Klotho suppresses tumor progression via inhibiting IGF-1R signaling in T-cell lymphoma

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Abstract. Klotho is a transmembrane protein and acts as an upstream modulator of insulin-like growth factor-1 receptor (IGF-1R) signaling, which was indicated to be involved in the pathogenesis of solid cancer and hematological malignancies, including T-cell lymphoma. Although Klotho was recently identified as a tumor suppressor in several types of human malignancies, the potential role of Klotho in T-cell lymphoma has not been reported. In the present study, we investigated the expression level and the molecular events of Klotho in T-cell lymphoma. Significantly lower expression of Klotho was observed in T-cell lymphoma patient samples compared to normal lymph nodes. Functional analysis after Klotho overexpression revealed significantly inhibited tumor cell viability in T-cell lymphoma. Moreover, apoptosis of T-cell lymphoma cells were induced after transfected with Klotho-overexpressing vectors. Forced expression of Klotho resulted in decline of activation of IGF-1R signaling, accompanied by decreased phosphorylation of its downstream targets, including AKT and ERK1/2. These data indicated that Klotho acts as a tumor suppressor via inhibiting IGF-1R signaling, thus suppressing the viability and promoting apoptosis in T-cell lymphoma. Taken together, Klotho may serve as a potential target for the therapeutic intervention of T-cell lymphoma.

Introduction

T-cell lymphoma is the main aggressive malignancy of T lymphocytes, which is characterized by highly heterogeneous molecular and clinical characteristics. This type of lymphoma is generally resistant to chemotherapy (1). Identification of the molecular abnormalities involved in the development of T-cell lymphoma will improve the classification and provide novel therapeutic strategy. Klotho was considered an anti-aging gene when it was originally identified (2). Klotho homozygous mutant mice (kl⁻/⁻) developed multiple premature aging syndromes, including hypogonadism, skin atrophy, and pulmonary emphysema (2,3). Klotho gene is located in chromosome 13q12 in human with the size of 50 kb (2). Klotho protein consists of an extracellular domain, a single transmembrane domain and an intracellular domain. The extracellular domain (secreted Klotho), consisting of KL1 and KL2, could be released into the serum to function as a circulating hormone to regulate the activity of oxidative stress and multiple growth factor receptors (4,5).

Accumulating evidence has implicated that Klotho is extensively downregulated in several solid malignancies (6-9). Klotho was revealed to modulate the activity of several signaling pathways, including the FGF signaling, insulin-like growth factor-I receptor (IGF-1R) and Wnt pathways (5,10-12). These pathways also play crucial roles in the development of T-cell lymphoma (13,14). However, the role of Klotho in T-cell lymphoma has not been reported.

In the present study, we hypothesized that Klotho was aberrantly expressed and involved in the pathogenesis in T-cell lymphoma. We observed decreased expression of Klotho in the patient tissues and cell lines of T-cell lymphoma. The anti-proliferative and pro-apoptotic effect of Klotho were further identified in T-cell lymphoma cells. Klotho could act as an upstream regulator of IGF-1R signaling in this disease. The data elucidated the potential tumor suppressive role of Klotho in the development of T-cell lymphoma. It may serve as a potent candidate for the target therapy of T-cell lymphoma.

Materials and methods

Patients and samples. The paraffin-embedded lymph node samples were collected from 35 newly diagnosed cases of T-cell lymphoma (21 males and 14 females; age range, 19-81 years; median age, 53 years) and 20 normal lymph nodes. Histological diagnoses were established according to the WHO classification (15). Normal peripheral blood mono-
nuclear cells (PBMCs) from healthy volunteers were isolated by Ficoll centrifugation (TBD Science, Tianjin, China). T cells were isolated with Nylon Wool Fiber Columns. This study was approved by the Medical Ethics Committee of Shandong Provincial Hospital Affiliated to Shandong University. All samples were obtained with informed consent in accordance with the Declaration of Helsinki.

**Cell lines.** Jurkat and Molt-3 (T-cell acute lymphocytic leukemia cell lines) were available from Typical Culture Preservation Commission Cell Bank (Chinese Academy of Science, Shanghai, China). MyLa 3676 (cuteaneous T-cell lymphoma, lymphoblast, Sezary syndrome) was retained by our laboratory. Karpas 299 (ALK+ ALCL cell line) was obtained from Shanghai Bioleaf Biotech Co., Ltd. All the above cell lines were maintained in PRMI-1640 (Gibco, Life Technologies, Rockville, MD, USA) containing 10% fetal bovine serum (FBS; Gibco, Life Technologies) and 1% penicillin/streptomycin mixture.

**Cell transfection.** Lentivirus vectors either encoding Klotho (LV-KL) or an empty lentiviral vector (LV-Con) were generated by GeneChem (Shanghai, China). Lentivirus transfection of T-cell lymphoma cells were performed according to the manufacturer's instructions. Infection efficiencies were assessed by green fluorescent protein (GFP) fluorescence through flow cytometry. The stably transfected cells were selected with 5 µg/ml puromycin (Sigma-Aldrich, USA).

**Immunohistochemistry (IHC).** IHC was carried out with primary rabbit anti-Klotho (Abcam, Cambridge, USA). 4-µM-paraffin sections were deparaffinized and hydrated. Antigen retrieval was performed in 0.01 M sodium citrate (pH 6.0) buffer under high-pressure. Endogenous peroxidase was blocked with 3% H2O2 for 15 min, followed by incubation overnight at 4°C with primary anti-Klotho (1:100). Then the tissue sections were treated with the second antibody from SP reagent kit (Zhongshan Golden Bridge Biotechnology Co., Ltd., Beijing, China) for 30 min at room temperature. Bound antibody was detected by secondary antibody and diaminobenzidine (DAB) kit (Zhongshan Golden Bridge Biotechnology). The stained slices were counterstained with hematoxylin and mounted. Negative control was performed with the primary antibody replaced by PBS. IHC staining was scored by two independent observers who were blinded to the patient clinical data. Cases with ≥10% positive cells were considered as positive.

**Real-time quantitative polymerase chain reaction (qRT-PCR).** For the qRT-PCR, total RNA was extracted with RNAiso Plus (Takara, Dalian, China). The mRNA level was detected by Nanodrop 2000 (Thermo Fisher Scientific, Waltham, MA, USA). Reverse transcription reaction was performed with reverse transcription reagents (Takara). Expression level of mRNA was quantified using SYBR Green Premix Ex Taq II kit (Takara) in LightCycler 480 real-time quantitative PCR system (Roche, Basel, Switzerland). Primers for qRT-PCR are listed in Table I. Glyceraldehyde-3-phosphate dehydrogenase (GAPDH) was used as an internal control and relative quantification was determined by the 2−ΔΔCt method.

<table>
<thead>
<tr>
<th>Gene</th>
<th>Primer sequence</th>
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<tbody>
<tr>
<td>Klotho</td>
<td>F, 5'-AGCAATCTGGTCTGAAACACTGG-G-3' R, 5'-CATGTTCACGCTGAAAGTTCAGG-3'</td>
</tr>
<tr>
<td>GAPDH</td>
<td>F, 5'-GGGAGAACCTGCGGGTATC-G-3' R, 5'-GAGTGGGTGTCGCTGGTA-G-3'</td>
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**Western blotting.** For western blot analysis, cell lysates were extracted by radio-immunoprecipitation assay (RIPA) buffer (Shenergy Biocolor, Shanghai, China) together with phosphatase inhibitor cocktail (PhosSTOP; Roche, Mannheim, Germany). The BCA assay (Shenergy Biocolor) was performed to detect protein concentration. Proteins (30 µg) were electrophoresed on SDS-PAGE and transferred to PVDF membranes (Millipore, Billerica, MA, USA). Membranes were blocked with 5% skim milk in Tris-buffered saline. After incubation with primary antibodies, membranes were incubated with HRP-conjugated secondary antibodies (Zhongshan Golden Bridge). Proteins were visualized by electro-chemiluminescence detection reagents (Amersham Imager; General Electric, USA). The following antibodies used for western blotting were purchased from Cell Signaling Technology: p-IGF-1R, t-IGF-1R, p-AKT, t-AKT, p-ERK1/2, t-ERK1/2, Mcl-1 and caspase-3. Klotho antibody was purchased from Abcam. The experiments were performed in triplicate with GAPDH (Zhongshan Golden Bridge) as endogenous control.

**CCK-8 proliferation assay.** The influence of Klotho on viability of T-cell lymphoma cells were assessed by Cell Counting Kit-8 (CCK-8; Dojindo, Kumamoto, Japan). Briefly, T-cell lymphoma cell lines stably transfected with LV-KL or LV-Con were seeded into 96-well plates (5,000 cells/100 µl/well, respectively) with PRMI-1640 medium supplemented with 10% FBS. Then the cells were incubated with 10 µl/well of CCK-8 and cell viability was detected by light absorption at 450 nm by Multiskan GO Microplate Reader (Thermo Scientific, Rockford, IL, USA).

**Cell apoptosis assay.** Apoptosis of T-cell lymphoma cells were tested by Annexin V-PE/7-aminoactinomycin (7AAD) kit (BD Biosciences, Bedford, MA, USA) according to the manufacturer's instructions. Cells with designed treatment were harvested and washed twice in cold PBS and resuspended in 1X binding buffer at a concentration of 1×105 cells/ml, then 100 µl was taken and incubated with 5 µl Annexin V-PE and 5 µl 7AAD for 15 min at room temperature in the dark for each tube. Afterwards, 400 µl of 1X binding buffer was added and cell apoptotic rates were detected by Navios Flow Cytometer (Beckman Coulter, CA, USA).

**Immunofluorescence.** T-cell lymphoma cells with designed treatment were seeded onto glass slides by liquid thin layer cell smear (Xiaogan Aohua, Xiaogan, China). Thereafter, the T-cell lymphoma cells were fixed in 4% PFA and permeabilized with 0.1% Triton X-100. The cells were blocked with
normal goat serum for 1 h. Then the slides were incubated with primary antibodies at 4˚C overnight. After washing with PBS, the DyLight 488-conjugated secondary antibodies (Abbkine, CA, USA) were added. Slides were washed and mounted with 4′6-diamino-2-phenylindole (DAPI; Boster, Wuhan, China). Images were examined and recorded by Nikon IX73 fluorescent microscope.

Statistical analysis. All statistical analyses were performed by using statistic software SPSS17.0 (SPSS Inc., Chicago, IL, USA). Data are presented as the mean ± standard deviation (mean ± SD) from three separate experiments. Differences between groups were analyzed by one-way analysis of variance (ANOVA) or t-tests. *p<0.05, **p<0.01.

Results

Klotho is downregulated in T-cell lymphoma. We examined Klotho expression for primary T-cell lymphoma samples including NK/T-cell lymphoma (n=12), peripheral T-cell lymphoma-not otherwise specified (PTCL-NOS, n=7), angio-immunoblastic T-cell lymphoma (AITL, N=6), cutaneous T-cell lymphoma (CTCL, n=6), and enteropathy associated T-cell lymphoma (EATL, n=4). Compared with normal lymph nodes, expression levels of Klotho were significantly lower in T-cell lymphoma tissues (Fig. 1A). Klotho positive rate was 14% (5 of 35) in T-cell lymphoma tissues whereas 75% (15 of 20) in normal lymph nodes. Cases with ≥10% positive cells were considered as positive. More detailed data are summarized in Table II.

Table II. Immunohistochemical expression of Klotho in T-cell lymphoma tissues.

<table>
<thead>
<tr>
<th>Klotho expression</th>
<th>NK/TCL (n=12)</th>
<th>AITL (n=6)</th>
<th>CTCL (n=6)</th>
<th>EATL (n=4)</th>
<th>PTCL-NOS (n=7)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Positive</td>
<td>0 (0%)</td>
<td>1 (16.7%)</td>
<td>2 (33.3%)</td>
<td>1 (25%)</td>
<td>1 (14.3%)</td>
</tr>
<tr>
<td>Negative</td>
<td>12 (100%)</td>
<td>5 (83.3%)</td>
<td>4 (66.7%)</td>
<td>3 (75%)</td>
<td>6 (85.7%)</td>
</tr>
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</table>

Figure 1. Expression of Klotho in the samples and cell lines of T-cell lymphoma. (A) Expression level of Klotho protein was lower in T-cell lymphoma cases than that in normal lymph nodes (original magnification, x400). (B) As detected by real-time quantitative PCR, Klotho mRNA levels were downregulated in T-cell lymphoma cell lines, compared to that in normal T cells (N1, N2 and N3). (C) Protein expression of Klotho was detected in DLBCL cells and normal T cells. *p<0.05, **p<0.01.
We next confirmed the expression of Klotho in T-cell lymphoma cell lines. As shown in Fig. 1B, T-cell lymphoma cells exhibited remarkably lower mRNA levels of Klotho compared to the peripheral blood T lymphocytes from healthy donors. Decreased protein levels of Klotho expression were also noted in T-cell lymphoma cell lines (Fig. 1C).

**Klotho restrains T-lymphoma cell proliferation.** To explore the function relevance of Klotho in the progression of T-cell lymphoma, T-cell lymphoma cell lines (Jurkat, Molt-3, Karpas 299 and MyLa 3676) were stably transfected with either Klotho-overexpression lentivirus vectors or empty vector control (Fig. 2A). Upregulated level of Klotho mRNA was confirmed by quantitative PCR (Fig. 2B). CCK-8 assay was employed to examine the viability of the above cells. Significantly decreased cell viability was noted in T-cell lymphoma cells transfected with LV-KL, compared with those transfected with LV-Con (Fig. 2C). These results indicate that Klotho inhibits the proliferation of T-cell lymphoma cells.

**Klotho promotes apoptosis of T-lymphoma cells.** To further investigate the mechanisms underlying the suppression of Klotho in cell proliferation by ectopic Klotho expression, Annexin V-based apoptotic assays were performed in Jurkat and Karpas 299 cell lines with stable LV-KL or LV-Con transfection. Cells transfected with LV-KL exhibited enforced apoptosis rates in T-cell lymphoma cell lines (Fig. 3A). In addition, apoptotic promotion effect of Klotho was confirmed by western blot analysis. Remarkable reduction of anti-apoptotic protein Mcl-1 and total caspase-3 protein was observed in Jurkat and Karpas 299 cell lines (Fig. 3B and C).

**Klotho suppresses the activation of IGF-1R signaling in T-cell lymphoma.** Having elucidated that Klotho could interfere with the viability and apoptosis of T-cell lymphoma cell lines, we next explored the involved molecular mechanisms responsible for the function of Klotho. The IGF-1R signaling plays significant role in the development of T-cell lymphoma and proliferation of T-cell lymphoma cell lines could be...
induced by IGF-1 exploration. CCK-8 assay was performed to determine the effect of Klotho overexpression on IGF-1-induced cell proliferation. T-cell lymphoma cell lines stably transfected with either Klotho overexpressing vector or empty control vector were incubated with IGF-1 or vehicle control in 0.5% FBS culture medium for 48 h. In the group without IGF-1 treatment, LV-KL transfection induced declined viability of T-cell lymphoma cell lines compared to the empty-vector group. Furthermore, we noted that the Klotho-induced inhibition of T-cell viability was less reversed by IGF-1 treatment. With IGF-1 addition, cell proliferation of LV-Con treated cells increased by up to 50%, whereas the only up to 30% enhancement of cell proliferation was found in cells with Klotho overexpression (Fig. 4A).

We next investigated the modulation effect of Klotho on IGF-1R signaling in T-cell lymphoma cell lines. Jurkat and Karpas 299 cells transfected with LV-KL revealed significantly decreased phosphorylation level of IGF-1R. Moreover, the downstream targets of IGF-1R signaling, including AKT and ERK1/2, were also inhibited by Klotho overexpression (Fig. 4B and C).

Discussion

In the present study, we provide the first evidence that Klotho as a potential tumor suppressor in the tumorigenesis of T-cell lymphoma. The decline of Klotho initiates anti-proliferative and pro-apoptotic effect through inhibiting IGF-1R signaling activation. This study illuminates the tumor suppressive effect of Klotho and highlights the potential application of Klotho in the targeted therapy in T-cell lymphoma.

The tumor suppressive function of Klotho has been reported in several human solid malignancies, but less in hematological cancers, especially in lymphoma (16-19). Lower Klotho expression was detected in T-cell lymphoma tissues whereas high expression level of Klotho was observed in normal lymphomas.
nodes, in accordance with recent studies showing decreased Klotho expression in human malignancies. Recently, accumulating evidence demonstrated several mechanisms involved in the aberrant expression of Klotho in solid cancers, including epigenetic mechanisms and autophagy (17,20,21). Additionally, Klotho decline has a significant prognostic value in cancers, such as large cell neuroendocrine carcinoma (LCNEC), small cell lung cancer (SCLC) and hepatocellular carcinoma (22-24). Usuda et al (22) reported that Klotho staining provided a novel biomarker for prognosis in patients with LCNEC, especially for those without lymphangioinvasion or lymph node metastasis. It suggested that evaluation of Klotho in cancer may improve the personalized treatment in human malignancies. T-cell lymphoma is characterized by heterogeneity in clinical, molecular and biological presentations. The clinical diagnosis of T-cell lymphoma is primary based on the biopsy histopathology (25). With the deepening of further investigations, lower Klotho expression may serve as a potential marker for the pathological diagnosis of T-cell lymphoma.

Furthermore, our data elucidated that the cell growth of T-cell lymphoma could be restrained by Klotho. Overexpression of Klotho inhibited the proliferation and induced apoptosis of T-cell lymphoma cells. The results are consistent with the studies in lung cancer (7,17). Moreover, it has been reported that soluble Klotho and conditioned medium from Klotho-
overexpressing cells could suppress the growth of pancreatic cancer and breast cancer cells (16,26). However, investigations with soluble Klotho and in vivo studies are still needed to be conducted to explore the effective domain of Klotho protein and evaluate the safety and efficacy, and pharmacokinetic mechanisms of Klotho in T-cell lymphoma.

Interestingly, we observed the suppressive effect of Klotho on the activation of IGF-1R signaling in T-cell lymphoma. Our findings are consistent with recent reports which elucidated that Klotho could serve as a regulator of IGF-1R signaling in cancer. Aberrant activation of IGF-1R signaling was involved in the development and progression of T-cell lymphoma (14,27). Moreover, structure-function analysis indicated that Klotho could interact with the IGF-1R (28). Klotho-induced suppression of IGF-1R pathway may acts as a novel mechanism involved in the pathogenesis of T-cell lymphoma. Additionally, several signaling pathways, such as Wnt signaling, FGF23 signaling and PI3K signaling, are also reported to participate in the biological mechanism of Klotho (16,29,30). Further investigations are still needed to clarify the detailed mechanism involved in the development different types of tumors.

In conclusion, this study identified that Klotho acted as a novel tumor suppressor in T-cell lymphoma. We elucidated the crucial role of Klotho in inhibiting tumor cell proliferation and inducing cell apoptosis in T-cell lymphoma. The tumor suppressive effect of Klotho may be mediated by inhibiting the activation IGF-1R signaling pathway. Being an endogenous circulating hormone, the secreted Klotho could function as an active form and inhibit the tumor growth safely and effectively in mice. These data suggested that Klotho may serve as a hopeful target for the development of novel therapeutic strategy of T-cell lymphoma.

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