

Detection of Circulating Gastrointestinal Cancer Cells in Conditionally Reprogrammed Cell Culture

CHUN-SEOK YANG¹, IN-HWAN KIM¹, HYUN-DONG CHAE¹, DAE-DONG KIM² and CHANG-HO JEON³

¹Department of General Surgery, Medical School,
Daegu Catholic University Medical Center, Daegu, Republic of Korea;

²Department of General Surgery, Medical School, Yonsei University, Seoul, Republic of Korea;

³Department of Laboratory Medicine, Medical School,
Daegu Catholic University Medical Center, Daegu, Republic of Korea

Abstract. *Background/Aim:* The aim of this study was to detect circulating tumor cells (CTC) in the peripheral blood of gastrointestinal cancer patients using conditionally reprogrammed cell (CRC) culture. *Materials and Methods:* We confirmed the sensitivity of the CRC culture method. Five ml of blood were obtained from 81 cancer patients (56 colorectal and 25 gastric). The collected mononuclear cells were cultured for 4 weeks in the CRC condition. Finally, cultured cells were characterized by RT-PCR for the expression of *hTERT* and *MAGE A1-6* mRNA. *Results:* The CRC method had a CTC detection limit of 6 cells for gastric cancer cells. After culture of 81 blood specimens, 38 formed visible cells, including 5 colonies. Among the 38 cells, 13 were *hTERT* positive and 4 were *MAGE A1-6* positive. The final CTC detection rate was 16.0%. *Conclusion:* The CRC culture may potentially be used to evaluate the metastatic cancer cells in the circulation.

Detection of circulating tumor cells (CTCs) in the blood has been investigated using various methods based on antibodies, transcripts or proteins (1, 2). Recently, many researchers utilized cell-free DNA (cfDNA) instead of CTCs (3-5). Regarding some solid tumors, cfDNA has produced promising results (6). However, the approach is limited because cfDNA yields the same information regarding mutations and epigenetic modifications as the tumor tissue

without providing any information about protein and RNA transcription profiles of the tumor phenotype (7).

The detection of CTCs has been refined by the progression from the conventional immunologic capture method using EpCAM and cytokeratin antibody to an *in vitro* cell culture method (8, 9). CTC culture consists of short-term culture, such as the epithelial immunospot approach, and long-term culture (1). Successful culture rates for patients with early lung cancer have been achieved using a three-dimensional co-culture system (10). However, CTC cultures for gastrointestinal cancers are rare (11-14).

The Georgetown University Medical Center (GUMC) has also developed a conditionally reprogrammed cell (CRC) culture method using 3T3 mouse fibroblasts (J2 cells) and a Rho-kinase inhibitor (Y27632) (15). The CRC culture method is very efficient for the culture of tumor cells *in vitro*.

In this study, we attempted to detect circulating colorectal and gastric cancer cells using the CRC culture. To characterize the cultured cells isolated from the peripheral blood, we used the Melanoma antigen-encoding gene (*MAGE*) and human telomerase reverse transcriptase (*hTERT*) mRNA as markers. *MAGE* is known to be a cancer-specific gene, but individual *MAGE* genes are poorly expressed in cancer cells (16, 17). Common *MAGE* primers that can detect *MAGE A1* to *A6* (*MAGE A1-6*) mRNA simultaneously were developed and have been used in various types of cancer (18, 19). The *hTERT* mRNA has also been studied in many types of cancer cells (20). Though the *hTERT* mRNA would be also expressed in activated lymphocytes (21), it has been utilized as a sensitive and specific tumor marker in the blood.

Materials and Methods

Sensitivity of CRC technique. To confirm the sensitivity of the CRC technique, we used SNU484 gastric cancer cells obtained from the Korean Cell Line Bank (Seoul, Republic of Korea), and colorectal

This article is freely accessible online.

Correspondence to: Chang-Ho Jeon, Department of Laboratory Medicine, Medical School, Daegu Catholic University Medical Center, 33 Duryugongwon-Ro 17-Gil Namgu, Daegu, 42732, Republic of Korea. Tel: +82 1038111034, e-mail: chjeon@cu.ac.kr

Key Words: Blood, neoplasm, culture, reverse transcription, *hTERT*, *MAGE*.

Table I. Gene-specific primers used in the polymerase chain reactions.

Gene	Sequences	Products	Ta (°C)	Cycles
MAGE A1-6	OF 5'-CTG AAG GAG AAG ATC TGC C	465 bp	60	30
	OR 5'-CCA GCA TTT CTG CCT TTG TGA			
	IF 5'-AAG GAG AAG ATC TGC CAG TG	262 bp	62	25
	IR 5'-GAG GCT CCC TGA GGA CTC T			
hTERT	OF 5'-CGG GCT GCT CCT GCG TTT GGT G	311 bp	68	30
	OR 5'-AGC CGC GGT TGA AGG TGA GAC TGG			
	IF 5'-TCA CCT CAC CCA CGC GAA AAC CTT	152 bp	66	25
	IR 5'-CGT GGG CCG GCA TCT GAA CAA A			
GAPDH	OF 5'-TCG GAG TCA ACG GAT TTG GTC GTA	320 bp	59	33
	OR 5'-CAA ATG AGC CCC AGC CTT CTC CA			

MAGE: Melanoma antigen-encoding gene; hTERT: human telomerase reverse transcriptase; GAPDH: glyceraldehyde-3-phosphate dehydrogenase; OF: outer forward; OR: outer reverse; IF: inner forward; IR: inner reverse.

cancer and gastric cancer cells isolated from cancer patients using CRC. J2 embryonic fibroblasts were kindly donated from the Georgetown University Medical Center (GUMC). Six, 12, 25, 50, 500 and 5,000 cells of the three cell types were incubated for 4 weeks in the CRC condition. The CRC condition was prepared using the same protocols of the GUMC (15).

Patient samples. Five ml of blood were obtained from each of the 81 cancer patients. The patients had been admitted at Daegu Catholic University Medical Center (Daegu, Korea) and diagnosed with gastric or colorectal cancer radiographically and pathologically. The 81 patients comprised 56 colorectal cancer and 25 gastric cancer patients. The stage of gastric and colorectal cancer was determined according to the American Joint Committee on Cancer 7th edition. Informed consent was obtained from each patient. The study protocol was approved by the Institutional Review Board (IRB) of Daegu Catholic University Medical Center. To collect CTCs from the blood specimens, red blood cells were lysed using RBC lysis buffer (Roche, Basel, Switzerland) and blood mononuclear cells were collected.

CRC culture and RNA extraction from the culture plates. The collected mononuclear cells were incubated for 4 weeks at 37°C in the CRC condition. Briefly, Swiss-3T3-J2 mouse fibroblasts were cultured to generate feeder cells and then, the isolated mononuclear cells were plated in F medium containing J2 feeder cells and Y-27632 (cat. no. 270-333M025, Enzo Life Sciences, Lausen, Switzerland). F medium was made by mixing Dulbecco's modified Eagle's medium (DMEM) (Invitrogen, Gaithersburg, MD, USA), 5% fetal bovine serum (Gibco; Thermo Fisher Scientific Inc., Waltham, MA, USA), L-glutamine (Gibco), penicillin/streptomycin mix (Gibco), F12 nutrient mix (Gibco), 25 ng/ml hydrocortisone (Sigma-Aldrich, St. Louis, MO, USA), 0.125 ng/ml epidermal growth factor (EGF) (Invitrogen), 5 µg/ml insulin (Sigma-Aldrich), 250 ng/ml amphotericin B (Gibco), 10 µg/ml gentamicin (Gibco) and 0.1 nM cholera toxin (Sigma-Aldrich). All cells were maintained at 37°C in a cell culture incubator with 95% humidity and 5% CO₂. At the end of incubation, some colonies, cell aggregations, J2 cells or unknown cells persisted on the plate. All 81 samples were treated with 1 ml Trizol (Invitrogen), RNA was extracted, and RNA quality and quantity were measured using a spectrophotometer (NanoDrop Technologies, Wilmington, DE, USA).

Real time RT-PCR. The mRNAs of *hTERT* and *MAGE A1-6* were amplified to determine the identity of the cultured cells. The mRNA of glyceraldehyde-3-phosphate dehydrogenase (*GAPDH*) was amplified to confirm the RNA integrity for all specimens. The *hTERT* and *MAGE A1-6* mRNAs were amplified with nested PCR, whereas *GAPDH* mRNA was amplified with a single round of PCR amplification, using specific primers for each gene and the PCR conditions presented in Table I. Cells were determined as cancer cells if they expressed one cancer related mRNA. The amplification of all mRNAs was performed using the LightCycler 480 (Roche, Basel, Switzerland). We interpreted the PCR products as positive results using fluorescent signals and melting temperature analysis.

STR (short tandem repeat) analysis of culture cells and cancer tissue. To verify the cultured cell, we used STR analysis. In one colorectal cancer patient, some cells proliferated to form a colony (CR1520). We extracted DNA using the remnant of RNA extraction procedure and STR markers were analyzed using PowerPlex 16 system (Promega, Madison, WI, USA) with ABI 3500xL Genetic Analyzer (Applied Biosystems, Waltham, MA, USA). The matched cancer tissue of CR1520 obtained from the biopsy was also used for the STR analysis.

IRB approval and statistical analysis. This study was conducted in accordance with the standards of the Declaration of Helsinki and was approved by the IRB of Daegu Catholic University Medical Center (CR-16-002-L). All patients have provided written informed consent for their information to be stored and used in the hospital database.

For the statistical analysis, Chi-squared test was performed using MedCalc Version 14.12.0 (Ostend, Belgium).

Results

Sensitivity of CRC culture. Sensitivity data of spiked cancer cells cultured in the CRC condition are summarized in Table II. Gastric cancer cells were cultured from 6 to 5,000 cells, and colorectal cancer cells were cultured from 25 to 5,000 cells. Interestingly, even the lowest number of gastric cancer cells (6) grew in culture.

Table II. Sensitivity of cancer cell culture using conditionally reprogrammed cell method.

Cell types	Culture result by cell number					
	6	12	25	50	500	5000
SNU 484	+	+	+	+	+	+
Colorectal cancer	–	–	+	+	+	+
Gastric cancer	+	+	+	+	+	+

Clinical data, CRC culture and RT-PCR results. Demographic data of cancer patients are summarized in Table III. Most gastric cancer patients were stage I and II, while colorectal cancer patients were stage II and III.

After 4 weeks of incubation, some cells were cultured in 38 of 81 blood samples. The cultured cells were observed as colonies, cell aggregations, J2 cells and unknown cells. Figure 1 shows the cultured colonies of gastric and colorectal cancer. Among the 38 cells, 13 cells showed positive results with *MAGE* A1-6 or *hTERT* RT-PCR analysis. The overall positive rates were 16.0% for *hTERT* and 4.9% for *MAGE* A1-6. The respective positive rates for each mRNA were 16.1% and 5.4% in colorectal cancer, and 16.0% and 4.0% in gastric cancer, respectively (Table III). All *MAGE* A1-6 positive cases were also positive for *hTERT* RT-PCR.

STR analysis and cancer cell determination. In the STR analysis, 16 STR markers of CR1520 were completely consistent with those of cancer tissue (Figure 2), demonstrating that the cultured cells were derived from the cancer tissue. We determined the 13 cultured cells as cancer cells based on the results of *hTERT* or *MAGE* A1-6 RT-PCR and STR analysis.

The positive rates of hTERT and MAGE A1-6 RT-PCR according to cancer stages. The association of RT-PCR positive cases with cancer stages were also analyzed (Table IV). Though the positive rates of RT-PCR were not statistically significantly associated with cancer stage, the positive rates of *hTERT* and *MAGE* RT-PCR were associated with increased cancer stage. The *hTERT* RT-PCR was the most efficient in identifying cancer cells in the CRC culture.

Discussion

CTCs are rare in the blood. Many methods have been used to increase CTC capture rates. These include immunomagnetic separation (22), microfluidic separation (23), microfiltration (24), electrophoresis (25), and acoustophoresis (26). Bobek *et al.* (27) and Zhang *et al.* (10) reported 66.7% and 73.7% detection rates from pancreatic and lung cancer patients,

Table III. Demographic data and circulating tumor cell culture results.

Classifications	Colorectal cancer	Gastric cancer	Total
Number	56	25	81
Age (Mean±SD)	68.4±11.8	64.8±13.6	67.3±12.4
Male:Female	35:21	17:8	52:29
Cancer Stage			
Stage I	10	13	23
Stage II	22	5	37
Stage III	20	6	26
Stage IV	4	1	5
CTC culture	N of Positive (%)		
Visible cells	23 (41.1)	15 (60.0)	38 (46.9)
Colony formation	3 (5.3)	2 (8.0)	5 (6.2)
<i>hTERT</i> RT-PCR	9 (16.1)	4 (16.0)	13 (16.0)
<i>MAGE</i> RT-PCR	3 (5.4)	1 (4.0)	4 (4.9)
<i>hTERT</i> or <i>MAGE</i> RT-PCR	9 (16.1)	4 (16.0)	13 (16.0)

respectively. Zhang *et al.* (10) successfully isolated CTCs from 14 of 19 early-stage lung cancer patients using a microfluidic three-dimensional co-culture system.

In this study, we did not use any special CTC isolation kit and culture device. We collected mononuclear cells after red blood cells lysis without the depletion of CD45 positive leukocytes to avoid cancer cell loss. The collected cells were inoculated into the CRC culture plate using a fibroblast feeder layer. The CRC culture system effectively supports the growth of normal and tumor cells of epithelial origin (15, 28).

We evaluated the sensitivity of the CRC culture method. In a prior study (15), the minimum number of primary keratinocytes capable of forming a colony was four. Presently, the minimum number was similar (six) but the cell number depended on the cell type. The results indicate the potential of the CRC culture method for the study of CTCs.

To determine whether the cultured cells were cancer cells, we assayed for *MAGE* and *hTERT* mRNA. Though the *hTERT* mRNA might be expressed in the cells of the CRC culture (15), its expression could differentiate human cancer cells from mouse fibroblast J2 cells. The results regarding *MAGE* mRNA revealed that this gene is a cancer specific oncogene (17, 29).

The CRC method allowed the growth of some cells from the 38 blood samples. Among them, 13 samples showed the expression of *MAGE* or *hTERT* mRNA. Using RT-PCR, the *hTERT* mRNA was more frequently amplified than the *MAGE* A1-6 mRNA. The general expression rates of *hTERT* and *MAGE* A mRNA in cancer cells were 85% (30) and 50% (31), respectively. This could explain the difference in RT-PCR positive rates between *hTERT* and *MAGE* A1-6

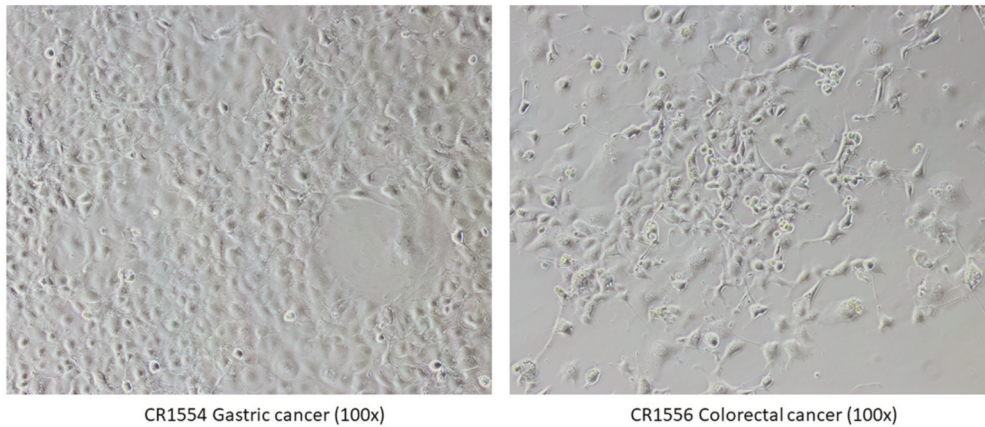


Figure 1. Light microscopic images of CR1554 and CR1556 cancer cells using conditionally reprogrammed cell culture.

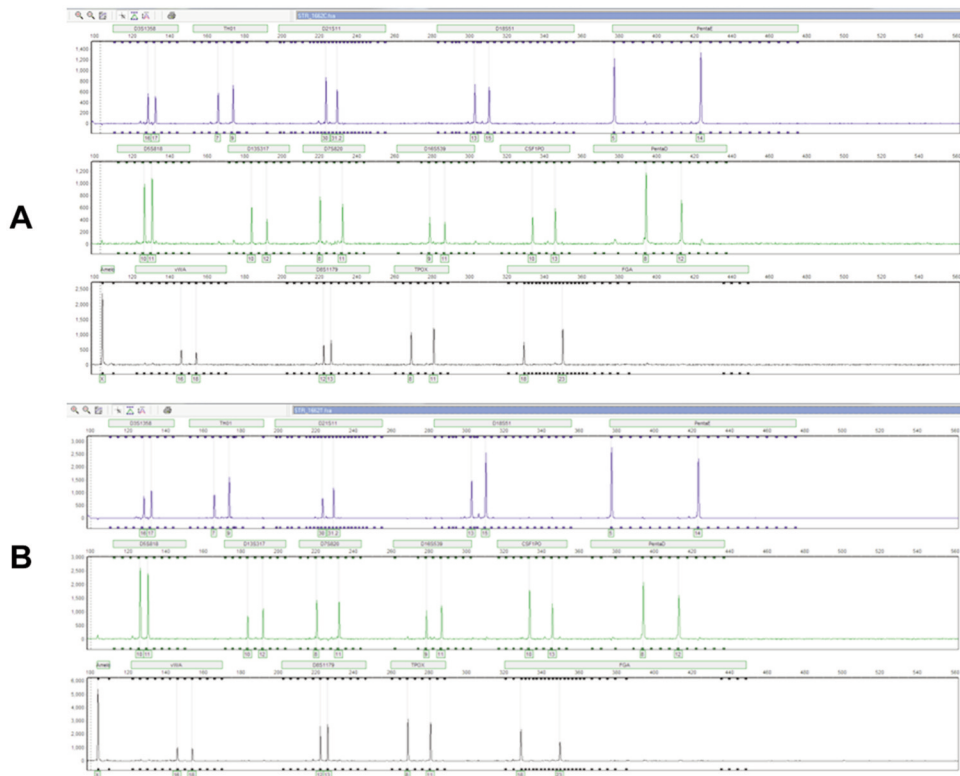


Figure 2. STR marker analysis of CR1520 disclosed the same pattern between cultured cell (A) and cancer tissue (B).

mRNAs. The *hTERT* mRNA expressed in all cancer cells and its positive rates were dependent on cancer stage. Therefore, *hTERT* RT-PCR might be the best marker for the identification of cancer cells in the CRC culture.

After one-year follow-up of the 13 patients with a positive culture, two had multiple metastases (CR1554 and CR1560) and four had metastases to adjacent organs (CR 1600,

CR1577, CR1546 and CR1632). Further studies are necessary to evaluate the clinical utility of CRC culture.

In the one case of positive culture, we isolated the tumor DNA from remnant of RNA extraction and analyzed the STR markers. In the CRC specimens, we could analyze DNA, RNA and protein levels. The CRC specimens can provide most of the information needed for cancer research.

Table IV. Positive rates of RT-PCR results by cancer stage.

	Stage 1	Stage 2	Stage 3-4	p-Value ^a
Number	23	27	31	
hTERT RT-PCR	2 (8.7)	3 (11.1)	8 (25.8)	0.165
MAGE RT-PCR	0 (0.0)	1 (3.7)	3 (9.7)	0.251

^aChi-squared test.

Therefore, the CRC culture might be the best method to study the metastatic event of cancer cells in the circulation.

The absence of a control group for comparison is a limitation of this study. The majority of circulating tumor cell studies using peripheral blood samples of cancer patients have not used a control group (27).

In summary, to isolate CTCs, we cultured blood cells of 81 cancer patients using the CRC culture method. The 38 cases displayed visible cells after 4 weeks. We characterized the culture cells using RT-PCR to detect *hTERT* and *MAGE A1-6* mRNA. Of these, 13 cases showed cancer cell growth. Our study proved the potential of CRC culture to detect CTCs.

Conflicts of Interest

All Authors have no conflicts of interest to declare regarding this study.

Authors' Contributions

Chun-Seok Yang and Dae-Dong Kim collected blood samples and clinical information from the patients of colorectal cancer. In-Hwan Kim and Hyun-Dong Chae collected blood samples and clinical information from the patients of colorectal cancer. Chun-Seok Yang wrote the article. Chang-Ho Jeon performed the entire experiment, collected the experimental data, and revised the version to be submitted. All Authors designed the concept of this study and analyzed the experimental results.

Acknowledgements

Supported by Basic Science Research Program through the National Research Foundation of Korea (NRF) funded by the Ministry of Education, Science and Technology (NRF-2013R1A1A2007189).

References

- 1 Ferreira MM, Ramani VC and Jeffrey SS: Circulating tumor cell technologies. *Mol Oncol* 10(3): 374-394, 2016. PMID: 26897752. DOI: 10.1016/j.molonc.2016.01.007.
- 2 Ilie M, Szafer-Glusman E, Hofman V, Long-Mira E, Suttman R, Darbonne W, Butori C, Lallée S, Fayada J, Selva E, Yu W, Marquette CH, Shames DS, Punnoose E and Hofman P: Expression of MET in circulating tumor cells correlates with expression in tumor tissue from advanced-stage lung cancer patients. *Oncotarget* 8(16): 26112-26121, 2017. PMID: 28212540. DOI: 10.18632/oncotarget.15345
- 3 Masuda T, Hayashi N, Iguchi T, Ito S, Eguchi H and Mimori K: Clinical and biological significance of circulating tumor cells in cancer. *Mol Oncol* 10(3): 408-417, 2016. PMID: 26899533. DOI: 10.1016/j.molonc.2016.01.010
- 4 Szilágyi M, Pös O, Márton É, Buglyó G, Soltész B, Keserű J, Penyige A, Szemes T and Nagy B: Circulating cell-free nucleic acids: Main characteristics and clinical application. *Int J Mol Sci* 21(18): 6827, 2020. PMID: 32957662. DOI: 10.3390/ijms21186827
- 5 Stewart CM and Tsui DWY: Circulating cell-free DNA for non-invasive cancer management. *Cancer Genet* 228-229: 169-179, 2018. PMID: 29625863. DOI: 10.1016/j.cancergen.2018.02.005
- 6 Jiang T, Ren S and Zhou C: Role of circulating-tumor DNA analysis in non-small cell lung cancer. *Lung Cancer* 90(2): 128-134, 2015. PMID: 26415994. DOI: 10.1016/j.lungcan.2015.09.013
- 7 Brock G, Castellanos-Rizaldos E, Hu L, Coticchia C and Skog J: Liquid biopsy for cancer screening, patient stratification and monitoring. *Transl Cancer Res* 4(3): 280-290, 2015. DOI: 10.3978/j.issn.2218-676X.2015.06.05
- 8 Kolostova K, Zhang Y, Hoffman RM and Bobek V: In vitro culture and characterization of human lung cancer circulating tumor cells isolated by size exclusion from an orthotopic nude-mouse model expressing fluorescent protein. *J Fluoresc* 24(5): 1531-1536, 2014. PMID: 25141982. DOI: 10.1007/s10895-014-1439-3
- 9 Halo TL, McMahon KM, Angeloni NL, Xu Y, Wang W, Chinen AB, Malin D, Strekalova E, Cryns VL, Cheng C, Mirkin CA and Thaxton CS: NanoFlares for the detection, isolation, and culture of live tumor cells from human blood. *Proc Natl Acad Sci U S A* 111(48): 17104-17109, 2014. PMID: 25404304. DOI: 10.1073/pnas.1418637111
- 10 Zhang Z, Shiratsuchi H, Lin J, Chen G, Reddy RM, Azizi E, Fouladdel S, Chang AC, Lin L, Jiang H, Waghay M, Luker G, Simeone DM, Wicha MS, Beer DG, Ramnath N and Nagrath S: Expansion of CTCs from early stage lung cancer patients using a microfluidic co-culture model. *Oncotarget* 5(23): 12383-12397, 2014. PMID: 25474037. DOI: 10.18632/oncotarget.2592
- 11 Kolostova K, Matkowski R, Gürlich R, Grabowski K, Soter K, Lischke R, Schützner J and Bobek V: Detection and cultivation of circulating tumor cells in gastric cancer. *Cytotechnology* 68(4): 1095-1102, 2016. PMID: 25862542. DOI: 10.1007/s10616-015-9866-9
- 12 Li Y, Zhang X, Ge S, Gao J, Gong J, Lu M, Zhang Q, Cao Y, Wang DD, Lin PP and Shen L: Clinical significance of phenotyping and karyotyping of circulating tumor cells in patients with advanced gastric cancer. *Oncotarget* 5(16): 6594-6602, 2014. PMID: 25026283. DOI: 10.18632/oncotarget.2175
- 13 Grillet F, Bayet E, Villeronce O, Zappia L, Lagerqvist EL, Lunke S, Charafe-Jauffret E, Pham K, Molck C, Rolland N, Bourgaux JF, Prudhomme M, Philippe C, Bravo S, Boyer JC, Canterel-Thouennon L, Taylor GR, Hsu A, Pascucci JM, Hollande F and Pannequin J: Circulating tumour cells from patients with colorectal cancer have cancer stem cell hallmarks in *ex vivo* culture. *Gut* 66(10): 1802-1810, 2017. PMID: 27456153. DOI: 10.1136/gutjnl-2016-311447
- 14 Kaifi JT, Kunkel M, Dicker DT, Joude J, Allen JE, Das A, Zhu J, Yang Z, Sarwani NE, Li G, Staveley-O'Carroll KF and El-Deiry WS: Circulating tumor cell levels are elevated in colorectal cancer patients with high tumor burden in the liver. *Cancer Biol Ther* 16(5): 690-698, 2015. PMID: 25785486. DOI: 10.1080/15384047.2015.1026508

- 15 Liu X, Ory V, Chapman S, Yuan H, Albanese C, Kallakury B, Timofeeva OA, Nealon C, Dakic A, Simic V, Haddad BR, Rhim JS, Dritschilo A, Riegel A, McBride A and Schlegel R: ROCK inhibitor and feeder cells induce the conditional reprogramming of epithelial cells. *Am J Pathol* 180(2): 599-607, 2012. PMID: 22189618. DOI: 10.1016/j.ajpath.2011.10.036
- 16 Sang M, Wu X, Fan X, Sang M, Zhou X and Zhou N: Multiple MAGE-A genes as surveillance marker for the detection of circulating tumor cells in patients with ovarian cancer. *Biomarkers* 19(1): 34-42, 2014. PMID: 24320162. DOI: 10.3109/1354750X.2013.865275
- 17 Joosse SA, Müller V, Steinbach B, Pantel K and Schwarzenbach H: Circulating cell-free cancer-testis MAGE-A RNA, BORIS RNA, let-7b and miR-202 in the blood of patients with breast cancer and benign breast diseases. *Br J Cancer* 111(5): 909-917, 2014. PMID: 24983365. DOI: 10.1038/bjc.2014.360.
- 18 Jheon S, Hyun DS, Lee SC, Yoon GS, Jeon CH, Park JW, Park CK, Jung MH, Lee KD and Chang HK: Lung cancer detection by a RT-nested PCR using MAGE A1—6 common primers. *Lung Cancer* 43(1): 29-37, 2004. PMID: 14698534. DOI: 10.1016/j.lungcan.2003.08.014
- 19 Kwon S, Kang SH, Ro J, Jeon CH, Park JW and Lee ES: The melanoma antigen gene as a surveillance marker for the detection of circulating tumor cells in patients with breast carcinoma. *Cancer* 104(2): 251-256, 2005. PMID: 15937912. DOI: 10.1002/cncr.21162
- 20 Ping B, Tsuno S, Wang X, Ishihara Y, Yamashita T, Miura K, Miyoshi F, Shinohara Y, Matsuki T, Tanabe Y, Tanaka N, Ogawa T, Shiota G and Miura N: Comparative study of ¹⁸F-FDG-PET/CT imaging and serum hTERT mRNA quantification in cancer diagnosis. *Cancer Med* 4(10): 1603-1611, 2015. PMID: 26275387. DOI: 10.1002/cam4.508
- 21 Chebel A, Rouault JP, Urbanowicz I, Baseggio L, Chien WW, Salles G and Ffrench M: Transcriptional activation of hTERT, the human telomerase reverse transcriptase, by nuclear factor of activated T cells. *J Biol Chem* 284(51): 35725-35734, 2009. PMID: 19843528. DOI: 10.1074/jbc.M109.009183
- 22 Andreopoulou E, Yang LY, Rangel KM, Reuben JM, Hsu L, Krishnamurthy S, Valero V, Fritsche HA and Cristofanilli M: Comparison of assay methods for detection of circulating tumor cells in metastatic breast cancer: AdnaGen AdnaTest BreastCancer Select/Detect™ versus Veridex CellSearch™ system. *Int J Cancer* 130(7): 1590-1597, 2012. PMID: 21469140. DOI: 10.1002/ijc.26111
- 23 Galletti G, Sung MS, Vahdat LT, Shah MA, Santana SM, Altavilla G, Kirby BJ and Giannakakou P: Isolation of breast cancer and gastric cancer circulating tumor cells by use of an anti HER2-based microfluidic device. *Lab Chip* 14(1): 147-156, 2014. PMID: 24202699. DOI: 10.1039/c3lc51039e
- 24 Kolostova K, Cegan M and Bobek V: Circulating tumour cells in patients with urothelial tumours: Enrichment and *in vitro* culture. *Can Urol Assoc J* 8(9-10): E715-E720, 2014. PMID: 25408812. DOI: 10.5489/cuaj.1978
- 25 Peeters DJ, De Laere B, Van den Eynden GG, Van Laere SJ, Rothé F, Ignatiadis M, Sieuwerts AM, Lambrechts D, Rutten A, van Dam PA, Pauwels P, Peeters M, Vermeulen PB and Dirix LY: Semiautomated isolation and molecular characterisation of single or highly purified tumour cells from CellSearch enriched blood samples using dielectrophoretic cell sorting. *Br J Cancer* 108(6): 1358-1367, 2013. PMID: 23470469. DOI: 10.1038/bjc.2013.92
- 26 Antfolk M, Antfolk C, Lilja H, Laurell T and Augustsson P: A single inlet two-stage acoustophoresis chip enabling tumor cell enrichment from white blood cells. *Lab Chip* 15(9): 2102-2109, 2015. PMID: 25824937. DOI: 10.1039/c5lc00078e
- 27 Bobek V, Gurlich R, Eliasova P and Kolostova K: Circulating tumor cells in pancreatic cancer patients: enrichment and cultivation. *World J Gastroenterol* 20(45): 17163-17170, 2014. PMID: 25493031. DOI: 10.3748/wjg.v20.i45.17163
- 28 Supryniewicz FA, Upadhyay G, Krawczyk E, Kramer SC, Hebert JD, Liu X, Yuan H, Chelvaraju C, Clapp PW, Boucher RC Jr, Kamonjoh CM, Randell SH and Schlegel R: Conditionally reprogrammed cells represent a stem-like state of adult epithelial cells. *Proc Natl Acad Sci U S A* 109(49): 20035-20040, 2012. PMID: 23169653. DOI: 10.1073/pnas.1213241109
- 29 Szatanek R, Drabik G, Baran J, Kolodziejczyk P, Kulig J, Stachura J and Zembala M: Detection of isolated tumour cells in the blood and bone marrow of patients with gastric cancer by combined sorting, isolation and determination of MAGE-1, -2 mRNA expression. *Oncol Rep* 19(4): 1055-1060, 2008. PMID: 18357396.
- 30 Ivancich M, Schrank Z, Wojdyla L, Leviskas B, Kuckovic A, Sanjali A and Puri N: Treating cancer by targeting telomeres and telomerase. *Antioxidants (Basel)* 6(1): 15, 2017. PMID: 28218725. DOI: 10.3390/antiox6010015
- 31 Kerkar SP, Wang ZF, Lasota J, Park T, Patel K, Groh E, Rosenberg SA and Miettinen MM: MAGE-A is more highly expressed than NY-ESO-1 in a systematic immunohistochemical analysis of 3668 cases. *J Immunother* 39(4): 181-187, 2016. PMID: 27070449. DOI: 10.1097/CJI.0000000000000119

Received January 8, 2021

Revised February 25, 2021

Accepted February 26, 2021