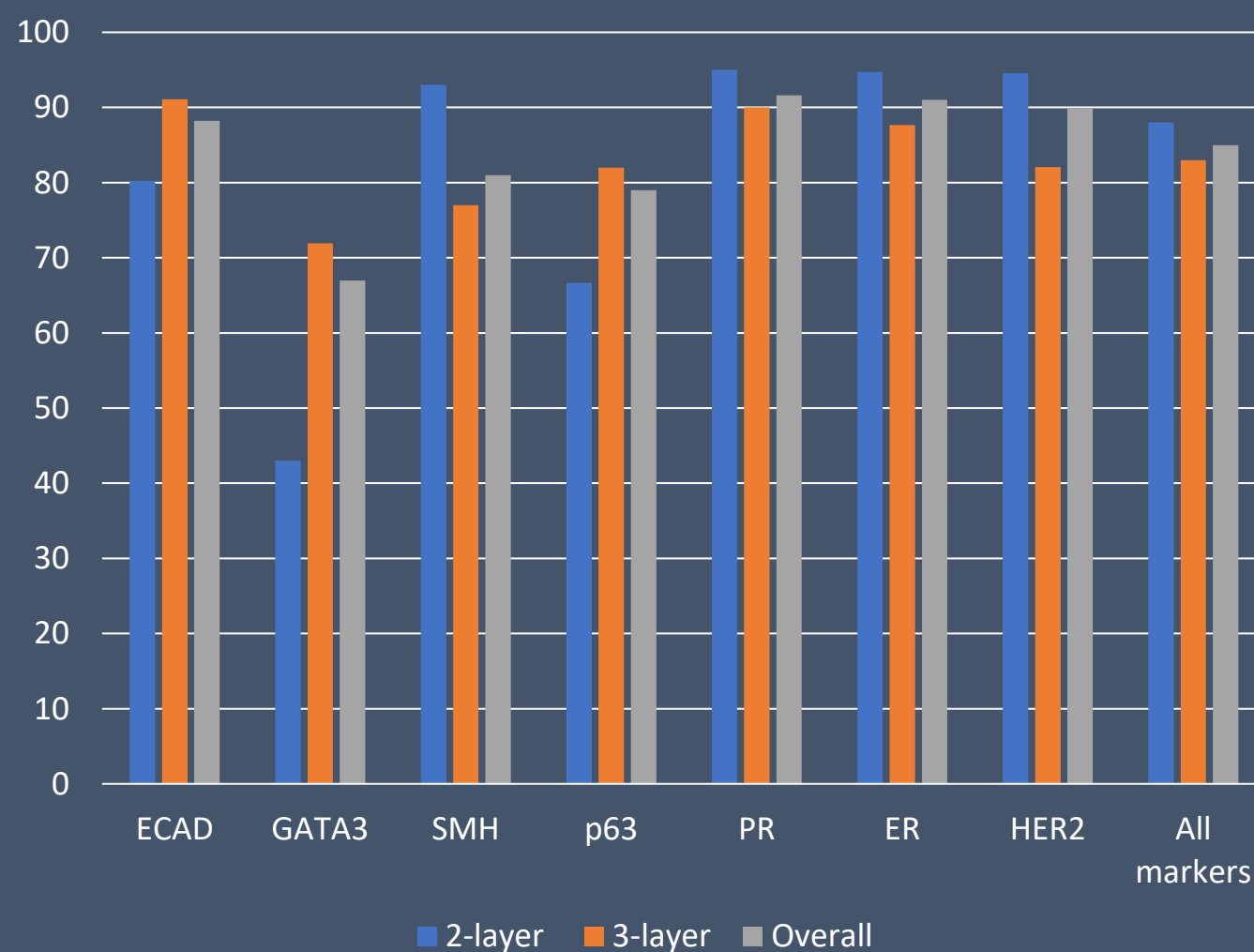
*Ranging from 50% for ECAD, GATA3 and SMH till 100% for PR\**

*\*15/15 participants used a low pH buffer, on a Leica or Dako platform.*



# KEY-POINTS FOR BEST PROTOCOLS

Detection system, pass rates



## Breast markers:

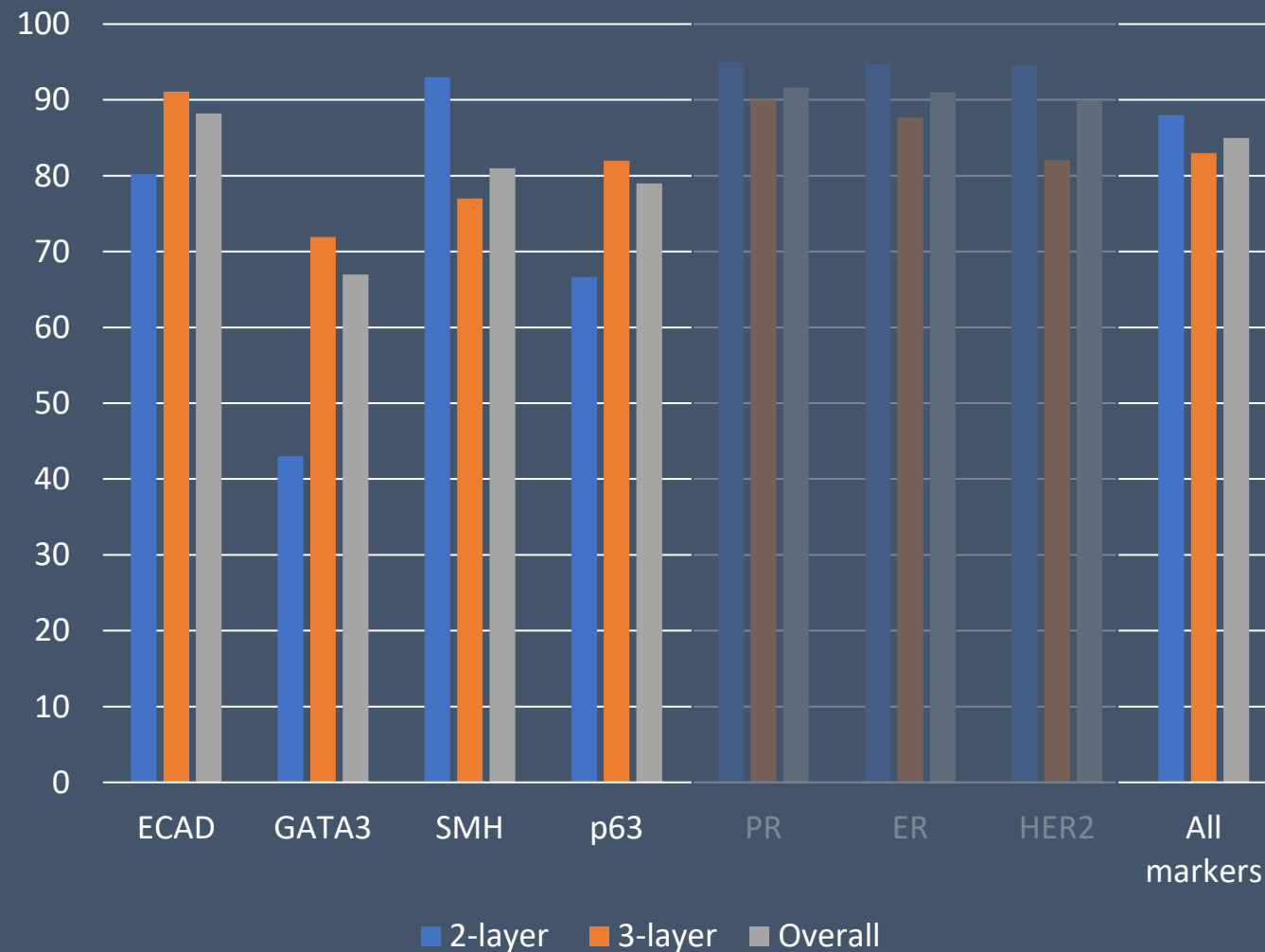
3-layer detection system: 83%

2-layer detection system: 88%



# KEY-POINTS FOR BEST PROTOCOLS

Detection system, pass rates



## Breast markers:

3-layer detection system: 83%

2-layer detection system: 88%

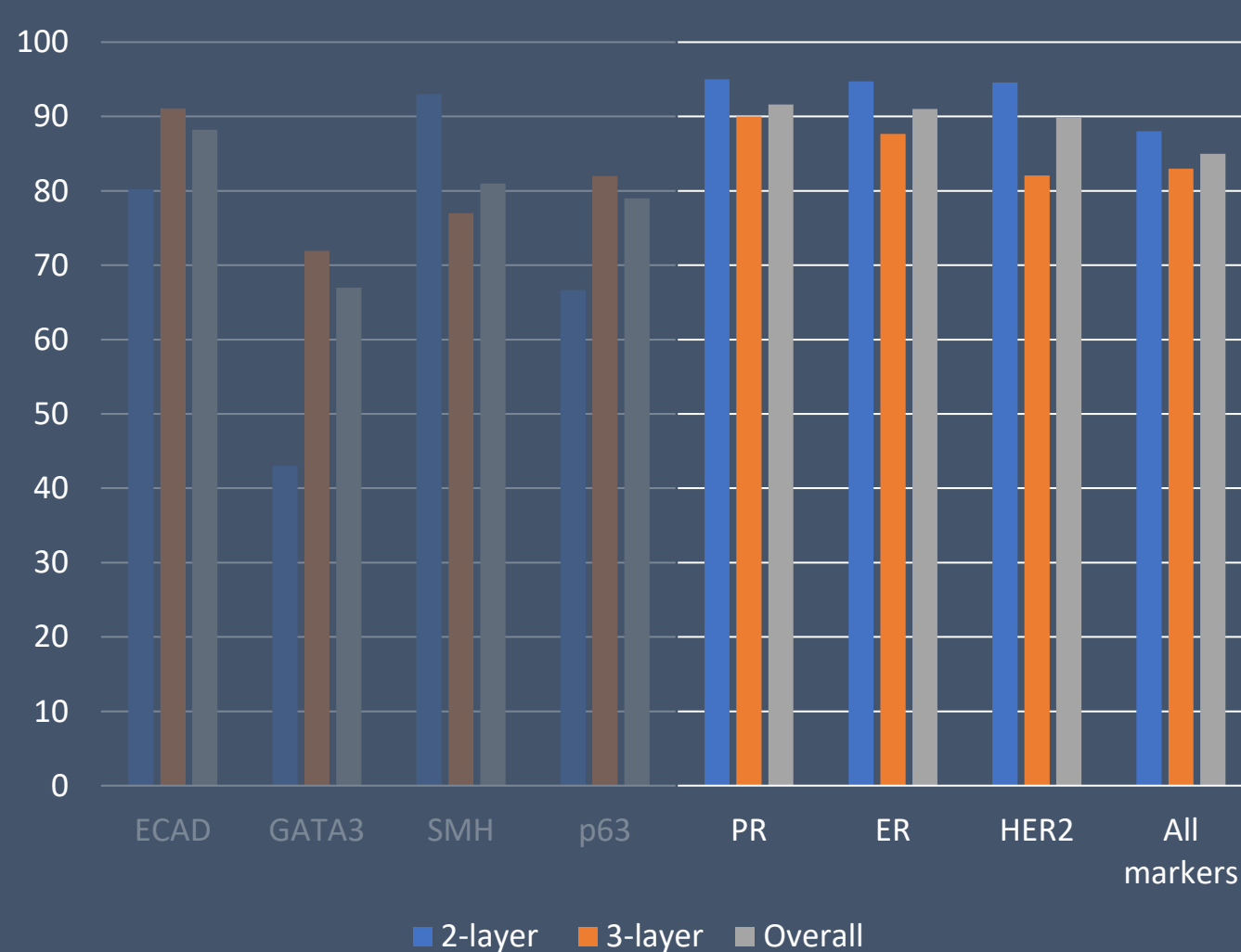
## Type I tests

3-layer detection system: 81%

2-layer detection system: 71%

# KEY-POINTS FOR BEST PROTOCOLS

Detection system, pass rates



## Breast markers:

3-layer detection system: 83%

2-layer detection system: 88%

### Type I tests

3-layer detection system: 81%

2-layer detection system: 71%

### Type II test

3-layer detection system: 87%

2-layer detection system: 95%



NOW TIME TO LOOK  
AT SOME SPECIFIC  
MARKERS

# GATA3 – PITFALLS/POINTS OF ATTENTION

Table 1. Antibodies and assessment marks for GATA3, Run 63

Concentrated antibodies	n	Vendor	Optimal	Good	Borderline	Poor	Suff. <sup>1</sup>	OR <sup>2</sup>
mAb clone <b>L50-823</b>	88	Cell Marque	31	40	33	24	56%	25%
	24	Biocare						
	4	BD Pharmingen						
	3	Zytomed Systems						
	3	Gennova						
	2	Bio-SB						
	2	Immunologic						
	1	Anacrom						
rmAb clone <b>EP368</b>	5	Cell Marque	4	-	1	1	67%	67%
	1	Quartett						
mAb clone <b>HG3-31</b>	2	Santa Cruz	-	-	-	2	-	-
rmAb clone <b>ZR65</b>	1	Zeta Corporation	-	-	1	-	-	-
Conc total	137		35	40	35	27	55%	26%
Ready-To-Use antibodies							Suff. <sup>1</sup>	OR. <sup>2</sup>
mAb clone <b>L50-823 760-4897<sup>3</sup></b>	56	Ventana/Roche	36	12	8	-	86%	64%
mAb clone <b>L50-823 760-4897<sup>4</sup></b>	67	Ventana/Roche	41	16	7	3	85%	61%
mAb clone <b>L50-823 390M-17,18,10</b>	42	Cell Marque	14	12	13	3	62%	33%
mAb clone <b>L50-823 PM 405AA</b>	12	BioCare Medical	5	3	2	2	67%	42%
mAb clone <b>L50-823 MAD-000632QD</b>	3	Master Diagnostica Vitro SA	1	2	1	-	-	-
mAb clone <b>L50-823 CGM-0130</b>	1	Celnovte	-	1	-	-	-	-
mAb clone <b>GATA3/6664 AMB89</b>	1	BioGenex	-	-	-	1	-	-
RTU total	183		97	46	31	9	78%	53%
Total	320		132	86	66	36		
Proportion			41%	27%	21%	11%	68%	

1) Proportion of sufficient results (optimal or good). (≥5 assessed protocols).

2) Proportion of Optimal Results (OR).

3) Vendor Recommended Protocol Settings (VRPS) to a specific RTU product applied on the vendor recommended platform(s) (≥5 assessed protocols).

4) Laboratory Modified Protocol Settings (LMPS) to a specific RTU product applied either on the vendor recommended platform(s), non-validated semi/fully automatic systems or used manually (indicated in percentage if ≥5 assessed protocols).

Table 2. Proportion of optimal results for GATA3 for the most commonly used antibodies as concentrate on the four main IHC systems\*

Concentrated antibody	Dako/Agilent Autostainer		Dako/Agilent Omnis		Ventana/Roche BenchMark XT / Ultra		Leica Biosystems Bond III / Max	
	TRS pH 9.0	TRS pH 6.1	TRS pH 9.0	TRS pH 6.1	CC1 pH 8.5	CC2 pH 6.0	BERS2 pH 9.0	BERS1 pH 6.0
mAb clone <b>L50-823</b>	1/12** (8%)	0/1	11/36 (31%)	0/1	15/46 (33%)	0/1	4/19 (21%)	-
rmAb clone <b>EP368</b>	1/1	-	2/2	-	0/1	-	0/1	-

\* Antibody concentration applied as listed above, HIER buffers and detection kits used as provided by the vendors of the respective systems.

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

No RTU products for Dako and Leica users.  
Use of conc. format of mAb L50-823 - and rmAb clone EP368 on Dako platforms - can obtain optimal results.

Recommended protocol settings:

- HIER in an alkaline buffer
- 40% pass rate for 2-step detection systems (8% optimal)
- 76% pass rate for 3-step detection systems (50% optimal)

# SMH - PITFALLS/POINTS OF ATTENTION

**Table 3. Proportion of sufficient and optimal results for SMH for the most commonly used RTU IHC systems**

RTU systems	Recommended protocol settings*		Laboratory modified protocol settings**	
	Sufficient	Optimal	Sufficient	Optimal
Dako AS mAb SMMS-1 <b>TR/IS066</b>	73% (8/11)	18% (2/11)	86% (6/7)	57% (4/7)
Leica BOND mAb S131 <b>PA0493</b>	100% (7/7)	100% (7/7)	3/3	2/3
VMS Ultra/XT mAb SMMS-1 <b>760-2704</b>	87% (13/15)	67% (10/15)	95% (38/40)	60% (24/40)

\* Protocol settings recommended by vendor – Retrieval method and duration, Ab incubation times, detection kit, IHC stainer/equipment.

\*\* Significant modifications: retrieval method, retrieval duration and Ab incubation time altered >25%, detection kit – only protocols performed on the specified vendor IHC stainer integrated.

**Table 2. Proportion of optimal results for SMH for the most commonly used antibody as concentrate on the 4 main IHC systems\***

Concentrated antibodies	Dako/Agilent Autostainer Link / Classic		Dako/Agilent Omnis		Ventana/Roche BenchMark GX / XT / Ultra		Leica Biosystems Bond III / Max	
	TRS pH 9.0	TRS pH 6.1	TRS pH 9.0	TRS pH 6.1	CC1 pH 8.5	CC2 pH 6.0	ER2 pH 9.0	ER1 pH 6.0
mAb <b>SMMS-1</b>	0/1**	-	6/7 (86%)	0/1	8/12 (67%)	-	6/8 (75%)	(1/1)

\* Antibody concentration applied as listed above, HIER buffers and detection kits used as provided by the vendors of the respective systems.

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

No RTU for Omnis is available. 13 laboratories used the Autostainer RTU on the Omnis; 15% pass rate, none optimal.

Limited data for concentrated formats on Omnis, but possible to achieve an optimal staining.

 <b>Nordic Immunohistochemical Quality Control</b> <small>Institute of Pathology, Aalborg University Hospital, Ladegaardsgade 3, P.O. Box 561, DK-9100 Aalborg, Denmark</small>	
<b>Recommended protocol for SMH</b> Obtained in run 66 12 Jul 2022	
<b>Immunostainer</b>	
Type:	Dako Omnis
<b>Primary antibody</b>	
Clone:	SMMS-1
Producer:	Cell Marque
Product no. / lot no.:	298M-14/15/16 / 0000144768
Diluent:	Antibody Diluent
Dilution factor:	1:400
Incubation time / temperature:	30 min. / 32°C
<b>Epitope retrieval, HIER</b>	
Device:	On Board / On Machine
Buffer:	Dako Omnis Target Retrieval Solution, High pH
Heating time at max. temp.:	30 min.
Maximum heating temp.:	97°C
<b>Visualization system</b>	
Producer:	Dako Omnis
Product / no.:	EnVision Flex / GV800/GV823
Linker:	Mouse LINKER
Incubation time linker:	10 min.
Incubation time polymer:	20 min.
Incubation temperature:	32°C
<b>Chromogen</b>	
Producer:	Dako Omnis
Product / no.:	DAB+ Substrate Chromogen System / GV825
Incubation time / temperature:	5 min. / 32°C
Enhancement:	None
<b>Disclaimer:</b> NordiQC makes every attempt to provide accurate and up-to-date information, yet NordiQC does not make any claim or warranty regarding the accuracy of the provided information nor does it represent that the contents of the web site and protocols reflect the most recent developments in immunohistochemistry at any point in time.	

# P63 - PITFALLS/POINTS OF ATTENTION

**Table 3. Proportion of sufficient and optimal results for p63 for the most commonly used RTU IHC systems**

RTU systems	Recommended protocol settings*		Laboratory modified protocol settings**	
	Sufficient	Optimal	Sufficient	Optimal
VMS Ultra/XT mAb 4A4 <b>790-4509</b>	57% (4/7)	0/7	88% (100/114)	52% (59/114)
Dako AS48 mAb DAK-p63 <b>IR662</b>	91% (11/12)	17% (2/12)	57% (4/7)	0/7
Dako Omnis mAb DAK-p63 <b>GA662</b>	85% (17/20)	25% (5/20)	100% (13/13)	62% (8/13)
Leica Bond mAb 7JUL <b>PA0103</b>	1/4	0/4	0/6	0/6

\* Protocol settings recommended by vendor – Retrieval method and duration, Ab incubation times, detection kit, IHC stainer/equipment.

\*\* Significant modifications: retrieval method, retrieval duration and Ab incubation time altered, detection kit – only protocols performed on the specified vendor IHC stainer integrated.

**Table 2. Proportion of optimal results for p63 for the most commonly used antibody as concentrate on the four main IHC systems\***

Concentrated antibodies	Dako/Agilent Autostainer		Dako/Agilent Omnis		Ventana/Roche BenchMark GX / XT / Ultra		Leica Bond III / Max	
	TRS pH 9.0	TRS pH 6.1	TRS pH 9.0	TRS pH 6.1	CC1 pH 8.5	CC2 pH 6.0	ER2 pH 9.0	ER1 pH 6.0
mAb clone <b>4A4</b>	0/3**	0/1	1/2	-	9/20 (45%)	-	1/7 (14%)	0/1
mAb clone <b>DAK-p63</b>	0/3	-	4/9 (44%)	0/1	17/24 (71%)	-	0/9	-
mAb clone <b>7JUL</b>	-	-	-	-	0/4	-	0/6	0/1

\* Antibody concentration applied as listed above, HIER buffers and detection kits used as provided by the vendors of the respective systems.

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

Vendor recommended protocol based on UltraView and 16-20 min. incubation of primary Ab.

Most common and successful modification was prolonging incubation time and use of OptiView or UltraView with amplification.

Vendor recommended protocol based on HIER in TRS Low pH.

Most successful modification was using HIER in TRS High pH.

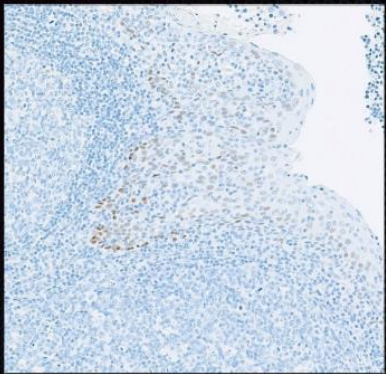
Less successful performance for 7JUL on the Bond platform.

Limited data for Bond users, but conc. 4A4 might be the best solution.

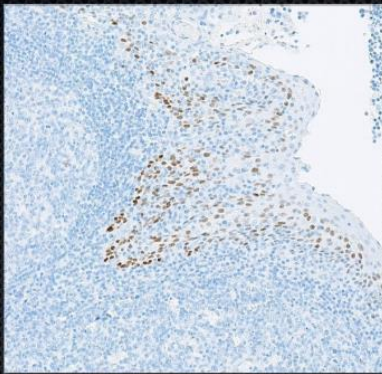


# P63 - PITFALLS/POINTS OF ATTENTION

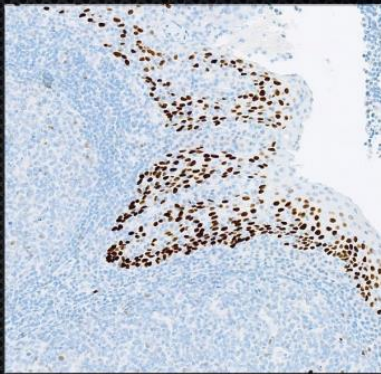
p63, 4A4 - OptiView (3-step) - Various HIER time NordiQC



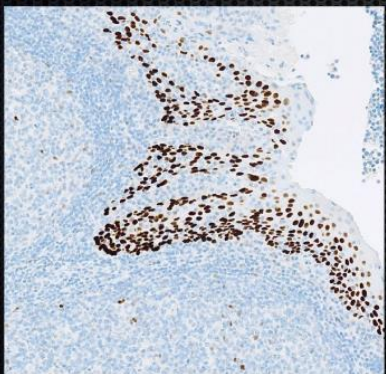
CC1\_8\_100°C



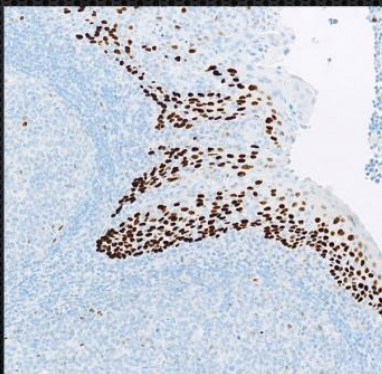
CC1\_16\_100°C



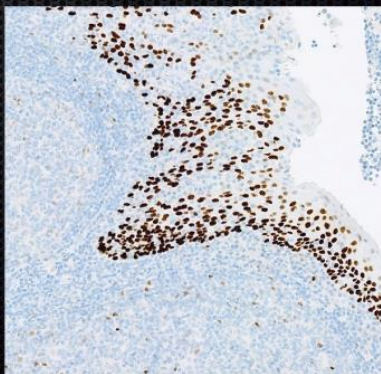
CC1\_32\_100°C



CC1\_48\_100°C

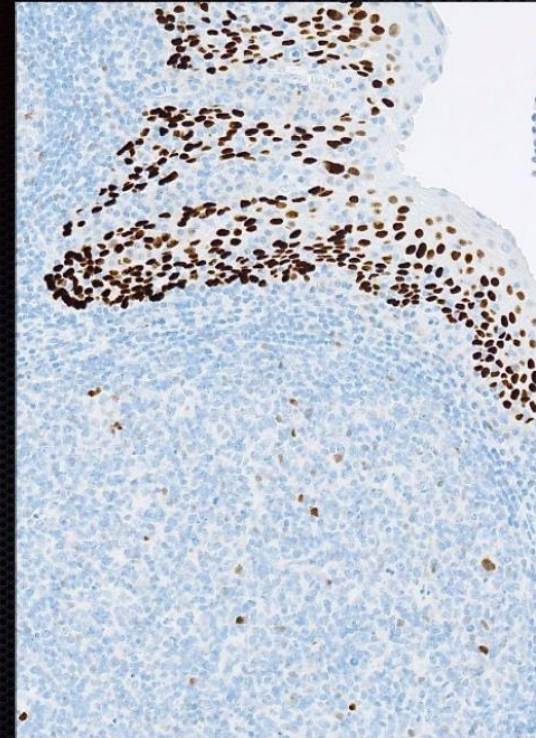


CC1\_64\_100°C

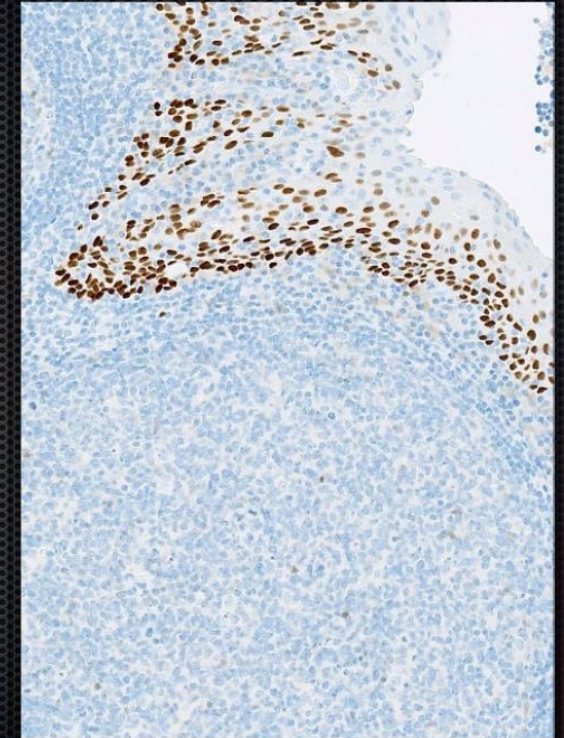


CC1\_92\_100°C

p63, 4A4 OptiView (3-step) vs UltraView (2-step) NordiQC



OptiView - HIER CC1\_48\_100



UltraView - HIER CC1\_52\_100



# ECAD - PITFALLS/POINTS OF ATTENTION

Table 1. Antibodies and assessment marks for ECAD, run 53

Concentrated antibodies	n	Vendor	Optimal	Good	Borderline	Poor	Suff. <sup>1</sup>	Suff. OPS <sup>2</sup>
mAb clone <b>NCH-38</b>	82	Agilent/Dako Immunologics	57	22	4	1	94%	98%
mAb clone <b>36</b>	1	BD Biosciences Biogenex	0	1	0	1	-	-
mAb clone <b>36B5</b>	13	Leica/Novocastra	2	10	1	0	92%	100%
mAb clone <b>4A2C7</b>	4	Life Tech./Invitrogen	2	2	0	0	-	-
mAb clone <b>BS38</b>	1	Nordic Biosite	0	1	0	0	-	-
mAb clone <b>DBM15.49</b>	1	Diagnostic BioSystems	1	0	0	0	-	-
mAb clone <b>ECH-6</b>	2	Zytomed Systems	1	0	1	0	-	-
mAb clone <b>HECD-1</b>	9	Life Tech./Invitrogen	4	5	0	1	90%	100%
mAb clone <b>GM016</b>	1	Genemed	1	0	0	0	-	-
mAb clone <b>SPM471</b>	1	Thermo S./Neomarkers	0	0	1	0	-	-
rmAb <b>EP700Y</b>	5	Cell Marque	0	4	1	0	-	-
rmAb <b>EP6</b>	1	Zeta Corporation	0	1	0	0	-	-
Ready-To-Use antibodies								
mAb clone <b>36 790-4497</b>	68	Roche/Ventana	54	11	3	0	96%	100%
mAb clone <b>GM016 8229-C010</b>	2	Sakura Finetek	2	0	0	0	100%	-
mAb clone <b>NCH-38 GA059</b>	31	Agilent/Dako	31	0	0	0	100%	100%
mAb clone <b>NCH-38 GA059<sup>3</sup></b>	6	Agilent/Dako	5	1	0	0	-	-
mAb clone <b>NCH-38 IS/IR059</b>	27	Agilent/Dako	26	1	0	0	100%	100%
mAb clone <b>NCH-38 IS/IR059<sup>3</sup></b>	6	Agilent/Dako	4	2	0	0	-	-
mAb clone <b>MX020 MAB-0738</b>	1	Maixin	0	1	0	0	100%	-
mAb clone <b>BS38 MAD-000643QD</b>	1	Master Diagnostica	1	0	0	0	100%	-
mAb clone <b>HECD-1 MAD-000761QD</b>	1	Master Diagnostica	1	0	0	0	100%	-
mAb clone <b>35B5 PA0387</b>	6	Leica/Novocastra	0	6	0	0	100%	-
rmAb clone <b>EP700Y 760-4440</b>	17	Roche/Ventana	0	2	15	0	13%	-
rmAb clone <b>EP700Y 246R-18</b>	6	Cell Marque	0	1	5	0	-	-
mAb clone <b>EP6 API3012</b>	1	Biocare Medical	0	1	0	0	100%	-
Total	298		192	72	31	3	-	
Proportion			65%	24%	10%	1%	89%	

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

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

3) Ready-to-use product developed for a specific semi/fully automated platform by a given manufacturer but inappropriately applied by laboratories on other non-validated semi/fully automatic systems or used manually.

Table 3. Proportion of optimal results for ECAD for the most commonly used antibody as concentrate on the 4 main IHC systems\*

Concentrated antibodies	Dako Autostainer Link / Classic		Dako Omnis		Ventana BenchMark XT / Ultra		Leica Bond III / Max	
	TRS pH 9.0	TRS pH 6.1	TRS pH 9.0	TRS pH 6.1	CC1 pH 8.5	CC2 pH 6.0	ER2 pH 9.0	ER1 pH 6.0
mAb clone <b>NCH-38</b>	8/10** (80%)	-	1/1	-	32/42 (76%)	-	6/6 (100%)	0/2

\* Antibody concentration applied as listed above, HIER buffers and detection kits used as provided by the vendors of the respective systems.

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

Concentrated format of mAb NCH-38 works on the main IHC Systems

Table 4. Proportion of sufficient and optimal results for ECAD for the most commonly used RTU IHC systems

RTU systems	Recommended protocol settings*		Laboratory modified protocol settings**	
	Sufficient	Optimal	Sufficient	Optimal
Dako AS mAb NCH-38 <b>IS/IR059</b>	100% (10/10)	100% (10/10)	100% (13/13)	100% (13/13)
Dako Omnis mAb NCH-38 <b>GA059</b>	100% (21/21)	100% (21/21)	(3/3)	(3/3)
VMS Ultra/XT/GX mAb 36 <b>790-4497</b>	100% (11/11)	72% (8/11)	95% (54/57)	81% (46/57)

\* Protocol settings recommended by vendor – Retrieval method and duration, Ab incubation times, detection kit, IHC stainer/equipment.

\*\* Significant modifications: retrieval method, retrieval duration and Ab incubation time altered >25%, detection kit – only protocols performed on the specified vendor IHC stainer were included.

RTU assays work as “plug-and-play” products.  
The majority of RTU assays obtain high pass rates  
– except assays based on rmAb EP700Y



# ER – PITFALLS / POINTS OF ATTENTION

Table 3. Comparison of pass rates for vendor recommended and laboratory modified RTU protocols

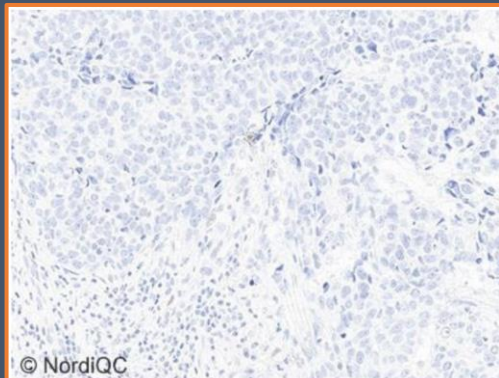
RTU systems	Vendor recommended protocol settings*		Laboratory modified protocol settings**	
	Sufficient	Optimal	Sufficient	Optimal
Dako AS48 rmAb EP1 <b>IR084/IS084</b>	3/3	1/3	18/21 (86%)	4/21 (19%)
Dako Omnis rmAb EP1 <b>GA084</b>	41/42 (98%)	29/42 (69%)	23/23 (100%)	12/23 (52%)
Leica Bond mAb 6F11 <b>PA009/PA0151</b>	2/2	0/2	12/19 (63%)	8/19 (42%)
VMS Ultra/XT/GX rmAb SP1 <b>790-4324/4325</b>	50/50 (100%)	30/50 (60%)	169/180 (94%)	121/180 (67%)

\* Protocol settings recommended by vendor – Retrieval method and duration, Ab incubation times, detection kit, IHC staining equipment.  
 \*\* Modifications included: retrieval method, retrieval duration, retrieval reagents, Ab incubation time, detection kit and use of amplification. Only protocols performed on the specified vendor IHC stainer are included.

Even with these successful results, changing RTU assays requires internal validation.

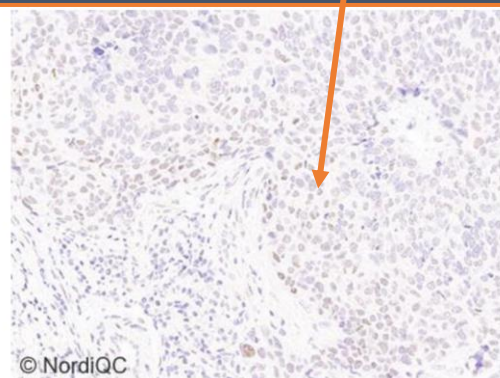
For Dako and Ventana products, the most common modification was using a 3-step detection system.

For Leica, modification in HIER – changing from low till high pH buffer was made the majority of participants.



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Fig. 6a  
Optimal ER staining of the breast carcinoma no. 5 expected to be negative using same protocol as in Figs. 1a – 5a. No nuclear staining reaction is seen and a high signal-to-noise ratio is observed.



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Fig. 6b  
Insufficient ER staining of the breast carcinoma no 5 with no ER expression. A weak but distinct nuclear staining reaction is seen in >10% of the neoplastic cells.  
 The insufficient result was only seen for the mAb clone 6F11 and likely was caused by performing HIER in an alkaline buffer in combination with other protocol settings inducing a too high level of technical/analytical IHC sensitivity compromising the diagnostic specificity.

# PR – PITFALLS / POINTS OF ATTENTION

Table 1. Antibodies and assessment marks for PR, run B31

Concentrated antibodies	n	Vendor	Optimal	Good	Borderline	Poor	Suff. <sup>1</sup>	OR <sup>2</sup>
mAb clone <b>16</b>	33	Leica Biosystems Monosan	20	12	-	2	94%	59%
mAb clone cocktail <b>16 + SAN27</b>	5	Leica Biosystems	2	2	1	-	80%	40%
rmAb clone <b>BP6081</b>	1	Biolyx	-	1	-	-	-	-
mAb clone <b>PgR 636</b>	13	Dako/Agilent Invitrogen	5	4	3	2	64%	36%
mAb clone <b>PgR 1294</b>	10	Dako/Agilent	8	1	1	-	90%	80%
mAb clone <b>PR88</b>	1	BioGenex	-	-	-	1	-	-
rmAb clone <b>SP2</b>	1	Diagnostic BioSystems Thermo Scientific	2	-	-	-	-	-
rmAb clone <b>SP42</b>	3	Zytomed	-	2	1	-	-	-
rmAb clone <b>YR85</b>	1	Fischer Scientific	-	1	-	-	-	-
rmAb clone <b>ZR4</b>	1	Zeta Corporation	1	-	-	-	-	-

Ready-To-Use antibodies	n	Vendor	Optimal	Good	Borderline	Poor	Suff. <sup>1</sup>	OR <sup>2</sup>
mAb clone <b>16 PA0312 (VRPS<sup>3</sup>)</b>	6	Leica Biosystems	6	-	-	-	100%	100%
mAb clone <b>16 PA0312 (LMPS<sup>4</sup>)</b>	12	Leica Biosystems	10	1	1	-	92%	83%
mAb clone <b>16 MAD-00670QD</b>	2	Master Diagnostica	-	-	2	-	-	-
mAb <b>PgR 636 IR/IS068 (VRPS<sup>3</sup>)</b>	4	Dako/Agilent	3	1	-	-	-	-
mAb <b>PgR 636 IR/IS068 (LMPS<sup>4</sup>)</b>	26	Dako/Agilent	21	3	-	2	92%	81%
mAb <b>PgR 1294 GA090 (VRPS<sup>3</sup>)</b>	33	Dako/Agilent	10	22	1	-	97%	30%
mAb <b>PgR 1294 GA090 (LMPS<sup>4</sup>)</b>	20	Dako/Agilent	11	5	4	-	80%	55%
rmAb clone <b>1E2 790-2223/4296 (VRPS<sup>3</sup>)</b>	53	Ventana/Roche	44	9	-	-	100%	83%
rmAb clone <b>1E2 790-2223/4296 (LMPS<sup>4</sup>)</b>	141	Ventana/Roche	108	23	9	1	93%	77%
mAb clone <b>IHC751 IHC751</b>	1	GenomeMe	1	-	-	-	-	-
rmAb clone <b>SP2 Kit-0013</b>	2	Maixin	2	-	-	-	-	-
rmAb clone <b>Y85 8360-C010</b>	4	Sakura Finetek	4	-	-	-	-	-
mAb <b>PgR 636 PM343</b>	1	Biocare Medical	-	1	-	-	-	-
<b>Total</b>	<b>377</b>		<b>258</b>	<b>88</b>	<b>23</b>	<b>8</b>		
<b>Proportion</b>			<b>68%</b>	<b>23%</b>	<b>6%</b>	<b>2%</b>	<b>92%</b>	

1) Proportion of sufficient results (optimal or good) (≥5 assessed protocols).

2) Proportion of optimal results (≥5 assessed protocols).

3) Vendor Recommended Protocol Settings (VRPS) to a specific RTU product applied on the vendor recommended platform(s).

4) Laboratory Modified Protocol Settings (LMPS) to a specific RTU product applied either on the vendor recommended platform(s) or other platforms.

Graph 1. Pass rate in the NordiQC assessments for PR

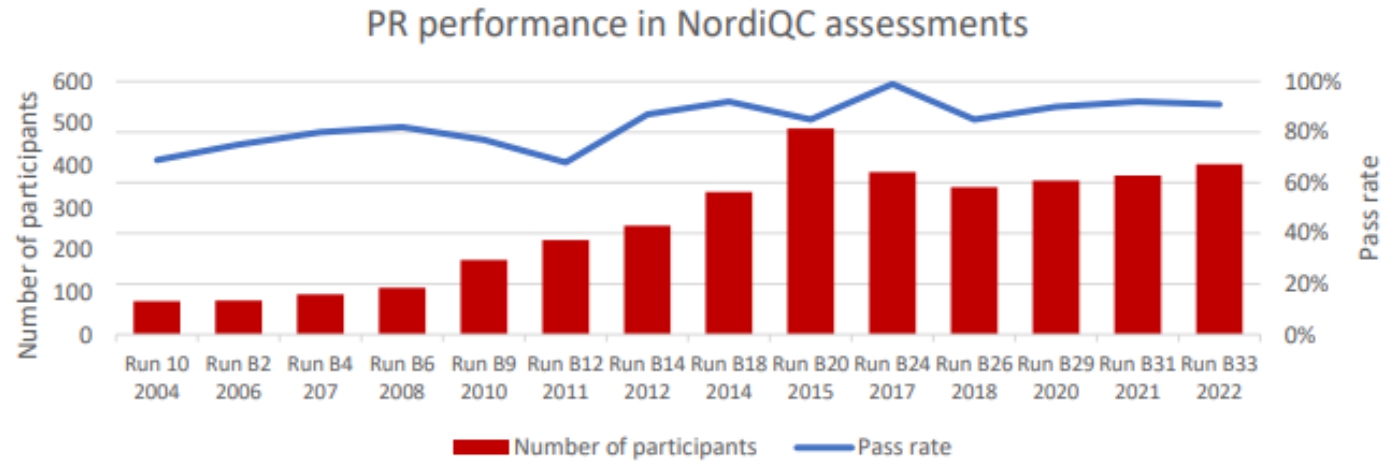


Table 3. Comparison of pass rates for vendor recommended and laboratory modified RTU protocols

RTU systems	Vendor recommended protocol settings*		Laboratory modified protocol settings**	
	Sufficient	Optimal	Sufficient	Optimal
Leica BOND MAX/ BOND III mAb 16 <b>PA0312</b>	9/9 (100%)	9/9 (100%)	11/11 (100%)	7/11 (64%)
Dako Autostainer+/ Autostainer Link mAb PgR 636 <b>IS068/IR068</b>	8/8 (100%)	6/8 (75%)	17/17 (100%)	15/17 (88%)
Dako Omnis mAb PgR 1294 <b>GA090</b>	33/41 (80%)	18/41 (44%)	22/23 (96%)	17/23 (74%)
Ventana BenchMark GX/XT/Ultra rmAb 1E2 <b>790-2223/790-4296</b>	62/63 (98%)	38/63 (60%)	128/142 (90%)	87/142 (61%)

\* Protocol settings recommended by vendor – Retrieval method and duration, Ab incubation times, detection kit, IHC stainer/equipment

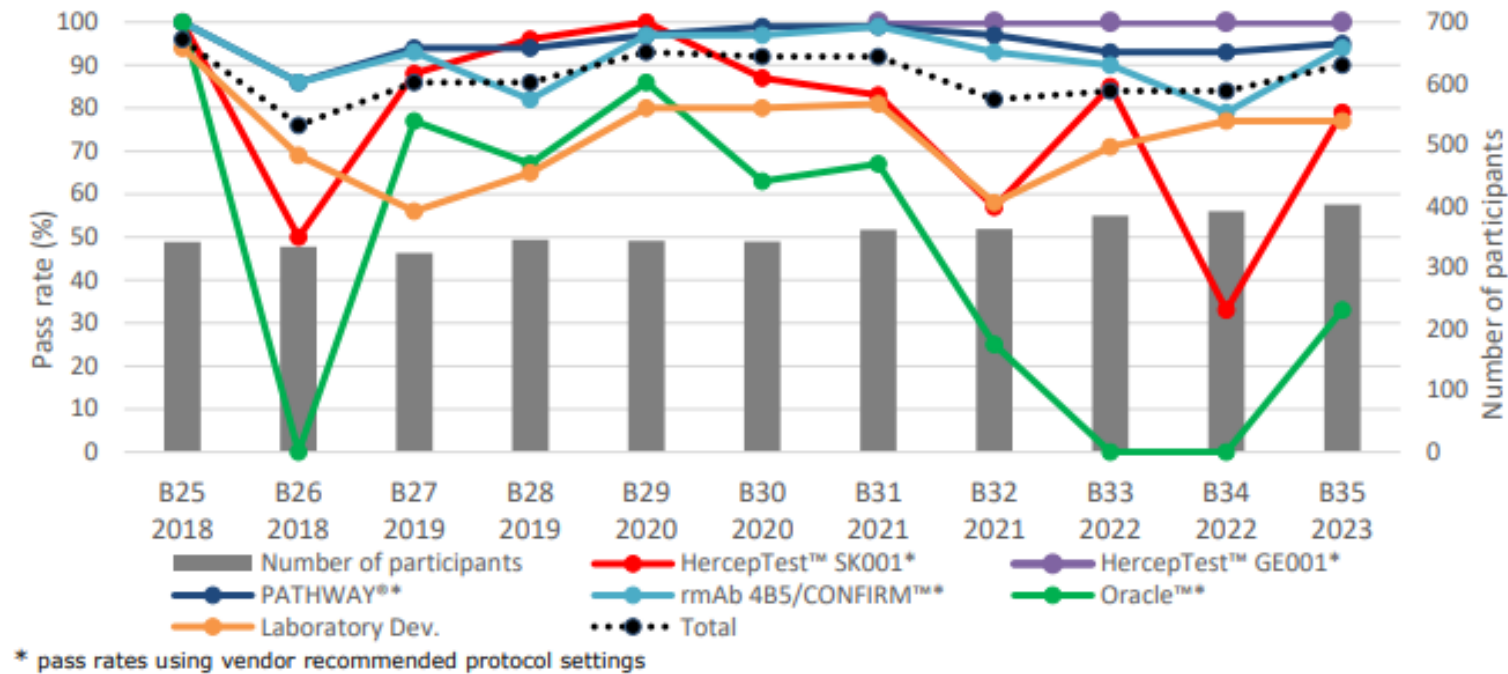
\*\* Modifications included: retrieval method, retrieval duration, retrieval reagents, Ab incubation time and detection kit. Only protocols performed on the specified vendor IHC stainer are included.

Autostainer RTU:  
If using Flex+ a pass rate of 100%, 90% optimal.

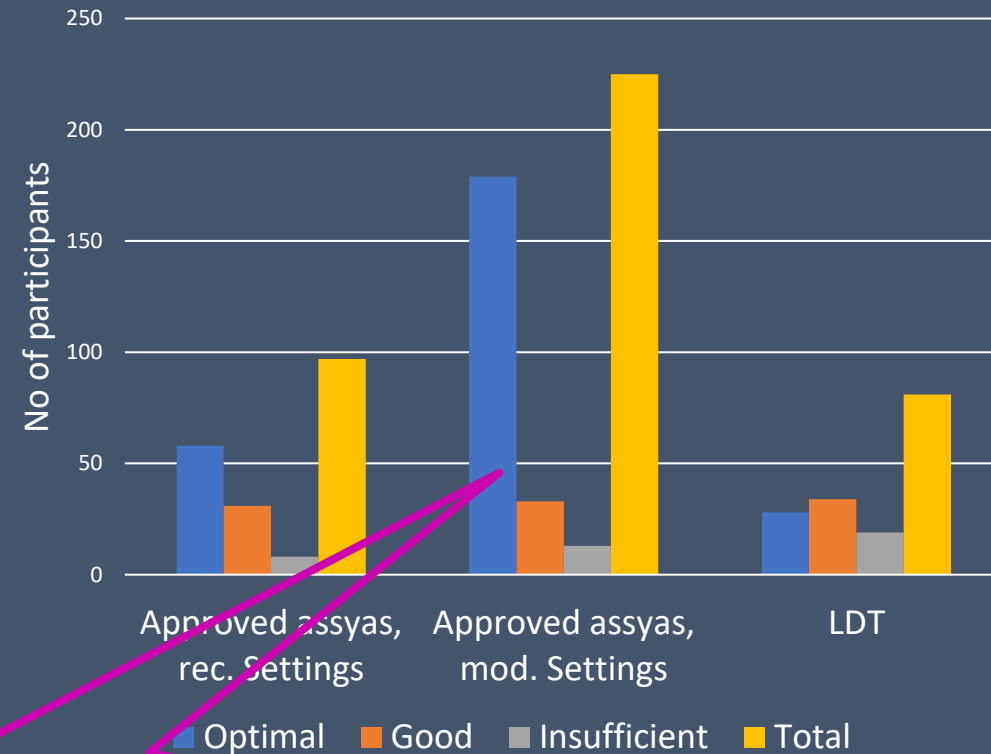
Omnis RTU:  
If using Flex+ a pass rate of 100%, 76% optimal.

# HER2 – PITFALLS / POINTS OF ATTENTION

Graph 1. Pass rates of the HER2 IHC assessments in the NordiQC breast cancer module 2018-2023



HER2 results, run B35

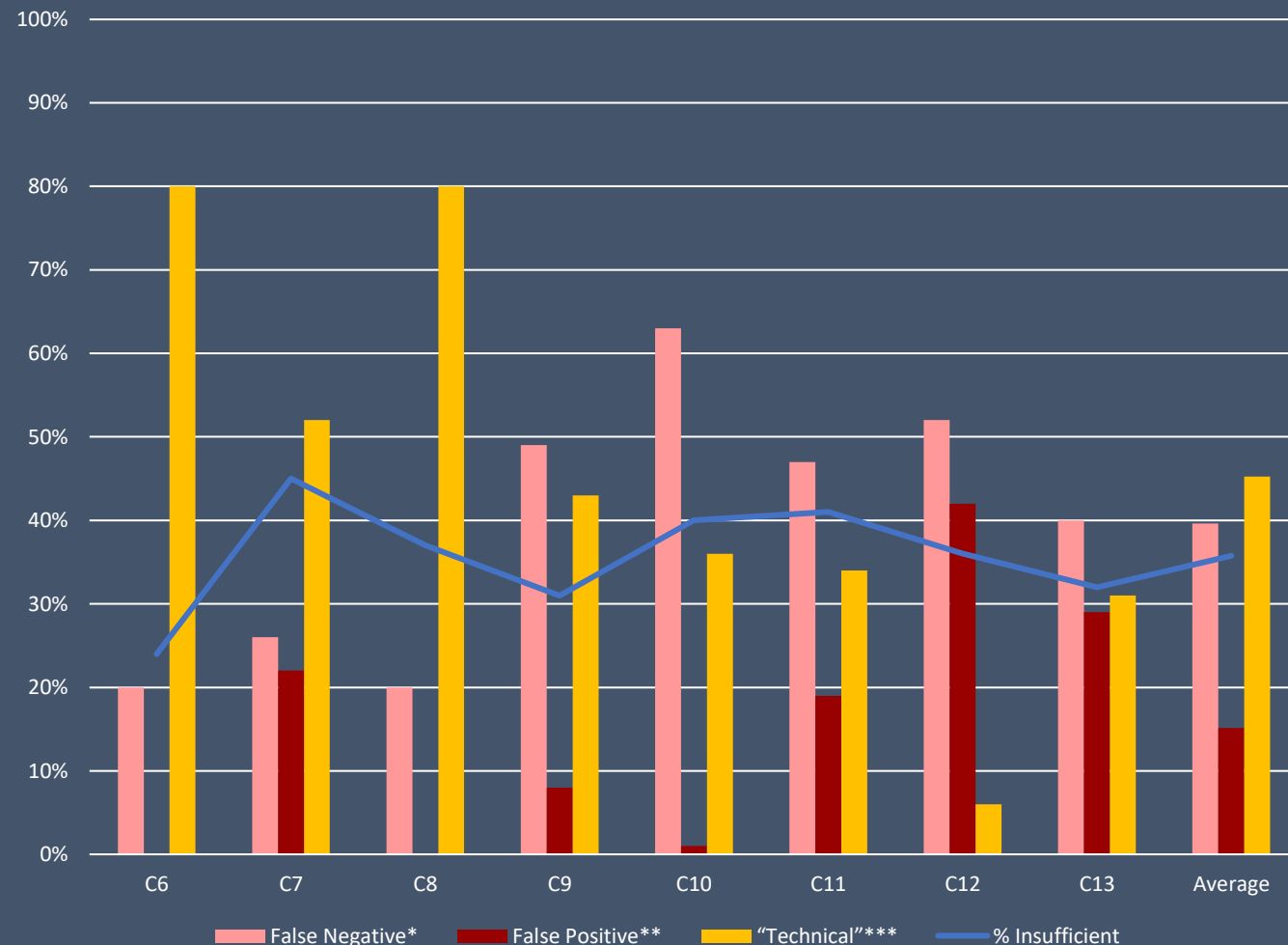


# PD-L1 IC – PITFALLS/POINTS OF ATTENTION

Table 2. Assessment marks for IHC assays and antibodies run C13, PD-L1 IC

CE-IVD / FDA approved PD-L1 assays		n	Vendor	Optimal	Good	Borderline	Poor	Suff. <sup>1</sup>	OR <sup>2</sup>
rmAb clone SP142, 741-4860 <sup>3</sup>		51	Ventana/Roche	24	18	8	1	82%	47%
rmAb clone SP142, 741-4860 <sup>4</sup>		1	Ventana/Roche	0	0	0	1	-	-
rmAb clone SP263, 741-4905 <sup>3</sup>		6	Ventana/Roche	0	3	2	1	50%	0%
rmAb clone SP263, 741-4905 <sup>4</sup>		1	Ventana/Roche	0	0	0	1	-	-
rmAb clone 28-8 pharmDX, SK005		1	Dako/Agilent	0	0	0	1	-	-
mAb clone 22C3 pharmDX, SK006		2	Dako/Agilent	0	1	1	0	-	-
mAb clone 22C3 pharmDX, GE006		3	Dako/Agilent	0	0	0	3	-	-
Antibodies <sup>7</sup> for laboratory developed PD-L1 assays, concentrated antibodies		n	Vendor	Optimal	Good	Borderline	Poor	Suff. <sup>1</sup>	OR <sup>2</sup>
mAb clone 22C3		4	Dako/Agilent	0	0	0	4	-	-
rmAb clone ZR3		1	Zeta Corporation	0	0	0	1	-	-
rmAb clone CAL10		4	Zytomed	0	0	4	0	-	-
rmAb clone E1L3N		2	Cell Signaling	0	0	2	0	-	-
rmAb clone QR001		1	Quartett	0	0	1	0	-	-
rmAb clone SP142		1	Abcam	0	1	0	0	-	-
Ready-To-Use antibodies <sup>8</sup>		n	Vendor	Optimal	Good	Borderline	Poor	Suff. <sup>1</sup>	OR <sup>2</sup>
rmAb clone SP142, 790-4860 (VRPS) <sup>5</sup>		21	Ventana/Roche	13	5	2	1	86%	62%
rmAb clone SP142, 790-4860 (LMPS) <sup>6</sup>		33	Ventana/Roche	17	9	6	1	79%	52%
rmAb clone SP263, 790-4905		1	Ventana/Roche	0	1	0	0	-	-
rmAb clone SP263, 790-4905 <sup>4</sup>		2	Ventana/Roche	0	0	2	0	-	-
rmAb clone SP142, RMA-0724		2	Fuzhou Maixin	0	2	0	0	-	-
rmAb clone AC37, PA168		1	Abcarta	0	0	0	1	-	-
mAb clone C9C9 CPM-0278		1	Celnovte	0	0	0	1	-	-
Total		139		54	40	28	17		
Proportion				39%	29%	20%	12%	68%	

Characteristics of insufficient results in the NordiQC PD-L1 IC assessments.





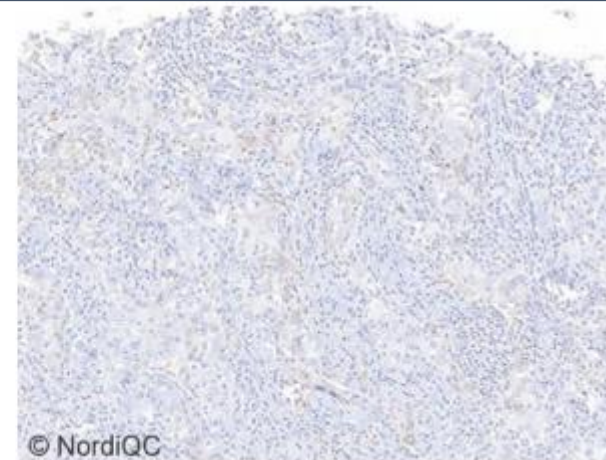
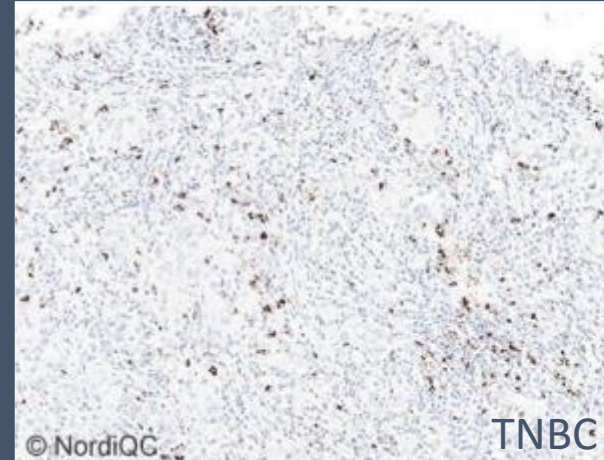
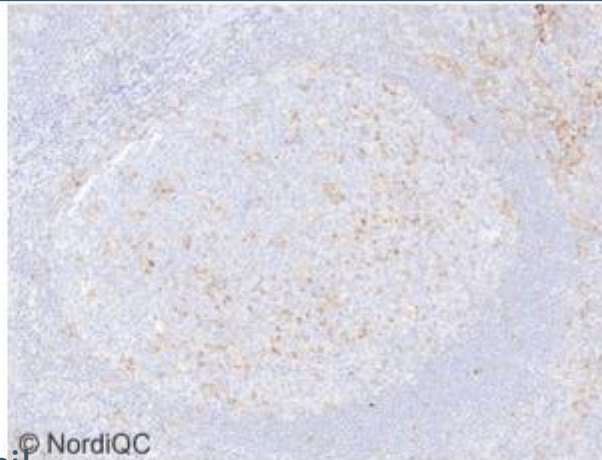
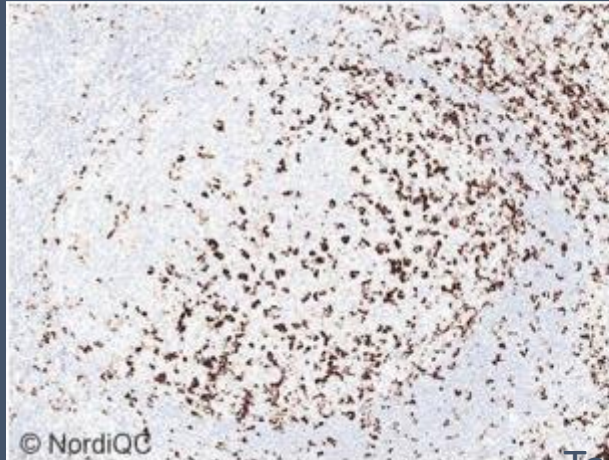
# PD-L1 IC – ICAPS

rmAb SP142

mAb 22C3

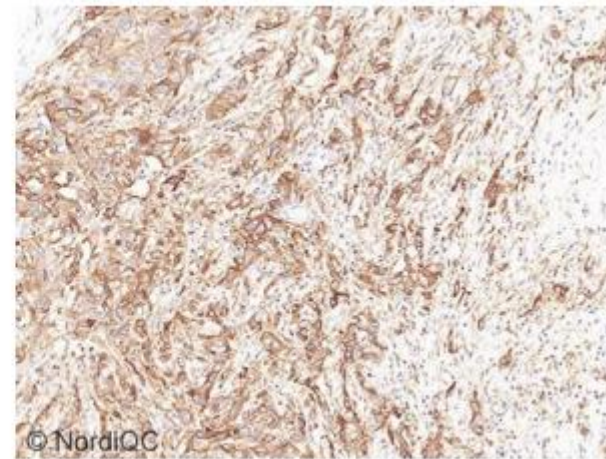
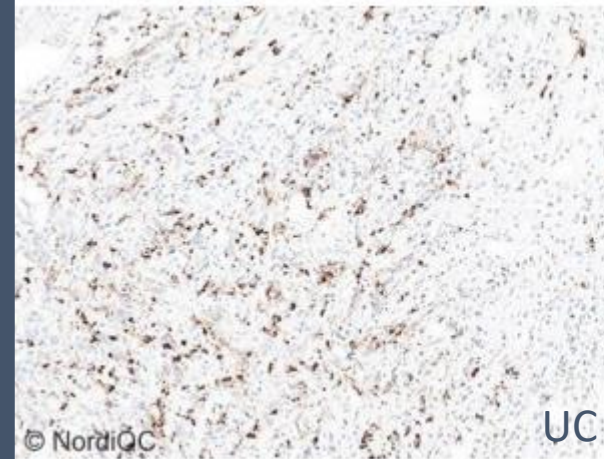
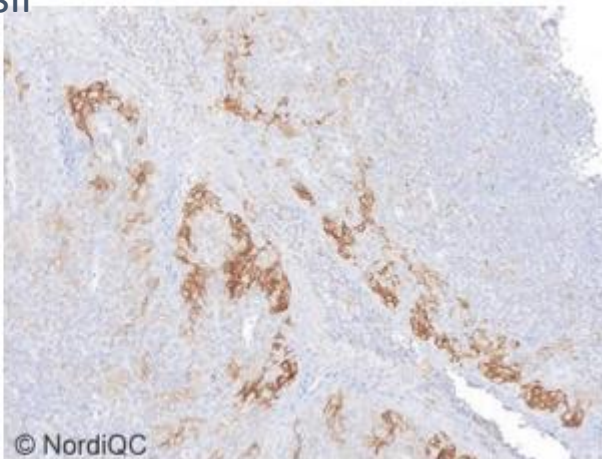
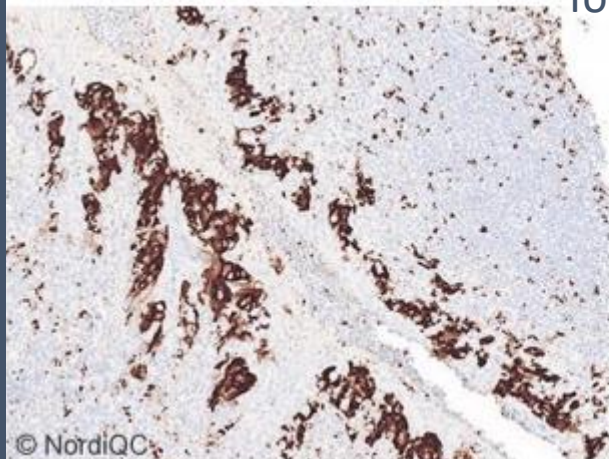
rmAb SP142

mAb 22C3



Tonsil

TNBC



UC

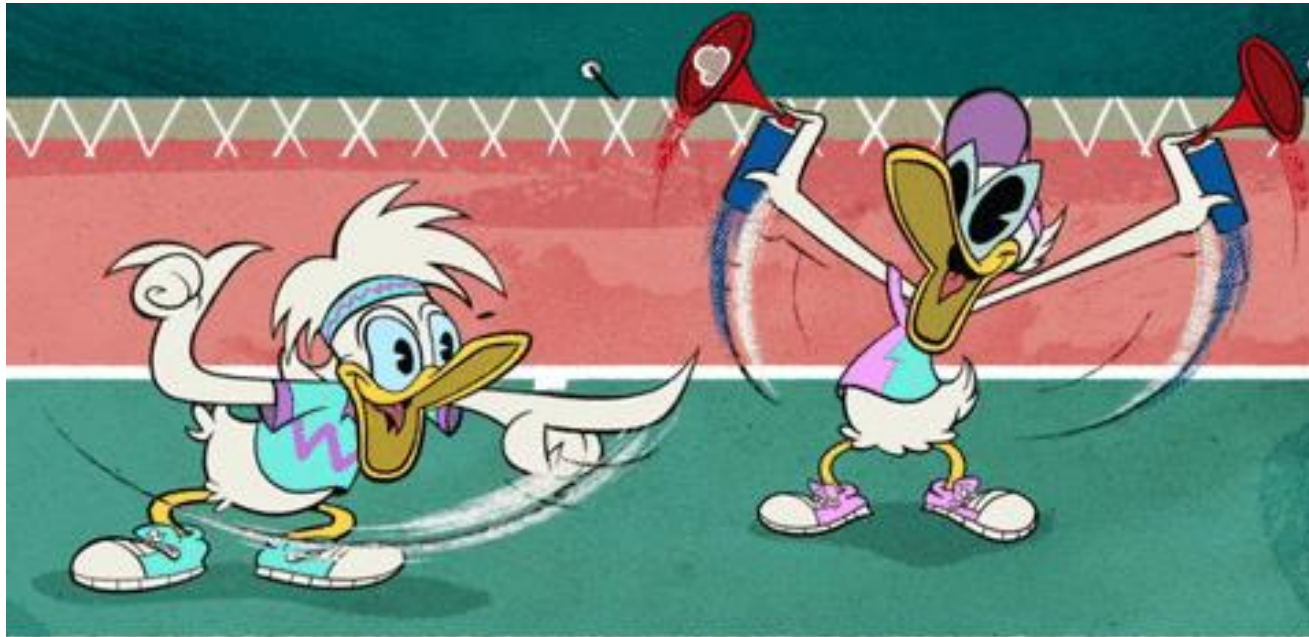
rmAb SP142

mAb 22C3

rmAb SP142

rmAb ZR3





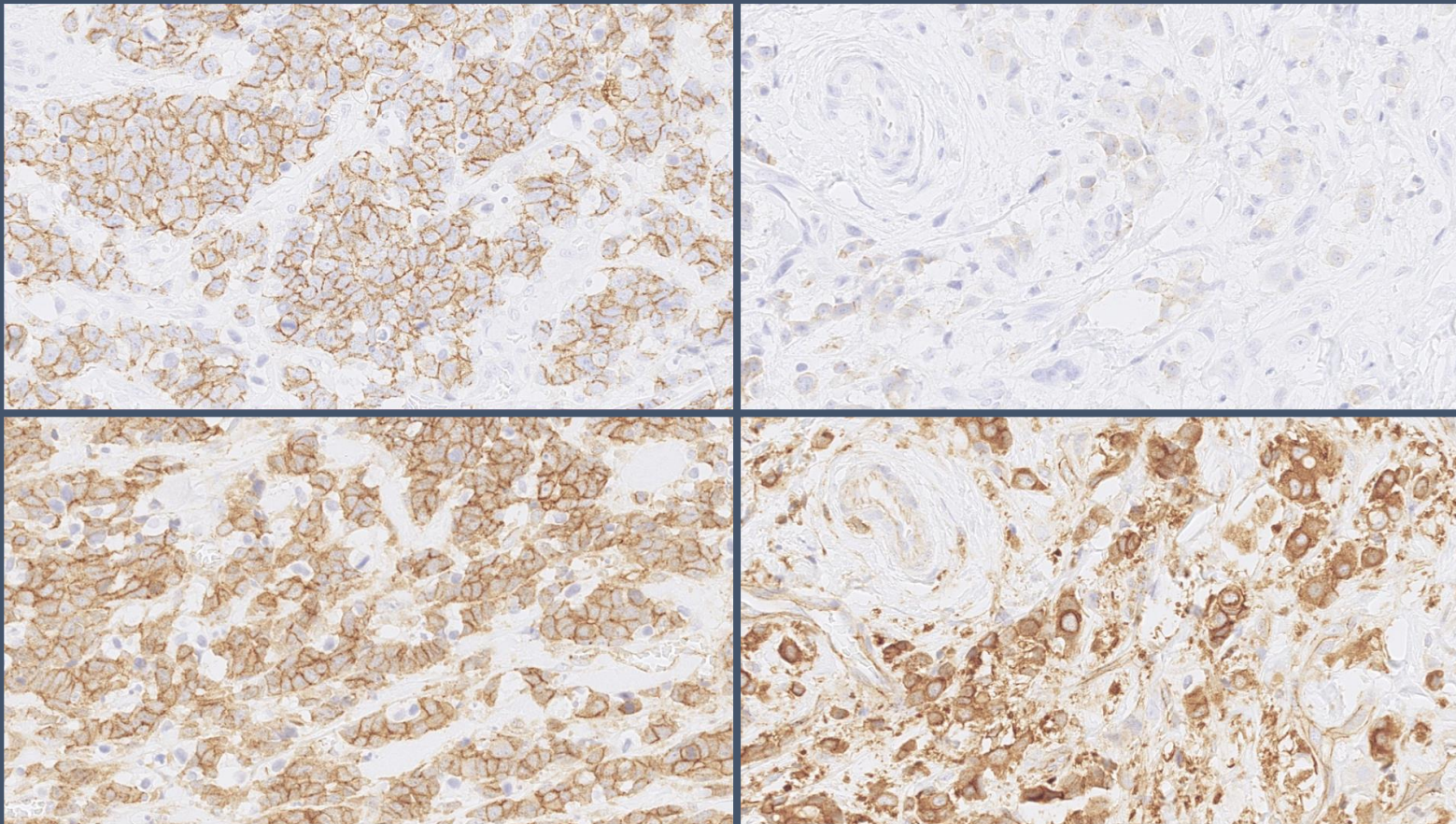
**CONGRATULATIONS!**

**...YOU SURVIVED!  
THANK YOU FOR  
YOUR ATTENTION**

# BONUS – P120

No NordiQC data available for p120 Catenin.

For the p120 stains below, a concentrated format of the mAb clone MRQ-5 is used.



Ductal breast carcinoma

Lobular breast carcinoma

**E-CAD**: membranous staining reaction in (most) ductal breast carcinomas, negative in (most) lobular breast carcinomas.

**p120**: membranous staining reaction in (most) ductal breast carcinomas, cytoplasmic staining reaction in (most) lobular breast carcinomas.

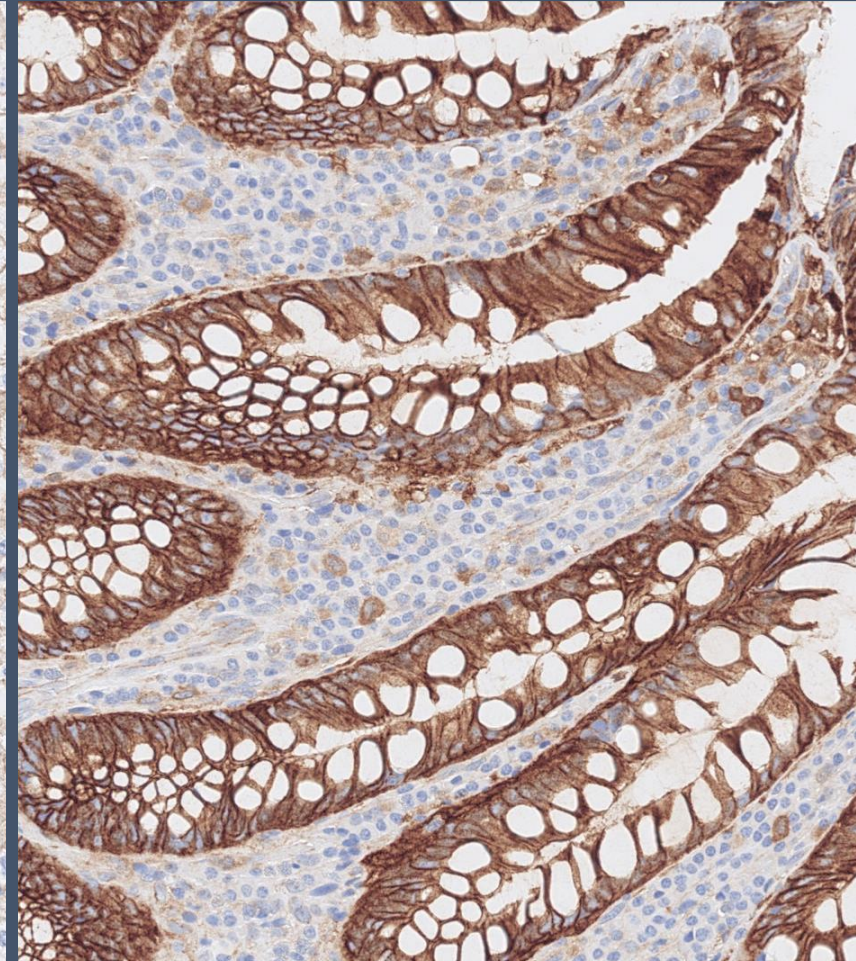
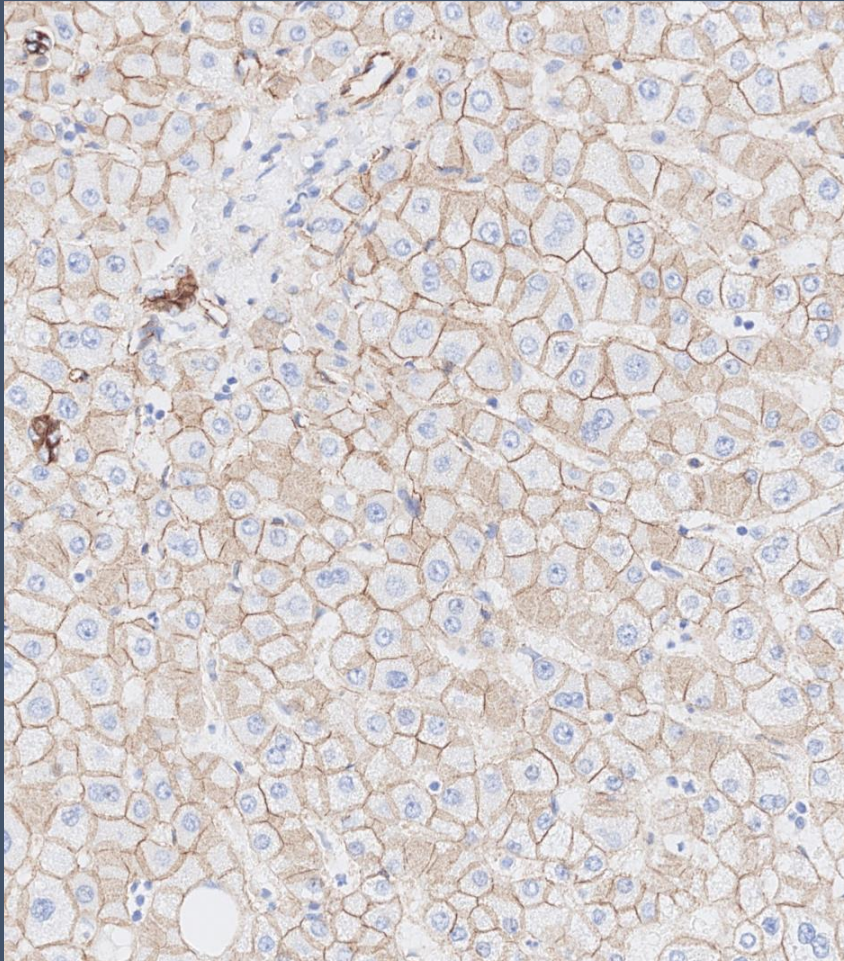


# BONUS – P120 ICAPS

No NordiQC data available for p120 Catenin.

For the p120 stains below, a concentrated format of the mAb clone MRQ-5 is used.

Liver:  
Hepatocytes must  
show a weak to  
moderate  
membranous  
staining



Appendix:  
Columnar  
epithelial cells  
must show a  
strong  
membranous  
staining.