

CORRESPONDENCE

Gene amplified in oesophageal cancer 1 (GAEC1) amplification in colorectal cancers and its impact on patient's survival

GAEC1 (gene amplified in oesophageal cancer 1) is located at 7q22.1, first identified in oesophageal cancer.¹ Initial work indicated that GAEC1 can act as an oncogene.² Our pilot study found ~80% of colorectal cancers showing amplification of GAEC1.³ In this research, we will study GAEC1 copy number in colon cancer cell lines and colorectal tissues, and its prognostic significance.

Two human colon cancer cell lines (SW480 and SW48) and one normal colonic epithelial cell line (FHC) were obtained from American Type Culture Collection. Culturing conditions for these cell lines were as published previously.⁴ Tissues were collected from 283 patients (213 Australian; 70 Japanese) diagnosed with colorectal cancers. Ninety surgically removed non-cancer colorectal tissues (diverticular diseases, hyperplastic polyps and volvulus) were used as controls. H&E

stained sections from each cancer were checked to select a block with sufficient cancer tissue and representative morphological features for each patient for DNA extraction.

Cancers were staged according to the Union for International Cancer Control (UICC) for TNM (tumour, node and metastasis) classification.⁵ Only primary adenocarcinomas were included. Actuarial survival rates of patients were calculated from the date of surgical resection of the colorectal cancers to the date of death or last follow-up. DNA extraction and PCR reactions for GAEC1 copy number were performed as in our previous studies.^{3,4} A ΔCt of <-1 was considered as gain of GAEC1 copies, Ct value >1 was considered as reduced GAEC1 copies and a Ct between these ranges was defined as normal/no change in GAEC1 copies.

GAEC1 DNA was detectable in all colon cell lines tested. The colon cancer cell lines (SW480 and SW48) showed reduced GAEC1 copies compared with normal colonic epithelial cells (FHC) (figure 1A). At the tissue level, GAEC1 gene copies were higher in colorectal cancer compared with non-cancer tissues ($p<0.0001$; table 1).

In Australian samples, there was no significant correlation between GAEC1 copy number and clinical/pathological

parameters (table 2). Japanese patients showed increased percentage of GAEC1 deletion in colorectal cancers with lymph node metastasis compared with cancers without any lymph node metastasis (10% vs 0%, $p=0.046$) (table 3). GAEC1 deletion was less common in early stage (Stages I and II) colorectal cancers compared with advanced stage (Stages III and IV) cancers in Japanese patients (0% vs 10%, $p=0.046$) (table 3). In addition, the proportion of cancers with GAEC1 amplification was slightly higher in the Japanese compared with the Australian populations (80% vs 59%). GAEC1 copy number distribution was significantly different, with GAEC1 copy number higher among Japanese cancers compared with Australian (p=0.003).

Median follow-up period for Australian and Japanese patients was 5 years. Survival data was available in 188 Australian patients (134 survived and 54 died of cancer). Among Japanese patients, 64 patients had follow-up data (43 patients survived and 21 died of cancer). Kaplan-Meier analysis indicated that mean survival of Australian and Japanese patients with colorectal cancer depended on pathological stage of cancer ($p<0.0001$). For Australian and Japanese patients with colorectal cancer, patients with high GAEC1 copies had longer survival compared with patients with low/

Figure 1 (A) Changes in GAEC1 copy number variation in colon cancer cell lines. Inverse ratio of GAEC1 versus HBD (control) (ct amplification ratio) was used to represent GAEC1 DNA levels in different cell lines. Cancer cell lines (SW480 & SW48) showed lower copies of GAEC1 DNA compared with normal epithelial cell line (FHC). (B and C) Correlation of GAEC1 copy number changes and patients' survival. Australian (B) and Japanese (C) patients diagnosed with colorectal cancers showed better prognosis when they had high GAEC1 copy numbers at the tissue level. On the contrary, patients with cancer having low GAEC1 copy number showed poorer prognosis than other patients. These findings were statistically significant in Australian ($p=0.05$) and Japanese populations ($p=0.02$).

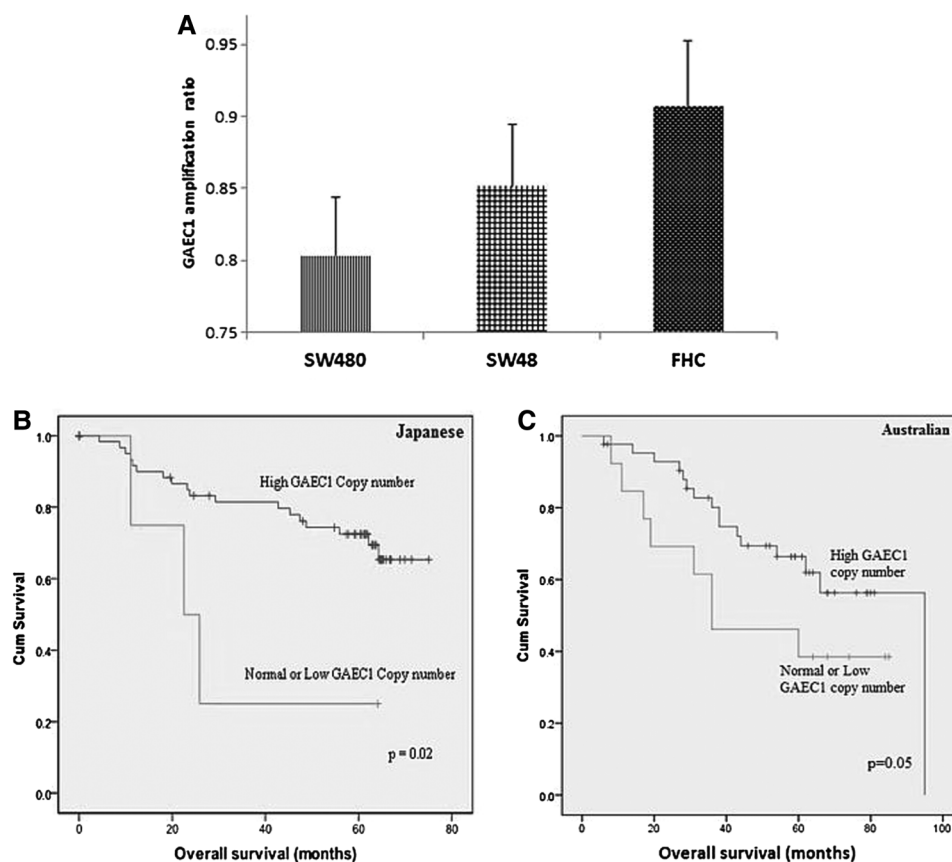


Table 1 DNA copy number variations of *GAEC1* in patients with colorectal cancers and non-neoplastic lesions

	Total	High <i>GAEC1</i>	Low <i>GAEC1</i>	No change	p Value
Australian cancer	213	125 (59%)	34 (16%)	54 (25%)	<0.0001
Japanese cancer	70	56 (80%)	3 (4%)	11 (16%)	
Non-neoplastic	90	18 (20%)	34 (38%)	38 (42%)	

All the statistical tests were done by χ^2 test.

normal *GAEC1* copies (Australians: mean survival time—71 months vs 49 months, $p=0.05$; Japanese: mean survival time—60 months vs 30 months, $p=0.02$; figure 1B,C).

In this study, Japanese patients showed a higher percentage of *GAEC1* amplification and fewer low copy number samples than Australian patients with colorectal cancer. This difference may reflect a real variance in characteristics or causes of colorectal cancer in the two populations. Also, these changes in *GAEC1* could be related to different environmental and genetic factors in these populations. Further studies on these parameters in

conjunction with *GAEC1* may be helpful in identifying exactly what forces are behind these differences. Japanese patients with colorectal cancer showed correlations between *GAEC1* copy number with pathological stages and presence of lymph node metastasis, with high *GAEC1* copies in cancers with earlier pathological stages or no lymph node metastasis. The findings imply that *GAEC1* may function more in early stages of colorectal cancer progression. At later pathological stages, *GAEC1*'s oncogenic roles may be carried on to potential interactors in molecular carcinogenesis. Further experiments including

matched primary and lymph node metastases in the same patient would help in understanding this phenomenon.

This study is the first to investigate the prognostic role of *GAEC1* in patients with colorectal cancer. In Australian and Japanese patients with colorectal cancer, increased *GAEC1* copy number was correlated with a longer survival time. This may indicate that patients with low *GAEC1* copies have more aggressive cancer, perhaps related to the association to pathological stage and lymph node metastasis seen in the Japanese patients. This finding supports the notion that *GAEC1* plays a role in cancer initiation rather than cancer progression.

In contrast with tissues, *GAEC1* showed loss of DNA copies in the colon cancer cell lines compared with normal colon cell lines. In cell lines, genes are less tightly regulated and they are genetically more complex than those in tissues.⁶ Studies have also proven that ex vivo culture conditions can result in the attainment of genomic modification that may not show in the in vivo situation.⁷ Thus, the difference in *GAEC1* copies could be due to environmental alterations in cell lines compared with solid cancer tissues. This phenomenon may indicate that *GAEC1* loses copies when cancer cells are more viable. This can be attributed to the factors influencing *GAEC1* deletion in lymph node metastases and cancer with poor patient prognosis.

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Contributors VG: preparation of manuscript and laboratory works. KY: laboratory works and clinical data for Japanese samples. SP: laboratory works. TT: laboratory works. ML: data management. C-TL: clinical management and follow-up data. JC-oT: supervision of the laboratory works. RAS: co-supervision of the project and proof reading of the manuscript. AK-YL: preparation of manuscript and overall supervision.

Table 2 *GAEC1* copy number variations and the relationships between clinical and pathological features of Australian patients with colorectal adenocarcinoma

Characteristics	Number	Amplification	Deletion	No change	p-Value
Age					0.814
≤60	36 (17%)	26 (72%)	7 (19%)	3 (8%)	
>60	177 (73%)	130 (73%)	28 (16%)	19 (11%)	
Gender					0.984
Male	122 (57%)	89 (73%)	20 (16%)	13 (11%)	
Female	91 (43%)	67 (74%)	15 (16%)	9 (10%)	
Size (mm)					0.186
≤50	160 (75%)	121 (75%)	22 (14%)	17 (11%)	
>50	53 (25%)	35 (66%)	13 (25%)	5 (9%)	
Histological subtypes					0.087
Conventional	195 (91%)	140 (72%)	33 (17%)	22 (11%)	
Mucinous	18 (9%)	16 (89%)	2 (11%)	0 (0%)	
Pathological grade					0.560
Well	64 (30%)	44 (69%)	14 (22%)	6 (9%)	
Mod	116 (54%)	86 (74%)	16 (14%)	14 (12%)	
Poor	33 (16%)	26 (79%)	5 (15%)	2 (6%)	
T-stage					0.750
T1 and T2	52 (24%)	36 (69%)	10 (19%)	6 (12%)	
T3 and T4	161 (76%)	120 (75%)	25 (15%)	16 (10%)	
Lymph node metastasis					0.378
Present	84 (39%)	62 (74%)	16 (19%)	6 (7%)	
Absent	129 (61%)	94 (73%)	19 (15%)	16 (12%)	
Distant metastasis					0.525
Present	34 (16%)	25 (74%)	4 (12%)	5 (15%)	
Absent	179 (84%)	131 (73%)	31 (17%)	17 (10%)	
Stage					0.738
Stages I and II	119 (56%)	86 (72%)	19 (16%)	14 (12%)	
Stages III and IV	94 (44%)	70 (75%)	16 (17%)	8 (8%)	

All the statistical tests were done by χ^2 test and Fisher's exact test.

Table 3 *GAEC1* copy number variations and correlations between clinical and pathological features of Japanese patients with colorectal adenocarcinoma

Characteristics	Number	Amplification	Deletion	No change	p Value
Age					
≤60	18 (26%)	18 (95%)	1 (5%)	0 (0%)	0.708
>60	51 (73%)	48 (94%)	2 (4%)	1 (2%)	
Gender					
Male	41 (59%)	38 (93%)	3 (7%)	0 (0%)	0.082
Female	29 (41%)	28 (96%)	0 (0%)	1 (4%)	
Size (mm)					
≤50	51 (73%)	49 (96%)	2 (4%)	0 (0%)	0.256
>50	19 (27%)	17 (90%)	1 (5%)	1 (5%)	
Histological subtypes					
Conventional	66 (94%)	63 (95%)	2 (3%)	1 (2%)	0.296
Mucinous	4 (6%)	3 (75%)	1 (25%)	0 (0%)	
Pathological grade					
Well	15 (21%)	15 (100%)	0 (0%)	0 (0%)	0.499
Mod	49 (70%)	46 (94%)	2 (4%)	1 (2%)	
Poor	6 (9%)	5 (83%)	1 (17%)	0 (0%)	
T-stage					
T1 and T2	20 (29%)	19 (95%)	0 (0%)	1 (5%)	0.104
T3 and T4	50 (71%)	47 (94%)	3 (6%)	0 (0%)	
Lymph node metastasis					
Present	31 (44%)	28 (90%)	3 (10%)	0 (0%)	0.046
Absent	39 (56%)	38 (97%)	0 (0%)	1 (3%)	
Distant metastasis					
Present	2 (3%)	2 (100%)	0 (0%)	0 (0%)	0.887
Absent	68 (97%)	64 (94%)	3 (4%)	1 (2%)	
Pathological stage					
Stages I and II	39 (56%)	38 (97%)	0 (0%)	1 (3%)	0.046
Stages III and IV	31 (44%)	28 (90%)	3 (10%)	0 (0%)	

All the statistical tests were done by χ^2 test and Fisher's exact test.

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Take-home messages

- ▶ *GAEC1* copy number altered significantly between patients with cancer and the normal population.
- ▶ *GAEC1* amplification was noticed in the cancer population and amplification of *GAEC1* was different in patients with cancer from distinctive ethnic backgrounds.
- ▶ *GAEC1* amplification was associated with negative lymph node metastasis and earlier pathological stages in the Japanese population.
- ▶ *GAEC1* amplification in colorectal cancer was associated with better prognosis.

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