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TITLE

NOT ALL ANTIBODIES HAVE BEEN CREATED EQUAL: Antibodies validated for routinely processed tissue seldom apply to frozen tissue sections.

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RUNNING TITLE

Re-validated antibodies on frozen sections.

CONFLICT OF INTERESTS

The Authors have no conflict of interests to declare.

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ABSTRACT

A customized context-dependent validation of antibodies for the prospective use, rather than a general stamp of validity, is required for reproducibility and data validity, besides the need of standardized reagents. In-situ antibody staining is a technique broadly used in experimental settings and human diagnostic practice. The first typically, but not exclusively uses lightly fixed material (cell smears, frozen sections), the second, routinely processed formalin-fixed, paraffin-embedded (FFPE) tissue. Differently from techniques based on tissue extraction, there is little awareness of the context-dependent constraints inherent with either type of in situ staining except that antigen masking associated with routine tissue processing limits the range of useful antibodies. We applied a panel of 126 antibodies validated for FFPE to lightly fixed (acetone, formalin) frozen sections and found that less than 30% performed conservatively with all fixations, 35% preferred one fixation over another, 13% gave non-specific staining, 23% did not stain at all. Individual antibody variegation of the paratope fitness for the differentially fixed antigen may be the cause. Re-validation of established antibody panels is required when applied to sections whose fixation and processing is different from the tissue where they have been initially validated.

INTRODUCTION

Over the last fifteen years, awareness of the frequent variability and unreliability of data caused by the use of non-standardized, not validated antibodies (Abs) in various types of immunoassays such as Western Blot (WB), Immunocytochemistry (ICC) and Immunohistochemistry/immunofluorescence (IHC/IF) (1-4), has suggested common guidelines for the correct use of these reagents. An International Working Group for Antibody Validation (IWGAV) convened in 2016 and published a proposal to ensure reproducibility and validity when using antibodies for immunolabeling (5). Five validation principles and criteria were proposed, based on: stain reduction upon genetic ablation of the target ("genetic"), comparison with an antibody-independent method ("orthogonal"), comparison of two different Abs against the same target ("independent antibody"), comparison with a "tagged protein expression" and mass spectrometry analysis of the immunocaptured protein ("IMS"). The suitability of each validation method for several applications, including in situ staining methods (ICC and IHC) was reported as well (5). Guidelines for antibody reporting in publications were published (3, 6) around the same time.

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These suggested guidelines apply to very diverse subjects in many science fields, including Pathology. Surgical Pathologists in particular are using IHC in an highly regulated diagnostic settings, where pre-analytical, analytical and reporting rules need to be strictly defined, because of the human health implications of these assays (7). Surgical Pathology Departments may routinely use >200 different primary antibodies and introduce about 20 new antibodies a year (7), however the focus of the antibody/assay validation is on technical reproducibility across the caseload rather than on the antibody validation as intended by the International Working Group mentioned above. The antibody specificity for the target is a given assumption, often provided by the company supplying the antibody or the assay. On the other hand, the type of tissue target in Surgical Pathology is overwhelmingly FFPE tissue sections, whose pre-analytical variables are also highly regulated (8). The modifications of the protein targets in FFPE material and the performance of antibodies in such conditions have been extensively published and are the basis for a successful use of FFPE tissue for diagnostic, prognostic and theragnostic purposes (9-11). Because of the robustness of tissue processing and immunoanalysis, archived FFPE human material and Surgical Pathology techniques are also the most abundant and diffuse tools for translational and basic science research.

The IWGAV recommendations are that "approaches for antibody validation must be carried out in an application- and context-specific manner." (5), suggesting that antibody validation must be independent for extractive (e.g. WB, immunoprecipitation) versus in-situ technique (ICC, IHC) on the basis of the different modifications that the protein target undergoes in each procedure (see Supplementary Table 1 in (5)). Although the IWGAV makes a distinction between ICC and IHC, reports on the requirement for an independent antibody validation on either type of tissue type are scanty at best, while the underlying assumption in the vast majority of publications is that an antibody good for FFPE must be good for a frozen section or a cell preparation as well and vice versa. The only caveat is that antigen masking in FFPE tissue makes it harder to find antibodies working on both types of tissues. A handful of groups (12-14) addressed in a systematic fashion the difference in antibody performance in FFPE versus lightly fixed frozen tissue section, these latter believed at that time to be the "golden standard" (12) to assess distribution, intensity and, ultimately, validity of an antibody. In those publications, the focus was on the differential antigen retention with each fixation protocol, rather than guestioning the antibody specificity, which was a given predefined value, also because coming from the Surgical Pathology practice.

We have previously investigated the epitope specificity and binding requirements of a large number of antibodies for human FFPE material by applying them to swine routinely fixed

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material (15). The results showed that about 50% did not react, despite complete identity of the epitope in both species. This may have occurred because of the fine requirement for some Abs of the conformation of the target side peptide chains not involved in the antigen binding (16) or because of non-identical juxtaposed adjacent proteins in the fixed complex. Of the Abs reacting with both species, we had evidence of species-specific negative effect of formalin fixation of BCL6, restricted to one antibody and not to others, while all did reacted on acetone-fixed frozen sections (15). In a separate investigation (11), we found that the epitopes for some antibodies were selectively destroyed by antigen retrieval (AR), possibly because of the conformational nature of the epitope.

These observations suggest that validated antibodies for FFPE sections may bear undisclosed paratope variations which may affect sensitivity, validity and usage when applied outside that context. Therefore we investigated a large panel of antibodies, validated on FFPE, on lightly fixed (acetone, formalin) frozen tissue sections. Antibodies for FFPE sections may detect linear epitopes (15) in a profoundly denatured protein mixture, thus we expected them to be working spectacularly on lightly fixed frozen material, but we found that this is quite often not the case.

MATERIALS AND METHODS

Human specimens

Fully anonymized human surgical specimen leftovers (pediatric tonsils, normal kidney, discarded serial sections from routinely processed formalin-fixed, paraffin-embedded (FFPE) material) were either snap-frozen in -80C chilled isopentane (Merck Life Science S.r.I.,Milano, Italy) or fixed overnight at RT in buffered 4% formaldehyde (Bio-Optica Milano Spa, Milano, Italy), processed through a graded ethanol gradient, then in xylene and embedded in molten paraffin for sectioning.

The study has been approved by the Institutional Review Board Comitato Etico Brianza, N. 3204, "High-dimensional single cell classification of pathology (HDSSCP)", October 2019. Patients consent was obtained or waived according to article 89 of the EU general data protection regulation 2016/679 (GDPR) and decree N. 515, 12/19/2018 of the Italian Privacy Authority.

Antibody validation

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A list with all the primary antibodies validated for FFPE use can be found in in Supplementary Table 1. The type of validation of each reagent was listed essentially according to Edfors et al. and Uhlén et al. (5, 17) and modified for tissue staining. The following criteria were used:

- <u>Orthogonal</u>: an antibody uniquely identifying a cell in the tissue whose high-dimensional phenotype (18, 19) corresponds to a cell described by single cell RNA sequencing (i.e transcriptomics) and bearing the transcript corresponding to the target is considered validated.
- <u>Independent Antibody</u>: A) two antibodies directed against two separate epitopes of the same protein and having an identical staining pattern and/or co-localizing by double IF or B) an antibody uniquely identifying a cell in the tissue whose high-dimensional phenotype corresponds to a cell whose phenotype is defined by a multidimensional flow cytometry panel, are considered validated
- <u>Genetic</u>: an antibody whose staining or absence of staining corresponds to a genetically engineered ectopic expression or absence of the target is considered validated
- <u>Peptide Microarray</u>: an antibody recognizing its unique peptide target on peptide microarrays (20, 21) is considered validated
- <u>Cross-species</u>: an antibody whose reactivity is conserved across genomic sequence variations in another species (15) is considered validated

We added an additional criterion, "historic", for antibodies whose widespread use in multiple applications (e.g. CD20) strongly suggests validity, despite lacking one of the criteria listed above.

Evidence of validation not present in the published literature, either peer-reviewed papers or documentation from producers, was produced in house. Some antibodies were validated according to multiple criteria. Validated, FFPE-proof antibodies directed against the same protein can be used exchangeably on routinely processed sections, besides obvious variations in host species or isotype (22).

Representative iconography of the FFPE staining for each antibody can be viewed in published papers ((19, 23)), in (24) or at the Human Protein Atlas website (https://www.proteinatlas.org/).

Frozen tissue fixation

Frozen tissue blocks were sectioned at 5 µm thickness in a Leica CM1850 cryostat (Leica Microsystems GmbH, Wetzlar, Germany) at -20C. Sections were collected on positively charged slides (Menzel-Glaser Superfrost Plus; Bio-Optica Milano Spa).

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Two fixation methods were chosen: in one, sections on slides were lifted from the cryostat, placed temporarily in a slide rack, immersed in acetone (Merck) at +4C for 5 min, allowed the acetone to evaporate, stored at RT overnight in a damp-proof box; subsequently, the sections were either used the same day or stored wrapped in plastic foil at -20C for later use. In the second method, slides with the frozen section attached were immediately immersed in 4% buffered formaldehyde (FA), fixed at RT for 18h (overnight), then washed in Tris-containing buffer for formaldehyde quenching (25). These sections were stored at +4C in 50 mM Tris HCl buffer pH 7.5, containing 0.01% Tween-20 (Merck) and 100 mM sucrose (TBS-Ts). Previous published experiments (11) showed that fixation in excess of 30 min is required for tissue stabilization and epitope rescue upon AR and 48h or 72h fixation do not add additional masking, compared to overnight fixation, therefore we chose this latter fixation time for all the experiments.

Frozen sections were dipped for 5 min in 2.5% horse serum in TBS (Vector Laboratories, Burlingame, CA, USA) before use for Fc receptor blocking.

Antigen retrieval

Antigen retrieval (AR) was performed placing the sections in a 800 ml glass container filled with the retrieval solutions (10 mM EDTA in Tris-buffer pH 8, Merck), irradiate in a household microwave oven at full speed for 8 min, followed by intermittent electromagnetic radiation to maintain constant boiling for the set time, and cooling the sections to about 50 °C before use. We found that 20 min of AR on FA-fixed frozen sections did not produce any additional retrieval compared to 1 min or 5 min (not shown), thus a protocol comprising the 8 min preheating and 1 min AR was used throughout.

Immunolabeling.

Sections were processed for indirect immunohistochemistry or immunofluorescent labeling as previously described in detail (19, 23).

Briefly, the sections were incubated for 30 min with optimally diluted primary antibodies in combination of up to four (22), washed and counterstained with specific distinct fluorochrome-tagged secondary antibodies (23). The slides, counterstained with DAPI and mounted, were scanned on an S60 Hamamatsu scanner (Nikon, Italia) at 20x magnification (23). Optimal exposure time for each antibody-fluorochrome combination was set in advance on FFPE and validated over multiple samples in a multiplex experiment on routinely processed material ((18) and not shown). Quadruple simultaneous IF staining guarantees internal control for effective

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staining. Dubious results were confirmed in single stain IHC. The vast majority of the tests were done on tonsil sections, which constitute the optimal positive control for a panel in which anti-leukocyte antibodies are overrepresented. Acetone-fixed, formalin-fixed and formalin-fixed AR treated frozen sections were all processed simultaneously.

Results were scored semi-quantitatively (neg, +/-, +, ++, +++) and qualitative staining features (diffuse, non-specific staining, etc.) were annotated.

Preparation of immunofluorescent images for image analysis.

Single .ndpi images for each case were saved as .tiff files and autofluorescence (AF) was subtracted (23). Grayscale images were inverted, brightness and contrast adjusted with the default ImageJ function (26), simultaneously processing a stack of three experimental sets (acetone, FA, FA>AR) before producing individual images.

RESULTS

The overwhelming majority of antibodies tested were selected for being reactive with AR-treated FFPE sections (118/126), yet 8 (6.3%) selectively failed to stain acetone fixed sections and additional 8 had minimal reactivity compared to FA fixed sections.

Fourteen, including these latter eight, required formalin fixation to be detected (11.1%) (Table 1, Fig. 1 and labelled "FA dependent" in Supplementary Table 1) and AR was able to enhance the staining in only three of them.

FA fixation did reduce epitope detectability in a quarter of the cases and abolished the reactivity in as many as 24 cases (19.8%) ("FA blocked" in Table 1 Fig. 1 and Supplementary Table 1); in all these instances, AR rescued the optimal reactivity.

The epitope composition of the "FA blocked" antibodies was available for 5 Abs only and short enough to be informative in one (CD3-12; ERPPPVPNPDYEPC). The epitope contained no lysine, the topmost aminoacid bound by formaldehyde, yet was sensitive to FA fixation on frozen sections.

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A group of 17 abs (13.5%) had immunoreactivity either selectively elicited by AR treatment (12 Abs) or abolished by the heat treatment. We defined those as "conformational" because of a possible disruption (or creation) of an epitope by altering the conformation of one or more protein loops generating the epitope (Table 1, Fig. 1 and Supplementary Table 1)

A group of 6 Abs (4.8%) we named "miscellaneous" (Table 1, and Supplementary Table 1) contained Abs behaving differently than the previous groups and were composed mostly of Abs reacting with acetone-fixed sections only. One antibody (GZMB) labelled selectively FA-fixed frozen sections.

Twenty nine Abs (22.8%) did not stain any type of lightly fixed sections.

The remaining Abs (35/126; 27.8%) had conserved reactivity across the three preparations: acetone fix, FA fix, FA>AR ("conserved"; Table 1, Fig. 1 and Supplementary Table 1). Whenever the protein distribution was shared with cells of diverse origin (e.g. podoplanin on endothelium, basal epithelium and follicular dendritic cells), differential response to the fixation protocols was noted according to the cell type. As an example, only the endothelial staining of Podoplanin (Fig. 1) conserved an equivalent intensity across the three fixation protocols. For some protein, a loss of the target from sections in acetone fixed tissue was noticed (e.g. LYZ, S100AB)

Spurious reactivity, either due to non-specific, background staining or to discrete unrelated targets (Fig. 1) was detected in 17 instances (13.4%), 10 of them (7.8%) in FA fixed sections.

In 19 instances, the same protein was tested with multiple independent antibodies which showed in all instances but two individual variations in staining pattern (Supplementary Table 1).

Because of the antibody panel choice, biased toward leukocyte antigens, most Abs did not react with swine FFPE tissue (15), thus no epitope variation information could be obtained.

DISCUSSION

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A minority of antibodies validated for the use on FFPE material apply for the use on frozen sections, formalin- or acetone fixed. The most likely reason for such unexpected results may lie in the subtle requirement of the paratope for an unique epitope binding environment, composed not just of the epitope, but also of the neighboring residues on the protein sequence (16) or of adjacent unrelated proteins and their post-translational modifications (27). Such epitope microenvironment is affected by the fixation method and, when applied, by further denaturation provided by high heath.

So far, the burden of the proof has been put on the antigen and its modification during fixation. Our investigation however started from anti-peptide antibodies validated on the new "gold standard", FFPE material (12), went back to the frozen tissue perceived as containing native proteins, and has unveiled how context-dependent the use of Abs for in situ detection can be.

Differently from previous published work (12-14), the use of multiple antibodies for the same target (35% of the panel) has disclosed a remarkably heterogeneous staining ability within the same target across the experimental conditions, suggesting that the source of variance is the paratope variegation, rather than epitope.

It has also shown evidence of the persistence of the antigen in the tissue with that particular fixation condition, indirectly re-validating the use of acetone-fixed or formalin-fixed frozen sections as "alternative gold standards" for antigen detection in tissue. Only a couple of antigens may have been lost from acetone-fixed sections, as previously published (12, 13).

The recommendation to validate and use antibodies limited to the technique employed (Western Blot, immunoprecipitation, immunohistochemistry, etc.) (5, 17) may now be made in a more detailed fashion also for immunological in situ techniques, distinguishing frozen sections from FFPE material, and within frozen sections, whether acetone of FA is used and if AR is required. Particularly worrisome for staining reliability across fixations is the presence, albeit small, of unrelated discrete reactivity (e.g. CD8 on epithelia; this report and (28, 29)) or the variegated effect of treatment on epitopes shared across diverse cell types (e.g. Podoplanin, PDL1) (Fig. 1).

The major bias of this study is the antibody choice, which was guided by the validity on routinely processed tissue sections, being only 9 of the selected Abs not reactive on FFPE. Another bias is the abundance of mouse non-IgG1, non-rabbit antibodies (42%), due to the multiplexing

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staining strategy (22), selection which however may have highlighted the problem by diversifying, besides the immunizing epitope, the host view of the immunogen.

Collectively, these data suggest that antibody binding may be very dependent on tissue fixation and processing in a subtle and unexpected fashion, affecting binding and specificity for the target.

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AUTHOR'S CONTRIBUTIONS

GC and MMB equally designed the experiments.MMB devised the image analysis algorithms, performed visual and digital image analysis as well as bioinformatic evaluation.FM and SB performed immunostaining experiments.FMB and RP provided essential reagents.

MMB, FMB and GC wrote the manuscript.

All authors have read and approved the final manuscript.

COMPETING INTEREST STATEMENT

The authors declare they have no competing interests.

TABLE LEGENDS

Table 1

Summary table of the staining results, grouped by category.

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Examples of antibody staining on lightly-

FIGURE LEGENDS

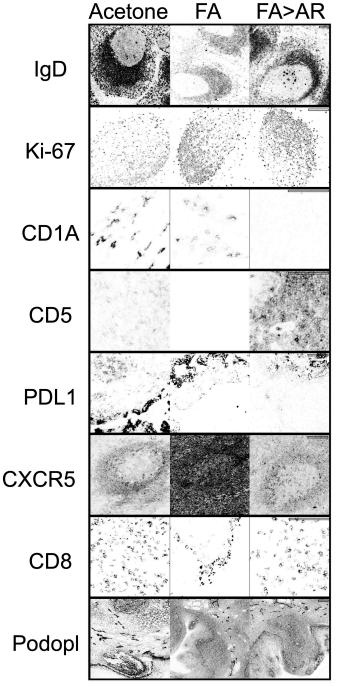


Figure 1

fixed frozen tonsil tissue. Representative examples of stainings obtained on acetone, formalin fixation (FA), and FA fixation followed by AR (FA>AR), left, middle and right columns respectively. From top to bottom: IgD (goat poly) and Ki-67 (UMAB107) are conserved in intensity across the fixations, the former reduced after FA fixation. CD1A (O10) staining is lost upon heating (AR). CD5 (CD5/54/F6) staining is lost upon FA fixation and regained after AR. PDL1: macrophage and epithelial cell staining for PDL1 (cocktail of rabbit MAbs) is maximal in Acetone-fixed section. Macrophage stain is all but gone in FA fixed, but partially retrieved after AR. Epithelial staining is preserved in FA and a basal layer staining is added, this latter lost upon AR. CXCR5: formalin fixation causes a diffuse non-specific staining for CXCR5 (#51505). Lymphocyte CD8 (C8/144B) staining is maintained and specific, except upon FA fixation, where is greatly reduced, while a non specific strong staining of the basal epithelial layer is produced.

Podoplanin (NZ-1.2) produces a bright staining of the basal epithelium, endothelial cells and follicular dendritic cells in acetone-fixed cells. Upon FA fixation, endothelial cells remain strong, while the other structures are quite weaker.

Scale bar = 50 μ m.

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SUPPLEMENTARY MATERIAL

SUPPLEMENTARY TABLE 1

List of primary antibodies.

Abbreviations:

Species: Rb= rabbit; RbMab= rabbit monoclonal ab; g1, g2a, g2b, g3= mouse isotypes.

FFPE: yes= working on routinely processed material.

RRID: Resources Identification Portal identifier (https://scicrunch.org/resources)

Classification: conformational= conformational epitope; conserved= staining conserved over fixation types; FA blocked= reactivity blocked by FA fixation; FA dependent= staining depends on FA fixation; negative= no staining; misc= miscellaneous behaviour, mostly Acetone only, not comprised in the previous classification. Spurious= spurious, non-specific reactivity present. Validation: type of antibody validation as described in M&M.

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TABLE 1

Reactivity	Acetone	FA	FA > AR	N. Abs	%	Note			
conserved	pos	pos	pos	35	27.8%	Reactivity preserved on all fixations			
FA blocked	pos	neg	pos	25	19.8%	Inactivated by FA fixation (rescued by AR)			
FA dependent	neg	pos	pos	14	11.1%	Requires FA fixation			
conformational	pos	pos	neg	17	13.5%	Heat-induced appearance/disappearance			
	neg	neg	pos						
misc	pos/neg	neg/pos	neg	6	4.8%	Mostly Pos in Acetone fixed only			
neg	neg	neg	neg	29	23.0%	No reactivity			
Total				126					

t Mab	Ab AID AXL	clone MAID-2 C89E7	Source Thermo CST	FFPE R yes	10669583 neg 11217435 +++	FA neg +	FA> AR neg +/-	Ac	FA snarse I ymphs, endoth	FA> AR	Classification neg misc	Spurious r	e: validation genetic
UBD	bcl-2	Bcl-2-100	Merck	yes yes	258541 +++	+ ++++	+/- +++	endothel +?	sparse Lymphs, endoth	Sparse lymphs	conserved		genetic
	BCL6 Blimp1/PRDM1	D-8 6D3	SCBT SCBT	yes yes	2063455 +++ 628168 note	+++ +/-	++ +++	background			conserved FA blocked		independent Ab independent Ab, historic
	BMI1 CD10	F6 NCL-CD10-L-270	Millipore Leica-Microystem	yes	309865 neg 442048 ++	neg	neg ++	5			neg conserved		cross-species
b	CD103	EP206	Bio SB	yes	+++	neg	+++				FA blocked		cross-species
	CD11c CD123	B-6 7G3	SCBT BD Pharmingen	yes no	626859 neg 395454 +++	neg +++	neg +++				neg conserved		orthogonal
	CD123 CD133	NCL-L-CD123 AC133	Leica-Microystem Milteny	yes yes	10555271 neg 244341 +++	neg	++++				conformational conserved		orthogonal, historic orthogonal
	CD14	5A3B11B5	SCBT	yes	781932 +/-	bckgnd	+++				conformational	spurious	independent Ab
	CD14 CD141	poly EPR4051	Merck Abcam/Epitomics	yes yes	1078467 neg 2201805 +++	neg	neg ++				neg FA blocked		independent Ab independent Ab
b	CD141 CD141	D-3	SCBT SCBT	yes yes	2201805 +++ 2201940 neg	neg neg	+++				conformational		independent Ab
	CD16	2H7	Leica-Microystem		563508 neg	neg	neg				neg		independent Ab
	CD163	GHI/61	SCBT	no	626933 +++	+++	neg				conformational FA blocked		independent Ab
	CD163 CD163	RM3/1 10D6	SCBT Thermo	no yes	626932 +++ 10982556 neg	neg neg	neg +				conformational		independent Ab
	CD1A	010	SCBT	yes	626957 +++	+++	neg				conformational		historic
	CD1c	2F4	Origene	yes	2623067 neg	neg	neg				neg		orthogonal
	CD1c CD1c	2D4 UMAB46	Origene Origene	yes yes	2623049 neg 2629056 neg	neg neg	neg neg				neg		orthogonal peptide microarray, orthogo
	CD10	B-8	SCBT	yes	2029050 neg 2073673 neg	neg	neg				neg		peptide microarray, orthogo
	CD20	L26	Dako	yes	2282030 +++	++++	+++				conserved		historic
	CD206	# 685645	R&D	yes	note	+++	+++	non-specific			FA dependent neg	spurious	
B	CD207 langerin CD21	2G9 + A3	Merck Thermo + SCBT	yes yes	1078467 neg +/-	neg neg	neg +++				FA blocked		historic historic
5	CD22	SP104	Merck	yes	++++	+++	+++				conserved		historio
	CD23	UMAB101	Origene	yes	2629143 neg	neg	++				conformational		
D	CD248	B1/35	BD Pharmingen	yes	11212259 +++	neg +++	++ +++				FA blocked conserved		
	CD27 CD271	EPR8569 NGFR 5	Abcam/Epitomics Neomarkers	yes yes	11155136 + 61258 +++	++++ +	+++ ++				conserved		historic
	CD3	poly	Merck	yes	259074 +	neg	neg				FA blocked		independent Ab, cross-spe
	CD3	CD3-12	SCBT	yes	1120361 +++	+/-	++		and that stores i		FA blocked		independent Ab
b	CD3 CD30	SP7 BerH2	Thermo SCBT	yes	149922 +++ 2335683 +++	note +	+++		endothel. stromal		FA blocked conserved	spurious	independent Ab, historic independent Ab, historic
	CD30 CD30	BerH2 CON6D/ C2	SCBT Thermo	yes yes	2335683 +++ 10988551 +++	+ +/-	+ +++				FA blocked		independent Ab, historic independent Ab
	CD303	AC144	Milteny	no	244165 +++	neg	neg				misc		independent Ab
	CD303	poly	Merck	yes	1078467 neg	neg	neg	non-specific	non-specific	non-specific	neg	spurious	independent Ab
b	CD303 CD30v	124b3.13 E4L4I	Dendritics CST	yes	note 2799462 neg	note	neg	panEpith	GC & DC (?)		neg	spurious	independent Ab independent Ab
-	CD30V CD31	JC70	SCBT	yes yes	2799462 neg 629040 +	neg ++	neg +++				conserved		independent Ab historic
	CD32abc	B-4	SCBT	yes	2103735 neg	note	note		nuclear bkgrnd	diffuse bckground		spurious	
	CD34	43A1	SCBT	yes	631133 +++	+++	+/-				conformational		independent Ab
ıb	CD39 CD4	IMG17B5F11 EPR6855	Abcam/Epitomics Abcam/Epitomics	yes yes	neg +	neg +/-	neg +++				neg conformational		independent Ab
U	CD4 CD4	MT310	SCBT	no	627055 +++	neg	nea				FA blocked		independent Ab
	CD45	Hle-1 2D1	BD Pharmingen	no	400154 +++	+++	neg				conformational		independent Ab, historic
	CD45+43	PD7/26 + 35-Z6`	BioLegend	yes	+++	+++	+++				conserved		independent Ab, historic
	CD45R0 CD5	UCHL-1 CD5/54/F6	SCBT SCBT	yes	627083 +++ 2075324 +/-	++	***				conserved FA blocked		independent Ab, historic
	CD5 CD56	123C3.D5	SCBT	yes yes	2075324 +/- 627127 note	neg note	note	endoth. basal sq	endothelium	endothelium	neg	spurious	cross-species, historic historic
	CD64	OTI3D3	Origene	yes	2623718 neg	note	+++		nucleolar, pancellular		conformational	spurious	independent Ab, historic
	CD68	KP1	SCBT	yes	627158 +++	++	+++				conserved		independent Ab, historic
	CD68	PGM1	Thermo	yes	10979558 neg	+/-	+				FA dependent		independent Ab
	CD69 CD7	poly poly	Merck Merck	yes yes	1078467 neg 1078467 +	neg neg	neg neg				neg misc		orthogonal
	CD74	UMAB231	Origene	yes	+/-	neg	+/-				FA blocked		orthogonal
	CD79a	JCB117	SCBT	yes	1120686 ++	+	+++				conserved		independent Ab, cross-spe
	CD79a CD8	HM47/A9	Merck SCBT	yes	10897816 +++	+++	+++		epith. basal layer, lymphs very weak		conserved conserved	spurious	independent Ab
	CD8 CD83	C8/144B F-5	SCBT SCBT	yes yes	1120718 +++ 831132 neg	note neg	+++ ++		opini, basai idyer, iyinipits very weak		conserved conformational	aparious	independent Ab, historic independent Ab
	CD85	poly	Merck	yes	1078467 neg	neg	neg				neg		
	CD8beta	F5	SCBT	yes	627211 +/-	neg	++				FA blocked		independent Ab
	CDKN1B p27 CLEC10A	DCS-72.F6 poly	Neomarkers Merck	yes yes	61869 +++ 1078467 +/-	+ nea	++ neg	non-specific	non-specific pan epithelial	non-specific pan epithelial	conserved neg	spurious spurious	orthogonal orthogonal
	CLEC10A CLEC9A	poly	Merck	yes ves	1078467 +/- 1078467 neg	neg neg	neg +		* · · · · · · · · · · · · · · · · · · ·	, sprotoliai	conformational	apurious	orthogonal
	CLEC9A	14N8D7	Thermo	no	2537779 neg	neg	neg				neg		orthogonal
	cMAF	poly	Merck	yes	1078467 neg	+/-	+/-				FA dependent		
	cREL CXCL13	poly	Merck Merck	yes	1078467 +/- 1078467 +++	neg +	+ ++				FA blocked conserved		orthogonal independent Ab
	CXCL13 CXCR5	poly #51505	Merck R&D	yes yes	1078467 +++ 2292654 +++	+ note	++ +++		interstitium		FA blocked	spurious	muependent Ab
	EOMES	poly	Merck	yes	1078467 neg	neg	neg				neg		
	FoxP3	236A/E7	Abcam/Epitomics	yes	445284 ++	neg	+++				FA blocked		independent Ab
	GATA3	poly	Merck	yes	1078467 neg	neg	neg				neg misc		bistoria
	GZMB GZMK	GRB7 GM6C3	BioLegend SCBT	yes no	neg +++	+++ ++	neg +++				misc conserved		historic
	Histone H3	1D8	Merck	yes	1841955 +/-	neg	+++				conformational		
	HLA-DR	L243	BD Pharmingen	no	400289 +++	+++	neg				conformational		orthogonal, historic
	HLA-DR	SPM288	SCBT	yes	1125217 +++	+++	+++	pan-cell			conserved neg		orthogonal
5 5	ID1 ID2	BCH-1/195-14 BCH-3/9-2-8	Biocheck Biocheck	yes yes	neg note	neg +++	neg ++	cytoplasmic pos			neg FA dependent	spurious	genetic genetic
5	IDO	D5J4E	CST	yes	2626818 neg	neg	neg				neg		J
	lgD	poly	Merck	yes	1078467 +++	++	+++				conserved		independent Ab
	IRF4	3E4	SCBT	yes	2127138 +/-	+	+/-				conserved FA dependent		independent Ab
	IRF4 IRF8	poly E-9	Merck SCBT	yes yes	1078467 +/- 10850401 +/-	++++ ++++	++++ ++++				FA dependent FA dependent		independent Ab, cross-spe independent Ab
	IRF8	E-9 poly	Merck	yes	1078467 neg	++++	+++				FA dependent		independent Ab, cross-spe
	kappa	poly	Merck	yes	1078467 +/-	+++	+/-				conserved		independent Ab, orthogona
b	Ki-67	SP6	Thermo	yes	2335745 ++	***	***				conserved		independent Ab, historic
	Ki-67 Ki-67	UMAB107 MIB3	OriGene J.Gerdes	yes yes	2629145 +++	+++ +++	+++ +++				conserved conserved		independent Ab independent Ab, historic
	KI-67 LAG3	MIB3 11E3	J.Gerdes Abcam/Epitomics	yes yes	+++ 776102 ++	+++ neg	neg				misc		independent Ab, historic independent Ab
	lambda	poly	Merck	yes	1078467 +/-	++	+++				conserved		independent Ab, orthogona
	LEF1	B-10	SCBT	yes	10991107 +++	neg	++	selected areas			FA blocked		
	LYVE1 LYZ	poly	Merck Merck	yes	1078467 +++ 1078467 +/-	+++ +++	***				conserved FA dependent		historic
	LYZ MX-1	poly	Merck Merck	yes yes	1078467 +/- 1078467 neg	+++ neg	+++ neg				neg		historic orthogonal
	MYC	EP121	Merck	yes	1078407 fileg +++	neg	++				FA blocked		independent Ab
b		E-10	CST	yes	neg	neg	neg				neg		orthogonal
b	NFkB1 p50	C-5	SCBT	yes	+/-	note	+		diffuse bckgnd		FA blocked	spurious	orthogonal
þ	NFkB2 p52		SCBT	yes yes	626897 +++ 2236668 +++	neg +/-	++ +++				FA blocked conserved		genetic, independent Ab independent Ab
b	NFkB2 p52 OX40	Ber-ACT35 1H9		,00		+/- note	+++		panEpith, low on Macs	Similar to Ac	conserved	spurious	peptide microarray, historic
þ	NFkB2 p52	Ber-ACT35 1H9 pool	Merck BioLegend	yes	+++								
b	NFkB2 p52 OX40 Pax5	1H9 pool UMAB197 + UMAB199	BioLegend BioLegend	yes yes	+++	neg	++				FA blocked		
ıb	NFkB2 p52 OX40 Pax5 PD-L1 PD1 pool PNAd	1H9 pool UMAB197 + UMAB199 MECA-79	BioLegend BioLegend SCBT	yes yes	+++ 627143 neg	neg ++	++				FA dependent		peptide microarray, indepen orthogonal
D	NFkB2 p52 OX40 Pax5 PD-L1 PD1 pool PNAd Podoplanin	1H9 pool UMAB197 + UMAB199 MECA-79 NZ-1.2	BioLegend BioLegend SCBT LsBio	yes yes yes	+++ 627143 neg 10920577 note	neg ++ note	++ note	BasalEp stronger			FA dependent conserved		peptide microarray, indepen orthogonal orthogonal
þ	NFkB2 p52 OX40 Pax5 PD-L1 PD1 pool PNAd	1H9 pool UMAB197 + UMAB199 MECA-79	BioLegend BioLegend SCBT	yes yes	+++ 627143 neg	neg ++	++	BasalEp stronger			FA dependent conserved neg FA blocked		peptide microarray, indepen orthogonal
Ь	NFkB2 p52 OX40 Pax5 PD-L1 PD1 pool PNAd Podoplanin PU1	1H9 pool UMAB197 + UMAB199 MECA-79 NZ-1.2 G148-74	BioLegend BioLegend SCBT LsBio BD Pharmingen	yes yes yes yes	+++ 627143 neg 10920577 note 395335 neg	neg ++ note neg	++ note neg	BasalEp stronger			FA dependent conserved neg		peptide microarray, indeper orthogonal orthogonal independent Ab, historic

Species	Ab	clone	Source	FFPE RRI	D avai	laple				FA> AR	Classification	Spurious re	a validation
g2A	SOX9	CL0639	Merck	yes	2665670 neg	neg	+++				conformational		
g1	TBET	O4-46	BD Pharmingen	yes	+/-	+++	+++	heterogeneous			FA dependent		
RbMab	TCF4	NCI-R159-6	Abcam/Epitomics	yes	2714172 +++	++	++				conserved		orthogonal
Gt	TCF7	poly	Merck	yes	1078467 +++	neg	++				FA blocked		
g1	TCRd	H-41	SCBT	yes	1130061 neg	neg	neg				neg		historic
Gt	TIM3	poly	Merck	yes	1078467 +++	+	++				conserved		orthogonal
Rat	TOX1	NAN448A	Abcam/Epitomics	yes	neg	++	+++				FA dependent		genetic
g2B	TP53	DO-7	Dako	yes	neg	neg	+++				conformational		independent Ab
RbMab	VISTA	D1L2G + D5L5T	CST	yes	++	neg	+/-				FA blocked		orthogonal
Gt	vWF	poly	Merck	yes	1078467 +++	+++	+++				conserved		cross-species, historic
Rb	ZEB1	poly	Merck	yes	1078467 +++	note	note		pan nuclear	pan nuclear	misc	spurious	cross-species
g2A	ZEB1	3G6	LsBio	yes	2572969 note	+++	++	non-specific			FA dependent	spurious	cross-species