Abstract. Aim: The present study investigated how well the results of integrative tumor-response assay (ITRA) compared to those of clinical response to chemotherapy in patients with metastatic colorectal cancer (CRC). Patients and Methods: A total of 129 patients with metastatic CRC were prospectively enrolled. ITRA consisted of two sequential histoculture drug-response assays (HDRA). First-stage HDRA was performed using 5-fluorouracil with leucovorin and oxaliplatin (FX), or with irinotecan (FR). Second-stage HDRA (ITRA) was performed for cells surviving after the first-stage HDRA, using FX, FR, and their combinations with bevacizumab and cetuximab. Results: Among 129 patients, 42 (32.6%) completed second-line chemotherapy, results that correlated with those of ITRA. The accuracy of ITRA for predicting response to second-line chemotherapy was 61.9% (26/42), with a sensitivity of 44.4% (8/18) and a specificity of 75% (18/24). Conclusion: Despite its relatively low accuracy, ITRA might be a useful technique for predicting therapeutic efficacy and selecting for appropriate first-line and second-line anticancer regimens for patients.

Recent trends in cancer treatment involve tailored medicine aimed to maximize individual responsiveness. The drug-response assay has been developed to increase therapeutic efficacy and reduce anticancer drug toxicity. The histoculture drug-response assay (HDRA) is representative of this effort. It is a methyl thiazolyl-diphenyl-tetrazolium bromide (MTT) assay with the advantage of more correctly reflecting the in vivo microenvironment using three-dimensional multicellular spheroids in tissue culture (1). Previous studies showed 66-92% accuracy of this assay in terms of correlation with clinical response to chemotherapy for solid tumors, including colorectal cancer (CRC) (2-5).

Approximately 20% of patients with CRC have stage IV disease. Of these, more than 75% have unresectable metastases and therefore need palliative treatment (6, 7). Most cases are incurable and have gradual progression of disease in spite of palliative chemotherapy. Our previous study demonstrated that HDRA was useful for predicting therapeutic efficacy and selecting the appropriate chemotherapy regimen for patients with metastatic CRC (5). However, an innate limitation of HDRA is that it is an in vitro assay devoid of the tumor microenvironment. Moreover, in cases of recurrent CRC, there is no usable assay to predict sensitivity to second-line chemotherapy.

We developed a novel in vitro chemosensitivity assay based on HDRA, the integrative tumor-response assay (ITRA, Patent No. 10-1235820; the Korean Intellectual Property Office, Seoul, Korea). It simultaneously predicts responsiveness to both first- and second-line chemotherapy. We investigated the chemosensitivity to clinically available regimens by ITRA using primary tumor tissues from patients with metastatic CRC and compared the results with the patients’ actual clinical responses.

Patients and Methods

Patient selection. A total of 129 patients with metastatic CRC were prospectively enrolled. All patients underwent palliative surgery between December 2011 and December 2013. Eligibility criteria included histologically-proven colorectal adenocarcinoma, Eastern Cooperative Oncology Group performance status ≤2, age ≥80 years,
and expectation of palliative chemotherapy. Patients were excluded if they had received preoperative chemotherapy or radiotherapy. We excluded nine cases in which ITRA results were incomplete because of insufficient tumor samples. Baseline demographic and clinicopathological characteristics are described in Table I. The median follow-up period was 28 months (range=0-64 months). By the end of the study period, only 42 patients had undergone second-line chemotherapy using regimens that were evaluated with ITRA. The outcomes of those who had both first-line and second-line chemotherapy using tested drug regimens were evaluated for correlation with ITRA results. ITRA results were correlated with clinical response (Figure 1). The study protocol was approved by the Institutional Review Board of Asan Medical Center (Registration No: 2012-0222), in accordance with the Declaration of Helsinki.

Investigated drug combinations and concentrations. The regimens tested included 5-fluorouracil (5-FU) with leucovorin and oxaliplatin (FX), 5-fluorouracil with leucovorin and irinotecan (FR), and these combinations with biologically targeted drugs. First-line chemotherapy regimens were FX or FR and second-line regimens were FX, FR, or one of these combined with a targeted agent (Table II). The targeted drugs used were bevacizumab (Avastin®; Roche, Basel, Switzerland) and cetuximab (Erbitux®; Merck, Munich, Germany). The combined regimens were FX with bevacizumab or cetuximab, and FR with bevacizumab or cetuximab. The doses initially used were based on the usual clinical dose as determined from pharmacokinetic and pharmacodynamic parameters obtained through phase I studies and empirical assays (8, 9). Median inhibitory concentrations (IC50) of tested drugs were determined as those causing a 50% dose-dependent reduction in cell viability in complete growth medium after 48 h of continuous exposure to the drug. The cutoff concentrations of the six anticancer drugs used to determine in vitro sensitivity and resistance were 50 μg/ml for 5-FU, 10 μg/ml for leucovorin, 20 μg/ml for irinotecan, 40 μg/ml for oxaliplatin, 20 μg/ml for bevacizumab, and 20 μg/ml for cetuximab, the same as in our previous study of HDRA (5).

Integrated tumor-response assay. ITRA was theoretically based on the HDRA procedure, but in vitro evaluation of resistance to the first-line anticancer drugs was sequentially repeated. In the following description of the procedure, the first three steps for ITRA were similar to those for HDRA. The details for HDRA were the same as in our previous work, except for the tumor volume (5, 10). The final two steps were specific for ITRA.

(i) Preparation of tissue samples. At least two sections of tumor tissue (1 cm in diameter) were obtained from fresh surgically resected specimens. This was a larger volume than we had previously collected for HDRA only. Tumor samples were aseptically washed in Hank’s balanced salt solution (HBSS; Gibco, Gaithersburg, MD, USA). Collagen sponge gels manufactured from pig skin (Gelfoam®) were purchased from Pharmacia & Upjohn (Kalamazoo, MI, USA). The cancerous portions of the specimen were minced with scissors into pieces approximately 1 mm in diameter. Necrotic and non-viable portions were excluded using 3-(4,5-dimethylthiazol-2-yl)-5-(3-carboxymethoxyphenyl)-2-(4-sulfopheny1)-2-tetrazolum (MTS; Sigma, St. Louis, MO, USA) staining. Viable cancer tissues were weighed on a chemical balance and placed on the prepared collagen gel surfaces (Gelfoam®, Pharmacia & Upjohn) in 96-well plates (Figure 2).
(ii) Tissue culture with first-line anticancer drugs. In this study, first-line chemotherapy regimens were confined to FX and FR. Three replicates were concurrently run for the control and treatment groups. The first-line anticancer drugs were dissolved in RPMI-1640 (Gibco) medium containing 10% fetal bovine serum (FBS, Biowest, Nuaille, France) in a humidified atmosphere containing 95% air and 5% CO₂. The drug solutions were added to each well (Figure 2A), and the plates were incubated for 3 days at 37°C.

(iii) Determination of sensitivity to first-line drugs. After histoculture, 900 μl of HBSS containing 0.1-mg/ml collagenase (type I; Sigma) and 100 μl of 0.2% MTT (Sigma) solution dissolved in phosphate-buffered saline was added to each culture well. Plates were incubated for another 4 h, the medium removed, and 0.5 ml of dimethyl sulfoxide (Sigma) was added to each well to extract MTT formazan. Extracts from each well (100 μl) were transferred to a 96-well plate and the absorbance was measured at 540 nm using a microplate reader (VersaMax, Sunnyvale, CA, USA). The rate of inhibition of cancer-cell viability (IR) was calculated using the following equation: IR (%) = (1 − mean absorbance of treated wells per gram of tumor/mean absorbance of control wells per gram of tumor) x100. The IR cutoff value for a positive response was previously determined to be ≥30% (11, 12) and this was used the present study.

(iv) Tissue culture with second-line anticancer drugs. After the measurement of IR for first-line drug regimens (HDRA), the culture media and first-line drugs were removed. Three replicates were concurrently run for the control and treatment groups. Second-line anticancer drugs were dissolved in RPMI-1640 medium (Gibco) containing 10% FBS (Biowest) in a humidified atmosphere containing 95% air and 5% CO₂. The drug solutions were added to each well (Figure 2B), and the plates were incubated for another 2 days at 37°C.

(v) Determination of sensitivity to second-line drugs. After the second histoculture, the IR for each drug regimen was measured in the same manner with an MTT end-point assay. An exemplary final ITRA report is shown in Figure 3.

Evaluation of tumor response. Each patient’s response to treatment was assessed every three or four chemotherapy cycles with consistent imaging techniques such as computed tomography or magnetic resonance imaging. According to the Revised RECIST (Response Evaluation Criteria in Solid Tumors) guideline (version 1.1, 2009) (13), complete response (CR) was defined as the disappearance of all metastatic lesions and a partial response (PR) as a reduction of at least 30% of the sum of the longest diameter of metastatic lesions, with no evidence of new lesions. Progressive disease (PD) was defined as an increase by at least 20% of the sum of the longest diameters of either metastatic or recurrent lesions. Stable disease was reported when the criteria for either PR or PD were not met. The primary endpoint was a correlation between the ITRA results and the clinical response in response rate (RR); this was defined as positive for tumor responses of CR or PR.

Statistical analysis. A cross-table analysis employing Pearson’s chi-square test or Fisher’s exact test, as appropriate, was used to compare categorical variables. A p-value of 0.05 was considered statistically significant. All calculations were carried out using SPSS software (version 21.0; IBM Corp., Armonk, NY, USA).

Results

ITRA tumor growth-inhibition ratios. Of 120 evaluated samples, IR with first-line regimens for FR and FX were 36% and 32.6%, respectively. IRs for second-line regimens were lower than those of first-line regimens (Figure 4A). The IR for first-line FR was significantly better than that for FX (p<0.001). The differences in IRs between first- and second-line regimens were significant for FR (p<0.001), but not for FX (p=0.077).

For second-line treatment, switching from FX to FR or from FR to FX and adding a targeted agent resulted in better IRs than adding a targeted drug to the first-line regimen (p<0.001). In terms of targeted agents, adding cetuximab to second-line
regimens resulted in significantly better IR than adding bevacizumab (p<0.001). The percentages of chemosensitive tumors were similar for each regimen (Figure 4B).

Clinical correlation of HDRA and ITRA. The clinical responses of 120 patients treated with first-line regimens are summarized in Table III. The overall RR was 50%. On repeated assessments, however, RR decreased. Out of 79 HDRAs of first-line chemotherapeutic regimens, the correlation with patient clinical response was 70.9% (56/79), with 73.2% (30/41) true-positives and 68.4% (26/38) true-negatives. The sensitivity was 71.4% (30/42) and specificity 70.3% (26/37). For the 42 patients completing second-line chemotherapy, the accuracy of ITRA for predicting response to second-line chemotherapy was 61.9% (Table IV).

We analyzed clinicopathological parameters that might predict better chemoresponsiveness to first- and second-line chemotherapy. Microsatellite instability (p=0.03), poorly differentiated or mucinous histology (p=0.005), and large tumor (size >6 cm, p=0.046) were significantly associated with poor chemoresponsiveness to first-line chemotherapy. Lymphovascular invasion was significantly related to poor chemoresponsiveness to second-line chemotherapy (p=0.049). However, multivariate analysis did not demonstrate any
significant predictor of chemoresponse to first- or second-line chemotherapy. Furthermore, clinicopathological parameters were not correlated with the accuracy of ITRA or HDRA.

Discussion
Pathways of inherent and acquired resistance account for the high rate of anticancer treatment failure. The mechanisms or pathways mediating resistance are explained by changes in intratumoral drug exposure or failure to elicit cytotoxicity (14). In terms of acquired resistance, sufficient exposure to a specific anticancer drug can elicit drug resistance through various pathways. In order to simulate the cellular environment in patients with recurrent CRC, we examined tumor cells that had survived first-line chemotherapy regimens, culturing them with second-line regimens. Our hypothesis on which ITRA was based was that tumor cells surviving after chemotherapy in our *in vitro* assay had equivalent genetic traits to those of *in vivo* recurrent CRC.

Although it was difficult to prove this, our results with ITRA showed a slightly positive correlation with clinical response to second-line chemotherapy.

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**Table III. Treatment response to first- and second-line chemotherapy among patients with colorectal carcinoma.**

<table>
<thead>
<tr>
<th>Clinical response</th>
<th>Total (n=120), n (%)</th>
<th>First-line (n=79), n (%)</th>
<th>Second-line (n=42), n (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td>CR</td>
<td>3 (2)</td>
<td>2</td>
<td>1</td>
</tr>
<tr>
<td>PR</td>
<td>57 (47)</td>
<td>40</td>
<td>17</td>
</tr>
<tr>
<td>SD</td>
<td>41 (34)</td>
<td>26</td>
<td>15</td>
</tr>
<tr>
<td>PD</td>
<td>20 (17)</td>
<td>11</td>
<td>9</td>
</tr>
<tr>
<td>Response rate</td>
<td>50%</td>
<td>53%</td>
<td>43%</td>
</tr>
</tbody>
</table>

CR, Complete response; PR, partial response; SD, stable disease; PD, progressive disease.

**Table IV. Correlation between integrative tumor-response assay (ITRA) and clinical response to second-line chemotherapy in patients with colorectal carcinoma.**

<table>
<thead>
<tr>
<th>ITRA, n</th>
<th>Positive</th>
<th>Negative</th>
<th>Total</th>
</tr>
</thead>
<tbody>
<tr>
<td>Positive</td>
<td>8</td>
<td>6</td>
<td>14</td>
</tr>
<tr>
<td>Negative</td>
<td>10</td>
<td>18</td>
<td>28</td>
</tr>
<tr>
<td>Total</td>
<td>18</td>
<td>24</td>
<td>42</td>
</tr>
</tbody>
</table>

Sensitivity: 44.4% (8/18); specificity: 75% (18/24); true-positive rate: 57.1% (8/14); true-negative rate: 64.3% (18/28); tccuracy: 61.9% (26/42); *complete response or partial response.*
ITRA is not simply a repetition of HDRA. It examines the sensitivity of resistant tumor cells. HDRA requires an increased volume of tumor sample when a greater number of chemotherapy regimens are evaluated. In the present study, we evaluated a total of twelve regimens of combination chemotherapy for all specimens, compared with seven regimens examined by HDRA in our previous work (5). Because ITRA used the same specimen volume (1 cm³) as HDRA, we were unable to perform ITRA in nine cases in the early period of this study. The specimen volume must be adjusted to the number of treatment regimens evaluated. Thus, a limitation of ITRA is that it cannot be performed in cases with small tumors that yield insufficient specimens.

Our twelve chemotherapy regimens were chosen considering every possible sequential combination of agents, assuming that targeted agents were used for second-line chemotherapy. Compared to the effort involved, the efficiency of ITRA was low in cases using targeted agents (6/42). As conditions vary in clinical practice, sequential combinations of chemotherapy regimens to be assayed can be adjusted. At the end of enrollment of the present trial, the Korean government extended economic support for targeted agents, allowing their use as first-line chemotherapy in patients with metastatic CRC. After that, we updated our assay to test seventeen possible sequential combinations of chemotherapy, any of which might be used in clinical practice. In terms of assay efficiency, ITRA might be most useful for the combination chemotherapy regimens of FX and FR without targeted agents.

The monoclonal antibody cetuximab, which targets the epidermal growth factor receptor (EGFR), has expanded the range of treatment options for metastatic CRC.(15) EGFR and downstream signaling pathways are activated by several mechanisms, including overexpression of the receptor, overexpression of the ligand, activation by a receptor mutation, or inactivation of tumor-suppressor genes (16). The presence of wild-type V-Ki-ras2 Kirsten rat sarcoma viral oncogene homolog (KRAS) is considered a predictor of response to EGFR inhibitor therapy (17, 18). The presence of a KRAS mutation is important to choose the appropriate targeted agent, especially for anti-EGFR therapy (19). It may be difficult to use cetuximab according to ITRA results in the presence of mutant form of KRAS. Bevacizumab is a monoclonal antibody against vascular endothelial growth factor (VEGF) and has shown clinical activity against metastatic CRC (20). Although our three-dimensional culture system has the advantage of more accurately reflecting the in vivo microenvironment, the performance of ITRA in vitro is difficult to compare with the in vivo cellular environment as it cannot account for vascular supply. Thus, the value of ITRA for evaluating response to bevacizumab is questionable because angiogenesis cannot be replicated in the in vitro cell-culture system. This may explain why ITRA was less successful in predicting response to the addition of bevacizumab than to that of cetuximab as second-line chemotherapy.

The RRs to first-line chemotherapy with FX and FR for patients with metastatic CRC are 40-50%, which further decrease with subsequent treatment (21). Recent combined chemotherapy using anti-EGFR and anti-VEGF agents enhanced RRs of chemotherapy and improved survival up to 30 months for patients with metastatic CRC (22). Although our ITRA was not highly predictive of response to these agents, it might be helpful in choosing optimal chemotherapy regimens and improving survival of patients with metastatic CRC.

This study has several limitations. We were only able to perform ITRA in one-third of the initially enrolled patients. Reasons for this varied, including patient choice, economic status, whether medical insurance covered the anticancer drugs, and an inadequate sequence of chemotherapy which was different from that of our study protocol. In addition, although ITRA results were positively correlated with clinical drug responses, the accuracy was relatively low. This might be because of the complexity of chemotherapy regimens, low drug concentrations, or a low cutoff value for IR. This is a fundamental limitation of an in vitro assay, as we suggested in our previous investigation of HDRA (5). Furthermore, an innate limitation of ITRA is its inability to test circulating tumor cells or cancer stem cells, which may be particularly resistant to different chemotherapeutic agents (23).

In conclusion, despite its relatively low accuracy, ITRA may be a feasible and useful technique for predicting therapeutic efficacy and selecting appropriate first-line and second-line anticancer regimens for individual patients with metastatic CRC.

Acknowledgements

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References


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