Serum biomarkers of chemoradiosensitivity in esophageal cancer is identified by the targeted metabolomics approach.

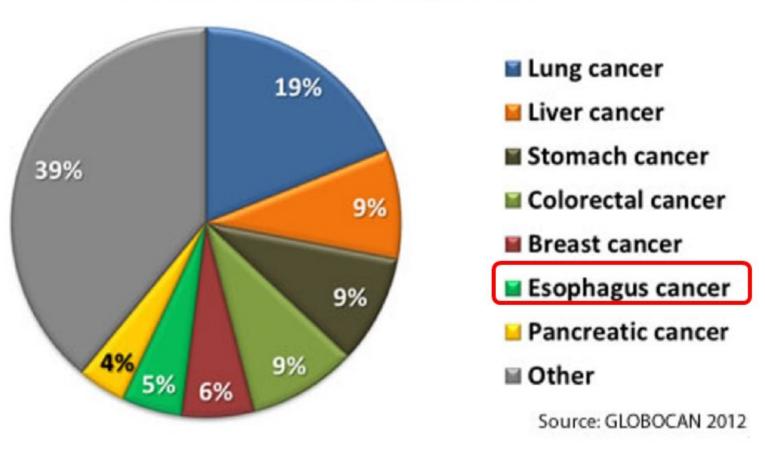


Division of Gastroenterology¹, Metabolomics Research², Kobe University Graduate School of Medicine, AMED-CREST³

Fujigaki S¹, Nishiumi S¹, Kobayashi T¹, Yoshida M^{1,2,3.}

Esophageal Cancer

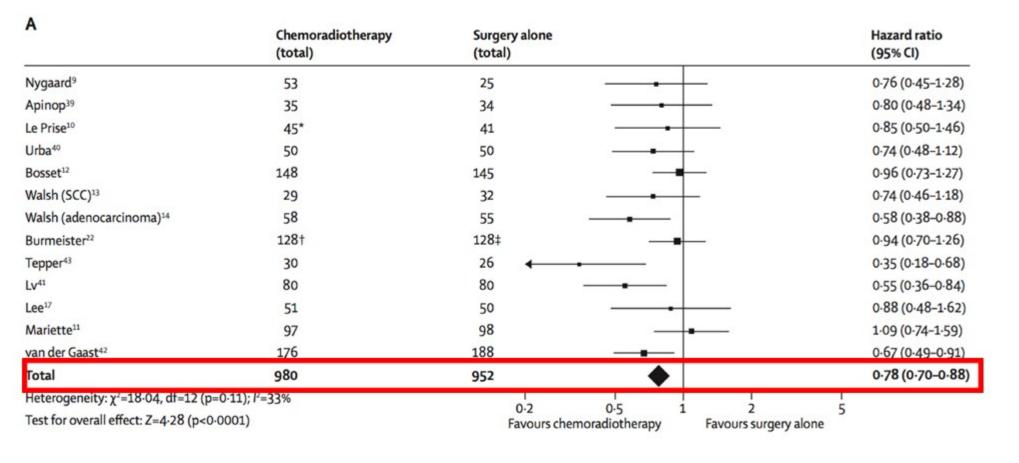
Most Common Causes of Cancer Death Worldwide in 2012



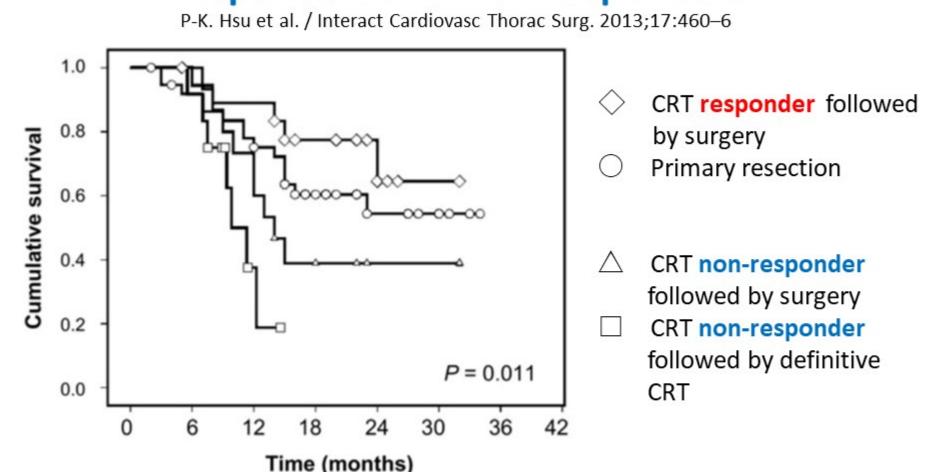
Esophageal Cancer

Sjoquist KM, Burmeister BH, Smithers BM et al. Lancet Oncol. 2011;12(7):681.

Pooled estimates for all-cause mortality for the trials that compared neoadjuvant chemoradiotherapy followed by surgery with surgery alone



Comparison of Neoadjuvant Chemoradiotherapy Responders and Non-responders



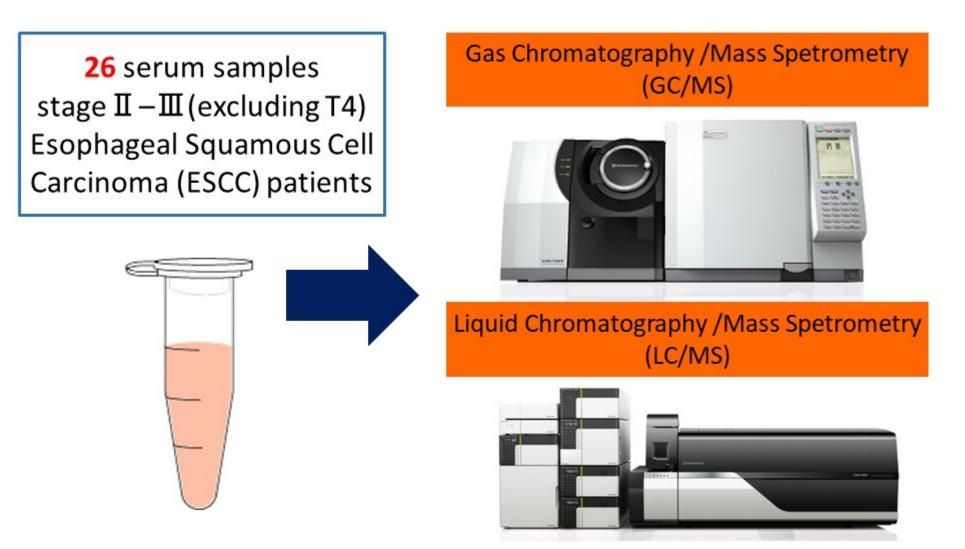
Primary resection vs CRT non-responder, P = 0.036

Aim

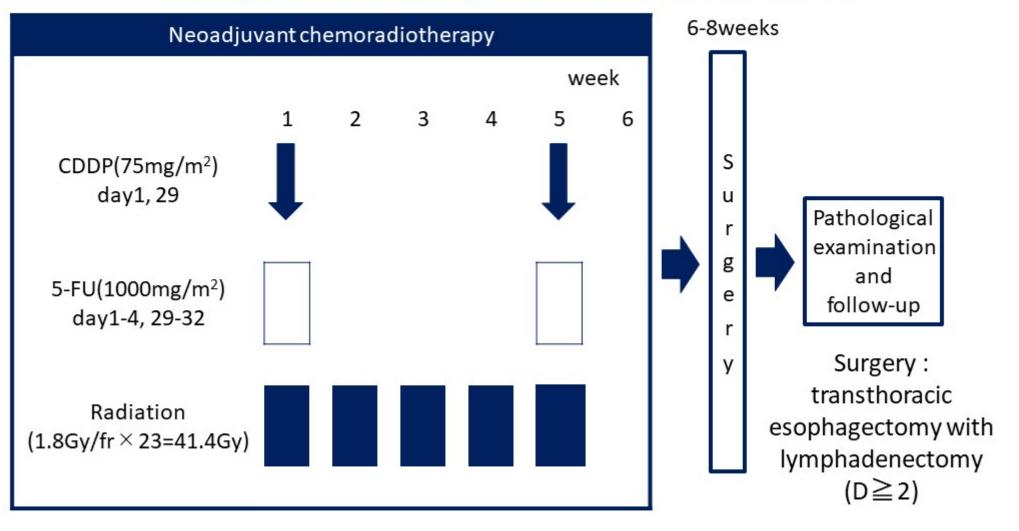
To avoid unfavorable outcomes unnecessary adverse events, reliable methods for predicting the response of esophageal cancer to chemoradiotherapy are desired.

Transcriptomics Gene-expression profiling Mitarion specific PCR MicroRNA-expression profiling DNA microarrays Multiplex PCR Multiplex PCR Multiplex PCR MicroRNA-expression profiling DNA microarrays Multiplex PCR MicroRNA-expression profiling DNA microarrays Multiplex PCR Proteomics Mass spectrometry Mass spectrometry Phosphoproteomic profiling Phosphoproteomic profiling Mass spectrometry Mass spectrometry

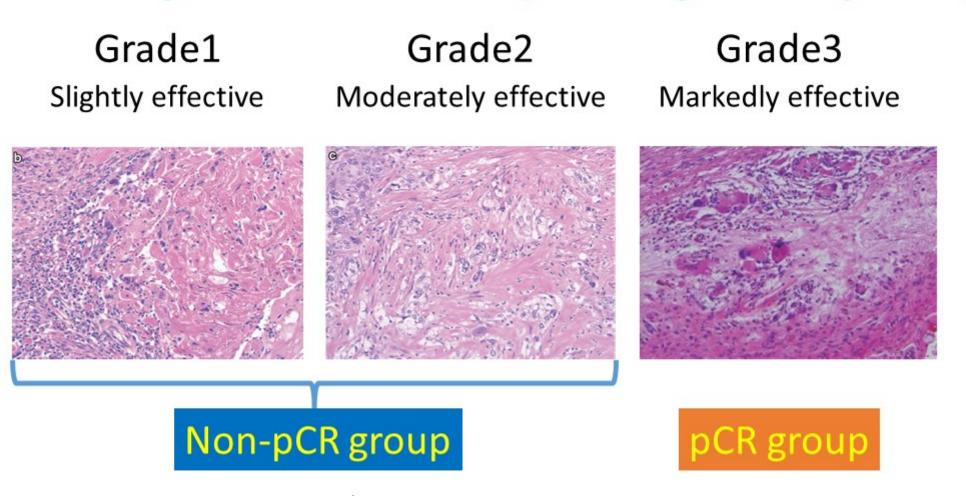
Multiplatform Metabolomics Approach



Feasibility Study of Neoadjuvant Chemoradiotherapy with Cisplatin plus 5-fluorouracil and Elective Nodal Irradiation for Stage II / III Esophageal Squamous Cell Carcinoma



Pathological Examination (Histological Response)



9th edition of the Japanese Classification of Esophageal Cancer

Characteristics of the Subjects

Manialala	Histological	<i>p</i> -value		
Variable	Non-pCR (n=13)	pCR (n=13)	(Fisher's exact t-test)	
Age				
< 65 years	10	7	0.411	
≧65 years	3	6		
Gender				
Male	10	13	0.22	
Female	3	0		
Tumor location				
Lt	4	7	0.428	
Mt	8	6		
Ut	1	0		
Clinical stage				
ΠA	1	0	0.529	
IIВ	2	5		
ША	6	4		
ШВ	4	4		

Serum Metabolites Associated with the Chemoradiosensitivity

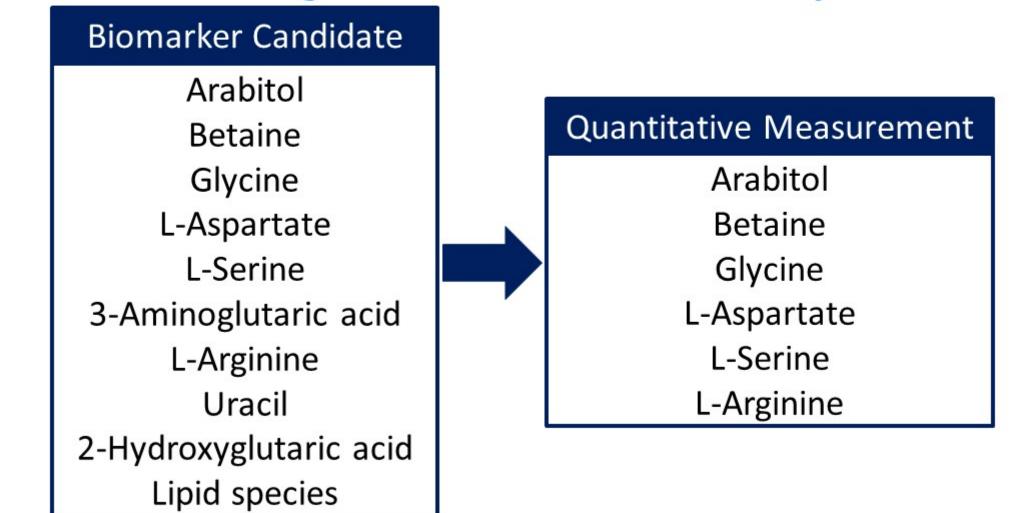
Metabolites	Subclass	Non-pCR (n=13)	pCR (n=13)	Fold	n value
		Mean	Mean	change*	<i>p</i> -value
Arabitol	Sugar alcohols	0.047	0.011	0.230	0.0066
3-Aminoglutaric acid	Amino acids	0.561	0.396	0.705	0.0313
Uracil	Pyrimidines	0.0012	0.0009	0.726	0.0378
Betaine	Amino acids	1.645	1.135	0.690	0.0103
Glycine	Amino acids	1.231	0.99	0.804	0.0103
L-Aspartate	Amino acids	1.249 ati	0.958	0.767	0.021
L-Serine	Amino acids	4.43	3.509	0.792	0.024
L-Arginine	Amino acids	5.386	4.147	0.770	0.0313
2-Hydroxyglutaric acid	Organic acids	0.023	0.016	0.726	0.0402

* Ratio of pCR to non-pCR

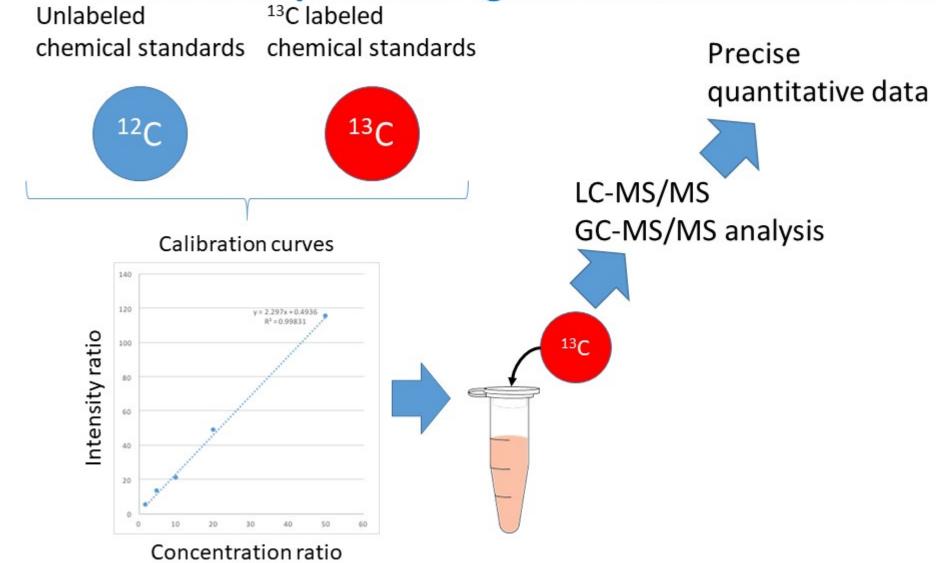
Serum Metabolites Associated with the Chemoradiosensitivity

	no Plot	Volca		Log ₂ (Fold	pCR (n=13)	Non-pCR (n=13)	Metabolites
	+		<i>p</i> -value	change*)	Mean	Mean	wetabolites
+ AC	×.			*******			(
■ FA	2 -		0.0056	-0.6251	0.1015	0.1566	PE(18:1/20:4)
▲ LPC	+ +						PE(18:2/20:4) PE(18:1/20:5)
× LPE	J 01		0.0089	-0.4904	0.5687	0.7989	PE(16:0/22:6) PE(16:1/22:5)
- PC	22	_					PE(20:2/18:4)
+ PE	+# + - ^	alue	0.0159	-0.5150	0.1497	0.2139	PE(16:0/20:4)
	+ +	^-d)0	0.0159	-0.7230	0.0185	0.0305	PE(18:1/20:3) PE(18:2/20:2)
-	+ + + 1 -	-log10(p-value	0.0355	-0.5569	0.1975	0.2906	PE(16:1/18:1) PE(16:0/18:2)
+	+ × + × ++		0.0355	-0.5938	0.1539	0.2323	PE(16:0/18:1)
	^*- **		0.0355	-0.4090	0.4338	0.5760	PE(18:0/20:4)
- <u>-</u> 	+ × + = ×0.5 -		0.0355	-0.5435	0.0436	0.0635	PE(18:1/22:6)
	* * * * * * * * * * * * * * * * * * *		0.0402	-0.4734	0.0371	0.0515	LPC(19:0) (sn-2)
			0.0402	-0.6383	0.0472	0.0735	PC(18:1/22:0)

Targeted Metabolomics Analysis



Stable Isotopic Labeling Assisted Metabolomics



Quantitative Measurement

Metabolit	-	R (n=13) pCR	(n=13)	Fold	n valva	
	M20000	n (μM) Mea	an (μM) cl	hange	<i>p</i> -value	
Arabitol	13.	298 3	.723	0.28	0.0086	
Betaine	62.	015 45	5.902	0.74	0.0613	
Glycine	360	.556 27	0.941	0.75	0.0345	
L-Asparta	te 42.	571 34	1.047	0.8	0.0734	
L-Serine	133	.991 96	5.960	0.72	0.0106	
L-Arginin	e 139	.611 10	8.265	0.78	0.0373	

Prediction Accuracy of pCR

	Metabolites	AUC (95% CI)	Sensitivity (%)	Specificity (%)	Cut-off value (μM)
	Arabitol	0.799 (0.616-0.981)	100	61.5	5.384
	Betaine	0.675 (0.437-0.912)	92.3	61.5	55.364
	Glycine	0.722 (0.517-0.927)	84.6	61.5	336.315
	L-Aspartate	0.692 (0.48-0.905)	84.6	61.5	39.582
	L-Serine	0.781 (0.57-0.992)	92.3	76.9	110.601
<u> </u>	L-Arginine	0.71 (0.504-0.917)	92.3	46.2	138.768

Association of High Serum L-Serine with Shorter Time to Progression (TTP) Cox regression model analysis of prognostic significance

Univariate analysis Variable HR 95% CI p-value ≥65 years/<65 years 0.096-2.24 0.339 0.46 Clinical stage **I**II A and **I**II B / **I**I A and **I**I B 5.33 0.97-99.3 0.0551 Serum Arabitol >5.384 μM/≤5.384 μM 3.08 0.097 0.81 - 12.5Serum Glycine >336.3 μM/≤336.3 μM 2.34 0.62 - 9.520.2056 Serum L-Serine >110.6 μM/≤110.6 μM 3.91 0.0463 1.02-18.7

Serum L-Serine was Significantly Correlated with Long-term Prognosis (Time to Progression and Cause Specific Survival)

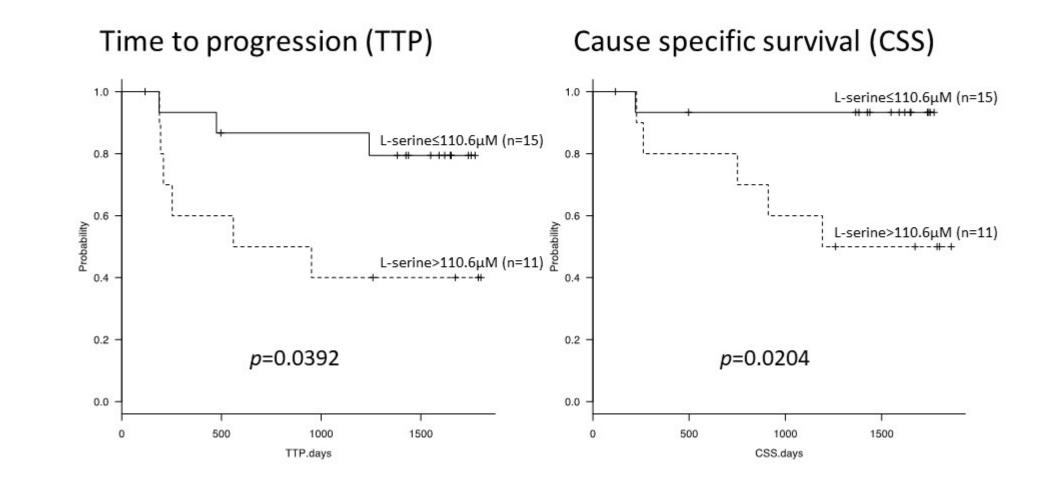
2.77

0.68 - 10.5

0.1463

Serum L-Arginine

>138.8 μM/≤138.8 μM



Conclusion

- In the comprehensive metabolomics analysis, it was confirmed that the serum metabolite profiles of the pCR group were different from those of the non-pCR group.
- The pCR group exhibited significantly lower serum concentrations of serine, glycine, arginine, and arabitol than the non-pCR group.
- The serum concentration of serine could be used to predict the prognosis of ESCC patients who received neoadjuvant chemoradiotherapy.