

Workshop in Diagnostic Immunohistochemistry Aalborg University Hospital, October 2023

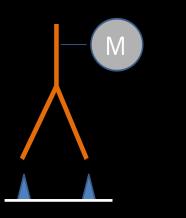
Technical aspects of immunohistochemistry & pitfalls Pre-Analytical - <u>Analytical (I & II)</u> - Post Analytical phase

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Immunohistochemistry – A simple technique ?

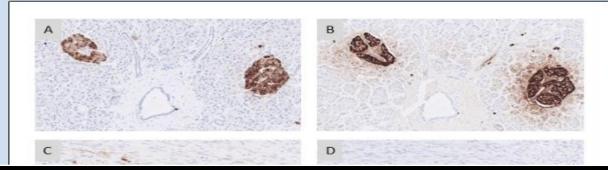


Infor Modules Assessments Protocols Controls Events <u>Login</u>

26-Apr-2024

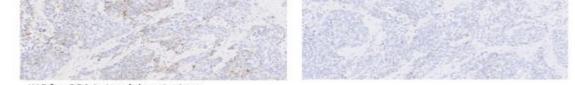
Results - Run 70, B37 and H25

↑ Top of page



External Quality Assurance programs

Staining quality varies significantly between different laboratories depending on the individual selection of methods



IHC for CGA in two laboratories:

Lab 1 (A+C+E): Optimal results in pancreas (A), appendix (C) and SCLC (E)

In the pancreas vast majority of endocrine islet cells show a strong and distinct cytoplasmic staining reaction. Normal ganglion cells and axons in the nerve plexus give a moderate and distinct, granular cytoplasmic staining reaction. The majority of neoplastic cells in the SCLC show a weak to moderate dot-like accentuation.

Lab 2 (B+D+F): Insufficient result due to inefficient HIER. In the pancreas the expected intense staining reaction is obtained, whereas both a reduced proportion, too weak staining and false negative staining reaction is seen in appendix and the SCLC.

The total test paradigm: Key elements in the IHC procedure

The analytic phase :

Begins with dewax of the cut slides and is completed with the coverslipping of the stained slides.

Unlike the pre-analytic factors, analytic parameters can easily be modified and controlled within the immunohistological laboratory.



Pre-analytic phase

Pre-fixation Fixation Post-Fixation/Decalcification Processing Dehydration & clearing Paraffin embedding Sectioning Drying/Storage



Analytic phase

Platform (manual/ Automated) Epitope retrieval Blocking Primary Antibody Detection system Chromogen Counterstain Mounting



Post-analytic phase

Design of controls Critical stain indicators Internal/External control Interpretation Positive/Negative Localization Quantification Cutt-of levels Reporting

Technical aspects of IHC and pitfalls – Analytical phase

Optimization of the IHC assay – issues to be addressed

- Purpose and/or "fit-for-purpose" (assay validated for intended use)
- How to establish "best practice protocol" of the IHC test (Calibration of the IHC assay with focus clone, antigen retrieval, titer & detection system)
 - Is the IHC test reproducible/robust (pre-analytic conditions)
 - Evaluation of the analytical sensitivity and specificity
- Identification of most robust controls providing information that the established level of detection is obtained in each test performed in daily practice.

Tissue materials are essential for all these processes (calibration, validation and controls)

Immunohistochemistry: Calibration of a biomarker/antibody depend on the type of marker and purpose of the test

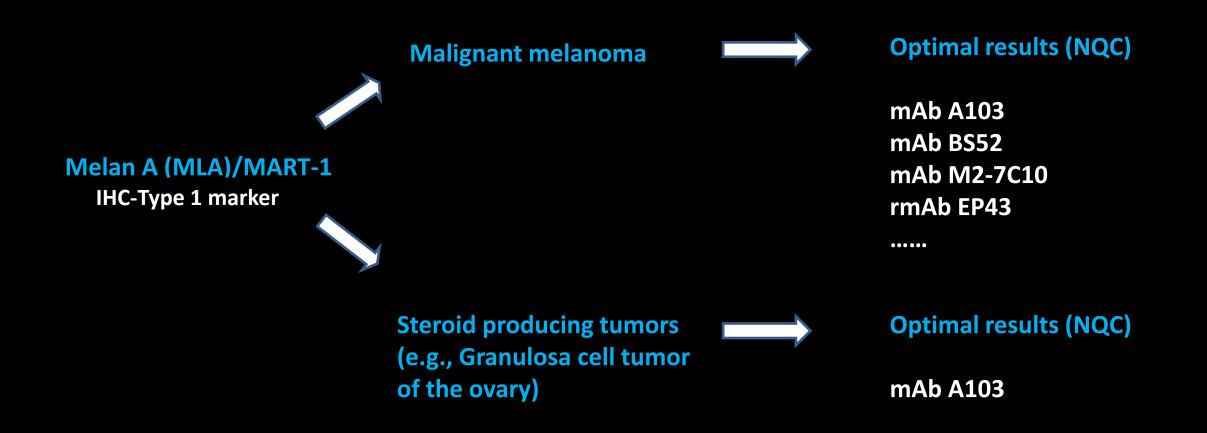
IHC-type 1 markers (Diagnostic)

Often calibrated to produced appropriate level of sensitivity and specificity (positive versus negative)

IHC-type 2 markers (Disease screening, predictive treatment & prognosis)

Predictive markers: The assays are calibrated to provide information of which patients will or will not benefit from a specific treatment (HER-2, PD-L1)

Purpose and Intended use



Concontrated antibadies		Vondor	Optimal	Cood	Renderline	Beer	Suff.1	Suff.
Concentrated antibodies n		Vendor	Optimal	Good	Borderline	Poor		OPS ²
mAb clone A103	69 19 5 1 1 1 1	Dako/Agilent Novocastra/Leica Cell Marque Diagnostic BioSystems Immunologic Monosan Thermo Scientific	22	43	24	8	67%	73%
mAb clone BS52	1	Nordic Biosite	1	0	0	0	-	-
mAb clone cocktail HMB45+MC-7310 +M2-9E3+T311	3	Biocare	1	2	0	0	-	-
mAb clone cocktail M2-7C10+M2-9E3	1	NeoMarkers	0	1	0	0	-	-
mAb clone M2-7C10	1	Zytomed	1	0	0	0	-	-
rmAb EP43	5 3 3	Nordic Biotite Cell Marque Epitomics	9	1	1	0	91%	100%
Ready-To-Use antibodies								
mAb clone A103 790-2990	87	Ventana/Roche	8	31	45	3	45%	66%
mAb clone A103, IR633/IS633	26	Dako/Agilent	10	14	2	0	92%	95%
mAb clone A103, 29 IR633/IS633 ³		Dako/Agilent	9	12	8	0	72%	-
mAb clone A103, 9 IR633/IS633 ⁴		Dako/Agilent	1	2	5	1	-	-
mAb clone A103, 9 PA0233		Novocastra/Leica	1	8	0	0	-	-
mAb clone A103, PA0233 ^s	1	Novocastra/Leica	0	0	1	0	-	-
mAb clone A103, 281M-87/281M-88		Cell Marque	1	0	2	0	-	-
mAb clone A103, API3114	1	Biocare	0	1	0	0	-	-
mAb clone A103, MAB-0275	1	Maixin	1	0	0	0	-	-
mAb clone cocktail HMB45+A103+T311, 904H-08	1	CellMarque	1	0	0	0	-	-
mAb clone cocktail HMB45+A103+T311, 790-4677	1	Ventana/Roche	0	1	0	0	-	-
mAb clone EP43, MAD- 000695QD-7/N	2	Master Diagnostica	0	1	1	0	-	-
rmAb clone EP43, 8319-C010	2	Sakura Finetek	1	1	0	0	-	-
Total	286		67	118	89	12	-	
			24%	41%	31%	4%	65%	

Proportion of sufficient stains (optimal or good)

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

 RTÚ system developed for the Dako/Agilent semi-automatic system (Dako Autostainer), but used by laboratories on the Da fully-automatic platform (Dako Omins)

 RTU system developed for the Dako/Agilent semi-automatic system (Dako Autostainer), but used by laboratories on differe platforms (e.g. Ventana BenchMark)

5) RTU system developed for the Leica Bond system, but used on the Ventana BenchMark system

Purpose and Intended use

Material

The slide to be stained for MLA comprised:

1. Skin, 2. Kidney, 3. Adrenal gland, 4-5. Malignant melanoma, 6. Granulosa cell tumour.



All tissues were fixed in 10% neutral buffered formalin.

Melan A <u>clone A103</u> (Run 56)

Melanoma/melanocyte assessment: 62% Sufficient/ 20% Optimal

Optimal results: Efficient HIER (alkaline buffer), high conc. of the primary Ab (app. 1:10-50) and a sensitive detection system (3-step)

Melanoma/melanocyte + <u>steroid hormone assessment</u>): Suff. 29%

Sufficient steroid hormone related A103 (cross)reaction can be very difficult to obtain on e.g., the Ventana BenchMark and Dako Omnis platforms.

Melan A (MLA) / MART-1:

263 participants ~ 92% used clone A103 as single reagent (no cocktails)

Table 1. Antibodies and Concentrated antibodies	n	Vendor	Optimal	Good	Borderline	Poor	Suff.1	OR ²
concentrated antibodies	57 19	Dako/Agilent Novocastra/Leica	Optimar	0000	Dordenine	1001	5011.	UK
mAb clone A103	6 1 1 1 1	Cell Marque Abcam Biocare Monosan Biogenex Zeta Corporation	21	57	9	0	90%	24%
mAb clone BS52	3	Nordic Biosite	3	0	0	0	-	-
mAb clone M2-7C10	1	Zytomed	1	0	0	0	-	-
mAb clone cocktail HMB45+MC-7310 +M2-9E3+T311	3	Biocare	2	1	0	0	-	-
mAb clone cocktail HMB45+A103+T311	1	Diagnostic Biosystems	0	0	0	1	-	-
mAb clone cocktail M2-7C10+M2-9E3	1	Thermo F. Scientific	1	0	0	0	-	-
rmAb clone EP1442Y	1	Abcam	1	0	0	0	-	-
rmAb clone EP43	9 9 1	Nordic Biotite Epitomics Cell Marque	18	1	0	0	100%	95%
Ready-To-Use antibodies								
mAb clone A103 790-2990 ³	3	Ventana/Roche	0	0	3	0	-	-
mAb clone A103 790-2990⁴	94	Ventana/Roche	6	74	11	3	85%	6%
mAb clone A103, IR633/IS633 ³	14	Dako/Agilent	1	13	0	0	100%	7%
mAb clone A103, IR633/IS633 ⁴	56	Dako/Agilent	12	36	7	1	85%	21%
mAb clone A103, PA0233/PA0044 ³	7	Leica Biosystems	2	5	0	0	100%	29%
mAb clone A103, PA0233/PA0044⁴	10	Leica Biosystems	6	4	0	0	100%	60%
mAb clone A103, 281M-87/281M-88	1	Cell Marque	1	0	0	0	-	-
mAb clone A103, API3114	1	Biocare	0	0	1	0	-	-
mAb clone M2-7C10, 281M-97/281M-98	2	Cell Marque	1	1	0	0	-	-
mAb clone cocktail HMB45+A103+T311, 904H-08	1	Cell Marque	0	0	1	0	-	-
mAb clone cocktail HMB45+A103+T311, 790-4677	1	Ventana/Roche	0	1	0	0	-	-
mAb clone cocktail HMB45+MC-7310 +M2-9E3+T311, PM165	1	Biocare	o	1	0	0	-	-
rmAb clone BP6086, [1064	1	Tuling Biotechnology	0	0	1	0		
mAb clone EP43, MAD-000695QD-7/N	1	Master Diagnostica	1	0	0	0	-	-
rmAb clone EP43, 8319-C010	3	Sakura Finetek	3	0	0	0	-	-
Total	312		80	194	33	5	-	
Proportion			26%	62%	11%	1%	88%	

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

Proportion of Optimal Results (OR).

3) Vendor Recommended Protocol Settings (VRPS) to a specific RTU product applied on the vendor recom asessed protocols)

4) Laboratory Modified Protocol Settings (LMPS) to a specific RTU product applied either on the validated semi/fully automatic systems or used manually (>5 asessed protocols)

Purpose and Intended use

Material

The slide to be stained for MLA comprised:

1. Kidney, 2. Skin, 3-4. Malignant melanoma, 5. Colon Adenocarcinoma.

All tissues were fixed in 10% neutral buffered formalin.

Melan A/MART-1 (Run 60)

MLA, A103 (melanoma/melanocyte assessment): 87% Sufficient / 18% Optimal.

Optimal results: Efficient HIER (alkaline buffer), high conc. of the primary Ab (app. 1:10-50) and a sensitive detection system (3-step).

Performance of MLA clone A103 is sensitive to the chosen platform e.g., Omnis or Benchmark.

Melan A (MLA) / MART-1:

273 participants ~ 88% used clone A103 as single reagent (no cocktails)

Is MLA, A103 the best primary Ab for detection of melanomas?

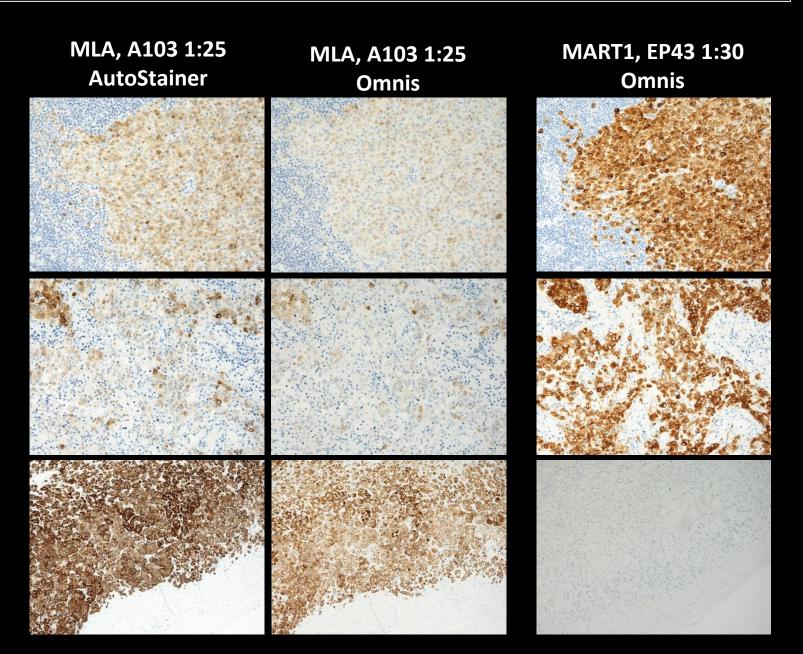
Melan A /MART1

Melanoma (Sentinel node)

Melanoma (Lymph node)

Adrenal gland

TRS High pH/Flex+ protocols



PRAME clone EPR20330 (Biocare Medical)

Purpose ?

Fit-for-purpose (validated for intended use)

PRAME: Purpose and "fit-for-purpose"

Antibody: PRAME	Controls (High. Low and None expression): HIGH: Malignant melanoma. The majoritity of the neoplastic cells should display an at least
Come: EPR20330 Company/Supplier: <u>Biocare</u> M. Date: 22-03-2023	LOW: Tonsil. Dispersed intragerminal centre lymphatic cells should show
	NONE: Appendix. Epithelial cells should be rep Remark: Subsets of lymphocytes/macrophages are positive in most of the tissue cores. In Remark: Subsets of lymphocytes/macrophages are positive in some cores. Only expression in normal
	addition, smooth muscle cells were assessed. epithelial or tumour cells were assessed.

		e Study (TMA Normal)		ined	Tota	al#	Not	es
ormal	Tissu	le Study (TWA Here		ined				
	Tis	sue		(+)		1		
				0	-	1		
a	Liv	ver		0		NA	Tis	sue core
b	S	oleen		NA	+	1	Di	spersed j
lc	P	lacenta		0	+	1	Di	spersed i
2a	T	onsil (5h)		0	+	1	F	ollicles ar
2a 2b	-	fonsil (29h)		0	+-	1	D	ispersed
			T	0	+	1	+	
2c			\top	0	+	1	+	
2d	+	in a acto-Celvia (22		0	+	1	+	
3a	+	un mino cervix (12011)	+	1	+	1	+	
3b	-+	Uterine endometrium		0	-	_	+	
3c	-+	Breast	-+-	0		1	+	
3d		Colon	+	0		1	-+	
4a		Pancreas	+	0		1	-+	
4b		Salivary Gland	-+	0		1		
4c		Placenta	-+	0		1		Few po
4d				1		1		Few pe
58	1	Kidney Adrenal gland		0		1		-
5	p	Adrenargiante				1		+
5	с	Prostate		0		1		
5	d	Testis		10		1		-
E	5a	Thyroid)	1		
	6b	Striated muscle			0		1	Norn
	6c	Cerebellum (240h)			0	1	_	
	6d	Skin		-				
	-							

Best practice protocol

Medullar carcinoma (Thyroid) 3b Follicular carcinoma (Thyroid) 0 3c Renal cell carcinoma 0 3d Clear cell carcinoma (Kidney) 0 3e-f Colon adenocarcinoma 0 4a Carcino-Sarcoma 0 (Uterus) 1 4b Endometroid Adc. (Uterus) 4c

1f

2a

2b

2c

2d

2e-f

3a

General tumour Study (TMA Tumour)

Breast carcinoma (Lobular)

Breast carcinomas

(Ductal)

	0	1 1 NA	Tissue core	Purpose: Discrir	ninate ma	lignant from k	penign	melanocytic lesions
	NA	1	Dispersed į	Specific Disease Tissue Study				
	0	1	Dispersed i Follicles ar	Tissue	# Stained (+)	# Stained (+) >76%1	Total #	Notes
	0	1	Dispersed	Nevus	3	0	10	Focal and/or dispersed cells
h)	0	1	Dispersee	Nevus atypical/dysplastic nevus	2	1	10	Both positive cases with focal stainin pattern.
20h) n	0 1 0 0 0 0	1 1 1 1 1 1		Melanoma In Situ	10	9	10	In two difficult cases, PRAME was po in dispersed melanocytic cells in one and in the second case, displayed for but diffuse reaction pattern, and thu regarded as positive using the criteri >76% positive neoplastic cells.
	0 0 1 0		Few po	Melanomas	16	16	17	4/16 (weak reaction of >76% of the neoplastic cells). The cores 2c and 4 without neoplastic cells, and thus, no assessed (NA).
h)			1 1 1 1 Norn	Rare isolated junctional melanocyte ¹ Threshold according to literature s Jul <u>;48</u> (7):856-862. <u>doi</u> : 10.1111/cup	supporting melanom			gn non- <u>lesional</u> skin. C compendium: J <u>Cutan Pathol</u> . 2021

Summery/Conclusion: 100% (10/10) Melanoma In Situ lesions and 94% (16/17) of melanomas (nos) expressed PRAME, whereas only 25% (5/20) of benign and atypical nevi were positive for PRAME - often in a minority of nuclei's and with focal staining pattern. This result is largely in agreement with published literature (see IHC Compendium) and RNA findings (Human protein Atlas (RNA seg/GTEx/Fantom5/TCGA dataset). Using test results without a threshold, the accuracy of the test is 87% (malignant vs benign melanocytic lesions) with a sensitivity of 96% and a specificity of 75%. Applying a threshold of >76% positive neoplastic cells supporting melanoma diagnosis (see selected references/IHC compendium), the accuracy of the test increased to 94% with a sensitivity of 96% (malignant vs benign melanocytic lesions) and a specificity of 95%. The assay is fit-for-purpose and thus, can be implemented as routine analysis. J Cutan Pathol. 2021 Jul;48(7):856-862

PReferentia	lly ex	bre	564	ed Antigen in MElanoma
tumour Study (TMA Tumour) Tissue			336	a Antigen in Mr.
rissue	Ch. i			
Liver	Stained	To	otal #	Notes
Lung SCC	(+)			Notes
Lung Adapase	1		1	
Lung Adenocarcinoma (Skin) Lung Adenocarcin	1		1	Weak reactions.
Lung Adenocarcinoma (Skin) intestine)	1	1	1	Faint to work
Breast carcinoma (Lobular)	1	1		Faint to weak reactions. Faint to weak reactions.
	0	<u> </u>		to weak reactions.
Ductal)	1	1		
ledullar carcinom (m		2	T	Faint reaction in dispersed cells
	0	1		dispersed cells
enal cell carcinoma (Thyroid)	0	1		
ear cell carcinoma (w.)	0	1		
	0	1		
cino-Sarcoma	0	2		
erus)	1			
ometroid Adc. (Uterus)		1	N	loderate to intense pueles
ometroid Ada (o	NΔ		+-	loderate to intense nuclear staining reaction of neoplastic cells.
				of reaction of neoplastic cells.
gn melanocytic l	esions		Fai	int and weak reaction in few dispersed neoplastic cells. ense nuclear staining reaction of neoplastic relis.
			Inte	ense nuclear staining reaction
			We	ak to moderate intensity.
# Notes				
Focal and/or dispersed of			Wea	ak staining intensity.
Both positive cases with	focal staining		Wea	ak to strong intensity.
pattern.				
In two difficult cases, PR			Faint	staining reactions.
in dispersed melanocytic				
and in the second case, but diffuse reaction patt			Core	6d (weak intensity) and core 6f (strong intensity). Core 6e not sed (lacking).
regarded as positive usin			assess	sed (lacking).
>76% positive neoplastic				tore be not
4/16 (weak reaction of >				
neoplastic cells). The co		are		
without neoplastic cells,				
assessed (NA).				
penign non-lesional skin.				
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s/IHC compendium: J Cutan Pat	hol. 2021			
mas (nos) expressed PRAME, w	haraas only 2	504		
and with focal staining pattern	. This result is		Det	
uman protein Atlas (RNA			not as	ssessed due to lack of tumour cells. Dispersed
the test is 87% (malignant vs be			sermir	hal centre lymphatic cells are made
d of >76% positive neoplastic ce				bessed due to lack of tumour cells. Dispersed all centre lymphatic cells are positive of normal follicles.
increased to 94% with a sensitiv				- Juicies.
oose and thus, can be implemen	ited as routine			
an Pathol. 2021 Jul;48	(7):856-86	2		

PRAME: Ready for accreditatio

IHCbase RS_V30 (SV25009)

Records

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Lick is PRAUE Lick is PRAUE Product is Aci 32238 Product is Aci 32238 Product is Version Bouline use Version Version Version Product is Version Bouline use Version Version Product Bouline use Version Version Product Bouline use Version Product Product Bouline use Version Product Product Bouline use Version Bouline use Version Product Product Bouline use Version Bouline use Vers	Lick Review Particular Lick Review Particular Particular Consign (Galaci) Particular Consign (Galaci) Particular Consign (Galaci) Redine use Yes Bit Cyre Type Tumor classification Particular Consign (Galaci) Bit Cyre Type Tumor classification Particular Consign (Galaci) Bit Cyre Type Tumor classification Particular Particular Consign (Galaci) Consign (Galaci) Particular Particular Particular Part	Antibody		Printing full version of the validation report (Buttons below): 1) Print report.	Ladegaardsgade 3, 9000 Aalborg, Denmark
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Platform Ominis (Dako) Routine use Yes Interpretention Outpretention Interpretention Spaces Mathia Compary Boacs Mathia Boacs Mathia Interpretention Interpretention Space Mathia	Platform Omna (Daka) Routine usa Ve Work We Wick type Type 1. Tumor classification Wick type type 2. Type 2. Tumor classification Type 2. Tumor classification Poincold The step 1. Tumor classification classification Wick type type 2. Tumor classification	Product no.	ACI 3252B	expressed PRAME, whereas only 25% (5/20) of benign and atypical nevi were	For more information, see "External Quality Assurance"
Routine use Yes HC type Type 1. Tumor classification Company Bocare Medical Company Bocare Medical Purposef The dignosity used to discrimited Melanoma In Sity Intended use Yes Intended use The dignosity used to discrimited Melanoma In Sity Mito Appendix Company Company Bocare Medical Intended use Purposef The dignosity used to discrimited Melanoma In Sity The dignosity used to discrimited Melanoma In Sity Mito Appendix To statis to the protocol, see PDF format. Mito Appendix Optimal dit. 1200 Antigen Retrieval TRS High pH Visualization System EnvFLEX+(RM) Usualization System EnvFLEX+(RM) Later protocol change	Routine us vs HC type Type 1, Tumor classification Company Boxeline disclassification Company Boxeline disclassification New Identification Protocol Protocol Protocol Protocol And Rear Refines and/or dentify Optimal dit. 1:200 Andigen Retrieval Visualization System Interpreter Angline Retrieval Mindom regin Statistic of the protocol change Visualization System Interpreter No stating of egibleial cells of the appendix.	Platform	Omnis (Dako)	positive for PRAME - often in a minority of nuclei's and with focal staining pattern.	
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Company Blocase Medical Purpose/ PPANE is prinarily used to discriminate Melanoma In Situ on information and prison and or discrimate Melanoma in Situ on information origin. Protocol Protocol Port details to the protocol, see PDF format. Dituent Renor Red (Biocase; PD904) Optimal dit. 1:200 Midig netrieval TRS High pH Visualization System HiP/DAB 5' (Omnig) System HiP/DAB 5' (Omnig) Similiary of the tonall.	Company Boares Madical Purpose/ FRAME: is primarity used to discriminate Melanoma in Situ (non-morigin-melanocytic lacions) and a specificity of 95% (malignant via being melanocytic lacions) and a specificity of 95% (malignant via being melanocytic lacions) and a specificity of 95% (malignant via being melanocytic lacions) and a specificity of 95% (malignant via being melanocytic lacions) and a specificity of 95% (malignant via being melanocytic lacions) and a specificity of 95% (malignant via being melanocytic lacions) and a specificity of 95% (malignant via being melanocytic lacions) and a specificity of 95% (malignant via being melanocytic lacions) and a specificity of 95% (malignant via being melanocytic lacions) and a specificity of 95% (malignant via being melanocytic lacions) and a specificity of 95% (malignant via being melanocytic lacions) and a specificity of 95% (malignant via being melanocytic lacions) and a specificity of 95% (malignant via being melanocytic lacions) and a specificity of 95% (malignant via being melanocytic lacions) and a specificity of 95% (malignant via being melanocytic lacions) and a specificity of 95% (malignant via being melanocytic lacions) and a specificity of 95% (malignant via being melanocytic lacions) and a specificity of 95% (malignant via being melanocytic lacions) and a specificity of 95% (malignant via being melanocytic lacions) and a specificity of 95% (malignant via being melanocytic lacions) and a specificity of 95% (malignant via being melanocytic lacions) and a specificity of 95% (malignant via being melanocytic lacions) and a specificity of 95% (malignant via being melanocytic lacions) and a specificity of 95% (malignant via being melanocytic lacions) and a specificity of 95% (malignant via being melanocytic lacions) and a specificity of 95% (malignant via being melanocytic lacions) and a specificity of 95% (malignant via being melanocytic lacions) and a specificity of 95% (malignant via being melanocytic lacion	IHC type	Type 1, Tumor classification	benign melanocytic lesions) with a sensitivity of 96% and a specificity of 75%.	Depart status Assessed Date
Purpose FRAME is primitivy used to discriminate Melanona In Situ Intended use is sensitivity of 95% (malignant vs beign melanocytic lesions) and a specificity of 95%. The assay is fit-for-purpose and thus, can be implemented as routine analysis. Protocol Por details to the protocol, see PDF format. Diluent Report Red (Blocare: FD904) Optimal dit. 1:200 Antigen Retrieval TRS High PH v 24 min 97C Detection Syste EnvFLEX++ (RM) v 10:10:20 min 32C Antigen Retrieval Kigh Expressor Antigen Retrieval Usualization Syste HRP/DAB 5' (Omnis) v Smin 32C Latest protocol change	Purpose/ PPANE is prinarity used to discimize Malaxona in Situ for prinarity of 90% (malignant use heigh melanocytic lesions) and a specificity of 90%. The assay is fit-for-purpose and fus, can be implemented as routine analysis. Protocol Protocol For details to the protocol, see PDF format. Dituent Penoir Red (Biccare: PD904) Optimal dit. 1:200 Antigen Retrieval TRS High pH 24 min 97C. Detection None Visualization None Latest protocol change Non Expressor No staining of epithelial cells of the appendix.	Company	Biocare Medical V	diagnosis (see selected references/IHC compendium: J Cutan Pathol. 2021 Jul;48	
Protocol For details to the protocol, see PDF format. Dituent Renoir Red (Biocare; PD904) Optimal dil. 1:200 High Expressor A strong, distinct nuclear staining reaction of virtually all neoplastic cells of the melanoma. Laboratory medical directores: Laboratory medical directores: Mik8 Melanoma Antigen Retrieval TRS High pH 24 min 97C Detection System EnvFLEX++ (RM) 10_10_20 min 32C Antigen Retrieval None 5 min 32C Latest protocol change An at least faint to weak, distinct nuclear staining reaction of scattered lymphocytes An at least faint to weak, distinct nuclear staining reaction of scattered lymphocytes	Protocol For details to the protocol, see PDF format. Diluent Renoir Red (Biocare; PD904) Optimal dil. 1200 High Expressor Antigen Retrieval TRS High pH 24 min 97C Detection System EnvFLEX++ (RM) 10_10_20 min 32C Low Expressor Ant least faint 1 weak, diatinct nuclear staining reaction of scattered lymphocytes In at least faint 1 weak, diatinct nuclear staining reaction of scattered lymphocytes In at least faint 1 weak, diatinct nuclear staining reaction of scattered lymphocytes Non Expressor Non Expressor Non Expressor Non Expressor Not Expressor Non Expressor Non Expressor Not Expressor Non Expressor Not E		from benign melanocytic lesions and/or identify melanomas in the diagnostic work-up among tumors of	95%. The assay is fit-for-purpose and thus, can be implemented as routine analysis.	Report approved by
Protocol Melanoma Adrenal Gl. Brain Tissue_7 Diluent Renoir Red (Biocare; PD904) Melanoma Adrenal Gl. Brain Tissue_7 Optimal dil. 1:200 Melanoma Adrenal Gl. Brain Tissue_7 Antigen Retrieval TRS High pH v 24 min 97C Astrong, distinct nuclear staining reaction of virtually all neoplastic cells of the melanoma. None Visualization System InvFLEX++ (RM) v 10_10_20 min 32C Low Expressor Nat least faint to weak, distinct nuclear staining reaction of scattered lymphocytes in secondary folicides of the tonsil. None v Nat least faint to weak, distinct nuclear staining reaction of scattered lymphocytes in secondary folicides of the tonsil. None v None v <t< th=""><th>Protocol Melanoma Adrenal Gl. Brain Tissue_7 Diluent Renoir Red (Biocare; PD904) Melanoma Adrenal Gl. Brain Tissue_7 Optimal dil. 1:200 High Expressor A strong, distinct nuclear staining reaction of virtually all neoplastic cells of the melanoma. Melanoma A strong, distinct nuclear staining reaction of virtually all neoplastic cells of the melanoma. Melanoma Me</th><th></th><th></th><th></th><th></th></t<>	Protocol Melanoma Adrenal Gl. Brain Tissue_7 Diluent Renoir Red (Biocare; PD904) Melanoma Adrenal Gl. Brain Tissue_7 Optimal dil. 1:200 High Expressor A strong, distinct nuclear staining reaction of virtually all neoplastic cells of the melanoma. Melanoma A strong, distinct nuclear staining reaction of virtually all neoplastic cells of the melanoma. Melanoma Me				
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Latest protocol change	Latest protocol change Non Expressor No staining of epithelial cells of the appendix.				
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	Non Expressor No staining of epithelial cells of the appendix.			×	
indi expressor			× [Non Expressor	
No staining of epithelial cells of the appendix.	Scattered lymphocytes might be positive (e.g., lamina propria mucosa).			No staining of epithelial cells of the appendix.	

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Technical aspects of IHC and pitfalls – Analytical phase

Optimization of the IHC assay – issues to be addressed

- Purpose and/or "fit-for-purpose" (assay validated for intended use)
- How to establish "best practice protocol" of the IHC test (Calibration of the IHC assay with focus clone, antigen retrieval, titer & detection system)
 - Is the IHC test reproducible/robust (pre-analytic conditions)
 - Evaluation of the analytical sensitivity and specificity
- Identification of most robust controls providing information that the established level of detection is obtained in each test performed in daily practice.

Tissue materials are essential for all these processes (calibration, validation and controls)

Calcitonin optimization (data sheets ?)

nti-Calcitonin (SP17), Rabbit Monoclonal	
rimary Antibody Bocho / Vontana / Coll M	
Catalog Number: 760-4705 Ordering Code: 76586554001	/larque
Quantity: 50 tests	
Controls: Medullary Carcinoma of	
Thyroid Lecture LeC	
Isotypes: IgG Clone Name: SP17	
Species: Rabbit	
Localization: Cytoplasmic	
Regulatory Status: IVD	
Recommended staining protocol with ultraView Procedure Type Method	
Deparaffinization Selected	
Cell Conditioning (Antigen Unmasking) Cell Conditioning 1, Mild	
Enzyme (Protease) Nor required	
BenchMark ULTRA instrumer	nt:
16 minutes, 36℃	
Antibody (Primary) BenchMark XT instrument:	
16 minutes, 37°C	
BenchMark GX instrument:	
16 minutes, 37°C	
optimizing an assay can be confusing	
ptimizing an assay can be contusing	
hinutes	
Vendor recommendations)	



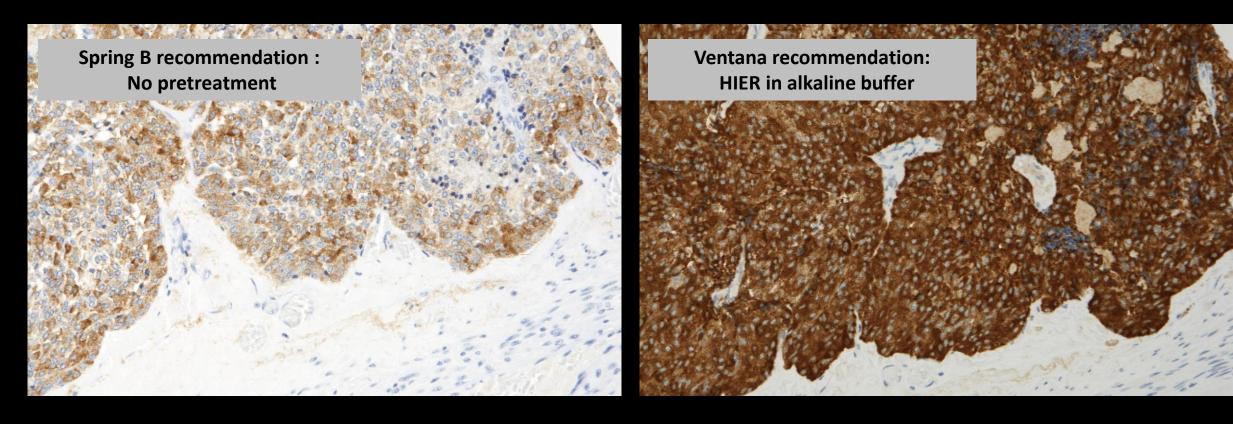
Rabbit Anti-Human Calcitonin Monoclonal Antibody (Clone SP17)

Spring Cat#	Roche P/N	Product Description	BREAK AND IN CONTRACTOR
M3170	05492769001	0.1 ml rabbit monoclonal antibody supplied as tissue culture supernatant in TBS/1% BSA buffer pH 7.5 with less than 0.1% sodium azide.	
M3172	06970419001	0.5 ml rabbit monoclonal antibody supplied as tissue culture supernatant in TBS/1% BSA buffer pH 7.5 with less than 0.1% sodium azide.	法法律公司
M3174	05298725001	 1.0 ml rabbit monoclonal antibody supplied as tissue culture supernatant in TBS/1% BSA buffer pH 7.5 with less than 0.1% sodium azide. 	
M3171	05298717001	7.0 ml pre-diluted rabbit monoclonal antibody supplied as tissue culture supernatant in TBS/1% BSA buffer pH 7.6 with less than 0.1% sodium azide. (For manual IHC only)	Human thyroid medullary carcinoma stained with anti-calcitonin antibody
INTENDED U	SE:	For Research Use Only. Not for use in diagnostic procedures.	
CLONE:		SP17	
IMMUNOGEN	1:	Synthetic human calcitonin 1-32 amino acid peptide.	
IG ISOTYPE:		Rabbbit IgG	
EPITOPE:		Not determined	
MOLECULAR	WEIGHT	15kDa	
SPECIES REA	ACTIVITY:	Human (tested). (See www.springbio.com for information on specie: sequence homology.)	s reactivity predicted by
DESCRIPTIO	N:	Calcitonin is a 32 amino acid peptide which can be demonstrated in hyperplastic thyroid. Staining for calcitonin may be used for the iden proliferative abnormalities ranging from C cell hyperplasia to invasiv in medullary carcinoma of the thyroid produces a fine granular patte deposits within the tumor may also exhibit varying degrees of calcit	ntification of a spectrum of C cell re tumors. Staining for calcitonin frm in the cytoplasm. Amyloid
APPLICATIO	NS:	Immunohistochemistry (IHC)	
IHC PROCED	URE:	Specimen Preparation: Formalin-fixed, paraffin-embedded tissues primary antibody.	s are suitable for use with this
		Deparaffinization: Deparaffinize slides using xylene or xylene alte Antibody Dilution: If using the concentrate format of this product, Antibody Diluent (Cat# ADS-125). The dilutions are estimates; actua veriability in methods and protocols.	dilute the antibody 1:100 in
		Antigen Retrieval: None Primary Antibody Incubation: Incubate for 30 minutes at room te	mperature.
		Slide Washing: Slides must be washed in between steps. Rinse s Detection: Detect the antibody as instructed by the instructions pro system.	lides with PBS/0.05% Tween.
POSITIVE CO	NTROL:	Thyroid medullary carcinoma	
CELLULAR L	OCALIZATION:	Cytoplasm	

Can we use the recommendations provided by the manufactures spec sheets?

Calcitonin: Vendor recommendations - spec sheets ?

Thyroid medullary carcinoma



Calcitonin, SP17 (1:800), Omnis, Flex+

Excellent IHC Staining. Always.

Make tissue arrays

Effectively recover epitopes
Eliminate non-specific staining

This is a site dedicated to achieving the excellence in immunohistochemistry staining. We offer unique products that allow you to get the best from your immunochemical research.

Our products are invented either by us or by our close partner companies.

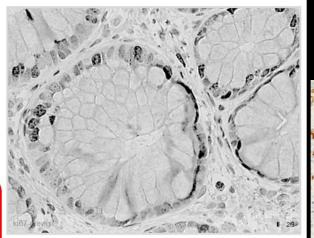
We help you to achieve standardization of IHC staining, first of all in research pathology, where many new and poorly studied antibodies are used.

Universal Buffer

We have done it: you do not need any more Tris, EDTA,

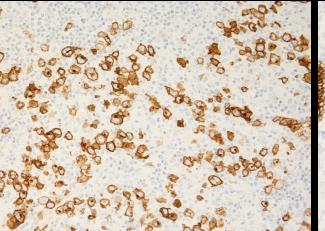
Citrate, Low of High pH buffers for epitope recovery. You do not even need proteas treatment of sections.

One R-Universal Buffer replaces all of them!

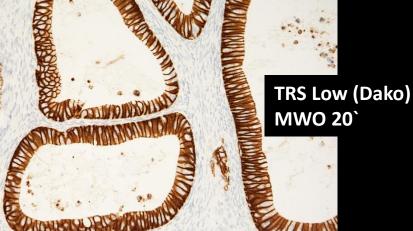


2100 Antigen Retriever. Probably the best epitope recovery unit for IHC on formalin-fixed sections. Used by hundreds of research groups and small pathology labs worldwide. Has

CD30, ConD6/D5 (1:50)

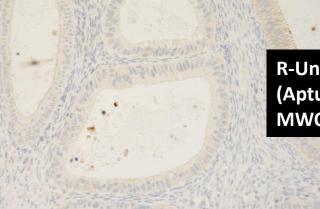


EP-CAM, MOC-31 (1:50)



Be critical – Do your own tests





R-Universal buffer (Aptum) MWO 20`

Hodgkin Lymphoma

Uterus / Endometrium

Optimizing IHC test (IHC-type 1 markers /conc. formats) - parameters to consider

Use a "Test Battery Approach" (TBA) (involves different pre-treatment regimes and dilution ranges)

Test more than one antibody clone against an antigen of interest before implementation for routine purpose

Test with robust, specific & sensitive detection system

Test/validate on normal and tumor tissue material with broad spectrum of antigen densities (specificity/sensitivity)

Compare results with external quality assurance programs, literature or colleagues

No antibody should be acquired without the basic knowledge of its performance characteristics and expected expression pattern Hadi Yaziji and Todd Barry – Adv Anat Pathol • Vol. 13, Number 5, September 2006

The "test battery approach" should be tailored to the individual IHC instruments or manual set-ups

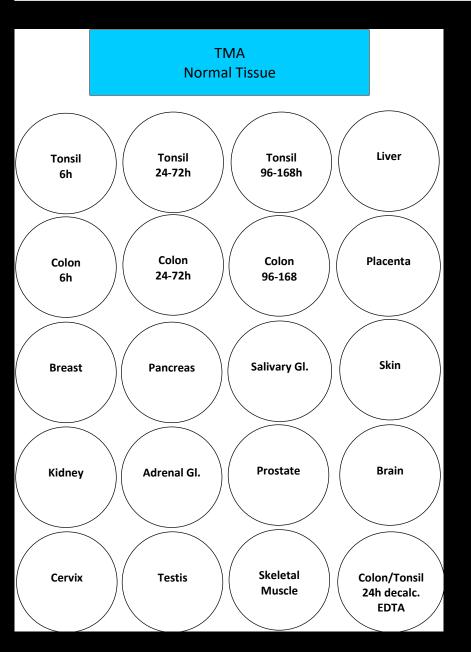
Omnis + Autostainer platforms

	Antibody Performa	Omnis		
	Dil. 1	Dil.2	Dil.3	Protocol A: 0.0 % Protocol B: 2.0 % Protocol C: 10.0 % Protocol D: 84.0 %
А	None	None	None	
В	Enzyme (1) 5 min.	Enzyme (1) 5 min.	Enzyme (1) 5 min.	Protocol E: 1.0 %
С	HIER TRS Low pH 6.1 (30`)	HIER TRS Low pH 6.1 (30`)	HIER TRS Low pH 6.1 (30`)	Protocol F: 3.0 %
D	HIER TRS High pH 9.0 (24`)	HIER TRS High pH 9.0 (24`)	HIER TRS High pH 9.0 (24`)	
E F	TRS Low (20`) + Pep (12`) * HIER TRS High pH 9.0 (30`)	TRS Low (20`) + Pep (12`) HIER TRS High pH 9.0 (30`)	TRS Low (20`) + Pep (12`) HIER TRS High pH 9.0 (30`)	Routine purpose : App 220 mark

kers

* Combined pre-treatment using Cytology Pepsin solution (Zytovision, #ES-0002-50)

Identify the protocol that discriminate between the desired (specific) positive staining and any unwanted (non-specific) background staining



Protocol set-up: Evaluate analytic sensitivity and specificity

Normal tissue including fixation and decalcification controls

Identification of the best practice protocol: Test Battery Approach (clone, titer, retrieval etc.)

SOX10, BS7 (dil. range 1:50-800): HIER High pH 24`/ 1:350 (after re-test) / EnvFlex+

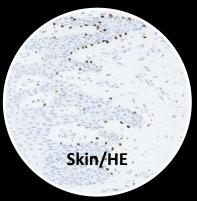
Establishing robustness of the IHC assay / pre-analytic parameter`s ?

SOX10, BS7; Robust to both fixation time in NBF and decalcification

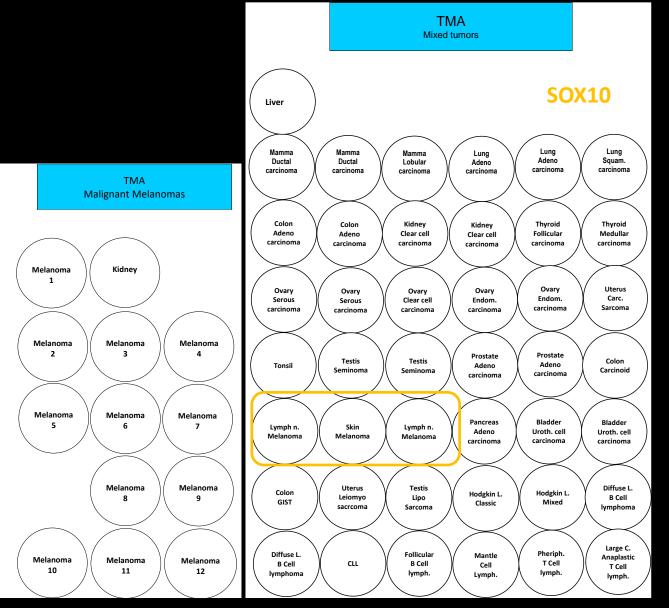
Identification of robust controls

SOX10, BS7; IHC Critical Assay performance Controls (high, low and non exp.)

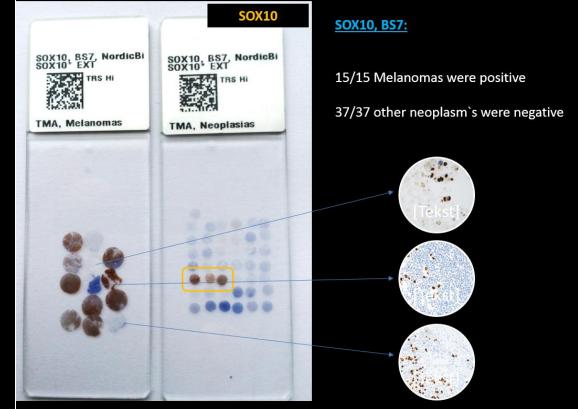








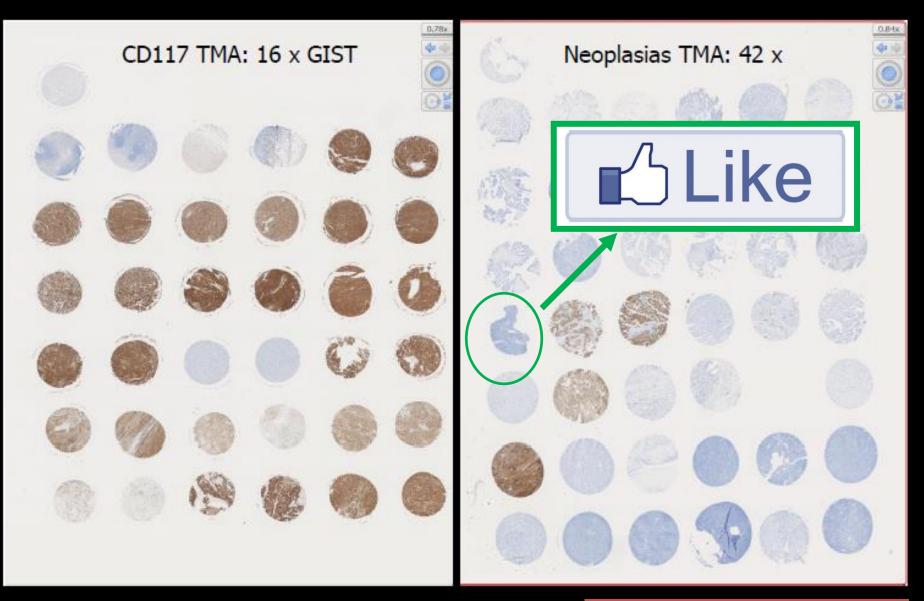
Diagnostic potential Analytical validation



Other SOX10+ tumours: Schwannoma's, neurofibroma's, myoepithelial carcinomas, triple negative breast cancer.....?

IHC – The Technical Test Approach





The technical test approach – Analytical phase

Analytical Validation - Evaluation of sensitivity and specificity

Recommendations for Improved Standardization of Immunohistochemistry

Neal S. Goldstein, MD, Stephen M. Hewitt, MD, PhD, Clive R. Taylor, MD, DPhil, Hadi Yaziji, MD, David G. Hicks, MD, and Members of Ad-Hoc Committee On Immunohistochemistry Standardization

Abstract: Immunohistochemistry (IHC) continues to suffer from variable consistency, poor reproducibility, quality assurance disparities, and the lack of standardization resulting in poor concordance, validation, and verification. This document lists the recommendations made by the Ad-Hoc Committee on Immunohistochemistry Standardization to address these deficiencies. Contributing factors were established to be underfixation and irregular fixation, use of nonformalin fixatives and ancillary fixation procedures divested from a deep and full understanding of the IHC assay parameters, minimal or absent IHC assay optimization and validation procedures, and lack of a standard system of interpretation and reporting. Definitions and detailed guidelines pertaining to these areas are provided.

Key Words: immunohistochemistry, pathology, assay, oncology, standardization, procedure, tissue, fixation

(Appl Immunohistochem Mol Morphol 2007;15:124-133)

IHC assay standardization was vital for reproducible and reliable results. Agencies, including the Biologic Stain Commission, CLSI (previously NACCLS), FDA, and the manufacturing sector established guidelines, standards, and recommendations for reagents and package inserts. These efforts have resulted in consistent, high-quality assay components and instruments on which contempor-

ary IHC is performed.¹⁻⁴ It has also a development and use of so-called black box II in which IHC assays have preset parameters manufacturer.⁵

Despite the improvements of reagents at tion, authors over the years have consistently inconsistent quality of IHC assays.⁶⁻¹¹ Unlik IHC-epochs, most of the causative responsis with the individual laboratory performing th specifically, the lack of standardization and a quality assurance programs.^{12,13} Prior consen ences identified the likely causative factors (

How many tissue samples are needed for the analytical validation process ?

Goldstein NS et al : Appl Immunohistochem Mol Morphol 2007 Mar; 15 : 124-133

25 tissue samples (Non-predictive markers/ IHC-type I: 10 high, 10 low and 5 non-expressors)

Principles of Analytic Validation of Immunohistochemical Assays

Guideline From the College of American Pathologists Pathology and Laboratory Quality Center

Patrick L. Fitzgibbons, MD; Linda A. Bradley, PhD; Lisa A. Fatheree, BS, SCT(ASCP); Randa Alsabeh, MD; Regan S. Fulton, MD, PhD; Jeffrey D. Goldsmith, MD; Thomas S. Haas, DO; Rouzan G. Karabakhtsian, MD, PhD; Patti A. Loykasek, HT(ASCP); Monna J. Marolt, MD; Steven S. Shen, MD, PhD; Anthony T. Smith, MLS; Paul E. Swanson, MD

 Context.—Laboratories must validate all assays before they can be used to test patient specimens, but currently there are no evidence-based guidelines regarding validation of immunohistochemical assays.

Objective.—To develop recommendations for initial analytic validation and revalidation of immunohistochemical assays.

Design.—The College of American Pathologists Pathology and Laboratory Quality Center convened a panel of pathologists and histotechnologists with expertise in immunohistochemistry to develop validation recommendations. A systematic evidence review was conducted to address key questions. Electronic searches identified 1463 publications, of which 126 met inclusion criteria and were extracted. Individual publications were graded for quality, and the key question findings for strength of evidence. Recommendations were derived from strength of evidence,

Res to hel Fitzgibbons PL et al : Arch Pathol Lab Med 2014;138:1432-1443

20 tissue samples (Non-predictive markers/IHC-type I: 10 positive and 10 negative cases apply including high & low expressors)

levels

revali

that ii 40 tissue samples (predictive markers/IHC-type 2: 20 positive and 20 negative cases

also provided for confirming assay performance when there are changes in test methods, reagents, or equipment. (*Arch Pathol Lab Med.* 2014;138:1432–1443; doi: 10.5858/arpa.2013-0610-CP)

Accepted for publication February 3, 2014. Published as an Early Online Release March 19, 2014 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

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

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 Jason L. Hornick, MD, PhD,## Merdol Ibrahim, PhD,*** Antonio Marchetti, MD, PhD,†††
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 Paul E. Swanson, MD, ¶¶¶ Mogens Vyberg, MD,§§§### Xiaoge Zhou, MD,###****
 and Clive R. Taylor, MD,†††
 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

Emina E. Torlakovic, MD, PhD,*†‡ 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,¶¶¶ Mogens Vyberg, MD,§§§### Xiaoge Zhou, MD,###**** Clive R. Taylor, MD,†††† and From the International Society for Immunohistochemistry and Molecular Morphology (ISIMM) and International Quality Network for Pathology (IQN Path)

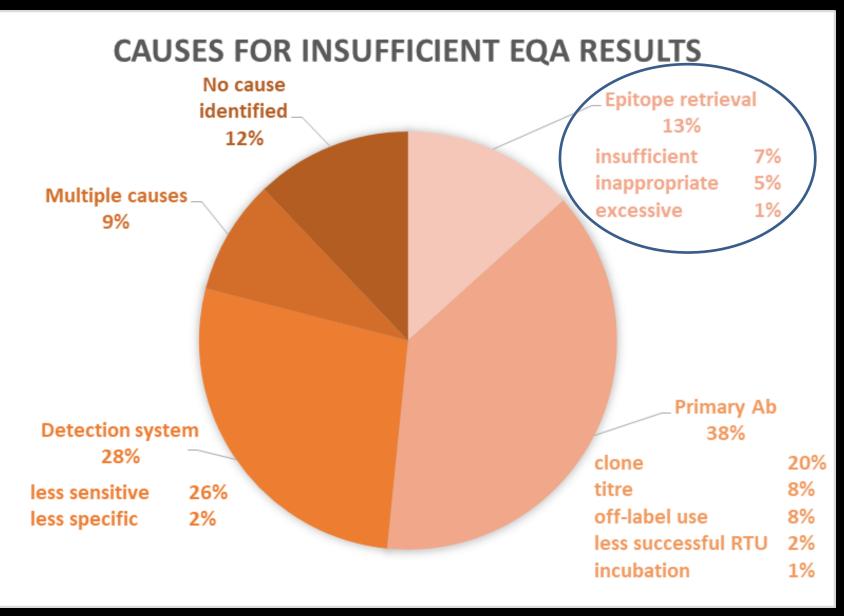
Evolution of Quality Assurance for Clinical Immunohistochemistry in the Era of Precision Medicine: Part 4: Tissue Tools for Quality Assurance in Immunohistochemistry

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Article sequence (part 1-4) published in Appl Immunohistochem Mod Morphol (2017) systematically describing/defining all aspects of the IHC test - what is required for a full technical validation and implementation of a new test.

Full technical validation

NordiQC External Quality Assurance program



App. 20-30% of all results are evaluated as insufficient

79 % of insufficient results are related to the choice/use of:

- Epitope retrieval procedure
- Primary Ab (including stainer platform)
- Detection system

False positive and/or negative results

Courtesy of Birgit Truumees et al (NordiQC): Poster presented at the annual USCAP meeting 2024

The purpose of antigen retrieval is to unmask antigenic determinants (epitopes) and recover immuno-reactivity

Antigen retrieval procedures for formalin fixed tissue:

- □ <u>Heat</u> Induced Epitope <u>Retrieval</u> (HIER)
- **Tissue digestion using proteolytic enzymes**
- **Combined pre-treatment (HIER with proteolytic digestion)**

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Vol. 39, No. 6, pp. 741-748, 1991 Printed in U.S.A.

Rapid Communication

Antigen Retrieval in Formalin-fixed, Paraffin-embedded Tissues: An Enhancement Method for Immunohistochemical Staining Based on Microwave Oven Heating of Tissue Sections

SHAN-RONG SHI, MARC E. KEY,¹ and KRISHAN L. KALRA

BioGenex Labora Shi et al. demonstrated that :

Received for publ

We describe a n formalin-fixed,

- Enzyme pre-digestion of tissue could be omitted.
- Incubation time with primary antibodies could be reduced, or dilutions of primary antibodies could be increased.
- Staining could be achieved on long-term formalin fixed tissue that failed to stain with conventional methods.
- Antibodies which where typically unreactive with formalin-fixed tissue gave excellent staining.



The mechanism of HIER is not completely understood, and several hypothesizes has been proposed:

Heating tissue sections in an appropriate buffer may unmask epitopes by :

- Hydrolysis/disruption of methylene cross-links formed by formalin fixation
- **Extraction of diffusible blocking proteins**
- **Precipitation/denaturation of proteins**
- **Rehydration of the tissue section allowing better penetration of the antibody**
- Removal of tissue-bound calcium ions by chelating substances
- **Other mechanism's ?**

ORIGINAL PAPER

Hypothesis for the mechanism for heat-induced antigen retrieval occurring on fresh frozen sections without formalin-fixation in immunohistochemistry

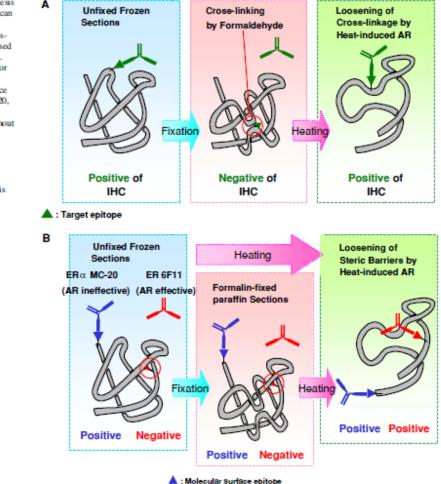
Kochi Kakimoto · Susumu Takekoshi · Katsuhiro Miyajima · R. Yoshiyuki Osamura

Demonstrated that: unfixed frozen sections, which did not show immunostaining with nine antibodies, were clearly stained after heating the sections

These results indicate that other mechanisms than breaking formalininduced cross-linkages may be present.

The authors proposed that : the accessibility to some target epitopes of antigenic proteins is limited by natural steric barriers even in the fresh state (frozen sections) caused by the antigenic protein itself

Fig. 7 Conventional hypothesis (A). Formaldehyde fixation can alter the three-dimensional structure of the epitope crosslinkages; these can be reversed by high-temperature heating. Our suggested mechanism for AR in IHC (B): Antibodies recognizing molecular surface epitopes, such as ERa MC-20, do not show increases in detection levels with or without heating whereas antibodies recognizing intramolecular epitopes, such as ER 6F11, show significantly increased detection levels because the three-dimensional structure is likely to be altered by heat denaturation



Intramolecularebitob

Target epitope

HIER buffers used by the vast majority of NordiQC participants

In house	Agilent Dako AS/Omnis	Roche Ventana Benchmark	Leica Biosystems Bond	Biocare	Thermo S LAB Vision
Low pH buffers					
Citric based buffers pH 6-6.7	TRS Low (3-in-1) pH 6.1	СС2 рН 6	BERS-1 pH 6	Diva Decloaker pH 6.2	
High pH buffer					
EDTA/EGTA based buffers pH 8-9	TRS High (3-in-1) pH 9	СС1 рН 8.5	BERS-2 pH 9	Borg Decloaker pH 9.5	HIER buffer H pH 9
	TRS High pH 9				

App. 95 % of all pretreatment protocols

Challenges:

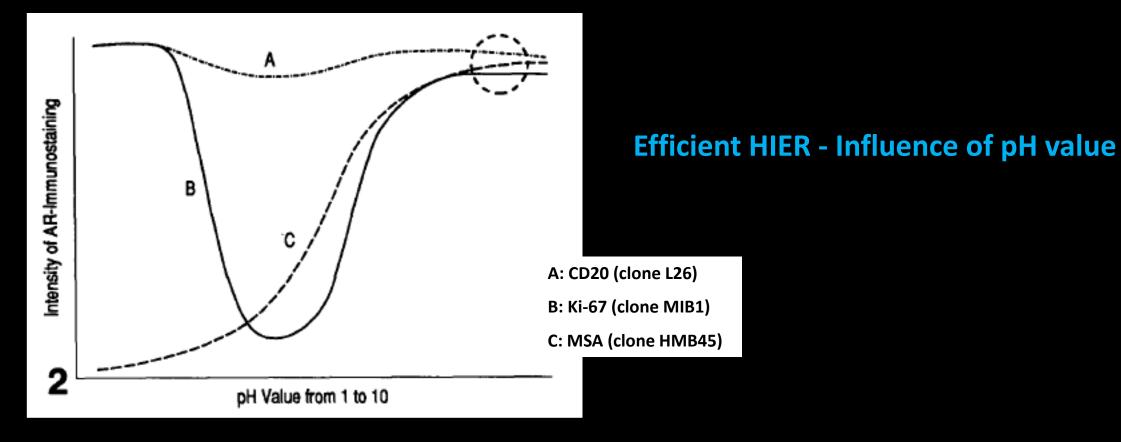
The platform often dictates the choice of HIER buffers

Efficient HIER depends on several parameters:

- **D** pH of the HIER buffer
- **Temperature**
- **Time**
- □ Chemicals/components of the HIER buffer (e.g., Citrate; TRIS; EDTA; TRIS-EDTA)

Less sensitive to routinely fixed tissue (formalin) compared to enzymatic pre-treatment

> 95% of all commonly used antibodies require HIER



Shi SR et al. J Histochem Cytochem 1995 43:193-201

Demonstrated that the performance of monoclonal antibodies were highly influenced by pH of the Antigen Retrieval buffer (AR).

Also, the results indicate the advantage of using an AR solution of higher pH value (8-9).

Efficient HIER - Influence of pH

HIER in TRS pH 6.1 (20 min at 97°C)

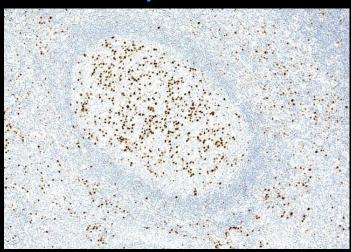
CD79, JCB117 (1:300)

MUM-1, MUM1p (1:400)



Autostainer: Flex+ as the detection system





HIER in TRS pH 9 (20 min at 97°C)

Tonsillar tissue fixed in 10% NBF (48h).

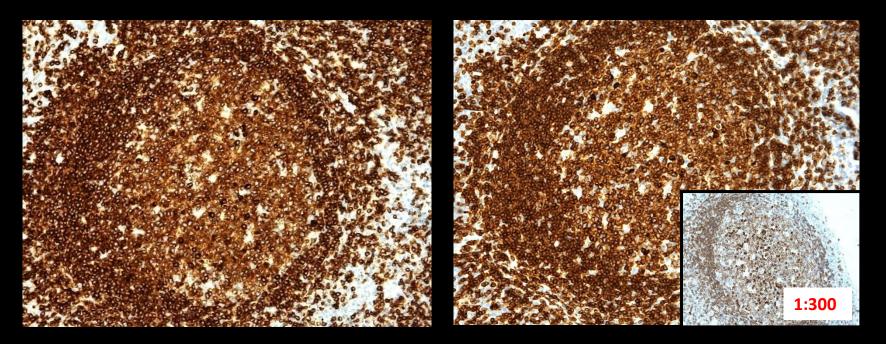
Efficient HIER - Influence of pH

HIER in TRS pH 9

HIER in TRS pH 6.1

CD79, JCB117 (1:300)

CD79, JCB117 (1:50)



Tonsillar tissue fixed in 10% NBF (48h). Flex+ as detection system

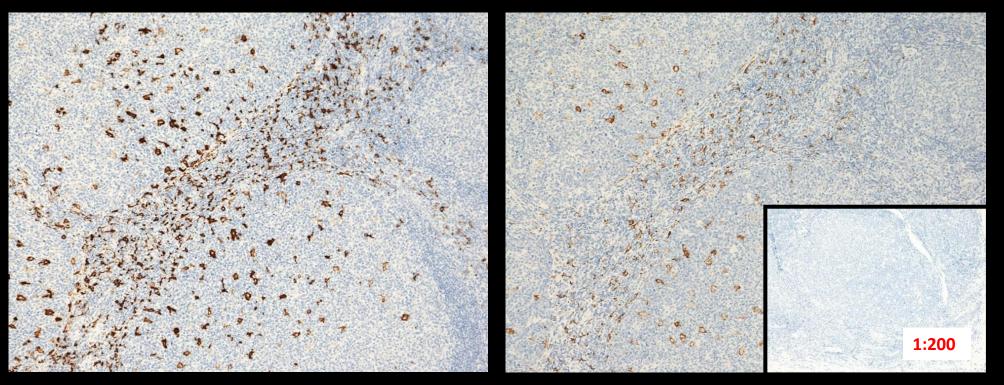
Efficient HIER - Influence of pH

HIER in TRS pH 9

CD163, MRQ-26 (1:200)/Flex+

HIER in TRS pH 6.1

CD163, MRQ-26 (1:25)/Flex+



For app. 85-90% of the epitopes, HIER in buffers at pH 8-9 is preferable to pH6

Ready To Use products: Recommendations to antigen retrieval provided by the manufactures ?

CK7 Run 62

Table 3. Proportion o RTU systems	Recom	imal results for CK7 mended of settings*	for the most commonly used RTU IHC sys Laboratory modified protocol settings**			
	Sufficient	Optimal	Sufficient	Optimal		
Dako AS mAb OV-TL 12/30 IR619	100% (12/12)	92% (11/12)	100% (11/11)	91% (10/11)		
Dako Omnis mAb OV-TL 12/30 GA619	100% (32/32)	97% (31/32)	100% (26/26)	92% (24/26)		
Leica Bond III/MAX mAb RN7 PA0942/PA0138	100% (6/6)	17% (1/6)	100% (11/11)	45% (5/11)		
VMS Ultra/XT rmAb SP52 790-4462	100% (16/16)	69% (11/16)	97% (98/101)	85% (86/101)		

* 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.

17% (1/6) BERS1 (citrate-based HIER buffer)

83% (5/6) BERS2 (alkaline based HIER buffer)

Efficient HIER - Influence of time and temperature

<u>Taylor CR et al : Applied Immunohistochemistry 1996; 4(3) : 144-166</u> - Temperature and time are inversely related :

Similar strong intensity of staining could be generated by the following heating conditions:

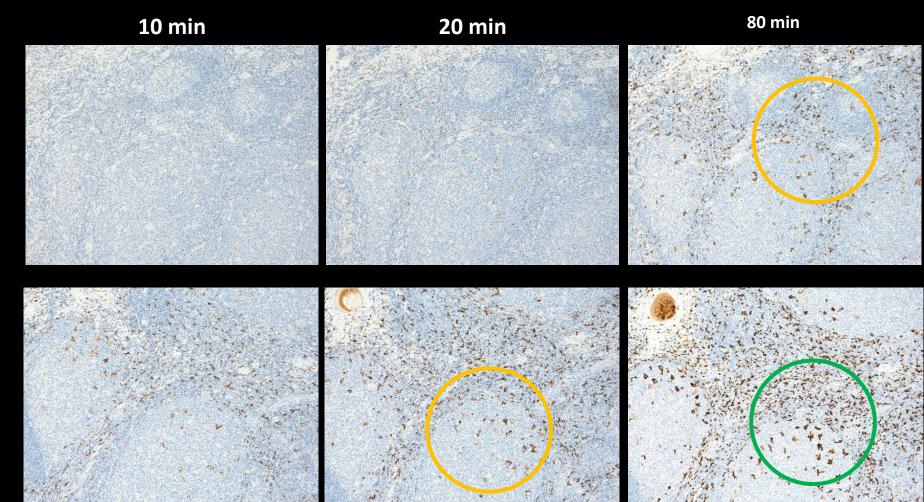
100°C for 20 min = 90°C for 30 min = 80°C for 50 min = 70°C for 10 h

Balaton AJ et al : Applied Immunohistochemistry 1996; 4(4) : 259 - 263

Optimal staining intensity could be generated by the following heating conditions:

Pressure cooker at 120°C for 3 min = MWO at 100°C for 20 min

HIER buffer - Influence of time and temperature



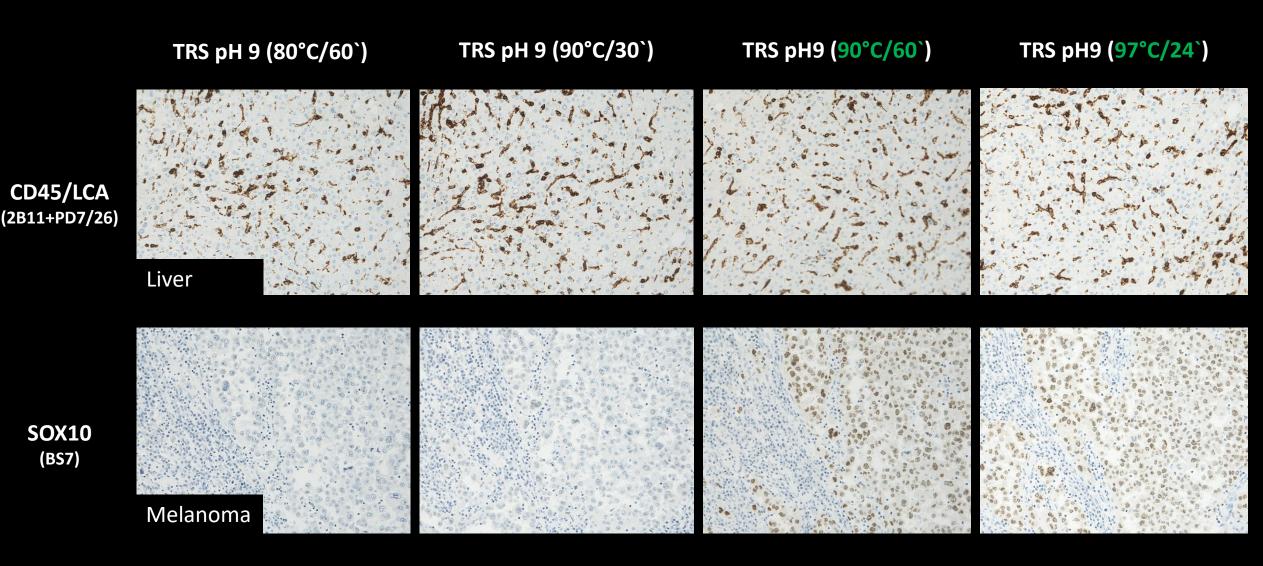
CD163, MRQ-26 (1:200)

HIER at 80°C TRS pH 9, Flex+

HIER at 97°C TRS pH 9, Flex+

Tonsillar tissue fixed in 10% formalin (48h).

Efficient HIER - Influence of time and temperature



ANTIGEN RETRIEVAL TECHNIQUES IN IMMUNOHISTOCHEMISTRY: COMPARISON OF DIFFERENT METHODS

STEFANO A. PILERI^{1*}, GIOVANNA RONCADOR¹, CLAUDIO CECCARELLI¹, MILENA PICCIOLI¹, ASPASIA BRISKOMATIS¹, ELENA SABATTINI¹, STEFANO ASCANI¹, DONATELLA SANTINI¹, PIER PAOLO PICCALUGA¹, ORNELLA LEONE¹, STEFANIA DAMIANI¹, CESARINA ERCOLESSI¹, FEDERICA SANDRI¹, FEDERICA PIERI¹, LORENZO LEONCINI² AND BRUNANGELO FALINI³

> ¹Second Service of Pathologic Anatomy and Haematopathology Section, Bologna University, Italy ²Institute of Pathologic Anatomy, Siena University, Italy ³Haematopathology Laboratory, Institute of Haematology, Perugia University, Italy

Chemical components of the HIER buffer`s

Standard low pH buffer`s (e.g., citrate based pH 6.0)

Standard high pH buffer`s (e.g., TE based pH 8-10)

Modified low pH buffers pH 6.1-6.2 : S1699/S1700 (Dako) or Diva Decloaker (Biocare)

Overall best performance:

HIER in EDTA pH 8.0 (compare with Tris-HCL pH 8.0)

118

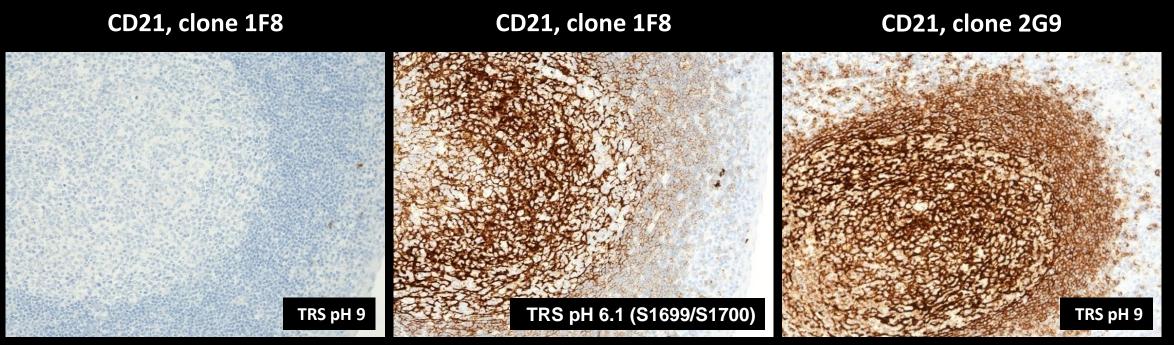
Table I—Results obtained under different methodological conditions with the antibodies most commonly used in the study of the haemolymphopoietic system and related disorders

Clone	Specificity	Source	Dilution	No AgR	PT	HBAR+ citrate	HBAR+ Tris-HCI pH8	HBAR+ EDTA pH8
O10	CD1a	Immunotech	1:40	_	_	+	++	++++
Poly	CD3	DAKO	1:300	_	++	+	++	++++
C8/144B	CD8	Dr Mason	1:6	_	_	++	++	++++
			1:400	_	_	+	+++-	++++
C3D-1	CD15	DAKO	1:6	+	_	+	++	++++
L26	CD20	DAKO	1:320 1:200	++		+++-	+++	++++
1.20	0.020	DAILO	1-3200	+	+	+++-	+++=	++++
IF8	CD21	DAKO	1:10	_	++++	-	-	-
IVIT IIVIO	CD23	DAKO	1.50			+	+++	++++
Ber-H2	CD30	Professor Stein	1:10	_	_	++	+++-	++++
			1:320	_	_	+	+++-	++++
QBEND-10	CD34	BioGenex	1:20	+	_	+++-	+++-	++++
			1:400	+	_	++	++	++++
BerMACDRC	CD35	DAKO	1:5	+	++++	_	+	+
MIAD03	CD40	Infinitiotech	1.100		++++			_
DF-T1	CD43	DAKO	1:200	+	_	+++-	+++-	++++
000000000000000000000000000000000000000	CD45	DAKO	1:1600	++	_	+++-	++++	++++
PD7/26+2B11	CD45	DAKO	1:200	_	+	+++-	++++	++++
LICHT 1	CD45D0	DAKO	1:4000	_	_	+	+++-	++++
UCHL-1 K1-B3	CD45R0 CD45R	Professor Parwaresch	1:120 1:80	+	++	++	++++	+++-
KI-B3	CD45R	Professor Parwaresch	1:320	++	+	+++-	++++	++++
4KB5	CD45RA	DAKO	1:20		+		++++-	
40.00	CD45RA CD57	Becton	1:20	++	++	++++	+++-	+++-
Y2/51	CD61	DAKO	1:5	++	+++-	+	+	+++-
12/31 NF1	CD01	DAKO	1.040		+++	+++++	+	++++
PG-MI	CD68	Professor Falini	1:20	+	++	++	++	++++
ICB117	CD79a	Dr Mason	1:10	+		+++-	+++-	++++
Kim-4p	Follicular dendritic cells		1:5	·	++++	++	++	+
DBA.44	Hairy cells	Professor Delsol	1:5	++	_	++++	+++-	++++
IC159	GlycophorinA	DAKO	1:320	+	_	++++	+++-	+++-
A TRACT		DAKO	1:10	++++	_	_	_	_
NP57	Neutrophilic elastase	DAINO						
M616	Neutrophilic elastase FVIII RAg	DAKO	1:6	+	++	++++	++	++++
					++ +++ -	++++ ++++	++ ++++	++++
M616	FVIII RAg	DAKO	1:6	+				
M616 Poly	FVIII RAg Lysozyme	DAKO DAKO	1:6 1:800	+++	+++-	++++	++++	++++
M616 Poly Poly	FVIII RAg Lysozyme IgA	DAKO DAKO DAKO	1:6 1:800 1:2000	+ ++ +	+++- +++-	++++ ++++	+++++ ++++	++++ ++++
M616 Poly Poly Poly	FVIII RAg Lysozyme IgA IgG	DAKO DAKO DAKO DAKO	1:6 1:800 1:2000 1:5000	+ ++ +	++++- ++++- +++++	++++ ++++ ++++	++++ +++- ++++	+++++ +++++ +++++
M616 Poly Poly Poly Poly	FVIII RAg Lysozyme IgA IgG IgM	DAKO DAKO DAKO DAKO DAKO	1:6 1:800 1:2000 1:5000 1:5000	+ ++ +	++++- ++++- +++++	++++ ++++ ++++ ++++	++++ +++- ++++ ++++	++++ ++++ ++++ ++++
M616 Poly Poly Poly Poly Poly	FVIII RAg Lysozyme IgA IgG IgM IgD	DAKO DAKO DAKO DAKO DAKO DAKO	1:6 1:800 1:2000 1:5000 1:5000 1:1000	+ ++ + ++ -	+++- +++- +++++ ++	++++ ++++ ++++ ++++	++++ +++- ++++ ++++	++++ ++++ ++++ ++++
M616 Poly Poly Poly Poly Poly Poly	FVIII RAg Lysozyme IgA IgG IgM IgD K-Ig light chain	DAKO DAKO DAKO DAKO DAKO DAKO DAKO	1:6 1:800 1:2000 1:5000 1:5000 1:1000 1:10 000	+ ++ ++ - +++	+++- ++++- +++++ +++ - ++++-	++++ ++++ ++++ ++++ ++++	++++ +++- ++++ ++++ +++- +++-	++++ ++++ ++++ ++++ ++++

CD-cluster of differentiation; No AgR-no antigen retrieval; PT-proteolytic treatment; HBAR-heat-based antigen retrieval; Poly-polyclonal antibody; FVIII RAg-Factor VIII-related antigen; MPO-myeloperoxidase.

In bold: overnight incubation of the primary antibody+SABC technique.

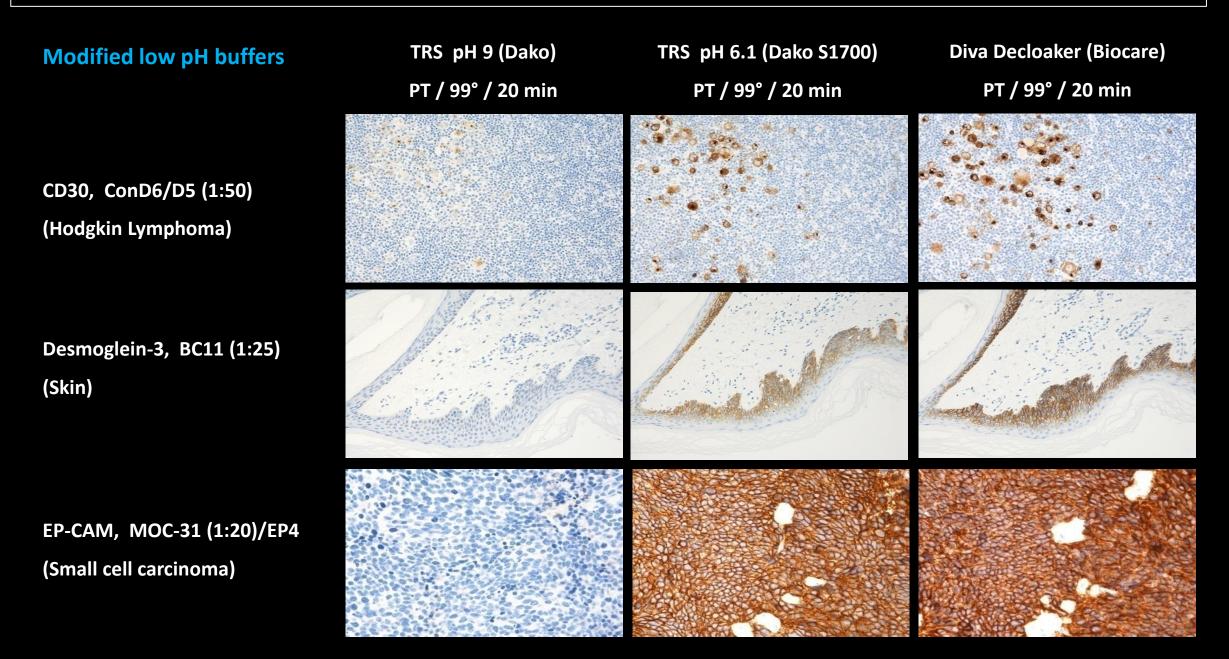
Modified low pH buffers



Markers requiring the TRS Low pH 6.1 (Dako, S1699/S1700) or Diva Decloaker pH 6.2 (Biocare, DV2004) :

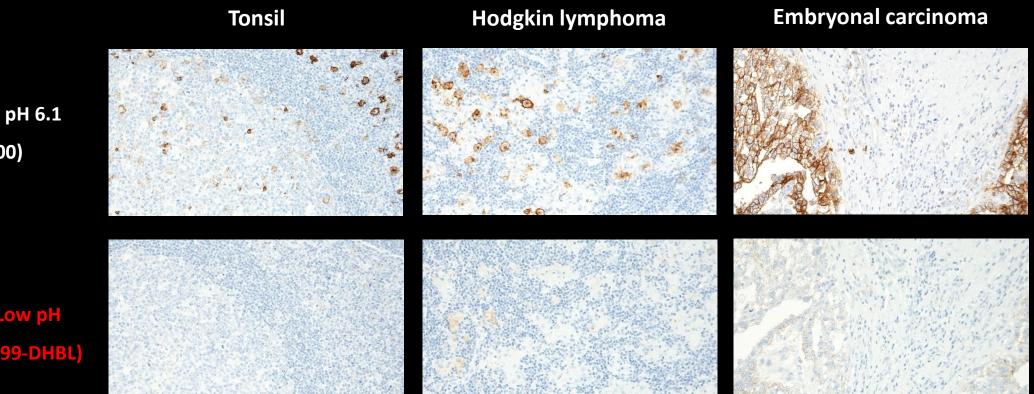
EPCAM (clone EP-4 or MOC-31 or "VU-1D9"); GP200 (clone SPM 314 or 66.4.C2); CD21 clone 1F8; CD61 clone Y2/51; NGFR clone MRQ-21; Desmoglein-3 clone BC11, PHH3 clone BC37 and

Mandatory for : CD7 clone CBC 3.7; CD30 clone ConD6/B5; CD5 clone Leu1.....



Modified low pH buffers

CD30 clone ConD6/B5



HIER buffer, TRS pH 6.1 (Dako S 1700)

HIER buffer, Low pH (LabVision TA-999-DHBL)

The purpose of antigen retrieval is to unmask antigen epitopes /restore antigenic determinants and recover immuno-reactivity

Antigen retrieval procedures for formalin fixed tissue:

- □ <u>Heat</u> Induced Epitope <u>R</u>etrieval (HIER)
- **Tissue digestion using proteolytic enzymes**



Combined pre-treatment (HIER with proteolytic digestion)

Proteolytic enzymes cleave more or less specific amino acid sequences within peptide chains.

→ Improves penetration of immuno-reagents into tissue structures and enhance accessibility to the epitopes of interest.

Enzymatic digestion - the problem?

A significant proportion of Labs still use enzymatic pre-treatment for e.g, "old" markers as cytokeratins and S100 introduced back in 1980-1990.

Only few markers require enzymatic digestion for "optimal performance".

022:				Vendor re	ecommenc	led protoc	ol setting	
Table 4. Pass epitope retri			e RTU DE-R-	11 antibody Pass rate	on the Bend	hmark platf	orm for diffe	erent
	Total		HIER		Prote	olysis	/ HIER + proteolysis	
	Protocols	Sufficient	Protocols	Sufficient	Protocols	Sufficient	Protocols	Sufficient
mAb DE-R-11 760-2513	139	98 (71%)	74	73 (99%)	49	11 (22%)	16	13 (81%)



"Optimal" enzymatic digestion depends on:

Enzyme type Concentration <u>Most common Enzymes</u> Time Proteinase K Pronase XIV Temperature Pronase XXIV Pepsin Fixation type & time Trypsin

Short time formalin fixation = gentle proteolysis Long time formalin fixation = prolonged proteolysis Difficult to control and to standardizes within routine LAB

Markers requiring enzymatic pretreatment :

FVIII (poly), LMV CK (CAM 5.2), PAN CK (MNF116), EGFR (various), TCR-β (8A3).....

Extracellulare matrix proteins (COLL-III (poly), Laminin (poly) and COLL-IV (CIV-22)

≤ 2% of all commonly used antibodies require enzymatic pre-treatment

Most common enzymes used in IHC:

Enzyme	Typical working conc.	Activation Temperature	Typical Incubation time	Cleavage nature
Proteinase K	0.1%, pH 8.0	25-37 °C	5-10 min.	Broad - all amino acids
Trypsin	0.1-0.25%, pH 7.6	37 °C	10 min.	Arginin / Lysin
Pepsin	0.2-0.4%, pH 2.0	37 °C	5-20min.	Broad ,favor peptides with aromatic amino-groups
Protease XXIV	0.05-0.1%, pH 7.6	37 °C	5-10 min.	Broad - all amino acids
Protease XIV	0.05-0.1%,pH 7.6	25-37 °C	10-30min.	Broad, favor peptides with aromatic residues

Choice of proteolytic enzyme

TCR-β, 8A3, 1:200 RR

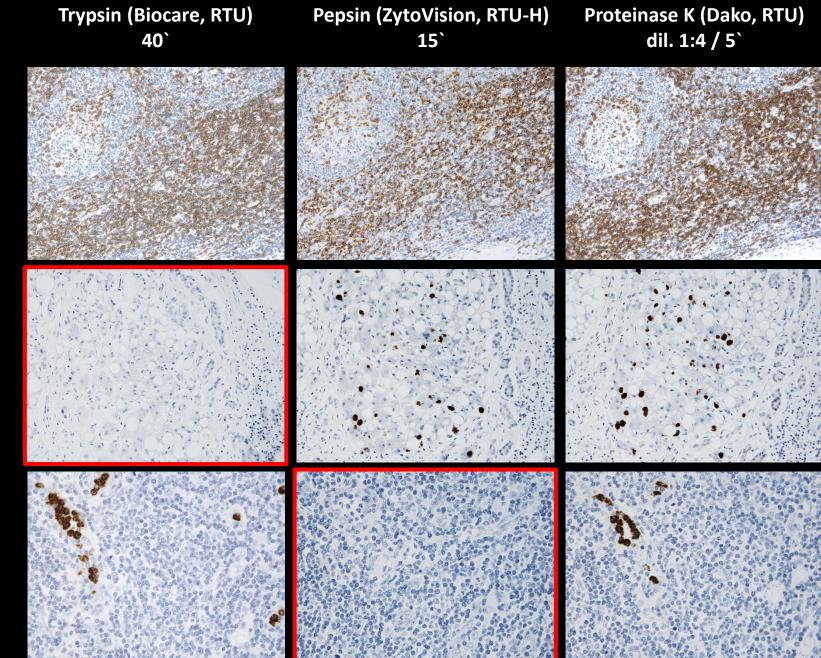
Tonsil

Ubiquitin, Ubi-1 1:750

Liver/ Mallory bodies

Neutrophil Elastase, NP57 1:1000

Tonsil



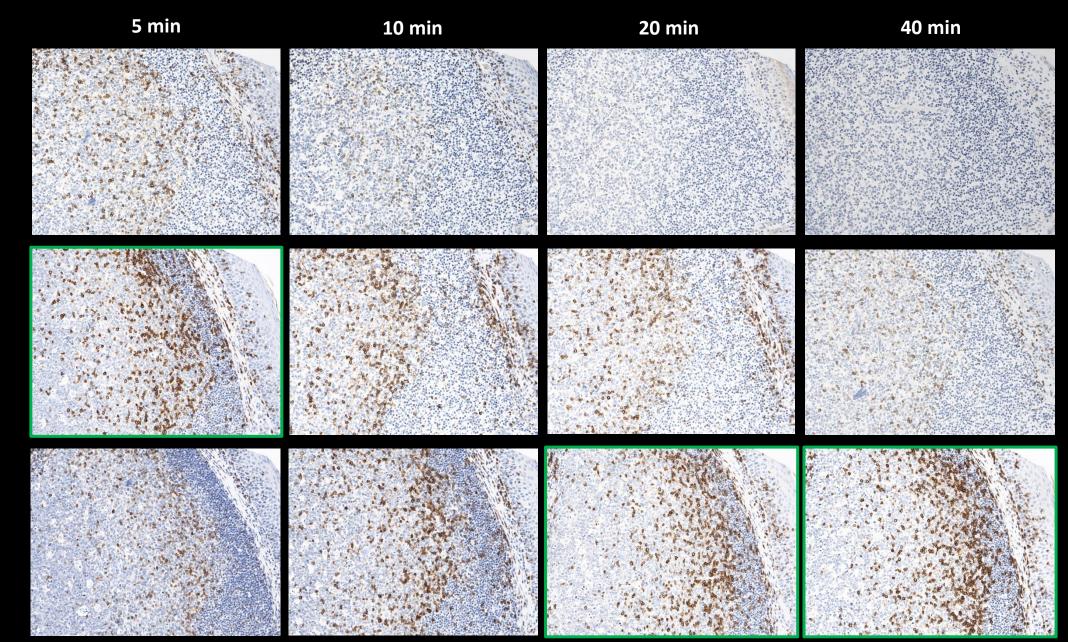
Proteolytic enzyme & digestion time ?

Tonsil NBF 48h Digestion temp. 32°C

Proteinase K (RTU S3020, Dako)

Proteinase K dil. 1:4 (RTU S3020, Dako)

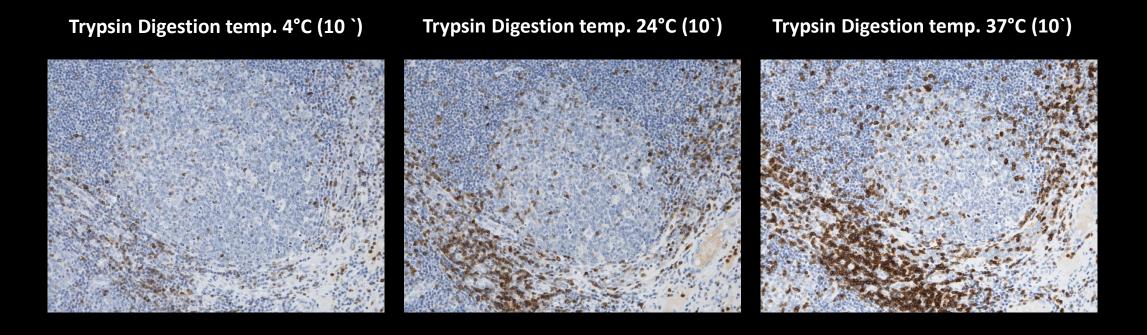
Trypsin (RTU, Biocare)



TCR-β clone 8A3 (1:200 RR) / Flex+ (Omnis)

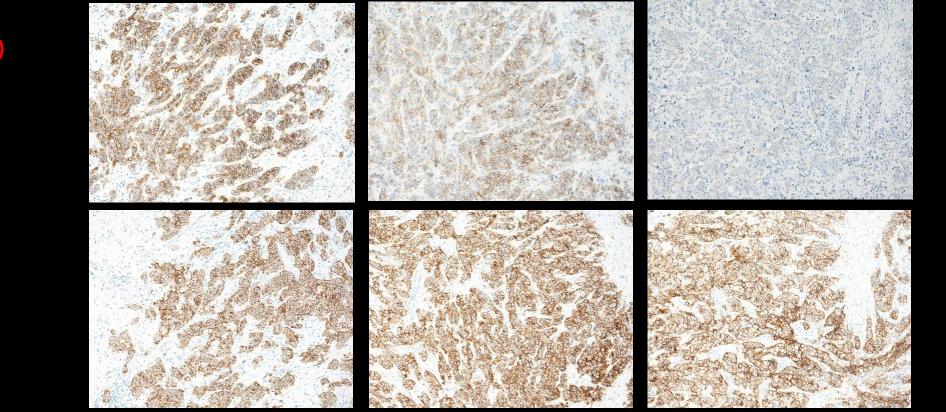
Proteolytic enzyme & digestion temperature ?

TCR β clone 8A3 (1:200 RR) / Flex+ (Tonsil NBF 48h)



Increased intensity of TCR β positive T-cells

Enzymatic digestion (Influence of fixation time)



EPCAM, clone MOC-31, dilution 1:20

NBF 24 h

NBF 48 h

NBF 120h

Adenocarcinoma (Breast) fixed in 10% Formalin

Pepsin / (Dako, S3002)

10 min/37°C

HIER, Low pH (S1700)

20 min / 97°C

The purpose of antigen retrieval is to unmask antigen epitopes /restore antigenic determinants and recover immuno-reactivity

Antigen retrieval procedures for formalin fixed tissue:

- □ <u>Heat</u> Induced Epitope <u>Retrieval</u> (HIER)
- **Tissue digestion using proteolytic enzymes**
- **Combined pre-treatment (HIER with proteolytic digestion)**

Only few markers require combined pre-treatment for "optimal performance".

CK-PAN

RTU (Ventana)

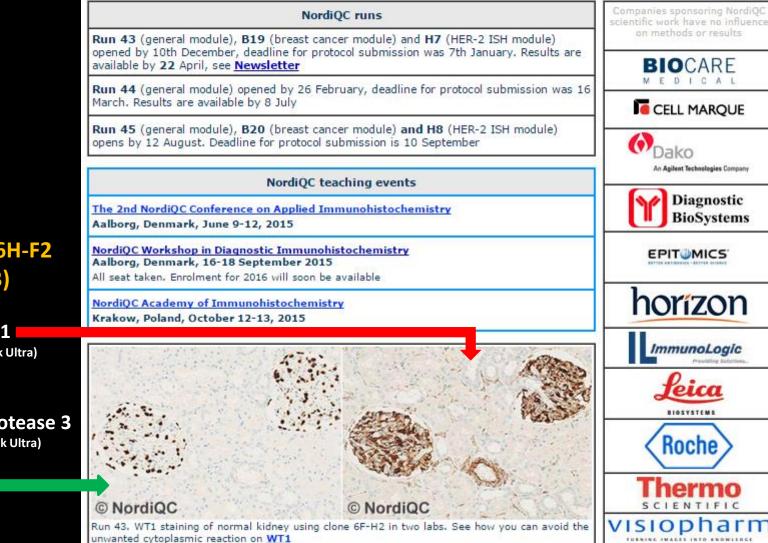
Table 4. Pass rates for antibody cocktails combined with epitope retrieval methods in the last three NordiQC runs

Pass rate for compiled data from run 54, 58 and 71											
	То	tal	HI	ER	Prote	olysis	HIER + proteolysi				
	Protocols	Sufficient	Protocols	Sufficient	Protocols	Sufficient	Protocols	Sufficient			
mAb AE1/AE3	609	449 (74%)	579	442 (76%)	12	1 (8%)	9	1 (11%)			
mAb AE1/AE3/5D3	21	12 (57%)	21	12 (57%)	-	-	-	-			
mAb AE1/AE3/PCK26	323	208 (64%)	39	22 (56%)	28	3 (11%)	254	181 (71%)			
mAb MNF116	27	2 (7%)	18	0 (0%)	7	2 (29%)	1	0			
mAb BS5	38	36 (95%)	38	36 (95%)	-	-	-	-			

Applying a combined pre-treatment protocol require a gentle/careful calibration of all parameters involved in the process (both HIER and proteolytic digestion steps)



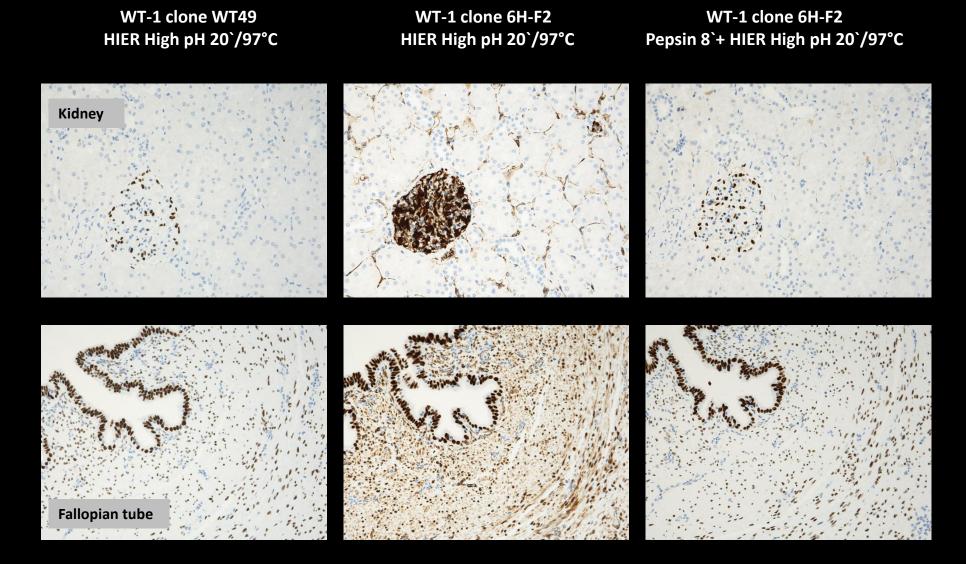
Home = Participation = Modules/tests = Assessments = Epitopes = Protocols = Techniques = Links Organization = Subscription = Newsletter = Accompanying letters = Seminars



WT1 clone 6H-F2 (RUN43)

HIER in CC1 (Ventana, Benchmark Ultra)

HIER in CC1 + Protease 3 (Ventana, Benchmark Ultra)

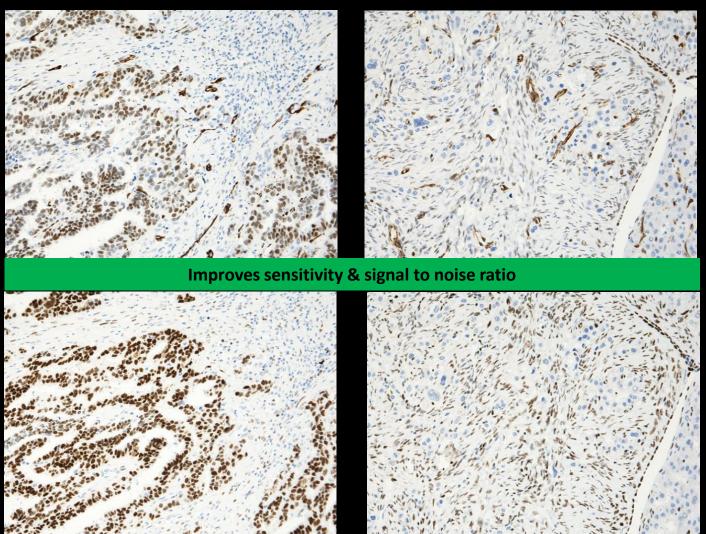


Proteolysis (Pepsin solution, RTU/Zytovision cat. no. ES-0001-50) followed by HIER

Combined pre-treatment (Enzymatic digestion followed by HIER)

Ovarian Serous Carcinoma

Ovarium/Peritoneum Metastasis (Hepatocellular carc.)



WT-1 clone 6H-F2 HIER High pH 20`/97°C

WT-1 clone 6H-F2 Pepsin 8`+ HIER High pH 20`/97°C

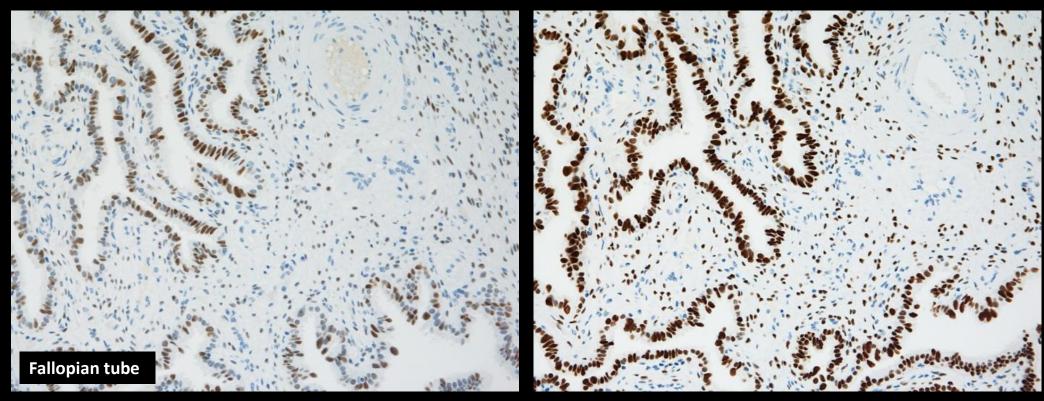
Proteolysis (Pepsin solution, RTU/Zytovision cat. no. ES-0001-50) followed by HIER

Combined pre-treatment (HIER followed by Enzymatic digestion)

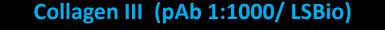
WT1, EP122 1:25 (Omnis)

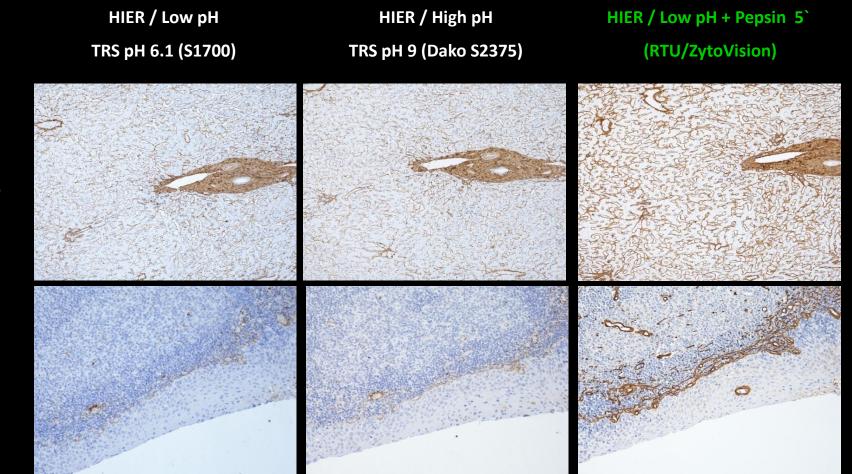
HIER TRS pH9 (24`/97°C)

HIER TRS pH9 (24`/97°C) + Pep © (3`)



Extracellulare matrix proteins: Combined pre-treatment (HIER followed by Enzymatic digestion)



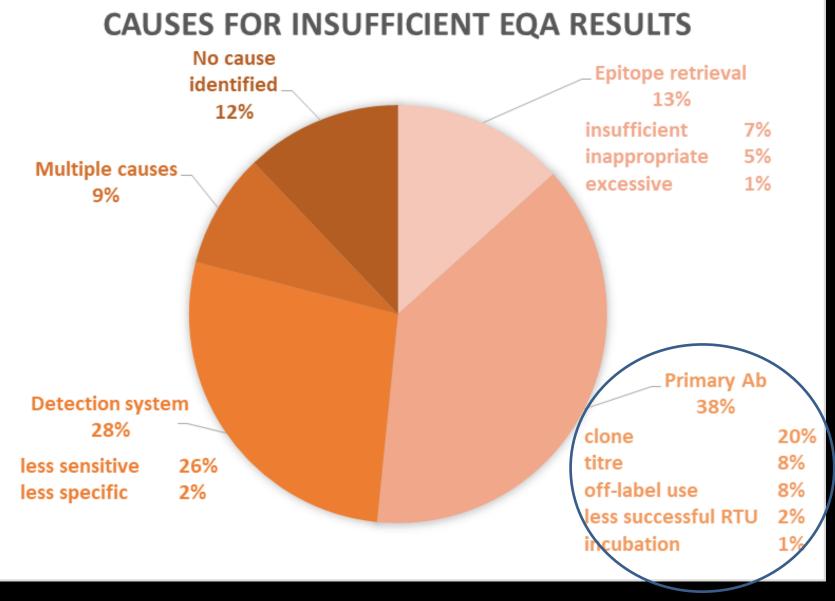


Hepar

Tonsil

Pause

NordiQC External Quality Assurance program



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79 % of insufficient results are related to the choice/use of:

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- Primary Ab (including stainer platform)
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False positive and/or negative results

Courtesy of Birgit Truumees et al (NordiQC): Poster presented at the annual USCAP meeting 2024

Antibody company (producer) - Quote:

In the world of next generation immuno-oncology research, having confidence in your immunoassay results is vital.

Unfortunately, 75% of antibodies in today's market are non-specific or simply do not work at all.

Volume 57(1): 7–8, 2009 Journal of Histochemistry & Cytochemistry http://www.jhc.org

PERSPECTIVE

Commercial Antibodies: The Good, Bad, and Really Ugly

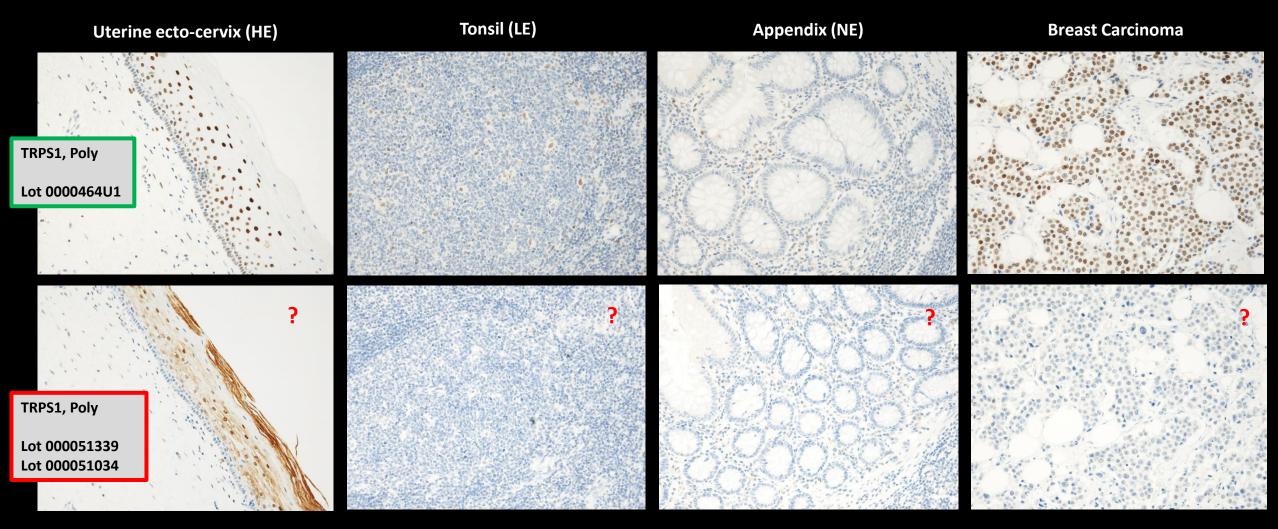
John R. Couchman

Biomedicine Institute, University of Copenhagen, Biocenter, Copenhagen, Denmark

SUMMARY The range of antibodies available commercially grows ever larger. Perhaps as a consequence, quality control is not always what it could and should be. Investigators must be aware of potential pitfalls and take steps to assure themselves that the specificity of each antibody is as advertised. Additionally, companies should provide the necessary information about the antigen and antibody to investigators, including references, so that the appropriate controls can be included. (J Histochem Cytochem 57:7–8, 2009)

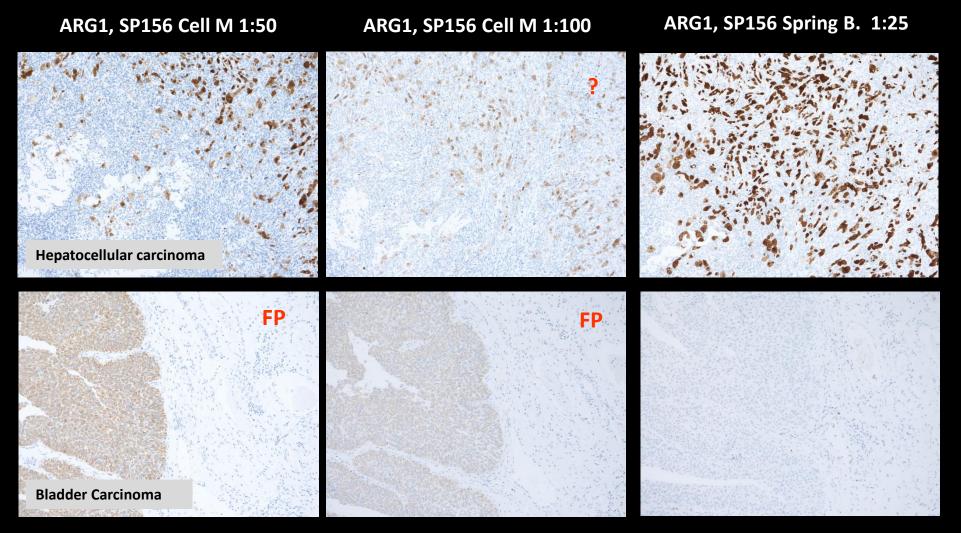
Primary Antibodies

Antibody choice: TRPS1, Poly (Invitrogen) 1:400 - Poor lots (same vendor)



HIER High pH 24`; Flex+ Rabbit linker

Antibody choice: Specificity (different vendors)

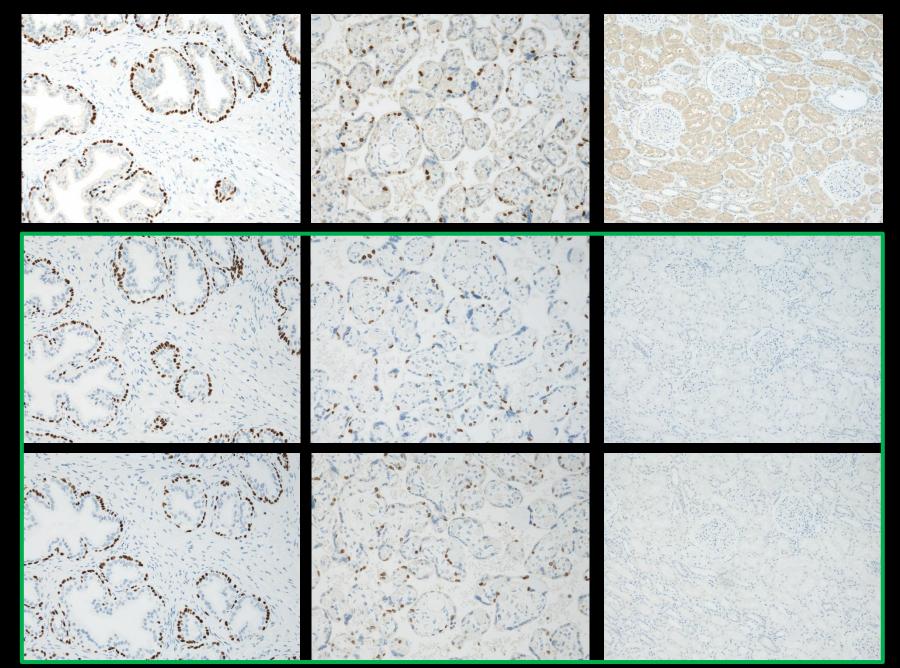


Dako Envision Flex+ - Poor Polymer-HRP batch(es) on the Omnis

P40 - Standard Envision Flex+ system (Poor HRP lot)

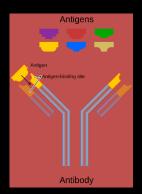
P40 - Standard Envision Flex+ system (Poor HRP reagent substituted with Envision K4003)

P40 - Quanto detection system (Thermo) (Substituting all Dako`s detection reagents)



Parameters affecting antibody-antigen reactions in tissue

Antibody choice – Sensitivity/Specificity Antibody Titer Antibody performance related to the chosen automated platform Antibody diluents



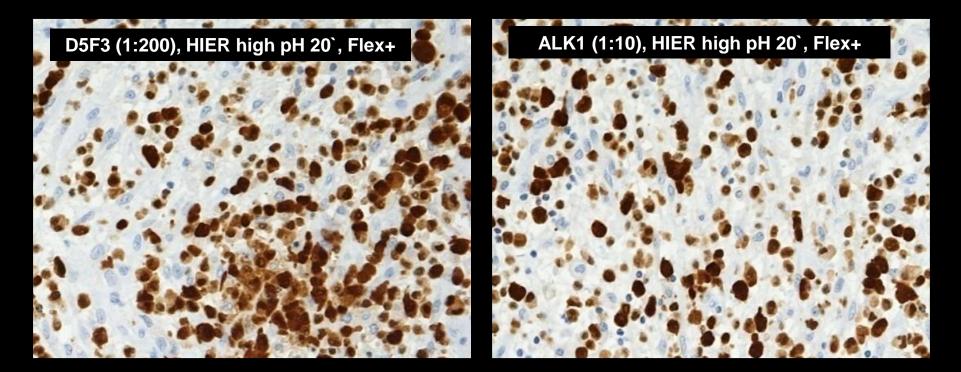
Incubation time Incubation temperature Sensitive to endogenous peroxidase blocking e.g., BCL6 (PG-B6p) and CD4 (1F6)

Storage of concentrated primary antibodies Storage of diluted primary antibodies

Problem: Primary antibody provides low sensitivity

Anaplastic lymphoma kinase (ALK)

Anaplastic large cell lymphoma (ALCL) (ALK-NPM rearrangement)



Anything wrong ?



Concentrated antibodies	n	Vendor	Optimal	Good	Borderline	Poor	Suff.1	OR ²
mAb clone 5A4	26 2 1 2 2 2 1	Leica Biosystems Monosan Abcam DBS Biocare Medical Zytomed Systems Invitrogen	8	9	14	4	49%	23%
mAb clone OTI1A4*	19 1 1 1	Origene Nordic Biosite Cell Signaling Zeta Corporation	16	6	0	0	100%	73%
mAb clone IHC509	1	GenomeMe	0	0	1	0	-	-
rmAb clone D5F3	19	Cell Signaling	7	9	3	0	84%	36%
rmAb clone ALK1	3 1	Dako/Agilent Cell Marque	0	0	0	4	-	-
rmAb clone QR017	1	Quartett	0	1	0	0	-	
rmAb clone SP8	1	BioGenex 0 0 0 1		-	-			
rmAb clone ZR305	1	Zeta Corporation	0	0	1	0	-	-
Ready-To-Use antibodies								
mAb clone 5A4 PA0306**/PA0831 (VRPS) ³	2	Leica Biosystems	1	1	0	0	-	-
mAb clone 5A4 PA0306*/PA0831 (LMPS) ⁴	10	Leica Biosystems	4	3	2	1	70%	40%
mAb clone 5A4 API3041	1	BioCare	0	0	1	0	-	-
mAb clone 5A4 CAM-0170	1	Celnovte	0	1	0	0	-	-
mAb clone 5A4 MAD-001720QD	1	Master Diagnostica	0	0	1	0	-	-
mAb clone ALK1 GA641	3	Dako/Agilent	0	0	0	3	-	-
mAb clone ALK1 IR641	4	Dako/Agilent	0	0	0	4	-	-
mAb clone ALK1 790/800-2918 (LMPS)⁴	10	Ventana/Roche	1	0	1	8	10%	10%
mAb clone 137E9E8 PA132	1	Abcarta	0	0	0	1	-	-
mAb clone OTI1A4 / 1A4 8344-C010	1	Sakura Finetek	1	0	0	0	-	-
mAb clone OTI1A4 / 1A4 GA785 (VRPS) ³	12	Dako/Agilent	12	0	0	0	100%	100%
mAb clone OTI1A4 / 1A4 GA785 (LMPS)⁴	4	Dako/Agilent	4	0	0	0	-	-
rmAb clone D5F3 790-4794 (VRPS) ³	73	Ventana/Roche	62	7	1	3	95%	85%
rmAb clone D5F3 790-4794 (LMPS)⁴	48	Ventana/Roche	36	9	3	0	94%	75%
rmAb clone SP8 RMPD007	1	Diagnostic BioSystems	0	0	0	1	-	-
Total	256		152	46	28	30		
			59%	18%	11%	12%	77%	

Proportion of sufficient stains (optimal or good) (\geq 5 assessed protocols).

Proportion of Optimal Results (>5 assessed protocols).

) Vendor Recommended Protocol Settings (VRPS) to a specific RTU product applied on the vendor recommended platform(s) (>5 seesed protocols)

) Laboratory Modified Protocol Settings (LMPS) to a specific RTU product (≥5 assessed protocols)

*) OTI1A4 is called 1A4 by some vendors

**) Product no. PA0306 has been terminated and replaced by PA0831

NQC Run 45, 51, 57 and 65 (ALK Lung)

87 protocols were based on clone ALK1:

Only two protocols (2%) were assessed as sufficient

NQC Run 65 (assessment spring/summer 2022)

1/21 protocols were assessed as sufficient

Don't use clone ALK1 to detect ALK rearranged lung adenocarcinomas - provides too low sensitive

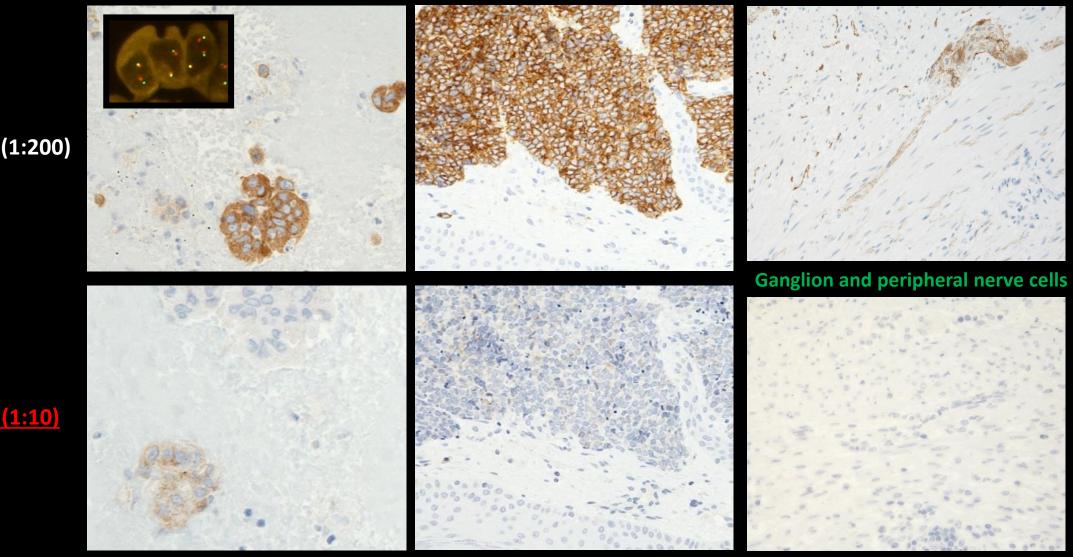
The clone ALK1 does not "fit-for-purpose"

D5F3, OTI1A4, 5A4

Lung AC (ALK-EML4)

MCC (Skin)

Appendix



ALK, D5F3 (1:200)

ALK, ALK1 (1:10)

HIER in high pH buffer 20`, Flex+

Clone ALK1 provides low sensitivity

Problem: Primary antibody provides low sensitivity

URO II/III

Concentrated antibodies	Reactivity	n	Vendor	Optimal	Good	Borderline	Poor	Suff.1	OR ³
mAb BC21	URO II	24 3	BioCare Medical Zytomed Systems	13	10	3	1	85%	37%
mAb AU-1	URO III	2	Cell Marque	-	-	-	2	-	-
rmAb SP73	URO III	7	Cell Marque Immunologic	-	-	1	7	0%	0%
rmAb ERP18799	URO II	1	ABCAM	-	1	-	-	-	-
rmAb EP321	URO III	1	Bio SB	-	-	1	-	-	-
rpAb AB82173	URO III	2	ABCAM	-	-	-	2	-	-
Ready-To-Use antibodies					_			Suff.1	OR.
mAb BC21 AVI 3051 KG		1	Biocare Medical	-	1	-	-	-	-
mAb BC21 API 3051 AA	URO II	1	Biocare Medical	-	1	-	-	-	-
mAb BC21 MAD-000773QD	URU II	1	Vitro SA	1	-	-	-	-	-
mAb BC21 MSG102		1	Zytomed Systems	-	1	-	-	-	-
mĀb BC21+BC17 API 3094 AA	URO II/III	3	Biocare Medical	-	2	1	-	-	-
mAb SP73 760-4533 (VRPS) ³		1	Roche/Ventana	-	-	1	-	-	-
rmAb SP73 760-4533 (LMPS)⁴	URO III	16	Roche/Ventana	-	-	2	14	0%	0%
rmAb SP73 345R-17/18		1	Cell Marque	-	-	-	1	-	-
Total		66		14	16	9	27		
Proportion				21%	24%	14%	41%	45%	

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

Proportion of Optimal Results (≥5 assessed protocols).

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

Laboratory Modified Protocol Settings (LMPS) to a specific RTU product (≥5 assessed protocols).

Urothelial carcinomas - The problem:

Abs raised exclusively against Uroplakin III show too low analytical sensitivity detecting urothelial cancers among carcinomas of unknown origin. RESEARCH ARTICLE

OPEN

Comparison of Antibodies to Detect Uroplakin in Urothelial Carcinomas

Heidi L. Kristoffersen, BMS, Rasmus Røge, MD, PhD, and Søren Nielsen, BMS

(Appl Immunohistochem Mol Morphol 2022;30:326-332)

	mmAb BC21 Uroplakin II			rm/	rmAb SP73 Uroplakin III			mmAb AU-1 Uroplakin III		
	n	n%	Mean H-score	n	n%	Mean H-score	n	n%	Mean H-scor	
Urothelial carcinomas (n = 58)	_			_			_			
Positive total	40	69	72	17	29	19	11	19	4	
High-expressor (H-score 150-300)	8	13	182	0	0	0	0	0	0	
Medium-expressor (H-score 10-149)	23	38	60	8	13	36	2	3	10	
Low-expressor (H-score 1-9)	9	15	6	9	15	3	9	15	3	
Negative (H-score <1)	18	31	0	41	71	0	47	81	0	
Nonurothelial carcinomas (n = 111)	_									
Positive total	3	3	18	0	0	0	0	0	0	
High-expressor (H-score 150-300)	0	0	0	0	0	0	0	0	0	
Medium-expressor (H-score 10-149)	2	2	27	0	0	0	0	0	0	
Low-expressor (H-score 1-9)	1	1	2	0	0	0	0	0	0	
Negative (H-score <1)	108	97	0	111	100	0	111	100	0	

mmAb indicates mouse monoclonal antibody; rmAb, rabbit monoclonal antibody.

Demonstrated that UPII klon BC21 outperforms UPIII Abs and at present should be the preferred choice of UP marker.

However, should be used in a panel with other Urothelial markers as e.g., GATA3 due to moderate analytical sensitivity (and due to positivity in a minority of nonurothelial carcinomas - app. 3%)

Appl Immunohistochem Mol Morphol • Volume 30, Number 5, May/June 2022

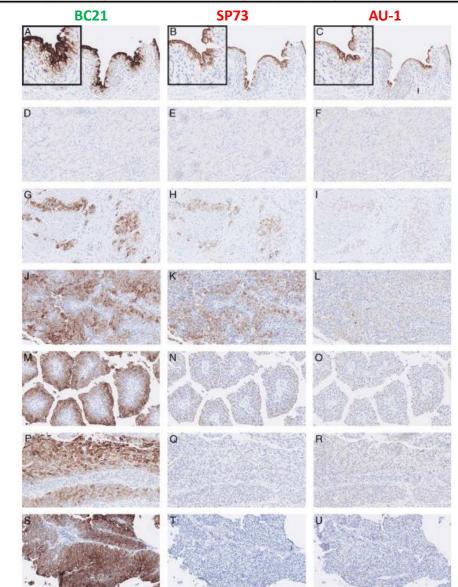


FIGURE 1. Examples of UP staining in normal tissue and UCs with various intensity using UPII mmAb clone BC21 (A, D, G, J, M, P, S), UPIII rmAb clone SP73 (B, E, H, K, N, Q, T), and UPIII mmAb clone AU-1 (C, F, I, L, O, R, U). A–C: normal urethra, insert high powerfield. D–F: normal kidney. G–I, J–L, M–O, P–R, and S–U: different UCs. mmAb indicates mouse monoclonal antibody; rmAb, rabbit mono-clonal antibody; UCs, urothelial carcinomas; UP, uroplakin.

Problem: Primary antibody provides low specificity and/or poor signal-to-noise ration

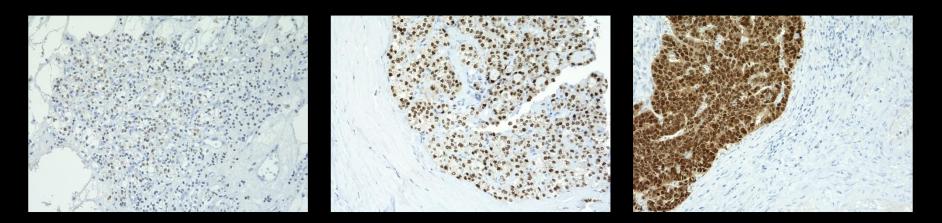
Which antibody ?

 Renal Cell Carcinoma (CC)
 Thyroid Carcinoma (Pa)
 Ovary Carcinoma (Se)

 Ovary Carcinoma (Se)
 Ovary Carcinoma (Se)

Pax-8, MRQ-50 (1:2000) HIER High pH/Flex+ (AS)

Pax-8, BC12 (1:150) HIER High pH/Flex+ (AS)



Problem: Primary antibody provides low specificity and/or poor signal-to-noise ration

Liau J-Y et al.: Appl Immunohistochem Mol Morphol . 2016 Jan;24(1):57-63

Demonstrated that neuroendocrine tumors (NET's) from a large variety of organs were immuno-reactive with the two less specific antibodies (pAb Proteintech & mAb MRQ-50) - cross-reacting with other PAX proteins

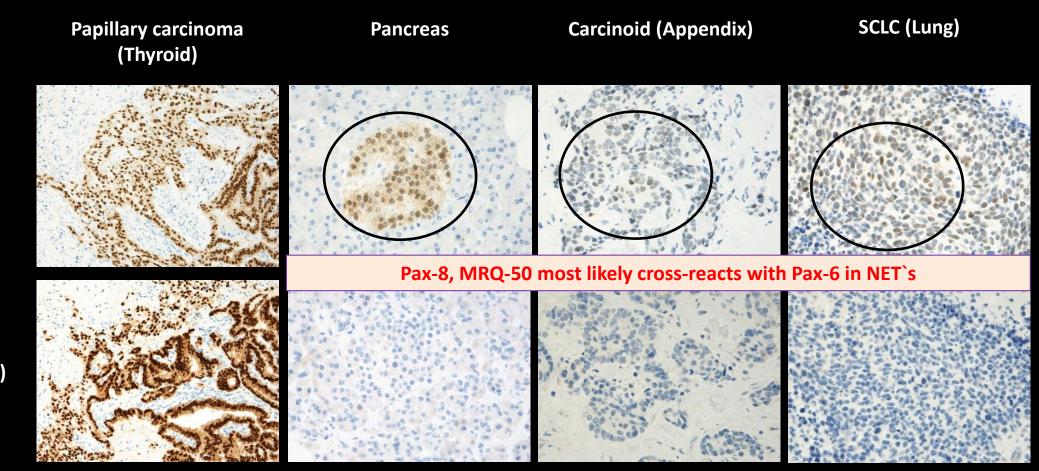
Also, all NET's were immuno-negative with the two monoclonal antibodies raised against the C-terminal part of PAX8 protein (PAX8R1 & BC12)

Moretti L et al. : Mod Pathol. 2012; 25 : 231-236

Demonstrated that an N-terminal PAX-8 polyclonal antibody cross-react with N-terminal region of PAX-5 and is responsible for reports of PAX-8 positivity in malignant lymphomas.

Also, PAX8 mRNA levels were not detected in any of the B-cell lymphoma cell lines studied. These results indicate that benign and malignant B-cells do not express PAX8.

Problem: Primary antibody provides low specificity and/or poor signal-to-noise ration



Pax8, MRQ-50 most likely raised against the N-terminal part of the PAX8 protein (cross-reacts with other PAX proteins)

Pax8, ZR1 or BC12 raised against the C-terminal part of the PAX8 protein (no cross-reacting with other PAX proteins)

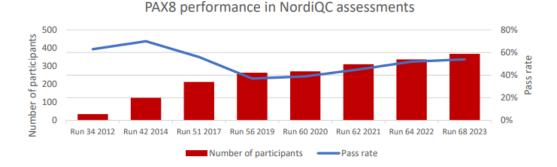
PAX-8, MRQ-50

PAX-8, ZR1 (BC12)

Problem: Primary antibody provides low specificity and/or poor signal-to-noise ration



Graph 1. Proportion of sufficient results for PAX8 in the eight NordiQC runs performed



Conclusion

Optimal staining results could be obtained with the rmAb clones **SP348. ZR-1. MXR013. GR002. OR016.** and **RM436.** Irrespective of the clone applied, efficient HIER, use of a sensitive 3-step polymer/multimer based detection system and careful calibration of the primary antibody were the most important prerequisites for an optimal staining result.

The rmAb clone **SP348** gave encouraging results and a high proportion of sufficient results on the main fully automated platforms and no cross-reaction with e.g. PAX5 was observed. No optimal results were however observed on the Bond Platform. The mAb clone **MRQ-50** provided a poor performance especially on the Ventana BenchMark and Dako Omnis platforms and at the same time also labelled PAX5 in B-cells. The **EP331** also provided a low pass rate due to aberrant nuclear staining reaction in non-PAX8 expressing cells and poor signal-to-noise ratio.

81% (101/125) of the protocols (Conc/RTU formats) showing crossreactivity with PAX5 was based on the mAb clone MRQ-50

- 20% (20/101) of the protocols provided sufficient result – none being optimal

PAX8 (Run 68, 2023)

Clone MRQ-50?

Cross-reactivity with PAX5 resulting in a distinct nuclear staining reaction of B-cells for antibodies raised against the N-terminal part of PAX8 was seen in 34% (125/368) of the returned slides (see Figs. 5a and 5b). This reaction applied for all polyclonal Abs and mAb clones MRQ-50, C12A32, IHC008, H5A8, PAX8R1 and rmAb 2774R. Within the last couple of years well-performing rmAbs without cross reactivity has been introduced to the market (see Table 1). Based on this, cross-reactivity with PAX5 was downgraded due to the risk of misinterpretation in the diagnostic work-up of CUP. The diagnostic challenges and different reaction profiles related to the choice of PAX8 Ab has e.g. been described by Kamaljeet Singh et al.; *AIMM 2020, Aug;28(7):558-561; Comparison of PAX8 Expression in Breast Carcinoma Using MRQ-50 and BC12 Monoclonal Antibodies* and Tacha D et al., *AIMM 2013, Jan;21(1):59-63; PAX8 mouse monoclonal antibody [BC12] recognizes a restricted epitope and is highly sensitive in renal cell and ovarian cancers but does not cross-react with b cells and tumors of pancreatic origin.*

FP/FN

FP/FN

Problem: Primary antibody provides low specificity and/or poor signal-to-noise ration

Concentrated antibodies	n	Vendor	Optimal	Good	Borderline	Poor	Suff.1	OR ²
mAb clone CL0276	5 2 1	Atlas Antibodies Sigma Aldrich Novus Biologicals	0	0	0	8	0%	0%
mAb clone CL0320	1	Atlas Antibodies	0	0	1	0	-	-
mAb clone SATBA4B10	3 2 2	Abcam Santa Cruz Zytomed Systems	0	0	2	5	0%	0%
mAb clone OTI5H7	1	ZSBio	1	0	0	0	-	-
rmAb clone EP281	30 12 1 1 1 1	Epitomics Cell Marque Immunologic BioSB Biocare Medical Unknown	22	14	4	6	78%	82%
rmAb clone SP281	4 1	Abcam Spring Bioscience	2	1	1	1	60%	40%
rmAb clone ZR167	1	Nordic Biosite	1	0	0	0	-	-
rmAb clone EPNCIR130A	5	Abcam	0	0	0	5	0%	0%
pAb HPA001042	5	Sigma Aldrich	0	0	2	3	0%	0%
pAb Ab69995	1	Abcam	0	0	0	1	-	-
Ready-To-Use antibodies							Suff. ¹	OR ²
rmAb clone EP281 384R-17/18	19	Cell Marque	7	10	1	1	89%	37%
rmAb clone EP281 PR/HAR239	2	PathnSitu	2	0	0	0	-	-
rmAb clone EP281 API3225	1	Biocare Medical	0	1	0	0	-	-
rmAb clone EP281 MAD-000747QD	1	Máster Diagnostica	0	0	1	0	-	-
rmAb clone EP281 BSB3199	2	BioSB	0	0	0	2	-	-
Total	105		35	26	12	32	-	
Proportion			33%	25%	11%	31%	58%	

Inferior clones (run 58): Display same reaction patterns in run 64

Table 1. Antibodies and assessment marks for SATB2, run 64									
Concentrated antibodies	n	Vendor	Optimal	Good	Borderline	Poor	Suff.1	OR ²	
mAb clone SATBA4B10	1 3 3	Abcam Santa Cruz Zytomed Systems	о	1	6	0	14%	0%	
rmAb clone EP281	13 66 1 3 1 5 3	Epitomics Cell Marque Diagnostic BioSystems BioSB Biocare Medical Gennova Scientific Zeta Corporation	48	28	8	8	83%	52%	
rmAb clone SP281	4 1	Abcam Zytomed Systems	о	4	0	1	80%	0%	
rmAb clone QR023	1	Quartett	1	0	0	0	-	-	
rmAb clone ZR167	1	Zeta Corporation	0	0	0	1	-	-	
pAb HPA001042	4	Sigma Aldrich	0	1	1	2	-	-	
pAb Ab69995	1	Abcam	0	0	0	1	-	-	

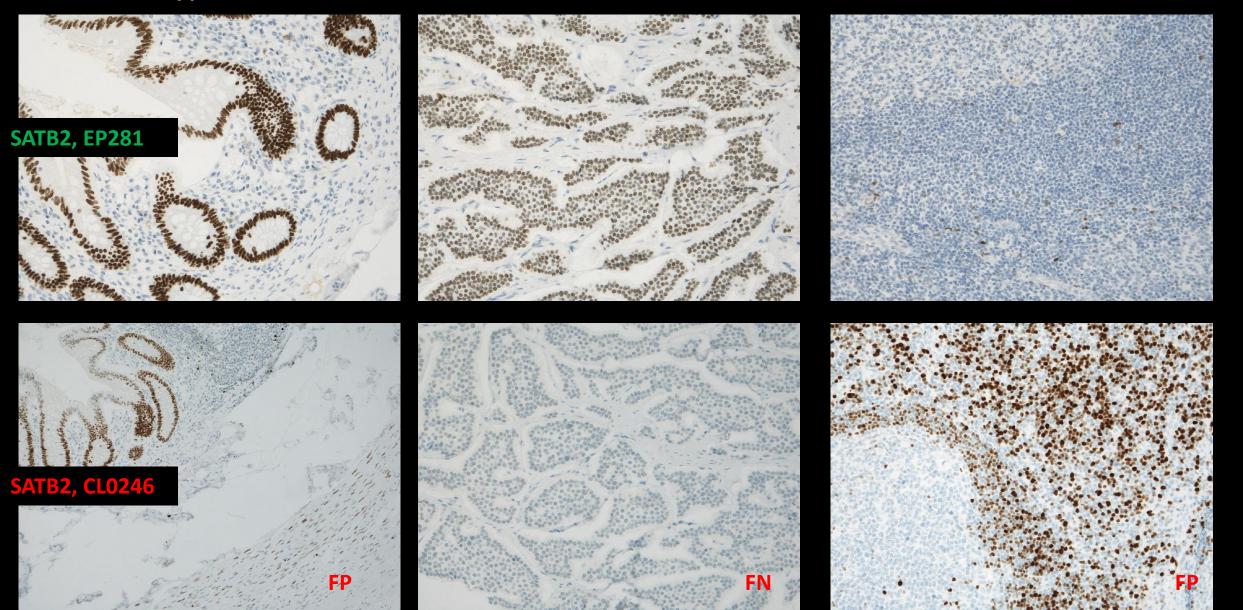
FP/FN + aberrant cytoplasmic staining FN + aberrant cytoplasmic staining

Impossible to calibrate correctly

Appendix

Neuroendocrine tumour (colon)

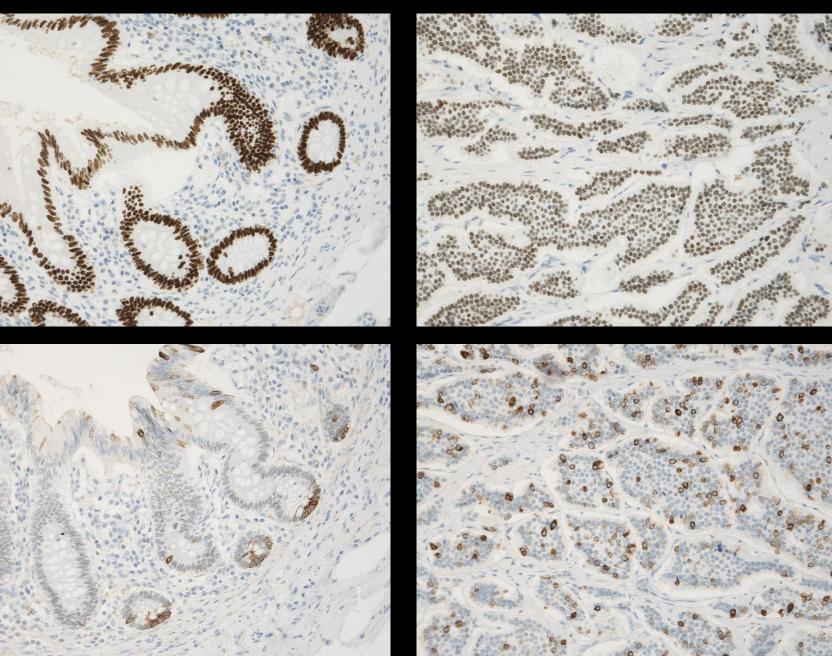
Tonsil



NordiQC results

Appendix

Neuroendocrine tumour (colon)



SATB2, EP281

States and

SATB2, pAb 69995

Micro anatomic localization

NordiQC results

Antibody choice: Sensitivity & Specificity

NordiQC results (2018-2024)

Abs providing low sensitivity	Abs providing low specificity and/or aberrant staining
SATB2 clone CL0276 & SATBA4310	SATB2 clone CL0276 & SATBA4310
SATB2 clone EPNCIR130A	PAX8 clone EP331 & MRQ-50
Uroplakin II+III clone AU-1 & SP73 & EP321 (all = Uro III)	MUM1 clone MRQ-43 & BC5
P16 clone G175-405	CK-HMW clone 34βE12
TTF1 clone 8G7G3/1	PR clone 1E12
ERG (Ets-Related-Gene) clone 9FY	ECAD clone EP700Y
ALK clone ALK1	PAX5 clone SP34
CK8/18 clone DC10 & C51 & CY90 (all = CK18)	SMAD4 clone RBT-SMAD4 & SP306
CK-PAN clone MNF116	MLH1 clone M1 & G168-728
CEA clone II-7	CD79a clone HM57
CGA clone DAK-A3 & 5H7	MSH6 clone 44
P63 clone 7JUL	Many pAbs (e.g., P40 and SOX10)
CD79a clone HM57 & 11E1	
ALK clone ALK1	
Many pAbs	

Problem: Poorly calibrated primary antibody (false negative or false positive)

The right primary antibody

The right protocol (AR procedure and detection system)

Poorly calibrated primary Ab ?

Tissue controls are important

Normal skin is the preferred positive control for GCDFP-15. The epithelial cells of the eccrine sweat glands must show an as strong as possible positive cytoplasmic staining reaction, while all other cells should be negative.

Normal breast tissue can also be used as control in which epithelial cells of the ductal glands must show an as strong as possible staining reaction.

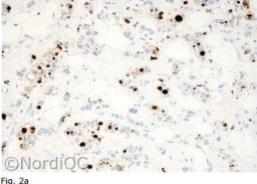
Background staining may be seen in vicinity of highly positive tissue structures (e.g. eccrine sweat glands)

Gross cystic disease fluid protein-15 (GCDFP-15)

Optimal staining for GCDFP-15 of the breast hyperplasia using the mAb clone 23A3 optimally calibrated as a concentrate, HIER in an alkaline buffer and a polymer based detection system. The majority of the ductal epithelia cells show a

compare with Fig. 2a - same protocol.

Insufficient staining for GCDFP-15 of the breast hyperplasia applying the mAb clone 23A3 as a concentrate using exactly the same protocol settings as used in Fig 1a, except for a 20 fold dilution of the primary antibody. The proportion and the intensity of the cells demonstrated are significantly reduced compared to the result in Fig. 1a. Also compare with Fig. 2b same protocol



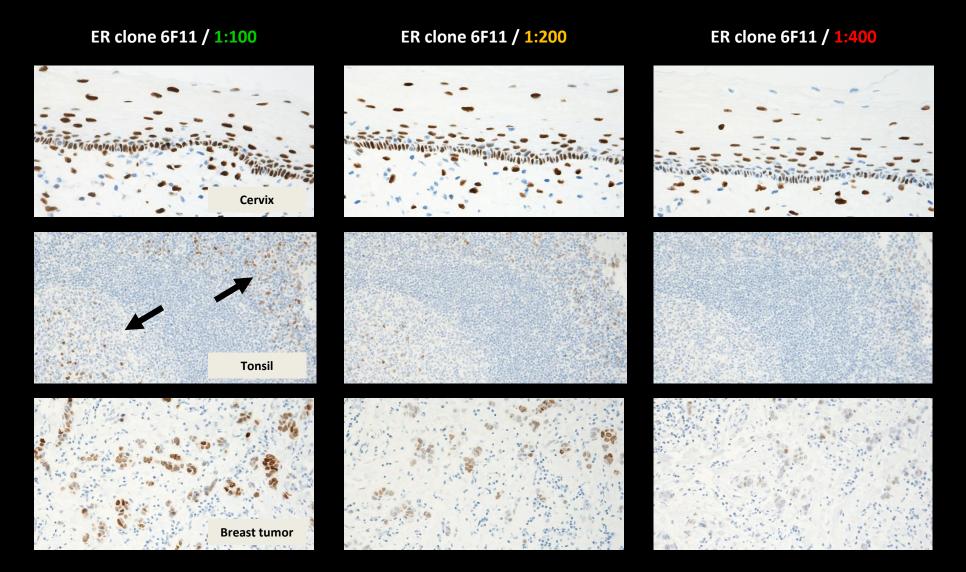
distinct moderate to strong cytoplasmic staining reaction. Also

Optimal staining for GCDFP-15 of the breast carcinoma no. 5 using same protocol as in Fig. 1a. The majority of the neoplastic cells show a moderate to strong dot-like cytoplasmic staining reaction

Insufficient staining GCDFP-15 of the breast carcinoma no. 5 using same protocol as in Fig. 1b. - same field as in Fig. 2a. Only scattered neoplastic cells show a faint dot-like reaction.

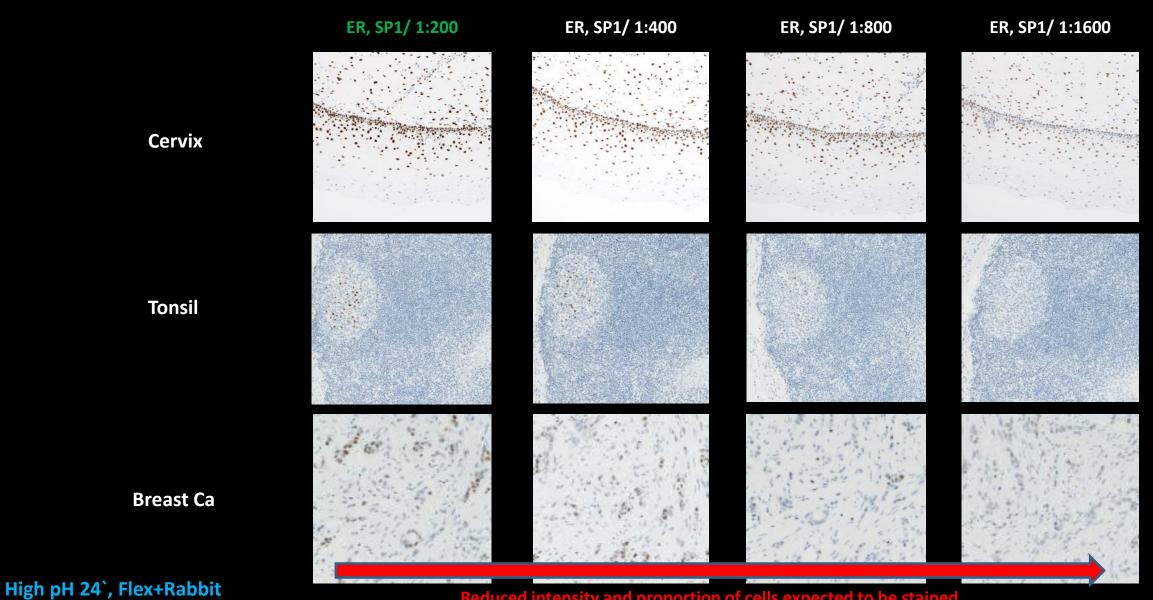


Antibody-Antigen reaction - Antibody Titer & Dilutions



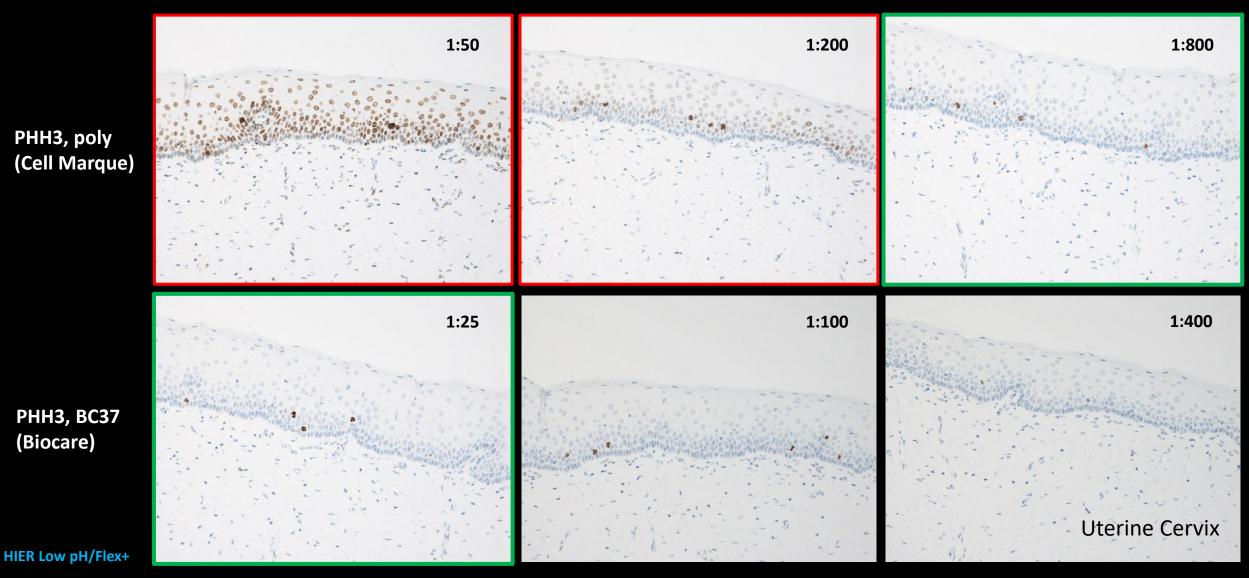
Staining indicators are extremely important - helping us to calibrate the IHC assay correctly

Problem: Primary antibody poorly calibrated (false negative)



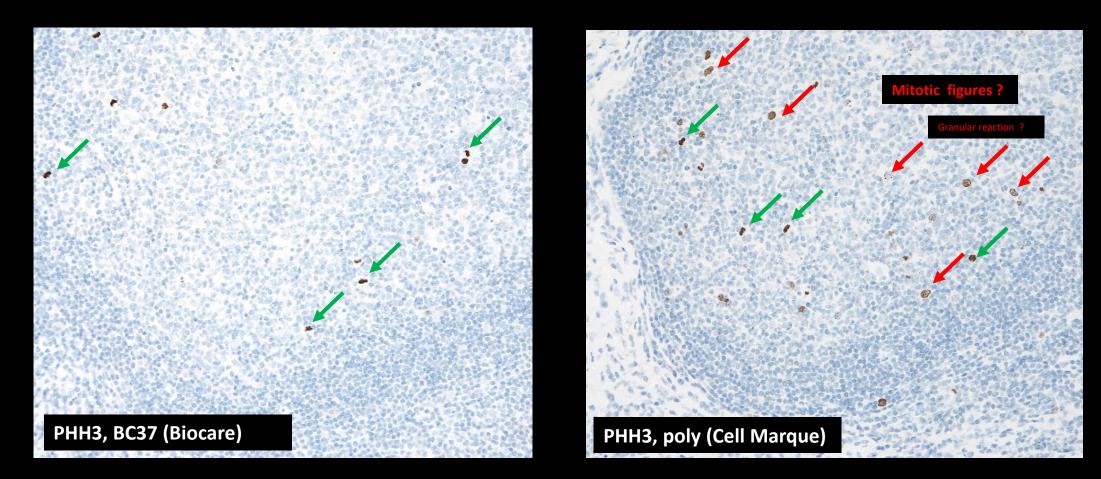
Reduced intensity and proportion of cells expected to be stained

Problem: Primary antibody poorly calibrated (false positive)



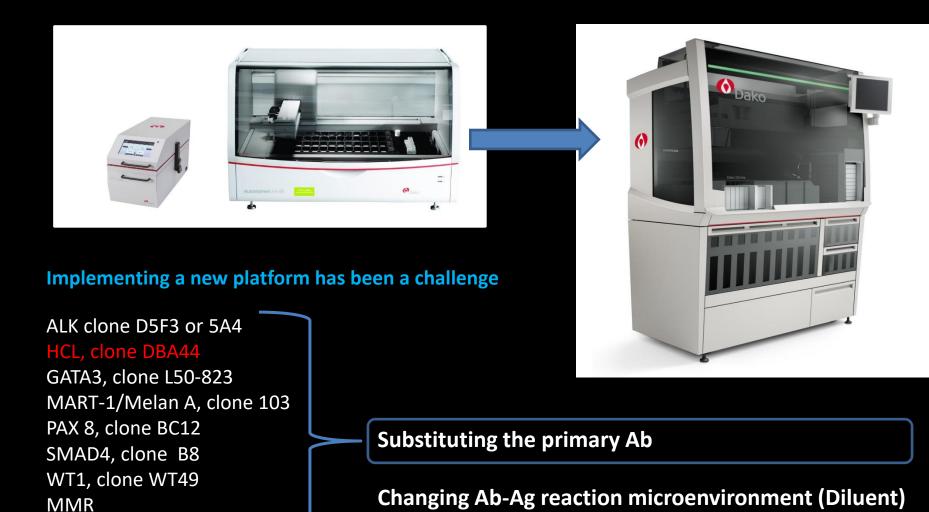
PHH3 should only be positive in cells in the late G2 and M phase (mitotic cells)

Problem: Primary antibody poorly calibrated (false positive)

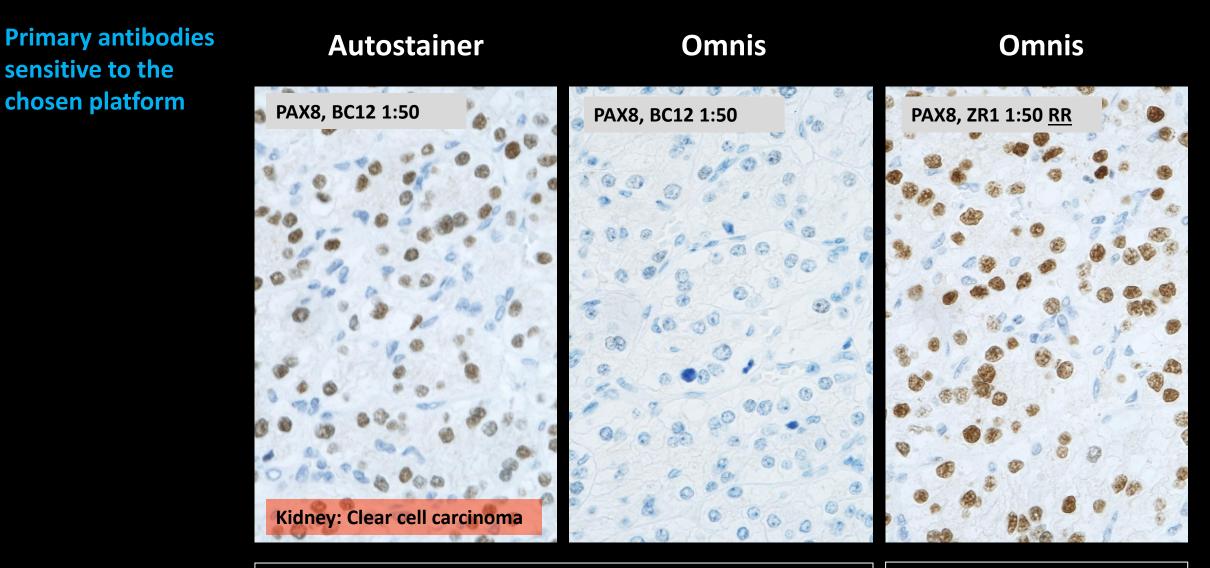


PHH3, poly might not be completely phospho-specific and might cross-reacts with cells that has not entered th late G2/M phase

Primary antibodies sensitive to the chosen platform



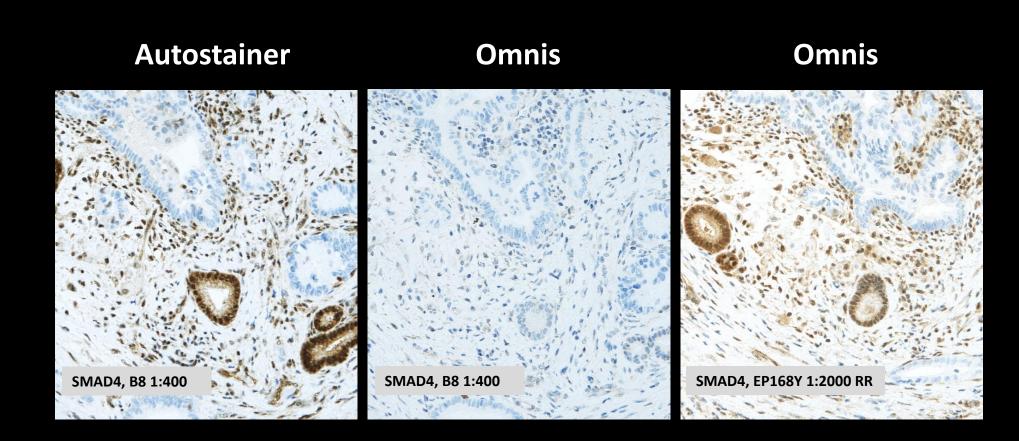
Low affinity primary antibodies



HIER High pH 20`, Flex+ (10+20)

HIER High pH 48`, Flex+ (10+20)

Primary antibodies sensitive to the chosen platform



HIER High pH 24`, Flex+ (10+20)

HIER High pH 20`, Flex+ (10+20)

Pancreatic Adenocarcinoma

Problem: Platform dependent antibodies (difficult markers)

NordiQC results (2018-2024)

Antibody	Clone	Platform(s)
CD4	4B12	BenchMark (Ventana), Omnis (Dako) and "BOND III (Leica)"
CD56	123C3	BenchMark (Ventana) and Omnis (Dako)
PMS2	A16-4	BenchMark
Melan A	A103	Benchmark and "Omnis"
P16 (RTU format`s Ventana)	E6H4	Omnis
Alpha Smooth Muscle Actin	1A4	Benchmark
SMAD4	B8	Benchmark and Omnis
CK8/18	5D3	Benchmark
EPCAM	BER-EP4	Benchmark and BOND (Leica)
BRAFmut	VE1	In general challenging on most platforms except for the Benchmark Ultra (Ventana)
CR	DAK-Calret1	Benchmark and Omnis
Desmin	D33	Omnis
Use of RTU-formats "Off-label"	E.g., Myosin, smooth muscle heavy chain	Often seen on the Omnis (RTU portfolio: IR vs GA products ?)

Problem: Does the automatic platform come with appropriate reagents fulfilling purpose and intended use of the assay e.g., <u>Antigen Retrieval Solutions</u>, Antibody diluents, Detection systems......



Assessment Run 56 2019

Epithelial cell-cell adhesion molecule (Ep-CAM)

Table 1. Antibodies and assessment marks for Ep-CAM, run 56									
Concentrated antibodies	n Vendor		Optimal	Good	Borderline	Poor	Suff. ¹	Suff. OPS ²	
mAb clone BS14	10	Nordic Biosite	9	1	0	0	100%	100%	
mAb clone Ber-Ep4	69 6 1	Dako Cell Marque Diagnostic Biosystems	14	13	21	28	36%	93%	
mAb clone MOC-31	23 5 1	Dako Cell Marque Diagnostic Biosystems	10	10	7	2	69%	71%	
mAb clone VU-1D9	5 3 1 1	 Merck Millipore Immunologic 		0	1	0	90%	100%	
rmAb clone EPR20532-225	1	Abcam	0	0	0	1	-	-	
Ready-To-Use antibodies									
mAb clone Ber-Ep4 760-4383	16	Ventana/Cell Marque	1	6	6	3	44%	100%	
mAb clone Ber-Ep4 248M-98	49	Cell Marque	5	13	16	15	37%	-	
mAb clone Ber-Ep4 IR/IS637	18	Dako	5	9	3	1	78%	87%	
mAb clone Ber-Ep4 IR/IS637 ³	6	Dako	1	2	2	1	-	-	
mAb clone Ber-Ep4 GA637	27	Dako	26	1	0	0	100%	100%	

OPS: Based on HIER in mod. Low pH buffers (Dako or Biocare)

Omitted from HIER in mod. Low pH buffers these buffers are not part of the reagent portfolio from the vendor

BS14 or VU-1D9 could be good alternatives compared to Ber-EP4 (and MOC-31) on platforms excluded from the use of modified low pH buffers as Diva Decloaker pH 6.2 (Biocare) or TRS pH 6.1 (Dako)

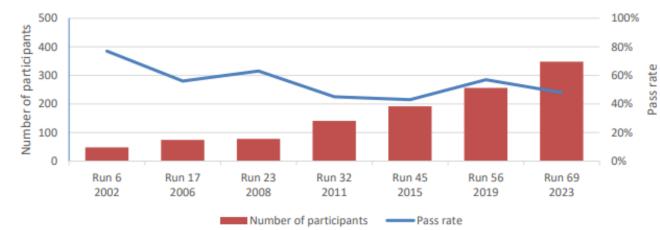


Assessment Run 69 2023 Epithelial cell-cell adhesion molecule (EpCAM)

Performance history

This was the seventh NordiQC assessment of EpCAM and, as shown in Graph 1, the pass rate decreased compared to the latest run 56, 2023.

Graph 1. Proportion of sufficient results for EpCAM in the seven NordiQC runs performed



EpCAM performance in NordiQC assessments

Latest assessment for EpCAM clone Ber-Ep4 - still a problem

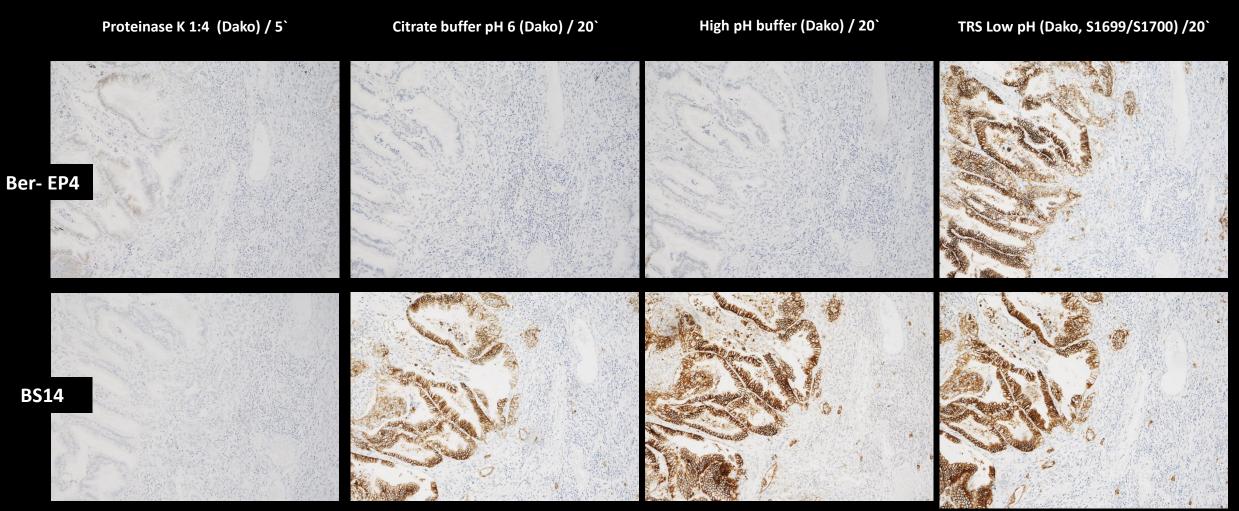
Substitute mAb clone Ber-Ep4 or apply modified Low pH buffer ?

Protocols mostly based on VRPS: 2-step detections system

Table 1. Antibodies and assessment marks for EpCAM, run 69

Concentrated antibodies	n	Vendor	Optimal	Good	Borderline	Poor	Suff.1	OR ²
mAb clone BS14	25	Nordic Biosite	19	6	0	0	100%	76%
mAb clone Ber-Ep4	40 8 1	Dako/Agilent Cell Marque Thermo Scientific	5	11	30	3	33%	10%
mAb clone MOC-31	16 1 2 1	Dako/Agilent Biocare Medical Cell Marque Leica Biosystems	1	14	4	1	75%	5%
mAb clone VU-1D9	9 1 4 1	Thermo Scientific Merck Millipore Diagnostic Biosystems Monosan	4	8	2	1	80%	27%
mAb clone C-10	2	Santa Cruz	0	0	2	0	-	-
mAb clone ZM131	1	Zeta Corporation	0	0	0	1	-	-
Ready-To-Use antibodies	Ready-To-Use antibodies							
mAb clone Ber-Ep4 760-4383 ³	13	Ventana/Roche	0	0	13	0	0%	0%
mAb clone Ber-Ep4 760-4383⁴	86	Ventana/Roche	1	20	60	5	24%	1%
mAb clone Ber-Ep4 248M-98	15	Cell Marque	0	1	13	1	7%	0%
mAb clone Ber-Ep4 IR/IS637 ³	1	Dako/Agilent	0	0	1	0	-	-
mAb clone Ber-Ep4 IR/IS637 ⁴	15	Dako/Agilent	3	2	10	0	33%	20%
mAb clone Ber-Ep4 GA637 ³	44	Dako/Agilent	12	30	2	0	95%	27%
mAb clone Ber-Ep4 GA637 ⁴	16	Dako/Agilent	3	8	5	0	69%	19%
PAD clone Ber-Ep4 PM107	3	Biocare Medical	0	0	1	2	-	-
mAb clone Ber-Ep4 MAD-001709QD	3	Master Diagnostica	0	0	2	1	-	-
mAb clone Ber-Ep4 PDM131	2	Diagnostic Biosystems	0	0	0	2	-	-
mAb clone Ber-Ep4 P-E002	1	Quartett	0	0	0	1	-	-
mAb clone Ber-Ep4 BMS048	1	Zytomed Systems	0	1	0	0	-	-
mAb clone Ber-Ep4 GM080402	2	Gene Tech	0	1	1	0	-	-

EPCAM clone Ber-EP4 vs BS14



EPCAM, BS14 (Nordic Biosite) is a better alternative compared to EPCAM Ber-EP4 (or MOC31) for automated platforms excluded from use of mod. low pH buffers.

Colon tumor

Problem: Platform dependent antibodies (difficult markers)

NordiQC results (2018-2024)

Antibody	Clone	Platform(s)
CD4	4B12	BenchMark (Ventana), Omnis (Dako) and "BOND III (Leica)"
CD56	123C3	BenchMark (Ventana) and Omnis (Dako)
PMS2	A16-4	BenchMark
Melan A	A103	Benchmark and "Omnis"
P16 (RTU format`s Ventana)	E6H4	Omnis
Alpha Smooth Muscle Actin	1A4	Benchmark
SMAD4	B8	Benchmark and Omnis
CK8/18	5D3	Benchmark
EPCAM	BER-EP4	Benchmark and BOND (Leica)
BRAFmut	VE1	In general challenging on most platforms except for the Benchmark Ultra (Ventana)
CR	DAK-Calret1	Benchmark and Omnis
Desmin	D33	Omnis
Use of RTU-formats "Off-label"	E.g., Myosin, smooth muscle heavy chain	Often seen on the Omnis (RTU portfolio: IR vs GA products ?)

Off-Label use: RTU product used on a platform to which the format has not been validated

Problem: Use of RTU-formats "Off-Label" exemplified by Myosin, smooth muscle heavy chain (SMH) NQC assessment Run 66

Autostainer (SMH, product IR066):

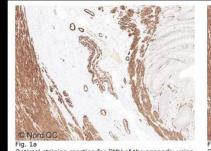
VRPS: Pass rate of 73% (18% optimal)

LMPS: Pass rate of 86% (57% optimal)

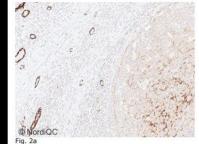
Omnis (SMH, Product IR066, Off-Label use)

Pass rate of 15% (no optimal results)

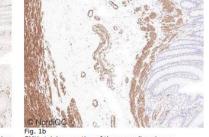
All the insufficient results were commented as weak, false negative or poor-signal-to-noise-ratio characterized by extensive background reaction combined with difficulties to demonstrate low-level antigen expressing structures. In this context, it was more successful to use the concentrated format of mAb clone SMMS-1 on Dako Omnis (see Table 2) as alternative to the off-label use of a RTU format not being developed and validated for the respective IHC platform.



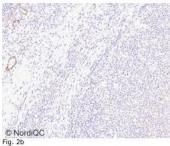
Optimal staining reaction for SMH of the appendix using the mAb clone SMMS-1 RTU (Ventana/Roche, 760-2704) within laboratory modified protocol settings, using OptiView as detection system /irtually all smooth muscle cells in vessels and lamina muscularis show a moderate to strong cytoplasmic staining reaction Also compare with Figs. 2a - 5a, same protocol.



Optimal SMH staining reaction of the tonsil using same protocol as in Fig. 1a. A weak to moderate staining reaction is seen in the follicular dendritic network in the germinal center. A high signal-to-noise ratio is observed.



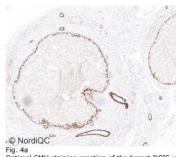
SMH staining reaction of the appendix using a insufficient protocol based on the mAb clone SMMS-RTU (Dako/Agilent, IR066) developed for the Autostainer, but applied on the Dako Omnis platform In appendix - same field as Fig. 1a, a weak to strong staining reaction is seen in virtually all smooth muscle cells as expected. Also compare with Figs. 2b - 5b, same protocol.



Insufficient SMH staining reaction of the tonsil using same protocol as in Fig. 1b - same field as Fig. 2a. The follicular dendritic network in the germinal center is virtually negative and only the smooth muscle cells of large vessels are demonstrated



Optimal SMH staining reaction of the breast hyperplasia using same protocol as in Figs. 1a and 2a. A moderate and distinct staining reaction is seen in virtually all myoepithelial cells lining the breast glands.

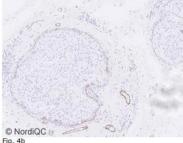


Optimal SMH staining reaction of the breast DCIS using same protocol as in Figs. 1a - 3a. A moderate, distinct and continuous staining reaction is

seen in the myoepithelial cells lining the breast DCIS component.



Insufficient SMH staining reaction of the breas hyperplasia using same protocol as in Figs. 1b and same field as Fig. 3a Only a faint staining reaction is seen in few myoepithelia cells lining the breast glands



Insufficient SMH staining reaction of the breast DCIS using same protocol as in Figs. 1b - 3b - same field as Fig. 4a.

Only a weak and disrupted staining reaction is seen in the myoepithelial cells lining the breast DCIS component.

Dako RTU IR-products providing poor results on the Omnis (2022-2024):

Desmin, D33/IR606 (Run 64, 2022) Calretinin, DAK-Calret1/IR627 (Run 64, 2022) CD56, 123C3/IR628 (Run 64, 2022) CD4, 4B12/IR649 (Run 67, 2023) CD5, 4C7/IR082 (Run 69, 2023)

Primary antibodies sensitive to the chosen platform



ALK clone D5F3 or 5A4 HCL, clone DBA44

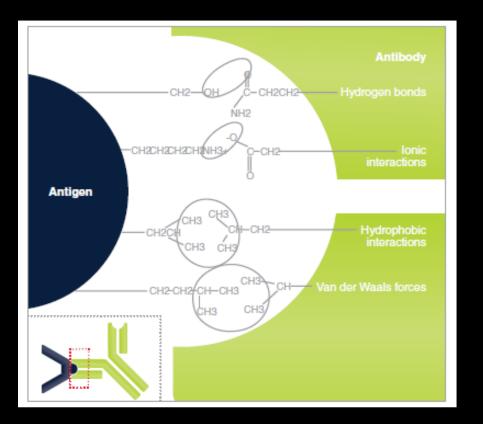
GATA3, clone L50-823 MART-1/Melan A, clone 103 PAX 8, clone BC12 SMAD4, clone B8 WT1, clone WT49 MMR ASMA, 1A4

- Changing the primary Ab

Changing Ab-Ag reaction microenvironment (Diluent)

Low affinity primary antibodies

Antibody Diluents: Antigen-Antibody reactions



The strength by which the primary Ab binding site binds to an antigenic epitope is called affinity

Antibodies are attracted initially through electrostatic interactions, and subsequently through weak forces

- Hydrogen bonds
- Hydrophobic interactions
- Van der Waals forces

Antibody diluent formulations can significantly alter stability and binding properties of antibodies affecting both epitope specificity and non-specific interactions

Applied Immunohistochemistry & Molecular Morphology 9(2): 176-179, 2001

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Formalin-Fixed and Heat-Retrieved Tissue Antigens: A Comparison of Their Immunoreactivity in Experimental Antibody Diluents

Thomas Boenisch, M.S.

Antibody diluents

Demonstrated that:

pH of the antibody diluent had a high impact of the final IHC result

Addition of NaCL (increasing the ionic strength) to the diluent negates most of the sensitivity gained through Antigen Retrieval (Table 3).

TABLE 3. Comparison of staining scores of 13
optimally diluted antibodies as a function of antigen
retrieval at pH 9.9, use of 0.05 M Tris (TB), pH 6.0 and
8.6, or Tris-buffered 0.15 M NaCl (TBS) of pH 6.0 and
8.6, and 0.02 M phosphate-buffered 0.15 M NaCl of pH
7.5 (PBS)

		Т	ТВ		TE	BS	PBS
Clone	pН	6.0	8.6		6.0	8.6	7.3
BLA.36 UCHL1 L26 PC10 N10/2 V9 TAL1B5 ER-PR-8 Ber-H2 4KB5 DF-T1 PD7/26 C3D-1		2 4 4 3 4 4 4 4 4 4 4 4 4 4	4 3 3 3 2 3 2 3 2 3 2 3 2 3 2 3 2 3 2 3		1 2 3 4 1 4 3 2 ND 4 2 ND ND	2 1 3 4 2 4 2 1 ND 2 0 ND ND	1 1 2 3 1 2 2 2 0 4 1 3 1
ND, not o	lone.			•			

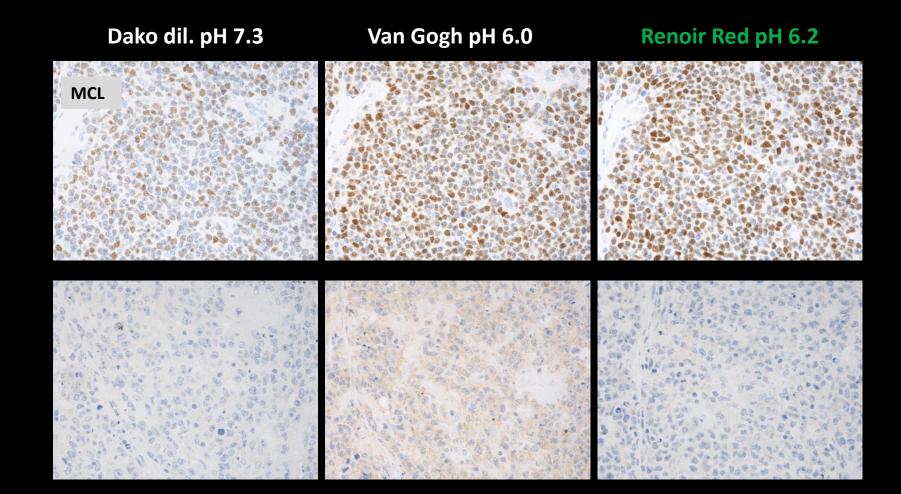
Antibody-Antigen reaction – Commercial Antibody Diluents

Biocare Medical's Antibody Diluent Offering

Antibody Diluent	Catalog Number	pН	Composition	Note
Da Vinci Green Diluent	PD900	7.3	Phosphate-based	May act as a "universal" diluent
Monet Blue Diluent	PD901	7.9	Tris-based	Uses Assure Technology with pH indicator. A change in diluent color indicates a change in pH, meaning optimal staining may be compromised.
Renaissance Background Reducing Diluent	PD905	7.3	Proprietary	Has potent background reducing agents; For antibodies with non-specific background
Renoir Red Diluent	PD904	6.0	Tris-based	Designed to work with Tris-based antibodies, rinsing buffers and/or detection systems that use alkaline phospha-tase
Van Gogh Yellow Diluent	PD902	6.0	Phosphate-based	Specifically formulated to enhance certain types of antibodies and may increase titers up to 2 to 4 times
Fluorescence Antibody Diluent	FAD901	7.3	Phosphate-based	Protease free and stabilizes fluorescent dyes for up to one month in a prediluted format
Revival Series Sampler	PD912H4	N/A	N/A	Includes 25 mls of each Da Vinci Green, Renoir Red, Van Gogh Yellow, Monet Blue

Standard Diluent pH 7.3 (Dako, K8006)

SOX11 clone MRQ-58 (1:200)

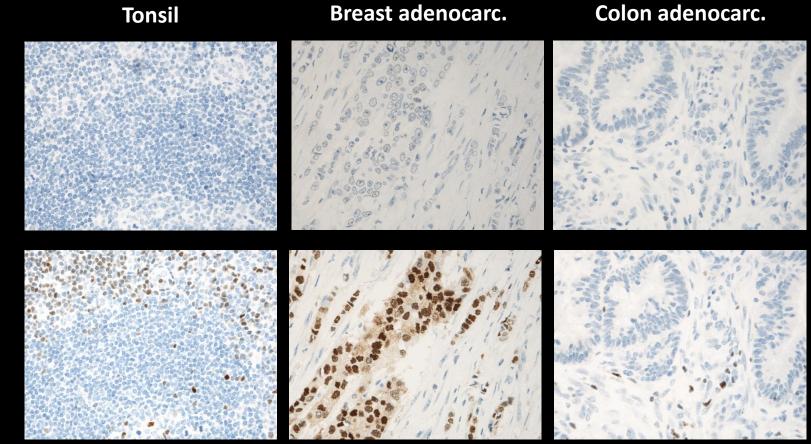


MCL

DLBCL, NOS

Antibody diluents

GATA3, L50-823 (Biocare) <u>1:800</u>



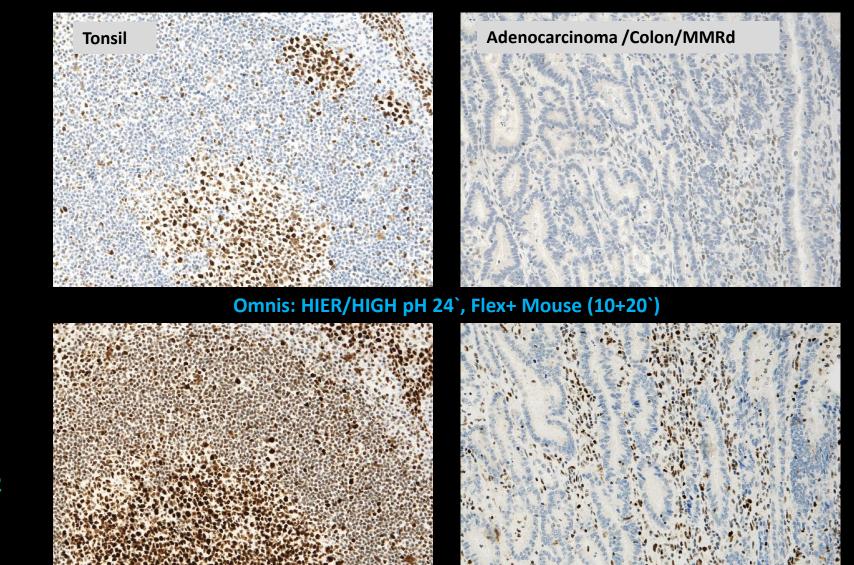
Dako Dil. pH 7.3

Biocare Renoir Red pH 6.2

Omnis: HIER/HIGH pH 24` at 97°C, Flex+ Mouse (10+20`)

Antibody diluents

MSH2, FE11 1:50 Dako dil. pH 7.3

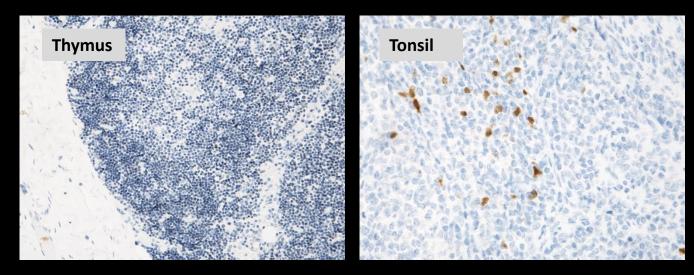


MSH2, FE11 1:50 Renoir Red pH 6.2

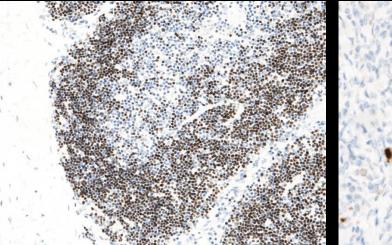
Antibody diluents

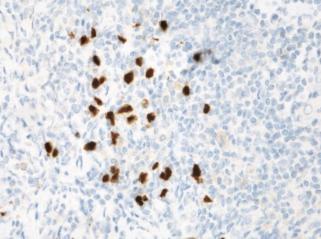
TdT, SEN28 1:50 Dako dil. pH 7.3

TdT, SEN28 1:50 Renoir Red pH 6.2



Omnis: HIER/HIGH pH 24`, Flex+ Mouse (10+20`)





Problem: Antibody diluent

CD10 clone 56C6 (1:50)/Dako Diluent pH 7.3



CD10 clone 56C6 (1:50)/Renoir Red pH 6.2



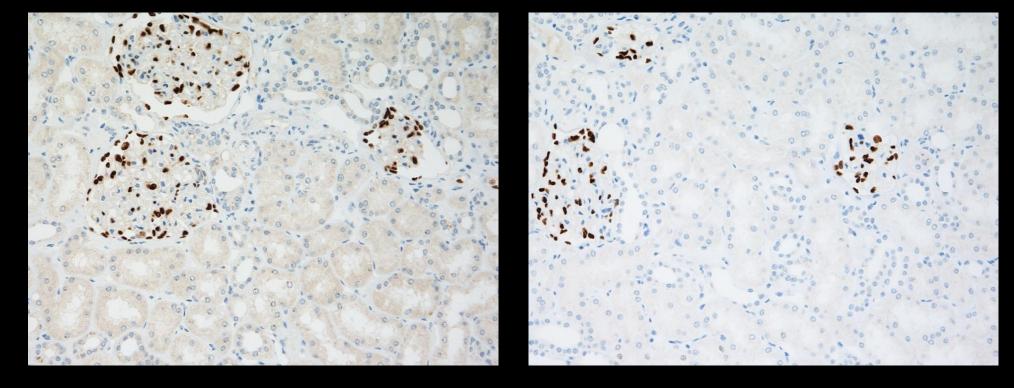
Renoir Red is not always the best antibody diluent Use a "antibody diluent test battery"

Omnis: HIER/HIGH pH 24`, Flex+ Mouse (10+20`)

Antibody diluents

WT1, EP122 1:25 Renoir Red (Biocare)

WT1, EP122 1:25 Background Sniper (Biocare)



The choice of antibody diluent can suppress unwanted/unspecific background staining

Kidney

HIER TRS pH9 (24` /97°C) + Pep © (3`)

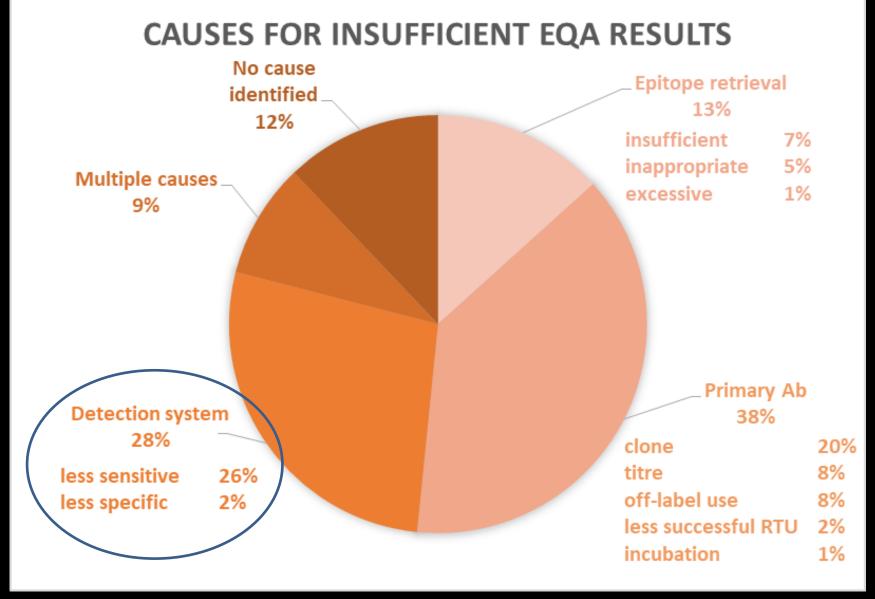
Antibody diluents (Experience from Dept. of surgical Pathology, Region Zealand, Denmark on the Omnis)

Antibody/Antigen reactions benefitting from diluting the primary Ab in Renoir Red pH 6.2 (improving signal): ALK (1A4), CR (CAL6), CD4 (EP204), CD5 (SP19), CMYC (EP121), GATA3 (L20-823), GPC3 (1G12), IMP3 (69.1), MLH1 (BC23, ES05 & GM011), MSH2 (G219-1129), MSH2 (FE11), MSH6 (EP49), NKX 3.1 (poly), SALL4 (6E3), PAX8 (ZR1), PMS2 (EP51), SOX10 (EP268), SOX11 (C1 & MRQ58), TdT (SEN28 & EP266), UP-II (BC21), WT1 (WT49) and

Antibody/Antigen reactions unaffected from diluting the primary Ab in Renoir Red pH 6.2 and compared to Dako diluent pH 7.3: BCL2 (124), BCL6 (LN22 & PG-B6p & GI191E/A3), CR (DAK-Calret1), <u>CD10 (56C6)</u>, CD163 (MRQ26), CD21 (2G9), CD5 (4C7), ER (SP1), <u>HHV8 (13B10)</u>, Mammaglobin (304-1A5), MUC5AC (CLH2), MUC6 (CLH5), and

Antibody/Antigen reactions benefitting from diluting the primary Ab in Background sniper (reduces background problems): Spirochete (poly), BORR (poly), WT1 (EP122) and

NordiQC External Quality Assurance program



App. 20-30% of all results are evaluated as insufficient

79 % of insufficient results are related to the choice/use of:

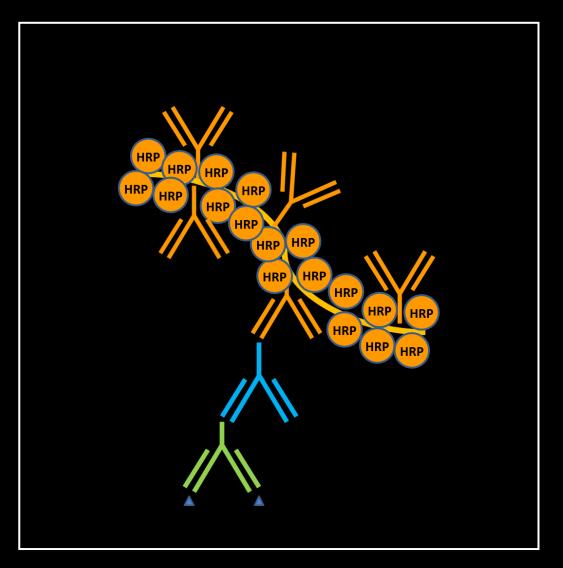
- Epitope retrieval procedure
- Primary Ab (including stainer platform)
- Detection system

False positive and/or negative results

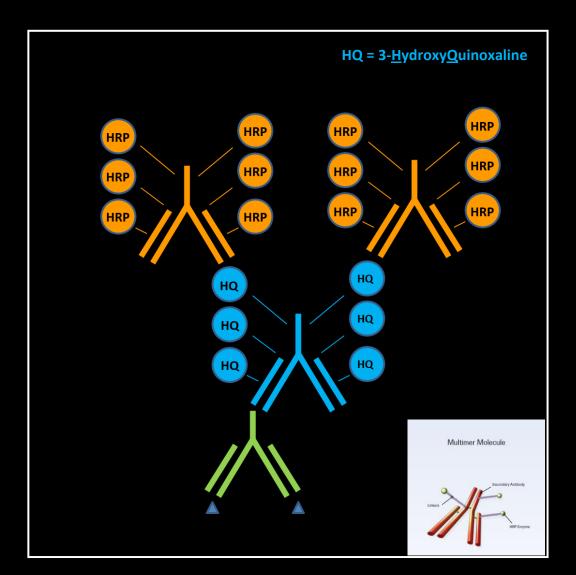
Considerations related to the choice of detection system:

- Sensitivity
- Specificity
- Enzyme conjugate (HRP or AP)
- Blocking of endogenous activity (PAX5 clone 1EW, CD4 clone 1F6, BCL6 clone PG-B6p)
- Turn around time (TAT)
- □ Automatic platform (open or closed system)
- **Price**

NordiQC: Virtually all laboratories use a polymer or multimer detections system

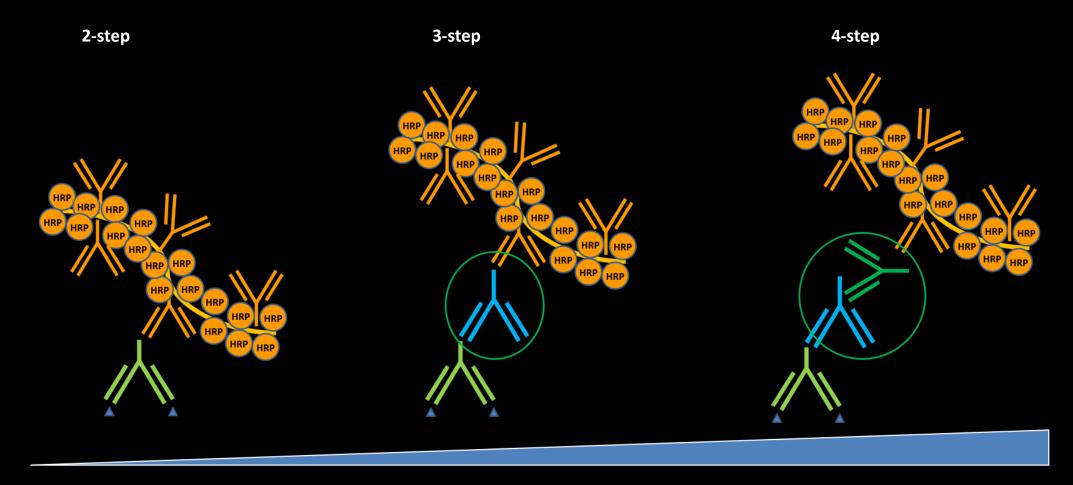






Multimer detection (Ultra View and OptiView)

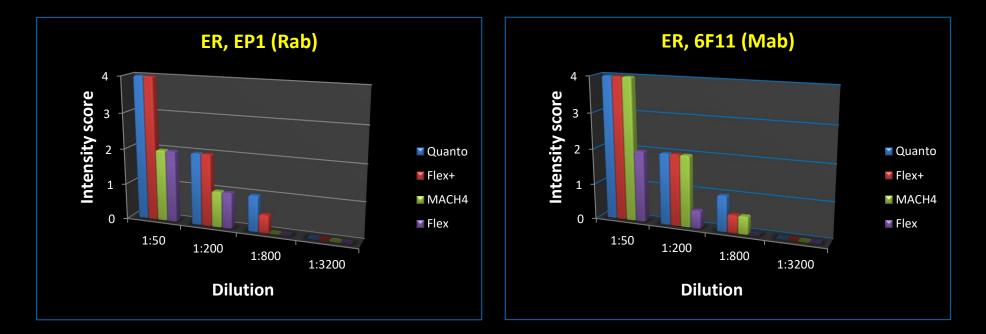
Detection systems and sensitivity



Increased sensitivity: Amplier A/B, HQ-Linker, Linker (Mouse/Rabbit), Enhancer, Universal Linker, Post Blocking

Know your detection system - strength and weakness

Detection systems - Performance Testing



<u>ER - Endpoint titration (some general remarks and important issues)</u>:

- **The 3-step polymer detection systems Quanto and Flex+** produced the overall highest intensity.
- **The 3-step polymer detection system MACH4 only enhances reactions with mouse monoclonal Abs (ER,6F11).**
- **Optimal staining**" was highly influenced by the concentration of the primary Abs and the nature of detection system.

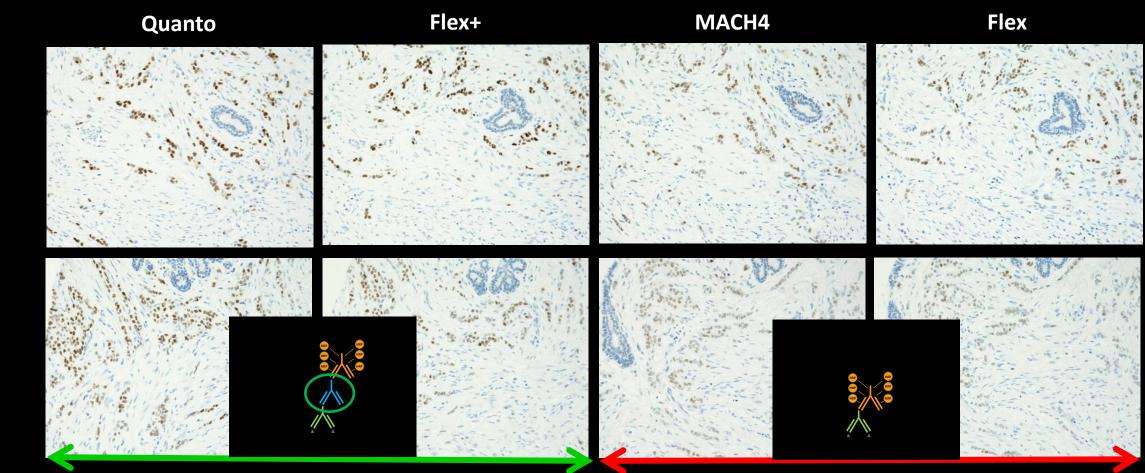
Performance testing of detection systems (Vendor recommended protocol settings)

Breast tumor

1:50

1:200

ER, EP1 (Rab)



High Intensity (3-step detection systems)

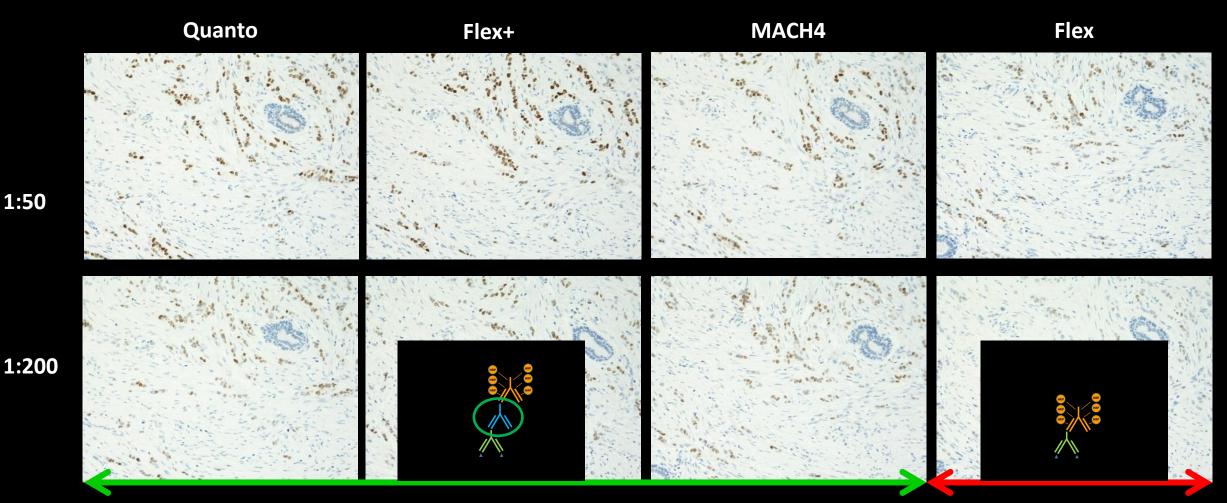
Low Intensity (2-step detection systems)

The technical test approach – Analytical phase

Performance testing of detection systems (Vendor recommended protocol settings)

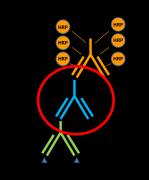
Breast tumor

ER, 6F11 (Mab)



High Intensity (3-step DS)

Low Intensity (2-step DS)



Polymer based detection systems

Without enhancement (second step)

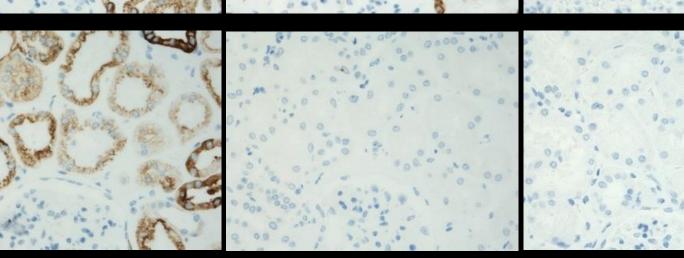
EnVision Flex (- Linker) MACH4 (- Mouse probe)

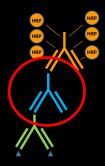
Quanto (- Amplifier)

CK8, EP1628Y (rabbit Ab)

 Kidney
 Image: Comparison of the second o

CK8, TS1 (mouse Ab)





Polymer based detection systems (Næstved LAB)

Expected reaction patterns

	Quanto LabVision TL-125-QHD	Quanto - Amplifier	Flex+ Dako K8012	Flex - linker	MACH4 Biocare M4U534	MACH4 - Mouse probe
Anti-X (Mab) Mouse primary Ab	+	-	+	(+)	+	-
Anti-X (Rab) Rabbit primary Ab	+	-	+	(+)	(+)	(+)

Detection system	Second step	HRP-Polymer	Enhances signal of
Quanto (LabVision)	Amplifier anti-Mab/Rab or Hapten conjugate ?	Quanto polymer (anti-unknown species or Hapten)	Both primary Mab or Rab
Flex+ (Dako)	Linker Mouse (Rab anti Mab) Linker Rabbit (Mab anti Rab)	Flex polymer anti-Mouse/Rabbit	Both primary Mab or Rab
MACH4 (Biocare)	Mouse probe anti-Mouse (Rab)	MACH4 polymer Anti-Rabbit	Only primary Mab
Flex*	-	Flex polymer anti-Mouse/Rabbit	-
* 2-step polymer system			

Know your detection system strength and weakness

Less sensitive detection systems

CYCD1, RUN 47 (2016)

the 3 main IHC	the 3 main IHC systems*								
Concentrated antibodies	Dak	Ventana BenchMark XT / Ultra			Leica Bond XI / Max				
	TRS pH 9.0	TRS pH 6.1	CC1 pH 8.5	Ν	CC2 pH 6.0		ER2 pH 9.0	Ν	ER1 pH 6.0
rmAb clone EP12	4/5** (80%)	-	3/5 (60%)		-		1/2		-
rmAb clone SP4	20/41** (64%)	0/1	11/31 (49%)		-		2/15 (13%)		0/1
 * Antibody concentration applied as listed above, HIER buffers and detection kips used as provided by the vendors of the respective systems. ** (number of optimal results/number of laboratories using this buffer) 									

Table 3. Proportion of optimal results for CyD1 for the most commonly used antibodies as concentrate on

Bond[™] Polymer Refine Detection

Catalog No: DS9800

Intended Use

This detection system is for in vitro diagnostic use.

Bond Polymer Refine Detection is a biotin-free, polymeric horseradish peroxidase (HRP)-linker antibody conjugate system for the detection of tissue-bound mouse and rabbit IgG and some mouse IgM primary antibodies. It is intended for staining sections of formalin-fixed, paraffin-embedded tissue on the Bond^{*} automated system.

The clinical interpretation of any staining or its absence should be complemented by morphological studies and proper controls. They should be evaluated within the context of the patient's clinical history and other diagnostic tests by a qualified pathologist. The Bond Polymer Refine Detection Kit must be used with laboratory best practice in the use of tissue controls. For assurance, laboratories should stain each patient sample in conjunction with positive, negative, and other tissue specific controls as needed.

Summary and Explanation

Immunohistochemical techniques can be used to demonstrate the presence of antigens in tissue and cells (see "Using Bond Reagents" in your Bond user documentation).

Bond Polymer Refine Detection utilizes a novel controlled polymerization technology to prepare polymeric HRP-linker antibody conjugates. The detection system avoids the use of streptavidin and biotip, and therefore eliminates non-specific staining as a result of endogenous biotin.

Bond Polymer Refine Detection works as follows:

- · The specimen is incubated with hydrogen peroxide to querch endogenous peroxidase activity.
- A user-supplied specific primary antibody is applied.
- · Post Primary IgG linker reagent localizes mouse antibodies.
- · Poly-HRP IgG reagent localizes rabbit antibodies.
- · The substrate chromogen, 3,3'-Diaminobenzidine tetrahydrochloride hydrate (DAB), visualizes the complex via a brown precipitate.
- · Hematoxylin (blue) counterstaining allows the visualization of cell nuclei.

Using Bond Polymer Refine Detection in combination with the Bond automated system reduces the possibility of human error and inherent variability resulting from individual reagent dilution, manual pipetting and reagent application.

The detection system Bond Refine acts by nature as a 2-step polymer system for detection of rabbit polyclonal or rabbit monoclonal primary antibodies

Only enhances reactions with mouse primary antibodies due to the Post Primary IgG linker (Rabbit antibody)

NQC Run 66, 2022: Synaptophysin clone DAK-Synap (RTU GA660)

The choice of the detection system can significantly influence the result

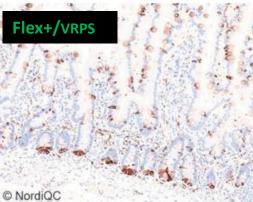
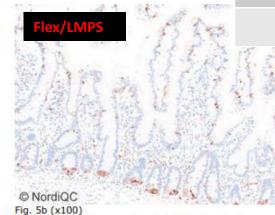


Fig. 5a (x100)

Optimal staining reaction for SYP in the duodenum using the mAb clone DAK-SYNAP (RTU GA660, Dako/Agilent), with vendor recommended protocol settings. The Paneth cells in the bottom of the crypt of Lieberkühn display a strong staining reaction. The goblet cells both in the mucin contained in the cytoplasm and in the cell membrane surrounding the mucin display a moderate to strong staining reaction without any staining of the other epithelia cells. Same protocol used in Fig. 6a.



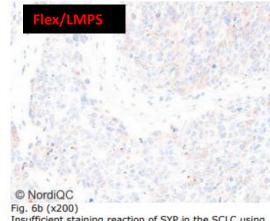
Insufficient staining reaction for SYP in the duodenum using the mAb clone DAK-SYNAP (RTU GA660, Dako/Agilent), with the same settings as in Fig. 5a but without linker. The Paneth cells in the bottom of the crypt of Lieberkühn display a weak to moderate staining reaction and only few goblet cells in the bottom of the crypts are weakly stained.

Same protocol used in Fig. 6b - same field as 1a- 1b and 5a.

Synaptophysin (RTU GA660, DAK-SYNAP)	Pass rate / Optimal results			
VRPS (FLEX+ with linker)	100% (34/34) / 85%			
LMPS (FLEX without linker)	0% (13/13) / 0%			

O NordiQC Fig. 6a (x200)

Optimal staining reaction of SYP in the SCLC using same protocol as in Fig. 5a. All majority of the neoplastic cells show a weak to strong cytoplasmic staining reaction.



Insufficient staining reaction of SYP in the SCLC using same protocol as in Fig. 5b - same field as Figs. 2a- 2b and 6a. The neoplastic cells display a too weak staining intensity with large parts of the tumor completely negative.

NQC Run 70, 2024: GATA3 clone L50-823 (RTU 760-4897)

The choice of the detection system can significantly influence the result

C NordiQC Fig. 1a (x100)

Optimal GATA3 staining reaction of the uterine cervix using the RTU system 760-4897 (Ventana/Roche), based on the mAb clone L50-823, applying vendor recommended protocol settings and OptiView as detection system. The squamous epithelial cells in the basal and intermediate layer of the surface epithelium display a weak to moderate, but distinct nuclear staining reaction, whereas the nuclei of superficial layers and stroma cells are negative. Same protocol settings as in Figs. 2a-4a.

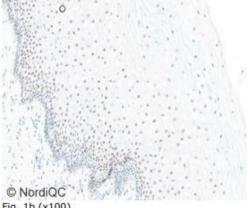


Fig. 1b (x100)

Insufficient GATA3 staining reaction of the uterine cervix using the same RTU system as in Fig. 1a, but with the vendor recommended protocol settings based on UltraView as the detection system. The proportion and intensity of cells expected to be demonstrated is significantly reduced, displaying only faint or false negative staining reaction. Same protocol settings as in Figs. 2b-4b. Same field as Fig 1a.

RTU system 760-4897 (Ventana/Roche) applying VRPS

OptiView vs UltraView ?

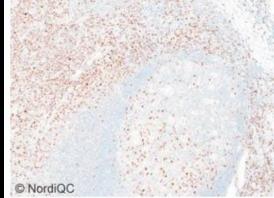


Fig. 2a (x100)

Optimal GATA3 staining reaction of the tonsil using the same protocol as in Figs. 1a-4a. The vast majority of T helper cells (Th2) display a moderate but distinct nuclear staining reaction, whereas the B-cells are negative.

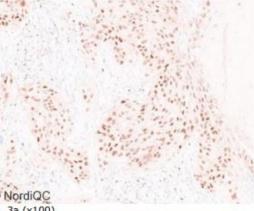


Fig. 3a (x100)

Optimal GATA3 staining reaction of the breast carcinoma using same protocol settings as in Figs. 1a-4a. A weak to strong nuclear staining reaction of virtually all neoplastic cells are seen.

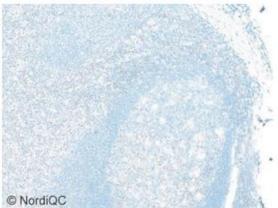
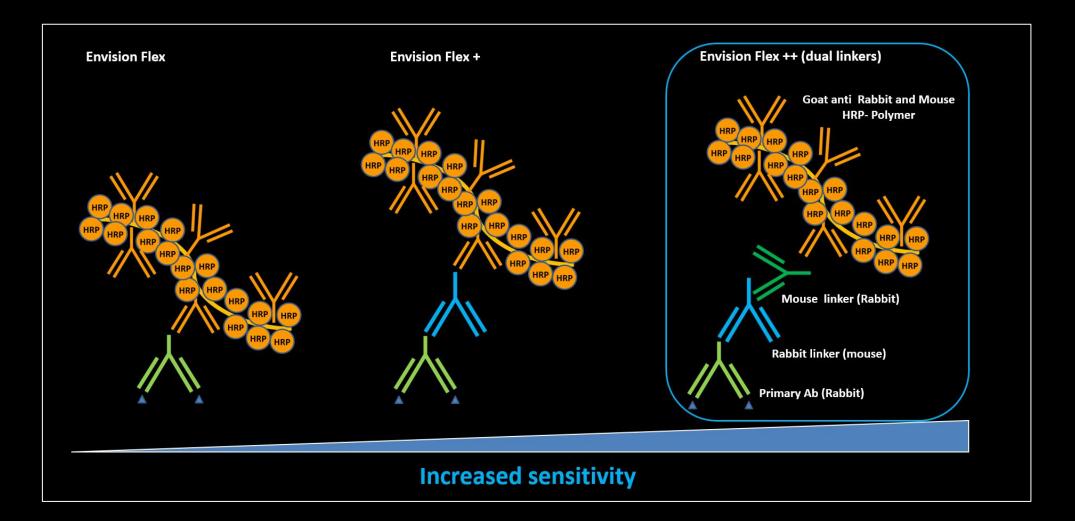


Fig. 2b (x100)

Insufficient GATA3 staining reaction of the tonsil using the same protocol as in Figs. 1b-4b. The vast majority of T helper cells (Th2) are false negative and only a fraction of germinal centre T-cells are weakly demonstrated compare with Fig. 2a, same field.

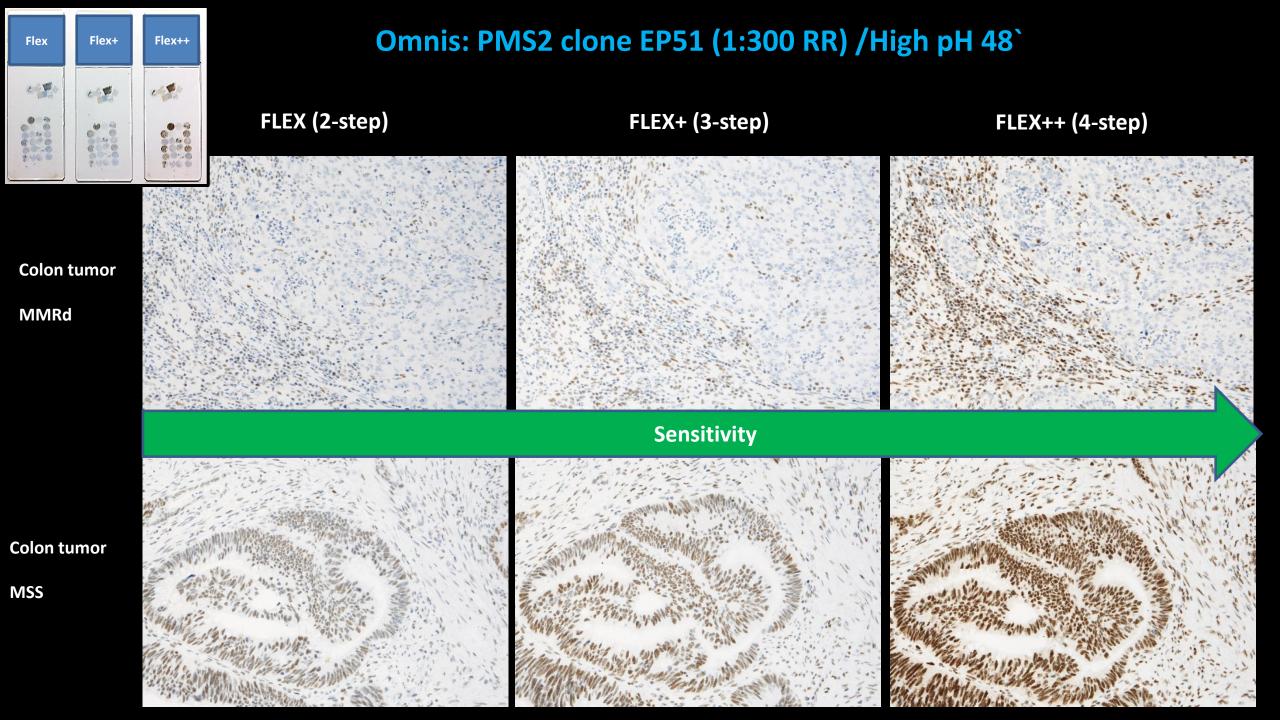


Insufficient GATA3 staining reaction of the breast carcinoma using the same protocol settings as in Figs. 1b-4b. Many of the neoplastic cells are false negative compare with Fig. 3a

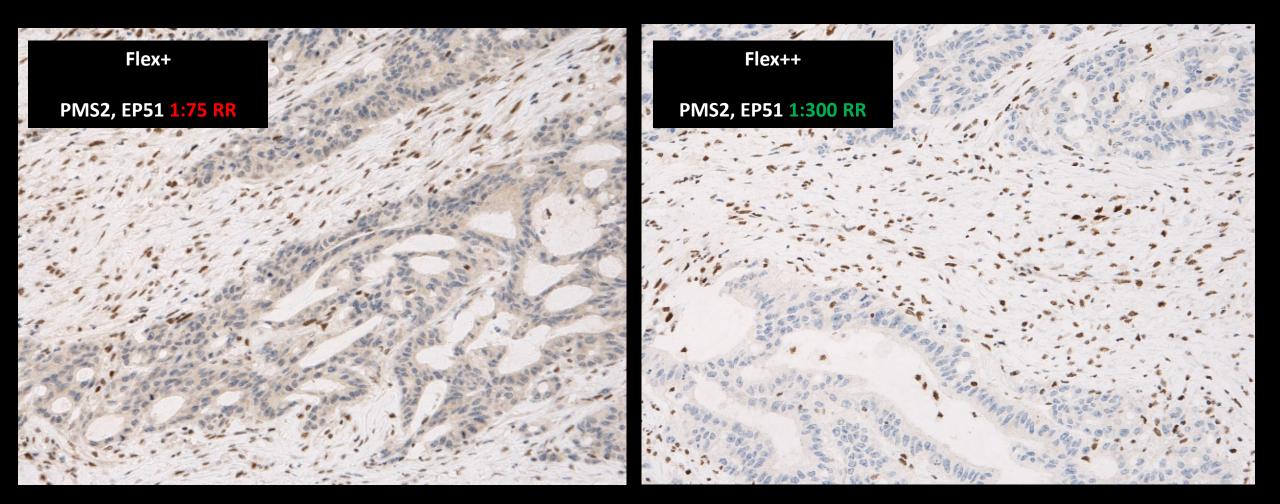


"New option" on the Omnis: Use of dual linkers

In general, works well with rabbit primary Abs but less efficient with primary mouse Abs



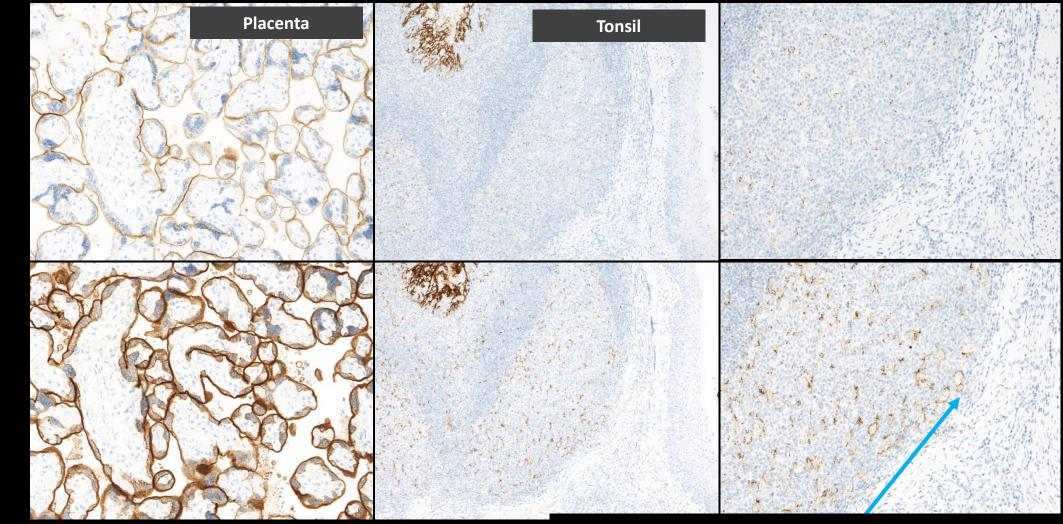
Colon tumor with loss of PMS2: Improves staining quality due to reduced background/noise



Omnis: HIER High pH 48`

Flex++ : 4-step polymer detection system (30-10-10-20`/Omnis)

rmAb PD-L1, CAL10 (rmAb 1:30RR); HIER in High pH 48`



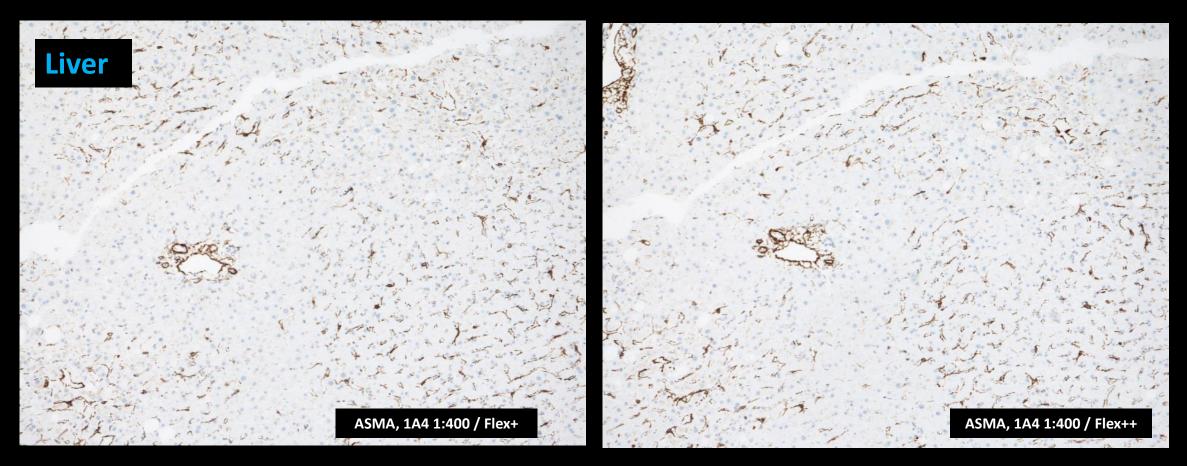
Flex+

Flex++

rmAbs: Dual linkers enhances the sensitivity of the assay

Flex++: 4-step polymer detection system (Omnis)

mAb ASMA clone 1A4; HIER in High pH 24`

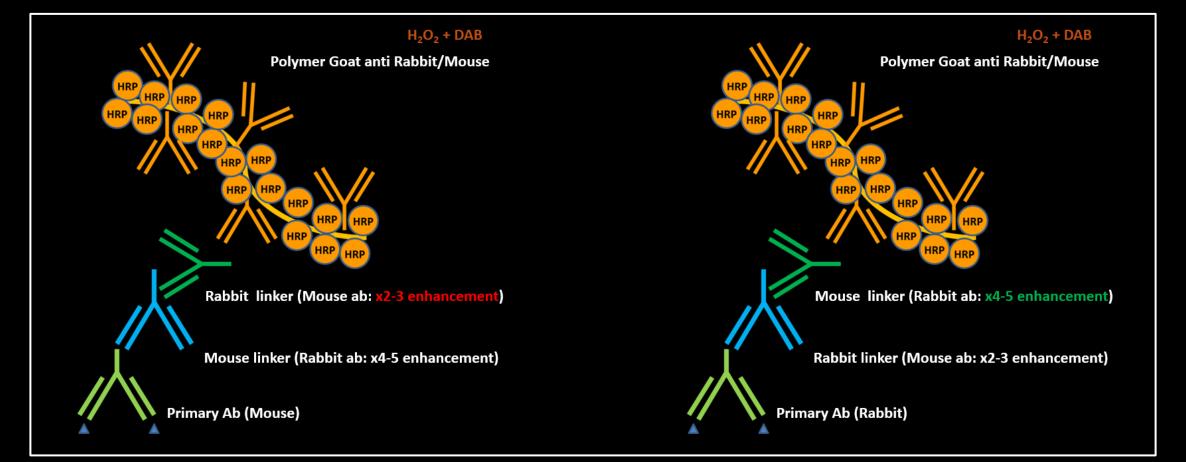


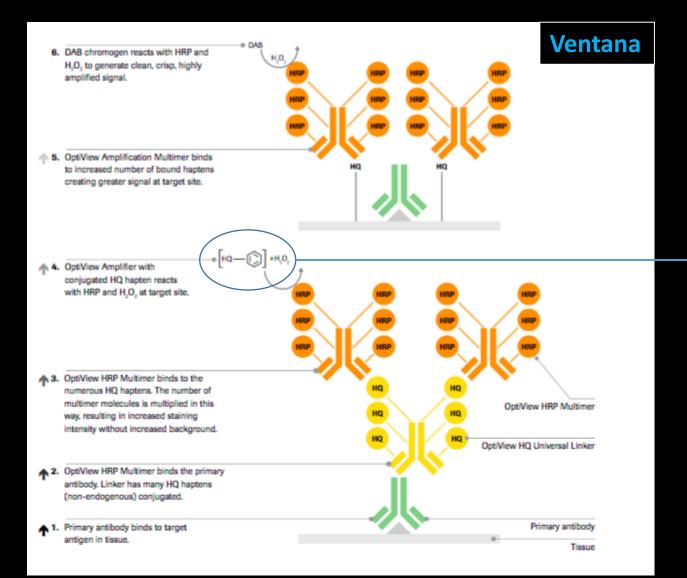
Mouse monoclonal antibodies: In most cases, no improvement in signal intensity using Flex++ compared to Flex+

Envision Flex with Dual Linkers: Mouse versus Rabbit primary Abs

EnVision FLEX+ Rabbit (LINKER) (Code K8009) may be applied for an optional signal amplification of rabbit primary antibodies. For signal amplification when using mouse primary antibodies, EnVision FLEX+ Mouse (LINKER) (Code K8021 or included in Code K8002) may be applied. As a guideline: EnVision FLEX+ Rabbit (LINKER) provides 2–3 fold signal amplification and EnVision FLEX+ Mouse (LINKER) provides 4–5 fold signal amplification.

EnVision FLEX /HRP visualization reagent (SM802) in the kit consists of a dextran backbone to which a large number of peroxidase (HRP) molecules and secondary antibody molecules have been coupled. A unique chemistry is used for the coupling reaction, which permits the binding of up to 100 HRP molecules and up to 20 antibody molecules per backbone.





Tyramide Signal Amplification

Mechanism of Tyramide amplification:

Introducing HRP (Optiview)Incubation with HQ-labelled Tyramide + H2O2

Tyramide, a phenolic compound, is converted into an short-lived extremely reactive intermediate

- Deposit HQ molecules in close vicinity of Ab/Ag reactions

Intermediates covalently binds rapidly to electron rich regions of adjacent proteins (esp. tyrosine)

Detection of HQ with anti-HQ / HRP MultimerVisualization with DAB

Tyramide Signal Amplification (TSA)

Concentrated antibodies	n	Vendor	Optimal	Good	Borderline	Poor	Suff. ¹	OR ²
mAb clone 5A4	2 1 1 2 2 1	Leica Biosystems Monosan Abcam DBS Biocare Medical Zytomed Systems Invitrogen	8	9	14	4	49%	23%
mAb clone OTI1A4*	19 1 1 1	Origene Nordic Biosite Cell Signaling Zeta Corporation	16	6	0	0	100%	73%
mAb clone IHC509	1	GenomeMe	0	0	1	0	-	-
rmAb clone D5F3	19	Cell Signaling	7	9	3	0	84%	36%
rmAb clone ALK1	3 1	Dako/Agilent Cell Marque	0	0	0	4	-	-
rmAb clone QR017	1	Quartett	0	1	0	0	-	-
rmAb clone SP8	1	BioGenex	0	0	0	1	-	-
rmAb clone ZR305	1	Zeta Corporation	0	0	1	0	-	-
Ready-To-Use antibodies								
mAb clone 5A4 PA0306**/PA0831 (VRPS) ³	2	Leica Biosystems	1	1	0	0	-	-
mAb clone 5A4 PA0306*/PA0831 (LMPS) ⁴	10	Leica Biosystems	4	3	2	1	70%	40%
mAb clone 5A4 API3041	1	BioCare	0	0	1	0	-	-
mAb clone 5A4 CAM-0170	1	Celnovte	0	1	0	0	-	-
mAb clone 5A4 MAD-001720QD	1	Master Diagnostica	0	0	1	0	-	-
mAb clone ALK1 GA641	3	Dako/Agilent	0	0	0	3	-	-
mAb clone ALK1 IR641	4	Dako/Agilent	0	0	0	4	-	-
mAb clone ALK1 790/800-2918 (LMPS)⁴	10	Ventana/Roche	1	0	1	8	10%	10%
mAb clone 137E9E8 PA132	1	Abcarta	0	0	0	1	-	-
mAb clone OTI1A4 / 1A4 8344-C010	1	Sakura Finetek	1	0	0	0	-	-
mAb clone OTI1A4 / 1A4 GA785 (VRPS) ³	12	Dako/Agilent	12	0	0	0	100%	100%
mAb clone OTI1A4 / 1A4 GA785 (LMPS) ⁴	4	Dako/Agilent	4	0	0	0	-	-
rmAb clone D5F3 790-4794 (VRPS) ³	73	Ventana/Roche	62	7	1	3	95%	85%
rmAb clone D5F3 790-4794 (LMPS)⁴	48	Ventana/Roche	36	9	3	0	94%	75%
rmAb clone SP8 RMPD007	1	Diagnostic BioSystems	0	0	0	1	-	-
Total	256		152	46	28	30		
Proportion			59%	18%	11%	12%	77%	

Proportion of sufficient stains (optimal or good) (≥5 assessed protocols)

Proportion of Optimal Results (25 assessed protocols).

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

Laboratory Modified Protocol Settings (LMPS) to a specific RTU product (≥5 assessed protocols).
 *) OTI1A4 is called 1A4 by some vendors

**) Product no. PA0306 has been terminated and replaced by PA0831.

<u>Lu-ALK</u>

For certain type of markers, the TSA system provides very good results but.....

Ready-To-Use antibodies and corresponding systems

mAb clone **5A4**, product no. **PA0306/PA0831**, Leica Biosystems, Bond III / Max: Protocols with optimal results were based on HIER using BERS2 (efficient heating time 20-40 min. at 99-100°C), 15-30 min. incubation of the primary Ab and Bond Polymer Refine (DS9800) as detection system. Using these protocol settings, 7 of 8 (88%) produced a sufficient staining result (optimal or good).

mAb clone OTI1A4, product no. GA785, Dako/Agilent, Omnis:

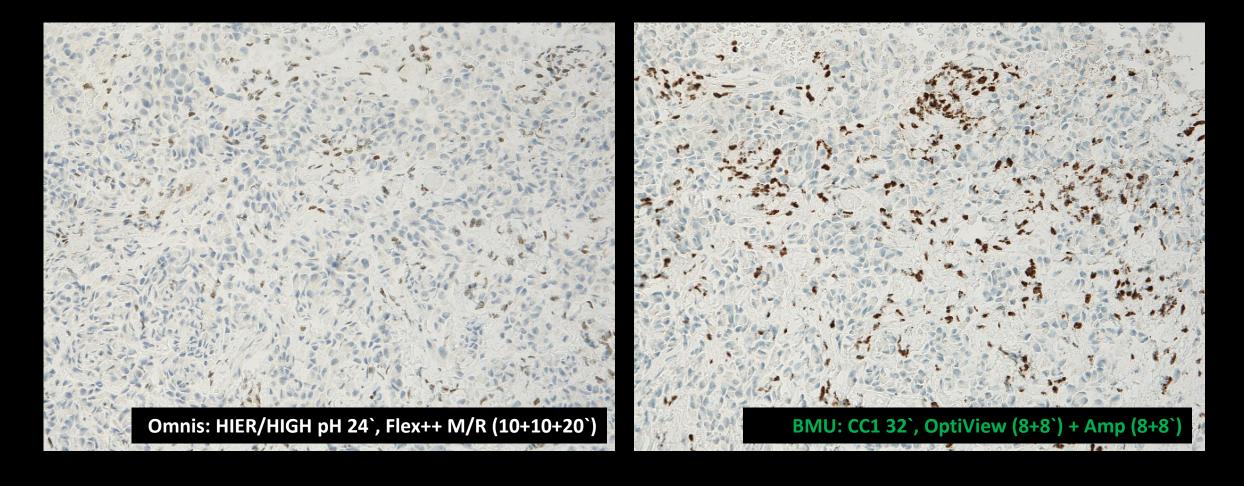
Protocols with optimal results were based on HIER using TRS High pH (efficient heating time 30 min. at 97-99°C), 20-30 min. incubation of the primary Ab and EnVision Flex (GV800/GV823+GV821) as detection system. Using these protocol settings, 16 of 16 (100%) produced a sufficient staining result.

rmAb clone **D5F3** product no. **790-4794**, Ventana/Roche, BenchMark GX, XT and Ultra: Protocols with optimal results were typically based on HIER using CC1 (efficient heating time 92 min.), 16 min. incubation of the primary Ab. and OptiView (760-700) + amplification kit (760-099) as detection system. Using these protocol settings, 110 of 117 (94%) laboratories produced a sufficient staining result.



Problem: Detection System

BAP1 clone C-4 (1:100) / Mesothelioma



The Histochemical Journal 31: 195–200, 1999. © 1999 Kluwer Academic Publishers. Printed in the Netherlands.

Concentration dependent and adverse effects in immunohistochemistry using the tyramine amplification technique

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Received 8 September 1998 and in revised form 1 December 1998

Summary

Although the tyramine amplification technique to enhance sensitivity in immunohistochemistry has been described in numerous methodological papers, it has not yet gained access to diagnostic immunohistochemistry. This is mainly due to problems and pitfalls occurring in adaptation of this method to routine application.

In this study a monoclonal antibody and a polyclonal antiserum (pan-cytokeratin and anti-myoglobin) were tested in tissues with different amounts of epitopes, using a checkerboard table and testing a total of 133 different dilution combinations of both the tyramide solution and the primary antibodies.

The specific tissue investigated, i.e. the amount of accessable epitope to be detected and the applied concentration of the tyramide solution mainly influenced the staining reaction. Several pitfalls such as an uneven distribution of the staining or dramatic overstaining (paradoxical overstaining) must be considered to achieve optimal results.

In conclusion, our data confirm methodological studies that the tyramine amplification technique is a powerful method to enhance immunohistochemical sensitivity. However, for reliable daily practice several pitfalls of the technique have to be circumvented.

TSA enhancement

Difficult to control causing:

- False positive staining

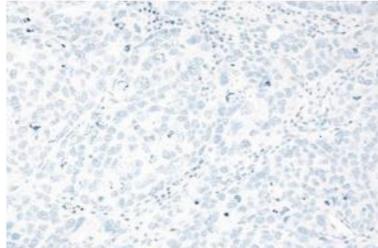
- Weak or false negative staining, unbalanced reaction of primary Ab and target epitopes giving a yes or no answer

- Uneven distribution of the TSA reaction product

NQC Run 66 (Napsin A)

Adenocarcinoma of the lung)

Problem with TSA Amp and false positive reactions (SCC of the lung)



© NordiQC Fig. 5a (x200)

Optimal staining reaction for Napsin A of the lung squamous cell carcinoma using the RTU system 760-4867 (Benchmark Ultra, Ventana/Roche) based on the mAb clone MRQ-60. Vendor recommended protocol settings were applied and OptiView was used as the detection system. All (8/8) protocols based on the same conditions, gave an optimal result. The protocol also provided the same reaction patterns as illustrated in Fig 1a-3a. and as expected, a negative result in the neoplastic cells of the lung squamous cell carcinoma.

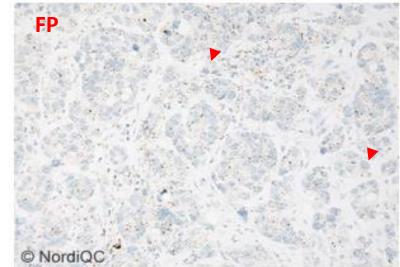


Fig. 5b (x200)

Insufficient staining reaction for Napsin A of the lung squamous cell carcinoma using the same RTU product as in Fig. 5a, but with extended HIER time in CC1 (64 min. at 98°C) extended incubation time in primary Ab (32 min.) and OptiView with amplification as detection

system. The neoplastic cells display a false positive and granular staining reaction. This problem was often seen with tyramide signal amplification. Laboratories should be cautious using this amplification step due to the granular deposit of the reaction product, that erroneously can be interpreted as a specific signal for Napsin A somnare with Fig 5a.

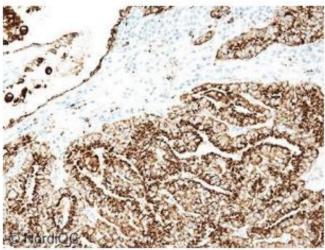


Fig. 3a (x200)

Optimal staining reaction for Napsin A staining of the lung adenocarcinoma (tissue core 3) using same protocol as in Figs. 1a and 2a. All the neoplastic cells display a strong and distinct granular cytoplasmic staining reaction.

TSA: Granular specific reactions vs granular unspecific reactions ?



Fig. 1a (x100)

Optimal PMS2 staining reaction of the tonsil using the GA087 RTU system for the Dako Omnis platform based on rmAb clone EP51, following the recommended protocol settings using dual linker. Virtually all mantle zone B-cells show a moderate and distinct nuclear staining reaction, while the germinal centre B-cells show a strong nuclear staining reaction. Also compare with Figs. 2a - 4a, same protocol.

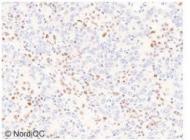


Fig. 4a (x200)

Optimal PMS2 staining reaction of the colon adenocarcinoma, tissue core no. 5, with loss of PMS2 expression using same protocol as in Figs. 1a - 3a. The neoplastic cells are negative, while stromal cells show a distinct nuclear staining reaction serving as internal positive tissue control.

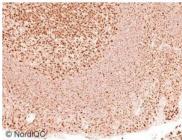


Fig. 1b (x100)

Insufficient PMS2 staining reaction of the tonsil using the mAb clone A16-4 on the Ventana Benchmark platform following the vendor recommended protocol settings with tyramide amplification same field as in Fig. 1a. Virtually all mantle zone B-cells show a moderate but granulated staining reaction of both nuclei and cytoplasm, while the germinal centre B-cells show a strong nuclear staining reaction. Also compare with Figs. 2b – 4b, same protocol

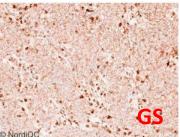


Fig. 4b (x200) Insufficient PMS2 staining reaction of the colon adenocarcinoma, tissue core no. 5, with loss of PMS2 expression using same protocol as in Figs. 1b - 3b, same field as in Fig. 4a. A diffuse granular staining reaction is seen in both the cytoplasmic and nuclear compartment complicating the interpretation of the neoplastic cells expected to be negative.

OptiView + TSA

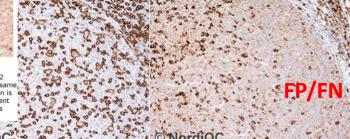


Fig. 1b (x200)

Fig. 1a (x200) Optimal CD4 staining of the tonsil using the rmAb clone SP35 as a concentrate, HIER in an alkaline buffer (CC1 pH 8.5) and a 3-step multimer based detection system (OptiView, Ventana) - same protocol used in Figs. 2a - 4a. The inducer/helper T-cells show a strong staining reaction, while the germinal centre macrophages show a moderate and distinct membranous staining reaction compare with Fig.1b.

Insufficient CD4 staining of the tonsil using the mAb 4B12 as a concentrate, HIER in an alkaline buffer (CC1 pH 8.5) and a multimer based detection system (OptiView with Tyramide amplification, Ventana) -same protocol used in Figs. 2b-4b. The protocol provided a too low sensitivity, but also poor signal-to-noise ratio and false positive staining. No staining of germinal centre macrophages is seen and simultaneously B-cells are labelled. The pattern of too weak staining reaction was observed with all protocols based on the mAb 4B12 performed on the Ventana BenchMark platform - compare with Fig. 1a (same field).

MSH6



Fig. 1a (X100)

control.

Optimal MSH6 staining reaction of the tonsil using the rmAb clone EP49, optimally calibrated, HIER in an alkaline buffer and a 3-step multimer based detection system (OptiView, Ventana). Virtually all mantle zone B-cells show a moderate and distinct nuclear staining reaction, while the germinal centre B-cells show a strong nuclear staining reaction. Also compare with Figs. 2a - 4a, same protocol.

adenocarcinoma no. 4 with loss of MSH6 expression using

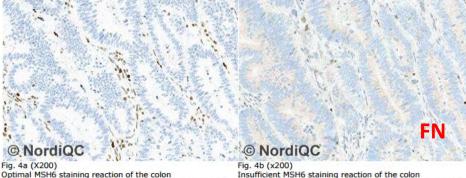
same protocol as in Figs. 1a - 3a. The neoplastic cells are

negative, while stromal cells show a distinct nuclear

staining reaction serving as internal positive tissue

Fig. 1b (X200)

Insufficient MSH6 staining reaction of the tonsil using the rmAb clone EP49 with a protocol providing a too low sensitivity- same field as in Fig. 1a. Only the germinal centre B-cells are distinctively demonstrated, while mantle zone B-cells expressing low level MSH6 virtually are unstained. The protocol was based on OptiView + amplification (Ventana) and the combination of a too low titre of the primary Ab and the use of tyramide based amplification provided an inadequate balance of the staining reaction. Also compare with Figs. 2b - 4b, same protocol.

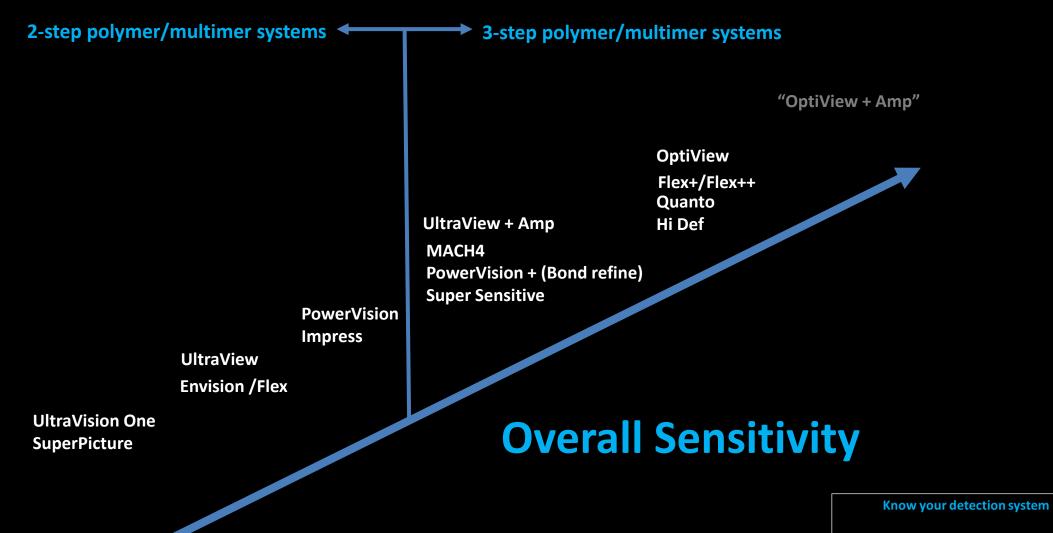


Insufficient MSH6 staining reaction of the colon adenocarcinoma no. 4 with loss of MSH6 expression using same protocol as in Figs. 1b - 3b - same field as in Fig. 4a. No nuclear staining reaction in the neoplastic cells is seen, but as only an equivocal nuclear staining reaction in the normal stromal cells is present, the staining pattern cannot reliably be interpreted. In addition a weak aberrant cytoplasmic staining reaction is seen.

CD4

The TSA detection system is not without problems and may provided either false positive or negative results.

Detection systems - Tested in Dept. of Pathology, Naestved, DK



Strength and weakness

The basal fundament for a technical optimal performance is :

Appropriate tissue fixation and processing

□ Appropriate and efficient epitope retrieval

- 97-98% of the primary Abs require HIER and app. 85-90% prefer high pH (alkaline) retrieval buffers.
- Use efficient HIER temperature and time (app. 100°C for 20 40min).

□ Appropriate choice of antibody / clone, diluent and dilution

- Compare different clones / Abs against the selected antigen of interest before implementation
- Calibrate the Ab concentration carefully

□ Appropriate and specific / sensitive detection system

- Use of a 3-step multimer/polymer system is preferable to a 2- step multimer/polymer system
- Don't use biotin-based detection systems

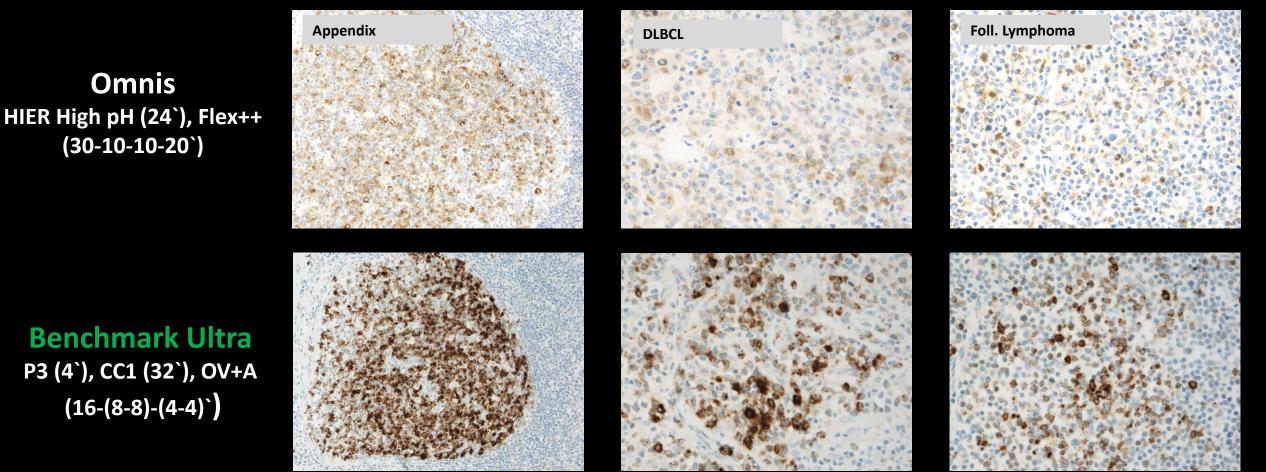
□ Appropriate choice of control material

Focus on Immunohistochemistry Critical Assay Performance Controls (iCAPCs)

Thank you for your attention



Problem: Detection systems



Serpin A9/GCET1 (clone 585302/1:2000RR or RAM341/1:200)

Benchmark Ultra P3 (4`), CC1 (32`), OV+A (16-(8-8)-(4-4)`)

Omnis

(30-10-10-20`)

Data & method kindly provided by Ole Nielsen, Dept. of Pathology, Odense, Denmark

Less sensitive detection systems

CD10, RUN 39 (2013)

Table 2. Optimal results for CD10 using concentrated antibodies on the 3 main IHC systems*								
Concentrated	Concentrated Dako Ventana Leica							
antibodies	Autostainer I	Link / Classic	BenchMark	XT / Ultra	Bond III / Max			
	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	64 %	0%	67 %		95 %	0 %		
56C6	14/22**	0/1	35/52	-	19/20	0/1		

* Antibody concentration applied as listed above, HIER buffers and detection kits used as provided by the venders of the respective platforms.
** (number of optimal results/number of laboratories using this buffer)

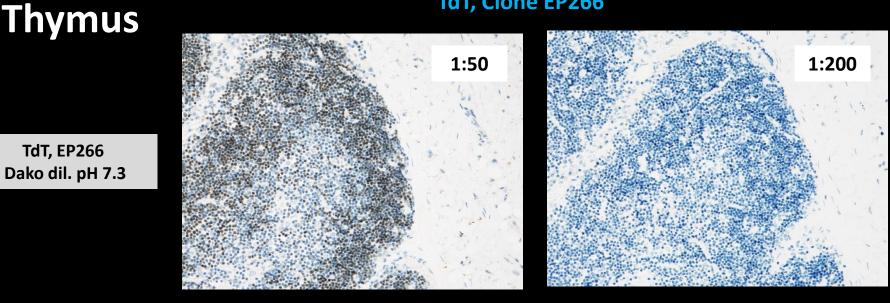
Protocols based on 2-step polymer/multimer detection systems, provided a sufficient staining result in 91% (n=42/46), and in 52% (n=24/46) an optimal result was obtained. If the same protocol settings were applied with a 3-step polymer/multimer based detection system (EnVision FLEX+ (Dako), Optiview (Ventana) or Bond Refine (Leica)), sufficient staining results were seen in 100% (n=58/58), and 86% (n=50/58) was evaluated as optimal. This is also reflected in the high proportion of optimal results (95%) using the mAb clone 56C6 as concentrate on the Leica IHC platforms (table 2), on which a 2-step polymer based detection system is used as standard.

2 & 3-step detection system (Flex/Flex+ or UV/OV

BOND refine (3-step detection system)

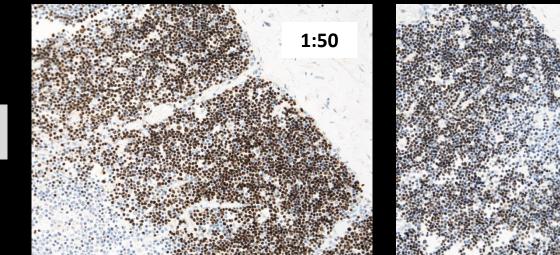
LD assay (mmAb clone 56C6) HIER in alkaline buffer and optimal dil. range	Detection system	Pass Rate`s (%)	Optimal (%)
2-step polymer/multimer system	Flex (Dako) or UltraView (Ventana)	91 (42 of 46)	52 (24 of 46)
3-step polymer/multimer system	Flex+ (Dako), OptiView (Ventana) or BOND Refine (Leica)	100 (58 of 58)	86 (50 of 58)

Antibody-Antigen reaction – Antibody Diluents



TdT, Clone EP266

Omnis: HIER/HIGH pH 24`, Flex+ Rabbit (10+20`)



1:200

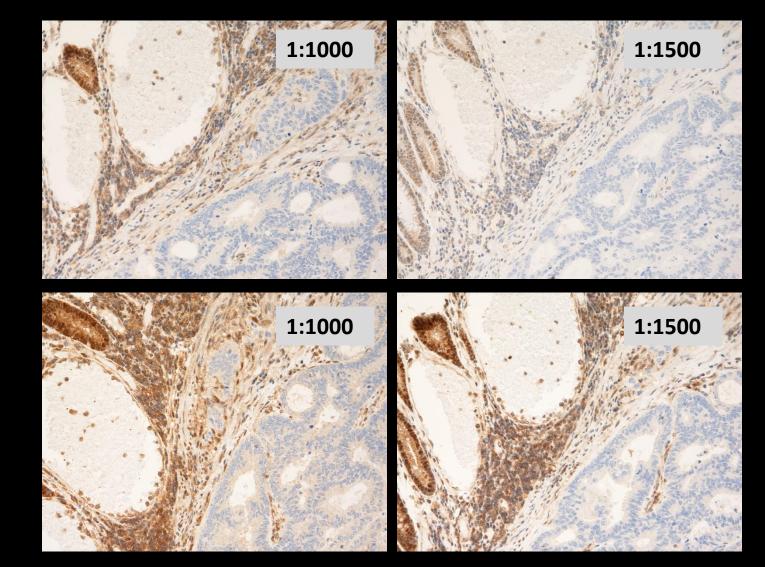
TdT, EP266 Renoir Red pH 6.2

TdT, EP266

Antibody diluents

SMAD4, Clone EP168Y

Colon Adenocarcinoma



Dako dil. pH 7.3

Renoir Red pH 6.2

Research Article

Performance of 4 Immunohistochemical Phosphohistone H3 Antibodies for Marking Mitotic Figures in Breast Cancer

Cornelia M. Focke, MD,* † Kai Finsterbusch, MSc,* Thomas Decker, MD,* and Paul J. van Diest, MD, PhD †

Conclusions: Performance and reliability varied significantly between the 4 tested antibodies. For faster identification of mitotic hot spots and as potential marker in digital image analysis, the Merck antibodies seem to be most suitable.

Key Words: PHH3, breast cancer, proliferation, mitotic activity, phosphohistone H3

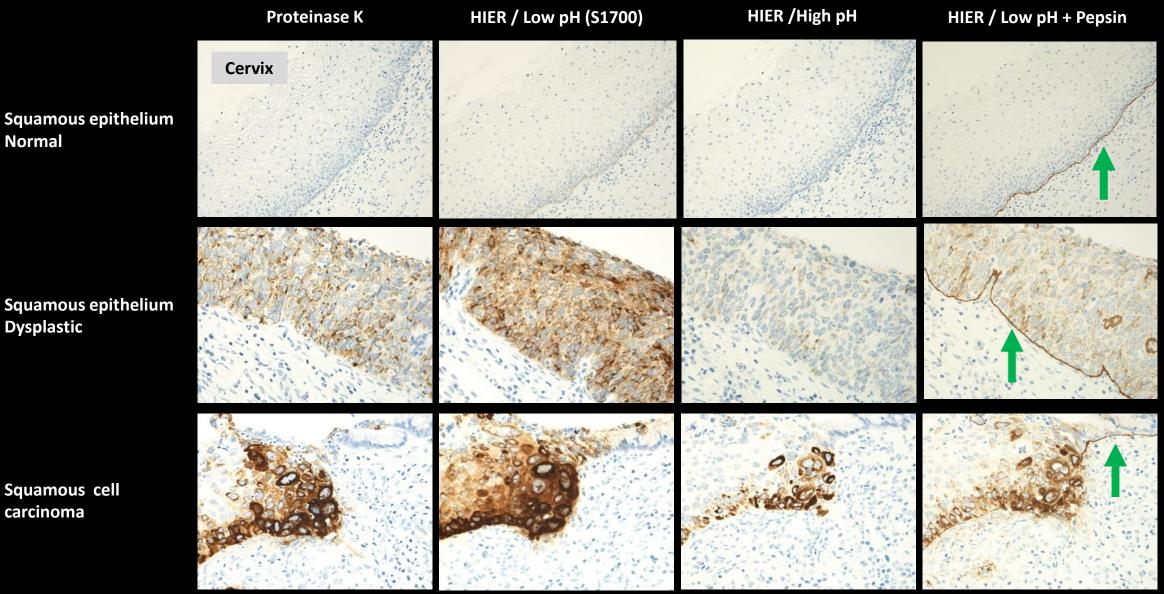
(Appl Immunohistochem Mol Morphol 2016;00:000-000)

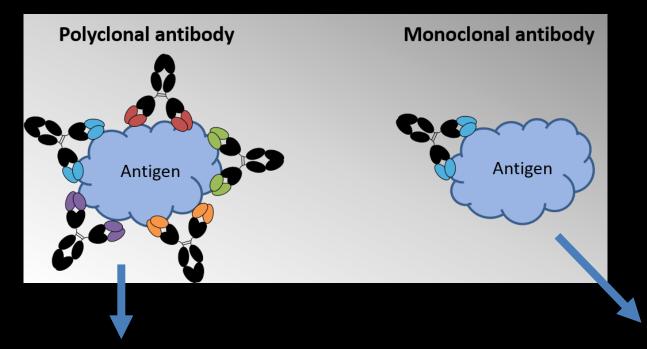
LAM-5 (γ2), D4B5

Normal

Dysplastic

carcinoma





Antibodies bind to antigen through the variable regions of the antibody.

The strength of the binding of an antibody to a specific epitope is called affinity.

High affinity antibodies will bind larger amounts of antigen in a given period of time and can be used at higher dilutions.

Heterogeneous population of antibodies reacting with different epitopes of an antigen

Not epitope specific

Might cross-reactivity with similar antigens

Background noise

Batch/lot variability

Many host species options: Normally rabbit

Homogeneous population of a specific antibody reacting with one epitope on a particular antigen

Epitope specific

No cross-reactivity with similar antigens

Less background noise

Less batch/lot variability

Host species options: Normally rabbit and mouse