

Overexpression of ASPH protein predicts poor outcomes in
retroperitoneal liposarcoma patients

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Abstract

Background: No druggable targets and prognostic factors are currently available for retroperitoneal liposarcoma (RPLS) due to very limited understanding of its molecular mechanisms and pathogenesis. This study aims to decipher expression profiling and prognostic value of aspartate- β -hydroxylase (ASPH), an activator of Notch signaling pathway, in RPLS.

Methods: Totally 138 patients with RPLS who received resection were recruited in this retrospective study. Immunohistochemistry was performed to decipher expression profiling of ASPH in archived specimens. Prognostic value of ASPH was evaluated by Kaplan-Meier plot and Cox proportional hazards model. Chi-square test was used to compare counting data.

Results: The overall positive rate of ASPH expression in RPLS was 90.6%. Well-differentiated liposarcoma had a similar positive rate to dedifferentiated liposarcoma (90.2% vs. 91.1%). A high level of ASPH expression correlated with reduced postoperative recurrence-free survival rate and overall survival rate ($P < 0.05$).

Conclusions: ASPH is an independent predictor for clinical outcome of patients with RPLS. A high level of ASPH expression confers poor post-operational prognosis.

Background

Liposarcoma is a common subtype of retroperitoneal sarcoma [1]. Due to the complicated anatomical heterogeneity of retroperitoneal liposarcoma (RPLS), either adjacent to or even invading into major vessels and important organs when diagnosed, massive bleeding frequently occurs during surgery. Thus, combined viscera resection is required, which makes the operation on RPLS much more challenging than other solid tumors, resulting in a much lower resection rate [2]. Currently, surgery is the most effective treatment for RPLS, and adjuvant therapies have shown little/no efficacy [3,4]. Chemotherapy has no definite therapeutic effect on RPLS [5], and radiotherapy is very limited due to serious toxicity to adjacent abdominal organs [6]. There is a paucity of research focusing specifically on molecular mechanisms underlying RPLS pathogenesis. Consequently, no diagnostic biomarkers, druggable targets or prognostic factors are available for RPLS.

Aspartate- β -hydroxylase (aspartyl- β -hydroxylase or asparaginyl- β -hydroxylase; ASPH) is highly conservative deoxidizing enzyme [7]. The ASPH is critical for embryonic development. Normally, ASPH is silenced during adulthood and only expressed in placenta [8,9]. Abnormal re-expression/upregulation of ASPH occurs in vast majority of malignancies, such as hepatocellular carcinoma, lung cancer, colorectal cancer, breast cancer and pancreatic cancer [10-12].

The enzymatic activity of ASPH depends on its catalytic domain [13,14], which hydroxylates aspartic acid or asparagine residues in EGF-like repeats of several proteins (such as Notch receptors and Notch ligands) in the presence of iron divalent, thus mediating cellular motility and differentiation [15,14,13]. Previously, correlation between ASPH mRNA transcription levels and insulin receptor substrates, growth factors, IGF receptors, Notch, Jagged, and HES was analyzed. Activation of

IGF-1/IGF-2 signal leads to upregulation of ASPH and thus Notch. Biologically, ASPH promotes tumorigenesis partially through activating Notch and SRC signaling pathways [16,17]. Notably, ASPH's oncogenic properties rely on its β -hydroxylase activity, which enhances proliferation, migration, invasion, and metastasis [18, 19].

The Notch signaling pathway is indicated to be highly activated in RPLS, and the growth of liposarcoma cells can be significantly blunted by knocking-down of Notch [20]. Hence, we explored if ASPH is associated with tumor characteristics and clinical outcome of RPLS. Here, in a large RPLS cohort, we performed immunohistochemistry to detect the expression profiling of ASPH and evaluate its potential in prognostics of RPLS.

Methods

Patient selection

After the ethical approved, 179 patients with RPLS were admitted to Department of retroperitoneal tumor surgery, Peking University International Hospital, from December 1, 2014 to February 28, 2018. Among them, 24 cases had surgical residues, 5 cases died due to serious postoperative complications, and 12 cases had extremely rare subtypes. These 41 cases were excluded from data analysis. Thus, 138 cases (71 males and 67 females) were eventually included in this study. All cases were confirmed by histopathologic and immunohistochemical examination. The mean age of patient population was 54.6 ± 11.5 . Histologically, 82 cases had well-differentiated liposarcoma (WD) and 56 had dedifferentiated liposarcoma (DD). All patients received no chemoradiotherapy or targeted therapy before surgery. The general information and FNCLCC scores of these patients were retrospectively collected from electronic medical record system.

Immunohistochemistry

Formalin fixation and paraffin-embedded (FFPE) tissue blocks from archived tissue-bank were applied for immunohistochemical staining. In brief, specimens were sectioned at 4 μ m-thick, deparaffinized in xylene, and rehydrated, followed by antigen retrieval in sodium citrate. Then slides were incubated with primary monoclonal antibody against ASPH (1:4000 dilution, courtesy of Professor Jack R. Wands, Brown University, USA) overnight at 4°C. Afterwards, biotin-labeled goat anti-mouse secondary antibody (Beijing Zhong Shan Jin Qiao biotechnology co. LTD) was added. Sections were then counterstained with hematoxylin, dehydrated, and mounted.

Immunohistochemical staining was evaluated and scored by 3 independent pathologists. Four high-power fields ($\times 200$) were randomly selected for each slide, and the number of positive cells in a single field was counted and scored according to staining distribution: score 0 (negative), score 1 (0-25%), score 2 (26%-50%), score 3 (51%-100%). The average score of all fields was calculated. The average score ≤ 0.5 was set as negative to low expression, and >0.5 as moderate to high expression.

Follow-up

All cases were followed up by telephone or outpatient. The follow-up period was from the patient's discharge to October 2019 or the death date. Median follow-up was 21.5 months (4-53 months). Among the 138 patients, 84 relapsed, 43 died (all due to tumor recurrence and metastasis), and 14 were out of contact. The follow-up rate was 89.8%.

Statistical analysis

Statistical software SPSS19.0 was used to analyze the data. Chi-square test was used for the analysis of counting data. The Kaplan-Meier method was used for survival analysis. COX regression model was used for the correlation analysis.

Comparisons made in which $P < 0.05$ were considered statistically significant.

Results

Expression of ASPH in RPLS

Compared to adjacent normal tissue, ASPH was overexpressed in RPLS tumor in most cases (Figure 1). The overall positive rate of ASPH staining was 90.6%. The positive rate [90.2% (74/82)] in well-differentiated RPLS was significantly lower than that in dedifferentiated RPLS [91.1% (51/56), $p = 0.028$].

Associations between ASPH expression and tumor characteristics of RPLS

There was no difference for ASPH expression between patients divided by age, gender, tumor number, P53 expression, and MDM2 expression ($P > 0.05$, Table 1). The patients with worse pathology subtypes ($P = 0.028$) and higher FNCLCC grades ($P = 0.007$) showed a higher ASPH expression level than others (Table 1).

ASPH is an independent prognostic factor for RPLS

Kaplan-Meier analysis showed that the overall survival (OS) of patients with high ASPH expression was much worse than that of patients with low expression (Figure 2A, $\chi^2 = 6.56$, $p = 0.010$). The relapse-free survival (RFS) of patients with high ASPH expression was also much worse than that of patients with low ASPH expression (Figure 2B, $\chi^2 = 7.17$, $p = 0.007$).

In Cox regression univariate model, expression level of ASPH and FNCLCC grade were identified as prognostic factors ($p < 0.05$, Table 2). In multivariate model, FNCLCC grade and expression level of ASPH were independent risk factors of RFS. Compared with lower expression level, the recurrence risk was 1.84 times higher in patients with higher expression level of ASPH (OR=1.84, 95% CI: 1.12-3.04, $p = 0.017$, Table 2).

Discussion

Compared with well-differentiated RPLS, dedifferentiated RPLS is much more malignant [21], with unfavorable histopathological and clinical characters such as invasion into adjacent tissues at an early stage and a higher FNCLCC grade [22]. In this study, ASPH expression in dedifferentiated RPLS was significantly higher than that in well-differentiated RPLS, and related to FNCLCC grade. Thus, ASPH may play a role in the progression of RPLS, especially in dedifferentiated RPLS.

Importantly, ASPH was an independent risk factor for relapse of RPLS. Patients with a higher level of ASPH expression were more likely to relapse. The effect of ASPH expression on OS is similar to RFS, except an intersection of curves in the period of 46 months after surgery. Considering the complexity and heterogeneity of RPLS patients' preoperative status, the unexpected intersection would be possibly explained by complicity after surgery. Some patients with RPLS may survive a relatively long time through receiving repeated or even multiple operations. Some patients give up further treatment after suffering from economic difficulties, psychological fear and complications after multiple surgical operations. All those factors make OS a less objective indicator than RFS.

It has been reported that ASPH promotes proliferation, migration, invasion in various cancer types, including hepatocellular carcinoma [23], breast carcinoma [24] and pancreatic cancer. Thus, ASPH acts as an oncogene and is expected to enhance aggressive/malignant cellular behaviors, possibly leading to a poor prognosis in RPLS. This hypothesis is under evaluation by cytological experiments. In this study, IHC data have provided supporting evidence for ASPH's oncogenic role in RPLS.

Previous studies have demonstrated that clinicopathological factors such as pathological subtypes and tumor size are associated with prognosis [25]. Some studies

have suggested that S-100, Ki-67 [26], MDM2 and P53 [27] are associated with the prognosis of RPLS. However, with a much larger cohort, neither MDM2 nor P53 is a prognostic factor. Instead, ASPH has been identified as a potential prognostic factor in RPLS. Considering the embryonic origins of RPLS are different from cancers originated from epithelium, very few tumor biomarkers are valid in RPLS. However, before a better understanding of molecular mechanisms of RPLS, ASPH with pan-cancerous verification would be the first choice in development of RPLS biomarkers.

The biological characteristics of RPLS are different from carcinomas. The RPLS tend to grow expansively and is prone to postoperative recurrence, but rarely exhibits distant metastasis [27]. Thus, RPLS patients receive repeated and multiple surgical treatment. It is extremely important for such patients to prolong overall/disease-free survival without recurrence after surgery, and to gain "time" and "space" for the next surgical resection. Our results suggested that patients with high-ASPH RPLS would be considered as high-risk of relapse, which should receive a much closer observation and longer follow-up.

It is reported that inhibiting expression or enzymatic activity of ASPH could undermine proliferation, migration, invasion, and metastasis of various tumor cells [28,29]. In addition, ASPH can serve as an immunotherapy target for liver cancer [30]. Further studies are needed to verify histopathological effects of ASPH on RPLS tumor cells, which would help to determine if ASPH can be used as a druggable target or an immunotherapeutic target for (neo) adjuvant treatment of RPLS.

Conclusions

In summary, ASPH is an independent predictor for clinical outcome of patients

with RPLS. A high level of ASPH expression confers poor post-operational prognosis. In particular, our study suggests that ASPH maybe a potential therapeutic target for of RPLS.

Abbreviations

RPLS: retroperitoneal liposarcoma

ASPH: aspartate- β -hydroxylase

OS: overall survival

RFS: relapse-free survival

Declarations

Ethics approval and consent to participate

This study was approved by the Ethics committee of Peking University International Hospital (Beijing, China) and all participants signed informed consent forms.

Availability of data and material

The datasets used and/or analysed during the current study available from the corresponding author on reasonable request.

Consent for publication

Not applicable

Competing interests

The authors declare that they have no competing interests.

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Authors' contributions

MMX and XBC carried out the main analysis. CHL conceived and designed the study. YQC, LHW, and XSR helped to collect and reformat the primary data. XSC helped to analyze data and revise the manuscript. MMX and XBC draft the manuscript. All authors have read and approved the manuscript.

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Figure Legend

Figure 1. Immunohistochemical staining for ASPH expression in retroperitoneal liposarcoma tissues.

(A-B) Negative staining of ASPH (100 \times , 200 \times , respectively); (C-D) Weak staining of ASPH where scored = 1 (100 \times , 200 \times , respectively, 0-25% positive cells); (E-F) Moderate staining of ASPH where scored = 2 (100 \times , 200 \times , respectively, 25-50% positive cells); (G-H) Strong staining of ASPH where scored = 3 (100 \times , 200 \times , respectively, 50-100% positive cells).

Figure 2. OS and RFS of retroperitoneal liposarcoma stratified by ASPH expression levels.

The OS stratified by ASPH expression levels. B. RFS stratified by ASPH expression levels.