Supplementary Information for "Systems pathology by multiplexed immunohistochemistry and whole-slide digital image analysis"

Authors

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Supplementary Methods

Immunohistochemistry.

NOTE 1: The full protocol requires a full working day (two days if overnight antibody incubation is included).

NOTE 2: All incubations are performed in a humid chamber.

DAB detection

Instruments

• PT Module, Thermo Fisher Scientific

Reagents

- Xylene, Fluka/FFChemicals
- EtOH, 96%, Altia
- EtOH, 99.8%, Altia
- 10 mM Tris-HCl buffered saline pH 7.4 (TBS), Biotop
- Tris-EDTA buffer (10 mM Tris-HCl, pH 9 + 1 mM EDTA (2-({2-

[Bis(carboxymethyl)amino]ethyl}(carboxymethyl)amino)acetic acid),

made in house

- H₂O₂, 30%, Fluka
- Normal Goat Serum (NGS), Gibco
- Tween-20, Thermo Fisher Scientific
- Mayer's Hematoxylin, Dako
- Pertex-mountant, HistoLab
- Secondary antibody, horseradish peroxidase (HRP)-conjugated goat antirabbit or anti-mouse antibody, Immunologic

- DAB (3,3'-Diaminobenzidine), Immunologic
- MilliQ-H2O, Millipore

Preparation of working solutions

- 1x TBS + 10 % NGS: Mix 5 ml NGS and 5 ml 10x TBS. Fill with water to 50ml. Filter through 0.22μm. (TBS-NGS)
- 1x TBS + Tween20 buffer: Mix 0.5 ml Tween-20 and 100 ml 10x TBS. Fill with water to 1L, mix well (TBST)

Procedure

Manually

- 1. Prior IHC, adhere paraffin sections to glass slides overnight at 37°C
- 2. Deparaffinize sections:
 - i. Xylene 3x5 min
 - ii. 99.8% EtOH 3x1 min
 - iii. 96% EtOH 2x1 min
 - iv. 70% EtOH 1x1 min
 - v. H_2O for 1x1 min

In PT Module

 Antigen retrieval: +99°C, 20 min in Tris-EDTA pH9 (buffer valid for two weeks or for 40 slides)

Manually

- 2. Wash in MilliQ-H_2O, 5 min and in 1x TBS, 5 min
- 3. Blocking peroxidase activity: Incubate 15 min in 0.9% $\rm H_2O_2$ solution in TBS
- 4. Wash 5 min, 1x TBST.

- 5. Protein blocking: Incubate 15 min in TBS-NGS. Tap the solution off the slide before primary antibody (no washing!)
- 6. Primary antibody: Diluted in TBS-NGS. Incubate on slides at room temperature (RT)
- 7. Wash 3 x 5 min, 1x TBST
- 8. Secondary Antibody (ready-to-use), incubate 30 min at RT
- 9. Wash 3 x 5 min, 1x TBST
- 10. DAB stain according to manufacturer's instructions (5 min incubation)
- 11. Wash 5 min, H_2O
- 12. Hematoxylin counterstain: 1 min in 10% Mayers Hematoxylin in H_2O
- 13. Rinse in running tap water, 5 min
- 14. Dehydration
 - i. 70 % EtOH 1x1 min
 - ii. 96 % EtOH 2x1 min
 - iii. 99.8 % EtOH 3x1 min
 - iv. Xylene 3x1 min
- 15. Mount with Pertex and coverslip

Fluorescence detection

Instruments

• PT Module, Thermo Fisher Scientific

Reagents

- Xylene, Fluka/FFChemicals
- EtOH, 96%, Altia
- EtOH, 99.8%, Altia

- 1x Tris-HCl buffered saline pH 7.4 (TBS), Biotop
- Tris-EDTA buffer (10 mM Tris-HCl, pH 9 + 1 mM EDTA (2-({2-[Bis(carboxymethyl)amino]ethyl}(carboxymethyl)amino)acetic acid), made in house
- H₂O₂, 30%, Fluka
- Normal Goat Serum (NGS), Gibco
- Tween-20, Thermo Fisher Scientific
- Hoechst 33342, Sigma-Aldrich
- Prolong Gold mountant, Thermo Fisher Scientific
- Secondary antibody, goat AlexaFluor555-conjugated anti-rabbit or antimouse antibodies, Thermo Fisher Scientific
- MilliQ-H2O, Millipore

Preparation of working solutions

- 1x TBS + 10 % NGS: Mix 5 ml NGS and 5 ml 10x TBS. Fill with water to 50ml. Filter through 0.22μm. (TBS-NGS)
- 1x TBS + Tween20 buffer: Mix 0.5 ml Tween-20 and 100 ml 10x TBS. Fill with water to 1L, mix well (TBST)

Procedure

Manually

- 1. Prior to IHC, adhere paraffin sections to glass slides overnight at 37°C
- 2. Deparaffinize sections:
 - vi. Xylene 3x5 min
 - vii. 99.8% EtOH 3x1 min
 - viii. 96% EtOH 2x1 min

- ix. 70% EtOH 1x1 min
- x. H_2O for 1x1 min

In PT Module

 Antigen retrieval: +99°C, 20 min in Tris-EDTA pH9 (buffer valid for two weeks or for 40 slides)

Manually

- 4. Wash in MilliQ-H $_2$ O, 5 min and in 1x TBS, 5 min
- 5. Blocking peroxidase activity: Incubate 15 min in 0.9% H₂O₂ solution
- 6. Wash 5 min, 1x TBST.
- 7. Protein blocking: Incubate 15 min in TBS-NGS. Tap the solution off the slide before primary antibody (no washing!)
- 8. Primary antibody: Diluted in TBS-NGS. Incubate on slides at room temperature (RT)
- 9. Wash 3 x 5 min, 1x TBST
- 10. Secondary antibody 1:300 in TBS-NGS + 1 μg/ml Hoechst, incubate 30 min at RT (protect from light)
- 11. Wash 3 x 5 min, 1x TBST, rinse 1 min in $\mathrm{H}_{2}\mathrm{O}$
- 12. Tap excess water off the slide and apply Prolong Gold mountant. Apply coverslip.

Multiplexed immunohistochemistry (mIHC)

NOTE 1: The full protocol requires three working days if overnight antibody incubations are included.

NOTE 2: If needed, the slides may be left overnight (+4°C) in protein blocking

buffer at steps 7, 18, or 29.

NOTE 3: All incubations are performed in a humid chamber.

Instruments

• PT Module, Thermo Fisher Scientific

Reagents

- Xylene, Fluka/FFChemicals
- EtOH, 96%, Altia
- EtOH, 99.8%, Altia
- 1x Tris-HCl buffered saline pH 7.4 (TBS), Biotop
- Tris-EDTA buffer (10 mM Tris-HCl, pH 9 + 1 mM EDTA (2-({2-[Bis(carboxymethyl)amino]ethyl}(carboxymethyl)amino)acetic acid), made in house
- H₂O₂, 30%, Fluka
- Normal Goat Serum (NGS), Gibco
- Tween-20, Thermo Fisher Scientific
- Secondary antibody, goat AlexaFluor555 or -647 conjugated anti-rabbit or anti-mouse antibodies, Thermo Fisher Scientific
- Tyramide signal amplification (TSA) kits for AlexaFluore488 and AlexaFluor555, PerkinElmer

- Secondary antibody, horseradish peroxidase (HRP)-conjugated goat antirabbit or anti-mouse antibody, Immunologic
- Secondary antibody, alkaline phosphatase (AP)-conjugated goat antirabbit or anti-mouse antibody, Immunologic
- Secondary antibody, goat AlexaFluor647- and AlexaFluor750-conjugated anti-rabbit or anti-mouse antibodies, Thermo Fisher Scientific
- VinaGreen (VG), Biocare Medical
- Liquid Permanent Red (LPR), Dako
- Mayer's Hematoxylin, Dako
- Hoechst 33342, Sigma-Aldrich
- Prolong Gold mountant, Thermo Fisher Scientific
- Pertex-mountant, HistoLab
- MilliQ-H2O, Millipore
- NaN₃, Sigma-Aldrich

Preparation of working solutions

- 1x TBS + 10 % NGS: Mix 5 ml NGS and 5 ml 10x TBS. Fill with water to 50ml. Filter through 0.22μm. (TBS-NGS)
- 1x TBS + Tween20 buffer: Mix 0.5 ml Tween-20 and 100 ml 10x TBS. Fill with water to 1 L, mix well (TBST)

Procedure

Manually

- 1. Prior to IHC, adhere paraffin sections to glass slides overnight at 37°C
- 2. Deparaffinize sections:

- xi. Xylene 3x5 min
- xii. 99.8% EtOH 3x1 min
- xiii. 96% EtOH 2x1 min
- xiv. 70% EtOH 1x1 min
- $xv. \quad H_2 0 \text{ for } 1x1 \min$

In PT Module

 Antigen retrieval: +99°C, 20 min in Tris-EDTA pH9 (buffer valid for two weeks or for 40 slides)

Manually

- 4. Wash in MilliQ-H₂O, 5 min and in 1x TBS, 5 min
- 5. Blocking peroxidase activity: Incubate 15 min in 0.9% $\rm H_2O_2$ solution in TBS
- 6. Wash 5 min, 1x TBST.
- 7. Protein blocking: Incubate 15 min in TBS-NGS. Tap the solution off the slide before primary antibody (no washing!)
- 8. Primary antibody: Diluted in TBS-NGS. Incubate on slides at room temperature (RT).
- 9. Wash 3 x 5 min, 1x TBST
- 10. HRP-conjugated secondary antibody 1:10 in TBST, incubate 45 min at RT
- 11. Wash 3 x 5 min, 1x TBST
- 12. TSA reaction 1 (and 2):
 - Prepare 150 μl TSA working solution (AlexaFluor488 or 555) 150
 μl per slide according to manufacturer's instructions
 - ii. Incubate exactly 15 min at RT (protect from light)
- 13. Wash 5 min, 1x TBST

- 14. Blocking peroxidase activity: Incubate 15 min in 0.9% H₂O₂ solution in TBS with 0.05% NaN₃ (NaN₃ is toxic, handle carefully!)
- 15. Wash 3 x 5 min, 1x TBST
- 16. REPEAT steps 7-13 for TSA reaction 2, then proceed to step 17
- 17. Antigen retrieval: +99°C, 20 min in Tris-EDTA pH9 (buffer valid for two weeks or for 40 slides)
- 18. Protein blocking: Incubate 15 min in TBS-NGS. Tap the solution off the slide before primary antibody (no washing!)
- 19. Two primary antibodies (rabbit + mouse) diluted in TBS-NGS. Incubate at RT (protect from light)
- 20. Wash 3 x 5 min, 1x TBST
- 21. Secondary antibodies (AlexaFluor647 and AlexaFluor750) mixed and diluted 1:300 in TBS-NGS + 1 μg/ml Hoechst, incubate 30 min at RT (protect from light)
- 22. Wash 3 x 5 min, 1x TBST
- 23. Tap excess water off the slide, mount with Prolong Gold and apply coverslip
- 24. IMAGING: Acquire fluorescence images within 4 hours after coverslipping. Continue to next steps as soon as possible after fluorescence imaging!
- 25. Detach the coverslip by incubating in 1x TBST as long as needed for the coverslip to slide off by gravity (do not force the coverslip off!)
- 26. Antigen retrieval: +99°C, 60 min in Tris-EDTA pH 9 (fresh buffer!)
- 27. Blocking peroxidase activity: Incubate 15 min in 0.9% H₂O₂ solution with 0.05% NaN₃ (NaN₃ is toxic, handle carefully!)

- 28. Wash 3 x 5 min, 1x TBST
- 29. Protein blocking: Incubate 15 min in TBS-NGS. Tap the solution off the slide before primary antibody (no washing!)
- 30. Two primary antibodies (rabbit + mouse) diluted in TBS-NGS. Incubate at RT (protect from light)
- 31. Secondary antibodies AP- and HRP-conjugated mixed 1:1 in TBS-NGS, incubate 30 min at RT (protect from light)
- 32. Wash 3 x 5 min, 1x TBST, rinse in H_2O
- 33. Prepare VG and LPR chromogen working solutions according to

manufacturers instructions

- i. Incubate VG 8 min
- ii. Wash $1 \min in H_2 O$
- iii. Incubate LPR 8 min
- iv. Wash $1 \min in H_2 O$
- 34. Hematoxylin counterstain: 30 sec in 10% Mayers Hematoxylin in H_2O
- 35. Rinse in tap water 3 min
- 36. Let slides dry at RT (do not expose to solvents!)
- 37. Mount with Pertex mountant and coverslip
- 38. IMAGING: Acquire brightfield images

Supplementary Figures



Supplementary Figure S1. Schematic workflow of multiplexed immunohistochemistry protocol. (a) The sectioned tissue is stained for four

targets using two tyramide signal amplification (TSA) reactions and four spectrally distinct fluorochromes with fluorescent nuclear counterstain. (b) The fluorescence image is acquired using high-resolution whole-slide imaging, after which the coverslip is detached. (c) A pair of antibodies is applied and detected using enzyme-linked secondary antibodies with red and green chromogenic substrates and blue counterstain. (d) High-resolution whole-slide transmitted light image is acquired, (e) and data from the fluorescence and the transmitted light images are used whole-slide image analysis. Scale bar 50 µm. Ab, antibody; IHC, immunohistochemistry; TSA, tyramide signal amplification. Relates to Figure 1 and Figure 2.



Supplementary Figure S2. Detailed workflow of the multiplexed detection. Steps 1 and 2: Sequential tyramide signal amplification reactions are performed after detecting the primary antibody with HRP-conjugated secondary antibody. The antibody complexes are denatured after the amplification reactions. Step 3: Two

primary antibodies from different hosts are used for a non-amplified fluorescent detection of two additional targets. The fluorescence image is acquired, and antibodies are denatured by heat. Step 4: Two primary antibodies from different hosts are used for an HRP-amplified chromogenic detection of two targets, after which the brightfield image is acquired. Nuclei are counterstained using Hoechst and haematoxylin. Relates to Figure 1 and 2.



Supplementary Figure S3. Antibody validation. Primary antibodies for (a) CK18, (b) CK8, (c) p63, (d) CK5, (e) E-cadherin, (f) PanCK clone AE1/3 (g) PanCK clone C11, (h) AR, (i) AMACR, (j) FoxP3, (k) CD4, (l) CD8, (m) CD45, (n) Ki67 were tested on lymph node (left panel) and prostate (right panel) FFPE tissues.

Antibody concentrations were the same as optimized for chromogenic or TSA amplified detection (see Table 1). Arrows in (m) indicate CD45+ immune cells in prostate sample.



Supplementary Figure S4. Heat-induced antibody detachment test. (a–d) Cytokeratin 5 (CK5) and (e–h) cytokeratin 8 (CK8) were detected using primary

antibodies and AlexaFluor555- or AlexaFluor647-conjugated secondary antibodies, respectively. The fluorescence images were acquired (a, e) before (=control) and (b, f) after heating the slides in hot Tris-EDTA (pH 9) and (c, g) subsequently in hot Glycine-SDS (pH 2). (d, h) Fluorescence intensity in cell object was measured in each condition. Median and quartiles are plotted with maximum and minimum outliers (indicated as cross) A.U., arbitrary unit. Scale bar 100 μ m. See Table 1 for information on the primary antibodies.







Supplementary Figure S5. Testing of antibodies for heat-induced denaturation. Primary antibodies were tested against (a) E-cadherin, (b) androgen receptor, (c) pan-cytokeratin (Invitrogen), (d) pan-cytokeratin (Abcam), (e) cytokeratin 8, (f) cytokeratin 18, (g) p63, (h) cytokeratin 5, (i) alpha-methylacyl-CoA racemase, (j) Ki67, (k) CD4, (l) FoxP3, (m) CD8, and (n) CD45. The antibodies were tested in two concentrations: Low concentration for amplified detection (TSA or chromogen detection) using HRP-conjugated secondary antibody and high concentration for non-amplified detection using fluorochrome conjugated secondary antibody. Chromogen DAB detection (brown) was used to detect primary antibodies. Any residual DAB signal implies incomplete denaturation of the primary antibody. Fluorescence detection (AlexaFluor555, green) confirmed that staining is adequate for non-amplified detection. (a–j) Prostate cancer and (k–n) lymph node were used as control tissues. N/A, not available. Scale bar 50 µm. See Table 1 for primary antibody details.



Supplementary Figure S6. Multiplex fluorescence signal-to-background analysis using mIHC panels with pairwise detection of (a) FoxP3 (AlexaFluor488/FITC) and CD4 (AlexaFluor555/CY3(1)), (b) CD4 (AlexaFluor555/CY3(2)) and CD45 (AlexaFluor647/CY5(1)) in patient 1, and (c) PanEpi (AlexaFluor647/CY5(2)) and CK5+p63 (AlexaFluor750/CY7) in patient 2. (d) Background-corrected fluorescence intensities were measured pairwise in neighbouring channels for FITC-CY3, CY3-CY5, and CY5-CY7 in two areas (area 1 and area 2), where the antibodies (and fluorochromes) are not co-localizing in the measured pixels (visual assessment). (e) Corresponding signal-to-background ratios (area1/area2 or area2/area1) with indicated exposure times. Scale bar 10 µm. S/B, signal-to-background; ms, millisecond.



Supplementary Figure S7. Image analysis workflow. (a) Whole slide images of all imaged channels were split into 2048x2048 pixel subregions, which were padded to 10240x10240 pixel image to enable segmentation of whole prostatic glands (used later for cell classification). Every marker in the padded image outside the whole slide image was set to zero, and the padded images were set as input for the image analysis pipeline. (b) Cells and glands were segmented from

padded input images. The intensity of each marker is measured inside every segmented cell (segmentation result of cells and glands is shown for visualization purposes only, the subregion is not extracted at this point in the workflow). (c) Every cell in the padded image is classified based on the marker profile and the gland segmentation. The central 2048x2048 pixel subregion of each padded image is extracted with its cell measurements (a cell belongs to the subregion where its centroid is located) and used to stitch the classified whole slide image with its cell measurements. Relates to Figure 1 and 2.

Supplementary tables

Supplementary Table S1. Fluorochromes and fluorescence imaging

specifications. LED light source was used for the excitation of all fluorochromes. ms, millisecond; mW, milliwatt.

Fluorochrome	Filter set	Excitation filter	Dichroic mirror	Emission filter	Excitation light source (power)	Exposure time, immune panel	Exposure time, epithelial panel
Hoechst 33342	DAPI cube (Zeiss Filter Set 02)	G365	FT395	LP420	390/18 (264 mW)	50 ms	50 ms
Tyramide- AlexaFluor488	FITC cube (Zeiss Filter Set 38 HE)	BP470/40 (HE)	FT495 (HE)	BP525/50 (HE)	475/28 (191 mW)	30 ms	20 ms
Tyramide- AlexaFluor555	Cy3 cube (Chroma Technology Corp 49004 ET CY3/R)	ET545/25x	T565lpxr	ET605/70m	542/27 (403 mW)	40 ms	100 ms
AlexaFluor647	Cy5 cube (Chroma Technology Corp 49006 ET CY5)	ET620/60x	T660lpxr	ET700/75m	633/22 (143 mW)	100 ms	50 ms
AlexaFluor750	Cy7 cube (Chroma Technology Corp 49007 ET CY7)	ET710/75x	T760lpxr	ET810/90m	740/40 (140 mW)	300 ms	150 ms

Supplementary Table S2. Normalized expression of markers in different cell classes using immune cell panel and epithelial antibody panel (range 0–1). AMACR, alpha-methylacyl-CoA racemase; AR, androgen receptor; ECad, epithelial cadherin; Pan-Epi, pan-epithelium; SD, standard deviation. Relates to Figures 3 and 4.

Immune cell panel	Epithel cells	ial	Epithelial leukocytes		Stromal leukocytes		Other cells		All cells		Positivity threshold
Marker	Mean	SD	Mean	SD	Mean	SD	Mean	SD	Mean	SD	
Pan-Epi	0.10	0.07	0.13	0.09	0.00	0.00	0.01	0.01	0.06	0.07	Global Otsu
CD45	0.02	0.00	0.07	0.05	0.13	0.06	0.03	0.01	0.06	0.06	Global Otsu
CD8	0.07	0.00	0.08	0.01	0.08	0.01	0.07	0.01	0.08	0.01	0.09
CD4	0.02	0.01	0.05	0.06	0.08	0.08	0.02	0.02	0.04	0.05	0.09
FoxP3	0.01	0.00	0.02	0.04	0.03	0.07	0.01	0.01	0.02	0.04	0.06
Ki67	0.01	0.03	0.01	0.02	0.01	0.02	0.01	0.02	0.01	0.03	0.10
	Cancer cells										
Epithelial cell panel	Cancer	cells	Benign Iumina	l cells	Basal c	ells	Stroma	al cells	All cell	s	Positivity threshold
Epithelial cell panel Marker	Cancer Mean	cells SD	Benign Iumina Mean	l cells SD	Basal c Mean	ells	Stroma Mean	al cells SD	All cell Mean	s SD	Positivity threshold
Epithelial cell panel Marker Pan-Epi	Cancer Mean 0.139	cells SD 0.062	Benign Iumina Mean 0.127	I cells SD 0.072	Basal c Mean	ells SD 0.061	Stroma Mean 0.021	SD 0.027	All cell Mean 0.079	s SD 0.080	Positivity threshold Adaptive Otsu
Epithelial cell panel Marker Pan-Epi CK5 + p63	Cancer Mean 0.139 0.040	cells SD 0.062 0.003	Benign Iumina Mean 0.127 0.048	l cells SD 0.072 0.008	Basal c Mean 0.176 0.104	ells SD 0.061 0.037	Stroma Mean 0.021 0.043	SD 0.027 0.010	All cell Mean 0.079 0.051	s SD 0.080 0.027	Positivity threshold Adaptive Otsu Global Otsu
Epithelial cell panel Marker Pan-Epi CK5 + p63 CK8	Cancer Mean 0.139 0.040 0.081	cells SD 0.062 0.003 0.022	Benign lumina Mean 0.127 0.048 0.071	l cells SD 0.072 0.008 0.023	Basal c Mean 0.176 0.104 0.061	ells SD 0.061 0.037 0.014	Stroma Mean 0.021 0.043 0.052	SD 0.027 0.010 0.013	All cell Mean 0.079 0.051 0.061	s 5D 0.080 0.027 0.021	Positivity threshold Adaptive Otsu Global Otsu 0.082
Epithelial cell panel Marker Pan-Epi CK5 + p63 CK8 CK18	Cancer Mean 0.139 0.040 0.081 0.200	cells SD 0.062 0.003 0.022 0.153	Benign lumina Mean 0.127 0.048 0.071 0.119	SD 0.072 0.008 0.023 0.136	Basal c Mean 0.176 0.104 0.061 0.075	SD 0.061 0.037 0.014 0.089	Stroma Mean 0.021 0.043 0.052 0.017	SD 0.027 0.010 0.013 0.017	All cell Mean 0.079 0.051 0.061 0.073	SD 0.080 0.027 0.021 0.115	Positivity threshold Adaptive Otsu Global Otsu 0.082 0.188
Epithelial cell panel Marker Pan-Epi CK5 + p63 CK8 CK18 AMACR	Cancer Mean 0.139 0.040 0.081 0.200 0.038	cells SD 0.062 0.003 0.022 0.153 0.052	Benign lumina 0.127 0.048 0.071 0.119 0.011	l cells SD 0.072 0.008 0.023 0.136 0.023	Basal c Mean 0.176 0.104 0.061 0.075 0.006	SD 0.061 0.037 0.014 0.089 0.007	Stroma Mean 0.021 0.043 0.052 0.017 0.005	SD 0.027 0.010 0.013 0.017 0.006	All cell Mean 0.079 0.051 0.061 0.073 0.013	s 5D 0.080 0.027 0.021 0.115 0.028	Positivity threshold Adaptive Otsu Global Otsu 0.082 0.188 0.041

Supplementary Table S3. Summary of publications for multiplexed IHC. None of the published methods fulfil all the specifications listed. Relates to Figure 5.

Reference	Publication year	Reference nr	>5-plex assay	Rapid implementation of new targets	High-resolution whole-slide image acquisition	High- resolution whole-slide image analysis	Open-source image analysis software
Camp, R. L., Chung, G. G. & Rimm, D. L.	2002	1	No	Yes	No	No	Yes
Wahlby, C., Erlandsson, F., Bengtsson, E. & Zetterberg, A.	2002	2	Yes	Yes	No	No	Yes
Xing, Y. et al.	2007	13	No	Yes	No	No	Yes
Mansfield, J. R.	2010	14	No	Yes	No	No	No
Peng, CW. et al.	2011	4	No	Yes	No	No	No
Gerdes, M. J. <i>et al.</i>	2013	5	Yes	No	Yes	No	Yes
van der Loos, C. M. <i>et al.</i>	2013	15	No	Yes	No	No	No
Angelo, M. <i>et al.</i>	2014	10	Yes	No	No	No	Yes
Brown, J. R. <i>et al.</i>	2014	6	No	Yes	No	No	No
Giesen, C. <i>et</i> al.	2014	11	Yes	No	No	No	Yes
Shipitsin, M. <i>et al.</i>	2014	7	No	Yes	No	No	No
Lin, JR., Fallahi- Sichani, M. & Sorger, P. K.	2015	17	Yes	No	No	No	Yes
Feng, Z. <i>et</i> al.	2015	16	Yes	Yes	No	No	No
Carstens, J. L. et al.	2017	9	Yes	Yes	No	No	No

Supplementary data on publications for antibodies used in this study. IHC, immunohistochemistry; ICC, immunocytochemistry; IF, immunofluorescence; LMA, lysate microarray.

CK18 (SantaCruz 6259, clone DC10)

Original publication

Lauerová L, Kovarik J, Bártek J, Rejthar A, Vojtěsek B. Novel monoclonal antibodies defining epitope of human cytokeratin 18 molecule. Hybridoma. 1988 Oct;7(5):495-504. PubMed PMID: 2461901.

IHC

Jiang LW, Chen H, Lu H. Using human epithelial amnion cells in human deepidermized dermis for skin regeneration. J Dermatol Sci. 2016 Jan;81(1):26-34. doi: 10.1016/j.jdermsci.2015.10.018. Epub 2015 Oct 31. PubMed PMID: 26596214.

ICC/IHC

Freyer N, Knöspel F, Strahl N, Amini L, Schrade P, Bachmann S, Damm G, Seehofer D, Jacobs F, Monshouwer M, Zeilinger K. Hepatic Differentiation of Human
Induced Pluripotent Stem Cells in a Perfused Three-Dimensional
Multicompartment Bioreactor. Biores Open Access. 2016 Aug 1;5(1):235-48. doi:
10.1089/biores.2016.0027. eCollection 2016. PubMed PMID: 27610270;
PubMed Central PMCID: PMC5003005.

CK8 (Invitrogen 18-0185, clone 5D3)

IHC

Duret C, Gerbal-Chaloin S, Ramos J, Fabre JM, Jacquet E, Navarro F, Blanc P, Sa-Cunha A, Maurel P, Daujat-Chavanieu M. Isolation, characterization, and differentiation to hepatocyte-like cells of nonparenchymal epithelial cells from adult human liver. Stem Cells. 2007 Jul;25(7):1779-90. Epub 2007 Apr 5. PubMed PMID: 17412893.

p63 (Abcam 124762, clone EPR5701)

IHC

Smirnova NF, Schamberger AC, Nayakanti S, Hatz R, Behr J, Eickelberg O. Detection and quantification of epithelial progenitor cell populations in human healthy and IPF lungs. Respir Res. 2016 Jul 16;17(1):83. doi: 10.1186/s12931-016-0404-x. PubMed PMID: 27423691; PubMed Central PMCID: PMC4947297.

CK5 (Abcam 52635, clone EP1601Y

IHC

Volkmer JP, Sahoo D, Chin RK, Ho PL, Tang C, Kurtova AV, Willingham SB,

Pazhanisamy SK, Contreras-Trujillo H, Storm TA, Lotan Y, Beck AH, Chung BI,

Alizadeh AA, Godoy G, Lerner SP, van de Rijn M, Shortliffe LD, Weissman IL, Chan
KS. Three differentiation states risk-stratify bladder cancer into distinct
subtypes. Proc Natl Acad Sci U S A. 2012 Feb 7;109(6):2078-83. doi:
10.1073/pnas.1120605109. Epub 2012 Jan 19. Erratum in: Proc Natl Acad Sci U
S A. 2012 Feb 28;109(9):3600. PubMed PMID: 22308455; PubMed Central
PMCID: PMC3277552.

E-cadherin (BD Biosciences 610182, clone 36)

Western blot / IF

Weng Z, Xin M, Pablo L, Grueneberg D, Hagel M, Bain G, Müller T, Papkoff J. Protection against anoikis and down-regulation of cadherin expression by a regulatable beta-catenin protein. J Biol Chem. 2002 May 24;277(21):18677-86. Epub 2002 Mar 19. PubMed PMID: 11904289.

IHC

Krebs AM, Mitschke J, Losada ML, Schmalhofer O, Boerries M, Busch H, Boettcher M, Mougiakakos D, Reichardt W, Bronsert P, Brunton VG, Pilarsky C, Winkler TH, Brabletz S, Stemmler MP, Brabletz T. The EMT-activator Zeb1 is a key factor for cell plasticity and promotes metastasis in pancreatic cancer. Nat Cell Biol. 2017 Apr 17. doi: 10.1038/ncb3513. [Epub ahead of print] PubMed PMID: 28414315.

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35

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CD4 (Abcam 133616, clone EPR6855)

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No publications.

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39