<u>FP3-2</u>

CO "c-met expression and molecular targeting therapy in triple-negative breast cancer"



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"Background"

➤ The mortality of breast carcinoma is decreasing because of recent developments in diagnostic techniques and therapies; however, the mortality of the triple-negative breast cancer (TNBC) remains poor. TNBC a poorly characterized subtype of tumor with no validated clinical assay to identify them.

➤ Reportedly, the prognosis of breast cancer is correlated with HGF/c-met coexpression and c-met expression. c-met signaling plays an important role in the proliferation of breast cancer cells. However, little is known about the c-met expression levels of TNBC.

➤ Here, we examined the correlation between TNBC and c-met expression and the effects of c-met inhibitors in TNBC cell lines.







*Materials and Methods" > Breast Cancer Cell Lines TNBC • MDA-MB 231; HR(-), HER2(-) • OCUB-2; HR(-), HER2(-) non-TNBC • MCF-7; HR(+), HER2(-) • OCUB-1; HR(-), HER2(+)

- ➤ c-met inhibitors
- SU11274
- PHA665752

c-met inhibitors on the proliferation of breast cancer cell lines were examined.

"Materials and Methods"

 A total of 1,036 patients who had undergone resection of a primary breast cancer at our institute were enrolled. TNBC 190 cases (18.3%) non-TNBC 846 cases (81.7%)

> Immunohistochemistry

antibody	clone	dilution	source	staining pattern	cutoff values
ER	clone 1D5	1:80	Dako Cytomation	Nucleus	0%
PR	clone PgR 636	1:100	Dako Cytomation	Nucleus	0%
HER2	-	1:300	Dako Cytomation	Membrane	10%
c-met	sc-162	1:100	SANTA CRUZ	Cytoplasm	30%

ER / PR / HER2 status and c-met expression were assessed by immunohistochemistry.



c-met kinase inhibitors inhibited the proliferation of TNBC cell lines. <u>(Proliferation assay)</u>





Breast cancer cell lines expressed c-met mRNA.

<u>c-met Expression</u>

<u>(RT-PCR)</u>





The quantitative RT-PCR showed that the expression level of c-met mRNA was significantly high in MDA-MB 231 and OCUB-2 cells, compared with the expression in OCUB-1 and MCF-7.







c-met kinase inhibitors inhibited the migration of TNBC cell lines.

(Wound healing assay)





control

HGF (30ng/ml)



MDA-MB 23.



c-met SiRNA inhibited the proliferation of TNBC cell lines.

Р





<u>c-met expression</u> (Immunohistochemistry)

positive







<u>Correlation between c-met positivene group and</u> <u>overall survival and disease-free interval.</u>

Triple negative breast cancers (n=190)



The prognosis of patients with c-met positive in 190 TNBC expression was significantly worse than that of those with c-met negative.



Multivariate analysis with respect to overall survival in 190 TNBC.

	Univarite analysis			Multivariate analysis			
	Odds ratio	95% CI	p value	Odds ratio	95% CI	p value	
Stage							
1 vs	2.54	1.04-6.22	0.041	0.37	0.06-2.39	0.298	
2, 3, 4							
Tumor size							
≤2 cm vs	2.46	1.11-5.45	0.027	2.59	0.60-11.22	0.202	
>2 cm							
Lymph node							
status	3.39	1.67-6.88	0.001	3.17	1.26-7.89	0.014	
N0 vs						0.014	
N1, N2, N3							
Lymphvascul							
ar invasion	1.84	0.94-3.58	0.074	1.31	0.65-2.66	0.448	
Negative vs	1.04	0171-0100		101	0.00 2.00	01110	
Positive							
c-met							
Negative vs	3.17	1.44-7.01	0.004	2.51	1.12-5.65	0.026	
Positive							

In TNBC, cases with c-met expression was an independent indicator of a poor prognosis by multivariate analyses.



"Summary"

➤ c-met was expressed in the TNBC cell lines, whose proliferation was enhanced by HGF. c-met kinase inhibitors and c-met SiRNA inhibited the proliferation of TNBC cell lines.

> Clinical samples were immunohistologically examined to demonstrate that c-met expression is an independent poor prognostic factor in TNBC.

"Conclusion"

c-met expression is a potential molecular target and useful in classifying TNBC.