

## Original Article

# Prognostic significance of stromal SMYD3 expression in colorectal cancer of TNM stage I-III

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**Abstract:** Background: SET and MYND domain-containing protein 3 (SMYD3) is a histone methyltransferases and it promotes progression of many kinds of cancers including lung cancer, ovarian cancer and gastric cancer. In colorectal cancer (CRC), SMYD3 is proved to stimulate the proliferation of cancer cells, but the clinical significance of SMYD3 in CRC has not been elucidated. Methods: In our study, we detected the expression of SMYD3 in CRC samples in TNM stage I-III with immunohistochemistry. The correlation between the expression of SMYD3 and the clinicopathological factors was analyzed with Chi-square test. The survival curve was displayed by Kaplan-Meier test and the statistical difference of subgroups was analyzed with log-rank test. Independent prognostic factors were identified by the Cox proportional hazards regression model. Results: The percentage of high SMYD3 expression and low expression accounts for 47.98% and 52.02% respectively. High expression of SMYD3 was significantly associated with advance T stage (P=0.006) and lower survival rates (P=0.010), and it could be identified as an independent prognostic factor indicating unfavorable prognosis of patients with CRC (P=0.032, HR=1.98, 95% CI=1.06-3.70). Conclusions: SMYD3 high-expression is a high risk for poorer prognosis of CRC in TNM stage I-III. Our findings suggested that detecting SMYD3 may help stratify patients by risk more precisely and help make the individual treatment strategy.

**Keywords:** SET and MYND domain-containing protein 3, colorectal cancer, tumor invasion, prognosis

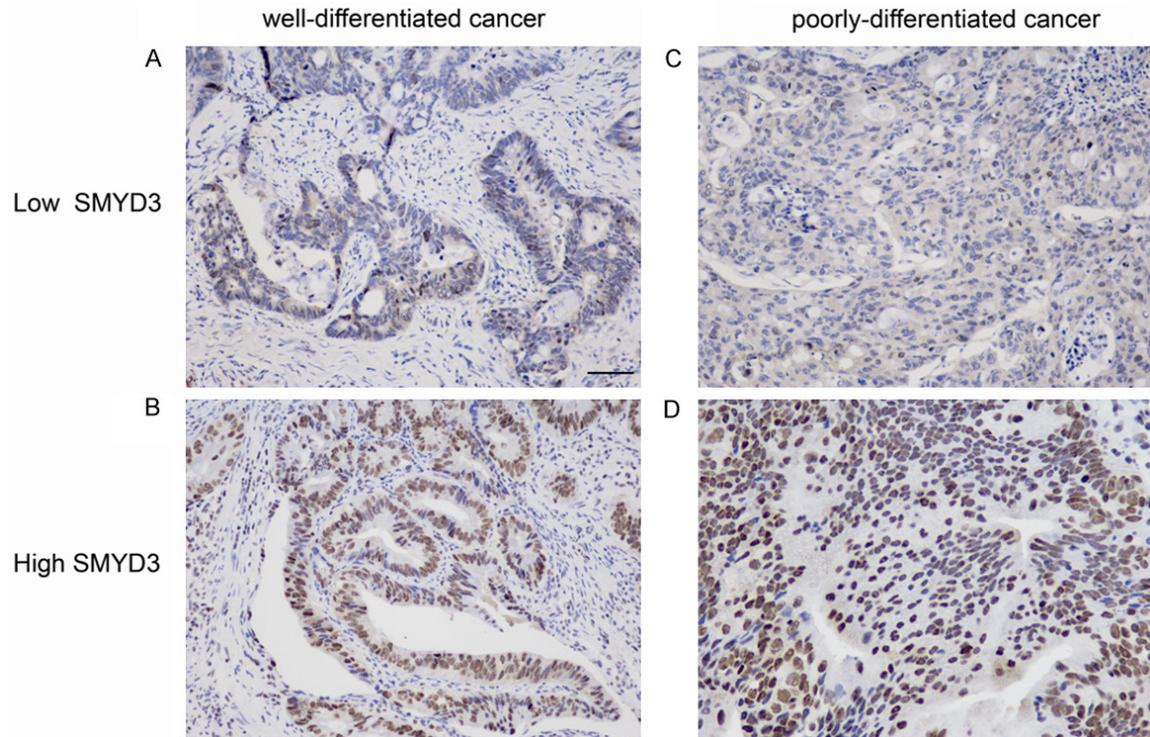
## Introduction

Colorectal cancer (CRC) is the third most common cause of cancer deaths with about 13 million new cancer cases and 694,000 cancer deaths in 2012 worldwide [1, 2]. In developing countries like China, the incidence of CRC is increasing significantly. Major high risk factors of CRC include genetic and environmental factors, a heat-treated high-fat diet, obesity, and a sedentary lifestyle [3]. Although the diagnosis and treatment equipment have developed rapidly in recent years, the 5-year overall survival rate of CRC remains about 30% in China [4, 5]. The reasons contributing to the unsatisfactory prognosis of CRC are its late tumor presentation, rapid progression and high recurrence [6]. About 20% of these patients suffer from tumor recurrence and metastasis even after radical surgery and systemic adjuvant chemotherapy [7]. So effective prognostic biomarkers are urgently needed to predict the prognosis and

define the high-risk patients for adjuvant chemotherapy.

SET and MYND domain-containing protein 3 (SMYD3) is a histone methyltransferases which methylates 'Lys-4' of histone H3 and induces di- and tri-methylation instead of monomethylation [8, 9]. The Smyd-family consists of 5 members (SMYD1, 2, 3, 4 and 5), containing a SET domain methyltransferases and an intervening MYND domain. The SET domain is essential to promote the dimethylation or trimethylation of chromosomal histone, resulting in a loosened state of the chromosome's spatial structure and affecting gene transcription [8, 10]. The MYND domain can enhance the methylation function of the SET domain by mediating protein-protein interactions [11, 12]. Overexpression of SMYD3 has been reported to be involved in the progression or prognosis in many kinds of cancers, including gastric cancer, leukemia, ovarian cancer, and so on [13,

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**Figure 1.** Representative immunohistochemical figure. The representative image for low-expression or high-expression of SMYD3 in well-differentiated (A and B) and poorly-differentiated (C and D) colon cancer. Scale bar: 100  $\mu$ m.

14]. It is demonstrated that SMYD3 is an oncoprotein via functioning as a transcriptional potentiator of multiple cancer-promoting genes, by which SMYD3 promotes cell proliferation and epithelial-mesenchymal transition in colon cancer [15]. However, the clinical significance of SMYD3 in CRC is still unknown.

In our study, we detected the expression of SMYD3 in CRC samples in TNM stage I-III with immunohistochemistry (IHC), and analyzed the correlation between SMYD3 expression, clinicopathological factors and overall survival rates. Moreover, we identified the independent prognostic factors of CRC.

### Materials and methods

#### *Patients and follow-up*

Our study was approved by the committee of Linyi Central Hospital. A total of 173 patients diagnosed as CRC from 2004 to 2010 were selected into our cohort according to the criteria (1) patients were in TNM stage I-III and underwent radical surgical resection, (2) available follow-ups more than 5 months and enou-

gh specimens for IHC, (3) the TNM stage of CRC was in stage I to III. All the specimens were obtained from the Department of Pathology with prior consents of patients. The diagnosis of CRC was double confirmed by two senior pathologists, and the TNM stage in our study was according to the 7th American Joint Committee on Cancer/Union for International Cancer Control.

#### *Immunohistochemistry*

The expression of SMYD3 was detected with IHC described previously in detail [16, 17]. The formalin-fixed and paraffin-embedded specimens were first deparaffinized at 55°C for 20 minutes and then washed with xylene. Rehydration of samples was realized by graded ethanol with concentration at 100%, 95%, and 80%. Slides were incubated in 3% hydrogen peroxide to block the endogenous peroxidase activity and in citrate buffer (pH=6.0) with a microwave oven for optimal antigen retrieval. Followed by blocking unspecific binding with incubation in 5% bovine serum albumin for 30 minutes, primary antibody of SMYD3 (#12,859, 1:500 dilution; Cell Signaling Technology, Danvers, MA,

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**Table 1.** Basic information of patients

| Characters          | Number | Percentage |
|---------------------|--------|------------|
| Sex                 |        |            |
| Male                | 81     | 46.82%     |
| Female              | 92     | 53.18%     |
| Age                 |        |            |
| <60                 | 72     | 41.62%     |
| ≥60                 | 101    | 58.38%     |
| Tumor diameter (cm) |        |            |
| ≤5                  | 69     | 39.88%     |
| >5                  | 104    | 60.12%     |
| T stage             |        |            |
| T1+T2               | 77     | 44.51%     |
| T3+T4               | 96     | 55.49%     |
| Lymph node invasion |        |            |
| No (N0)             | 105    | 60.69%     |
| Yes (N1/2)          | 68     | 39.31%     |
| TNM stage           |        |            |
| I                   | 50     | 28.90%     |
| II                  | 55     | 31.79%     |
| III                 | 68     | 39.31%     |
| Differentiation     |        |            |
| Good                | 97     | 56.07%     |
| Moderate            | 48     | 27.75%     |
| Poor                | 28     | 16.18%     |
| Adjuvant therapy    |        |            |
| No                  | 54     | 31.21%     |
| Yes                 | 112    | 64.74%     |
| SMYD3               |        |            |
| Low                 | 90     | 52.02%     |
| High                | 83     | 47.98%     |

Abbreviations: SMYD3 = SET and MYND domain-containing protein 3.

USA) was used to incubate the specimen at 4°C overnight. After rinsing the slide with phosphate buffered saline for 3 times, the biotin-labeled secondary antibody (Beyotime Institute of Biotechnology, Shanghai, China) and streptavidin-peroxidase (Beyotime Institute of Biotechnology, Shanghai, China) were used to incubate the specimen. The visualization of slides was achieved by interaction with 3,3'-diaminobenzidine substrate.

### Evaluation of IHC results

The results of IHC were blindly evaluated by two senior pathologists unaware of the clinical data. According to previous reports [18], the score

**Table 2.** Correlations between clinicopathological factors and SMYD3 expression

| Characters          | SMYD3 |      | P*    |
|---------------------|-------|------|-------|
|                     | Low   | High |       |
| Sex                 |       |      |       |
| Male                | 41    | 40   | 0.728 |
| Female              | 49    | 43   |       |
| Age                 |       |      |       |
| <60                 | 31    | 41   | 0.046 |
| ≥60                 | 59    | 42   |       |
| Tumor diameter (cm) |       |      |       |
| ≤5                  | 41    | 28   | 0.112 |
| >5                  | 49    | 55   |       |
| T stage             |       |      |       |
| T1+T2               | 49    | 28   | 0.006 |
| T3+T4               | 41    | 55   |       |
| Lymph node invasion |       |      |       |
| No (N0)             | 59    | 46   | 0.173 |
| Yes (N1/2)          | 31    | 37   |       |
| TNM stage           |       |      |       |
| I                   | 33    | 17   | 0.060 |
| II                  | 26    | 29   |       |
| III                 | 31    | 37   |       |
| Differentiation     |       |      |       |
| Good                | 44    | 53   | 0.140 |
| Moderate            | 29    | 19   |       |
| Poor                | 17    | 11   |       |

\*means calculated by  $\chi^2$  test. Abbreviations: SMYD3 = SET and MYND domain-containing protein 3.

system of IHC consists of two parts: the score of IHC positive cell percentage and the score of staining intensity. The score of IHC positive cell percentage was defined as follows: 1, less than 25% of positive cells; 2, 25%-50% of positive cells; 3, 50%-75% of positive cells; and 4, more than 75% of positive cells. The score of staining intensity was described as: 0 for negative staining, 1 for weak staining, 2 for moderate staining, and 3 for strong staining. The receiver operating characteristic (ROC) curve was fitted to identify the point with the highest sum of sensitivity and specificity, which was defined as the cut-off of IHC score [19]. The cut-off divided the patients into high- and low-expression of SMYD3.

### Statistical analysis

All data were analyzed with software SPSS 22.0 (IBM cooperation, USA). The correlation

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**Table 3.** Correlation between SMYD3 and clinicopathological factors

| Characters          | 5-year survival rate | P*    |
|---------------------|----------------------|-------|
| Sex                 |                      |       |
| Male                | 71.8                 | 0.414 |
| Female              | 61.1                 |       |
| Age                 |                      |       |
| <60                 | 59.5                 | 0.574 |
| ≥60                 | 70.8                 |       |
| Tumor diameter (cm) |                      |       |
| ≤5                  | 72.5                 | 0.312 |
| >5                  | 58.8                 |       |
| T stage             |                      |       |
| T1+T2               | 78.9                 | 0.003 |
| T3+T4               | 52.5                 |       |
| Lymph node invasion |                      |       |
| No (N0)             | 74.0                 | 0.008 |
| Yes (N1/2)          | 54.7                 |       |
| TNM stage           |                      |       |
| I                   | 87.6                 | 0.002 |
| II                  | 56.5                 |       |
| III                 | 54.7                 |       |
| Differentiation     |                      |       |
| Good                | 70.4                 | 0.059 |
| Moderate            | 66.6                 |       |
| Poor                | 50.6                 |       |
| Adjuvant therapy    |                      |       |
| No                  | 44.0                 | 0.015 |
| Yes                 | 73.2                 |       |
| SMYD3               |                      |       |
| Low                 | 75.8                 | 0.010 |
| High                | 50.4                 |       |

\*means calculated by log-rank test. Abbreviations: SMYD3 = SET and MYND domain-containing protein 3.

between the expression of SMYD3 and the clinicopathological factors was analyzed with Chi-square test. The survival curve was displayed by Kaplan-Meier test and the statistical difference of subgroups was analyzed with log-rank test. Independent prognostic factors were identified by the Cox proportional hazards regression model.  $P < 0.05$  was considered as statistically significant.

### Results

#### *Expression of SMYD3 in CRC tissues*

The expression and location of SMYD3 in CRC tissues was explored by the IHC. As a histone

methyltransferases, SMYD3 was mainly observed in the nucleus in our experiment, which is corresponding to its function. According to the expression of SMYD3, the cohort was divided into subgroups of high SMYD3 expression and low expression. The representative images for SMYD3 low-expression group and high-expression were displayed in **Figure 1A** and **1B**. The percentage of high SMYD3 expression and low expression accounts for 47.98% and 52.02% respectively (**Table 1**).

#### *Correlation between SMYD3 and clinicopathological factors of CRC*

The  $\chi^2$  test was performed to analyze the correlation between SMYD3 expression and clinicopathological factors of CRC (**Table 2**). In our study, the T stage was significantly associated SMYD3 expression ( $P = 0.006$ ). Patients with high SMYD3 expression had a more possibility to have advanced T stage compared with the patients with low SMYD3 expression. Similarly, advanced TNM stage tended to be substantially associated with high expression of SMYD3, but the statistical significance was not so remarkable ( $P = 0.060$ ). Intriguingly, younger patients (<60 years old) seemed to have a change to have high expression of SMYD3, which might be partially because of the higher metabolism requiring SMYD3 function.

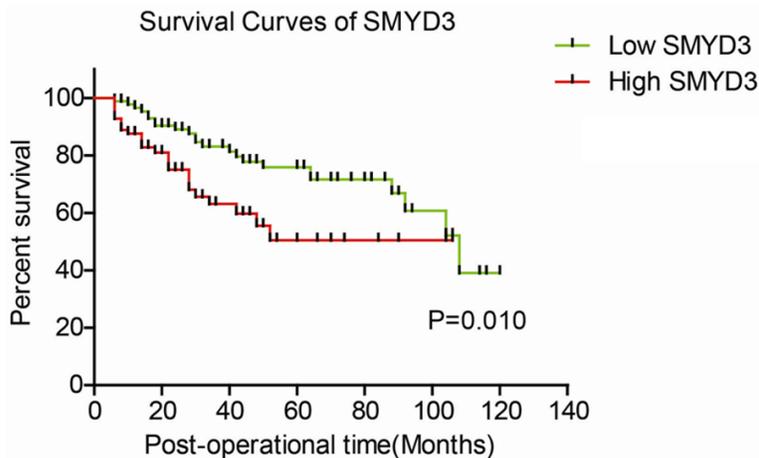
#### *Association between SMYD3 expression and the survival rates*

The univariate analysis was first carried out to examine the significant factors associated with the overall survival rates (**Table 3**). In our experiments, high expression of SMYD3 was significantly associated with lower survival rates ( $P = 0.010$ ). The 5 year overall survival rates of high expression and low expression of SMYD3 were 50.4% and 75.8% (**Figure 2**). Moreover, advanced T stage and TNM stage, positive lymphatic invasion and no adjuvant therapy were all proved to be remarkably associated with the overall survival rates of TNM I-III CRC ( $P = 0.003$ ,  $0.002$ ,  $0.008$  and  $0.015$ , respectively). Levels of tumor differentiation also seemed to be a prognostic factor with a statistical insignificant value ( $P = 0.060$ ).

#### *Independent prognostic factor of CRC*

The multivariate analysis was also performed to identify the independent prognostic factors

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**Figure 2.** Overall survival curve of SMYD3. The overall survival curve of low-expression and high-expression of SMYD3 was drawn with Kaplan-Meier method and the statistical difference was calculated by log-rank test. Patients with high SMYD3 expression had significantly low survival rates compared with those with low expression of SMYD3.

**Table 4.** Multivariate analysis

| Characters          | HR   | 95% CI    | P*    |
|---------------------|------|-----------|-------|
| T stage             |      |           |       |
| T1+T2               | 1    |           |       |
| T3+T4               | 2.23 | 1.17-4.24 | 0.015 |
| Lymph node invasion |      |           |       |
| No (N0)             | 1    |           |       |
| Yes (N1/2)          | 1.65 | 0.92-2.97 | 0.096 |
| Differentiation     |      |           |       |
| Good                | 1    |           |       |
| Moderate            | 1.44 | 0.72-2.88 | 0.308 |
| Poor                | 3.04 | 1.41-6.56 | 0.005 |
| Adjuvant therapy    |      |           |       |
| Yes                 | 1    |           |       |
| No                  | 2.23 | 1.19-4.17 | 0.012 |
| SMYD3               |      |           |       |
| Low                 | 1    |           |       |
| High                | 1.98 | 1.06-3.70 | 0.032 |

\*means calculated by Cox-regression model. Abbreviations: SMYD3 = SET and MYND domain-containing protein 3. HR = hazard ratio, 95% CI = 95% confidence interval.

of CRC. All the defined prognostic parameters were enrolled into the Cox-regression hazard model, and tumor differentiation was also selected because of its tendency to affect the prognosis (**Table 4**). TNM stage was excluded because it had obvious interaction with T stage and lymphatic status. High expression of SMYD3 was identified to be an independent

prognostic factor of CRC in TNM stage I-III (P=0.032, HR=1.98, 95% CI=1.06-3.70). Additionally, the advanced T stage, poor differentiation and no adjuvant therapy was confirmed to be high risk for prognosis of patients with CRC (P=0.015, 0.005 and 0.032, respectively).

### Discussion

According previous studies, SMYD3 could promote cancer progression by activating the transcription of multiple target genes via making chromatin more accessible via catalyzing histone methylation, or interacting the promoter of target genes and initiating transcrip-

tion by associating with RNA polymerase II and RNA helicase [15, 20]. In carcinogenesis and cancer progression, there are many genes regulated by SMYD3. Many downstream genes of SMYD3 have been discovered including 15-LOX-1, Nkx2.8, RIZ1, c-Met, WNT10B, MMP9, and androgen receptor [21-23]. Their transcription was regulated by the function and expression of SMYD3. On the contrary, SMYD3 expression was also reported to be elevated in presence of KRAS mutation in several kinds of cancers such as rectal cancer and lung adenocarcinoma [24]. In CRC, previous studies demonstrated that SMYD3 could promote the cell proliferation of CRC cell lines [8]. In our study, we demonstrated that SMYD3 overexpression was an independent biomarker indicating poorer prognosis of CRC. The genes and proteins regulated by SMYD3 and the underlying mechanisms are still unknown. We hope more interests could be focused on SMYD3 onco-function in CRC, helping explore the signaling network downstream of SMYD3.

CRC is still the leading cause of cancer deaths in developed countries. The application of adjuvant therapies like chemotherapy increases the overall survival rate of patients with CRC remarkably. More effective drugs especially target drugs are still in urgent need for the treatment of CRC. The aberrant SMYD3 level could regulate the epigenetic transcription of target genes and exhibit the tumorigenic effects, mak-

ing SMYD3 a hotspot in cancer treatment. Based on this oncogenic function of SMYD3, it is considered as a drug target for novel anticancer agents [25]. The inhibitor of SMYD3 such as BCI-121 has been demonstrated to impair cell growth of SMYD3-overexpressing solid cancer [26]. Several other compounds like EPZ030456 or GSK2807 were also proved to inhibit the SMYD3 catalytic activity with experiments in vitro. However, more animal experiments or clinical trials should be carried out to push the proceeding of SMYD3 to be the drug target.

In summary, we detected the expression of SMYD3 in 173 patients with CRC in TNM stage I-III, and demonstrated SMYD3 high-expression to be a high risk for poorer prognosis of CRC. Our findings suggested that detecting SMYD3 may help stratify patients by risk more precisely and help make the individual treatment strategy.

#### Disclosure of conflict of interest

None.

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#### References

- [1] Ferlay J, Soerjomataram I, Dikshit R, Eser S, Mathers C, Rebelo M, Parkin DM, Forman D and Bray F. Cancer incidence and mortality worldwide: sources, methods and major patterns in GLOBOCAN 2012. *Int J Cancer* 2015; 136: E359-386.
- [2] Liu H, Xu Y, Zhang Q, Yang H, Shi W, Liu Z, Li K, Gong Z, Ning S, Li S and Chen Y. Prognostic significance of TBL1XR1 in predicting liver metastasis for early stage colorectal cancer. *Surg Oncol* 2017; 26: 13-20.
- [3] Akin H and Tozun N. Diet, microbiota, and colorectal cancer. *J Clin Gastroenterol* 2014; 48 Suppl 1: S67-69.
- [4] Lee YC, Lee YL, Chuang JP and Lee JC. Differences in survival between colon and rectal cancer from SEER data. *PLoS One* 2013; 8: e78709.
- [5] Liu H, Liu Z, Li K, Li S, Song L, Gong Z, Shi W, Yang H, Xu Y, Ning S, Ismail S and Chen Y. TBL1XR1 predicts isolated tumor cells and micrometastasis in patients with TNM stage I/II colorectal cancer. *J Gastroenterol Hepatol* 2017; [Epub ahead of print].
- [6] Liu H, Xu Y, Zhang Q, Li K, Wang D, Li S, Ning S, Yang H, Shi W, Liu Z and Chen Y. Correlations between TBL1XR1 and recurrence of colorectal cancer. *Sci Rep* 2017; 7: 44275.
- [7] Wolpin BM and Mayer RJ. Systemic treatment of colorectal cancer. *Gastroenterology* 2008; 134: 1296-1310.
- [8] Hamamoto R, Furukawa Y, Morita M, Iimura Y, Silva FP, Li M, Yagyu R and Nakamura Y. SMYD3 encodes a histone methyltransferase involved in the proliferation of cancer cells. *Nat Cell Biol* 2004; 6: 731-740.
- [9] Van Aller GS, Reynoird N, Barbash O, Huddleston M, Liu S, Zmoos AF, McDevitt P, Sinnamon R, Le B, Mas G, Annan R, Sage J, Garcia BA, Tummino PJ, Gozani O and Kruger RG. Smyd3 regulates cancer cell phenotypes and catalyzes histone H4 lysine 5 methylation. *Epigenetics* 2012; 7: 340-343.
- [10] Xu S, Wu J, Sun B, Zhong C and Ding J. Structural and biochemical studies of human lysine methyltransferase Smyd3 reveal the important functional roles of its post-SET and TPR domains and the regulation of its activity by DNA binding. *Nucleic Acids Res* 2011; 39: 4438-4449.
- [11] Rea S, Eisenhaber F, O'Carroll D, Strahl BD, Sun ZW, Schmid M, Opravil S, Mechtler K, Ponting CP, Allis CD and Jenuwein T. Regulation of chromatin structure by site-specific histone H3 methyltransferases. *Nature* 2000; 406: 593-599.
- [12] Gottlieb PD, Pierce SA, Sims RJ, Yamagishi H, Weihe EK, Harriss JV, Maika SD, Kuziel WA, King HL, Olson EN, Nakagawa O and Srivastava D. Bop encodes a muscle-restricted protein containing MYND and SET domains and is essential for cardiac differentiation and morphogenesis. *Nat Genet* 2002; 31: 25-32.
- [13] Liu Y, Deng J, Luo X, Pan Y, Zhang L, Zhang R and Liang H. Overexpression of SMYD3 was associated with increased STAT3 activation in gastric cancer. *Med Oncol* 2015; 32: 404.
- [14] Liu TT, Xu H, Gao WP, Zhang SX, Zhou XH, Tang J and Liu QN. SET and MYND domain-containing protein 3 (SMYD3) polymorphism as a risk factor for susceptibility and poor prognosis in ovarian cancer. *Med Sci Monit* 2016; 22: 5131-5140.
- [15] Sarris ME, Moulos P, Haroniti A, Giakountis A and Talianidis I. Smyd3 is a transcriptional potentiator of multiple cancer-promoting genes and required for liver and colon cancer development. *Cancer Cell* 2016; 29: 354-366.
- [16] Liu N, Zhang J, Sun S, Yang L, Zhou Z, Sun Q and Niu J. Expression and clinical significance of fibroblast growth factor 1 in gastric adenocarcinoma. *Onco Targets Ther* 2015; 8: 615-621.

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- [17] Xu YF, Yang XQ, Lu XF, Guo S, Liu Y, Iqbal M, Ning SL, Yang H, Suo N and Chen YX. Fibroblast growth factor receptor 4 promotes progression and correlates to poor prognosis in cholangiocarcinoma. *Biochem Biophys Res Commun* 2014; 446: 54-60.
- [18] Xu Y, Yang X, Li Z, Li S, Guo S, Ismail S, Liu H, Huang Z, Zhang Z, Chen Y and Sun Q. Sprouty2 correlates with favorable prognosis of gastric adenocarcinoma via suppressing FGFR2-induced ERK phosphorylation and cancer progression. *Oncotarget* 2017; 8: 4888-4900.
- [19] Xu YF, Ge FJ, Han B, Yang XQ, Su H, Zhao AC, Zhao MH, Yang YB and Yang J. High-mobility group box 1 expression and lymph node metastasis in intrahepatic cholangiocarcinoma. *World J Gastroenterol* 2015; 21: 3256-3265.
- [20] Kim H, Heo K, Kim JH, Kim K, Choi J and An W. Requirement of histone methyltransferase SMYD3 for estrogen receptor-mediated transcription. *J Biol Chem* 2009; 284: 19867-19877.
- [21] Zou JN, Wang SZ, Yang JS, Luo XG, Xie JH and Xi T. Knockdown of SMYD3 by RNA interference down-regulates c-Met expression and inhibits cells migration and invasion induced by HGF. *Cancer Lett* 2009; 280: 78-85.
- [22] Liu C, Xu D, Han H, Fan Y, Schain F, Xu Z, Claesson HE, Bjorkholm M and Sjoberg J. Transcriptional regulation of 15-lipoxygenase expression by histone h3 lysine 4 methylation/demethylation. *PLoS One* 2012; 7: e52703.
- [23] Cock-Rada AM, Medjkane S, Janski N, Yousfi N, Perichon M, Chaussepied M, Chluba J, Langsley G and Weitzman JB. SMYD3 promotes cancer invasion by epigenetic upregulation of the metalloproteinase MMP-9. *Cancer Res* 2012; 72: 810-820.
- [24] Colon-Bolea P and Crespo P. Lysine methylation in cancer: SMYD3-MAP3K2 teaches us new lessons in the Ras-ERK pathway. *Bioessays* 2014; 36: 1162-1169.
- [25] Rajajeyabalachandran G, Kumar S, Murugesan T, Ekambaram S, Padmavathy R, Jegatheesan SK, Mullangi R and Rajagopal S. Therapeutical potential of deregulated lysine methyltransferase SMYD3 as a safe target for novel anticancer agents. *Expert Opin Ther Targets* 2017; 21: 145-157.
- [26] Peserico A, Germani A, Sanese P, Barbosa AJ, di Virgilio V, Fittipaldi R, Fabini E, Bertucci C, Varchi G, Moyer MP, Caretti G, del Rio A and Simone C. A SMYD3 small-molecule inhibitor impairing cancer cell growth. *J Cell Physiol* 2015; 230: 2447-2460.