An Anticancer Effect of Curcumin Mediated by Down-Regulating Phosphatase of Regenerating Liver-3 Expression on Highly Metastatic Melanoma Cells

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ABSTRACT

Phosphatase of regenerating liver-3 (PRL-3) has been suggested as a potential target for anticancer drugs based on its involvement in tumor metastasis. However, little is known about a small-molecule inhibitor against PRL-3. In this study, we report that curcumin, the component of the spice turmeric, shows its antitumor effect by selectively down-regulating the expression of PRL-3 but not its family members PRL-1 and -2 in a p53-independent way. Curcumin inhibited the phosphorylation of Src and stat3 partly through PRL-3 down-regulation. Cells with PRL-3 stably knocked down show less sensitivity to curcumin treatment, which reveals that PRL-3 is the much further upstream target of curcumin. Curcumin treatment also remarkably prevented B16BL6 from invading the draining lymph nodes in the spontaneous metastatic tumor model, which is probably of relevance to PRL-3 down-regulation. Our results reveal a novel capacity of curcumin to down-regulate oncogene PRL-3, raising its possibility in therapeutic regimen against malignant tumor.

The phosphatase of regenerating liver (PRL) as a tyrosine phosphatase family includes 3 members: PRL-1, PRL-2, and PRL-3. In 2001, PRL-3 was first reported to be overexpressed in metastatic lesions derived from colorectal cancers, but it was expressed at lower levels in primary tumors and normal colorectal epithelium (Saha et al., 2001). The elevated PRL-3 expression was then found in other highly metastatic cancers such as gastric carcinomas (Miskad et al., 2004), Hodgkin’s lymphoma (Schwering et al., 2003), melanomas (Wu et al., 2004), and breast (Polato et al., 2005) and ovarian tumors (Zeng et al., 2003), suggesting that PRL-3 may be a molecular marker for metastatic tumor cells. Indeed, several in vitro and in vivo studies support a causal link between PRL-3 and tumor metastasis. Overexpressing PRL-3 promotes motility and invasion of both tumor cell lines and normal cell lines (Zeng et al., 2003; Wu et al., 2004), whereas knocking down endogenous PRL-3 with small interfering RNA attenuates cancerous cell motility and metastatic tumor formation (Qian et al., 2007). Treatment with monoclonal antibody of PRL-3 massively inhibited the tumor growth in vivo (Li et al., 2005; Guo et al., 2008). Therefore, PRL-3 is considered a tractable target for anticancer drugs, and regulating its expression and function may become a new strategy to prevent or treat tumor metastasis. However, there is no report on the natural small-molecule compounds that can regulate PRL-3 expression.

Curcumin is a polyphenol derived from dietary spice turmeric. It possesses wide-ranging anti-inflammatory and anticancer properties (Sharma et al., 2005). The abilities of curcumin to induce apoptosis of cancer cells and to inhibit angiogenesis and cell adhesion contribute to its chemotherapeutic potential in the treatment of cancer. Several phase I and phase II clinical trials indicate that curcumin is quite safe and may exhibit therapeutic efficacy in patients with progressive advanced cancers (Dhillon et al., 2008). Although inhibition of several cell signaling pathways involving Akt (Woo et al., 2003), nuclear factor-κB (Aggarwal et al., 2006), activator protein-1 (Balasubramanian and Eckert, 2007), or c-Jun N-terminal kinase (Chen and Tan, 1998) have been implicated in the biological effects of curcumin, its direct molecular target and mechanism of inhibition in tumor metastasis remain to be well clarified.

ABBREVIATIONS: PRL, phosphatase of regenerating liver; DMEM, Dulbecco’s modified Eagle’s medium; DMSO, dimethyl sulfoxide; FBS, fetal bovine serum; GAPDH, glyceraldehyde-3-phosphate dehydrogenase; PBS, phosphate-buffered saline; PCR, polymerase chain reaction; siRNA, small interfering RNA; RT-PCR, reverse transcription-polymerase chain reaction; bp, base pair.
In the present study, we showed a novel activity of curcumin in this first report of specific down-regulation of PRL-3 expression, which contributes to the in vivo antimetastatic effect of curcumin. Such activity of curcumin was further demonstrated to be at the transcriptional level without affecting the stability of either PRL-3 mRNA or protein and to result in the inhibition of the phosphorylation of Src and Stat3.

Materials and Methods

Reagents. Curcumin (>98%) was purchased from Shanghai R&D Centre for Standardization of Traditional Chinese Medicine (Shanghai, China), and the stock solution was prepared with dimethyl sulfoxide (DMSO). Cycloheximide and actinomycin D were purchased from Sigma-Aldrich (St. Louis, MO), and dissolved at 5 mg/ml in PBS and cycloheximide dimethyl sulfoxide, respectively.

Animals. C57BL/6J mice (6–8 weeks old) were obtained from the Shanghai Laboratory Animal Center (Shanghai, China). Throughout the experiments, mice were maintained with free access to pellet food and water in plastic cages at 21 ± 2°C and kept on a 12-h light/dark cycle. Animal welfare and experimental procedures were performed strictly in accordance with the Guide for the Care and Use of Laboratory Animals (Institute of Laboratory Animal Resources, 1996) and the related ethical regulations of China (Ministry of Science and Technology, 2006). All efforts were made to minimize the animals’ suffering and to reduce the number of animals used.

Cell Culture. B16 and B16BL6 mouse melanoma cells, EMT-6 mouse breast carcinoma cells, PC3 human prostate cancer cells, and MCF-7 human breast carcinoma cells were purchased from the American Type Culture Collection (Manassas, VA). Murine embryonic fibroblasts were generated from embryonic day 13.5 embryos. All of the cells used were maintained in Dulbecco’s modified Eagle’s medium (DMEM) (Invitrogen, Carlsbad, CA) supplemented with 10% fetal bovine serum (FBS) (Invitrogen), 100 U/ml penicillin, and 100 μg/ml streptomycin and incubated at 37°C in a humidified atmosphere containing 5% CO2.

RT-PCR and Real-Time PCR. RNA samples were treated by DNase and subjected to semiquantitative RT-PCR. First-strand cDNAs were generated by reverse transcription using oligo(dT). The cDNAs were amplified by PCR for 28 cycles (94°C for 30 s, 59°C for 30 s, and 72°C for 30 s) using Taq DNA polymerase (Promega, Shanghai, China). The PCR products were electrophoresed on a 2% agarose gel and visualized by ethidium bromide staining. The Gel Imaging and Documentation DigiDoc-It System (version 1.1.23; UVP, Inc., Upland, CA) was used to scan the gels, and the intensity of the bands was assayed using Labworks Imaging and Analysis Software (UVP, Inc.). Quantitative PCR was performed with the ABI Prism 7000 sequence detection system (Applied Biosystems, Foster City, CA) using SYBR Green I dye (Biotium, Inc., Hayward, CA), and threshold cycle numbers were obtained using ABI Prism 7000 SDS software version 1.0. Conditions for amplification were 1 cycle of 94°C for 5 min followed by 35 cycles of 94°C for 30 s, 59°C for 35 s, and 72°C for 45 s. The primer sequences used in this study were as follows: PRL-1: forward, 5′-CAACCAATGGGACATCTGA-3′; reverse, 5′-CATGGAATCATGCGACTCC-3′; PRL-2: forward, 5′-ATTGGCCAATAATGAGCGAC-3′; reverse, 5′-ACAGGCGCTTCTCCTAAT-3′; PRL-3: forward, 5′-CTCTCTCATCAAACCATACCC-3′; reverse, 5′-TACATGACCAGCATCTG-3′; GAPDH: forward, 5′-CATGGCTTCCGTGTTCCTA-3′; reverse, 5′-GCCGCACTCAGTCCA-3′. The resultant PCR products were 472 (PRL-1), 339 (PRL-2), 468 (PRL-3), and 191 bp (GAPDH).

Cell Proliferation Assay. Cells (3 × 10^4/well) were prepared in 96-well plates in serum-containing medium. After 24 h, various concentrations of curcumin were added to the plates. The cells were treated with 20 μM curcumin for various hours. At the indicated time, the mRNA level of PRL-3 was measured by real-time PCR by RT-PCR three times, and the electrophoresis presented is the representative one. GAPDH was used as an invariant control. B, B16BL6 cells were treated with 20 μM curcumin for various hours. At the indicated time, the mRNA level of PRL-3 was measured by real-time PCR and calculated by using GAPDH as an invariant control. B, B16BL6 cells were treated with different concentrations of curcumin for 12 h. Total RNA was prepared for analyses of PRL-3 expression by RT-PCR three times, and the electrophoresis presented is the representative one. GAPDH was used as an invariant control.

Fig. 1. Curcumin selectively inhibited PRL-3 expression in multiple cell lines. A, different cell lines were grown in six-well plates and were treated with different concentrations of curcumin for 12 h. Total RNA was prepared for analyses of PRL-3 expression by RT-PCR three times, and the electrophoresis presented is the representative one. GAPDH was used as an invariant control. B, B16BL6 cells were treated with 20 μM curcumin for various hours. At the indicated time, the mRNA level of PRL-3 was measured by real-time PCR and calculated by using GAPDH as an invariant control. C, B16BL6 cells were treated with various concentrations of curcumin for 12 h. The mRNA level of PRL-1, PRL-2, and PRL-3 were measured by real-time PCR and calculated by using GAPDH as an invariant control. Data are the mean ± S.E.M. of three independent experiments. *p < 0.05, and **p < 0.01 versus cells cultured without curcumin.
and normalized to that for the transfected for 12 h with the full length of mouse PRL-3 promoter with the primers as follows: PRL-3 promoter sense, 5'-tides of 5'-H11032 Dual Luciferase System (Promega). A fragment of 1986-bp nucleotide of 1240 Wang et al. was cloned into the luciferase reporter plasmid PGL-3 (Promega). With antisense, 5'-CGCAAGCTTCAACAGGCACTCAGTCAAGC-3', and pRNA-U6.1/Neo luciferase siRNA (5'-GCU-UACCGUGAGUCUCGUA-3') was used as the monitor and a negative control. B16BL6 melanoma cells were cultured to 60% confluence in a 35-mm plate and transfected with luciferase siRNA or PRL-3 siRNA using the Lipofectamine 2000 (Life Technologies). After transfection (48 h), cells were passaged to a 100-mm dish, and the cell adhesion assay was performed as described previously (Wu et al., 2004). In brief, 96-well flat-bottomed culture plates were coated with 50 µl of fibronectin (10 µg/ml) (Calbiochem, San Diego, CA) in PBS overnight at 4°C, respectively. Plates were then blocked with 0.2% bovine serum albumin for 2 h at room temperature followed by washing three times with DMEM. The cells were harvested with trypsin/EDTA, washed with ice-cold PBS twice, and resuspended in DMEM. Cells (2 x 10^4/well) were added to each well in triplicate and incubated for 30 min at 37°C. Plates were then washed three times with DMEM to remove unbonded cells. Cells remaining attached to the plates were quantified with 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium assay as described above.

Luciferase Assay. Luciferase assays were performed with the Dual Luciferase System (Promega). A fragment of 1986-bp nucleotides of 5'-untranscriptional region of PRL-3 promoter was amplified with the primers as follows: PRL-3 promoter sense, 5'-CGGCCCTAGCGGGCAAATCTCCAGTCATG-3', and antisense, 5'-GGGTAGACTCAGGATGTGGGCGATGCCCTTTTTGGCC-3', which was subcloned into the luciferase reporter plasmid PGL-3 (Promega). With the β-galactosidase reporter as an internal control, cells were co-transfected for 12 h with the full length of mouse PRL-3 promoter vectors. The cells were then exposed to curcumin for 12 h. The luciferase activity of the PRL-3 promoter reporters was determined and normalized to that for the β-galactosidase reporter. The luciferase levels of the remaining cells were calculated according to the optical density value.

Cell Adhesion Assay. The cell adhesion assay was performed as described previously (Wu et al., 2004). In brief, 96-well flat-bottomed culture plates were coated with 50 µl of fibronectin (10 µg/ml) (Calbiochem, San Diego, CA) in PBS overnight at 4°C, respectively. Plates were then blocked with 0.2% bovine serum albumin for 2 h at room temperature followed by washing three times with DMEM. The cells were harvested with trypsin/EDTA, washed with ice-cold PBS twice, and resuspended in DMEM. Cells (2 x 10^4/well) were added to each well in triplicate and incubated for 30 min at 37°C. Plates were then washed three times with DMEM to remove unbonded cells. Cells remaining attached to the plates were quantified with 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium assay as described above.

Electrophoretic Mobility Shift Assay. Nuclear proteins were extracted from B16BL6 melanoma cells using NE-PER nuclear and cytoplasmic extraction reagents (Pierce, Rockford, IL) with a protease inhibitor cocktail (Sigma-Aldrich). An oligonucleotide probe containing the p53-binding motif of the PRL-3 promoter (5'-GGGTAGACTCAGGATGTGGGCGATGCCCTTTTTGGCC-3') was prepared as was done previously by Basak et al. (2008) and then labeled with biotin by Invitrogen (Shanghai, China). Detection of p53-oligonucleotide complex was performed using a LightShift chemiluminescent electrophoretic mobility shift assay kit (Pierce). In brief, nuclear protein (10 µg) was incubated with 20 fmol biotin-labeled oligonucleotide for 20 min at room temperature in binding buffer consisting of 10 mM Tris at pH 7.5, 50 mM KCl, 1 mM dithiothreitol, 2.5% glycerol, 5 mM MgCl2, 50 ng of poly(dA-dT), and 0.05% Nonidet P-40. The specificity of the p53 DNA binding was determined in competition reactions in which a 50-fold molar excess (1 µmol) and 100-fold molar excess (2 µmol) of unlabeled oligonucleotide were added to the binding reaction, respectively. Products of binding reactions were resolved by electrophoresis on a 6% polyacrylamide gel using 0.5× Tris-borate EDTA buffer (Invitrogen). p53-oligonucleotide complex was electroblotted to a nylon membrane (Invitrogen). After incubation in blocking buffer for 15 min at room temperature, the membrane was incubated with streptavidin-horseradish peroxidase conjugate for 30 min at room temperature. The membrane was incubated with chemiluminescent substrate for 5 min and allowed to expose radiographic film.

Stably Interfering PRL-3 in B16BL6 Melanoma Cells. We followed methods described previously (Qian et al., 2007) to generate the pRNA-U6.1/Neo PRL-3 siRNA (5'-GCUCCACCUCCUGGAGAAGUA-3'), and pRNA-U6.1/Neof luciferase siRNA (5'-GCU-UACCGUGAGUCUCGUA-3') was used as the monitor and a negative control. B16BL6 melanoma cells were cultured to 60% confluence in a 35-mm plate and transfected with luciferase siRNA or PRL-3 siRNA using the Lipofectamine 2000 (Life Technologies). After transfection (48 h), cells were passaged to a 100-mm dish, and G418 sulfate (Geneticin; Sigma) was added to final concentration of 400 µg/ml. Resistant cells were allowed to grow for 10 days. Individual G418-resistant colonies were selected and screened for mono colony in 96-well plates by limited dilution. P1 and P9 were clones 1 and 9 of B16BL6 cells stably transfected with PRL-3 siRNA. L10 and L13 were selected time, cell lysates were collected, and the protein level of PRL-3 and tubulin were determined by immunoblotting at least three times; representative data are shown. The bands were semiquantified by grayscale scanning. C, B16BL6 cells were transfected with the PRL-3 promoter luciferase reporter plasmid. After transfected for 12 h, cells were incubated with different concentrations of curcumin for an additional 12 h. Luciferase activity was expressed as relative units after β-galactosidase normalization. Data are mean ± S.E.M. of three independent experiments. B, B16BL6 cells were treated with 20 µM curcumin in the presence of 5 µM/ml actinomycin D for different hours, and total RNA was extracted. The mRNA level of PRL-3 was measured by real-time PCR and calculated by using GAPDH as an invariant control. Data are the mean ± S.E.M. of three independent experiments. B, B16BL6 cells were treated with 20 µM curcumin in the presence of 5 µM/ml actinomycin D for different hours, and total RNA was extracted. The mRNA level of PRL-3 was measured by real-time PCR and calculated by using GAPDH as an invariant control. Data are the mean ± S.E.M. of three independent experiments. B, B16BL6 cells were treated with 20 µM curcumin in the presence of 5 µM/ml actinomycin D for different hours, and total RNA was extracted. The mRNA level of PRL-3 was measured by real-time PCR and calculated by using GAPDH as an invariant control. Data are the mean ± S.E.M. of three independent experiments. B, B16BL6 cells were treated with 20 µM curcumin in the presence of 5 µM/ml actinomycin D for different hours, and total RNA was extracted. The mRNA level of PRL-3 was measured by real-time PCR and calculated by using GAPDH as an invariant control. Data are the mean ± S.E.M. of three independent experiments.
were clones 10 and 13 of B16BL6 cells stably transfected with luciferase siRNA.

Western Blot. The Western blot was performed as described previously (Qian et al., 2007). The cells were collected and lysed (50 mM Tris, pH 8.0, 150 mM NaCl, 1% Nonidet P-40, 0.1% SDS, 5 mM EDTA, 0.1 mM phenylmethylsulfonyl fluoride, 0.15 μM aprotinin, 1 μg/ml pepstatin, and 10% glycerol). Anti-phosphorylation of Src (Tyr416 or Tyr527), anti-Src, anti-stat3, anti-phosphorylation of stat3 (Tyr705), anti-cyclin D1, anti-cyclin D3 (Cell Signaling Technology, Danvers, MA), anti-tubulin, anti-p53, anti-GAPDH (Santa Cruz Biotechnology, Santa Cruz, CA), and anti-PRL-3 antibodies (ProSci Inc., Poway, CA) were used for Western blot.

Cell Migration Assay. Cell migration assay was performed using 8.0-μm pore-size Transwell inserts (Corning Life Sciences, Lowell, MA). In brief, the undersurface of the membrane was coated with fibronectin, laminin (10 μg/ml; Sigma-Aldrich) in phosphate-buffered saline, pH 7.4, for 2 h at 37°C. The membrane was washed in PBS to remove excess ligand, and the lower chamber was filled with 0.6 ml of DMEM with 10% FBS. Cells were serum-starved overnight (0.5% FBS), harvested with trypsin/EDTA, and washed twice with serum-free DMEM. Cells were resuspended in the medium (DMEM with 0.5% FBS), and 1 × 10⁵ cells in 0.1 ml were added to the upper chamber. After 6 h at 37°C, the cells on the upper surface of the membrane were removed using cotton tips. The migrant cells attached to the lower surface were fixed in 10% formalin at room temperature for 30 min and stained for 20 min with a solution containing 1% crystal violet and 2% ethanol in 100 mM borate buffer, pH 9.0. The number of migrated cells on the lower surface of the membrane was counted under a microscope in five fields.

In Vivo Metastasis Assay. The B16BL6 tumor cells metastasis model was performed as reported previously (Qian et al., 2007). Twelve days after injection, the mice were distributed into three groups with six mice each according to tumor size. Curcumin, dissolved in olive oil, was given by intraperitoneal injection at dose of 50 and 100 mg/kg every 2 days, respectively. Tumor volumes were measured every 2 days from day 12 to 26 and calculated by the following formula: \(0.5236 \times L_1 \times (L_2)^{2}\), where \(L_1\) is the long axis and \(L_2\) is the short axis of the tumor. At the end of the experiment, mice were killed and the right footpads, draining popliteal lymph nodes were resected, and photos were taken.

Statistical Analysis. Data are expressed as mean ± S.E.M. Student’s t test was used to evaluate the difference between two groups.
Kaplan-Meier method was used to evaluate the survival test. \( p < 0.05 \) was considered to be significant.

**Results**

**Curcumin Selectively Down-Regulates PRL-3 Expression in Multiple Tumor Cell Lines.** Several mouse and human cell lines were treated with various concentrations of curcumin and tested for the PRL-3 mRNA level. As shown in Fig. 1A, curcumin decreased PRL-3 mRNA level in mouse melanoma B16BL6, B16, EMT-6, human breast cancer MCF-7, and human prostate cancer PC3 cells. Then, B16BL6 cells with high PRL-3 expression were chosen for further use, and a time-dependent inhibition of PRL-3 level was confirmed in curcumin-treated cells (Fig. 1B). It is noteworthy that the expression of PRL-1 and PRL-2, which have been identified as homologous to PRL-3, was hardly affected by curcumin (Fig. 1C).

**Curcumin Inhibits PRL-3 Expression at the Transcriptional Level in a p53-Independent Way.** The stability of PRL-3 mRNA in B16BL6 cells was examined after treatment with or without 20 \( \mu \)M curcumin in the presence of 5 \( \mu \)g/ml actinomycin D. Curcumin had no impact on degradation of PRL-3 mRNA (Fig. 2A). There was also no appreciable difference in PRL-3 protein stability between B16BL6 cells exposed to curcumin and those exposed to the solvent, DMSO (Fig. 2B). To evaluate whether the inhibitory effect of curcumin is at the transcriptional level, B16BL6 cells were transiently transfected with the PRL-3 promoter luciferase reporter plasmid pGL-3, in which a fragment of 5'-untranscriptional region of PRL-3 promoter was subcloned. Compared with the untreated control, curcumin decreased the PRL-3 transcriptional activity in a dose-dependent manner (Fig. 2C). Next, we examined the effect of curcumin on p53, an important transcription factor associated with PRL-3 expression (Basak et al., 2008). After B16BL6 cells were exposed to various concentrations of curcumin, no reduction in p53 protein amount was observed (Fig. 2D).

**The Anticancer Effects of Curcumin Are Related to PRL-3.** The expression of PRL-3 was verified in mouse melanoma cell lines. The highly metastatic B16BL6 cells express PRL-3 mRNA approximately 3-fold more than the lowly metastatic B16 cells, which is consistent with our previous result (Wu et al., 2004), and so does the protein level (Fig. 3A). To detect whether PRL-3 is a key target of curcumin, we generated B16BL6 cells with PRL-3 stably knocked down (P1 and P9), in which the PRL-3 level was approximately 10% of those cells transfected with the control siRNA (L10 and L13). Likewise, curcumin strongly inhibited adhesion (Fig. 3B) and migration (Fig. 3C) of cells with high PRL-3 expression, whereas the cells with low PRL-3 expression, both endogenous and stably knocked down, proved to be less sensitive to the treatment of curcumin. Similar results were obtained for the inhibitory effect of curcumin on cell proliferation (Fig. 3D).

**Curcumin Blocks the Src/stat3 Pathway through Down-Regulating PRL-3 Expression.** Curcumin is able to inhibit the phosphorylation of stat3 (Bharti et al., 2003; Blasius et al., 2006). In addition, PRL-3 can induce Src activation, which initiates a number of signal pathways cumulating in the phosphorylation of extracellular signal-regulated kinase 1/2, stat3, and p130Cas (Liang et al., 2007). To evaluate the involvement...
ment of PRL-3 in the inhibitory effect on stat3 phosphorylation by curcumin, B16BL6 cells were treated with 10, 20, or 40 μM curcumin for 24 h followed by immunoblotting. Curcumin reduced PRL-3, phosphorylated Src (Tyr416), and phosphorylated stat3 (Tyr705) at the protein level, whereas phosphorylated Src (Tyr527), total Src, and stat3 were not affected (Fig. 4A). After B16BL6 cells were transfected with various amount of PRL-3-expressing plasmids, overexpression of PRL-3 reversed the inhibition of the phosphorylation of Src (Tyr416) and stat3 (Tyr705) by curcumin in a dose-dependent manner (Fig. 4B). As a direct transcript target of stat3, cyclin D1 but not cyclin D3 was also reversed (Fig. 4B). To exclude the possibility caused by the nonspecific effect of overexpression system, L13 and P1 cells were treated with curcumin for 24 h followed by immunoblotting. As shown in Fig. 4C, Tyr416 phosphorylation of Src kinase was reduced by approximately 73 and 28%, and Tyr705 phosphorylation of stat3 was reduced by approximately 53 and 14% in L13 and P1 cells, respectively.

Curcumin Inhibits the Tumor Growth and Spontaneous Metastasis of B16BL6 Cells in Vivo. C57BL/6J mice were injected subcutaneously with B16BL6 cells into right footpads and treated with 50 and 100 mg/kg curcumin by intraperitoneal injection every 2 days. Compared with olive oil, curcumin significantly inhibited the tumor growth in a dose-dependent manner (Fig. 5A). After 26 days of tumor cell inoculation, the tumor tissues were removed. As shown in Fig. 5B, the mRNA and protein levels of PRL-3 were significantly inhibited in the tumor by curcumin treatment. Moreover, curcumin dose-dependently reduced the metastatic potential of the tumor cells (Fig. 5C). To investigate whether curcumin treatment influences lymphatic metastasis, the popliteal lymph nodes were collected and photographed. The control group showed 100% visible metastasis. In contrast, lower ratios of metastasis were found in curcumin-treated mice with approximately 50% in 50 mg/kg group, and 17% in 100 mg/kg group, respectively (Fig. 5D). Histological analysis revealed that all lymph nodes from the control group showed 100% visible metastasis.
group contained metastases. But curcumin-treated mice had developed lymph node metastases at lower frequency (Fig. 5D).

**Discussion**

It is well accepted that PRL-3 is a metastasis-associated gene. In the initial study, we observed that curcumin inhibited cell proliferation and adhesion of mouse melanoma B16BL6, in which high invasive and metastatic activity is closely correlated with its high level of PRL-3 expression. These findings implicate that curcumin might show anticancer effects at least partially by regulating PRL-3. Indeed, curcumin decreased PRL-3 mRNA of B16B16 in a dose- and time-dependent manner, and the inhibitory effect occurred at the transcriptional level. The cells with PRL-3 stably knocked down were less susceptible to curcumin inhibition. PRL-3 expression in other cell lines was also inhibited by curcumin, suggesting that this mechanism is not unique to B16B16 highly expressing PRL-3. It is noteworthy that curcumin had no effect on the expression of PRL-1 and PRL-2, which share a high degree (>75%) of amino acid sequence identity. It is likely that curcumin down-regulates PRL-3 transcription through a pathway different from PRL-1 or PRL-2. The 5′-noncoding regions of mouse PRLs are much more divergent, and the expression pattern of the PRLs differs among tissues, which supports the possibility of differential transcription regulation (Zeng et al., 1998).

The exact mode of PRL-3 transcription regulation is unclear. Although recent study has demonstrated that PRL-3 is a p53 target gene and induces G1 cell-cycle arrest in a p53-dependent manner in primary cells (Basak et al., 2008), we found that curcumin had no effect on p53 despite the PRL-3 down-regulation. Upon exposure to curcumin, no changes were detected in either p53 protein level or p53 binding to the corresponding site in the promoter of PRL-3 (Supplemental Fig. 1A). Moreover, curcumin still decreased PRL-3 even in p53-deficient murine embryonic fibroblasts (Supplemental Fig. 1B). Curcumin has ever been reported to accelerate the p53 accumulation in some tumor cell lines, such as MCF-7 (Choudhuri et al., 2002). In our test, PRL-3 down-regulation in MCF-7 cells seemed to be less sensitive to curcumin treatment than in B16BL6 cells. That is, the PRL-3 level of MCF-7 cells showed no significant change after exposure to 20 μM curcumin for 12 h (Fig. 1A), which may be caused by the accumulation of wild type of p53. However, such accumulation of p53 did not up-regulate PRL-3, because the target transcription factor of curcumin could be more active than p53 in the transcription regulation of PRL-3. Taken together, these results suggest that there are other transcription factors, rather than p53, involving in regulating PRL-3 transcription in tumor cells, and they are targeted by curcumin.

Some tumor cells were consistently not arrested by up-regulated PRL-3 in a p53-dependent manner like primary cells (Ryan et al., 2001). On the other hand, the p53 tumor suppressor plays a critical role in protecting organisms from developing cancer (Liang et al., 2007). Degrading wild-type p53 might lead to the accumulation of DNA-damaged cells by inhibiting their p53-induced apoptosis (Ryan et al., 2001; Vousden and Lu, 2002). Therefore, it seems unreasonable to down-regulate PRL-3 expression by targeting the degradation of p53 in tumor cells. Further study is in progress to elucidate the mechanism of PRL-3 transcription regulation by using curcumin as a tool.

Considerable studies suggest that curcumin shows wide-ranging anti-inflammatory and anticancer properties and is able to affect multiple targets (Anand et al., 2008). In this study, we demonstrated that PRL-3 is not only a normal target of curcumin, but a trigger one. Elevated PRL-3 will lead to Src activation through down-regulating the synthesis of C-terminal Src kinase protein, which in turn leads to tyrosine phosphorylation of a number of proteins in human embryonic kidney 293 cells (Liang et al., 2007). In highly metastatic melanoma cells, we were surprised to find decreases in tyrosine phosphorylation in PRL-3 stably knocked down cell lines (P1 and P9), compared with L1 and L13 cells, which stably express luciferase siRNA. However, unlike that in human embryonic kidney 293 cells, we have not noticed significant change of the Tyr527 but the Tyr416 phosphorylation of Src (Supplemental Fig. S2). This result indicates that there might be other relationships between PRL-3 and Src activation that are independent of C-terminal Src kinase.

As an important substrate of Src, stat3 can be also elevated by PRL-3, and the Src/stat3 pathway has been demonstrated to be implicated in tumor metastasis, including proliferation, invasion, and motility (Darnell, 2002) As shown in Fig. 3C, the inhibition of the activity of Src and stat3 made by curcumin was approximately 45 and 39% through down-regulating PRL-3 expression, respectively. Cells with PRL-3 stably knocked down by siRNA proved to be less susceptible to the anticancer effect of curcumin. These findings suggest that PRL-3 is the much further upstream target of curcumin. This study is the first to reveal the relationship between the inhibition effect of curcumin on stat3 phosphorylation and PRL-3 and provide a possible mechanism by which curcumin inhibits the metastasis of different cancers.

In vivo study showed that curcumin dose-dependently inhibited the tumor growth and prevented B16BL6 cells in primary tumor from invading the draining lymph nodes. As expected, the PRL-3 expression in the tumor tissues was remarkably decreased by curcumin. These results are similar to those in our previous study using PRL-3 siRNA (Qian et al., 2007). Because of the high level of PRL-3 mRNA in heart, therapeutic targeting of PRL-3 might exhibit cardiotoxicity (Stephens et al., 2005). However, we detected no changes either of PRL-3 mRNA level in the cardiac muscle tissues or of myocardial function in the mice treated with curcumin. Moreover, we did not detect visible protein of PRL-3 in the heart and muscle of adult mice like the tumor cell lines, which indicates that the role of PRL-3 protein synthesis system is unique in normal tissues, and it also supports the idea that targeting the expression of endogenous PRL-3 by curcumin is safe and feasible as a novel therapy for cancer.

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