

Abstract

Uptake of estrogen precursors is important for cell proliferation in estrogen receptor (ER)-positive breast cancer. Estrone sulfate (E1S) is known as the main precursor of estradiol (E2). Since E1S is a hydrophilic compound, it likely requires a solute carrier to cross plasma membrane. The transporter for E1S therefore seems to be a good candidate as a new therapeutic target against ER-positive breast cancer. The aim of this study is to clarify the relationship between the expression level of the transporter for E1S and cell proliferation in ER-positive breast cancer. Organic anion transporting polypeptide (OATP, *SLCO*) is reported to be involved in uptake of various organic anions, including E1S. The expression of *SLCO1A2*, 2B1 and 3A1 mRNAs was detected in normal breast tissues, malignant breast tissues and MCF-7 cells (a breast cancer-derived cell line). The expression level of *SLCO2B1* mRNA in malignant breast tumors was significantly higher than that in normal breast tissues. Significant positive correlations were observed between the expression level of *SLCO2B1* mRNA and histological grade, expression of Ki-67 protein and STS mRNA in breast malignant tumors. Overexpression of OATP2B1 caused enhancements of E1S uptake, E2 secretion, ER-signal transduction and cell proliferation in MCF-7 cells. The uptake of E1S resulted in down-regulation of ER and induction of Ki-67 in MCF-7 cells. The present study has suggested that the transporter for E1S, such as OATP2B1, affects progression of ER-positive breast cancer, and inhibition of E1S uptake may be enumerated as a new therapeutic target against ER-positive breast cancer.

Objectives

To investigate the relationship between the expression of the transporter for E1S and cell proliferation in ER-positive breast cancer.

Experimental Methods

Cell-based assay

• Cell proliferation assay, reporter gene assay by using reporter plasmid containing estrogen response element and ELISA for E2 were conducted in MCF-7 cells with or without overexpression of OATP2B1.

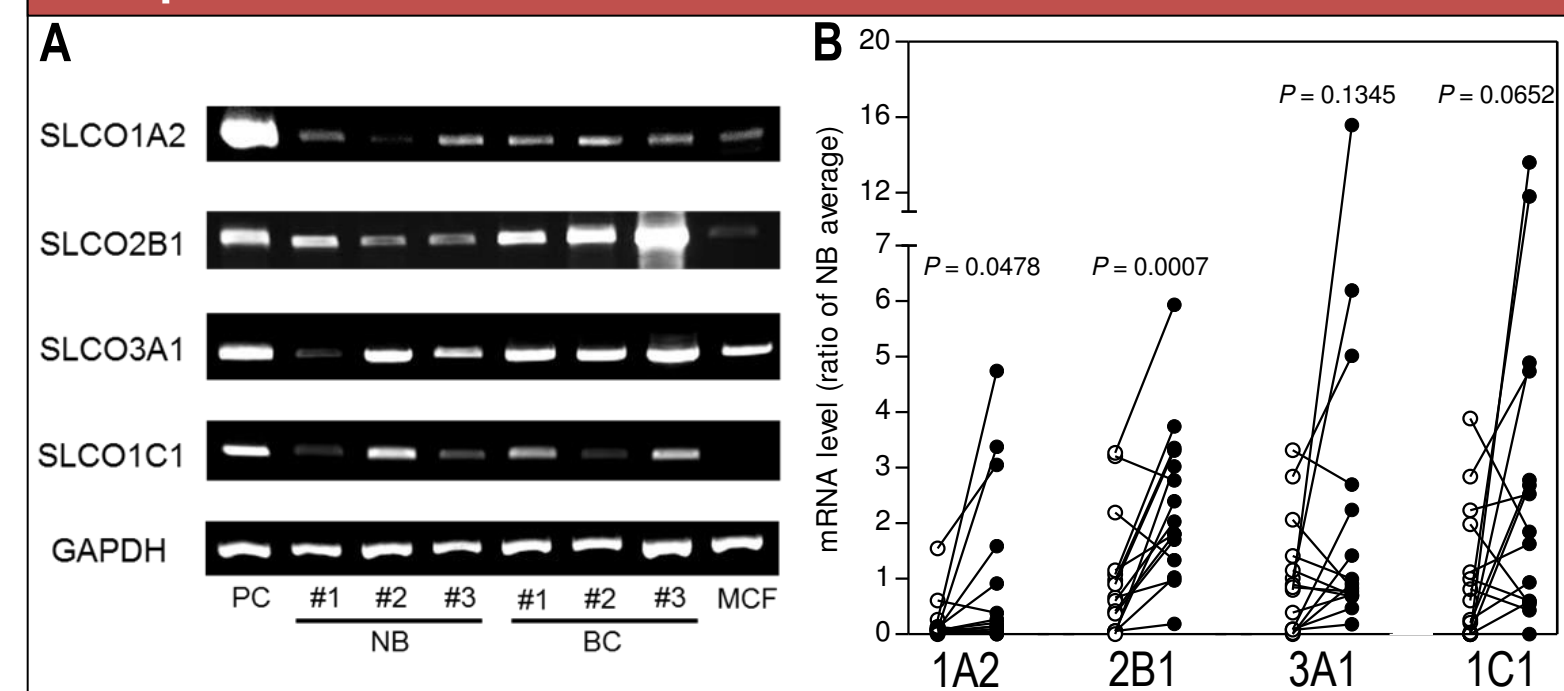
• E1-S uptake experiment was carried out by using [³H]E1-S.

Clinical Samples

• Normal breast and cancer tissues were collected from 16 and 49 patients, respectively. All patients provided written informed consent.

• This study was approved by the ethics committee of Graduate School of Medicine, Chiba University (approval No. 100).

1. Expression of *SLCO* mRNAs in breast cancer tissues

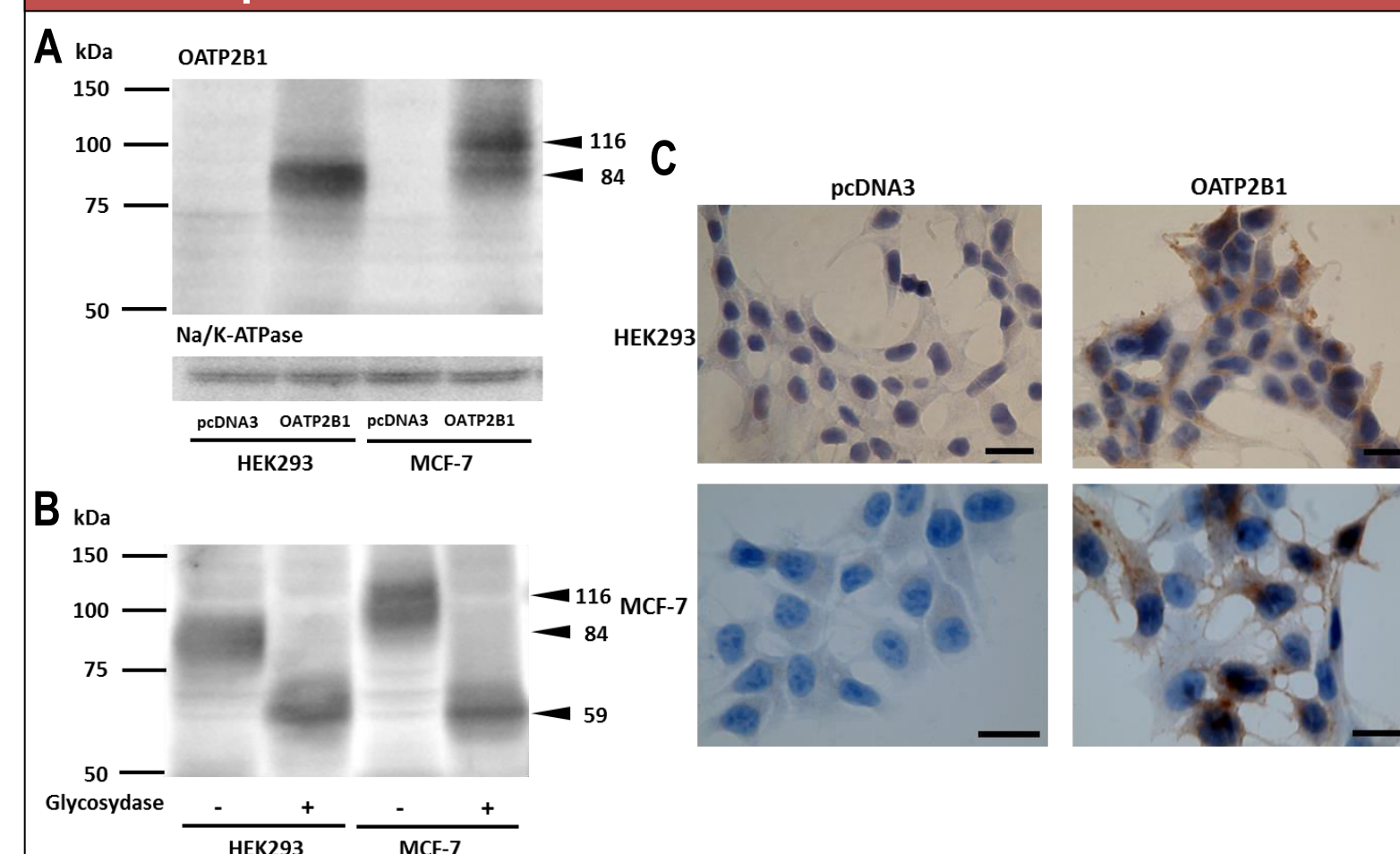


A, Expression of *SLCO* mRNAs in normal breast, cancer tissues and MCF-7 cells. NB, normal breast; BC, breast cancer. B, Differences in the expression levels of *SLCO* mRNAs between NB and BC.

2. The relationship between the expression of *SLCO2B1* mRNA and clinicopathological parameters

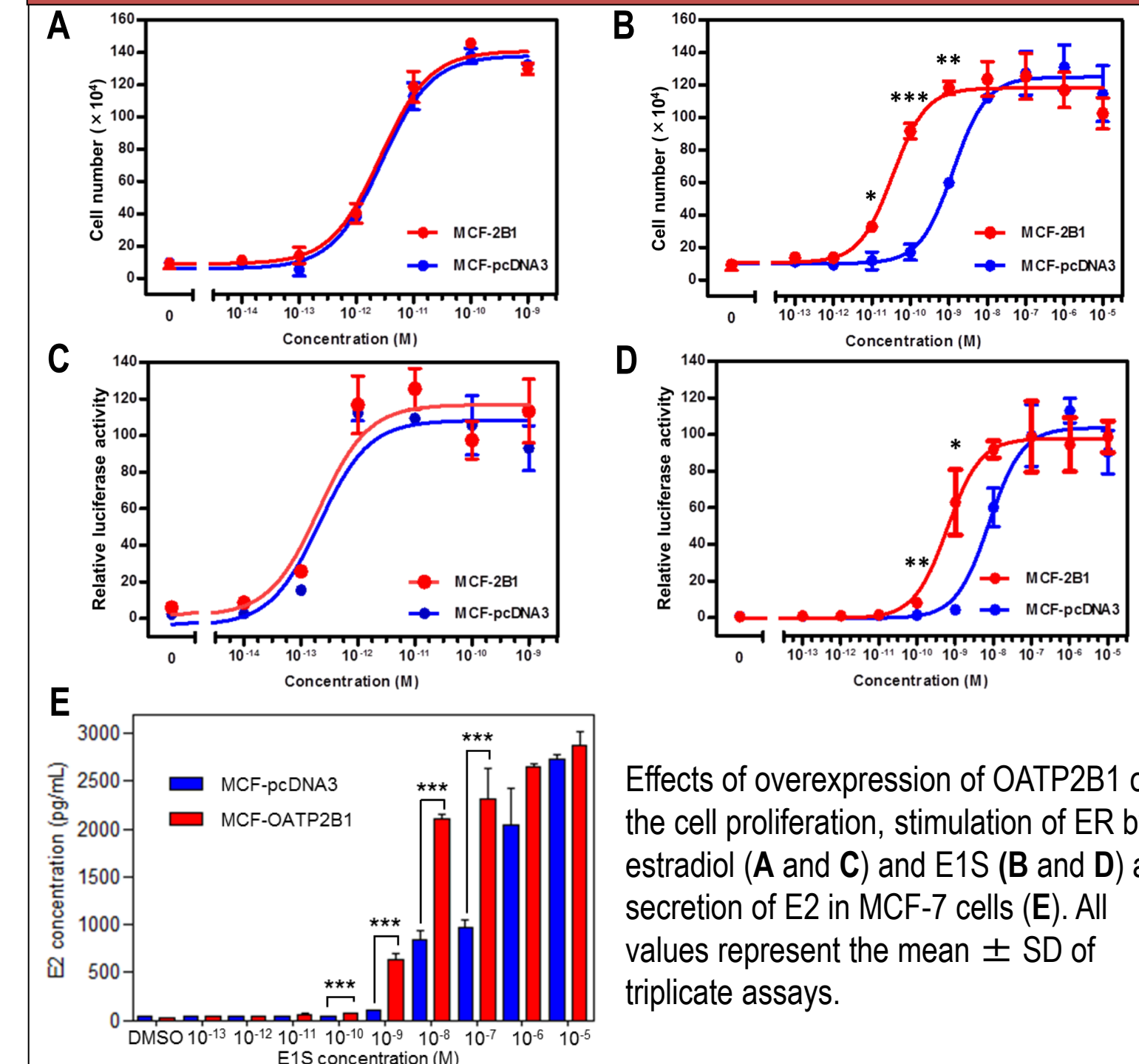
Clinical parameter	n	<i>SLCO2B1</i> mRNA expression (mean ± SE)	P value
Age			
<50 years	23	2.30 ± 0.47	0.220
≥50 years	26	3.60 ± 0.93	
Tumour size			
≤2 cm	18	3.16 ± 0.75	0.795
>2 cm	31	2.89 ± 0.75	
Lymph node status			
Positive	27	2.19 ± 0.41	0.132
Negative	22	3.97 ± 1.08	
Histological grade			
1	14	1.66 ± 0.26	0.022
2 + 3	35	3.52 ± 0.74	
ER			
Positive	40	3.71 ± 1.35	0.563
Negative	9	2.83 ± 0.60	
PgR			
Positive	37	5.10 ± 1.88	0.169
Negative	12	2.30 ± 0.34	
HER2			
Positive	7	2.67 ± 0.70	0.693
Negative	42	3.04 ± 0.62	
Subtype			
Luminal A-like	16	1.79 ± 0.28	0.072
Luminal B-like	19	4.19 ± 1.29	
Ki-67 labelling index	49	r = 0.335	0.019
Estrogenic genes			
Steroid sulfatase	49	r = 0.442	0.001
ERα	49	r = -0.030	0.840
Cyclin D ₁	49	r = 0.177	0.223
Aromatase	49	r = 0.009	0.954

3. Overexpression of OATP2B1 in MCF-7 cells

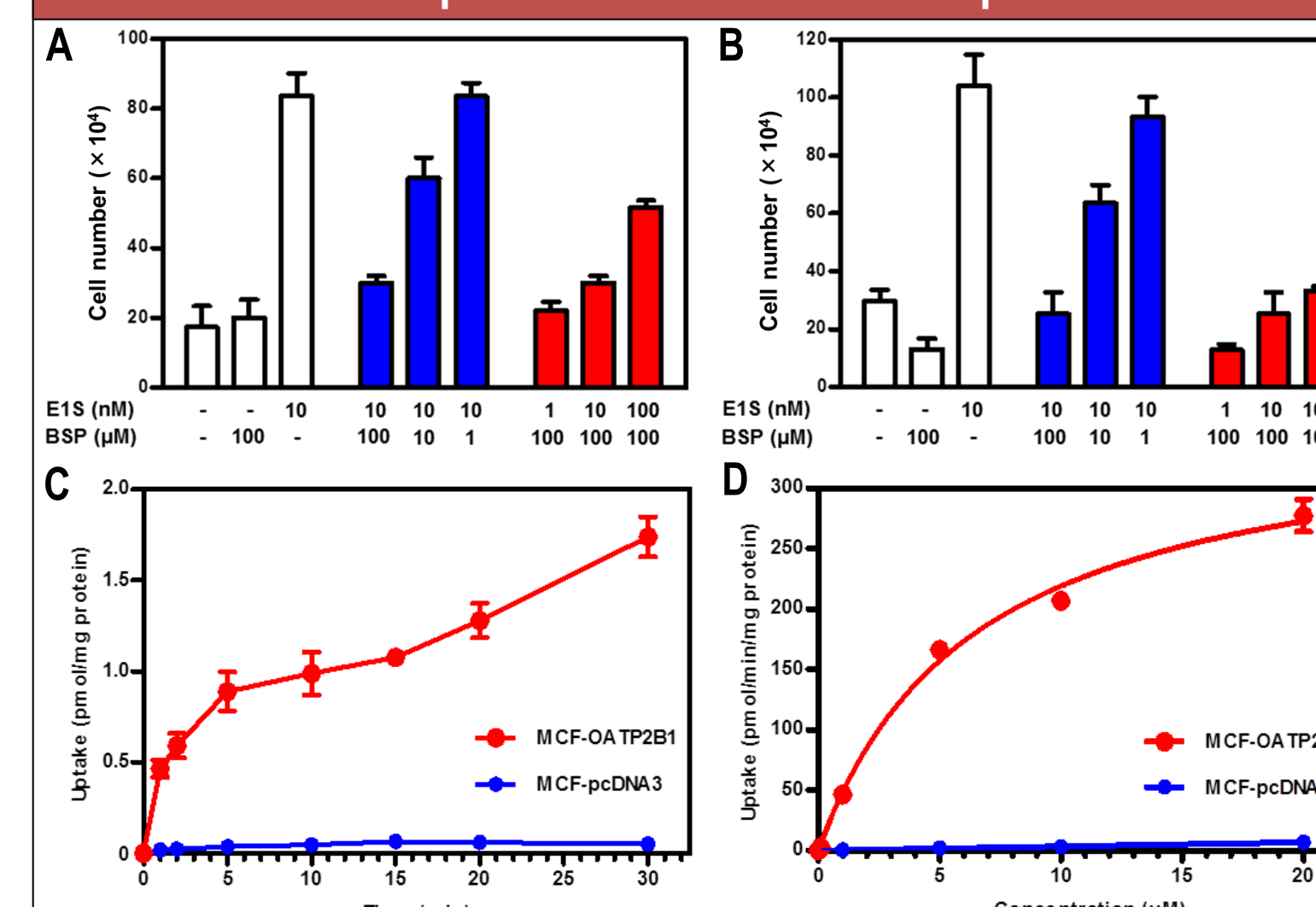


Detection of intact OATP2B1 (A) and deglycosylated OATP2B1 (B) expressed in MCF-7 cells. Na/K-ATPase was used as a loading control. Immunocytochemistry of OATP2B1 expressed in MCF-7 (C). Brown, OATP2B1; blue, nuclei. HEK293 cells stably expressing OATP2B1 were used as a positive control for OATP2B1 specific detection.

4. Effects of overexpression of OATP2B1 on MCF-7 cell proliferation

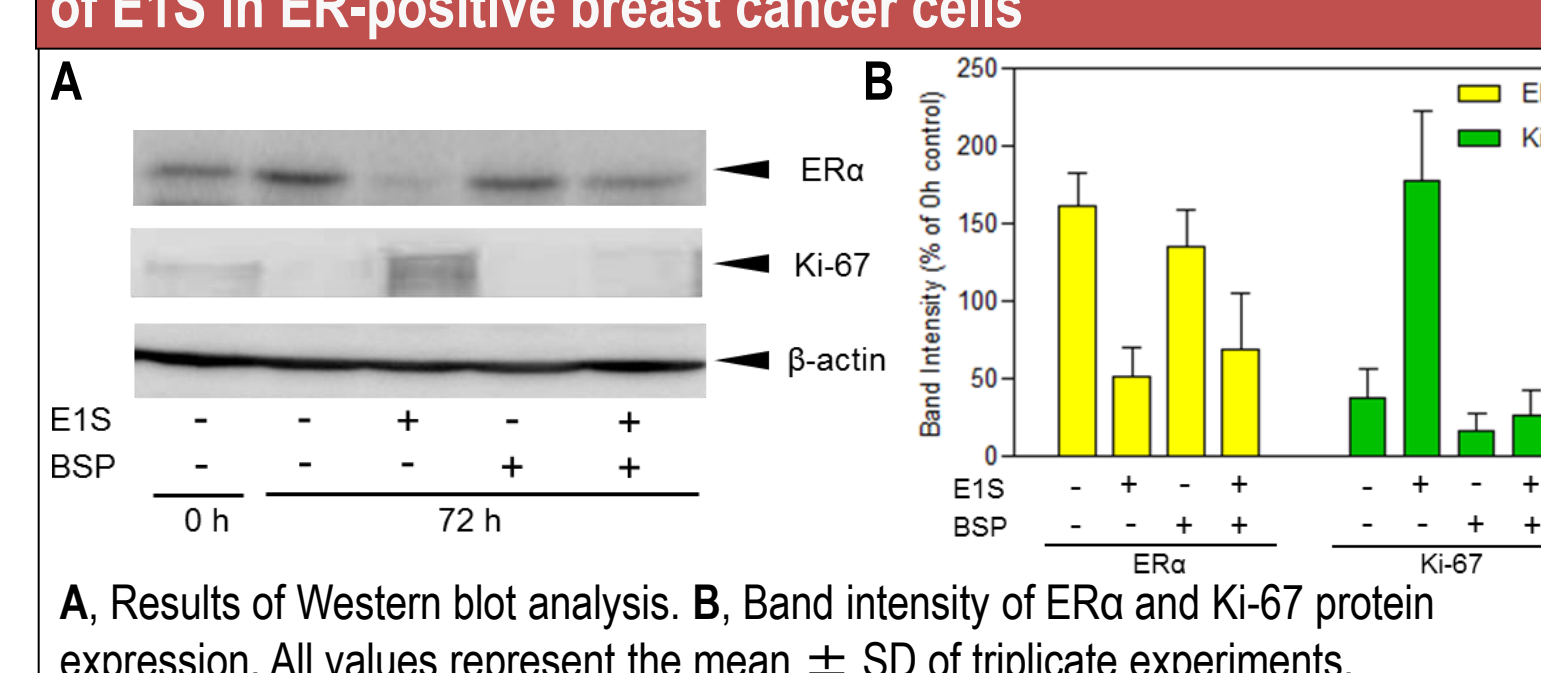


5. Effects of overexpression of OATP2B1 on uptake of E1S



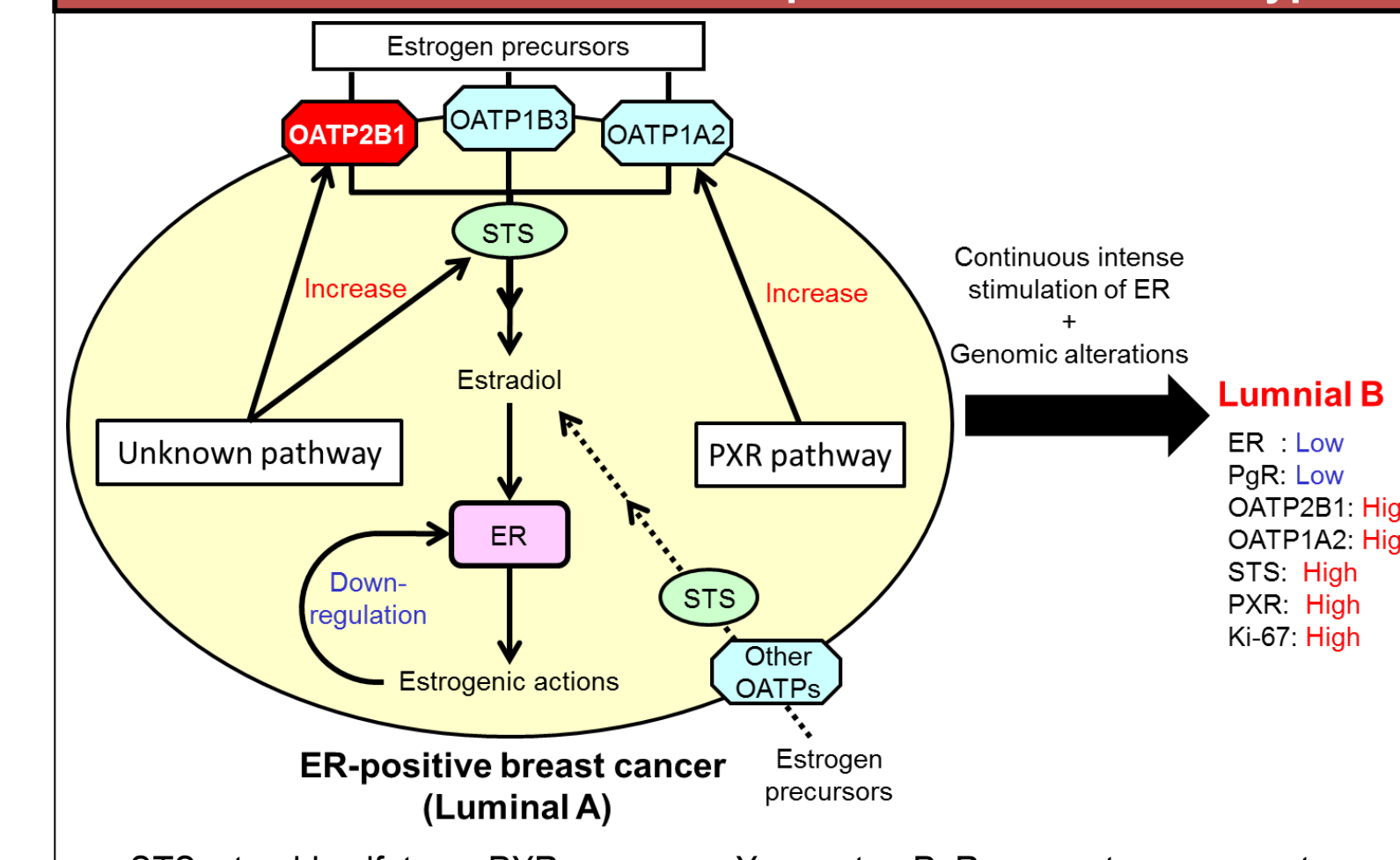
Effects of inhibition of E1S uptake by bromosulphthalein (BSP, a specific inhibitor against OATPs) on the cell proliferation (A, MCF-pcDNA3; B, MCF-OATP2B1). Time course of [³H]E1S (20 nM) uptake (C) and dose-concentration plot of E1S uptake by MCF-7 cells (D). Data represent the mean ± SD of triplicate experiments.

6. Alteration of the expression levels of ERα and Ki-67 by uptake of E1S in ER-positive breast cancer cells



A, Results of Western blot analysis. B, Band intensity of ERα and Ki-67 protein expression. All values represent the mean ± SD of triplicate experiments.

7. Putative relation between E1S uptake and luminal subtypes



STS, steroid sulfatase; PXR, pregnane X receptor, PgR, progesterone receptor

Results

- The expression level of *SLCO2B1* mRNA in breast cancer tissues was significantly higher than that in normal breast tissues, and OATP2B1 was therefore investigated further.
- The expression level of *SLCO2B1* mRNA was positively correlated with Ki-67 labeling index and STS mRNA expression.
- Cell-based assays showed that difference in the expression level of OATP2B1 affected the cell proliferation of ER-positive breast cancer by changing amount of estrogen in the cancer cells.
- The expression level of *SLCO2B1* mRNA in Luminal B tended to be higher than that in Luminal A, and uptake of E1S caused down-regulation of ER and induction of Ki-67.

Discussion

- OATP2B1 may play a key role in supplying ER-positive breast cancer cells with E1S and supporting cell proliferation in ER-positive breast cancer.
- OATP2B1 might be useful as a novel therapeutic target against ER-positive breast cancer, especially luminal B-like cancer.

Conclusion

OATP2B1 may be involved in proliferation of ER-positive breast cancer cells by increasing the amount of estrogens in breast cancer cells.

Contact information

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