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<th>Overexpression of Interleukin 2 Receptor, Thymidine Kinase and Immunoglobulin-Associated Alpha-1 Messenger RNA in a Clinical Case of Enzootic Bovine Leukosis</th>
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<tr>
<td>Citation</td>
<td>Journal of Veterinary Medical Science, 74(9): 1203-1206</td>
</tr>
<tr>
<td>Issue Date</td>
<td>2012</td>
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<td>URL</td>
<td><a href="http://ir.obihiro.ac.jp/dspace/handle/10322/4096">http://ir.obihiro.ac.jp/dspace/handle/10322/4096</a></td>
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<td>日本獣医学会</td>
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Bovine leukosis is one of the most common types of neoplasms detected in dairy cattle, and is generally divided into enzootic bovine leukemia (EBL) and sporadic bovine leukemia (SBL). EBL is caused by bovine leukemia virus (BLV), while the cause of SBL is unknown [1]. BLV infection is clinically silent, occurring as an aleukemic state or as persistent lymphocytosis. Approximately 5% of animals infected with BLV develop B-cell lymphoma in various lymph nodes and organs after a long latent period [4]. Tumor cells often infiltrate many organs, including the abomasum, heart, uterus, retrobulbar space, and epidermal region of the central nervous system [21]. Therefore, clinical signs in cattle with lymphoma are generally non-specific and include weight loss, decreased appetite, and decreased productivity [1, 2].

Bovine lymphosarcoma can be easily identified by direct physical examination and diagnosed by cytological evaluation of swollen lymph nodes when lymphadenopathy and obvious neoplastic changes in target organ are present. However, it is more difficult to suspect and diagnose when located in more specific areas such as the spinal cord or abomasum and when lymph node enlargement and/or lymphocytosis is not evident [4]. While ultrasound-guided fine needle aspiration (FNA) or biopsy of vascular masses located in the body cavity, retrobulbar space, or heart is helpful [4], such diagnostic tools are not always available. Serum lactate dehydrogenase (LDH) activity may be elevated in some cattle with lymphosarcoma [7], but its specificity is insufficient to confirm diagnosis [4]. Serum thymidine kinase (TK) activity was recently reported as a potential biomarker of bovine lymphosarcoma, but is not easy to determine as it requires a radioimmunoassay [17]. In humans, gene expression profiling has been used to assess biomarkers of lymphosarcoma and leukemia. For example, Wilms’ tumor gene 1 (WT1) expression is considered a sensitive biomarker for monitoring residual disease in acute myeloid leukemia [18, 19]. Some other genes, including interleukin 2 receptor (IL2R), TK and immunoglobulin-associated alpha-1 (Mbh1 or CD79a) genes were also known to be markers of hematopoietic neoplasia [3, 7, 10]. In the present report, we describe a clinical case of EBL, and the expression of candidate genes as potential biomarkers of lymphosarcoma was measured.

A 49-month-old Holstein cow with anorexia, tachypnea, and difficulty standing up was 1st examined by a local veterinarian. At examination, the cow had a body temperature of 39.9°C, heart rate of 80 beats/min (bpm), and respiratory rate of 52 breaths/min. Despite antibiotic treatment via injection and infusion, the general condition of the cow did not improve. On day 7, enlargement of the subiliac lymph nodes was observed, and bovine leukemia was suspected. The cow was then taken to the Veterinary Teaching Hospital at Obihiro University of Agriculture and Veterinary Medicine. Depression, emaciation, and tachypnea were observed on admission. Rectal temperature, heart rate, and respiratory rate were 40.5°C, 92 bpm, and 96 breaths/min, respectively. Swelling of multiple peripheral lymph nodes,
including superficial cervical (R: 12 × 7 × 3 cm, L: 15 × 9 × 3 cm), subiliac (R: 18 × 5 × 3 cm, L: 12 × 3 × 3 cm), mandibular (R and L: diameter 4 cm), and mammary lymph nodes (R and L: diameter 8 cm) was observed. Several masses were identified in the pelvic cavity by rectal palpation. Cytological finding of FNA of the right subiliac lymph node revealed the presence of large lymphoid cells with obvious cellular atypia and several mitotic cells, indicating lymphosarcoma (Fig. 1).

Hematologic examination showed microcytic and normochromic anemia and lymphocytosis [RBC: 4.96 × 10⁶/µl, hemoglobin: 7.8 g/dl, hematocrit: 20.3%, mean corpuscular volume: 43.3 fl, mean corpuscular hemoglobin: 38.4 g/dl, WBC: 107,000/µl, neutrophils: 1,070/µl (1%), and lymphocytes: 105,930/µl (99%)]. More than 90% of lymphocytes in peripheral blood were microscopically atypical with indented nuclei and finely stippled chromatin. Serum biochemical analysis showed increased LDH activity (2,380 IU/l). LDH isozyme analysis showed elevated activities for LDH-2 (807 IU/l) and LDH-3 (455 IU/l). Serum TK activity was also measured using a commercial radioenzyme TK-assay kit with ¹²⁵I-iododeoxyuridine as a tracer (Kishimoto Clinical Laboratory, Inc., Obihiro, Japan). Extremely higher activities of serum TK activity (1,000 IU/l) were recorded compared with normal cattle [17]. Antibodies against BLV were detected by agar-gel immunodiffusion (Kitasato Institute Research Center for Biologicals, Saitama, Japan).

Messenger RNA (mRNA) expression of IL2R, TK and Mb1 in tumor tissues was examined by reverse transcription-polymerase chain reaction (RT-PCR) using the following primer sets: 5'-acg-cca-tgt-tca-agg-tct-tc -3' (IL2R forward) and 5'-gtt-ctg-cgc-ctc-tgt-gtg-tt -3' (IL2R reverse), 5'-cca-agt-cat-tga-tgg-caa-ga-3' (MB1 forward) and 5'-gat-ata-gcc-ccc-gaa-ttt-ca -3' (MB1 reverse), and 5'-cca-ggt-tgc-gta-caa-gt-3' (TK1 forward) and 5'-tct-cgc-aga-act-cca-aaa-3' (TK1 reverse). Beta-actin gene (Actb) expression was examined as an internal control using the following primer set: 5'-ctt-tcc-agc-ctc-cct-tcc-t-3' (ACTB forward) and 5'-ggg-cag-tga-tct-ctt-tct-g-3' (ACTB reverse). RT-PCR was performed on swollen subiliac lymph node tissue obtained by FNA from the case animal and two healthy cows without lymphoadenopathy as controls. Total RNA was extracted using the RNeasy Mini Kit (QIAGEN, Germantown, MD, U.S.A.) according to manufacturer’s instructions. cDNA was synthesized using 2 μg total RNA and the SuperScript™ III 1st-strand synthesis system (Invitrogen, Carlsbad, CA, U.S.A.). Our results showed that IL2R, TK1 and Mb1 genes were highly expressed in tumor tissue corresponding to the case compared to control cows (Fig. 2).

The cattle was euthanized and necropsy was performed on day 15. Gross examination revealed swelling of systemic lymph nodes, including superficial cervical, mandibular, mammary, medial, iliac, and renal lymph nodes. Furthermore, the spleen was enlarged and swollen (70 × 20 × 8 cm). Yellowish white tissue was found in the sternal bone marrow. Histopathological examination revealed neoplastic lymphoid cell infiltration in enlarged lymph nodes, liver, spleen, bone marrow, uterus (especially in the endometrium), and lamina propria of the urinary bladder, abdomen and intestine (Fig. 2). Immunohistochemical examination showed that tumor cells within enlarged lymph nodes were stained positive for BLA-36 and negative for CD3 antibodies (data not shown).
These findings suggested that the tumor cells were B-cell origin.

In the present case, we diagnosed a 49-month-old Holstein cow with EBL, based on clinical symptoms, hematological and biochemical examinations, cytology, pathological examination, and immunohistochemical analysis. The reported biomarkers of lymphosarcoma of cattle were examined in the present case [7, 9, 17]. Higher activities of total LDH, LDH-2 and LDH-3 activities (reference range; total LDH: 692–1445, LDH-2: 187–390 IU/l and LDH-3: 10–260 IU/l) [6] were observed in the present case. We also observed extremely higher TK activity in the case animal (more than 1,000 IU/l vs. 5.4 IU/l) [17]. These higher activities of both LDH and TK suggest aggressive proliferation of tumor cells in lymphoid organs and peripheral blood.

In the present study, gene expressions known as biomarkers of hematopoietic neoplasia in human were also examined. IL2R is a heterotrimetric protein expressed on the surface of immune cells, including lymphocytes and natural killer cells, and is the receptor for interleukin 2 [14, 22]. Close association of aberrant expression of IL2R with the infection of human T-cell leukemia virus (HTLV-1) was reported [20]. IL2R is thought to be directly or indirectly activated by viral products, and the aberrant expression of gene might be involved in some stages of HTLV-1-infected lymphocytes [11, 24]. TK is a cellular enzyme involved in a DNA synthesis salvage pathway, and its levels have been shown to correlate directly with the proliferative activity of tumor cells [5, 8]. Increased TK expression is often associated with increased expression of cell proliferation markers [12, 15, 23]. Both IL2R and TK1 gene overexpression have both been reported in several human leukemia cases [3, 10]; however, there are no reports available for bovine leukemia. Mb1 is a well-known B-cell specific gene, and it has been reported that it is a useful marker for B-cell neoplasms in human [13]. This gene also plays a key role in B-cell development, stabilization, and function [16]. Mb1 in vitro overexpression in BLV-induced bovine B-cell lines has been also reported [25]. In the present report, we observed increased expression of IL2R, TK1, and Mb1 genes in the case animal compared to control cows. Increased tumor cell proliferation in neoplastic lymph nodes may contribute to the overexpression of these genes. Although only one clinical case of EBL was examined in the present study, our results underscore the use of IL2R, TK1 and Mb1 gene expression as biomarkers of bovine leukosis as well as hematopoietic neoplasia in human. However, more EBL cases should be examined to confirm the validity of using the expression of these genes as biomarkers of bovine leukosis. Detail of gene expression in different stages of the disease should be also clarified by using more reliable quantitative real-time PCR in the future study.

ACKNOWLEDGMENT. We thank Dr. Noboru Kobayashi of the Tokachi Agricultural Mutual Aid Association for introducing this clinical case.

REFERENCES


