

## Supplementary Information

### Profiling surface glycans on live cells and tissues using quantum dot-lectin nanoconjugates

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**Table S1.** List of lectins used in this study

Abbreviation	Origin	Glycan Specificity
SBA	<i>Glycine max</i>	N-acetylgalactosamine (N-GalNAc), galactose
ConA	<i>Canavalia ensiformis</i>	$\alpha$ -mannose
WGA	<i>Triticum unlgari</i>	N-acetylglucosamine (N-GlcNAc)
LTL	<i>Lotus tetragonolobus</i>	$\alpha$ -linked L-fucose, Sia-Lex, Lex
SNA	<i>Sambucus nigra</i>	Sia( $\alpha$ -2,6)Gal/ GalNAc
MAA	<i>Maackia amurensis</i>	Sia( $\alpha$ -2,3)Gal

**Table S2.** List of cell line.

Designations	Origin	Morphology	Organ	Disease
MCF-7	Human	Epithelial	Breast	Adenocarcinoma
HEK293	Human	Epithelial	Kidney	None
ACHN	Human	Epithelial	Kidney	Renal cell adenocarcinoma
A549	Human	Epithelial	Lung	Carcinoma
HeLa	Human	Epithelial	Cervix	Adenocarcinoma
HCT116	Human	Epithelial	Colon	Colorectal carcinoma
HepG2	Human	Epithelial	Liver	Hepatocellular carcinoma
HDF	Human	Fibroblast-like	Epithermis	None

**Table S3.** Information of breast cancer tissue specimen

No.	Age	Sex	Organ	Diagnosis	pTNM	Stage
1	37	F	Breast	Infiltrating duct carcinoma	T2N0M0	II A
2	48	F	Breast	Infiltrating duct carcinoma	T2N3aM0	III C
3	36	F	Breast	Infiltrating duct carcinoma	T3N0M0	II B
4	40	F	Breast	Infiltrating duct carcinoma	T2N0M0	II A
5	51	F	Breast	Infiltrating duct carcinoma	T2N0M0	II A
6	54	F	Breast	Infiltrating duct carcinoma	T2N0M0	II A
7	43	F	Breast	Infiltrating duct carcinoma	T2N0M0	II A
8	58	F	Breast	Infiltrating duct carcinoma	T2N3aM0	III C
9	37	F	Breast	Infiltrating duct carcinoma	T2N3aM0	III C

**Table S4.** The stage of breast cancer tissue

Stage	T	N	M
II A	T0-T2	N0	M0
II B	T2	N1	M0
	T3	N0	M0
III C	Any T	N3	M0

**T : Primary tumor stage**

**T0** : No evidence of primary tumor

**T1** : Tumor  $\leq 20$  mm in greatest dimension

**T2** : Tumor  $\geq 20$  mm but  $\leq 50$  mm in greatest dimension

**T3** : Tumor  $> 50$  mm in greatest dimension

**N : Regional lymph nodes stage**

**N0** : No regional lymph node metastasis identified histologically.

**N1** : Micrometastases; or metastases in 1-3 axillary lymph nodes; and/or in internal mammary nodes with metastases detected by sentinel lymph node biopsy but not clinically detected.

**N2** : Metastasis in 4-9 axillary nodes; or in clinically detected internal mammary lymph nodes in the absence of axillary lymph node metastasis.

**N3** : Metastasis in 10 or more axillary lymph nodes, or in infraclavicular (level III axillary) lymph nodes, or in clinically detected ipsilateral internal mammary lymph nodes in the presence of 1 or more positive level I, II axillary lymph nodes; or in more than 3 axillary lymph nodes and in internal mammary lymph nodes with micrometastases or macrometastases detected by sentinel lymph node biopsy but not clinically detected/or in ipsilateral supraclavicular lymph nodes

**N3a** : Metastasis in 10 or more axillary lymph nodes (at least one tumor deposit greater than 2.0 mm); or metastases to the infraclavicular (level III axillary) node.

**M : Distant metastasis stage**

**M0** : No clinical or radiographic evidence of distant metastases.

**M1** : Distant detectable metastases as determined by classic clinical and radiographic means and/or histologically proven larger than 0.2 mm.

**Table S5.** Information of colon cancer tissue

No.	Age	Sex	Organ	Diagnosis	pTNM	Stage
1	44	M	Rectum	Adenocarcinoma, moderately differentiated	T3N2aM1a	IV A
2	47	F	Sigmoid colon	Adenocarcinoma, moderately differentiated	T3N1bM1b	IV B
3	64	M	Sigmoid colon	Adenocarcinoma, well differentiated	T3N1aM0	III B
4	67	M	Colon	Mucinous adenocarcinoma	T2N2bM0	III C
5	60	M	Rectum	Adenocarcinoma, moderately differentiated	T3N0M0	II A
6	59	M	Colon	Adenocarcinoma, well differentiated	T3N0M0	II A
7	58	F	Rectum	Adenocarcinoma, moderately differentiated	T3N2bM0	III C
8	64	M	Rectum	Adenocarcinoma, moderately differentiated	T3N2aM0	III B
9	68	M	Colon	Mucinous carcinoma	T4aN1bM0	III B

**Table S6.** The stage of colon cancer tissue

Stage	T	N	M
II A	T3	N0	M0
III B	T3-T4a	N1/N1c	M0
	T2-T3	N2a	
	T1-T2	N2b	
III C	T4a	N2a	M0
	T3-T4a	N2b	
IV A	T4b	N1-N2	M1a
	Any T	Any N	
IV B	Any T	Any N	M1b

**T : Primary tumor stage**

**T0** : Primary tumor cannot be assessed

**T1** : Tumor invades submucosa

**T2** : Tumor invades muscularis propria

**T3** : Tumor invades through the muscularis propria into pericolorectal tissues

**T4a** : Tumor penetrates to the surface of the visceralperitoneum

**T4b** : Tumor directly invades or is adherent to other organs or structures

**N : Regional lymph nodes stage**

**N0** : No regional lymph node metastasis

**N1** : Metastasis in 1-3 regional lymph nodes

**N1a** : Metastasis in one regional lymph node

**N1b** : Metastasis in 2-3 regional lymph nodes

**N2** : Metastasis in four or more regional lymph nodes

**N2a** : Metastasis in 4-6 regional lymph nodes

**N2b** : Metastasis in seven or more regional lymph nodes

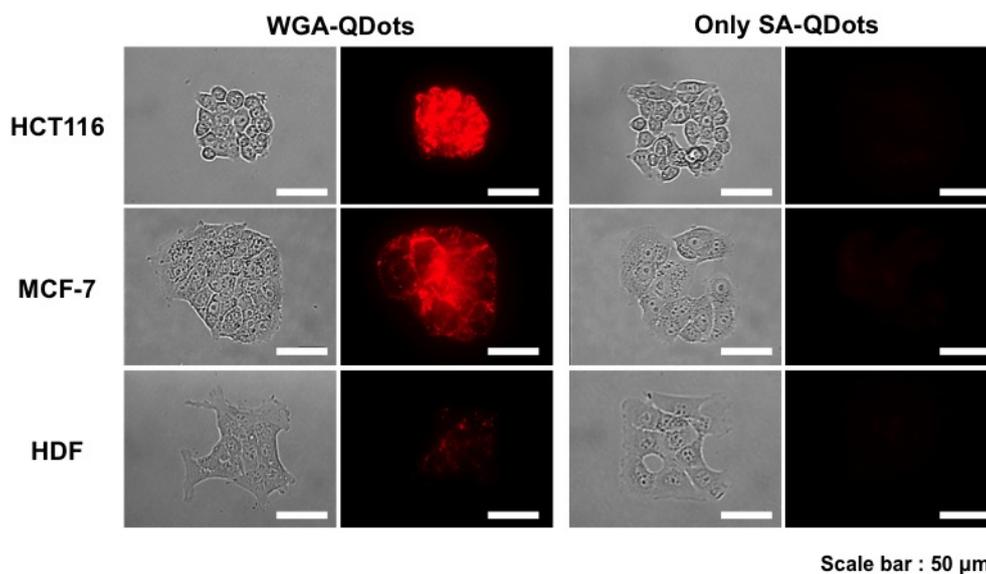
**M** : Distant metastasis stage

**M0** : No distant metastasis

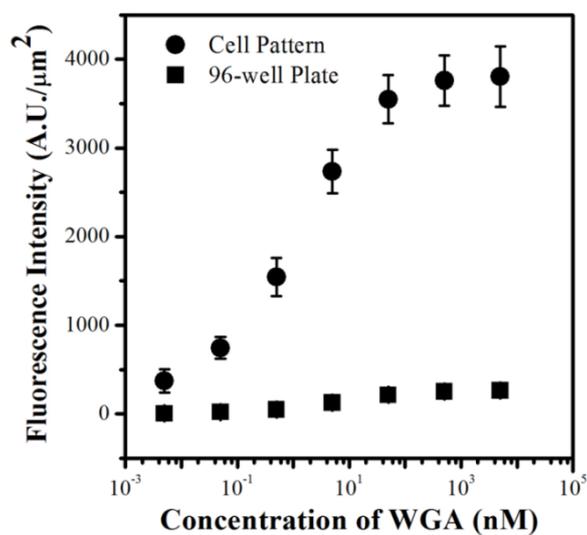
**M1** : Distant metastasis

**M1a** : Metastasis confined to one organ or site (e.g., liver, lung, ovary, nonregional node)

**M1b** : Metastases in more than one organ/site or the peritoneum

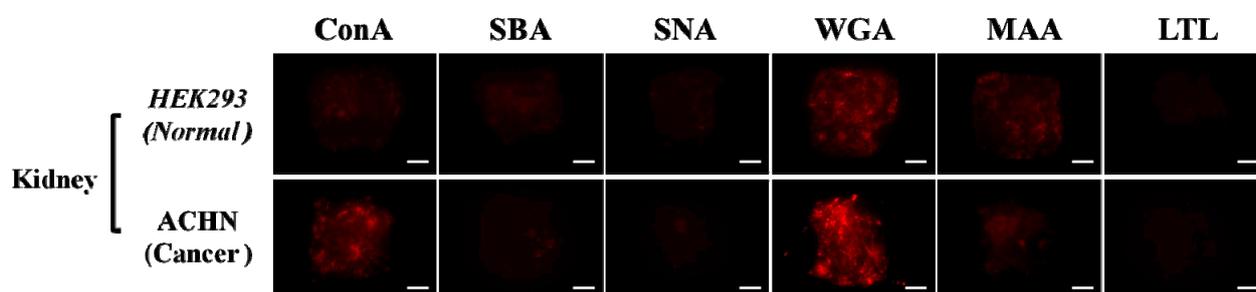


**Figure S1.** The investigation of non-specific binding with streptavidin-QDots (SA-Qdots) and comparison of fluorescence images between WGA-Qdots and only SA-Qdots..



**Figure S2.** Comparison of cell pattern approach with conventional 96-well plate method. MCF-7 are incubated with various concentration of WGA.

(a)



(b)



**Figure S3.** Binding profile of kidney cancer cell (ACHN) and normal kidney cell (HEK293) to 6 lectins. One representative cell pattern for each lectin and cell line is selected. (a) Representative fluorescence images of HEK293 and ACHN cell line. Scale bars indicate 40 μm. (b) Heat map with hierarchical clustering for the 6 types of lectins examined.

Figure S3 shows distinctive binding patterns of both ACHN and HEK293. ConA (3 folds,  $p < 0.01$ ) and WGA (2 folds,  $p < 0.01$ ) show strong signal intensity against the cancer ACHN cell line, although SBA, SNA, and LTL also exhibit similar fluorescence intensity. This result suggests that mannose and GlcNAc are over-expressed on ACHN cells compared to HEK293.