

Supplementary Materials

Quantitative diagnosis of breast tumors by morphometric classification of microenvironmental myoepithelial cells using a machine learning approach

Yoichiro Yamamoto, § 1,2,3,4 Akira Saito, § 4,5,6 Ayako Tateishi,¹ Hisashi Shimojo,¹ Hiroyuki Kanno,¹ Shinichi Tsuchiya,⁷

Ken-ichi Ito,⁸ Eric Cosatto,⁹ Hans Peter Graf,⁹ Rodrigo R. Moraleda,^{10,11} Roland Eils^{2,3} and Niels Grabe^{12,13}

§ Both authors contributed equally to this work

¹ Department of Pathology, Shinshu University School of Medicine, Nagano, Japan.

² Division of Theoretical Bioinformatics, German Cancer Research Center (DKFZ), Heidelberg, Germany.

³ Department for Bioinformatics and Functional Genomics, Institute of Pharmacy and Molecular Biotechnology (IPMB) and Bioquant, University of Heidelberg, Heidelberg, Germany.

⁴ RIKEN Center for Advanced Intelligence Project, Pathology Informatics Unit, Tokyo, Japan.

⁵ Quantitative Pathology & Immunology, Tokyo Medical University, Shinjuku, Tokyo, Japan.

⁶ Department of Molecular Pathology, Tokyo Medical University, Shinjuku, Tokyo, Japan.

⁷ Diagnostic Pathology, Ritsuzankai Iida Hospital, Nagano, Japan.

⁸ Division of Breast and Endocrine Surgery, Shinshu University School of Medicine, Nagano, Japan.

⁹ Department of Machine Learning, NEC Laboratories America, NJ, USA.

¹⁰ Applied Tumor Immunity Clinical Cooperation Unit, National Center for Tumor Diseases, German Cancer Research Center, Heidelberg, Germany.

¹¹ Department of informatics, Technical University Federico Santa Maria Valparaiso, Chile.

¹² Department of Medical Oncology, National Center for Tumor Diseases, University of Heidelberg, Heidelberg, Germany.

¹³ Hamamatsu Tissue Imaging and Analysis Center, Bioquant, University of Heidelberg, Heidelberg, Germany.

Table S1. Characteristics of objects.

Total number of nuclei: 11661

	Cells (p63)	Cells (HE)	ROIs (p63)	ROIs (HE)	Cases	Age
Normal	2749	1289	14	14	7	65.4±13.9
UDH	2307	1073	16	16	5	56.6±7.6
LG-DCIS	1819	666	21	21	5	65.0±15.6
HG-DCIS	1203	555	19	19	5	63.8±11.6

(Mean±S.D.)

UDH: Usual ductal hyperplasia

LG-DCIS: Low Grade Ductal carcinoma in situ

HG-DCIS: High Grade Ductal carcinoma in situ

Table S2. Analyzed ROIs

	Total Cases	ROIs						
		Case1	Case2	Case3	Case4	Case5	Case6	Case7
Normal	7	2	2	2	2	2	2	2
UDH	5	4	2	3	3	4		
DCIS G1	5	4	7	4	4	2		
DCIS G3	5	4	3	5	3	4		

Table S3. Morphological features**Area & Shape**

Area	Compactness	Eccentricity	Form Factor	Major Axis Length
Maximum Feret Diameter	Maximum Radius	Mean Radius	Median Radius	Minimum Feret Diameter
Minor Axis Length	Perimeter	Solidity	Ratio of Feret Diameter	Ratio of Axis Length
FracAtD 1of4*	FracAtD 2of4*	FracAtD 3of4*	FracAtD 4of4*	

Nuclear texture (GLCM)

Angular Second Moment	Contrast	Correlation	Difference Entropy	Difference Variance
Entropy	Info. Measure of Correlation1	Info. Measure of Correlation2	Inverse Difference Moment	Sum Average
Sum Entropy	Sum Variance	Variance		

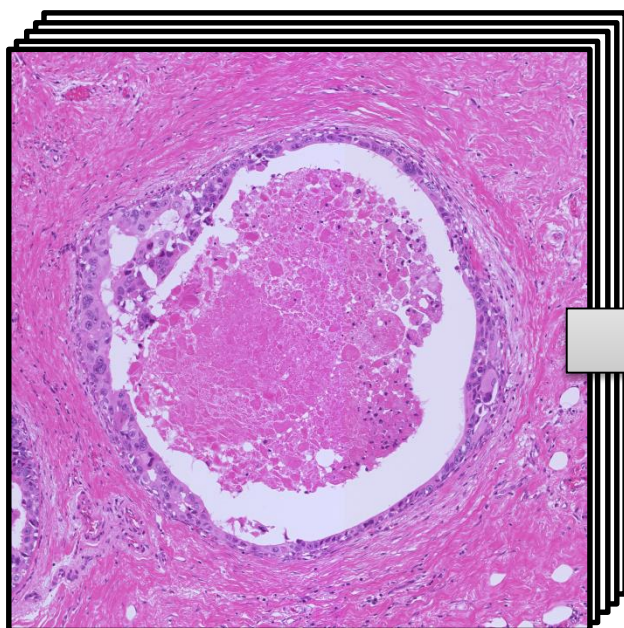
* FracAtD: Fraction of total stain in an object at a given radius. The distribution is measured from the center of the object, where the center is defined as the point farthest from any edge. The numbering is from 1 (innermost) to 4 (outermost).

Table S4. Possible candidate molecules of crosstalk between luminal cells and myoepithelial cells.

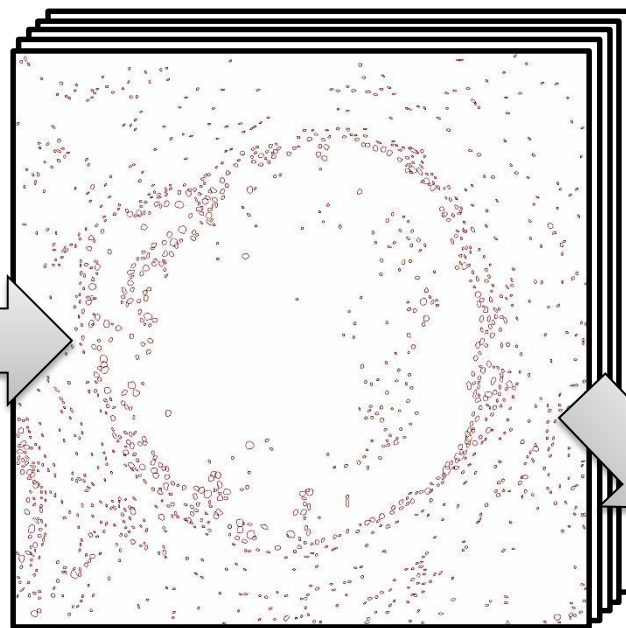
Ligand (Luminal cell)	Receptor (Myoepithelial cell)
ADAM15, ADAM17, COL18A1	ITGB1
CDH1	ITGB7
DUSP18	ITGA3
DUSP18	ITGA7
FN1	SDC2
IL15	IL15RA
SLIT2	ROBO1
SHH	SMO
WNT7B	FZD1

Figure S1

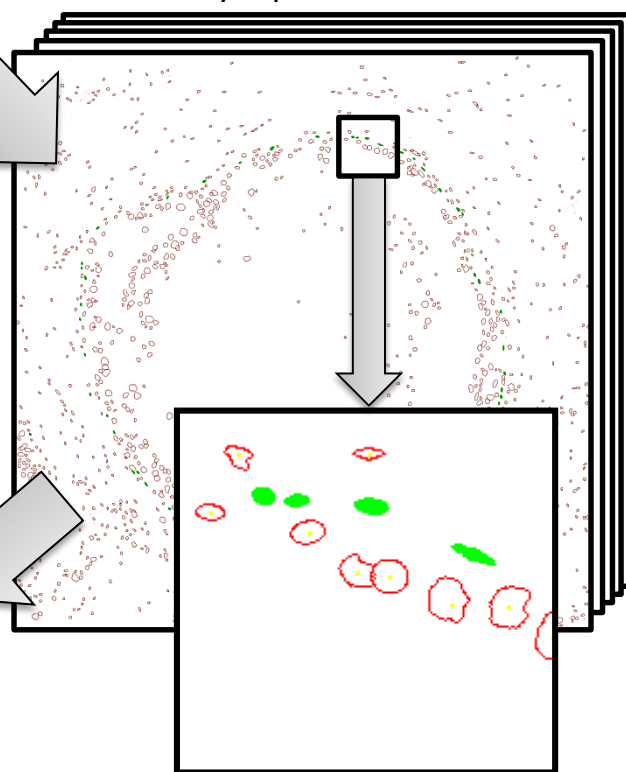
1. Raw data (Serial sections of p63 IHC slides)



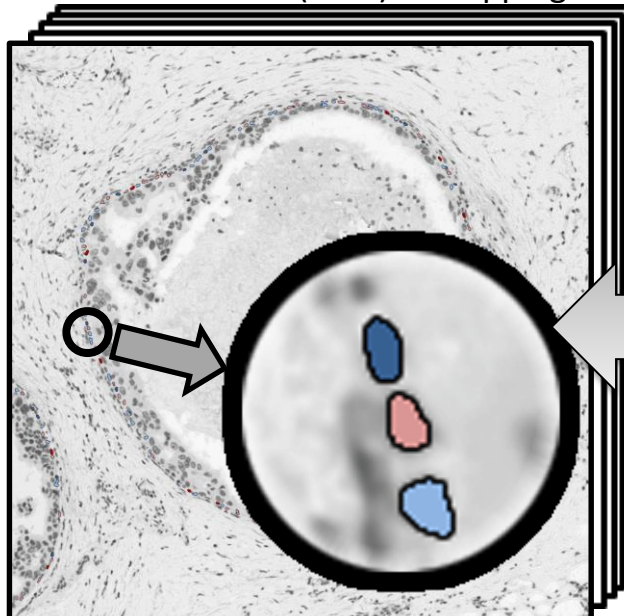
2. Segmentation of all cells by program of NEC



3. Manually picking up myoepithelial cells



5. Classification (SVM) & Mapping



4. Measurement by CellProfiler

No.	Area	Compactness	Eccentricity	Extent	FormFactor	MajorAxis	MaxFeretD
1	153	1.774397	0.948938	0.68	0.537721	25.12586	24.18877
2	156	1.264323	0.86517	0.619048	0.738498	20.09701	19.72308
3	134	1.708976	0.911633	0.553719	0.389348	22.39436	21.0238
4	147	1.648768	0.932858	0.480392	0.566413	23.42876	23.34524
5	126	1.441525	0.913324	0.7875	0.660322	19.98287	19.41649
6	87	1.542556	0.914052	0.639706	0.59216	17.20261	16.12452
7	85	1.681075	0.90786	0.664063	0.56573	17.66234	15.52417
8	142	1.623574	0.91986	0.44375	0.379272	22.61481	21.47091
9	86	1.661746	0.933837	0.632353	0.55903	18.03145	16.12452
10	94	2.959781	0.980617	0.29375	0.404999	26.16462	23.6008
11	107	2.002178	0.960625	0.607955	0.453488	22.56063	21.37758
12	134	2.143986	0.968961	0.408061	0.487632	26.256	24.6981
13	129						24.1660
14	129						21.3775
15	227						24.6981
16	81						22.47221
17	89						17.02939
18	128						23.85372
19	119						22.36068
20	162						17.80449
21	87						17.26268
22	152						17.80449
23	102						18.78829
24	132						16.12452
25	138						18.60108
26	87	0.988378	0.930842	0.621923	0.521284	20.20003	19.23538
27	90	2.050249	0.962401	0.592105	0.553329	20.97888	18.68154
28	107	1.619323	0.936211	0.509524	0.627779	19.88177	19.10497
29	128	1.47595	0.904151	0.547009	0.625903	20.23429	18.86796
30	110	1.674226	0.918816	0.482456	0.505745	20.20779	18.24829
31	179	1.059069	0.614502	0.699219	0.775235	17.31132	17.20465
32	192	2.348707	0.971567	0.347988	0.44822	27.37905	24.83948
33	86	1.187822	0.816857	0.735043	0.766295	14.04825	13.41841
34	86	1.132375	0.730476	0.735043	0.763446	13.08082	12.16553
35	159	1.178992	0.827222	0.623529	0.801271	19.11294	17.80449
36	94	1.248771	0.851312	0.696296	0.745863	15.38524	14.86607
37	156	1.264323	0.86517	0.619048	0.738498	20.09701	19.72308
38	134	1.708976	0.911633	0.553719	0.389348	22.39436	21.0238
39	147	1.648768	0.932858	0.480392	0.566413	23.42876	23.34524



Figure S2

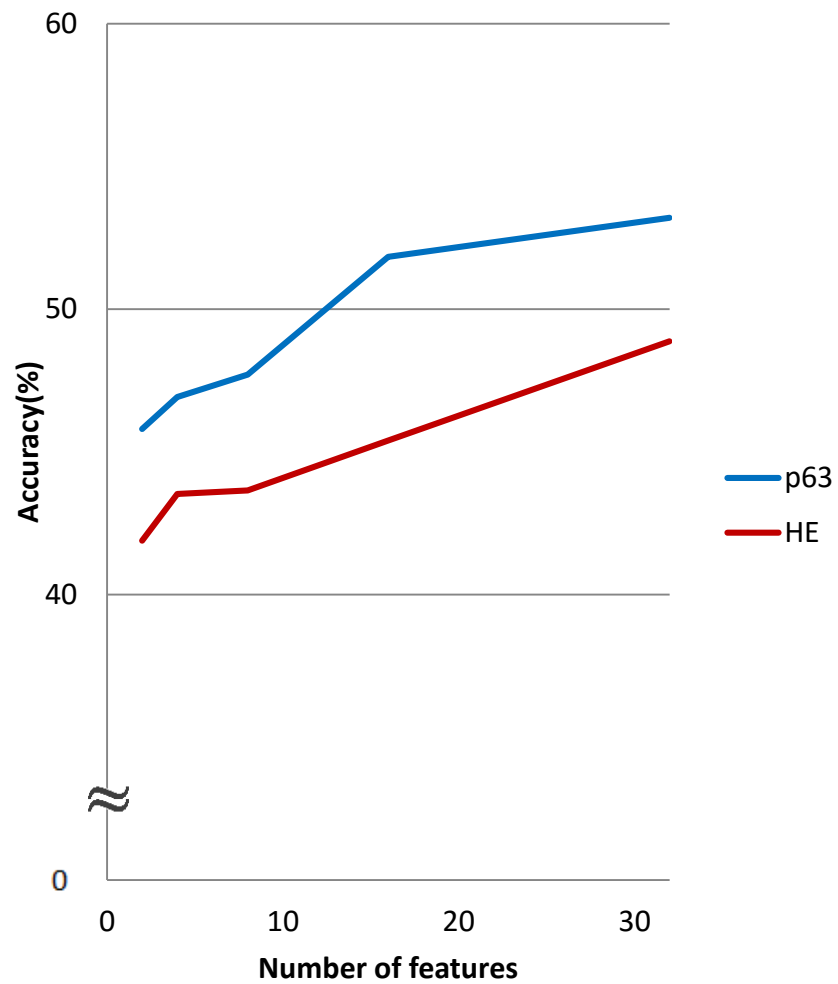


Figure S3

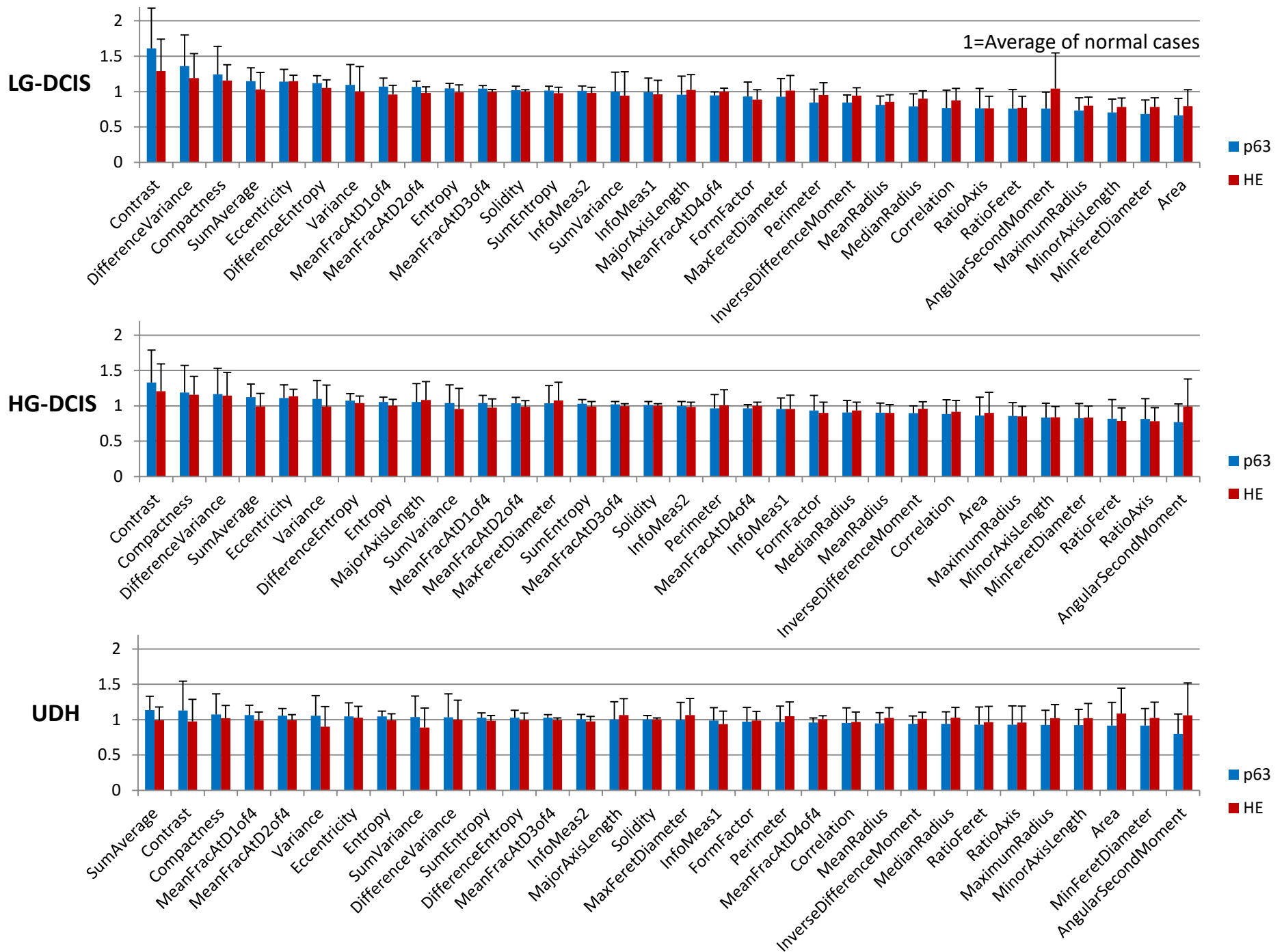
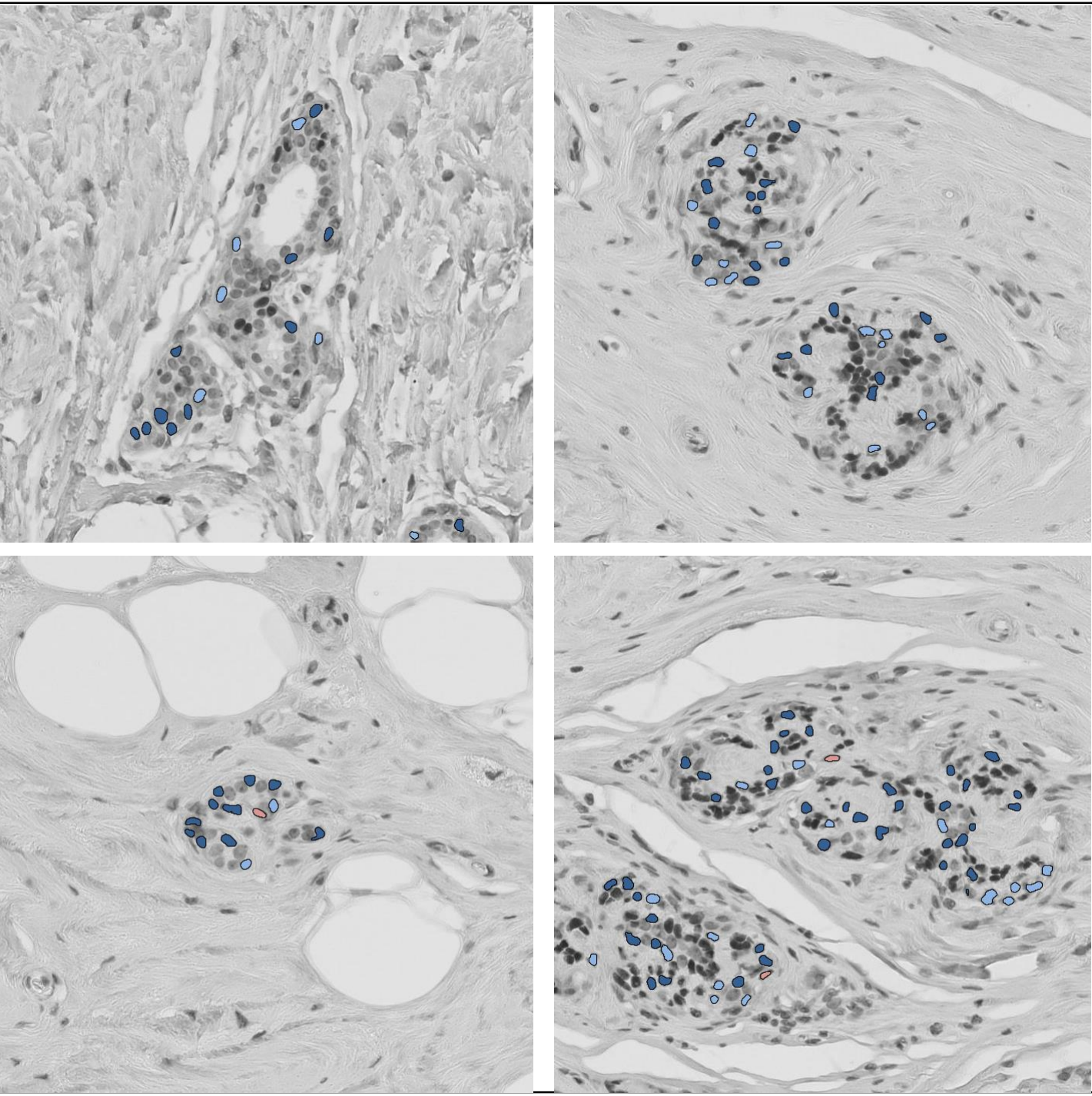
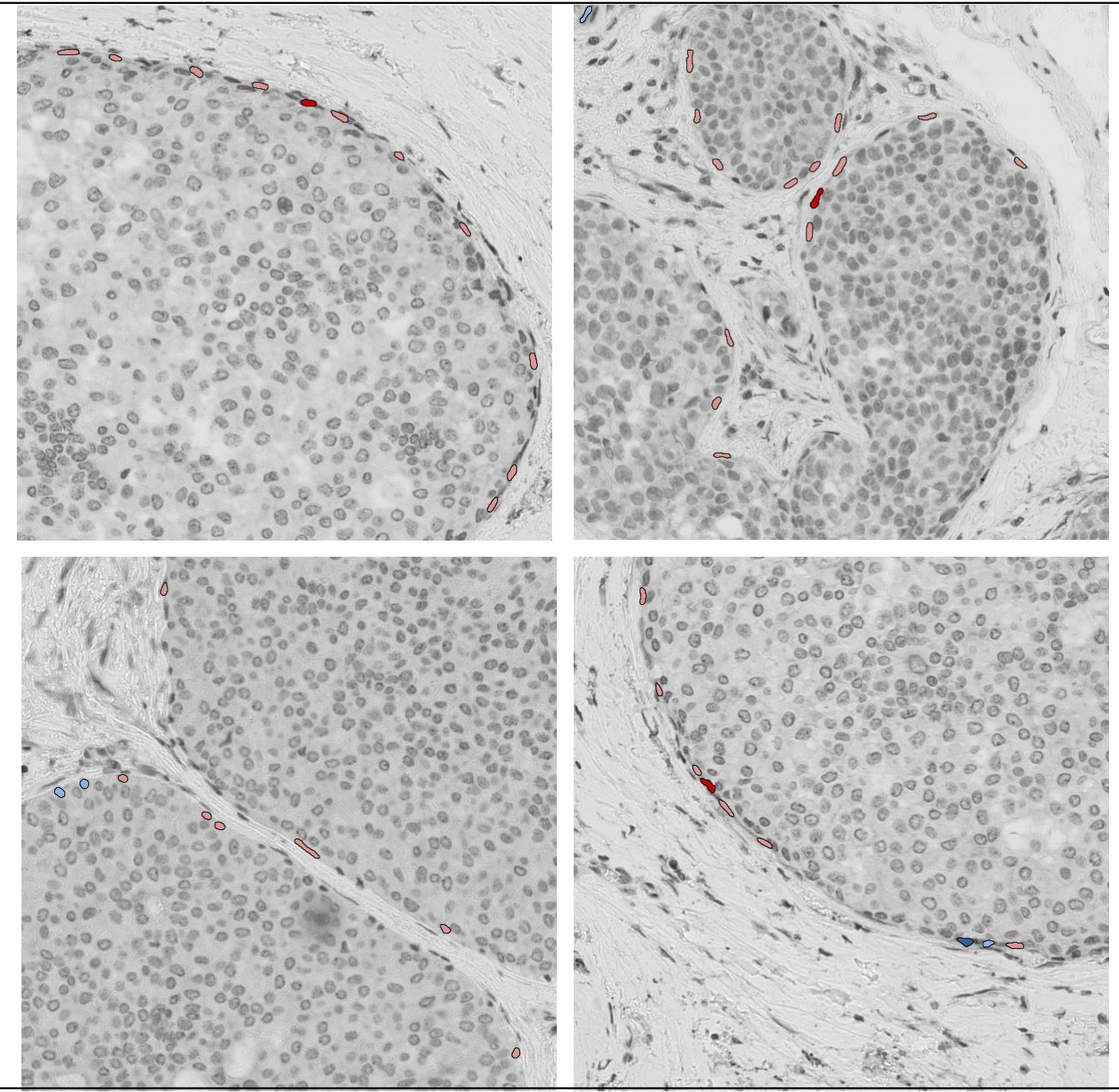


Figure S4

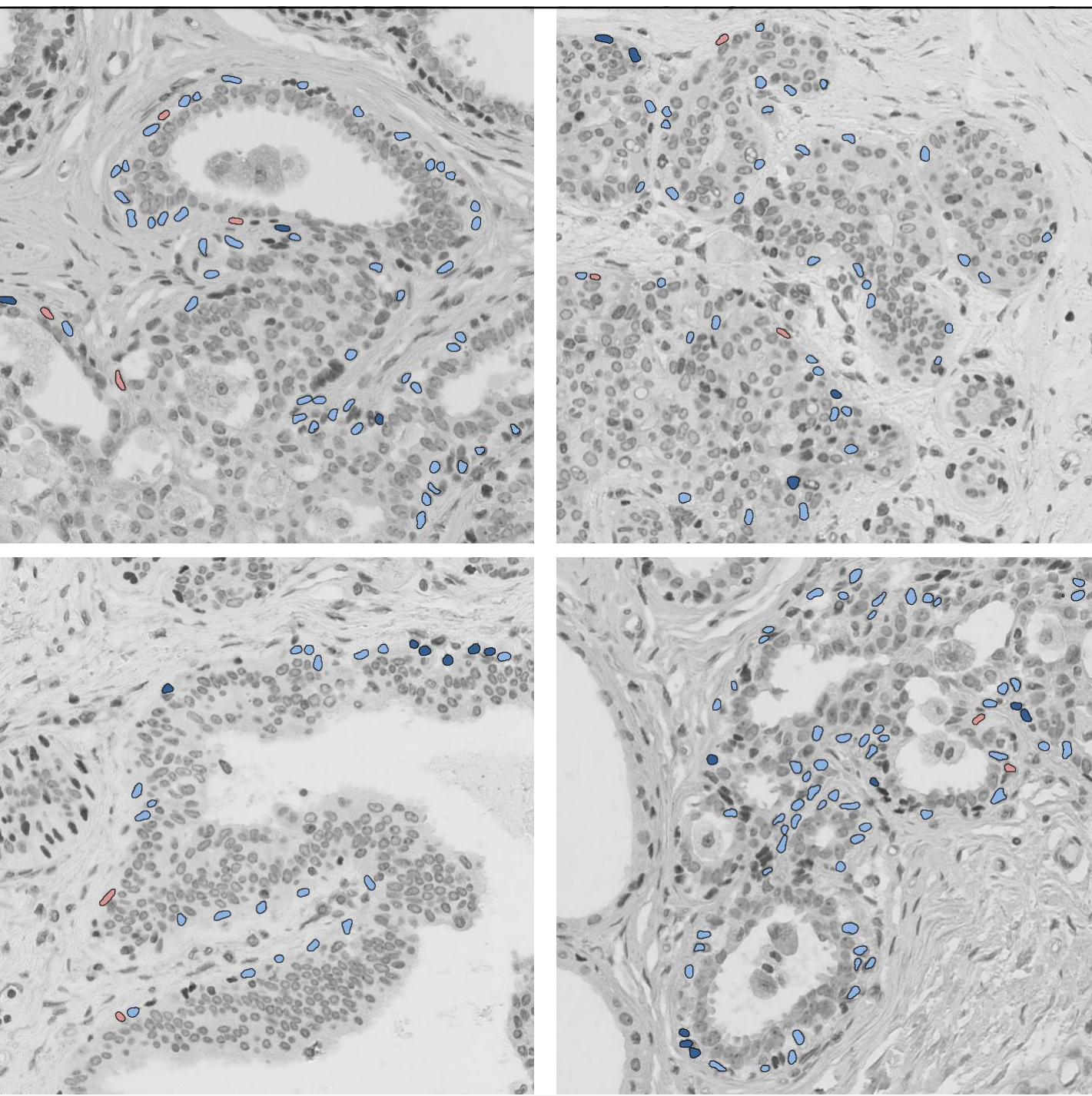
Normal



LG-DCIS



UDH



HG-DCIS

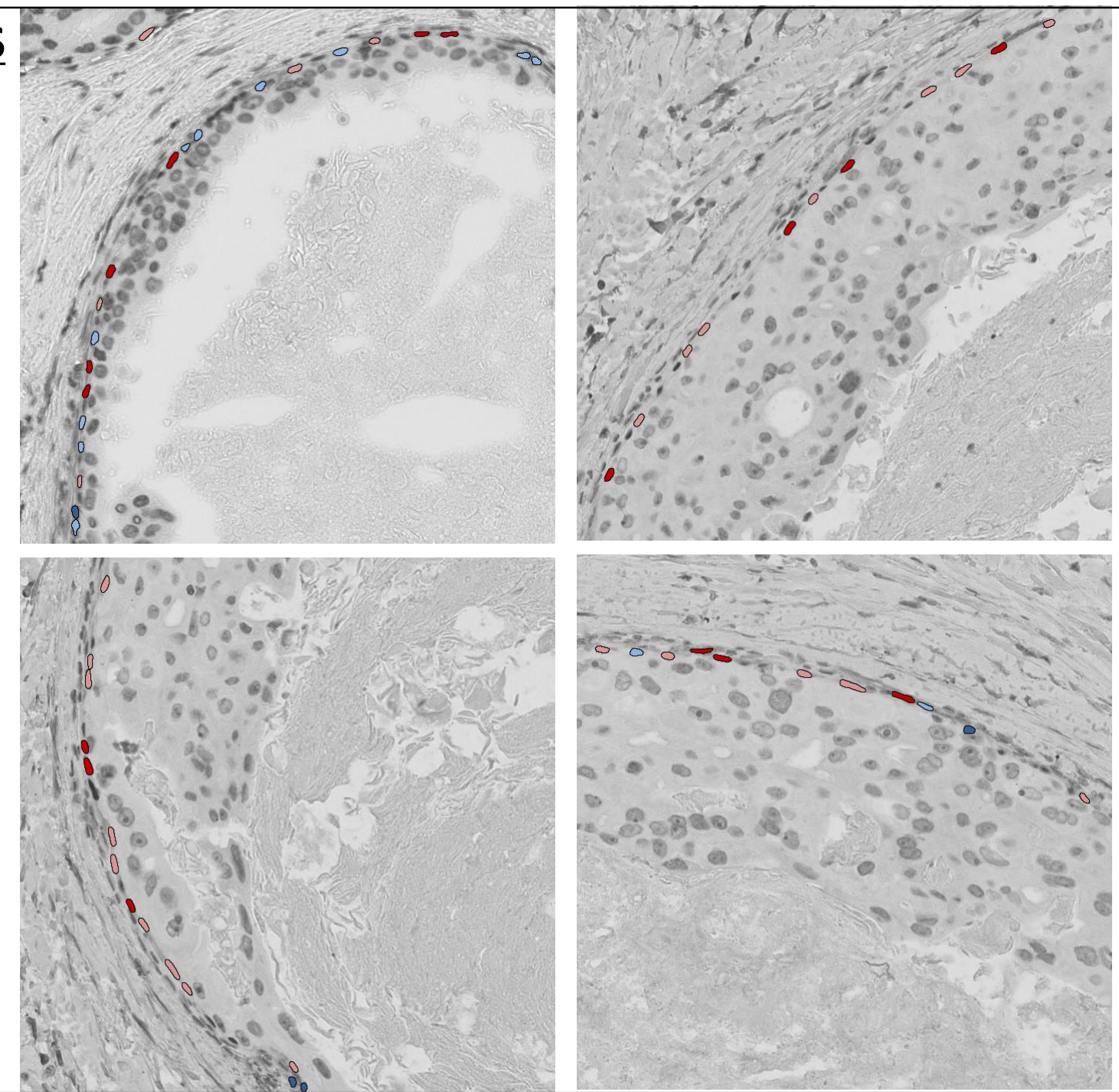
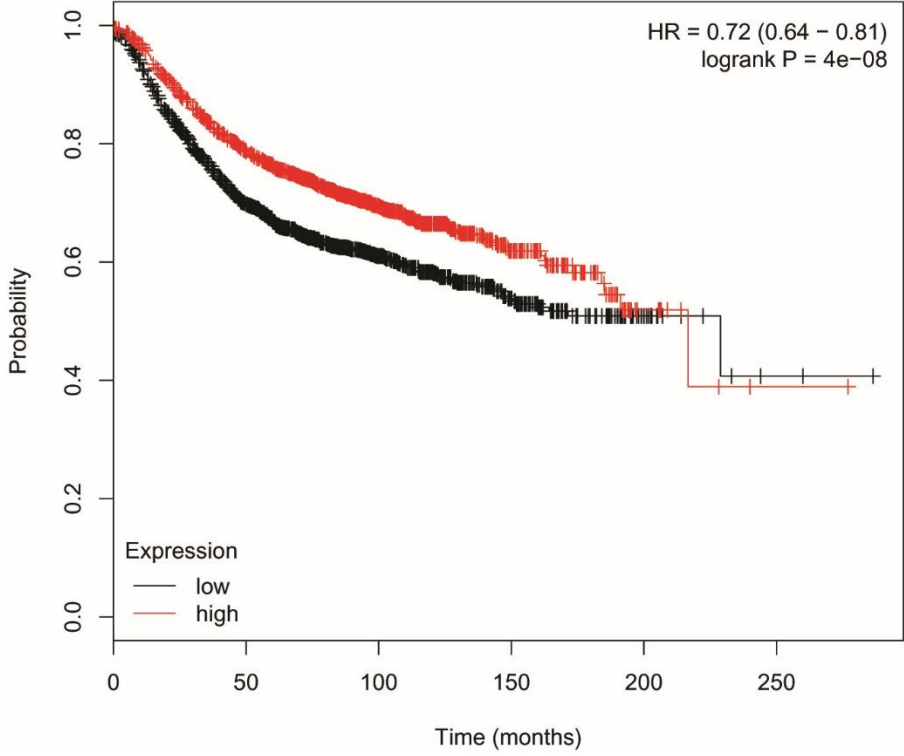


Figure S5

A

SHH



B

SLIT2

