Abstract. Long non-coding (Inc)RNA heart and neural crest derivatives expressed 2-antisense RNA 1 (HAND2-AS1), an IncRNA antisense transcript adjacent to the heart and neural crest derivatives expressed 2 protein (HAND2), inhibits the development of several types of cancer; however, its role in melanoma is currently unknown. In the present study, it was revealed that lncRNA HAND2-AS1 was downregulated, whereas Rho-associated protein kinase 1 (ROCK1) mRNA was upregulated in tumor tissues when compared with the healthy tissues of patients with melanoma. Expression levels of lncRNA HAND2-AS1 and ROCK1 were inversely correlated in tumor tissues, but not in healthy tissues. LncRNA HAND2-AS1 expression levels were affected by tumor thickness, but not by tumor metastasis. LncRNA HAND2-AS1 overexpression led to inhibited melanoma cancer cell proliferation and an increased expression level of ROCK1. ROCK1 overexpression caused no evident effects on lncRNA HAND2-AS1 expression, but promoted cancer cell proliferation and decreased the effects of lncRNA HAND2-AS1 overexpression on cancer cell proliferation. Thus, it is possible that lncRNA HAND2-AS1 overexpression leads to inhibited cancer cell proliferation in melanoma cells through the downregulation of ROCK1.

Introduction
Melanoma is a type of malignant tumor that develops from melanocytes, which are known as pigment-containing cells (1). Melanoma is one of the most common types of cancer in the USA with an incidence rate of 1 in 100 (2). By contrast, the incidence rate of the disease in China is relatively low, with only 3 in 20,000 developing this malignancy during their lifetime (3). Survival conditions of patients with early-stage melanoma following surgical resection are generally satisfactory (4); however, those patients with melanoma at advanced stages of the disease typically exhibit a high prevalence of brain metastases, which is a major cause of mortality (5,6). Furthermore, surgical resection of the primary tumor affects the appearance of patients, which in turn affects their quality of life (7).

As a member of the Rho-associated protein kinase (ROCK) family, ROCK1 regulates cell contraction by regulating calcium ion intake (8). Previous studies have indicated that ROCK1 also serves a critical role in the development of different types of cancer, including melanoma (9,10). As a result, inhibition of ROCK1 expression is considered to be a promising therapeutic target for cancer treatment (10,11). Previous studies have indicated that ROCK1 participates in cancer biology not only by interacting with proteins, but also through crosstalk with non-coding RNAs, such as long non-coding (Inc)RNAs (12,13). LncRNA heart and neural crest derivatives expressed 2-antisense RNA 1 (HAND2-AS1), an IncRNA antisense transcript adjacent to the heart and neural crest derivatives expressed 2 protein (HAND2), is a well-characterized tumor suppressor in several types of malignancy (14-16); however, its involvement in melanoma is currently unknown. Therefore, the present study aimed to investigate the involvement of HAND2-AS1 in melanoma.

Materials and methods
Cell lines and human tissues. The two human melanoma cell lines C32 and SK-MEL-28 were purchased from the American Type Culture Collection (ATCC; Manassas, VA, USA). The two cell lines were cultured in ATCC-formulated Eagle's minimum essential medium (catalog no. 30-2003) supplemented with fetal bovine serum to a concentration of 10% in an incubator at 37°C containing 5% CO₂.

Tumor tissue and adjacent healthy tissue (within 2 cm around tumors) biopsies were obtained from 56 patients with melanoma who were admitted to The First Hospital of Jilin University (Changchun, China) between May 2015 and June 2018. Tumor tissues and adjacent healthy tissues
were examined and confirmed by three experienced pathologists. The inclusion criteria were as follows: i) Patients diagnosed via pathological biopsies; ii) patients had not been treated before admission; and iii) patients understood the experimental principle and were willing to participate. Exclusion criteria were: i) Patients diagnosed with multiple diseases, such as chronic diseases; and ii) patients refused to donate tissue samples. The 56 patients included 31 men and 25 women, and had an age range of between 26 and 67 years, with a mean age of 47.2±5.1 years. There were 12 cases in stage I, 16 in stage II (no metastasis), 18 in stage III (lymph node metastasis) and 10 in stage IV (distant metastasis). All patients provided written informed consent. The present study was approved by the ethics committee of The First Hospital of Jilin University.

Reverse transcription-quantitative polymerase chain reaction (RT-qPCR). RNAzol® RT RNA Isolation Reagent (GeneCopoeia, Inc., Rockville, MD, USA) was used to extract total RNA from tumor tissues, adjacent healthy tissues and *in vitro* cultured cells. RNA concentration was determined using a NanoDrop™ 2000 spectrophotometer (Thermo Fisher Scientific, Inc., Waltham, MA, USA) and reverse transcription was performed using a RevertAid RT Reverse Transcription kit (Thermo Fisher Scientific, Inc., Waltham, MA, USA) and reverse transcription protocol. Cells treated only with Lipofectamine 3000 without vectors were used as control cells. Cells transfected with empty vectors were used as negative control cells. Expression of HAND2-AS1 and ROCK1 was detected using *in vitro* transfection.

**Vectors and cell transfection.** Vectors expressing HAND2-AS1 and ROCK1 were provided by GeneCopoeia, Inc. All cell transfections were performed using Lipofectamine® 3000 (Invitrogen; Thermo Fisher Scientific, Inc.) with all operations performed in strict accordance with the manufacturer’s protocol. Cells treated only with Lipofectamine 3000 without the presence of vectors were used as control cells. Cells transfected with empty vectors were used as negative control cells. Expression of HAND2-AS1 and ROCK1 was detected using RT-qPCR 24 h after transfection as aforementioned.

**Cell proliferation assay.** At 24 h following transfection, cell proliferation ability was tested using a Cell Counting Kit-8 (CCK-8) assay (Beyotime Institute of Biotechnology, Haimen, China). Briefly, cell suspensions (3×10⁴ cells/ml) were prepared using ATCC-formulated Eagle's minimum essential medium containing 10% fetal bovine serum. In cases of ROCK1 inhibitor treatment, cells were incubated with 10 µM ROCK1 inhibitor AT13148 (cat. no. S7563, Selleck Chemicals, Houston, TX, USA) for 24 h at 37°C before use. Each well of a 96-well plate was filled with 0.1 ml cell suspension containing 3×10⁴ cells. Cells were cultured in an incubator at 37°C, containing 5% CO₂ and 10 µl CCK-8 solution was added after 24, 28, 72 and 96 h. Cells were cultured under the same conditions for a further 4 h, followed by determining optical density values at 450 nm using a microplate reader (Bio-Rad Laboratories, Inc.).

**Western blot analysis.** Protein Extraction kit (catalog no. NP02-37853; Novus Biologicals LLC, Littleton, CO, USA) was used to extract total protein from *in vitro* cultured cells. A bicinechonic acid Protein Quantiﬁcation kit (catalog no. ab102536; Abcam, Cambridge, UK) was used to determine protein concentrations with all steps performed in strict accordance with the manufacturer’s protocol. Protein samples were denatured and subjected to SDS-PAGE (10% gels; 45 µg per lane). Following protein transfer onto polyvinylidene difluoride membranes, membranes were blocked in 5% fat-free milk in PBS at room temperature for 2 h. The primary antibodies used were rabbit anti-human ROCK1 (1:1,400; catalog no. ab97592; Abcam) and rabbit anti-human GAPDH (1:1,400; catalog no. ab9485; Abcam) and the incubation was performed at 4°C for 15 h. A secondary incubation was performed at 24°C for 2 h using horseradish peroxidase-conjugated goat anti-rabbit immunoglobulin G secondary antibody (1:1,200; catalog no. MBS345036; MyBioSource, Inc., San Diego, CA, USA). An ECL™ Western Blotting Analysis system (Sigma-Aldrich; Merck KGaA, Darmstadt, Germany) was used to develop signals. ImageJ software (version 1.46; National Institutes of Health, Bethesda, MD, USA) was used to normalize the data.

**Statistical analysis.** All experiments were performed in triplicate. Data were expressed as the mean ± standard deviation and processed by GraphPad Prism version 6 (GraphPad Software, Inc., La Jolla, CA, USA). Correlation analyses were performed using Pearson’s correlation coefficient. Comparisons between tumor tissues and adjacent healthy tissues were performed using a paired t-test. Comparisons among multiple groups were performed using one-way analysis of variance, followed by Tukey’s post hoc test. P<0.05 was considered to indicate a statistically significant difference.

**Results**

**LncRNA HAND2-AS1 and ROCK1 mRNA expression are altered in tumor tissues compared with in adjacent healthy tissues.** Expression of lncRNA HAND2-AS1 and ROCK1 mRNA in tumor tissues and adjacent healthy tissues of 56 patients with melanoma was detected using RT-qPCR. When compared with tumor tissues, expression levels of lncRNA HAND2-AS1 were significantly increased in adjacent healthy tissues (P<0.05; Fig. 1A). In contrast, expression levels of ROCK1 mRNA were significantly increased in tumor tissues compared with healthy control tissues (P<0.05; Fig. 1B). In addition, the lncRNA HAND2-AS1 level slightly decreased (Fig. 1C), whereas the ROCK1 level slightly increased (Fig. 1D) with the increase in tumor stage.
Expression levels of lncRNA HAND2-AS1 and ROCK1 are inversely correlated in tumor tissues. Pearson's correlation coefficient analyses indicated that expression levels of lncRNA HAND2-AS1 and ROCK1 mRNA were significantly and inversely correlated in tumor tissues (Fig. 2A). In contrast, the correlation between expression levels of HAND2-AS1 and ROCK1 mRNA was not significant in adjacent healthy tissues (Fig. 2B).

Expression levels of lncRNA HAND2-AS1 in tumor tissues are affected by tumor thickness but not by tumor metastasis. There were 22 cases of patients with a primary tumor thickness of <2 mm, 18 between 2 and 4 mm, and 16 with >4 mm. As presented in Fig. 3A, expression levels of lncRNA HAND2-AS1 in tumor tissues decreased with the increase in tumor thickness (P<0.05). There were 28 cases of non-metastasis, 18 cases of lymph node metastasis and 10 cases of distant metastasis. As presented in Fig. 3B, there were no significant differences in expression levels of lncRNA HAND2-AS1 among these three groups.

LncRNA HAND2-AS1 overexpression results in ROCK1 downregulation in cells of human melanoma cell lines. Following transfection, overexpression of HAND2-AS1 and
ROCK1 was achieved in the C32 and SK-MEL-28 human melanoma cell lines (P<0.05; Fig. 4A). Compared with the control group and negative control groups, HAND2-AS1 overexpression resulted in significantly downregulated ROCK1 expression in the two cell lines (P<0.05; Fig. 4B). In contrast, no significant differences in expression levels of lncRNA HAND2-AS1 were identified among the control, negative control and ROCK1 overexpression groups (P<0.05; Fig. 4C).

lncRNA HAND2-AS1 overexpression results in inhibited proliferation of melanoma cells possibly through ROCK1. Compared with the control and negative control groups, HAND2-AS1 overexpression and treatment with ROCK1 inhibitor AT13148 resulted in clear inhibition, whereas ROCK1 overexpression led to significantly increased proliferation of C32 (Fig. 5A) and SK-MEL-28 (Fig. 5B) human melanoma cell lines (P<0.05). In addition, cells with HAND2-AS1 and ROCK1 overexpression exhibited significantly promoted
proliferation compared with cells with HAND2-AS1 overexpression alone (P<0.05).

Discussion

The functionality of lncRNA HAND2-AS1 has been characterized in osteosarcoma, colorectal cancer and endometrioid endometrial carcinoma (14-16). It is generally considered that lncRNA HAND2-AS1 serves a role in tumor suppression in those cancer types by inhibiting cancer progression through interactions with different signaling pathways. The key result of the present study is that lncRNA HAND2-AS1 is also a tumor suppressor in melanoma and inhibits cancer cell proliferation. The actions of lncRNA HAND2-AS1 are likely to be mediated by interactions with ROCK1.

As an oncogene, ROCK1 is typically upregulated during cancer development (18,19). Consistent with previous studies (18,19), the results of the present study indicated that expression of ROCK1 was significantly increased in tumor tissues compared with in adjacent healthy tissues of patients with melanoma. It has been reported that overexpression of ROCK1 is essential for the growth of certain types of tumor (20). In the present study, it was observed that ROCK1 overexpression significantly promoted the proliferation of melanoma cells, further confirming the oncogenic role of ROCK1 in melanoma.

Previous studies have indicated that differentially expressed lncRNAs are key factors in the progression of human cancer types, including melanoma (21,22). As a tumor suppressor, lncRNA HAND2-AS1 has been revealed to be downregulated in osteosarcoma, colorectal cancer and endometrioid endometrial carcinoma (14-16). In the present study, it was identified that lncRNA HAND2-AS1 was significantly downregulated in tumor tissues when compared with the adjacent healthy tissues of patients with melanoma. Notably, expression of lncRNA HAND2-AS1 in tumor tissues was affected by tumor thickness, but not by tumor metastasis, indicating that lncRNA HAND2-AS1 is unlikely to be a key regulator of cancer metastasis in melanoma. However, lncRNA HAND2-AS1 was reported previously to inhibit cancer cell invasion and metastasis in endometrioid endometrial carcinoma (16). Therefore, lncRNA HAND2-AS1 may serve a number of roles in different types of human malignancy.

In vitro cell experiments revealed that lncRNA HAND2-AS1 and ROCK1 serve opposite roles in melanoma cell proliferation. In addition, lncRNA HAND2-AS1 is likely to be an upstream inhibitor of ROCK1 in the regulation of cancer cell proliferation; however, the molecular mechanism underlying the inhibitory effects of lncRNA HAND2-AS1 on ROCK1 is currently unknown. We hypothesize that lncRNA HAND2-AS1 may inhibit ROCK1 expression through disease-associated factors due to the observation that expression levels of lncRNA HAND2-AS1 on ROCK1 were significantly and inversely correlated only in tumor tissues, but not in adjacent healthy tissues.

In conclusion, in the present study, lncRNA HAND2-AS1 was identified to be downregulated, and ROCK1 upregulated in melanoma. HAND2-AS1 overexpression may inhibit cancer cell proliferation in melanoma cells by downregulating ROCK1. Therefore, HAND2-AS1 overexpression may serve as a potential therapeutic target for melanoma.

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Availability of data and materials

The datasets used and/or analyzed during the present study are available from the corresponding author on reasonable request.
Authors' contributions

PL and XY designed the experiments; PL and RD performed the experiments and analyzed all of the data. XY drafted the manuscript and all authors approved the final manuscript.

Ethics approval and consent to participate

Ethical approval was obtained from the Ethics Committee of The First Hospital of Jilin University, Changchun, China. All procedures performed in the parts of the study involving human participants were in accordance with the ethical standards of the institutional and national research committee and with The Declaration of Helsinki (1964) and its later amendments or comparable ethical standards.

Patient consent for publication

The present study followed the tenets of The Declaration of Helsinki, and informed written consent was obtained from all patients and controls following explanation of the nature and possible consequences of the study.

Competing interests

The authors declare that they have no competing interests.

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