Hollow spheroids beyond the invasive margin indicate the malignant potential of colorectal cancer


INTRODUCTION

An isolated cancer cell or a cancer cluster of fewer than five cancer cells was previously defined as a budding.1 The correlation between budding and metastasis and poor prognosis in colorectal cancer has been reported.1–5 It had been shown that tumour budding was associated with lymphatic invasion and lymph node metastasis in rectal cancer.1 It was also reported that tumour budding was an important predictive factor for recurrence and poor prognosis in patients with colorectal cancer.5 Moreover, it has been shown that tumour budding was significantly correlated with lymph node metastasis, local recurrence, distant metastasis and shorter survival in advanced colorectal cancers.6–8 However, the progression of tumour budding, whether an isolated cancer cell or a cluster of less than five cancer cells, has still to be elucidated. Some reports...
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MATERIALS AND METHODS

Patients

Three hundred and fourteen consecutive patients with colorectal cancer who underwent surgery between January 2002 and August 2005 in Wakayama Medical University Hospital (WMUH) were enrolled in the present study. Forty-six patients had 46 stage I tumours, 125 stage II, 87 stage III, and 56 stage IV based on the UICC TNM classification. The mean age of the patients was 64.9±8.3 years, and there were 185 males and 129 females. The tumours were resected from either the colon (n=175) or the rectum (n=129). The follow-up period was 5 years. Patients with stage III and IV tumours received postoperative chemotherapy based on 5-fluorouracil, whereas most stage I and II patients received no chemotherapy.

Immunohistochemistry

Immunohistochemical staining was performed using paraffin-embedded tissue sections. In brief, sections were de-paraffinised and autoclave-treated at 121°C for 7 min in 0.1 M citrate buffer (pH 6.5). Endogenous peroxidase activity was blocked by incubation in 10% hydrogen peroxide for 5 min at room temperature. After washing with Tris buffer solution and incubation with 5% bovine serum albumin, the sections were incubated with the primary monoclonal antibody to 29H2 (ab94510, diluted 1:100; Abcam, Cambridge, MA, USA), CEACAM1-L polyclonal antibody (diluted 1:1000; Roche Applied Science, Rotkreuz, Switzerland), CEACAM1-S polyclonal antibody (diluted 1:1000; Roche) or M30 cytoDEATH monoclonal antibody (diluted 1:50; Roche) at 4°C for 18 h. M30 monoclonal antibody detects caspase-cleaved cytokeratin 18, which is specific to apoptotic cells.20 27 A biotinylated secondary antibody and peroxidase-conjugated streptavidin were each applied at room temperature for 90 min. Finally, the sections were incubated in 3’-diaminobenzidine for 5 min, followed by haematoxylin counterstaining and mounting. All specimens were blindly reviewed twice by four individuals including a pathologist (KT, JI, SY and YN). In case of difference, the specimen was viewed again using a multihead microscope and the disagreement resolved by consensus.

Cells

Two human colorectal cancer cell lines were used in the present study. LS174T and HT29 cells were purchased from the American Type Culture Collection (Manassas, Virginia, USA). LS174T and HT29 cells were maintained in Eagle’s Minimum Essential Medium (EMEM; Wako, Tokyo, Japan) or McCoy’s 5A (Gibco, Grand Island, New York, USA) medium containing 10% fetal bovine serum (FBS; Gibco), 100 U/ml of penicillin G, 100 μg/ml of streptomycin, and 0.25 μg/ml of amphotericin B (Gibco) at 37°C in a 5% humidified CO2 atmosphere.

Transfection and RNA interference

Transfection of plasmid coding CEACAM1-4L or -4S and short hairpin RNA (shRNA) plasmids targeted to CEACAM1 was performed as previously described.25 Briefly, CEACAM1-4L or -4S cDNA cloned into the pH-β actin vector20 (0.8 μg) was mixed with 1 μl of Lipofectamine 2000 (Invitrogen, Carlsbad, California, USA) in a final volume of 100 μl of Opti-MEM medium, and added to LS174T and HT29 cells grown to 40% confluence in 24-well plates. At 48 h after transfection, G418 (Roche, Basel, Switzerland) was used to isolate stably transfected cells. Short hairpin RNA plasmids targeted to CEACAM1 were synthesised by SABiosciences (Frederick, Maryland, USA) as follows: insert sequence CCA CAA ATG ACA CTG GAA TCT, and gga atc tca ttc gat gca tac (negative control). Each plasmid (0.8 μg) was transfected in the same way as outlined above. The expression level of CEACAM1 was confirmed as previously described.25

demonstrated that the histological morphogenesis of cancer cells can be replicated in 3D culture.9 10 Accordingly, it is speculated that an isolated cancer cell progresses to form budding cells, next a spheroid, and then a hollow spheroid with a lumen. In clinical specimens, hollow spheroids are often regarded as a part of tube formation and a continuous main tumour. In the current study, we report that hollow spheroids formed by polarised luminal cells with an inner empty space exist beyond the tumour margin in colorectal cancer tissue. Moreover, we show that the presence of hollow spheroids is more strongly correlated than tumour budding with metastasis and poor prognosis.

Carcinoembryonic antigen-related cell adhesion molecule 1 (CEACAM1) plays an important role in this process, including lumen formation and inner cell apoptosis in breast cancer cells.9–12 However, the role of CEACAM1 in colorectal lumen formation in 3D culture has not been discussed. CEACAM1 is a type I transmembrane glycoprotein expressed in epithelial cells, and has three or four extracellular domains and either the long (71–73 amino acids, CEACAM1-L) or short (8–10 amino acids, CEACAM1-S) cytoplasmic domain isoform. CEACAM1, previously known as biliary glycoprotein I or CD66a, is a member of the carcinoembryonic antigen family and the immunoglobulin superfamily.9 15 CEACAM1 is down-regulated in malignant tissues derived from the colon,14 15 liver,16 breast17 and prostate,18 whereas high expression of CEACAM1 is associated with the progression of gastric cancer,19 malignant melanoma20 and lung cancer.21 22 In colorectal cancer, it is known that down-regulation of CEACAM1 occurs at the adenoma or early cancer stage,14 15 and CEACAM1 is re-expressed in advanced colorectal cancers.23 24 Recently, we showed that re-expressed CEACAM1, especially with long cytoplasmic domain dominance, in advanced colorectal carcinoma is associated with tumour metastasis and shorter survival.25 In the present study, we demonstrate that CEACAM1 isoform balance with long cytoplasmic domain dominance plays an important role in hollow spheroid formation in colorectal cancer.
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Matrigel culture
Matrigel (200 μl, BD Biosciences, Bedford, MA, USA) was plated on 24-well plates and allowed to solidify for 15 min at 37°C. After the Matrigel solidified, 0.5 ml of the cell suspension in the culture medium containing 2×10⁴ cells/ml was directly seeded into each well and incubated at 37°C in a 5% humidified CO₂ atmosphere for 2 h. After incubation, Matrigel (200 μl, 50% concentration) was added to the upper layer and allowed to solidify for 15 min at 37°C. Complete culture medium was then added, and the plates were incubated at 37°C in a 5% humidified CO₂ atmosphere. During culture, cells were regularly monitored for cell growth and lumen formation, which was scored by the presence or absence of a lumen. Culture medium was changed every 3 days until day 15.

Statistical analysis
The Student t test, the χ² test or Fisher’s exact test were used for statistical evaluation, and clinicopathological factors determined to be significant by the χ² test were analysed by multivariate logistic regression, and an OR with 95% CI was calculated for each factor. The Kaplan–Meier method was used to estimate postoperative survival and the log-rank test was used to determine statistical significance. A Cox proportional hazards model was used to assess the risk ratio under simultaneous contributions from several covariates. p Values of <0.05 were considered to be statistically significant. Statistical calculations were performed using StatView V5.0 software (SAS Institute).

RESULTS
Hollow spheroids beyond the invasive margin of colorectal cancer
To clarify that the colorectal cancer nest had a hollow spheroid structure and that CEACAM1 and central cell apoptosis were involved in hollow spheroid formation, we performed immunohistochemistry with CEACAM1 and M30. A field with more than one hollow spheroid beyond the invasive margin was regarded as positive under ×100 magnification. Of the 91 paraffin-embedded primary tumour tissue sections stained with H&E, 96 sections (30.6%) showed cancer nests with a spheroid hollow beyond the invasive border (figure 1). CEACAM1 and CEACAM1-L, but not CEACAM1-S, were expressed on the luminal inner surface and in the central apoptotic cells (figure 1B). The central cells in the lumen were identified as apoptotic cells as shown by M30 staining (figure 1B). To clarify that the cancer nest had a hollow spheroid structure, serial 2 μm sections of paraffin-embedded tumour tissue were subjected to immunohistological examination with CEACAM1 (29H2). The serial colorectal cancer tissue sections revealed a hollow spheroid structure, and not a tube formation (figure 1C). Budding was counted under ×200 magnification. A field with less than five budding foci was viewed as negative and a field with more than five was viewed as positive.³ A total of 131 sections (41.7%) were positive for budding at the invasive margin.

Hollow spheroids and the malignant potential of colorectal cancer
The χ² test or Fisher’s test for lymph node metastasis revealed that the presence of hollow spheroids, budding, depth of tumour invasion, lymphatic permeation, venous permeation and age >65 were statistically significant (table 1). Multivariate logistic regression analysis for lymph node metastasis showed that the presence of hollow spheroids (p<0.001) and lymphatic permeation (p=0.001) were independent risk factors (table 2). The χ² test or Fisher’s test for distant metastasis revealed that the presence of hollow spheroids, budding, venous permeation, lymphatic permeation and depth of tumour invasion were statistically significant (table 1). The independent risk factors in multivariate logistic regression analysis for distant metastasis were hollow spheroids (p<0.001) and budding (p=0.005) (table 2).

Depth of tumour invasion, lymphatic permeation, venous permeation, lymph node metastasis, distant metastasis, budding and hollow spheroids were of prognostic value for survival in univariate analyses (table 3). Multivariate analyses with the seven parameters which in univariate analyses were significant regarding prognosis, revealed that the presence of hollow spheroids (p=0.001) and distant metastasis (p<0.001) were significant independent prognostic factors (table 3). The Kaplan–Meier method used for overall survival analysis showed that hollow spheroids were significantly associated with shorter survival. The 5-year survival of the hollow spheroid group was 44.8% compared to 88.9% for the non-hollow spheroid group (p<0.0001) (figure 2A). The 5-year survival of the budding group was 60.3% compared to 86.3% for the non-budding group (p<0.0001) (figure 2B).

Association between CEACAM1 and hollow spheroid formation
Hollow spheroid formation of CEACAM1-4L or -4S transfected HT29 cells
Since CEACAM1 is known to have two cytoplasmic domain isoforms, in order to elucidate the effect of CEACAM1 overexpression on colorectal cancer hollow spheroid formation in 3D culture, HT29 cells moderately expressing CEACAM1 were transfected with a plasmid encoding the CEACAM1-4L or -4S isoform. Stable CEACAM1-4L or -4S transfected HT29 cells had significantly elevated CEACAM1 mRNA and protein levels, which was confirmed by RT-PCR and western blot analyses, respectively, as previously described.²⁹ CEACAM1-4L transfected HT29 cells formed a lumen, whereas CEACAM1-4S transfected HT29 cells formed a solid sphere (figure 3A). HT29 cells overexpressing CEACAM1-4L exhibited significantly more lumen formation (p<0.05) (figure 3B). Conversely, CEACAM1-4S transfected HT29 cells exhibited significantly less lumen formation (p<0.01) (figure 3B).
To address whether CEACAM1 knock-down inhibits hollow spheroid formation by colorectal cancer cells, CEACAM1-4L overexpressing LS174T cells were transfected with shRNA against CEACAM1. Stable CEACAM1 shRNA transfected LS174T cells had significantly reduced CEACAM1 mRNA and protein levels, which was confirmed by RT-PCR and western blot analyses, respectively, as previously described. Vector control transfected LS174T cells formed a lumen, whereas CEACAM1 shRNA silencing LS174T cells formed a solid sphere (figure 3A). CEACAM1 knock-down cells exhibited significantly less lumen formation (p<0.01) (figure 3C).

Hollow spheroid formation of CEACAM1-4S transfected LS174T cells overexpressing CEACAM1-4L

To determine whether disturbance of CEACAM1 cytoplasmic isoform balance affected colorectal cancer hollow spheroid formation, CEACAM1-4L overexpressing LS174T cells were transfected with plasmid encoding CEACAM1-4S. Stable CEACAM1-4S transfected LS174T cells overexpressing CEACAM1-4L hollow spheroid formation (figure 3A).

Figure 1  Immunohistochemical staining of hollow spheroids in colorectal cancer tissue. (A) Hollow spheroid, (arrows) beyond the invasive margin expressed CEACAM1 in the luminal surface (29H2) (original magnification, ×100). (B) The luminal surface and centre of the lumen were stained with CEACAM1 (left upper, CEACAM1 (29H2); original magnification, ×400), CEACAM1-L (right upper, CEACAM1-L specific antibodies; original magnification, ×400), CEACAM1-S (left lower, CEACAM1-S specific antibodies, staining of endothelial cells and neutrophils were used as an internal positive control; original magnification, ×400) and M30 cytoDEATH (right lower; original magnification, ×400). (C) Serial sections (i, ii, iii, iv, v, vi) from top (i) to bottom (vi) of a hollow spheroid stained with CEACAM1 (29H2). The lumen was a part of the hollow spheroid and was not a tube structure (original magnification, ×400).
cells had significantly reduced CEACAM1-4L mRNA and protein levels, which was confirmed by RT-PCR and western blot analyses as previously described. The CEACAM1 and CEACAM1-4L protein levels of these cells were significantly suppressed compared to those of the parental or vector-transfected cells (control).

CEACAM1-4S overexpressing cells reduced the number of colorectal hollow spheroids formed (p<0.01) (figure 3D).

**DISCUSSION**

Most investigations for colorectal cancer have shown that tumour budding beyond the invasive margin is a risk factor for lymph node metastasis. In the present study, hollow spheroids are more correlated with lymph node involvement, distant metastasis and overall survival than tumour budding, and multivariate analysis showed that the presence of hollow spheroids is an independent risk factor for metastasis and shorter survival. In particular, multivariate analysis of overall survival demonstrated that independent predictive factors for shorter survival are distant metastasis and hollow spheroids, not tumour budding, indicating that the presence of hollow spheroids is strongly predictive of the malignancy of colorectal cancer. We have previously shown that CEACAM1-L dominance in colorectal cancer is correlated with shorter survival. In the present study, we provided data showing that hollow spheroids expressed CEACAM1-L, not CEACAM1-4S, in the luminal surface, and that their formation is promoted by CEACAM1-4L expression. Kaplan–Meier analysis demonstrated that the hollow spheroid group had

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<th>Table 1</th>
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<td>Present</td>
<td>72</td>
</tr>
<tr>
<td>Absent</td>
<td>59</td>
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* Moderate: moderately differentiated adenocarcinoma; well: well differentiated adenocarcinoma.
† Others: poorly differentiated adenocarcinoma and mucinous adenocarcinoma.

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<th>Table 2</th>
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<td>Hollow spheroid</td>
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<tr>
<td>Distant metastasis</td>
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shorter survival. These results suggested that CEACAM1-L is involved in hollow spheroid formation leading to poor prognosis. A cohort validation study is needed to address the importance of hollow spheroids for colorectal cancer malignancy.

Tumour budding is defined as the presence of individual free cells or undifferentiated cancer cells. This histological structure is also termed ‘sprouting’, ‘focal dedifferentiation’ or ‘budding’. However, it remains unclear whether tumour budding cells survive or undergo apoptosis. Based on 3D culture experiments that replicate the histological morphogenesis of cancer cells, it was speculated that surviving and static dedifferentiated cells proliferate and form spheroids, followed by central cell apoptosis. In the present study, we clarified the existence of hollow spheroids beyond the invasive margin of colorectal cancer, indicating that the same phenomenon of hollow spheroid formation in 3D culture may also occur in clinical colorectal cancer tissue. Further investigations are required to determine if hollow spheroids are derived from tumour budding, although hollow spheroids do seem to be one of the advanced forms of tumour budding.

The mRNA level of CEACAM1 was shown to be down-regulated in colorectal adenoma and early adenocarcinoma. However, some reports have demonstrated that CEACAM1 protein expression is upregulated in advanced adenocarcinomas, and that the intensity of CEACAM1 protein expression is correlated with TNM stage but is not significantly correlated with either overall survival or disease-free survival. Recently, we demonstrated that CEACAM1 was re-expressed at the invasive margin of advanced colorectal cancers, and that CEACAM1-L dominant expression was associated with lymph node involvement, distant metastasis and shorter survival. The isoform balance of CEACAM1 is crucial in the discussion of CEACAM1 function. In the current study, it was demonstrated that CEACAM1-L, not -S, was expressed on the luminal surface of hollow spheroids. The result suggested that CEACAM1-L is involved in hollow spheroid formation leading to poor prognosis.

Our results showed that CEACAM1-L plays an important role in colorectal lumen formation, in contrast to some reports that breast cancer lumen formation is due to CEACAM1-S. Previous breast cancer studies have already demonstrated the discrepancy that either long or short isoforms can promote lumen formation under different conditions. In 3D culture experiments, CEACAM1-4S mediates lumen formation in MCF7 breast cancer cells, whereas CEACAM1-4L induced lumen formation in a mouse model.

### Table 3: Univariate and multivariate analyses of survival

<table>
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<th>Multivariate analysis</th>
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<tr>
<td></td>
<td>HR (95% CI)</td>
<td>p Value</td>
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<tr>
<td>Age (&lt;65)</td>
<td>1.228 (0.785 to 1.919)</td>
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<tr>
<td>Gender (male)</td>
<td>0.974 (0.618 to 1.536)</td>
<td>0.911</td>
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<tr>
<td>Histologic type (others)*</td>
<td>1.522 (0.480 to 4.828)</td>
<td>0.476</td>
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<tr>
<td>Depth (T3, T4)</td>
<td>9.309 (2.285 to 37.920)</td>
<td>0.002</td>
</tr>
<tr>
<td>Lymphatic permeation</td>
<td>3.307 (1.746 to 6.262)</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>Venous permeation</td>
<td>2.200 (1.252 to 3.868)</td>
<td>0.006</td>
</tr>
<tr>
<td>Lymph node metastasis</td>
<td>4.323 (2.635 to 7.093)</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>Distant metastasis</td>
<td>9.126 (5.787 to 14.391)</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>Budding</td>
<td>3.562 (2.209 to 5.743)</td>
<td>&lt;0.001</td>
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<tr>
<td>Hollow spheroid</td>
<td>6.881 (4.240 to 11.166)</td>
<td>&lt;0.001</td>
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*Others: poorly differentiated adenocarcinoma and mucinous adenocarcinoma.

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Figure 2 Kaplan–Meier plots of overall patient survival times. (A) The survival times of the hollow spheroid group and the non-hollow spheroid group were analysed. The 5-year survival of the hollow spheroid group was 44.8% compared to 88.9% for the non-hollow spheroid group (p<0.0001). (B) The survival times of the budding group and the non-budding group were analysed. The 5-year survival of the budding group was 60.3% compared to 86.3% for the non-budding group (p<0.0001).
According to mutation analysis of CEACAM1-4S cytoplasmic domain, phosphorylation of the cytoplasmic domain is crucial for lumen formation. Therefore, both the long and short cytoplasmic domain isoforms of CEACAM1, according to phosphorylation status, can induce lumen formation. The molecular mechanism of CEACAM1 in lumen formation remain to be investigated in greater detail.

Previous studies have shown that enforced overexpression of CEACAM1 in normal and cancerous breast cells induced inner cell apoptosis and hollow spheroid formation in 3D culture and a mouse model. CEACAM1-4S can mediate the reversion of breast cells from a malignant to a non-malignant morphology with a well differentiated structure in 3D culture. However, the association between hollow spheroids in colorectal cancer and CEACAM1 remains to be elucidated. Our clinical investigation showed that the luminal surface of hollow spheroids was stained with CEACAM1-L, but not the short isoform, and central cell apoptosis of hollow spheroids was shown by M30 (cleaved cytokeratin 18) staining. This finding, that hollow spheroids in colorectal cancer are formed by central cell apoptosis, is consistent with the breast cancer 3D culture model. Our 3D culture experiments showed that enforced CEACAM1-4L expression of colorectal cancer cells promoted hollow spheroid formation in 3D culture, and that hollow spheroid formation was suppressed by shRNA targeted to CEACAM1. These results indicated that CEACAM1 expression is important for hollow spheroid formation in colorectal cancer. We also showed that hollow spheroid formation was inhibited by overexpression of CEACAM1-4S, indicating that CEACAM1 isoform balance plays an important role in hollow spheroid formation.

CEACAM1-L contains two tyrosines in immunoreceptor-tyrosine-based inhibitory motifs (ITIMs) that can...
be phosphorylated by Src kinases.\textsuperscript{31} and once phosphorylated, can bind the SHP-1 and SHP-2 phosphatases.\textsuperscript{32} Two cytoplasmic tyrosines of CEACAM1-L affect in vivo tumour growth-inhibitory properties and intercellular adhesion.\textsuperscript{33} It has been shown that CEACAM1-L is able to bind actin and tropomyosin, and that CEACAM1-S can also bind actin, tropomyosin, annexin II and calmodulin.\textsuperscript{34} In normal colonic epithelial cells, CEACAM1-L constitutes 15\%--20\% of the total CEACAM1 protein.\textsuperscript{35} A significant overexpression of the CEACAM1-L isoform or reversion of the normal ratio of CEACAM1-L/S isoforms reverses the tumour inhibition phenotype, whereas CEACAM1-L exhibits tumour inhibition potential when the CEACAM1 isoforms are balanced normally.\textsuperscript{35} We herein demonstrated that CEACAM1-4L promoted hollow spheroid formation in colorectal cancer, while CEACAM1-4S inhibited hollow spheroid formation. The difference may be explained by the different functions of the long and short cytoplasmic domains.

Cytoplasmic domain splicing isoform balance affects the function of CEACAM1. The balance of CEACAM1 isoform expression has been evaluated in epithelial cells.\textsuperscript{35--37} Both long and short CEACAM1 cytoplasmic domain isoforms coexist in most normal tissues. The long isoform is usually expressed in smaller amounts than the shorter isoform.\textsuperscript{35} In normal colonic epithelial cells, CEACAM1-L accounts for 15\%--20\% of total CEACAM1 expressed.\textsuperscript{35} The long isoform has a tumour inhibition phenotype when expressed at physiological concentrations and in the appropriate ratio relative to the short isoform. However, significant overexpression of the long isoform or disruption of the normal L/S ratio abolished the tumour suppressive function.\textsuperscript{35} Therefore, adjustment of the relative ratio of the CEACAM1 isoforms may cause reversion to the normal phenotype. Although further investigation is required to clarify the appropriate isoform balance, CEACAM1 splicing control may be a therapeutic option.

It is important to discuss the function of the carcinoembryonic antigen (CEA) family in molecular and cellular biology. It was shown that only carcinoembryonic antigen-related cell adhesion molecule 6 (CEACAM6) expression, and not CEA or CEACAM1, is an independent prognostic factor for resectable colorectal cancer.\textsuperscript{35} Therefore, CEACAM6 may also be involved in hollow spheroid formation. The association between hollow spheroid formation and other CEA family members such as CEACAM6 and CEA remains to be investigated.

It is speculated that hollow spheroids may form in glandular sites distant from the original lesion. Therefore, hollow spheroids may have acquired tumour-initiating properties. Tumour-initiating cells are essential for tumour growth in a distant lesion. CD133 is one of the promising tumour-initiating markers. It has been reported that the CD133-positive cells in colorectal cancer are tumour initiating and have high tumourigenic potential.\textsuperscript{38 39} Several reports have recently shown that expression of CD133 correlated with poor prognosis in colorectal cancer patients.\textsuperscript{40 41} We hypothesised that hollow spheroids at the invasive margin of colorectal cancers express CD133 as a tumour-initiating marker, and behave as a tumour-initiating phenotype. In the immunohistochemical staining of our preliminary experiments, the staining pattern of CD133 expression resembled that of CEACAM1 expression, indicating that there may be a molecular association between CEACAM1 and CD133 (see supplementary data). CD133 expressing hollow spheroids were observed in all patients with hollow spheroids, suggesting that some of the CEACAM1-expressing hollow spheroids may have tumour-initiating ability. Although further investigation for other tumour-initiating cell markers is required to address the association between hollow spheroids and tumour initiation, hollow spheroids beyond the invasion margin may be predictive of potent tumour-initiating growth at metastatic sites.

In conclusion, the present study demonstrated that hollow spheroids existed beyond the invasive margin and were an independent risk factor for lymph node involvement, distant metastases and shorter survival in patients with colorectal cancer. We also showed that CEACAM1 isoform balance plays an important role in central cell apoptosis in hollow spheroids. Identification of hollow spheroids beyond the invasive margin is important for recognition of the malignant potential of colorectal cancer.

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Competing interests None.

Ethics approval The study was approved by the Human Ethics Review Committee of Wakayama Medical University (#802).

Contributors KT performed immunohistochemistry, conducted cellular and molecular experiments, and carried out statistical analysis. SY conceived and designed the study. JI, SK and TN performed immunohistochemistry. KT, TH and KM carried out surgery. YO, NY and TW performed statistical analysis. YN confirmed the results of immunohistochemical analysis. JS helped in the conception and design of the study and in the production of antibodies. HY helped in the conception and design of the study. All authors drafted and critically revised the manuscript.

Provenance and peer review Not commissioned; externally peer reviewed.

Data sharing statement Immunohistochemistry and 3D culture data will be shared.

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