Effects of Delay in the Snap Freezing of Colorectal Cancer Tissues on the Quality of DNA and RNA

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Purpose: The success of basic molecular research using biospecimens strongly depends on the quality of the specimen. In this study, we evaluated the effects of delayed freezing time on the stability of DNA and RNA in fresh frozen tissue from patients with colorectal cancer.

Methods: Tissues were frozen at 10, 30, 60, and 90 minutes after extirpation of colorectal cancer in 20 cases. Absorbance ratio of 260 to 280 nm (A₂₆₀/A₂₈₀) and agarose gel electrophoresis were evaluated. In addition, the RNA integrity number (RIN) was assayed for the analysis of the RNA integrity.

Results: Regardless of delayed freezing time, all DNA and RNA samples revealed A₂₆₀/A₂₈₀ ratios of more than 1.9, and all DNA samples showed a discrete, high-molecular-weight band on agarose gel electrophoresis. The RINs were 7.53 ± 2.04, 6.70 ± 1.88, 6.47 ± 2.58, and 4.22 ± 2.34 at 10, 30, 60, and 90 minutes, respectively. Though the concentration of RNA was not affected by delayed freezing, the RNA integrity was decreased with increasing delayed freezing time.

Conclusion: According to the RIN results, we recommend that the collection of colorectal cancer tissue should be done within 10 minutes for studies requiring RNA of high quality and within 30 minutes for usual RNA studies.

Keywords: Colorectal neoplasms; Tissue banks; DNA; RNA; Quality control

INTRODUCTION

With recent rapid development of genomics and proteomics, as well as increased demands for studies on the usefulness of biological markers, the trend of studies on diseases is changing. As a result of such changes, in the investigation of diseases, the demand for tissues samples reflecting the characteristics of diseases well and for added clinical information is on the increase more and more. Therefore, in several institutions in Korea, as well as other countries, efforts have been made to secure sufficiently good quality tissue samples by operating tissue sample banks. Particularly, in the future, the possibility that investigators could collect tissue samples from several institutions through networks and conduct studies is high; thus, the need to provide good quality tissue samples is higher. Nevertheless, the system for collecting and storing tissue samples has not yet been standardized, and many diverse systems, depending on the conditions at the institutions, exist.

Multiple factors may determine the quality of tissue samples; however, the most important factor is the time spent to collect the tissue samples. During surgical resection of solid tumors, major blood vessels are ligated, which induces tissue ischemia and causes protein degradation and denaturation until the fixation in formalin or in liquid nitrogen. Furthermore, it has been reported that the expression of genes varies with the elapse of the ischemic time after surgical resection and that with the delay of the fixation time, the usefulness of mRNA is reduced [1, 2]. Therefore, good quality tissue samples could be obtained by fixing tumor tissues as soon as possible after surgical resection. The interval from surgical tissue extraction

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to freezing is 30 minutes in the standard operating procedure of Tuba Frost, and the guideline of the MD Anderson Cancer Center recommends that the interval be 10 minutes [3, 4]. Nevertheless, regarding RNA degradation, the standard for the time interval from tissue extraction to tissue freezing is not clear. In this study, through an examination of the change in the stability of DNA and RNA with changing time interval from surgical resection to freezing of colorectal cancer tissues, a time interval that must be observed to provide good quality tissue samples suitable to molecular biological studies is suggested.

**METHODS**

**Collection and storage of fresh frozen colorectal cancer tissues**

From among patients who underwent surgery for colorectal cancer at the Chonbuk National University Hospital from July 2009 to October 2009, the subjects of this study were 20 patients from whom fresh frozen tissues were collected by the Chonbuk National University Hospital (National Biobank of Korea). The subjects were patients who were not treated with chemotherapy or radiation therapy prior to surgery. Consent for the donation of specimens was obtained prior to surgery, and tissue samples were collected only from patients who signed a consent form for the donation of specimens. Tissues extracted in operating rooms were placed in an icebox maintained at 4°C by adding ice in advance and were transported to the gross examination room. After gross examination, tumor tissues were collected in a range that would not affect diagnosis. The tumors were sectioned in the midline, and while starting from the left side and advancing to the right side, 4 consecutive tissues 0.5 × 0.5 × 0.5 cm³ in size were collected (Fig. 1). The tissues collected 10 minutes, 30 minutes, 60 minutes, or 90 minutes after extraction were rapidly frozen with liquid nitrogen by adding it to isopentane chilled in advance and were subsequently stored in a liquid-nitrogen tank.

**Preparation of the mirror image slide of fresh frozen tissues and evaluation**

To examine the adequacy of fresh frozen tissues for use in experiments, we prepared hematoxylin-eosin slides of the mirror image area of fresh frozen tissues, and we evaluated the adequacy of the frozen samples. For the preparation of precise mirror image slides, sections were obtained from tissues sharing the cross section of the area harvested by the fresh frozen tissues. The left end of the tissue surface opposite to tissues frozen after 10 minutes was marked with red ink and paired with the collected tissues, and the side without contact to tissues harvested at each time point was marked with black ink, and during paraffin embedding, the orientation was maintained to allow the side without marking with black ink to be microsected (Fig. 1). Paraffin blocks of mirror image tissue sections were prepared, and their adequacy was evaluated by hematoxylin-eosin staining. Concerning the categories for the evaluation of adequacy and the standard, by referring to the result of the standardization of the Korean human sample management guideline, the quantity of included tumors (less than 50% is not adequate), contamination of tissues (tissues without contamination of other tissues are excellent), the quantity of necrosis (more than 50% area of the entire slide is not suitable), the quantity of extracellular mucin (tissue with a quantity of mucin more than 50% of the entire slide is not suitable), and inflammatory reaction in the interstitial space were evaluated [5].

**Evaluation of the stability of DNA and RNA of fresh frozen tissues**

DNA and RNA were extracted from fresh frozen tissues, and the stability of fresh frozen tissues was evaluated by using a quantitative analysis and an analysis of integrity. Fresh frozen tissues were divided and added to 1.5-mL microcentrifuge tubes, and DNA was extracted by using a QIAamp DNA mini kit (Qiagen, Hilden, Germany) according to the method suggested by the manufacturer. RNA was extracted using the RNeasy protect mini kit (Qiagen, Hilden, Germany) according to the method suggested by the manufacturer. For the quantitative analysis of the extracted DNA and RNA, the ratio of A_{260}/A_{280} was measured through the use of the NanoDrop (Thermo Fisher Scientific Inc, Waltham, MA, USA). Cases with a ratio of A_{260}/A_{280} higher than 1.8 were evaluated as excellent, cases with a ratio between 1.6 and 1.8 were adequate, and cases with a ratio of less than...
1.6 were inadequate. The integrity of DNA was evaluated by using agarose gel electrophoresis, and if distinct high-molecular-weight bands were observed in electrophoresis, it was evaluated to be adequate. The integrity of RNA was evaluated by measuring the 28s and the 18s ribosomal RNA (rRNA) bands shown on agarose gel electrophoresis by using the Multi Gauge V3.0 program (FujiFilm, Tokyo, Japan) and by measuring the RNA integrity number (RIN) through the use of the Agilent 2100 Bioanalyzer (Agilent Technologies, Santa Clara, CA, USA). On electrophoresis, the 28s and the 18s rRNA bands were examined, and the ratio of the 28s to the 18s rRNA band was recorded as a mean ± standard deviation. A RIN higher than 7 was evaluated to be ‘excellent,’ a RIN between 4 and 7 was ‘good,’ and a RIN lower than 4 was ‘bad’ [5-9]. The Agilent 2100 Bioanalyzer is an automated bioanalyzer that applies microfluidics techniques, and it consists of microfluidic chips, gel-filled channels, equipment for classifying voltage-induced molecular weights in the gel-filled channel, and a laser-induced fluorescence detection system. When a small amount of RNA sample is classified according to molecular weight within microfibrillar chips and detected by laser-induced fluorescence detection, the result is presented as electropherograms.

The RIN is calculated by using the “RIN software algorithm” and is shown as a number from 1 to 10. Complete RNA is shown as RIN 10. As RNA is degraded, the RIN becomes lower, and completely degraded RNA is shown as RIN 1 [8]. The RIN is presented as numbers; thus, it has advantages in that it provides objective evaluation data on RNA integrity and suggests the range of the use of RNA; nonetheless, it has the shortcoming of requiring expensive equipment.

Statistical analysis
Statistical analysis was performed by using SPSS ver.15.0 (SPSS Inc., Chicago, IL, USA). The differences in the A260/A280 ratio of DNA, the A260/A280 ratio of RNA, the ratio of the 28s to the 18s rRNA band, and the RIN value between the groups was validated by using the one-way ANOVA test, and a P-value less than 0.05 was considered to be statistically significant.

RESULTS

Evaluation of the adequacy of the collection area of fresh frozen tissues
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Table 1. Histopathologic evaluation of the sampling adequacy for fresh frozen tissue

| No. | Diagnosis               | Tumor size (cm) | Contained tumor volume (%) | Contamination | Necrosis | Extracellular mucin | Stromal inflammation |
|-----|-------------------------|-----------------|----------------------------|---------------|----------|---------------------|----------------------|
| 1   | Adenoca, MD             | 4.1 × 3.3       | 65                         | No            | No       | No                  | Moderate             |
| 2   | Adenoca, WD             | 5.3 × 3.8       | 65                         | No            | No       | No                  | Moderate             |
| 3   | Adenoca, WD             | 10.5 × 6.1      | 80                         | No            | No       | No                  | Mild                 |
| 4   | Adenoca, MD             | 4.1 × 3.2       | 20                         | No            | Yes (30%)| No                  | Mild                 |
| 5   | Adenoca, MD             | 2.6 × 2.2       | 70                         | No            | No       | Yes (5%)            | Moderate             |
| 6   | Adenoca, MD             | 4.8 × 4.5       | 70                         | No            | Yes (5%)| No                  | Severe               |
| 7   | Adenoca, MD             | 4.6 × 3.7       | 80                         | No            | Yes (10%)| Yes (5%)            | Mild                 |
| 8   | Adenoca, MD             | 5.3 × 4.6       | 85                         | No            | Yes (10%)| No                  | Mild                 |
| 9   | Adenoca, WD             | 4.8 × 3.7       | 90                         | No            | No       | No                  | Mild                 |
| 10  | Adenoca, MD             | 4.5 × 3.9       | 85                         | No            | No       | No                  | Mild                 |
| 11  | Adenoca, MD             | 5.3 × 3.2       | 85                         | No            | No       | No                  | Moderate             |
| 12  | Mucinous adenoca        | 9.1 × 6.4       | 40                         | No            | No       | Yes (60%)           | Severe               |
| 13  | Adenoca, MD             | 5.1 × 4.3       | 50                         | No            | No       | No                  | Mild                 |
| 14  | Adenoca, MD             | 6.6 × 5.0       | 80                         | No            | Yes (10%)| No                  | Moderate             |
| 15  | Adenoca, MD             | 6.3 × 4.8       | 60                         | Yes*(25%)     | No       | No                  | Moderate             |
| 16  | Adenoca, MD             | 4.4 × 4.1       | 80                         | No            | No       | Yes (30%)           | Moderate             |
| 17  | Adenoca, WD             | 2.1 × 1.4       | 60                         | Yes*(25%)     | No       | Yes (5%)            | Mild                 |
| 18  | Adenoca, WD             | 9.2 × 6.3       | 95                         | No            | No       | Yes (10%)           | Moderate             |
| 19  | Adenoca, WD             | 5.6 × 4.3       | 85                         | No            | No       | No                  | Mild                 |
| 20  | Mucinous adenoca        | 2.9 × 2.1       | 65                         | Yes*(20%)     | No       | Yes (55%)           | Mild                 |

Adenoca, adenocarcinoma; MD, moderately differentiated; WD, well differentiated.

*Presence of normal mucosal tissue.
lected by the fresh frozen tissues are shown in Table 1. Two samples (10%) contained less than 50% tumor cells, 3 samples (15%) showed tissue contamination, no samples had more than 50% necrosis, 2 samples (10%) showed 50% extracellular mucus, and 2 samples (10%) showed a severe interstitial inflammatory reaction. The results for the adequacy of the area collected by the fresh frozen tissues show that in a total 20 cases of colorectal cancer samples, 15 cases (75%) were determined to be adequate. Even for samples for which the collection area was not adequate, this was thought not to affect the stability of the fresh frozen tissues; thus, the stabilities of DNA and RNA of all 20 cases were evaluated.

Evaluation of the stability of DNA for fresh frozen tissues
DNA was extracted from fresh frozen tissues, and the A_{260}/A_{280} ratio was measured. The ratio for tissues frozen 10 minutes after extraction was 1.95 ± 0.04, that for tissues frozen 30 minutes after extraction was 1.92 ± 0.06, that for tissues frozen after 60 minutes was 1.93 ± 0.05, and that for tissues frozen 90 minutes after extraction was 1.92 ± 0.05 (Fig. 2A); no statistical differences were detected (P = 0.258). Regardless of the time interval from extraction to freezing, the A_{260}/A_{280} ratio of all tissues was higher than 1.8 and this ratio was determined to be ‘excellent’. In agarose gel electrophoresis of DNA, all tissues showed distinct high-molecular-weight bands; thus, regardless of the elapsed time from extraction to freezing, the integrity of all tissues was evaluated to be ‘adequate’ (Fig. 2B).

Evaluation of the stability of RNA for fresh frozen tissues
RNA was extracted from fresh frozen tissues, and the A_{260}/A_{280} ratio was measured. The 28s/18s ratio gradually decrease with increasing delayed freezing time. (Fig. 3C)

**Fig. 2.** Purity and integrity of DNA in 20 cases of colorectal carcinoma tissues. (A) The means of the A_{260}/A_{280} ratio are over 1.8 regardless of delayed freezing time, and all DNA samples are judged excellent. (B) In agarose gel electrophoresis, all cases of DNA samples reveal a discrete high molecular weight band. In each case, the lanes are in order of delayed time 10, 30, 60, and 90 minutes, respectively.

**Fig. 3.** Purity and integrity of RNA in 20 cases of colorectal carcinoma tissues. (A) The means of the A_{260}/A_{280} ratio are over 1.8 regardless of delayed freezing time, and all RNA samples are judged excellent. (B) In agarose gel electrophoresis, 28s and 18s ribosomal RNA bands are shown. In each case, the lanes are in order of delayed time 10, 30, 60, and 90 minutes, respectively. (C) The means of the 28s/18s ratio gradually decrease with increasing delayed freezing time.
ratio was measured. The ratio of tissues frozen 10 minutes after extraction was 2.08 ± 0.03, that of tissues frozen 30 minutes after extraction was 2.07 ± 0.04, that of tissues frozen 60 minutes after extraction was 2.07 ± 0.03, and that of tissues frozen 90 minutes after extraction was 2.06 ± 0.04 (Fig. 3A); no statistical differences were detected (P = 0.182). Regardless of the time from extraction to freezing, the A_{260}/A_{280} ratio of all tissues was higher than 1.8, and this ratio was determined to be ‘excellent’. In RNA agarose gel electrophoresis, in tissues frozen 10 minutes after extraction, the 28S and the 18S rRNA bands were observed in 19 cases, and the bands could not be distinguished in 1 case. In tissues frozen 30 minutes and 60 minutes after extraction, in all 20 cases, the 28S and the 18S rRNA bands were observed. In tissues frozen 90 minutes after extraction, the 28S and the 18S rRNA band were observed in 16 cases, and the bands could not be distinguished in 4 cases (Fig. 3B). The ratio of the 28S to the 18S rRNA band of samples frozen 10 minutes after extraction was 1.80 ± 0.68, that of tissues frozen 30 after extraction was 1.68 ± 0.52, that of tissues frozen 60 minutes after extraction was 1.60 ± 0.37, and that of tissues frozen 90 minutes after extraction was 1.34 ± 0.57 (Fig. 3C); no statistical differences were observed. Nonetheless, in tissues frozen 10 minutes after extraction, the ratio of the 28S to the 18S rRNA band showed a tendency to be higher in comparison with tissues frozen 90 minutes after extraction (P = 0.058).

The RIN of tissues frozen 10 minutes after extraction was 7.53 ± 2.04, that of tissues frozen 30 minutes after extraction was 6.70 ± 1.88, that of tissues frozen 60 minutes after extraction was 6.47 ± 2.58, and that of tissues frozen 90 minutes after extraction was 4.22 ± 2.34 (Fig. 4A). The RIN according to the interval from extraction to freezing was statistically different. The RIN of tissues frozen 10 minutes after extraction was higher than that of tissues frozen 90 minutes after extraction (P = 0.000), the RIN of tissues frozen 30 minutes after extraction was also higher than that of tissues frozen 90 minutes after extraction (P = 0.004), and the RIN of tissues frozen 60 minutes after extraction was also higher than that of tissues frozen 90 minutes after extraction (P = 0.011). On the other hand, for each time interval, 10, 30, 60, and 90 minutes, the numbers of ‘excellent’ specimen were 15 cases, 8 cases, 12 cases, and 2 cases, respectively. The numbers of ‘good’ specimens was 3, 11, 3, and 5, respectively. The numbers of ‘poor’ specimens were 2, 1, 5, and 13, respectively (Fig. 4B).

**DISCUSSION**

In our study, when fresh frozen tissues were collected from colorectal cancer, the adequacy of the harvested area was evaluated, and according to the interval from surgical extraction to freezing, the change in the stability of DNA and RNA was evaluated. Since fresh frozen tissues are used primarily for the extraction of DNA, RNA and protein, an evaluation of the adequacy of frozen tissues, such as whether the tissues harvested from tumors reflect the characteristics of the harvested area, whether the tissues are contaminated, etc., must be performed prior to the use of the tissue samples.

The mirror image tissues adjacent to the fresh frozen tissues were embedded in paraffin, prepared as slides, and read under light microscope and the adequacy of the harvested area was evaluated. Among 20 cases, 15 cases were adequate, and 5 cases were inadequate. It is a level comparable to other reports. Sandusky et al. [10] reported that among 480 specimens collected from various cancer tissues, samples containing less than 65% tumor components were 25%, and particularly, in colorectal cancer tissues, specimens containing less than 65% tumor components were 32.2%. In our study, two specimens had a quantity of tumor cells less than 50% and were, thus, determined to be inadequate. Among them, one contained substantially more interstitial components than tumor cells and it was an adenocarcinoma associated with substantial necrosis (case 4). The other case was a mucinous carcinoma (case 12). A mucinous carcinoma is defined as a tumor that has more than a 50% mucinous component; thus, in regard to the case, whether it is reasonable to determine the case as inadequate because the
quantity of tumor cells is small must be addressed.

Three samples were determined to be inadequate because of the contamination of tissues, and all of them contained more than 20% of normal mucosal tissues. Among them, the long diameters of the tumors of 2 cases was smaller than 3 cm; thus, in comparison with other cases, the size of tumors was small (cases 17 and 20), and due to this, normal mucosal tissues were contained. Therefore, for the collection of tumor tissues, more comprehensive attention and skilled gross examination are required. It is thought that a flexible guideline to regulate the number of tissues to be harvested, depending on the size of tumors, is required. In the two cases, the quantity of extracellular mucin was more than 50%; thus, the samples were determined to be inadequate, but both cases were mucinous carcinomas (cases 12 and 20). Even if tissues are determined to be inadequate due to a small amount of tumors, contamination with other tissues, necrosis, or large amount of extracellular mucin, the tissues could be used for investigations by the application of the laser capture microdissection method, which allows tumor cells to be harvested selectively; thus, a more precise and detailed description is required to read the mirror image slides.

When DNA and RNA are extracted from fresh frozen tissues, a purity test, as well as integration test, is performed. Nucleic acid has the characteristic of absorbing ultraviolet light in specific patterns. When 260-nm wavelength ultraviolet light is irradiated, as the concentration of nucleic acid in samples becomes higher, the ultraviolet absorbance also becomes higher. In nucleic acid extraction samples, contamination with protein, organic components and other molecules is frequent, and proteins absorb the 280-nm wavelength. Therefore, the purity test applies the ratio of the absorbance at a 260-nm wavelength to that at a 280-nm wavelength. The lower the A\text{260}/A\text{280} ratio is, the higher the protein contamination. In our study, regardless of the elapsed time from tissue extraction to freezing, the A\text{260}/A\text{280} ratio of all samples was higher than 1.8, and the samples were determined to be ‘excellent.’

The methods evaluating the integrity of DNA are electrophoresis, Southern analysis, gene-specific polymerase chain reaction (PCR), multiple gene-specific PCR, randomly amplified polymorphic DNA (RAPD)-PCR, etc. [11]. Among them, electrophoresis is a method examining DNA fragmentation. Since it is inexpensive in comparison with other methods, it has been used widely [11]. In our study, the electrophoresis method was applied to evaluate DNA integrity, and regardless of the elapsed time from tissue extraction to freezing, all tissues were determined to be ‘adequate.’ Therefore, the DNA of colorectal cancer tissues was found to be stably maintained up to 90 minutes after extraction.

There are two methods for evaluating the integrity of RNA. One is the method assessing the ratio of the 28s to the 18s RNA band after agarose gel electrophoresis, and the other is the method measuring the RIN by using a bioanalyzer. rRNA accounts for more than 80% of the entire RNA, so it can be assessed readily. On the other hand, mRNA is only 1-3% of the entire RNA, so even if sensitive methods are applied, mRNA detection is not easy. Based on many studies inferring that the quality and the quantity of rRNA reflect the quality and the quantity of mRNA, the quality of mRNA has been evaluated by using electrophoresis of the entire RNA. Generally, when the ratio of the 28s to the 18s rRNA band is 2.0, RNA is considered to be intact. RNA is degraded readily by unskilled treatments of specimens and by destruction due to ischemia, apoptosis, necrosis, etc. The 28s rRNA is bigger than the 18s rRNA, and the structure of 28s rRNA is more unstable; thus, 28s rRNA is degraded more than 18s rRNA. Therefore, it is very difficult to obtain a ratio of the 28s to the 18s rRNA band of 2.0, and in the study reported by Sandusky et al. [10] in 75% of 1,550 specimens, the ratio of the 28s to the 18s band was found to be in the range of 1.5-1.8. In addition, Dumur et al. [12] concluded that for the analysis of a gene array, samples with a ratio of the 28s to the 18s rRNA band of higher than 1.4 were adequate, and some studies showed that the ratio of the 28s to the 18s rRNA band was not adequate to evaluate the quality and the usefulness of RNA [1, 13, 14]. In our study, the A\text{260}/A\text{280} ratios for various elapse times from colorectal cancer tissue extraction to freezing were not different, which shows that the purity level of the extracted RNA was high, without protein contamination. In addition, the ratio of the 28s to the 18s rRNA band of samples frozen 10 minutes after the extraction of colorectal cancer tissues was highest. Since 28s rRNA is degraded more than 18s rRNA with time, the ratio of the 28s to the 18s rRNA band is decreased, and by freezing the samples within a short time after extraction, good quality RNA could be obtained. On the other hand, in agarose gel electrophoresis, among samples frozen 10 minutes after extraction, the 28s and the 18s rRNA bands could not be distinguished in 1 case (case 12), a mucinous carcinoma with severe infiltration of inflammatory cells. Therefore, inflammation, mucin, etc. are thought to have effects on the stability of RNA. Nonetheless, in other cases showing inflammation, mucin, etc., the 28s and the 18s rRNA bands could be distinguished; thus, the possibility of carelessness during the specimen treatment and RNA extraction is higher than the effect of inflammation and mucin.

The intensity of the rRNA band is affected by the condition of electrophoresis, the amount of loaded RNA, and the saturation level of ethidium bromide; thus, subjective results are obtained by using a gross evaluation of the ratio of the 28s to the 18s band on the agarose gel. Hence, recently, studies evaluating RNA integrity by using a bioanalyzer to measuring the RIN have been on the rise. Fleige et al. [15] measured the RINs of various bovine organs, and they reported that in comparison with other organs, lower values were obtained from gastrointestinal tissues, which was due to their containing a large...
quantity of connective tissues; thus, a large amount of RNA was degraded during the tissue collection and extraction processes. In addition, depending on the organ, the RIN shows diverse results from 4.56 to 9.62, depending on whether tissues are hard (whether tissues contain connective tissues or adipose tissues), the difference of RNase enzyme activity, etc [15]. Human tissue with a RIN higher than 7 (‘excellent’) could be used for studies requiring high-quality RNA, such as gene array assays, and that with a RIN between 4 - 7 (‘good’) could be used for studies applying quantitative RT-PCR [9]. In our study, the RIN was highest in samples frozen 10 minutes after colorectal tissue extraction, and with increasing elapsed time, the RIN was decreased. Thus, if samples were frozen within a short time after tissue extraction, good quality RNA could be obtained. Particularly, at the four time points, 10, 30, 60, and 90 minutes, the numbers of specimens for which the RIN was determined to be ‘excellent’ were 15 cases, 8 cases, 12 cases, and 2 cases, respectively. Hence, for gene array assay, it is better to freeze samples within 10 minutes after extraction. The number of specimens determined to be ‘good’ or ‘excellent’ were 18 cases, 19 cases, 15 cases, and 7 cases, respectively; thus, for studies applying quantitative RT-PCR, it is better to freeze samples within 30 minutes after extraction. For specimens frozen 30 minutes - 60 minutes after extraction, it would be better to use them after confirmation by RNA quantification and integrity analysis. Our results on the adequacy of times from extraction to freezing were comparable to those of several other studies. Spruessel et al. [16] reported that for freezing 5-8 minutes after colorectal resection, early changes in the expression of genes and proteins were observed, and after 30 minutes, 20% of detectable genes and proteins showed important changes in expression. Huang et al. [1] reported that the stable expression of genes was shown within 20 minutes of ischemic time, and after 40 minutes, important changes in the expression of genes were shown.

In colorectal tissues, the change in the stability of DNA and RNA with the elapsed time from surgical extraction to freezing was examined. DNA was found to be stable for up to 90 minutes after extraction. The stability of RNA, on the other hand, decreased with increasing time elapsed after extraction. Based on the results, there were differences in the suitable elapsed time between extraction and freezing according to RNA research method, and for gene array assay, freezing within 10 minutes is thought to be required, and for studies applying quantitative RT-PCR, freezing within 30 minutes is thought to be required. Nevertheless, the stability of RNA becomes lower as the time until freezing is delayed, so tissues should be collected and treated within as short a time as possible, and high quality RNA can be obtained by paying more attentions to the RNA extraction process.

CONFLICT OF INTEREST

No potential conflict of interest relevant to this article was reported.

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