

ORIGINAL ARTICLE

Preclinical development of a humanized neutralizing antibody targeting HGF

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Hepatocyte growth factor (HGF) and its receptor, cMET, play critical roles in cell proliferation, angiogenesis and invasion in a wide variety of cancers. We therefore examined the anti-tumor activity of the humanized monoclonal anti-HGF antibody, YYB-101, in nude mice bearing human glioblastoma xenografts as a single agent or in combination with temozolomide. HGF neutralization, The extracellular signal-related kinases 1 and 2 (ERK1/2) phosphorylation, and HGF-induced scattering were assessed in HGF-expressing cell lines treated with YYB-101. To support clinical development, we also evaluated the preclinical pharmacokinetics and toxicokinetics in cynomolgus monkeys, and human and cynomolgus monkey tissue was stained with YYB-101 to test tissue cross-reactivity. We found that YYB-101 inhibited cMET activation *in vitro* and suppressed tumor growth in the orthotopic mouse model of human glioblastoma. Combination treatment with YYB-101 and temozolomide decreased tumor growth and increased overall survival compared with the effects of either agent alone. Five cancer-related genes (TMEM119, FST, RSPO3, ROS1 and NBL1) were overexpressed in YYB-101-treated mice that showed tumor regrowth. In the tissue cross-reactivity assay, critical cross-reactivity was not observed. The terminal elimination half-life was 21.7 days. Taken together, the *in vitro* and *in vivo* data demonstrated the anti-tumor efficacy of YYB-101, which appeared to be mediated by blocking the HGF/cMET interaction. The preclinical pharmacokinetics, toxicokinetics and tissue cross-reactivity data support the clinical development of YYB-101 for advanced cancer.

Experimental & Molecular Medicine (2017) 49, e309; doi:10.1038/emm.2017.21; published online 24 March 2017

INTRODUCTION

Hepatocyte growth factor (HGF), also known as scatter factor, is a multifunctional cytokine composed of an amino-terminal domain and four kringle domains in the alpha chain (54–65 kDa) and a serine protease homology domain in the beta chain (31–35 kDa).¹ The binding of HGF to its receptor, cMET, activates intracellular signal transduction pathways that regulate cell proliferation, motility, invasion, angiogenesis, and anti-apoptosis.^{2,3} Thus, aberrant activation of the HGF/cMET pathway triggers growth and metastasis in a variety of human cancers.^{4,5} Upregulation of HGF and the overexpression and

activation of cMET are observed in a number of human cancers such as breast, head and neck, lung, prostate, renal, colorectal, and hepatocellular as well as myeloma, glioblastoma and sarcomas.^{6–8} Furthermore, a high blood level of HGF is associated with poor prognosis in gastric and ovarian cancer.^{6,9} For example, HGF and cMET expression levels correlate with tumor invasiveness, metastasis, and overall survival in breast cancer^{10–12} and with poor survival rates in non-small-cell lung cancer.^{13,14} Given that the interaction of HGF and cMET is involved in tumorigenesis and metastasis, both proteins are promising targets for therapeutic agents. HGF inhibitors bind

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Received 8 December 2016; accepted 23 December 2016

to HGF to prevent its interaction with cMET and the subsequent activation of the HGF/cMET pathway. We previously generated a rabbit-human antibody that effectively neutralizes the *in vitro* activity of HGF.¹⁵ In a colorectal cancer cell xenograft model, this antibody effectively suppressed innate irinotecan resistance induced by fibroblast-derived HGF.¹⁶

In this study, we tested a humanized version of the anti-HGF antibody (YYB101) using *in vitro* HGF neutralization assays and an orthotopic mouse model of human glioblastoma. We then carried out tissue cross-reactivity, pharmacokinetic, and toxicity studies of the monoclonal antibody (mAb) compliant with good laboratory practice. In this preclinical study, we evaluated the dose response of YYB-101 and compared the efficacy of YYB-101 alone versus combination therapy with YYB-101 and temozolomide (TMZ), the standard-of-care chemotherapy drug.

MATERIALS AND METHODS

ERK phosphorylation assay

After a 24-h incubation in serum-free medium, human liver carcinoma HepG2 cells were treated for 5 min with 128 pM HGF (ProSpec, Rehovot, Israel) along with either YYB-101 (0.1, 0.3, 1, 3, or 10 nM) or human IgG (10 nM; Sigma-Aldrich, St Louis, MO, USA). The cells were washed with ice-cold phosphate-buffered saline (PBS) twice to stop the reaction, lysed in a lysis buffer consisting of 20 mM Tris (pH 7.4) 137 mM NaCl, 1% Triton X-100, protease inhibitor cocktail, and phosphatase inhibitor cocktail, and then clarified by centrifugation at 15 000 × *g* for 5 min at 4 °C. The cell lysate proteins were separated by sodium dodecyl sulfate polyacrylamide gel electrophoresis (NuPAGE 4–12% Bis-Tris; Invitrogen, Carlsbad, CA, USA) under reducing conditions and transferred to a nitrocellulose membrane, as previously described.¹⁷ After blocking with 5% skim milk in PBS containing 0.02% Tween-20 (blocking buffer) for 30 min, the membranes were incubated overnight at 4 °C with either an anti-ERK1/2 rabbit antibody (Santa Cruz Biotechnology, Inc., Santa Cruz, CA, USA) or an anti-phospho-ERK1/2 rabbit antibody (Cell Signaling Technology, Inc., Beverly, MA, USA) diluted in blocking buffer (1:1000). The membranes were then incubated with horseradish peroxidase-conjugated mouse anti-rabbit antibody (The Jackson Laboratory, Bar Harbor, ME, USA) diluted in blocking buffer (1:1000) for 1 h at room temperature. Protein bands were visualized using an enhanced chemiluminescence system (Thermo Fisher Scientific, Waltham, MA, USA) following the manufacturer's instructions.

Scattering assay

MDCK-2 cells were incubated for 20 h in Dulbecco's modified Eagle's medium containing 67 pM HGF alone or in combination with either control IgG or YYB-101, as previously reported.¹⁵ The cells were fixed with 100 µl paraformaldehyde (4%) for 30 min at room temperature and washed with PBS. Photographs were taken of individual colonies.

Orthotopic mouse model of human glioblastoma

Human glioblastoma U-87 MG cells were washed, resuspended in PBS, and then injected stereotactically into the left striatum of 6-week-old female Balb/c nude mice (1 × 10⁵ per mouse). The injection coordinates were 2.2 mm to the left of the midline and 0.2 mm posterior to the bregma at a depth of 3.5 mm. After 1 week, YYB-101 or control IgG was administered by intraperitoneal injection (5, 10, or 30 mg kg⁻¹; *n* = 10 per group) three times per week. TMZ (5 mg kg⁻¹) or 2.5% cremophor (TMZ control) was administered by oral gavage twice a

week. After 3 weeks, the mice were killed, and tumor tissue was collected and immediately frozen in liquid nitrogen. All animal experiments were conducted in accordance with protocols approved by the Institutional Animal Care and Use Committee at the National Cancer Center, Republic of Korea.

Magnetic resonance imaging

Magnetic resonance imaging (MRI) studies were carried out using a 7.0 T magnet (BioSpin, Bruker, Germany). After localizer imaging on three orthogonal axes, T2-weighted images of the entire mouse brain were acquired using a rapid acquisition with refocused echoes sequence, with TR and TE set to 2500 and 35 ms, respectively. Other parameters used were a 2-cm field of view and a 256 × 256 matrix in four averages, resulting in a total scan time of 4 min.

RNA preparation

Snap-frozen brain tumor tissue of mice bearing human glioblastoma xenografts was pulverized in liquid nitrogen using a frozen mortar and pestle. Total RNA was extracted using TRIzol reagent (Invitrogen) and purified using RNeasy columns (Qiagen, Valencia, CA, USA) according to the manufacturers' protocol.

Microarray

To generate biotinylated cRNA, total RNA was amplified and purified using the Ambion Illumina RNA amplification kit (Ambion, Austin, TX, USA) according to the manufacturer's instructions. Briefly, total RNA was reverse transcribed using a T7 oligo(dT) primer. Second-strand cDNA was synthesized, *in vitro* transcribed, and labeled with biotin-NTP. After purification, the cRNA was quantified using a spectrophotometer (NanoDrop ND-1000, Wilmington, DE, USA). Labeled cRNA samples were hybridized to each Human-HT-12 v4 Expression BeadChip for 16–18 h at 58 °C, according to the manufacturer's instructions (Illumina, Inc., San Diego, CA, USA). Signal detection was carried out using Amersham Fluorolink Streptavidin-Cy3 (GE Healthcare Bio-Sciences, Little Chalfont, UK), and the arrays were scanned with an Illumina Bead Array Reader confocal scanner according to the manufacturer's instructions. Array data export processing and analysis was performed using Illumina BeadStudio v3.1.3 (Gene Expression Module v3.3.8).

Real-time PCR

Total RNA (1 µg) extracted from brain tumor tissue was used to synthesize cDNA using M-MLV reverse transcriptase (Invitrogen). Real-time PCR assays were performed using a LightCycler 480 SYBR Green I Master reaction mix (Roche Molecular Systems, Pleasanton, CA, USA) and the following gene-specific primers: transmembrane protein 119 (TMEM119) sense 5'-GCCTCCTCATCCTTCTGTG-3' and antisense 5'-TATCCCATCCAGGAAGTTGG-3'; follistatin (FST), sense 5'-GTTTTCTGTCCAGGCAGCTCCAC-3' and antisense 5'-GCAAGATCCGGAGTGCTTTACT-3'; R-spondin 3 (RSPO3), sense 5'-AATACATCGGCAGCCAAAACGCC-3' and antisense 5'-TG TCAAGGCACTTCCAAGGTG-3'; proto-oncogene tyrosine-protein kinase ROS (ROS1), sense 5'-GGTGACAGTGCTTATAACG-3' and antisense 5'-AAGGTTGGAATGAGCTGGATA-3'; neuroblastoma 1 (NBL1), sense 5'-CCAAGTCCATCCAGAACAGG-3' and antisense 5'-CTCAGCCCCCTCTTCTCT-3'; and glyceraldehyde 3-phosphate dehydrogenase (GAPDH), sense 5'-ACCACAGTCCATGCCATCAC-3' and antisense 5'-TCCACCACCCTGTTGCTGTA-3'.

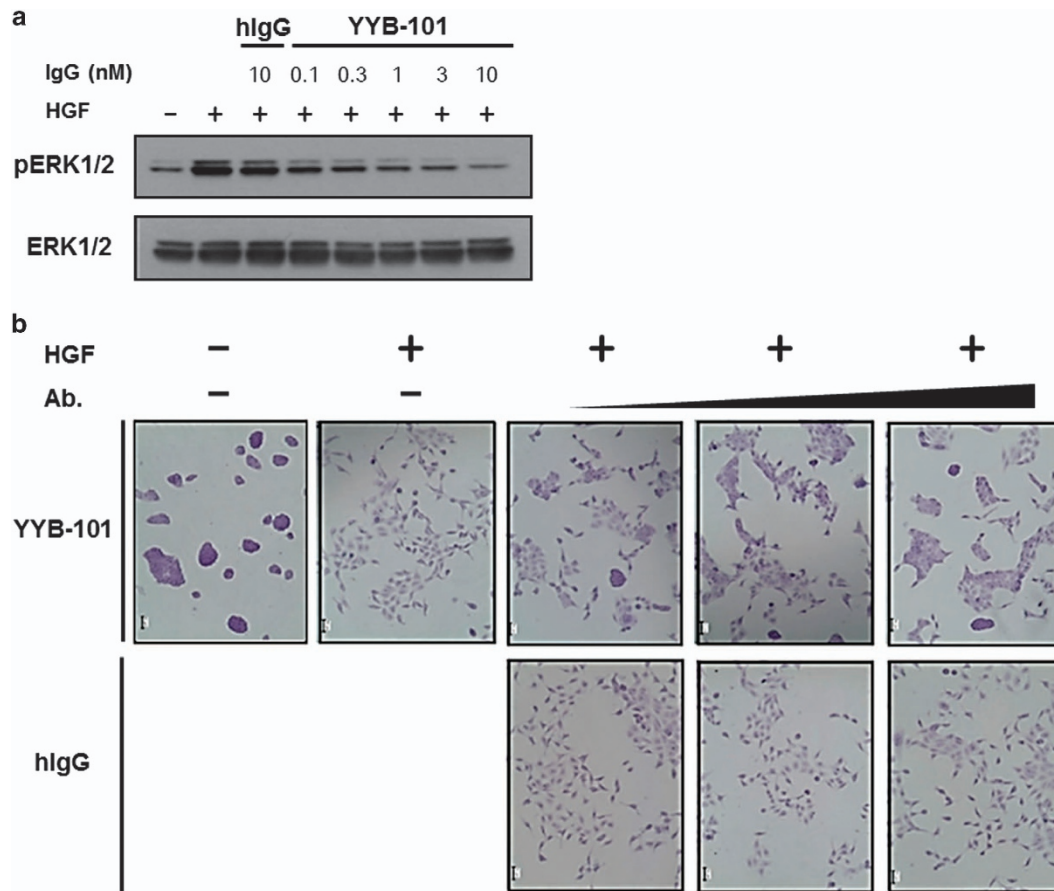


Figure 1 Neutralization of hepatocyte growth factor (HGF) by YYB-101. (a) HepG2 cells were treated with 130 pM HGF alone or in combination with human IgG (hIgG) or YYB-101, and HGF/cMET signaling was evaluated by immunoblot using anti-phosphoERK1/2 and anti-ERK1/2 antibodies. (b) HGF-induced scatter was evaluated by incubating MDCK-2 cells with 67 pM HGF alone or in combination with either hIgG or YYB-101 (1, 10 or 100 nM).

Tissue cross-reactivity of YYB-101

The cross-reactivity of YYB-101 was evaluated in cryosections of 40 different tissue samples from three human donors, kidney and liver tissue samples from nine human donors, and 10 tissue samples from three cynomolgus monkeys. The study was designed to meet the requirements of the US Food and Drug Administration and the European Medicines Agency. In brief, YYB-101 and the control human IgG4 (Abcam, Cambridge, MA, USA) were labeled using the ImmunoProbe Biotinylation Kit (Sigma-Aldrich) according to the manufacturer's instructions. Tissue sections were fixed in acetone and tested with 0.02–10 $\mu\text{g ml}^{-1}$ of biotinylated YYB-101 and biotinylated human IgG4. Three concentrations of biotinylated YYB-101 were selected for antibody titration: 1.25, 0.63 and 0.31 $\mu\text{g ml}^{-1}$. Slides were incubated with either biotinylated YYB-101 or biotinylated human IgG4 diluted in 0.2% bovine serum albumin at room temperature for 1 h, followed by a 45-min incubation with the secondary biotinylated antibody labeled with peroxidase (Vectastain Elite ABC kit, Vector Laboratories, Burlingame, CA, USA). The tissue sections were then stained with 3,3'-diaminobenzidine tetrahydrochloride (Vector Laboratories) and counterstained with hematoxylin. The intensity of the specific immunohistochemical staining was graded on a scale of 1–5, where 1 = negative and 5 = intense reactivity. Cross-reactivity was reported as the mean score (the sum of the values was divided by the number of tissue samples evaluated).

Pharmacokinetics, toxicokinetics and anti-drug antibodies of YYB-101 in cynomolgus monkeys

This study was conducted by the Test and Control Article Department of Medicilon Preclinical Research, LLC, (Shanghai, China) in accordance with regulations outlined in the USDA Animal Welfare act and conditions specified in The Guide for the Care and Use of Laboratory Animals.¹⁸ Four male cynomolgus monkeys (3–5 years old), originally obtained from Guangxi Grandforest Scientific Primate Company, Ltd. (Guangxi, China), were used in this study. At study termination, all animals were transferred to the stock colony.

YYB-101 was diluted in buffer containing 20 mM sodium phosphate, 150 mM NaCl (pH 7.0), and 0.01% Polysorbate 20 (final concentration, 0.6 mg ml^{-1}), sterilized using a 0.22- μm filter, and then administered via intravenous infusion for 2 h. Blood samples were collected from the femoral vein pre-dose and 2, 4, 8, 24, 48, 96, 168, 240, 336, 504 and 672 h after initiation of the infusion. The blood samples were allowed to clot and then centrifuged at 3500 rpm for 10 min at 4 °C. The serum was stored at -80 °C.

The concentration of YYB-101 in the serum samples was determined by using a sandwich enzyme immunoassay. The microtiter plate (Corning, Oneonta, NY, USA) was pre-coated with recombinant human HGF (R&D Systems, Minneapolis, MN, USA) before adding serum samples and standards to the wells. After incubation and washing, analytes were detected by using horseradish peroxidase-conjugated goat anti-human IgG (kappa chain specific) and the

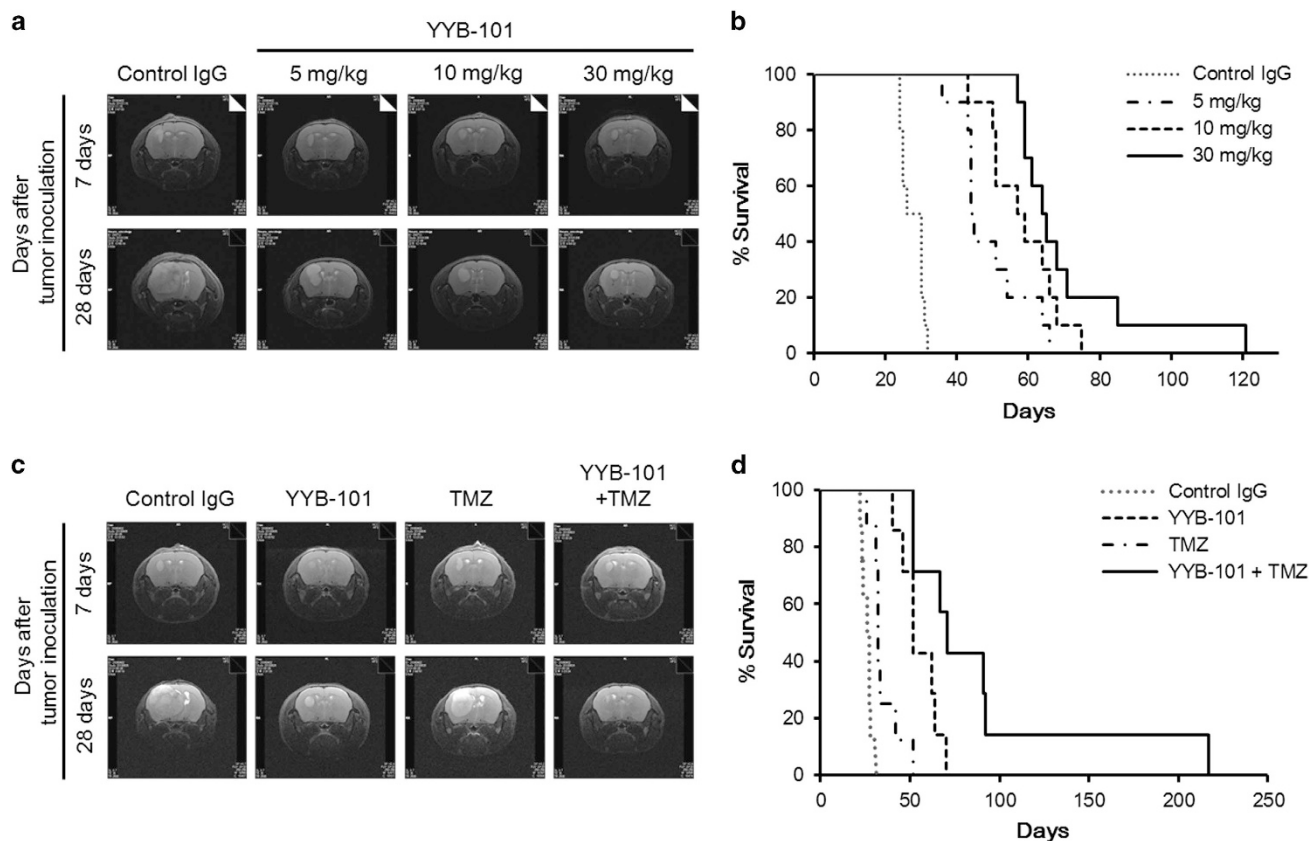


Figure 2 Effect of YYB-101 with or without temozolomide on tumor growth and survival in mice bearing human glioblastoma xenografts. On day 7 after tumor inoculation, the tumor-bearing mice began treatment with IgG (10 mg kg^{-1}) or YYB-101 ($5, 10$ or 30 mg kg^{-1} , $n=10$ per group), which was administered three times per week for 3 weeks. (a) MRI brain scans performed at 7 and 28 days. (b) A Kaplan–Meier curve was constructed, and survival was analyzed by log-rank test (control versus each YYB-101-treated group, $P<0.001$; 5 mg kg^{-1} YYB-101 versus 30 mg kg^{-1} YYB-101, $P=0.0024$). On day 7 after tumor inoculation, the tumor-bearing mice received 10 mg kg^{-1} IgG ($n=8$), 10 mg kg^{-1} YYB-101 ($n=7$), 5 mg kg^{-1} temozolomide (TMZ, $n=8$), or YYB-101+TMZ ($n=7$). (c) MRI brain scans were performed at 7 and 28 days. (d) A Kaplan–Meier curve was constructed, and survival was analyzed by log-rank test (control versus TMZ-treated groups, $P=0.005$; TMZ versus combination treatment, $P=0.0001$).

substrate o-phenylenediamine (Thermo Fisher Scientific). The enzyme-substrate reaction was terminated by adding 1 N H_2SO_4 solution, and the color change was measured spectrophotometrically at 492 nm. Estimates of non-compartmental pharmacokinetic parameters such as time of maximum observed serum concentration (T_{max}), maximum observed serum concentration (C_{max}), area under the serum concentration-time curve from zero to infinity ($\text{AUC}_{0-\infty}$), terminal exponential half-life ($t_{1/2z}$), and clearance were calculated with WinNolin software (version 5, Pharsight Corporation, Cary, NC, USA).

Blood samples were collected from all surviving animals. On days 1 and 22, blood was obtained before infusion and approximately 2, 4, 8, 24, 48, and 96 h after the start of infusion (SOI). On days 8 and 15, blood was collected before infusion and at approximately 2 h from the SOI. Blood was also collected once daily on days 29, 36, 43, 57, 71, and 85 (converted to hours relative to SOI on day 22). The samples obtained 2 h from the SOI were collected within 2 min prior to the end of infusion. Blood samples were also collected at all time points from control animals; however, only the samples collected 2 h from the SOI were analyzed. For each animal, the following toxicokinetic parameters were determined: maximum observed serum concentration (C_{max}), time of maximum observed serum concentration (T_{max}), and area under the serum concentration-time curve (AUC). The

AUC from time 0 to 168 h ($\text{AUC}_{0-168\text{h}}$) and AUC from time 0 to the time of the final quantifiable sample (AUC_{last}) were calculated by the linear trapezoidal method.¹⁹ The accumulation ratio (R) was calculated for each dose group as follows:

$$R = \text{AUC}_{0-168\text{h Day 22}} \div \text{AUC}_{0-168\text{h Day 1}}$$

An enzyme immunoassay was used to detect anti-YYB-101 antibodies in the sera of cynomolgus monkeys. Briefly, a streptavidin-coated plate (Meso Scale Discovery, Gaithersburg, MD, USA) was blocked with blocking buffer (3% bovine serum albumin in PBS). HGF mAb master mix containing YYB-101-biotin, YYB-101-SULFO-TAG, and assay buffer (0.5% bovine serum albumin, 0.05% Tween 20, 0.01% sodium azide in PBS) was then added to the wells and incubated at room temperature for 30 min. After washing the plate, the samples were added to the wells and incubated at room temperature for 2 h. The plate was washed, $2 \times$ Read Buffer T (Meso Scale Discovery) was added to the wells, and the plates were read within 20 min.

Serum human HGF levels in a human HGF knock-in transgenic mouse after intravenous injection of YYB-101

YYB-101 was administered via single intravenous injection ($10, 50$, or 200 mg kg^{-1}) to *Hgf^{tm1.1(HGF)Aveo Prkdc^{scid}/J}* mice (Stock #014543,

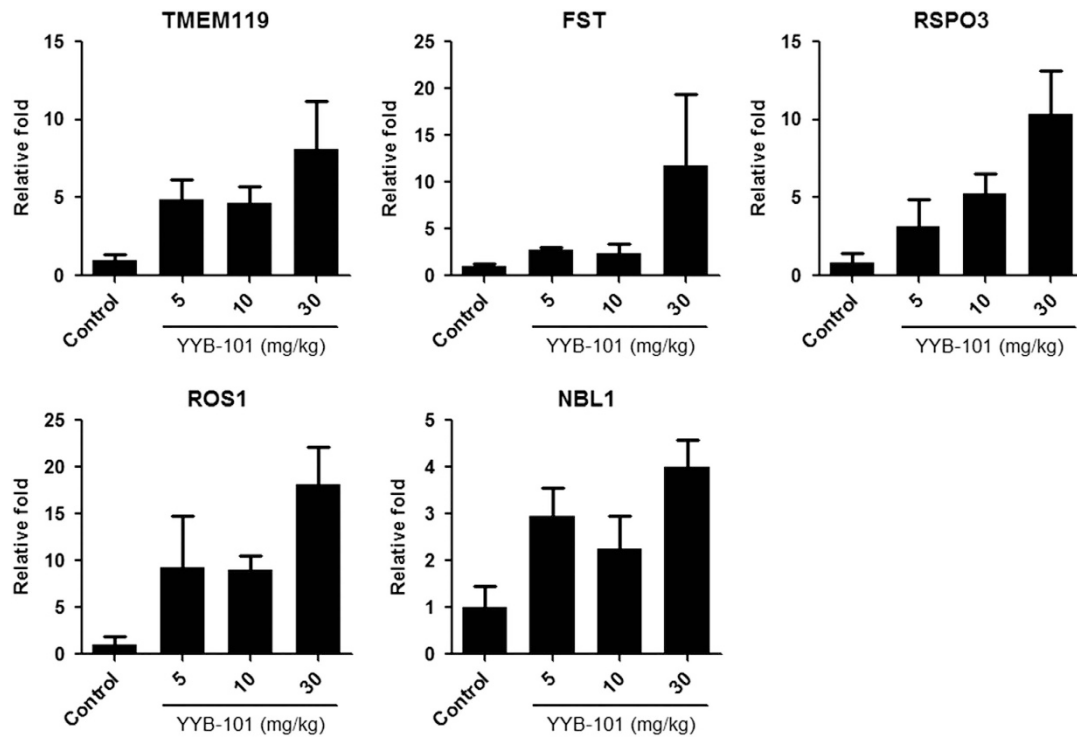


Figure 3 Effect of hepatocyte growth factor (HGF) inhibition on the expression of carcinogenesis-related genes. The relative expression of five carcinogenesis-related genes (TMEM119, FST, RSPO3, ROS1 and NBL1) was determined by real-time PCR in tumor tissue of mice treated with 10 mg kg⁻¹ IgG (control) or 5, 10 or 30 mg kg⁻¹ YYB-101 (each group, *n*=2). The results are expressed as the mean ± s.d. from three independent experiments (control versus each YYB-101-treated group; *P*<0.03).

The Jackson Laboratory, Bar Harbor, ME, USA), which possess a knock-in mutation of the human *HGF* gene. Blood was collected before the injection and 0.5, 4, 24, 48, 168, 336 and 504 h after the injection. Serum was prepared as described earlier. The concentration of human HGF in the serum samples was determined using an enzyme immunoassay kit (Human HGF Quantikine ELISA kit, R&D Systems, Minneapolis, MN, USA) according to the manufacturer's instructions.

Measurement of serum HGF levels in sarcoma patients

This study included 73 patients diagnosed with 22 different sarcomas at Seoul National University Hospital. Blood samples were obtained from September 2013 to July 2015. The serum was separated, aliquoted, and stored at -80 °C until analysis. HGF concentrations in the serum were determined using an enzyme-linked immunosorbent assay kit (Human HGF Quantikine ELISA Kit, R&D Systems) following the manufacturer's instructions.

RESULTS

Therapeutic efficacy of the HGF neutralizing antibody YYB-101

To evaluate the effect of YYB-101 on HGF/cMET signaling, HepG2 cells were treated with 130 pM HGF along with YYB-101 or IgG. The results of the immunoblot analysis show that YYB-101 decreased ERK1/2 phosphorylation in a dose-dependent manner (Figure 1a). In addition, YYB-101 neutralized the scattering of HGF-treated MDCK-2 cells in a dose-dependent manner, whereas the human IgG control did not show any effect at the same concentrations (Figure 1b).

To determine the efficacy of YYB-101 against tumor growth and survival *in vivo*, nude mice bearing human glioblastoma xenografts were treated with YYB-101 (5, 10, or 30 mg kg⁻¹; *n* = 10 per group) three times a week for 3 weeks. MRI scans of the mouse brain showed that tumor growth was strongly inhibited by YYB-101, and the tumor growth rate negatively correlated with antibody dosage (Figure 2a). Overall survival was longer in YYB-101-treated mice than in IgG-treated control mice (*P*<0.0001; Figure 2b). In consideration of clinical practice, we also tested YYB-101 in combination with TMZ, the standard-of-care chemotherapy drug used for glioblastoma. Our results showed that combination treatment inhibited tumor growth more effectively than YYB-101 alone (*P*=0.0164) or TMZ alone (*P*=0.0001; Figure 2c) and increased overall survival compared with the survival observed with either agent alone (Figure 2d).

However, as with other treatments, brain tumors treated with YYB-101 eventually showed regrowth. To determine which genes are involved in tumor regrowth, we analyzed microarray data from tumor tissue treated with YYB-101 or IgG (control) and identified 30 genes that were highly expressed in YYB-101-treated tumors compared with expression in controls (Supplementary Table S1). Five of these genes (TMEM119, FST, RSPO3, ROS1 and NBL1) are associated with carcinogenesis. We analyzed the expression of these genes by real-time PCR, which confirmed the microarray data (Figure 3). Our results showed that expression of the carcinogenesis-related genes was

Table 1 Tissue cross-reactivity of YYB-101

	Human			Cynomolgus monkey		
	YYB-101 concentration ($\mu\text{g ml}^{-1}$)					
	1.25	0.63	0.31	1.25	0.63	0.31
Liver ^a	1.89	1.11	0.56	–	–	–
Lung	1.67	1	0.67	–	–	–
Breast	1.67	1.67	0.67	–	–	–
Colon	1	0.33	–	–	–	–
Heart	0.33	–	–	–	–	–
Kidney ^a	1.33	0.78	0.56	–	–	–
Prostate	2.33	2	1	–	–	–
Urinary bladder	1.33	1	0.67	0.33	–	–
Pituitary	2.67	2	1.67	–	–	–
Placenta	4	3.67	2.67	–	–	–

– No positive staining.
^aNumber of human donors = 9.

significantly higher in YYB-101-treated tissue than expression in controls (each comparison, $P < 0.03$), suggesting the involvement of these genes in tumor regrowth due to YYB-101 resistance.

Cross-reactivity of YYB-101 with human and cynomolgus monkey tissue

Tissue cross-reactivity studies were assessed in tissue from human donors and cynomolgus monkeys (Table 1). In human tissue, specific staining for YYB-101 was observed in blood vessel walls, epithelial cells, urothelium, connective tissue, stromal tissue, and smooth muscle. In addition, staining for YYB-101 was observed in the brain (Purkinje cells of the cerebellum), lens of the eye, kidney (tubule walls and Bowman's capsule), liver (hepatocytes), and scattered cells in the lung (data not shown).

In the cynomolgus monkey tissue, specific staining for YYB-101 was observed in blood vessel walls within the urinary bladder only. In the 10 tissue samples examined as part of this study, no concordance was observed between humans and cynomolgus monkeys, possibly because of the lower affinity of YYB-101 for monkey HGF (data not shown).

Pharmacokinetics of YYB-101

The pharmacokinetics of YYB-101 was investigated in four male cynomolgus monkeys after a single intravenous injection (10 mg kg^{-1}). The serum concentration versus time curves and pharmacokinetic parameters of YYB-101 are presented in Figure 4. The YYB-101 serum T_{max} was 2 h, C_{max} was $221.57 \mu\text{g ml}^{-1}$, and $\text{AUC}_{(0-\infty)}$ was $94\,802.96 \mu\text{g ml}^{-1} \text{h}^{-1}$. The $t_{1/2z}$ was ~ 21.7 days and clearance was $0.11 \text{ ml kg}^{-1} \text{h}^{-1}$.

Toxicokinetics of YYB-101 and anti-drug antibodies

Mean YYB-101 serum concentration-time profiles and toxicokinetic parameters are shown in Table 2. Systemic exposure to YYB-101 was independent of sex (data not shown). Following

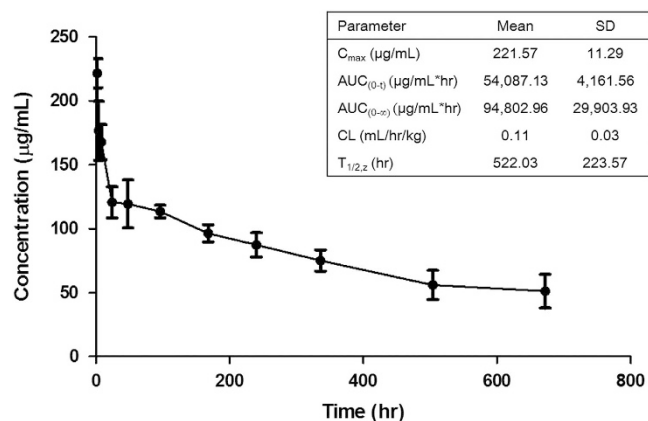


Figure 4 Pharmacokinetic profile of YYB-101 in cynomolgus monkeys. Serum concentration-time curve of YYB-101 in male cynomolgus monkeys following a 2-h intravenous infusion at 10 mg kg^{-1} . The inserted table describes selected pharmacokinetic parameters. The results are expressed as the mean \pm s.d. ($n = 3$).

intravenous administration of YYB-101, the mean systemic exposure ($\text{AUC}_{0-168 \text{ h}}$) and C_{max} values of YYB-101 increased proportionally with dose. The mean peak and trough serum concentrations of YYB-101 appeared to approach steady state following the four weekly infusions of YYB-101 (Table 2). Serum concentrations were quantifiable in recovered animals 63 days after the last dose ($\sim 2.8\%$ of C_{max} from day 22). Systemic exposure ($\text{AUC}_{0-168 \text{ h}}$) increased with repeated intravenous administration of YYB-101, with accumulation ratios ranging from 2.38 to 2.95. Anti-drug antibodies were detected on day 1 in one female monkey at 50 mg kg^{-1} per day YYB-101 but were not detected in samples collected from this animal on day 29 or 85. The C_{max} and AUC values of this animal were similar to those of animals in which no anti-drug antibodies were detected (data not shown).

Changes in serum human HGF levels in the human HGF knock-in mouse

Because YYB-101 does not bind to mouse HGF, we used human HGF knock-in mice ($Hgf^{tm1.1(HGF)Aveo} Prkdc^{scid/J}$) to evaluate the serum HGF level after injection of YYB-101. In this mouse model, exons 2–18 of the human HGF gene were inserted into exons 3–6 of the endogenous mouse Hgf gene by homologous recombination. The serum HGF concentration was monitored after a single injection of YYB-101 (Figure 5). Before treatment, the mean serum human HGF concentrations were 59, 54, and 71 ng ml^{-1} in mice treated with the low dose (10 mg kg^{-1}), medium dose (50 mg kg^{-1}), and high dose (200 mg kg^{-1}), respectively. The serum human HGF level dropped below 1 ng ml^{-1} at 30 min after the injection in each treatment group. Serum HGF levels were still below 50% of pre-dose values at 48 h in the low-dose group, 168 h in the medium-dose group, and 336 h in the high-dose group. The mean human HGF levels were higher than that of pre-dose values after 168 h in the low-dose group and after 504 h in the medium-dose group.

Table 2 Toxicokinetic parameters on days 1 and 22 following intravenous infusion of YYB-101 at 10, 50, or 200 mg kg⁻¹ perday in cynomolgus monkeys (males and females combined)

Dose (mg kg ⁻¹ day ⁻¹)	Day	Statistic	C _{max} (ng ml ⁻¹)	C _{max} /Dose (kg × ng ml ⁻¹ mg ⁻¹)	T _{max} ^a (h)	T _{last} ^a (h)	AUC _{Tlast} (ng × h ml ⁻¹)	AUC _{0-168hr} (ng × h ml ⁻¹)	AUC _{0-168hr} /Dose (kg × ng × h ml ⁻¹ mg ⁻¹)	R ^b
10	1	N	6	6	2 (2-2)	168 (168-168)	6	6	6	NA
		Mean	363 000	3 6300			31 000 000	31 000 000	3 100 000	NA
		s.d.	24 200	2420			2 980 000	2 980 000	298 000	NA
	22	CV%	7	7			10	10	10	NA
		N	6	6	2 (2-2)	168 (168-168)	6	6	6	6
		Mean	738 000	73 800			77 200 000	77 200 000	7 720 000	2.49
50	1	s.d.	44 300	4430			10 700 000	10 700 000	1 070 000	0.286
		CV%	6	6			14	14	14	11
		N	6	6	2 (2-2)	168 (168-168)	6	6	6	NA
	22	Mean	1 950 000	39 000			163 000 000	163 000 000	3 250 000	NA
		s.d.	289 000	5770			13 100 000	13 100 000	262 000	NA
		CV%	15	15			8	8	8	NA
200	1	N	6	6	2 (2-2)	168 (168-168)	6	6	6	6
		Mean	3 650 000	73 000			387 000 000	387 000 000	7 730 000	2.38
		s.d.	75 200	1500			34 800 000	34 800 000	696 000	0.119
	22	CV%	2	2			9	9	9	5
		N	10	10	2 (2-2)	168 (168-168)	10	10	10	NA
		Mean	7 330 000	36 600			666 000 000	6 660 000 000	3 330 000	NA
200	1	s.d.	670 000	3350			61 500 000	61 500 000	308 000	NA
		CV%	9	9			9	9	9	NA
		N	10	10	2 (2-96)	168 (168-1512)	10	10	10	10
	22	Mean	17 800 000	89 200			3 180 000 000	1 980 000 000	9 910 000	2.95
		s.d.	4 350 000	21 700			1 160 000 000	652 000 000	3 260 000	0.841
		CV%	24	24			36	33	33	29

Abbreviations: AUC, area under the curve; CV, coefficient of variation; NA, not applicable.

^aMedian (minimum—maximum), median value only reported if actual collection interval.

^bR = AUC_{0-168 h Day 22}/AUC_{0-168 h Day 1}.

Serum HGF level in cancer patients

The distribution of serum HGF level of the 73 patients included in this study was shown in Figure 6. The median serum HGF level was 1158 pg/ml (range, 477–13 474), which was considerably greater than the serum HGF level in a normal population (260–390 pg ml⁻¹).²⁰ The most common sarcoma was osteosarcoma (*n* = 18), and the median serum HGF level in these patients was 1223 pg ml⁻¹ (range, 692–13 474).

DISCUSSION

Dysregulated HGF-cMET signaling promotes cell survival, proliferation, and invasion. High HGF levels are detected in the serum and tumor tissue of patients with various cancers and are correlated with poor prognosis. Thus, HGF is considered a promising target for cancer therapy, and several HGF inhibitors have been evaluated in clinical studies. Rilotumumab, a fully human anti-HGF antibody, is the most extensively tested HGF inhibitor; however, two phase III clinical studies were halted because of patient deaths.²¹ Clinical trials of combination treatment with the humanized anti-HGF antibody ficlatuzumab and gefitinib in pulmonary adenocarcinoma patients failed to show better efficacy than gefitinib monotherapy.²² The results of a phase I study of the humanized anti-HGF antibody HuL2G7 in patients with advanced solid malignancies showed some adverse events but no dose-limiting toxicities.²³ To date, none of these HGF neutralizing antibodies have shown efficacy in a clinical setting; therefore, we investigated YYB-101 as a potential therapeutic agent.

In this study, we showed that YYB-101 inhibits cMET activation and cell scattering *in vitro* and suppresses tumor growth in mice bearing human glioblastoma xenografts, which

is consistent with results in mouse models of colorectal cancer.²³ In addition, combination treatment with YYB-101 and TMZ in our study and with irinotecan in the colorectal cancer model was more effective than treatment with YYB-101 alone. We also identified genes that were overexpressed after tumor relapse in the mouse xenograft model, which may be useful in the selection of patients who would be more likely to benefit from YYB-101 treatment.

Rilotumumab binds the beta chain of HGF, which binds cMET with low affinity, leaving the alpha chain free to engage cMET, resulting in partial antagonism.²⁴ In contrast, YYB-101 binds to the alpha chain of HGF (manuscript under preparation), which binds cMET with high affinity.²⁵ For this reason, YYB-101 may completely block HGF binding to cMET and be more effective in a clinical setting. One of the unique findings of this study was the rapid decrease in serum HGF levels in human HGF knock-in mice, which recovered in a dose-dependent manner. In contrast, injection of rilotumumab increased the total plasma level of HGF (free HGF and HGF bound to rilotumumab).²⁶ We discovered that YYB-101 competes with one of the antibodies used in the sandwich enzyme immunoassay. Although it is not clear whether the total HGF level was decreased by YYB-101 injection, the amount of HGF free to bind cMET was reduced.

A recent study reported that serum HGF level is a biomarker that can predict the outcome and resistance to treatment with trastuzumab in HER2-positive patients with metastatic gastric cancer.²⁷ Our result that YYB-101 can markedly decrease serum HGF level suggests that the combination of trastuzumab and YYB-101 may provide clinical benefit in these patients. Similarly,

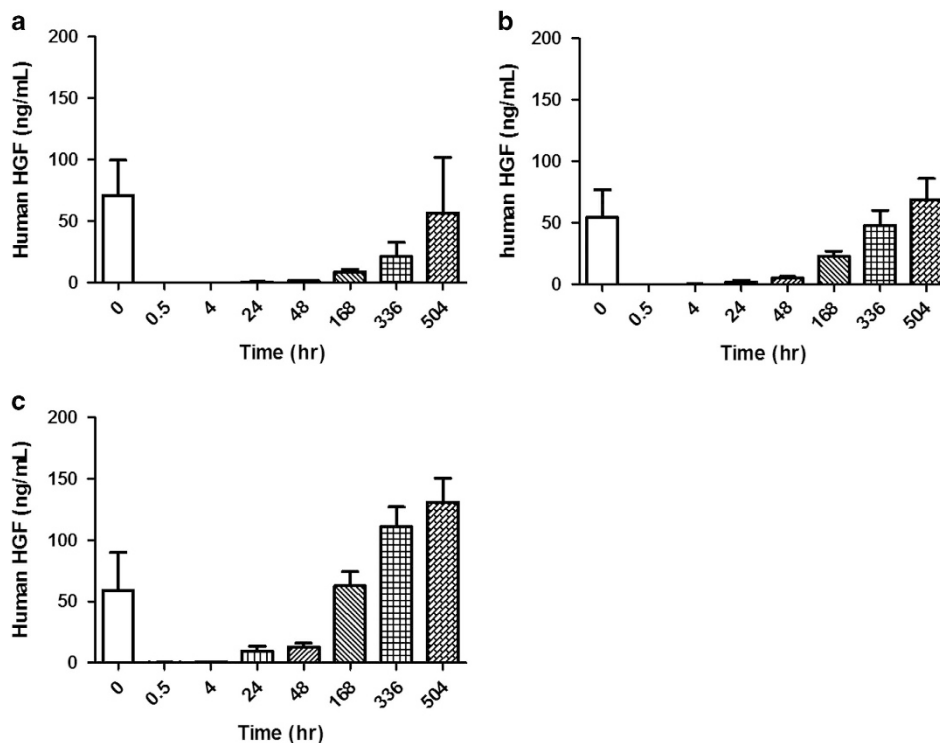


Figure 5 Serum hepatocyte growth factor (HGF) concentration after a single intravenous bolus dose of YYB-101 in human HGF knock-in SCID mice. The mice were treated with (a) 200 mg kg⁻¹ YYB-101, (b) 50 mg kg⁻¹ YYB-101, or (c) 10 mg kg⁻¹ YYB-101. Results are expressed as the mean and \pm s.d. (each group, $n=5$).

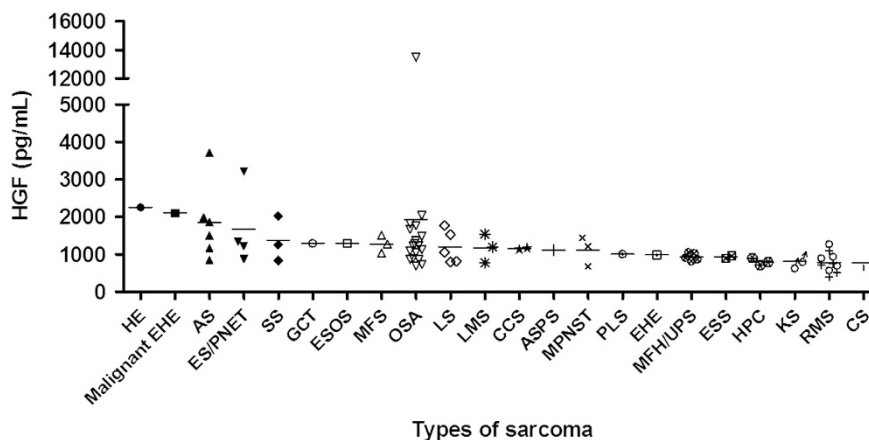


Figure 6 Serum concentration of HGF in sarcoma patients. Each point represents an individual patient, and horizontal lines represent mean values. AS, angiosarcoma; ASPS, alveolar soft part sarcoma; CCS, clear cell sarcoma; CS, chondrosarcoma; EHE, epithelioid hemangioendothelioma; ES/PNET, Ewing sarcoma/primitive neuroectodermal tumor; ESOS, extraskeletal osteosarcoma; ; ESS, endometrial stromal sarcoma GCT, giant cell tumor; HE, hemangioendothelioma; HPC, hemangiopericytoma; KS, Kaposi sarcoma; LMS, leiomyosarcoma; LS, liposarcoma; MFH/UPS, malignant fibrous histiocytoma/pleomorphic undifferentiated sarcoma; MFS, myxofibrosarcoma; MPNST, malignant peripheral nerve sheath tumor; OSA, osteosarcoma; PLS, pleomorphic liposarcoma; RMS, rhabdomyosarcoma; SS, synovial sarcoma.

YYB-101 may be useful in the treatment of ovarian cancer given that an increased serum HGF level is a marker for poor prognosis in this disease,⁹ and the peritoneal spreading of ovarian cancer is enhanced by ascites-mediated activation of cMET.^{28,29}

There were no unusual findings regarding the pharmacokinetic and toxicokinetic properties of YYB-101 in cynomolgus monkeys. The peak serum concentration of YYB-101 was

observed at 2 h after infusion, followed by a steep decrease until 24 h and then a slow elimination, which is comparable to that of other clinically approved antibodies. Anti-drug antibodies were detected in only one animal at a single time point, and little cross-reactivity to normal tissue was observed. On the basis of these results, a phase I clinical study is ongoing in patients with advanced solid tumors (NCT02499224).

CONFLICT OF INTEREST

The authors declare no conflict of interest.

ACKNOWLEDGEMENTS

This study was supported by the National OncoVenture Program (No. H111C1191) and research grants from the National Cancer Center Grants (NCC0911670 and NCC1111370). The study protocol was approved by the Institutional Review Board of Seoul National University Hospital (H1412-040-632). Written informed consent was obtained from all patients in accordance with the Declaration of Helsinki.

- Lokker NA, Mark MR, Luis EA, Bennett GL, Robbins KA, Baker JB *et al*. Structure-Function Analysis of Hepatocyte Growth-Factor—Identification of Variants That Lack Mitogenic Activity yet Retain High-Affinity Receptor–Binding. *EMBO J* 1992; **11**: 2503–2510.
- Trusolino L, Comoglio PM. Scatter-factor and semaphorin receptors: cell signalling for invasive growth. *Nat Rev Cancer* 2002; **2**: 289–300.
- Gherardi E, Birchmeier W, Birchmeier C, Vande Woude G. Targeting MET in cancer: rationale and progress. *Nat Rev Cancer* 2012; **12**: 89–103.
- Singh-Kaw P, Zarnegar R, Siegfried JM. Stimulatory effects of hepatocyte growth factor on normal and neoplastic human bronchial epithelial cells. *Am J Physiol* 1995; **268**(6 Pt 1): L1012–L1020.
- Zhang YW, Vande Woude GF. HGF/SF-met signaling in the control of branching morphogenesis and invasion. *J Cell Biochem* 2003; **88**: 408–417.
- Maulik G, Shrikhande A, Kijima T, Ma PC, Morrison PT, Salgia R. Role of the hepatocyte growth factor receptor, c-Met, in oncogenesis and potential for therapeutic inhibition. *Cytokine Growth Factor Rev* 2002; **13**: 41–59.
- Birchmeier C, Birchmeier W, Gherardi E, Vande Woude GF. Met, metastasis, motility and more. *Nat Rev Mol Cell Biol* 2003; **4**: 915–925.
- Christensen JG, Burrows J, Salgia R. c-Met as a target for human cancer and characterization of inhibitors for therapeutic intervention. *Cancer Lett* 2005; **225**: 1–26.
- Aune G, Lian AM, Tingulstad S, Torp SH, Forsmo S, Reseland JE *et al*. Increased circulating hepatocyte growth factor (HGF): a marker of epithelial ovarian cancer and an indicator of poor prognosis. *Gynecol Oncol* 2011; **121**: 402–406.
- Edakuni G, Sasatomi E, Satoh T, Tokunaga O, Miyazaki K. Expression of the hepatocyte growth factor/c-Met pathway is increased at the cancer front in breast carcinoma. *Pathol Int* 2001; **51**: 172–178.
- Yamashita J, Ogawa M, Yamashita S, Nomura K, Kuramoto M, Saishoji T *et al*. Immunoreactive hepatocyte growth factor is a strong and independent predictor of recurrence and survival in human breast cancer. *Cancer Res* 1994; **54**: 1630–1633.
- Maemura M, Iino Y, Yokoe T, Horiguchi J, Takei H, Koibuchi Y *et al*. Serum concentration of hepatocyte growth factor in patients with metastatic breast cancer. *Cancer Lett* 1998; **126**: 215–220.
- Siegfried JM, Weissfeld LA, Singh-Kaw P, Weyant RJ, Testa JR, Landreneau RJ. Association of immunoreactive hepatocyte growth factor with poor survival in resectable non-small cell lung cancer. *Cancer Res* 1997; **57**: 433–439.
- Kasahara K, Arao T, Sakai K, Matsumoto K, Sakai A, Kimura H *et al*. Impact of serum hepatocyte growth factor on treatment response to epidermal growth factor receptor tyrosine kinase inhibitors in patients with non-small cell lung adenocarcinoma. *Clin Cancer Res* 2010; **16**: 4616–4624.
- Kim K, Hur Y, Ryu EK, Rhim JH, Choi CY, Baek CM *et al*. A neutralizable epitope is induced on HGF upon its interaction with its receptor cMet. *Biochem Biophys Res Commun* 2007; **354**: 115–121.
- Woo JK, Kang JH, Kim B, Park BH, Shin KJ, Song SW *et al*. Humanized anti-hepatocyte growth factor (HGF) antibody suppresses innate irinotecan (CPT-11) resistance induced by fibroblast-derived HGF. *Oncotarget* 2015; **6**: 24047–24060.
- Burnette WN. ‘Western blotting’: electrophoretic transfer of proteins from sodium dodecyl sulfate–polyacrylamide gels to unmodified nitrocellulose and radiographic detection with antibody and radioiodinated protein A. *Anal Biochem* 1981; **112**: 195–203.
- Institute of Laboratory Animal Research CoLS, National Research Council *Guide for the Care and Use of Laboratory Animals*. National Academy Press: Washington, DC, USA, 1996.
- M. Gibaldi DP. *Pharmacokinetics*, 2nd edn, Taylor & Francis: Oxfordshire, UK, 1982.
- Funakoshi H, Nakamura T. Hepatocyte growth factor: from diagnosis to clinical applications. *Clin Chim Acta* 2003; **327**: 1–23.
- Cunningham DTN, Davidenko I, Murad AM. Phase III, randomized, double-blind, multicenter, placebo (P)-controlled trial of rilotumumab (R) plus epirubicin, cisplatin and capecitabine (ECX) as first-line therapy in patients (pts) with advanced MET-positive (pos) gastric or gastroesophageal junction (G/GEJ) cancer: RILOMET-1 study. *J Clin Oncol* 2015; **33**: (suppl:abstr 4000).
- Mok TS, Geater SL, Su WC, Tan EH, Yang JC, Chang GC *et al*. A randomized phase 2 Study comparing the combination of ficlatuzumab and gefitinib with gefitinib alone in Asian patients with advanced stage pulmonary adenocarcinoma. *J Thorac Oncol* 2016; **11**: 1736–1744.
- Woo JK, Kang JH, Kim B, Park BH, Shin KJ, Song SW *et al*. Humanized anti-hepatocyte growth factor (HGF) antibody suppresses innate irinotecan (CPT-11) resistance induced by fibroblast-derived HGF. *Oncotarget* 2015; **6**: 24047–24060.
- Greenall SA, Adams TE, Johns TG. Incomplete target neutralization by the anti-cancer antibody rilotumumab. *MABs* 2016; **8**: 246–252.
- Kirchhofer D, Yao X, Peek M, Eigenbrot C, Lipari MT, Billeci KL *et al*. Structural and functional basis of the serine protease-like hepatocyte growth factor beta-chain in Met binding and signaling. *J Biol Chem* 2004; **279**: 39915–39924.
- Zhang Y, Doshi S, Zhu M. Pharmacokinetics and pharmacodynamics of rilotumumab: a decade of experience in preclinical and clinical cancer research. *Br J Clin Pharmacol* 2015; **80**: 957–964.
- Rimassa L, Abbadessa G, Personeni N, Porta C, Borbath I, Daniele B *et al*. Tumor and circulating biomarkers in patients with second-line hepatocellular carcinoma from the randomized phase II study with tivantinib. *Oncotarget* 2016; **7**: 72622–72633.
- Nakamura M, Ono YJ, Kanemura M, Tanaka T, Hayashi M, Terai Y *et al*. Hepatocyte growth factor secreted by ovarian cancer cells stimulates peritoneal implantation via the mesothelial-mesenchymal transition of the peritoneum. *Gynecol Oncol* 2015; **139**: 345–354.
- Matte I, Lane D, Laplante C, Garde-Granger P, Rancourt C, Piche A. Ovarian cancer ascites enhance the migration of patient-derived peritoneal mesothelial cells via cMet pathway through HGF-dependent and -independent mechanisms. *Int J Cancer* 2015; **137**: 289–298.



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