



THE UNIVERSITY *of* EDINBURGH

Edinburgh Research Explorer

DNA mismatch repair and CD133-marked cancer stem cells in colorectal carcinoma

Citation for published version:

Cheah, PL, Jing, L, Looi, L-M, Teoh, K-H, Ong, DB-L & Arends, M 2018, 'DNA mismatch repair and CD133-marked cancer stem cells in colorectal carcinoma' PeerJ. DOI: 10.7717/peerj.5530

Digital Object Identifier (DOI):

[10.7717/peerj.5530](https://doi.org/10.7717/peerj.5530)

Link:

[Link to publication record in Edinburgh Research Explorer](#)

Document Version:

Publisher's PDF, also known as Version of record

Published In:

PeerJ

General rights

Copyright for the publications made accessible via the Edinburgh Research Explorer is retained by the author(s) and / or other copyright owners and it is a condition of accessing these publications that users recognise and abide by the legal requirements associated with these rights.

Take down policy

The University of Edinburgh has made every reasonable effort to ensure that Edinburgh Research Explorer content complies with UK legislation. If you believe that the public display of this file breaches copyright please contact openaccess@ed.ac.uk providing details, and we will remove access to the work immediately and investigate your claim.



DNA mismatch repair and CD133-marked cancer stem cells in colorectal carcinoma

Phaik-Leng Cheah¹, Jing Li¹, Lai-Meng Looi¹, Kean-Hooi Teoh¹, Diana Bee-Lan Ong¹ and Mark J. Arends²

¹Division of Anatomical Pathology, Department of Pathology, Faculty of Medicine, University of Malaya, Kuala Lumpur, Malaysia

²Division of Pathology, Edinburgh Cancer Research Centre, University of Edinburgh, Edinburgh, United Kingdom

ABSTRACT

Background. Except for a few studies with contradictory observations, information is lacking on the possibility of association between DNA mismatch repair (MMR) status and the presence of cancer stem cells in colorectal carcinoma (CRC), two important aspects in colorectal carcinogenesis.

Methods. Eighty (40 right-sided and 40 left-sided) formalin-fixed, paraffin-embedded primary CRC were immunohistochemically studied for CD133, a putative CRC stem cell marker, and MMR proteins MLH1, MSH2, MSH6 and PMS2. CD133 expression was semi-quantitated for proportion of tumor immunopositivity on a scale of 0–5 and staining intensity on a scale of 0–3 with a final score (units) being the product of proportion and intensity of tumor staining. The tumor was considered immunopositive only when the tumor demonstrated moderate to strong intensity of CD133 staining (a decision made after analysis of CD133 expression in normal colon). Deficient MMR (dMMR) was interpreted as unequivocal loss of tumor nuclear staining for any MMR protein despite immunoreactivity in the internal positive controls.

Results. CD133 was expressed in 36 (90.0%) left-sided and 28 (70.0%) right-sided tumors ($p < 0.05$) and CD133 score was significantly higher in left- (mean \pm SD = 9.6 ± 5.3 units) compared with right-sided tumors (mean \pm SD = 6.8 ± 5.6 units) ($p < 0.05$). dMMR was noted in 14 (35%) right-sided and no (0%) left-sided CRC. When stratified according to MMR status, dMMR cases showed a lower frequency of CD133 expression (42.9%) and CD133 score (mean \pm SD = 2.5 ± 3.6 units) compared with pMMR tumors on the right (frequency = 84.6%; mean score \pm SD = 9.2 ± 5.0 units) as well as pMMR tumors on the left (frequency = 90.0%; mean score \pm SD = 9.6 ± 5.3 units) ($p < 0.05$). Interestingly, frequencies of CD133 immunoreactivity and CD133 scores did not differ between pMMR CRC on the right versus the left ($p > 0.05$).

Conclusion. Proficient MMR correlated with high levels of CD133-marked putative cancer stem cells in both right- and left-sided tumors, whereas significantly lower levels of CD133-marked putative cancer stem cells were associated with deficient MMR status in colorectal carcinomas found on the right.

Subjects Gastroenterology and Hepatology, Pathology

Keywords Tumor side, Colorectal carcinoma, CD133, DNA mismatch repair protein, Cancer stem cells

Submitted 15 March 2018
Accepted 7 August 2018
Published 11 September 2018

Corresponding author
Phaik-Leng Cheah,
cheahpl@ummc.edu.my

Academic editor
Lanjing Zhang

Additional Information and
Declarations can be found on
page 11

DOI 10.7717/peerj.5530

© Copyright
2018 Cheah et al.

Distributed under
Creative Commons CC-BY 4.0

OPEN ACCESS

How to cite this article Cheah et al. (2018), DNA mismatch repair and CD133-marked cancer stem cells in colorectal carcinoma. *PeerJ* 6:e5530; DOI 10.7717/peerj.5530

INTRODUCTION

Colorectal carcinoma (CRC) is a leading cause of death worldwide and the most common malignancy in Malaysian males and the third most common in Malaysian females (*Lim, Ganesanathan & Palaiyan, 2008*). While there have been major advances in understanding the pathogenesis of this important tumor since the CRC tumorigenic model was proposed by *Fearon & Vogelstein (1990)*, much still remains unclear. In recent years, interest has been mounting over the cancer stem cell hypothesis of carcinogenesis in which cancer stem cells are viewed as distinct cells within a tumor that possess the capacity to self-perpetuate, hence maintaining the immortality of the tumor and eventually giving rise to the heterogeneous lineages of the cancer (*Clarke et al., 2006; Lopez et al., 2012*). The cancer stem cell theory would implicate cancer stem cells as central to most tumor development, including CRC, and it would therefore be important to evaluate cancer stem cells and factors that may influence their function and roles to provide insight into this potentially important area.

The identification of the microsatellite instability (MSI) pathway, due to deficient DNA mismatch repair (dMMR) has been incriminated as an alternative in the development of CRC in families with Lynch syndrome (accounting for 2–4% CRC) and in ~15% of sporadic cancers (*Aaltonen et al., 1993; Ionov et al., 1993; Thibodeau, Bren & Schaid, 1993; Szyllberg et al., 2015; Thompson & Spurdle, 2015*). Deficient MMR arises due to failure of the DNA mismatch repair system that results in “instability” of microsatellites; microsatellites being tandem repeats of 1–6 nucleotides scattered in coding and non-coding regions, and the microsatellites together make up 3% of the genome (*Hearne, Ghosh & Todd, 1992; Parsons et al., 1993; Subramanian et al., 2003*). Insertions or deletions of the microsatellites can result in a “mutator” phenotype, affecting genes eg PTEN, BAX, TGFBR2, EGFR etc, that have critical functions in cell signaling, apoptosis and proliferation (*Iacopetta, Grieu & Amanuel, 2010*).

CD133 identified by the AC133 antigenic epitope, is the human homologue of murine Prominin-1, and is one of the most touted markers for the putative cancer stem cells in CRC (*Lapidot et al., 1994; Bonnet & Dick, 1997*). Identifying DNA mismatch repair (MMR) status and correlating this with CD133 expression could provide important information regarding the relationship between defective MMR and colorectal cancer stem cell function. Although some work has been carried out in this area, information remains limited and results are somewhat contradictory, with *Park et al. (2014)* and *Neumann et al. (2012)* demonstrating microsatellite stability with its implicit accompanying proficient DNA mismatch repair system in CRC being associated with high CD133, and *Huh et al. (2010)* the reverse. As deficient MMR is generally reported to occur more commonly in right-sided CRC (*Yearsley et al., 2006; Iacopetta, Grieu & Amanuel, 2010; Benedix et al., 2012; Koi, Tseng-Rogenski & Carethers, 2018*), the study was also designed to stratify the CRC cases according to location of the tumor, to determine if CD133 expression is associated with tumor side.

MATERIAL AND METHODS

Colorectal carcinomas were categorized based on anatomical site viz those proximal to the splenic flexure as right-sided and those at the splenic flexure as well as distal to it as left-sided. Forty consecutive right-sided and 40 left-sided primary CRC, diagnosed between 1st January 2005 and 31st December 2007, were retrieved from the archives of the Department of Pathology, Faculty of Medicine, University of Malaya. All cases were histologically reviewed and only re-confirmed cases were entered into the study. Relevant clinical information was obtained from the surgical pathology examination request forms. The study was approved by the Institutional Review Board (IRB) of the University of Malaya Medical Centre (MEC 794.75) and carried out in compliance with the Declaration of Helsinki. Informed, written consent to undergo resection of the colorectal cancer was obtained from all patients. The Institutional Review Board of the University of Malaya Medical Centre accepts that additional consent from patients is not required for retrospective studies on archived material and which are observational in nature. Confidentiality of patients' personal information was maintained by restricting access of data to the authors of the present work. Five 4 μ m sections were cut from the formalin-fixed, paraffin-embedded primary tumor tissue block selected during histopathological review and placed on platinum coated slides (Matsunami Glass Industries, Kishiwada, Japan) for subsequent immunohistochemical staining.

CD133

Immunohistochemical staining with a rabbit polyclonal antibody to CD133 (1:200; Abcam ab19898, Cambridge, United Kingdom) and detection by the ultraView universal DAB detection kit (Ventana Medical Systems Inc., Tucson, AZ, USA) was carried out on the Ventana Benchmark XT autostainer (Ventana Medical Systems Inc., Tucson, AZ, USA). A case of glioblastoma, previously shown to be immunopositive for CD133 was used as a positive control and run with each batch. CD133 immunostaining was looked for at the glandular luminal surface (Ong *et al.*, 2010). Expression was semi-quantitated for (1) proportion of CD133 immunopositivity in the CRC as 0 (negative), 1 (<5% tumor staining for CD133), 2 (6–25% staining), 3 (26–50% staining), 4 (51–75% staining) and 5 (>75% staining) and (2) intensity of staining 0 (negative), 1 (weak), 2 (moderate) and 3 (strong) (Chew, Teoh & Cheah, 2012). The final CD133 immunopositivity score was calculated as proportion of CD133 immunopositivity multiplied by intensity of staining and arbitrarily measured as “units”. The determination of the cut-off immunostaining positivity would be determined after comparing the CD133 immunostaining in CRC with that of the normal colonic mucosa, in those cases where the latter was available on the same tissue section as the CRC. After determination of the cut-off, the scores would be subcategorized as negative (0 units), low (1–5 units), intermediate (6–10 units) and high (11–15 units).

DNA mismatch repair protein

Based on the understanding that the expression of DNA MMR gene products, MLH1, MSH2, MSH6 and PMS2, are known to fairly reflect MSI status (Pinol *et al.*, 2005; Niessen *et al.*, 2006; Yuan *et al.*, 2015), immunohistochemical staining of the CRC cases was carried

out for DNA MMR on the Ventana Benchmark XT autostainer (Ventana Medical Systems Inc., Tucson, AZ, USA) using monoclonal antibodies to MLH1 (1:100; BD Pharmingen, clone G168-728), MSH2 (1:800; BD Pharmingen, clone G219-1129), MSH6 (1:500; BD Transduction Laboratories, clone 44/MSH6) and PMS2 (1:100; BD Pharmingen, clone A16-4). Detection was via the ultraView universal DAB detection kit (Ventana Medical Systems Inc., Tucson, AZ, USA). Proficient MMR (pMMR) protein expression was defined as unequivocal tumor nuclear immunostaining in the presence of immunopositivity of the internal positive controls (lymphocytes, fibroblasts or normal enterocytes in the vicinity of the tumor). The tumor was classified as having deficient MMR (dMMR) when there was unequivocal loss of tumor nuclear staining for one or more of the MMR proteins in the presence of immunoreactivity in the internal positive controls. The case would be withdrawn whenever the internal controls failed.

Statistical analysis was performed using SPSS (IBM SPSS Statistics version 22; Armonk, NY, USA). For this study, categorical variables would be tested by Chi-square or Fisher's exact test while the Independent Samples T-test or One-way Anova would be used for parametric, and Mann-Whitney U or Kruskal-Wallis for non-parametric continuous variables. Statistical significance was set as $p < 0.05$.

RESULTS

The demographic data and clinical characteristics of the CRC entered into this study are shown in [Table 1](#). Except for a slight preponderance of older patients ($p = 0.043$) and poorly differentiated tumors on the right ($p = 0.025$), there was no significant difference noted in gender and ethnicity of the patients, or size of tumor and TNM staging between the right- and left-sided tumors. Seventy-four of the CRC had some adjacent normal colonic mucosa. Of these, 30 (40.5%) demonstrated weak CD133 staining in the luminal borders of the normal colonic crypts ([Fig. 1A](#)), coupled occasionally with membranous staining of the crypt cells. No staining was observed in 44 of the cases. When staining was present, it was observed in few to majority (>75%) of the normal colonic crypts. Based on these observations, it was decided that weak staining intensity in whatever percentage of the tumor would be considered equivocal and interpreted as negative. Cases without CD133 staining were similarly negative. The CD133 score of the case in either of the above situations would then be calculated to be zero (0) units. Therefore, only staining of at least moderate intensity would be used in the calculation of the score i.e., proportion of CD133 immunopositivity multiplied by intensity of staining. Although when expressed, CD133 decorated the luminal surface of the tumor glands when the glandular structures could be clearly deciphered ([Fig. 1B](#)), tumors with poor differentiation commonly demonstrated CD133 as tiny blobs which probably represent abortive glands, in the tumor cellular cytoplasm ([Fig. 1C](#)) ([Mia-Jan et al., 2013](#)). [Table 2](#) shows the expression of CD133 and MMR status as per size, stage and differentiation of the CRC. CD133 was expressed in 80.0% (64/80) of the CRC. CD133 scores of the cases ranged between 0–15 units with mean \pm SD = 8.2 ± 5.6 units. The size, stage and differentiation of the CRCs were not linked to CD133 expression in terms of frequency of immunopositivity and CD133 score. For the

Table 1 Demographic data and clinical characteristics. Comparison between cases of right-sided and left-sided colorectal carcinoma.

Case characteristics		Right-sided (n = 40)	Left-sided (n = 40)	p
Age (yrs)	Mean	64.6	60.2	0.043*
	Range	15–87	32–81	
Sex	Male	18 (45.0%)	26 (65.0%)	0.072
	Female	22 (55.0%)	14 (35.0%)	
	M:F	1.0:1.2	1.0:0.5	
Race	Chinese	25 (62.5%)	17 (42.5%)	0.089
	Indian	9 (22.5%)	8 (20.0%)	
	Malay	4 (10.0%)	13 (32.5%)	
	Others	2 (5.0%)	2 (5.0%)	
Tumor size	0–5 cm	18 (45.0%)	21 (52.5%)	0.493
	6–10 cm	20 (50.0%)	19 (47.5%)	
	11–15 cm	2 (5.0%)	0 (0%)	
TNM stage	I	4 (10.0%)	3 (7.5%)	0.900
	II	14 (35.0%)	13 (32.5%)	
	III	20 (50.0%)	20 (50.0%)	
	IV	2 (5.0%)	4 (10.0%)	
Differentiation	Well	1 (2.5%)	2 (5.0%)	0.025*
	Moderate	27 (67.5%)	35 (87.5%)	
	Poor	12 (30.0%)	3 (7.5%)	

MMR status, dMMR (Fig. 1D) was noted in 14/80 (17.5%) of the cases, while a majority (82.5%) of cases were pMMR. Of the dMMR tumors, 10 had concomitant loss of MLH1 and PMS2, two concomitant MSH2 and MSH6 loss and one each of isolated loss of either PMS2 or MSH6. dMMR tumors tended to be larger and less differentiated compared with pMMR ones.

When the cases were stratified according to tumor location and MMR status (Table 3), CD133 immunopositivity was identified in 28 of the 40 (70.0%) right-sided and 36 of the 40 (90.0%) left-sided CRC ($p = 0.025$). CD133 scores ranged between 0–15 units on both sides, with right-sided scores (mean \pm SD = 6.8 ± 5.6 units) being lower than left (mean \pm SD = 9.6 ± 5.3 units) ($p = 0.021$). Taking MMR status into account, only 6 of the 14 dMMR CRC (42.9%) were CD133 immunopositive in comparison with 58 of the 66 pMMR cases (87.9%) ($p = 0.0007$). CD133 scores of dMMR cases (range = 0–10 units; mean \pm SD = 2.5 ± 3.6 units) were also lower ($p = 0.00004$) than pMMR CRC (range = 0–15 units; mean \pm SD = 9.4 ± 5.1 units). Frequency of CD133 immunopositivity did not differ ($p = 0.702$) between pMMR tumors on the right (84.6%; 22/26) and the left (90.0%; 36/40). Similarly, CD133 scores of the right-sided pMMR CRC (range = 0–15; mean \pm SD = 9.2 ± 5.0 units) did not differ ($p = 0.625$) from those on the left (range = 0–15; mean \pm SD = 9.6 ± 5.3 units). When CD133 scores were subcategorized to negative, low, intermediate and high, the frequency of dMMR cases in the higher score category was lower than pMMR cases both on the right ($p = 0.0002$) and left ($p = 0.00006$). Interestingly, there

Table 2 CD133 expression and DNA mismatch repair (MMR) status. The CD133 expression and MMR status as per size, TNM stage and differentiation of the colorectal carcinoma ($n = 80$).

Colorectal carcinoma characteristics	CD133									MMR status				
	Immunoexpression			Score						pMMR	dMMR	p		
	Pos n (%)	Neg n (%)	p	Units	p	Neg	Low n (%)	Intermediate n (%)	High n (%)	p	n (%)		n (%)	
Size (cm)	0–5	35 (54.7)	4 (25.0)	0.071	9.5 ± 4.8	0.084	4 (25.0)	3 (37.5)	15 (60.0)	17 (54.8)	0.087	36 (54.5)	3 (21.4)	0.004*
	6–10	28 (43.8)	11 (68.8)		7.3 ± 6.0		11 (68.8)	4 (50.0)	10 (40.0)	14 (45.2)		30 (45.5)	9 (64.3)	
	11–15	1 (1.6)	1 (6.3)		2.0 ± 2.8		1 (6.3)	1 (12.5)	0 (0)	0 (0)		0 (0)	2 (14.3)	
TNM Stage	I	7 (10.9)	0 (0)	0.250	11.4 ± 3.2	0.356	0 (0)	0 (0)	3 (12.0)	4 (12.9)	0.353	6 (9.1)	1 (7.1)	0.889
	II	19 (29.7)	8 (50.0)		7.2 ± 6.1		8 (50.0)	3 (37.5)	7 (28.0)	9 (29.0)		22 (33.3)	5 (35.7)	
	III	32 (50.0)	8 (50.0)		8.5 ± 5.6		8 (50.0)	3 (37.5)	12 (48.0)	17 (54.8)		32 (48.5)	8 (57.1)	
	IV	6 (9.4)	0 (0)		7.5 ± 3.8		0 (0)	2 (25.0)	3 (12.0)	1 (3.2)		6 (9.1)	0 (0)	
Differentiation	Well	3 (4.7)	0 (0)	0.353	9.3 ± 3.1	0.123	0 (0)	0 (0)	2 (8.0)	1 (3.2)	0.078	3 (4.5)	0 (0)	0.008*
	Mod	51 (79.7)	11 (68.8)		8.8 ± 5.5		11 (68.8)	4 (50.0)	20 (80.0)	27 (87.1)		55 (83.3)	7 (50.0)	
	Poor	10 (15.6)	5 (31.3)		5.5 ± 5.6		5 (31.3)	4 (50.0)	3 (12.0)	3 (9.7)		8 (12.1)	7 (50.0)	

Notes.

n, number; pMMR, proficient DNA mismatch repair; dMMR, deficient DNA mismatch repair; pos, positive; neg, negative; mod, moderate.

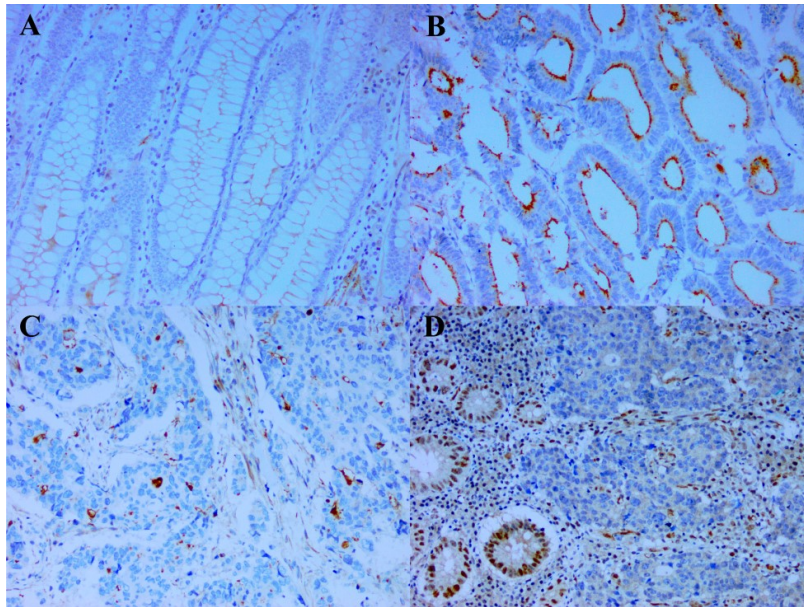


Figure 1 CD133 expression and deficient DNA mismatch repair. (A) The weak CD133 staining seen at the luminal border of some normal colonic crypts is considered equivocal and interpreted as negative (x200). In contrast, (B) is a case of colorectal carcinoma with strong CD133 staining of the glandular luminal surface (x200) and (C) a case of poorly differentiated colorectal carcinoma demonstrating CD133 staining as tiny cytoplasmic blobs which represent abortive glands (x200). (D) Colorectal carcinoma with deficient DNA mismatch repair (represented by loss of MLH1). Note the lack of immunopositivity of MLH1 in the colorectal carcinoma nuclei in the presence of immunopositivity in the internal positive controls (stromal cells, lymphocytes and nearby normal enterocytes) (x200)

Full-size  DOI: [10.7717/peerj.5530/fig-1](https://doi.org/10.7717/peerj.5530/fig-1)

was a marginal increase of left-sided pMMR CRCs in the higher score category compared with pMMR cases on the right ($p = 0.034$).

DISCUSSION

Except for a slight preponderance of older patients and poorly differentiated tumors on the right, the right- and left-sided CRCs showed neither difference in patients' gender and ethnicity, nor in size and stage of the tumors, prompting the notion that both the right- and left-sided CRCs in this study were reasonably matched. Immunopositivity for CD133 was noted in 80.0% (64/80) of the CRC in this study, which falls into the range previously described in other studies (*Yang et al., 2011; Del Rio et al., 2013*). The expression was not associated with size, stage or differentiation of the CRC, as was also noted by *Hong et al. (2015)*. For DNA mismatch repair status, 66 (82.5%) cases were pMMR. In comparison, dMMR was detected in 17.5% of our cases, at a rate comparable to the 6 to 20% reported

Table 3 DNA mismatch repair protein (MMR) status and CD133 expression. CD133 immunoeexpression and score being stratified by MMR status in right-sided and left-sided colorectal carcinomas.

	Right-sided colorectal carcinoma						Left-sided colorectal carcinoma						
	Immuno-expression		CD133				Immuno-expression		CD133				
	Positive	Negative	Units	Low n (%)	Intermed n (%)	High n (%)	Positive	Negative	Units	Low n (%)	Intermed n (%)	High n (%)	
dMMR	6 (21.4)	8 (66.7)	2.5 ± 3.6	3 (100.0)	3 (18.8)	0 (0)	dMMR	0 (0)	0 (0)	0 (0)	0 (0)	0 (0)	0 (0)
pMMR	22 (78.6)	4 (33.3)	9.2 ± 5.0	0 (0)	13 (81.3)	9 (100.0)	pMMR	36 (100.0)	4 (100.0)	9.6 ± 5.3	5 (100.0)	9 (100.0)	22 (100.0)

Notes.

n, number; NA, not applicable; intermed, intermediate; dMMR, deficient DNA mismatch repair; pMMR, proficient DNA mismatch repair.

in other populations (Kim & Kang, 2014). Of the dMMR tumors, 10 had concomitant loss of MLH1 and PMS2, 2 concomitant MSH2 and MSH6 loss and one each of isolated loss of either PMS2 or MSH6. As a further point, in the interpretation of MMR findings, it is necessary to consider the dictates of the obligatory partnerships of eukaryotic MMR proteins (Sinicrope & Sargent, 2012). In the instances of concomitant loss of MLH1 and PMS2 in 10 cases and concurrent loss of MSH2 and MSH6 in two cases, the loss of PMS2 and MSH6 are most likely secondary events consequent upon loss of their respective obligatory binding partners i.e., MLH1 and MSH2. On the contrary, the isolated MSH6 loss in one CRC and loss of PMS2 in another appear to be independent events without loss of their binding partners. The tendency for dMMR tumors to be larger and less differentiated as noted here has also been observed and reported (Iacopetta, Grieco & Amanuel, 2010; Batur et al., 2016).

In this study, dMMR was observed exclusively in right-sided tumors. The predominance of dMMR on the right has been previously reported (Yearsley et al., 2006; Iacopetta, Grieco & Amanuel, 2010; Benedix et al., 2012) although as Cushman-Vokoun et al. (2013) cautions, studies on colorectal carcinoma selected for familial Lynch syndrome screening may show dMMR as high as 48% in left-sided tumors. Our population was unselected and it is not surprising that dMMR was not detected in any of the left-sided tumors. Interestingly, CD133 expression, was in terms of frequency of expression (70.0% right and 90.0% left) and scores (right: mean \pm SD = 6.8 \pm 5.6 units; left: mean \pm SD = 9.6 \pm 5.3 units on left) significantly lower in right-sided compared with left-sided tumors. Thus, pMMR and high level CD133 expression appear to be more prominent in left-sided tumors, indicating a possible association between CD133-marked putative cancer stem cells and proficient MMR. Although proper outcome assessment of pMMR and CD133-positive tumors would require larger study populations and longer follow-up, which was not included as part of this study, expression of CD133 (Bin et al., 2014; Kashiwara et al., 2014) and proficient MMR or MSS (microsatellite stability) (Lin et al., 2014; Seppala et al., 2015) in CRC are generally associated with poorer outcomes. Our study results may therefore indirectly imply that left-sided tumors could be encumbered with two negative-prognostic impact markers. Notwithstanding the above, a large series has shown a bias for a better outcome in left-sided CRC (Loupakis et al., 2015; Price et al., 2015) making the argument that other factors, which have not been interrogated here, also influence the final outcome of cases of CRC.

Stratification of the tumors according to MMR status demonstrated a significantly lower frequency of CD133 immunopositivity as well as CD133 score (42.9%; 2.5 \pm 3.6 units) in dMMR or microsatellite instability-high (MSI-H) CRC compared with pMMR (MSS) cases, both on the right (84.6%; 9.2 \pm 5.0 units) as well as the left (90.0%; 9.6 \pm 5.3 units). At this juncture, it should be evident that the ratio of dMMR to pMMR cases will determine the final frequency and score of a cohort. Hence, the lower CD133 observed in the CRCs on the right compared with those on the left as discussed earlier must take this into cognizance. No difference was observed in the frequency of CD133 immunopositivity or CD133 scores between pMMR cases occurring on the right and left. The observation of a marginal increase of left-sided pMMR cases in the higher CD133 score category compared

with right-sided pMMR cases is noteworthy. While the lack of difference in frequency of positivity and scores leads to the reasoning that CD133 is less likely to be associated with side but more likely with MMR status of the tumor, the observation that more left-sided pMMR cases appeared to be in the high CD133 score category will need larger studies to confirm whether the finding is spurious or has its implications. Nevertheless, this does not detract from the main conclusion that putative CD133-marked stem cells seem to be associated with proficient DNA mismatch repair.

The finding of lower CD133 in dMMR CRC in this study would appear to be more akin to those of *Neumann et al.*'s (2012) and *Park et al.*'s (2014) rather than that of *Huh et al.* (2010). *Neumann et al.* (2012) studied CD133 and MLH1 immunohistochemically in right-sided colonic cancers and noted that CD133 positivity and pMMR were both more common in cases with distant metastasis. *Park et al.* (2014) demonstrated that high CD133 immunopositivity was seen more frequently in MSS CRC. On the contrary, Huh et al. found that CD133 mRNA in their cases of CRC was significantly correlated with MSI-H and by inference dMMR. Notwithstanding the above, the authors also noted that only three out of five cases with CD133 positive immunohistochemical staining, as defined by higher CD133 expression in the cancer compared with the normal colon, showed similar higher CD133 mRNA compared with the normal. *Sgambato et al.* (2010) had also found that CD133 protein evaluated by flow cytometry did not correlate with its mRNA expression level. This then raises the question as to the whether CD133 mRNA should be assumed to equate with CD133 protein expression in CRC tissues. If not, this may explain *Huh et al.*'s (2010) observation that CD133 is associated with MSI-H, which is in contradistinction to the conclusions of *Neumann et al.* (2012), *Park et al.* (2014) and this study. Possible reasons for the association of CD133 with proficient DNA mismatch repair remain unclear. Suggestions of a link via epithelial-mesenchymal transition (EMT), whereby pMMR supports EMT and increased "stemness" of cells; or a situation when increased methylation of promoter CpG islands of CD133 and MLH1 may occur concurrently are possibilities to be considered (*Park et al.*, 2014). Although there is little work done in this area, it is plausible that colorectal cancer stem cells may significantly benefit from a proficient DNA mismatch repair system with a stable microsatellite milieu that promotes continued viability and self-renewal characteristics. Whether the increased capacity for DNA repair noted in human pluripotent stem cells in a study by *Luo et al.* (2012) applies to the MMR system in CRC stem cells will need further investigation.

CONCLUSIONS

This study which addresses the DNA mismatch repair status and colorectal cancer stem cells of 40 right- and 40 left-sided colorectal carcinomas showed that proficient MMR correlated with high levels of CD133-marked putative cancer stem cells in both right- and left-sided tumors, whereas significantly lower levels of CD133-marked putative cancer stem cells were associated with deficient MMR status in colorectal carcinomas found on the right.

ADDITIONAL INFORMATION AND DECLARATIONS

Funding

This study was funded by University of Malaya's research grants RP016C-13HTM and RP016D-13HTM. The funders had no role in study design, data collection and analysis, decision to publish, or preparation of the manuscript.

Grant Disclosures

The following grant information was disclosed by the authors:
University of Malaya's: RP016C-13HTM, RP016D-13HTM.

Competing Interests

The authors declare there are no competing interests.

Author Contributions

- Phaik-Leng Cheah conceived and designed the experiments, performed the experiments, analyzed the data, contributed reagents/materials/analysis tools, prepared figures and/or tables, authored or reviewed drafts of the paper, approved the final draft.
- Jing Li performed the experiments, prepared figures and/or tables, approved the final draft.
- Lai-Meng Looi conceived and designed the experiments, performed the experiments, analyzed the data, contributed reagents/materials/analysis tools, authored or reviewed drafts of the paper, approved the final draft.
- Kean-Hooi Teoh performed the experiments, contributed reagents/materials/analysis tools, approved the final draft.
- Diana Bee-Lan Ong contributed reagents/materials/analysis tools, approved the final draft.
- Mark J. Arends authored or reviewed drafts of the paper, approved the final draft.

Human Ethics

The following information was supplied relating to ethical approvals (i.e., approving body and any reference numbers):

The study was approved by the Institutional Review Board (IRB) of the University of Malaya Medical Centre (MEC 794.75).

Data Availability

The following information was supplied regarding data availability:

The raw data are provided in the [Supplemental File](#).

Supplemental Information

Supplemental information for this article can be found online at <http://dx.doi.org/10.7717/peerj.5530#supplemental-information>.

REFERENCES

- Aaltonen LA, Peltomaki P, Leach FS, Sistonen P, Pylkkanen L, Mecklin JP, Jarvinen H, Powell SM, Jen J, Hamilton SR, Petersen GM, Kinzler KW, Vogelstein B, De la Chapelle A. 1993. Clues to the pathogenesis of familial colorectal cancer. *Science* 260:812–816 DOI 10.1126/science.8484121.
- Batur S, Vuralli Bakkaloglu D, Kepil N, Erdamar S. 2016. Microsatellite instability and B-type Raf proto-oncogene mutation in colorectal cancer: clinicopathological characteristics and effects on survival. *Bosnian Journal of Basic Medical Sciences* 16:254–260 DOI 10.17305/bjbms.2016.1238.
- Benedix F, Meyer F, Kube R, Kropf S, Kuester D, Lippert H, Roessner A, Kruger S. 2012. Influence of anatomical subsite on the incidence of microsatellite instability, and KRAS and BRAF mutation rates in patients with colon carcinoma. *Pathology, Research and Practice* 208:592–597 DOI 10.1016/j.prp.2012.07.003.
- Bin Z, Guangbo Z, Yan G, Huan Z, Desheng L, Xueguang Z. 2014. Overexpression of B7-H3 in CD133+ colorectal cancer cells is associated with cancer progression and survival in human patients. *The Journal of Surgical Research* 188:396–403 DOI 10.1016/j.jss.2014.01.014.
- Bonnet D, Dick JE. 1997. Human acute myeloid leukemia is organized as a hierarchy that originates from a primitive hematopoietic cell. *Nature Medicine* 3:730–737.
- Chew MF, Teoh KH, Cheah PL. 2012. CD133 marks for colorectal adenocarcinoma. *The Malaysian Journal of Pathology* 34:25–28.
- Clarke MF, Dick JE, Dirks PB, Eaves CJ, Jamieson CH, Jones DL, Visvader J, Weissman IL, Wahl GM. 2006. Cancer stem cells—perspectives on current status and future directions: AACR workshop on cancer stem cells. *Cancer Research* 66:9339–9344 DOI 10.1158/0008-5472.CAN-06-3126.
- Cushman-Vokoun AM, Stover DG, Zhao Z, Koehler EA, Berlin JD, Vnencak-Jones CL. 2013. Clinical utility of KRAS and BRAF mutations in a cohort of patients with colorectal neoplasms submitted for microsatellite instability testing. *Clinical Colorectal Cancer* 12:168–178 DOI 10.1016/j.clcc.2013.04.005.
- Del Rio P, Bonati E, Crafa P, Campanini N, Montana Montana C, Bezer L, Dell'Abate P, Sianesi M. 2013. CD133 is a selective marker of CRC? *Minerva Chirurgica* 68:87–95.
- Fearon ER, Vogelstein B. 1990. A genetic model for colorectal tumorigenesis. *Cell* 61:759–767.
- Hearne CM, Ghosh S, Todd JA. 1992. Microsatellites for linkage analysis of genetic traits. *Trends in Genetics* 8:288–294.
- Hong I, Hong SW, Chang YG, Lee WY, Lee B, Kang YK, Kim YS, Paik IW, Lee H. 2015. Expression of the cancer stem cell markers CD44 and CD133 in colorectal cancer: an immunohistochemical staining analysis. *Annals of Coloproctology* 31:84–91 DOI 10.3393/ac.2015.31.3.84.

- Huh JW, Park YS, Lee JH, Kim HR, Shin MG, Kim YJ. 2010. CD133 mRNA expression and microsatellite instability in colorectal carcinoma. *Journal of Surgical Oncology* 102:765–770 DOI 10.1002/jso.21734.
- Iacopetta B, Grieu F, Amanuel B. 2010. Microsatellite instability in colorectal cancer. *Asia-Pacific Journal of Clinical Oncology* 6:260–269 DOI 10.1111/j.1743-7563.2010.01335.x.
- Ionov Y, Peinado MA, Malkhosyan S, Shibata D, Perucho M. 1993. Ubiquitous somatic mutations in simple repeated sequences reveal a new mechanism for colonic carcinogenesis. *Nature* 363:558–561 DOI 10.1038/363558a0.
- Kashihara H, Shimada M, Kurita N, Iwata T, Sato H, Kozo Y, Higashijima J, Chikakiyo M, Nishi M, Matsumoto N. 2014. CD133 expression is correlated with poor prognosis in colorectal cancer. *Hepatogastroenterology* 61:1563–1567.
- Kim JH, Kang GH. 2014. Molecular and prognostic heterogeneity of microsatellite-unstable colorectal cancer. *World Journal of Gastroenterology* 20:4230–4243 DOI 10.3748/wjg.v20.i15.4230.
- Koi M, Tseng-Rogenski SS, Carethers JM. 2018. Inflammation-associated microsatellite alterations: mechanisms and significance in the prognosis of patients with colorectal cancer. *World Journal of Gastrointestinal Oncology* 10:1–14 DOI 10.4251/wjgo.v10.i1.1.
- Lapidot T, Sirard C, Vormoor J, Murdoch B, Hoang T, Caceres-Cortes J, Minden M, Paterson B, Caligiuri MA, Dick JE. 1994. A cell initiating human acute myeloid leukaemia after transplantation into SCID mice. *Nature* 367:645–648 DOI 10.1038/367645a0.
- Lim KG, Ganesanathan S, Palaiyan K. 2008. In: Lim GCC, Rampal S, Halimah Y, eds. Cancer incidence in Peninsular Malaysia 2003–2005: the third report of the national cancer registry. Kuala Lumpur: National Cancer Registry, Ministry of Health, 83–86.
- Lin CC, Lin JK, Lin TC, Chen WS, Yang SH, Wang HS, Lan YT, Jiang JK, Yang MH, Chang SC. 2014. The prognostic role of microsatellite instability, codon-specific KRAS, and BRAF mutations in colon cancer. *Journal of Surgical Oncology* 110:451–457 DOI 10.1002/jso.23675.
- Lopez J, Ruiz G, Organista-Nava J, Gariglio P, Garcia-Carranca A. 2012. Human papillomavirus infections and cancer stem cells of tumors from the uterine cervix. *The Open Virology Journal* 6:232–240 DOI 10.2174/1874357901206010232.
- Loupakis F, Yang D, Yau L, Feng S, Cremolini C, Zhang W, Maus MK, Antoniotti C, Langer C, Scherer SJ, Muller T, Hurwitz HL, Saltz L, Falcone A, Lenz HJ. 2015. Primary tumor location as a prognostic factor in metastatic colorectal cancer. *Journal of the National Cancer Institute* 107:Article dju427 DOI 10.1093/jnci/dju427.
- Luo LZ, Gopalakrishna-Pillai S, Nay SL, Park SW, Bates SE, Zeng X, Iverson LE, O'Connor TR. 2012. DNA repair in human pluripotent stem cells is distinct from that in non-pluripotent human cells. *PLOS ONE* 7:e30541 DOI 10.1371/journal.pone.0030541.
- Mia-Jan K, Jung SY, Kim IY, Oh SS, Choi E, Chang SJ, Kang TY, Cho MY. 2013. CD133 expression is not an independent prognostic factor in stage II and III colorectal

- cancer but may predict the better outcome in patients with adjuvant therapy. *BMC Cancer* 13:166 DOI 10.1186/1471-2407-13-166.
- Neumann J, Horst D, Kriegl L, Maatz S, Engel J, Jung A, Kirchner T. 2012.** A simple immunohistochemical algorithm predicts the risk of distant metastases in right-sided colon cancer. *Histopathology* 60:416–426 DOI 10.1111/j.1365-2559.2011.04126.x.
- Niessen RC, Berends MJ, Wu Y, Sijmons RH, Hollema H, Ligtenberg MJ, De Walle HE, De Vries EG, Karrenbeld A, Buys CH, Van der Zee AG, Hofstra RM, Kleibeuker JH. 2006.** Identification of mismatch repair gene mutations in young patients with colorectal cancer and in patients with multiple tumours associated with hereditary non-polyposis colorectal cancer. *Gut* 55:1781–1788 DOI 10.1136/gut.2005.090159.
- Ong CW, Kim LG, Kong HH, Low LY, Iacopetta B, Soong R, Salto-Tellez M. 2010.** CD133 expression predicts for non-response to chemotherapy in colorectal cancer. *Modern Pathology* 23:450–457 DOI 10.1038/modpathol.2009.181.
- Park JJ, Kwon JH, Oh SH, Choi J, Moon CM, Ahn JB, Hong SP, Cheon JH, Kim TI, Kim H, Kim WH. 2014.** Differential expression of CD133 based on microsatellite instability status in human colorectal cancer. *Molecular Carcinogenesis* 53(Suppl 1):E1–E10 DOI 10.1002/mc.21971.
- Parsons R, Li GM, Longley MJ, Fang WH, Papadopoulos N, Jen J, De la Chapelle A, Kinzler KW, Vogelstein B, Modrich P. 1993.** Hypermutability and mismatch repair deficiency in RER+ tumor cells. *Cell* 75:1227–1236.
- Pinol V, Castells A, Andreu M, Castellvi-Bel S, Alenda C, Llor X, Xicola RM, Rodriguez-Moranta F, Paya A, Jover R, Bessa X, Gastrointestinal Oncology Group of the Spanish Gastroenterological Association. 2005.** Accuracy of revised Bethesda guidelines, microsatellite instability, and immunohistochemistry for the identification of patients with hereditary nonpolyposis colorectal cancer. *JAMA* 293:1986–1994 DOI 10.1001/jama.293.16.1986.
- Price TJ, Beeke C, Ullah S, Padbury R, Maddern G, Roder D, Townsend AR, Moore J, Roy A, Tomita Y, Karapetis C. 2015.** Does the primary site of colorectal cancer impact outcomes for patients with metastatic disease? *Cancer* 121:830–835 DOI 10.1002/cncr.29129.
- Seppala TT, Bohm JP, Friman M, Lahtinen L, Vayrynen VM, Liipo TK, Ristimaki AP, Kairaluoma MV, Kellokumpu IH, Kuopio TH, Mecklin JP. 2015.** Combination of microsatellite instability and BRAF mutation status for subtyping colorectal cancer. *British Journal of Cancer* 112:1966–1975 DOI 10.1038/bjc.2015.160.
- Sgambato A, Puglisi MA, Errico F, Rafanelli F, Boninsegna A, Rettino A, Genovese G, Coco C, Gasbarrini A, Cittadini A. 2010.** Post-translational modulation of CD133 expression during sodium butyrate-induced differentiation of HT29 human colon cancer cells: implications for its detection. *Journal of Cellular Physiology* 224:234–241 DOI 10.1002/jcp.22124.
- Sinicrope FA, Sargent DJ. 2012.** Molecular pathways: microsatellite instability in colorectal cancer: prognostic, predictive, and therapeutic implications. *Clinical Cancer Research* 18:1506–1512 DOI 10.1158/1078-0432.ccr-11-1469.

- Subramanian S, Madgula VM, George R, Mishra RK, Pandit MW, Kumar CS, Singh L. 2003.** Triplet repeats in human genome: distribution and their association with genes and other genomic regions. *Bioinformatics* **19**:549–552.
- Szylberg L, Janiczek M, Popiel A, Marszalek A. 2015.** Serrated polyps and their alternative pathway to the colorectal cancer: a systematic review. *Gastroenterology Research and Practice* **2015**:Article 573814 DOI [10.1155/2015/573814](https://doi.org/10.1155/2015/573814).
- Thibodeau SN, Bren G, Schaid D. 1993.** Microsatellite instability in cancer of the proximal colon. *Science* **260**:816–819.
- Thompson BA, Spurdle AB. 2015.** Microsatellite instability use in mismatch repair gene sequence variant classification. *Genes* **6**:150–162 DOI [10.3390/genes6020150](https://doi.org/10.3390/genes6020150).
- Yang ZL, Zheng Q, Yan J, Pan Y, Wang ZG. 2011.** Upregulated CD133 expression in tumorigenesis of colon cancer cells. *World Journal of Gastroenterology* **17**:932–937 DOI [10.3748/wjg.v17.i7.932](https://doi.org/10.3748/wjg.v17.i7.932).
- Yearsley M, Hampel H, Lehman A, Nakagawa H, De la Chapelle A, Frankel WL. 2006.** Histologic features distinguish microsatellite-high from microsatellite-low and microsatellite-stable colorectal carcinomas, but do not differentiate germline mutations from methylation of the MLH1 promoter. *Human Pathology* **37**:831–838 DOI [10.1016/j.humpath.2006.02.009](https://doi.org/10.1016/j.humpath.2006.02.009).
- Yuan L, Chi Y, Chen W, Chen X, Wei P, Sheng W, Zhou X, Shi D. 2015.** Immunohistochemistry and microsatellite instability analysis in molecular subtyping of colorectal carcinoma based on mismatch repair competency. *International Journal of Clinical and Experimental Medicine* **8**:20988–21000.