KPT-330 has antitumour activity against non-small cell lung cancer

H Sun*1,5, N Hattori2,5, W Chien2, Q Sun2, M Sudo2, G L E-Ling2, L Ding2, S L Lim2, S Shacham3, M Kauffman3, T Nakamaki4 and H P Koeffler1,2

1Department of Hematology and Oncology, Cedars-Sinai Medical Center, Los Angeles, CA, USA; 2Cancer Science Institute of Singapore, National Cancer Institute, NUS, Singapore, Singapore; 3Karyopharm Therapeutics, Boston, MA 01760, USA and 4Division of Hematology, Department of Medicine, Showa University School of Medicine, Tokyo, Japan

**Background:** We investigated the biologic and pharmacologic activities of a chromosome region maintenance 1 (CRM1) inhibitor against human non-small cell lung cancer (NSCLC) cells both in vitro and in vivo.

**Methods:** The in vitro and in vivo effects of a novel CRM1 inhibitor (KPT-330) for a large number of anticancer parameters were evaluated using a large panel of 11 NSCLC cell lines containing different key driver mutations. Mice bearing human NSCLC xenografts were treated with KPT-330, and tumour growth was assessed.

**Results:** KPT-330 inhibited proliferation and induced cell cycle arrest and apoptosis-related proteins in 11 NSCLC cells lines. Moreover, the combination of KPT-330 with cisplatin synergistically enhanced the cell kill of the NSCLC cells in vitro. Human NSCLC tumours growing in immunodeficient mice were markedly inhibited by KPT-330. Also, KPT-330 was effective even against NSCLC cells with a transforming mutation of either exon 20 of EGFR, TP53, phosphatase and tensin homologue, RAS or PIK3CA, suggesting the drug might be effective against a variety of lung cancers irrespective of their driver mutation.

**Conclusions:** Our results support clinical testing of KPT-330 as a novel therapeutic strategy for NSCLC.

Chromosome region maintenance 1/exportin-1 (CRM1, XPO1) is an important nuclear protein export receptor, which recognises the leucine-rich nuclear export signal on proteins (Fornerod et al., 1997; Fukuda et al., 1997; Ossareh-Nazari et al., 1997). Of the six known nuclear export proteins, CRM1 is a sole exporter of the nuclear export signal-containing cargo proteins such as p53, p73, CDKN1A, Rb, BRCA1, and IkB (Rodriguez et al., 1999; Huang et al., 2000; Takenaka et al., 2004; Turner et al., 2012). During initiation or progression of cancers, malignant cells appear to acquire the ability to export key nuclear proteins that influence survival and proliferation (Turner et al., 2012). Overexpression of CRM1 in cancer cells is correlated with poor survival (Noske et al., 2008; Shen et al., 2009; van der Watt et al., 2009; Kojima et al., 2013; Tai et al., 2013). High levels of CRM1 can enhance nuclear export of tumour suppressive proteins and proteins associated with drug resistance (Kau and Silver, 2003; Turner et al., 2009). Thus, inhibition of CRM1 may be an effective anticancer therapy (Turner et al., 2012; Zhang and Wang, 2012). Leptomycin B (LMB) specifically binds and blocks CRM1 and potentially inhibits growth of various cancer cell lines, but its clinical development was discontinued due to significant toxicity (Newlands et al., 1996; Mutka et al., 2009; Pathria et al., 2012). Nevertheless, blocking CRM1 nuclear export activity either by small interfering RNA or several CRM1 inhibitors (LMB and ratjadone) can restore tumour cell sensitivity to chemotherapeutic drugs such as doxorubicin (Turner et al., 2009; Lu et al., 2012), etoposide (Turner et al., 2009), cisplatin (Takenaka et al., 2004), and imatinib mesylate (Aloisi et al., 2006). Novel, orally bioavailable small molecules, ‘selective inhibitors of nuclear export’ (SINEs) have recently been developed. These compounds induce apoptosis and block proliferation of cancer cells, and show potent antileukaemic activity in mice (Lapalombella et al., 2012; Ranganathan et al., 2012;...
Azmi et al., 2013a,b; Etchin et al., 2013a,b; Kojima et al., 2013; Salas Fragrameni et al., 2013; Tai et al., 2013; Zhang et al., 2013). SINEs synergize with doxorubicin to inhibit growth of multiple myeloma cells (Turner et al., 2012).

Non-small cell lung cancer (NSCLC) represents about 85% of lung cancers, mainly consisting of adenocarcinoma, squamous cell, and large-cell carcinomas. Overexpression of epidermal growth factor receptor (EGFR) occurs in more than 60% of metastatic NSCLC, which is correlated with a poor prognosis (Sharma et al., 2007). Lung cancer patients with mutations in the tyrosine kinase (TK) domain (exons 19 or 21) of the EGFR gene respond poorly to an EGFR TK inhibitor (TKI, gefitinib or erlotinib). Indeed, acquired resistance to EGFR TKI therapy eventually occurs in nearly all patients (Ohashi et al., 2013). Resistance to EGFR TKI develops due to abnormalities in several prominent pathways including a mutation at exon 20 of EGFR, phosphatidylinositol-4, 5-bisphosphate 3-kinase, catalytic subunit alpha (PI3KCA) mutation, as well as, loss-of-function mutations of the phosphatase and tensin homologue gene leading to the constitutive activation of Akt. K-ras mutations can result in the constitutive activation of EGFR signalling pathways. Cells with these mutations will acquire resistance to EGFR TKI (Sharma et al., 2007; Stewart, 2010; Johnson et al., 2012). Therefore, because of the mark heterogeneity of NSCLC, besides manipulating components of the EGFR signalling pathways, molecular therapeutic approaches that involve simultaneously targeting several distinct pathways such as using SINEs may have therapeutic benefits (Sharma et al., 2007; Stewart, 2010; Johnson et al., 2012).

Mutations of p53 are very common in lung cancer cell lines, as well as 40–90% of fresh NSCLC tumours (Stewart, 2010). The presence of p53 mutations is a poor prognostic marker in patients with adenocarcinoma of the lung (Stewart, 2010). Previously, LMB was shown to induce cell death in cervical carcinoma cell lines; these cells likely expressed papilloma virus E6 associated with inactivation of p53 (Freedman and Levine, 1998; Lecane et al., 2003; Hoshino et al., 2008). The potential of a CRM1 inhibitor as a therapeutic agent against lung cancer including those with p53 alterations has not been fully investigated.

Here, we provide evidence that the SINE KPT-330 has antitumour activity both in vitro and in vivo against NSCLC cells irrespective of p53 mutational status.

### MATERIALS AND METHODS

**Reagents and antibodies.** KPT-330 was obtained from Karyopharm Therapeutics (Natick, MA, USA). Gefitinib (product number G-4408), Dasatinib (product number D-3307), Docetaxel (product number D-1000), Paclitaxel (product number P-9600), Ganciclovir (product number G-4408), Dasatinib (product number D-3307), Docetaxel (product number D-1000), Paclitaxel (product number P-9600), Wortmannin (product number R0395), Actinomycin D (product number A1410), and Streptomycin (Invitrogen, Grand Island, NY, USA) at 37°C and 5% CO2.

Mutational status of the NSCLC cell lines is shown in Supplementary Table S1. The cells’ identity was verified by STR analysis (ATCC). Proliferation assays were performed in 96-well plates using 3 x 104 cells per well in 100 μl for 72 h. Cell numbers were evaluated with the use of the 3-[4, 5-dimethylthiazol-2-yl]-2, 5-diphenyltetrazolium bromide (MTT) reagents (Promega, Madison, WI, USA). The absorbance of wells at 570 nm was measured with a microplate reader (Infinite 200; Tecan, San Jose, CA, USA). Dose-response curves were plotted to determine half-maximal inhibitory concentrations (IC50) for the compounds using the GraphPad Prism4 (GraphPad Software, San Diego CA, USA). To determine the synergy of two drugs, they were combined at a constant ratio (based on the individual drug’s IC50) at nine concentrations. Synergy was determined using the CalcuSyn software (BioSoft, Cambridge, UK). The extent of drug interaction between the two drugs was presented using the combination index (CI) for mutually exclusive drugs. Different CI values were obtained when solving the equation for different concentrations of drugs. A CI of 1 indicates an additive effect, whereas a CI of <1 denotes synergy. For the cell viability assays, NSCLC cells were seeded into 96-well plates and treated for 72 h with drugs.

**Apoloptosis and cell cycle assays.** Cells were seeded at 50% confluency in 6 cm plates. After overnight incubation, culture media were removed and replaced with media containing either diluent control or various concentrations of indicated drugs.

| Table 1. Quantitative RT–PCR primer sequences |
|---------------------------------------------|
| **Oligonucleotide name** | **Sequence** |
|-------------------------|--------------|
| p53-F | 5'-CTGAGGTGTGCTGACTGTACCACCTATCC-3' |
| p53-R | 5'-CTATTCACTGCTTGCCAACTCTGGAGGC-3' |
| BCL-CL-F | 5'-GCAGGCGAGAGTGGTAACT-3' |
| BCL-CL-R | 5'-GTGCTCTTGATCTCCAGTCGTA-3' |
| Bax-F | 5'-TGCAATGTATTGGCTAGCGGAC-3' |
| Bax-R | 5'-GGAGGCTTGAAGGTCTCACC-3' |
| Noxa-F | 5'-GGATGCTCCTGGAGAAGGGG-3' |
| Noxa-R | 5'-ACGTCGACCCTCCTAGAAA-3' |
| P53A1-P1-F | 5'-AGCTTACCTGCCAGAAGCTCCTGCTC-3' |
| P53A1-P1-R | 5'-GCAATCCGAGGAGTCCCTGCT-3' |
| PUMA-F | 5'-AGCCACTCAGCGCAGGCTCAG-3' |
| PUMA-R | 5'-AGGAGCTCCGACATCCGAGTCT-3' |
| p73-F | 5'-TTCACCGAGAGATGACTG-3' |
| p73-R | 5'-CAGGATTCATCCACATCTG-3' |
| CRM1-F | 5'-CTCTTGCCAGCAGCATCGGA-3' |

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After 24 h, cells were washed with phosphate-buffered saline, trypsinized, washed once with phosphate-buffered saline and resuspended in annexin-binding buffer (BD Biosciences, San Jose, CA, USA) at 1 × 10^6 cells per ml. Cells were stained with propidium iodide (BD Biosciences) and annexin V (BD Biosciences) according to the manufacturer’s protocol and assayed on a LSRII flow cytometer (BD Biosciences). Cell cycle analyses were performed by propidium iodide staining (Sigma-Aldrich) for DNA content and flow cytometric analysis.

**Proteins, lysates, fractionation, immunoblotting, immunofluorescence, and immunohistochemistry.** Protein lysates from cells were extracted using ProteoJET Mammalian Cell Lysis Reagent (Thermo Scientific, Waltham, MA, USA). Protein concentrations were determined by bicinchonic acid assay (Thermo Scientific). Protein lysates were resolved by sodium dodecyl sulphate–polyacrylamide gel electrophoresis, transferred to Immobilon-P PVDF membrane (Merck Millipore, Billerica, MA, USA), and membranes were blocked for 1 h with TBS containing 5% non-fat dry milk and 0.5% Tween 20 (TBST). After blocking, the PVDF membrane was incubated with primary antibody at 4°C overnight.

The following day, the membrane was washed with T-TBS and incubated for one hour with a secondary antibody-HRP conjugate at room temperature. After washing the PVDF membranes with T-TBS, proteins were visualised with SuperSignal chemiluminescent hpr substrates (Thermo Scientific). To prepare nuclear and cytoplasmic extracts, cells were lysed in 10 mM HEPES (pH 7.9), 10 mM KCL, 0.1 mM EDTA, 0.1 mM EDTA, 0.6% NP-40, 1 mM DTT, and protease inhibitors (Thermo Scientific) and centrifuged at 16 000 g for 10 min to collect soluble fractions (cytosolic extracts). Insoluble material was washed with the lysis buffer (20 mM HEPES (pH 7.9), 0.4 M NaCl, 1 mM EDTA, 1 mM EDTA, 0.6% NP-40, 1 mM DTT, and protease inhibitors) and centrifuged at 16 000 g for 10 min to collect the nuclear extracts. The nuclear and cytoplasmic fractions were subjected to immunoblot analysis using either anti-hnRNP A1 antibody or anti-α-tubulin monoclonal antibody.

For immunohistochemistry, tumour tissues were fixed in formalin, embedded in paraffin, cut into 5.0 μm sections. Endogenous peroxidase on these slide preparations were quenched using 3% H2O2 in distilled water. Sections were blocked for 1 h at room temperature in phosphate-buffered saline containing 3%
normal serum and incubated overnight at 4°C with primary antibody for either TUNEL assay or against Ki-67. Biotin-conjugated secondary antibody (1:200) and avidin-biotin peroxidase complex were then applied to the sections. Data from three randomly selected fields of view per section were captured using a microscope (Olympus, Tokyo, Japan) at ×200 magnifications. TUNEL- and Ki-67-positive cells were counted using ImageJ software (NIH, Bethesda, MD, USA) and expressed as a percentage of total cells counted.

Quantitative RT–PCR. Total RNA from cells was extracted using AxyPrep Multisource Total RNA Miniprep Kit (Axygen, Tewksbury, MA, USA) according to the manufacturer’s protocol. First-strand cDNA was synthesised from total RNA using Maxima First Strand cDNA Synthesis Kit for RT–PCR (Thermo Scientific). Gene expression levels were measured by 7500 Fast Real-Time PCR System (Applied Biosystems, Beverly, MA, USA) using the KAPA SYBR FAST qPCR Kits (Kapa Biosystems, Woburn, MA, USA). Expression of each gene was normalised to β-actin as a reference. The forward and reverse primer sequences have been previously described (Leong et al., 2007). The conditions for all quantitative RT–PCR reactions are as follows: 2 min at 95°C followed by 15 s at 95°C and 30 s at 60°C for 40 cycles. All PCR products were confirmed by the presence of a single peak upon melting curve analysis and by gel electrophoresis. Primers used for all quantitative RT–PCR were listed on Table 1.

shRNA and lentiviral infections. The shRNA constructs in the pLKO.1 lentiviral vector with the following target sequences were used. shCRM1, 5′-GCTCAAGAAGTACTGACACAT-3′ shp53-1, 5′-CATCATCCACTACAATCACAT-3′ shp53-2, 5′-CGGCAGACAGGAAAGAAT-3′ Viral particles were made by transfection of 293 T cells. Target cells were incubated with lentiviruses in the presence of 8 mg·mL⁻¹ polybrene for 16 h followed by replacement of the lentivirus-containing media with fresh media. Two days after infection, Puromycin (2 mg·mL⁻¹) was added for 3 days to eliminate uninfected cells.

Animal models. NOD/SCID mice were purchased from Cancer Science Institute of Singapore (Dr Shin Leng). All animal studies were conducted according to protocols approved by the National University of Singapore Institutional Animal Care and Use Committee (IACUC). Seven-week-old male NOD/SCID mice were inoculated subcutaneously in both flanks with a suspension of H1975 (1.0 × 10⁶ cells) with matrigel. Tumour volumes were measured thrice weekly with calipers and were calculated using the following formula: volume (mm³) = (width (mm))² × length (mm)/2. Mice were randomly divided into two groups (five mice per group): orally treated with vehicle (0.6% w/v aqueous Pluronic F-68 (Karyopharm Therapeutics)) or KPT-330 (10 mg·kg⁻¹, thrice weekly for 4 weeks). Mice were given Nutri-Cal (tomlyn) during therapy to allow robust nutrition.

Statistics. Differences between conditions were analysed using either paired or unpaired two-tailed Student’s t-test. One-way analysis of variance was used for comparisons among multiple groups. Differences were considered significant for P <0.05.

RESULTS

KPT-330 suppresses the growth of NSCLC cells. The antiproliferative activity of KPT-330 against 11 NSCLC cell lines was examined using MTT assay. KPT-330 (0.1–1000 nM) induced a

Figure 2. KPT-330-treated NSCLC cells: measurement of cell cycle and apoptosis. (A, B) Cell cycle of NSCLC cells (A549, H460, H1975, PC14, H1299, and H23) was analysed by flow cytometry after treatment with KPT-330 (1 μM, 24 h) vs diluent control. Representative tracings of cell cycle of A549 and PC14 are displayed in panel A. (C, D) Cells were analysed by flow cytometry for apoptosis (annexin V/propidium iodide positivity) after exposure to either KPT-330 (1, 10, 100, and 1000 nM) or diluent control for 24 h. Representative tracing of apoptosis analysis of A549 and PC14 is shown in panel C.
KPT-330 promotes G1 arrest and apoptosis. Cell cycle distribution was determined by propidium iodide staining of six NSCLC cell lines (A549, H460, H1975, PC14, H1299, and H23) exposed to KPT-330 (1 μM, 24 h) (Figure 2A and B). All six NSCLC cell lines had an increase of the G1 cell cycle (range: 56–82%), and a decrease in the S (range: 8–20%) and G2/M (range: 2–20%) phases in response to KPT-330 (Figure 2B). Apoptosis was examined using annexin V combined with propidium iodide staining (Figure 2C and D). A clear dose-dependent increase in the percent of annexin V/propidium iodide co-positive cells occurred in the six NSCLC cell lines after cultured in KPT-330 (1, 10, 100, and 1000 nM for 24 h) (Figure 2D). The range of apoptosis induced by KPT-330 was 22–54%.

Effect of KPT-330 on wild type (wt) and mutant (mut) p53 NSCLC cells. p53 wild type (p53-wt, A549) and mutant (p53-mut, PC14) NSCLC cells treated with KPT-330 (1 μM, 24 h) resulted in dramatic change in key cell cycle and cell death pathways in both cell lines (Figure 3A and B). KPT-330 stimulated activation of caspase-3 and -9, causing PARP cleavage associated with increased levels of pro-apoptotic mediators (Bax, Bim, and Puma) and slightly decreased levels of anti-apoptotic proteins (Bcl-2 and Bcl-xL). Cell cycle inhibitors CDKN1A and CDKN1B were upregulated, while CCND1 and c-Myc were downregulated in the presence of KPT-330 (Figure 3A). These changes in protein expression ± KPT-330 were similar when comparing p53-wt (A549) and mut-p53 (PC14) NSCLC cells. p53 and its family member (e.g. p73) are active either in cell cycle arrest and DNA

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**Figure 3.** Cellular events after exposure of NSCLC cells to KPT-330. A549 cells (p53-wt) and PC14 cells (p53-mut) were exposed to KPT-330 (1 μM, 24 h), harvested followed by either (A) whole-cell protein extractions or (B) nuclear (confirmed, hnRNP A1) and cytoplasmic (confirmed, α-tubulin) protein extractions, or (C) total RNA. Protein was immunoblotted and probed with antibodies specific for various proteins. RNA was analysed for apoptosis-related genes (quantitative RT–PCR). (D) A549 cells were stably infected with either one of two shp53 vectors (shp53-1, shp53-2) or scrambled vector. p53-knockdown efficiency was evaluated by quantitative RT–PCR. (E) Growth curves of A549 NSCLC cells stably containing either shp53 or scrambled vectors following their exposure (72h) to different concentrations of KPT-330 (0, 10, 100, and 1000 nm). Cell viability (MTT assay) was determined after 72-h treatment. (F) Decrease expression of p53 mediated by pifithrin-α. A549 cells were exposed to either pifithrin-α (5 μM, 48 and 72 h) or diluent control, harvested and whole-cell extractions prepared for immunoblot. (G) Dose-response assay of A549 cultured in either diluent (WT) or pifithrin-α (5 μM) plus various concentrations of KPT-330 (0, 100, 200, 400, 800, and 1600 nm). Cell viability was determined using annexin V combined with propidium iodide staining (Figure 2C and D). A clear dose-dependent increase in the percent of annexin V/propidium iodide co-positive cells occurred in the six NSCLC cell lines after cultured in KPT-330 (1, 10, 100, and 1000 nM for 24 h) (Figure 2D). The range of apoptosis induced by KPT-330 was 22–54%.
repair, or apoptosis if the DNA damage of the cell is too severe to be repaired. The p53 family member, p73, can help to regulate p53-dependent genes in p53-deficient cells (Vilgelm et al., 2008) and has become a potential therapeutic target for treating p53-mutant cancers (Irwin et al., 2003). KPT-330 increased levels of wild-type p53 in A549 and levels of p73 in PC14 (mutant p53) (Figure 3A and B). Bax, p53AIP, Noxa, and Puma are pro-apoptotic mediators of cell death and are known targets of both p53 and p73. KPT-330 (1 μM, 24 h) induced their expression in both cell lines, albeit more profoundly in the p53-wt cells (A549) (Figure 3C). To identify whether p53 is important for KPT-330-dependent antiproliferative effects in p53-wt NSCLC cells, we stably expressed two lentivirus vector-based shp53 (shp53-1 and shp53-2) in A549 cells. Compared with scramble vector control, shp53-1 and shp53-2 stably infected cells achieved 31% and 49% knockdown efficiency as measured by western blotting, respectively (Figure 3D). p53-knockdown A549 cells (shp53-1 and shp53-2) developed significant cell resistance to several concentrations of KPT-330 (100 and 1000 nM, Figure 3E). Also, pifithrin-α is a potent agonist of p53, which can decrease both the nuclear stability and the basal DNA-binding activity of p53 in many cells (Komarov et al., 1999; Murphy et al., 2004; Sohn et al., 2009). A549 cells exposed to pifithrin-α (5 μM) had decreased levels of p53 (48 h and 72 h) (Figure 3F), p21<sup>swp1</sup>, and Mdm2 (Supplementary Figure S1). Similar to p53-knockdown cells, A549 cells exposed to pifithrin-α (5 μM) were more resistant to the antiproliferative effect of KPT-330 (Figure 3F). The apoptosis induced by KPT-330 (cleavage of caspase-3 and -9, as well as polyADP ribose polymerase (PARP) (Figure 3A)) was inhibited by caspase inhibitor Z-VAD-FMK (10 μM) in the A549 cells (Figure 3H). As a positive control, Z-VAD-FMK decreased apoptosis induced by Actinomycin D (Figure 3H).

Figure 4. Drug combinations with NSCLC cells. (A) Upper and middle panel: H1975 cells were cultured with either KPT-330 (30 nM), diluent (control), experimental drug, or experimental drug + KPT-330 (+ KPT) for 72 h, and cell viability was determined by MTT. Lower panel: combination of KPT-330 with cisplatin cultured with each of six NSCLC cell lines. In each group of four: 1st rectangle is control, 2nd, KPT-330, 3rd, cisplatin, and 4th, combination of KPT-330 and cisplatin. Concentrations of KPT-330 and cisplatin were based on the IC50 of each drug for each NSCLC cell lines. (B) Combinatorial effect of KPT-330 and cisplatin on H1975 displayed as the CI. CI defines the interaction between KPT-330 and cisplatin as plotted against the fraction of cell viability. CI values: 1, additive; <1, synergistic; >1, antagonistic. Table to the right defines each tested concentration of KPT-330 and cisplatin. (C) Cells analysed by flow cytometry for apoptosis (annexin V/propidium iodide positivity) after exposure to either KPT-330 (500 nM); cisplatin (1 μg ml<sup>−1</sup>); KPT-330 + cisplatin, or DMSO diluent (control) for 24 h. Numbers in the figure represent percentage of apoptotic cells (annexin V alone + PI alone + double positive).
KPT-330 and cisplatin synergy. H1975 NSCLC cells were treated with KPT-330 (30 nM) and 11 additional therapeutic drugs (wortmannin (10 nM), cisplatin (1 μg/ml), doxorubicin (1 μM), dasatinib (50 nM), gefitinib (1 μM), panobinostat (10 nM), bortezomib (10 nM), palitaxel (2 nM), gemcitabine (50 nM), 17-DAG (10 nM), and rapamycin (1 μM)) (Figure 4A, upper and middle panels). The concentrations of the drugs were based on dose-response proliferation experiments against H1975 cells (data not shown). The combination of cisplatin and KPT-330 had greater antiproliferative activity than either drug alone against six NSCLC cell lines (Figure 4A, lower panel). Cisplatin is often given to patients with NSCLC either as adjuvant therapy or those with progressive disease. Therefore, the combinatorial effect of KPT-330 with cisplatin was evaluated using six different paired combinations of KPT-330 (40 and 80 nM) and cisplatin (0.5, 1, and 2 μg/ml) against H1975 NSCLC cell lines. The data were placed on a CI plot (Figure 4B). Synergistic effects were observed with KPT-330 at either 40 or 80 nM and cisplatin at either 0.5, 1, or 2 μg/ml (CI values of 0.816, 0.648, 0.514, 0.441, and 0.418 (No. 2, 3, 4, 5, and 6, respectively). In addition, the combination of KPT-330 and cisplatin increased apoptosis of H1975 > H460 > A549, which was greater than either drug alone (Figure 4C).

Silencing CRM1 sensitises NSCLC cells to cisplatin. As proof of principal, CRM1 was silenced in order to determine whether the observed synergistic activity of KPT-330 and cisplatin could be produced in another model system. H1975 cells were stably infected with the pLKO.1 lentiviral vector expressing shCRM1. Efficient shRNA was confirmed by immunoblot (60% knockdown) (Figure 5A) and quantitative RT–PCR (66% knockdown) (Figure 5B). CRM1-knockdown cells had significantly slower growth than those stably carrying a scrambled vector (Figure 5C). The CRM1 knockdown of NSCLC cells and the scramble vector carrying control cells were treated with different concentrations of cisplatin (0.01–10 μg/ml) for 72 h, and cell numbers were assayed by MTT. Compared with control cells, CRM1-knockdown cells cultured with cisplatin had enhanced antiproliferative activity compared with scrambled carrying cells also exposed to cisplatin (IC50, shCRM1, 1.62 μg/ml vs scramble, 8.1 μg/ml (Figure 5D)). Moreover, upon exposure to cisplatin (1 μg/ml, 24 h), the CRM1-knockdown cells had elevated levels of Bim and Puma compared with control cells (scramble control + cisplatin) (Figure 5E). Likewise, mRNA expression of Puma was significantly enhanced in the CRM1-knockdown cells cultured with cisplatin compared with a similar treatment of the control cells carrying the scramble vector (Figure 5F).

shp73 reduces sensitivity to KPT-330 in NSCLC cells. The p73 was stably silenced (shp73) to explore the importance of p73 in mediating the antiproliferative effect of KPT-330 on NSCLC cells.
having a mutant p53. H1975 cells were stably infected with the pLKO.1 lentiviral vector expressing shp73. Efficient shRNA was confirmed by immunoblot (p73, 60% knockdown) (Figure 6A) and by quantitative RT–PCR (64% knockdown) (Figure 6B). Growth of the p73-knockdown NSCLC cells was slightly faster than the vector control cells (stable scramble vector) (Figure 6C). Also, the p73-knockdown cells were more resistant to the growth inhibitory effects of KPT-330 (IC50, scramble, 98 nm vs shp73, >1000 nm) (Figure 6D). Transiently silence of p73 (44% knockdown, Supplementary Figure S2A) in PC14 cells were also more resistant the treatment of KPT-330 compared with the vector control cells (IC50, scramble, 197 nm vs shp73, 318 nm) (Supplementary Figure S2B). In addition, p73-knockdown cells exposed to KPT-330 had decreased apoptosis (Figure 6E), decreased levels of cleaved PARP and caspase-3, as well as lower levels of BimEL (Figure 6F) compared with the scramble vector + KPT-330. Likewise, mRNA expression of Noxa and Puma was lower in the p73-knockdown cells cultured with KPT-330 compared with cells cultured with the scramble vector + KPT-330 (Figure 6G).

Figure 6. Stable silencing of p73 using shRNA in H1975 cells plus addition of KPT-330. H1975 cells were stably infected with either a p73-specific shRNA (shp73) or scrambled shRNA (scramble, control). p73-knockdown efficiency was evaluated by immunoblot (A) (densitometry shows 64% silencing of p73 in cells having shRNA p73) and quantitative RT–PCR (B). (C) Cell proliferation was measured by MTT assay after 1–4 days of culture. (D) Growth curves of H1975 cells stably carrying shp73 cells or scramble vector cultured with KPT-330 (0, 1, 3, 10, 30, 100, 320, and 1000 nM, 3 days). (E) H1975 cells with either stable p73 knockdown or scrambled vector were exposed to KPT-330 (50 nM, 24 h) or DMSO and analysed by flow cytometry for apoptosis (annexin V/propidium iodide positivity). Numbers in the figure represent percentage of apoptotic cells (annexin V alone + PI alone + double positive). (F) Immunoblots of whole-cell extracts of H1975 cells either with or without stably containing shp73 ± KPT-330 (1000 nm, 24 h). (G) mRNA levels measured by quantitative RT–PCR of cells as described in panel F. Results for panels B, C, D, and G represent the mean ± s.d. of three experiments done in triplicate. *P<0.05.
Antitumour activity of KPT-330 against human NSCLC xenografts growing in vivo. H1975 NSCLC xenografts were established in NOD/SCID mice (outlined in the Materials and Methods). These cells have the T790M EGFR mutation rendering them resistant to inhibition by the TKIs gefitinib and erlotinib. Single-agent KPT-330 resulted in a marked inhibition of tumour growth when compared with vehicle-treated controls (Figure 7A and B). Immunohistochemistry analysis showed decreased Ki-67-positive cells (measure of cell growth) and an increased percent of TUNEL-positive cells (increased apoptosis) produced by KPT-330 (Figure 7C). These findings established the efficacy of KPT-330 against NSCLC cells in vivo. Importantly, this therapy did not have a significant effect on body weight compared with the vehicle group (mean relative body weight for vehicle versus KPT-330 cohorts: 106% and 105% body weight compared with their mean starting weight, respectively) (Figure 7D). In addition, we analysed peripheral blood after the 28-day treatment (Table 2). KPT-330-treated mice had no significant alterations of their total white blood cells as well as neutrophils and platelet counts, haematocrit, as well as their serum levels of albumin, alkaline phosphatase, ALT, and creatinine compared with vehicle group. These results suggest that the KPT-330 has promising therapeutic applications for the treatment of NSCLC patients associated with minimal toxicity.

## DISCUSSION

Previous reports demonstrated that the natural product LMB, a CRM1 inhibitor, had anticancer activity. But because of its significant toxicity, trials with this drug were discontinued (Newlands et al, 1996). Also, significant animal toxicity was noted with LMB (Mutka et al, 2009). However, the marked gastrointestinal toxicity of LMB was not based on its ability to inhibit CRM1 as LMB derivatives with improved pharmacologic properties, but retaining CRM1 inhibitory capacity displayed much less toxicity (Mutka et al, 2009; Lapalombella et al, 2012). Both classes of CRM1 inhibitors (LMB and SINE compounds) bind to CYS 539 in the cargo-binding groove of SC-CRM1. However, LMB fills most of the groove, while SINE compounds occupy only around 40% of the groove. Sun et al (2013) also demonstrated that LMB, but not
SINE, undergoes hydrolysis following the binding to CRM1, resulting in the decreased affinity of LMB to CRM1. In addition, the authors showed that LMB binding to CRM1 is not reversible even following 24 h of incubation, while SINE binding is reversible at 24 h. Therefore, the improved toxicity profile of SINE compounds compared with LMB may result from their reversibility of binding to CRM1, allowing ‘normal’ cells to recover.

p53 is mutated in more than half of all human cancers, leading to a variety of biological effects and often associated with a poor clinical outcome (Freedman and Levine, 1998; Lecane et al, 2003; Hoshino et al, 2008). Nuclear export of this protein is mediated exclusively through CRM1 (Stommel et al, 1999). Previous studies showed that LMB induced p53 activation, suggesting this was the key to inducing growth arrest associated with apoptosis in prostate cancers (Lecane et al, 2003), neuroblastomas (Smart et al, 1999), and melanoma (Pathria et al, 2012). Recently, a study showed that p53 was a critical mediator of KPT-induced differentiation and apoptosis of acute myeloid leukaemia cells (Ranganathan et al, 2012). Interestingly, we observed that KPT-330 induced apoptosis in both p53-wt and -mut NSCLC cells. We observed that KPT-330 increased the levels of p73. As a transcription factor, p73 shares structural and functional similarities with p53. In cancer cells that express p53-wt, p73 cooperates with p53 to induce apoptosis; whereas in p53-mutant cancer cells, p73 has been reported to cause apoptosis via activation of p53-inducible genes (Irwin, 2004; Moll and Slade, 2004; Vayssade et al, 2005; Chung and Irwin, 2010). p73, similar to p53, is upregulated in response to a subset of DNA-damaging agents including cisplatin, inducing expression of apoptosis regulated genes including Bax, Bin, Noxa, and Puma (Moll & Slade, 2004; Vilgelm et al, 2008; Lunghi et al, 2009; Zawacka-Pankau et al, 2010). In addition, tumours often have a p53 missense mutation at R175H, R248H, R249S, or R273H in human cancers. These p53-mutant proteins retain the ability to bind to p73 and transactivate endogenous CDKN1A, suppressing cell proliferation (Willis et al, 2003). We showed that KPT-330 induced apoptosis in H1975 cells (R273H, mutation of p53) associated with expression of CDKN1A. Silencing of p73 (shRNA p73) blunted the ability of KPT-330 to slow the growth of these NSCLC cells. p73 has been shown to be an important determinant of chemosensitivity in humans cancers (Irwin et al, 2003) and has an important role in cisplatin-induced apoptosis of tumours (Al-Bahlani et al, 2011). Taken together, p73 may have a significant role in the antiproliferative activity of KPT-330 wild-type and p53-mutant NSCLC cells.

In our study, two of the NSCLC cell lines have a T790M mutation in exon 20 of EGFR, making them resistant to both TKIs (gefitinib and erlotinib) (Sharma et al, 2010). But these cells were sensitive to the antiproliferative effects of KPT-330 showing that KPT-330 can inhibit growth of NSCLC cells even after their loss of responsiveness to a TKI. Furthermore, 5 of 11 of the NSCLC cell lines had either a K- or N-RAS mutation, 2 of 11 NSCLC had PIK3CA mutations. KPT-330 had antiproliferative activity against each of these NSCLC cell lines. Thus, KPT-330 might have unique abilities to inhibit growth of tumour cells containing mutant tumour suppressor genes or activated oncogenes.

In conclusion, the present study shows for the first time that KPT-330 has prominent in vitro and in vivo antitumour activity against NSCLC cells. KPT-330 inhibited cell proliferation and induced apoptosis of NSCLC cells in vitro and in vivo. The drug was effective regardless of the mutational status of either p53, EGFR, RAS, PI3K, or phosphatase and tensin homologue. The drug also reduced levels of CRM1 protein, and produced a significant nuclear accumulation of p53 and p73. Combination therapy with KPT-330 and cisplatin displayed synergistic antiproliferative activity in vitro. Treatment of mice bearing NSCLC tumours with oral KPT-330 significantly reduced the size of their tumours. These data suggest that KPT-330 either alone or in combination with platinum-based regimens may be effective in the treatment of NSCLC patients despite the heterogeneity of this disease.

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**CONFLICT OF INTEREST**

MK and SS are employees of Karyopharm Therapeutics, a biopharmaceutical company that develops selective inhibitors of nuclear export-targeted therapeutics.

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