



A phase 2 basket trial of combination therapy with trastuzumab and pertuzumab in patients with solid cancers harboring human epidermal growth factor receptor 2 amplification (JUPITER trial)

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Abstract

Introduction: Human epidermal growth factor receptor 2 (*HER2*) gene amplification and mutations have emerged as oncogenic drivers and therapeutic targets not limited to breast and gastric cancers, but also in a variety of cancers. However, even if an actionable gene alteration is found, the incidence of *HER2* amplification in these cancers is less than 5%. It is too difficult to conduct a conventional randomized, controlled trial in a rare fraction. Therefore, we have designed a organ-agnostic basket study, which covers a variety of solid cancers harboring *HER2* amplification, in 1 study protocol.

Methods/Design: This trial is a multicenter, single-arm, basket phase 2 study in Japan. Patients with solid cancers harboring *HER2* amplification that have progressed with standard treatment, or rare cancers for which there is no standard treatment, will be eligible. Target cancers include bile duct, urothelial, uterine, ovarian, and other solid cancers where *HER2* amplification is detected by comprehensive genomic profiling using next-generation sequencing technology. A total of 38 patients will be treated with combination therapy with trastuzumab and pertuzumab every 3 weeks until disease progression, unmanageable toxicity, death, or

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patient refusal. The primary endpoint is the objective response rate, and secondary endpoints are progression-free survival, overall survival, and duration of response.

Discussion: The aim of this trial is to evaluate the safety and efficacy of combination therapy with trastuzumab and pertuzumab in patients with locally advanced or metastatic, solid cancers harboring *HER2* amplification. Instead of focusing on 1 organ type, our trial design uses a basket study focusing on *HER2* amplification, regardless of the site or origin of the cancer. The results of our study will advance clinical and scientific knowledge concerning the treatment of locally advanced, rare solid cancers harboring *HER2* amplification, using the combination of trastuzumab and pertuzumab.

Trial registration: This trial was registered in Japan Registry of Clinical Trials (jCRT) on February 25, 2019, as jRCT2031180150.

Abbreviations: AE = adverse event, CI = confidence interval, HER = human epidermal growth factor receptor, ICR = central radiology imaging review, ORR = objective response rate, OS = overall survival, PFS = progression free survival, RECIST = response evaluation criteria in solid tumors, SAF = safety analysis set.

Keywords: basket trial, human epidermal growth factor receptor 2 amplification, pertuzumab, phase 2 trial, solid cancers, trastuzumab

1. Introduction

In the past decade, next-generation sequencing and comprehensive genomic profiling have been developed to facilitate detailed classification of cancers, and efficient and unbiased detection of clinically actionable mutations. [1-3] Genetic alterations underlying the cancer pathology reflect cancer biology and might predict response to therapy better than histology. [4,5] It is well established that human epidermal growth factor receptor 2 (HER2) is an oncogene, associated with higher risk of recurrence and a poor prognosis in breast cancer. Past studies showed that molecular targeted therapies against HER2 amplification improved prognosis of HER2 positive breast and gastric cancer patients. [6] Gene amplification and overexpression of this protein are seen in approximately 20% to 30% of breast cancer cases.^[7] Trastuzumab, the first HER2-directed monoclonal antibody, is a milestone in the treatment of HER2-positive breast cancer at all stages of the disease and in all treatment lines.[8] Four anti-HER2 agents have been currently used for the treatment of breast cancer in Japan. Among these, the combination of trastuzumab with the second HER2 monoclonal antibody pertuzumab has shown substantial effect in HER2-amplifed breast cancer. [9,10] The HER2 pathway also plays a key role in pathogenesis of gastric cancer, in which 1.2% to 9% of patients are HER2-positive. [11] This second indication for trastuzumab has been approved in Japan for HER2-positive advanced or recurrent gastric cancer. HER2 amplification and mutations have emerged as oncogenic drivers and therapeutic targets not limited to breast and gastric cancer, but also in a variety of cancers^[12] such as lung,^[13] colorectal,^[14] bladder,^[15] and biliary,^[16] urothelial carcinoma, [17] gynecologic, [18] and head and neck cancers. [19] Even if an actionable gene mutation is found in these cancers, the incidence of HER2 amplification is less than 5%. [4] Despite its considerable therapeutic potential, the evidence is not matured yet to use the treatment in routine clinical practice. In order to address this unmet clinical need, MyPathway study, a phase 2a study that combines multiple basket studies under an adaptable master protocol to evaluate the efficacy of treatments that target molecular alterations in HER2 (trastuzumab plus pertuzumab), BRAF(vemurafenib), the Hedgehog pathway (vismodegib), or EGFR (erlotinib) in patients with tumor types outside current labeling for these treatment regimens, is in progress.^[20] Interestingly, Hainsworth et al reported that 30 of 114 patients (26%; 95% confidence interval [CI], 19%–35%) with HER2 amplification/overexpression had objective responses to treatment

with trastuzumab plus pertuzumab.^[20] The objective responses were seen in 9 primary tumor types: colorectal, bladder, biliary, salivary gland, non-small cell lung, pancreas, ovary, prostate, and skin cancers.

In consideration of the MyPathway study, we designed a histology-independent phase 2 basket trial of combination therapy with trastuzumab and pertuzumab in Japanese patients with cancer types other than breast and gastric cancers harboring *HER2* amplification.

1.1. Objective

The objective of this trial is to evaluate the safety and efficacy of combination therapy with trastuzumab and pertuzumab in patients with locally advanced or metastatic solid cancers harboring *HER2* amplification such as bile duct, urothelial, uterine, ovarian, and other solid cancers.

2. Methods

2.1. Trial organization and ethical matters

JUPITER trial (A Japanese basket trial using comprehensive genomic profiling informed tumor-agonistic therapy) is an investigator-initiated clinical trial. The study protocol (Ver. 1.0, 2019/1/7) was designed by the study initiators at the Cancer Center, Tokyo Medical and Dental University, Tokyo, Japan, and an additional 6 study sites will participate (Supplemental Digital Content Table S1, http://links.lww.com/MD/E656). The study protocol and the informed consent form were approved by the institutional review board at Tokyo Medical and Dental University (approval number 2018-1005) and at each participating study sites. All participating patients will provide written, informed consent before initiation of any study-specific procedures (Supplemental material). The study is conducted in accordance with the ethical principles originating in or derived from the Declaration of Helsinki and good clinical practice guidelines. The trial was registered in jCRT (Japan Registry of clinical Trials) as jRCT2031180150 (Date of registration February 25, 2019).

2.2. Study design and participants

This trial is a Japanese multicenter, single-arm, basket phase 2 study in patients with solid cancers harboring *HER2* amplification that have progressed with standard treatment, or rare cancers for

Table 1

Participant eligibility criteria.

Inclusion criteria

- 1. Aged 20 yr or over.
- Histologically proven, locally advanced or metastatic solid cancers such as bile duct, urothelial, uterine, ovarian, and other solid cancers excluding breast, gastric and colorectal cancers, where a response to standard therapy is not expected, or assessed by the investigator as having no available or standard therapy.
- 3. HER2 amplification detected by next-generation sequencing technology at a clinical laboratory, certified according to an international standard such as a CLIA-certified and CAP-accredited laboratory. Sampling time is not limited, and liquid biopsy is allowed.
- 4. Measurable or evaluable lesions according to RECIST v1.1.
- Tumor tissue specimen is available. Tumor tissue in which HER2 amplification is identified by FoundationOne CDxTM is excluded.
- 6. Eastern Cooperative Oncology Group (ECOG) performance status of 0 to 1.
- All of the following laboratory parameters must be within the ranges specified:
 - (a) Hemoglobin ≥9.0 g/dL.
 - (b) Neutrophil count ≥1500/mm³.
 - (c) Platelet count ≥75,000/mm³.
 - (d) Total bilirubin <2.0 mg/dL.
 - (e) AST and ALT < 2.5 x the institutional upper limit of normal (ULN).
 - (f) Serum creatinine <1.5 mg/dL.
- 8. Women of child-bearing potential or men who have a woman of child-bearing potential as a partner must agree to use proper contraception from initiation of treatment with the study drug to 28 wk after treatment.
- 9. Written informed consent.

Exclusion criteria

- Suffering from Grade ≥2 toxicity (Grade ≥3 neuropathy) due to previous treatment during 4 wk before enrollment.
- History of treatment with either trastuzumab or pertuzumab for the same tumor as in this study.
- Genomic variations seen in tumors are known to constitute resistant mutations to trastuzumab or pertuzumab.
- 4. Breast cancer with HER2 amplification or overexpression.
- 5. Gastric cancer with HER2 amplification or overexpression.
- 6. Colorectal cancer with HER2 amplification or overexpression.
- 7. History of anti-cancer therapy including chemotherapeutic agents, biological agents, radiotherapy, or endocrine therapy other than the study drugs during 4 wk before enrollment. However, treatment with megestrol acetate and bisphosphonates, and medical castration for the treatment of prostate cancer as a supportive therapy are allowed.
- 8. Primary brain tumor.
- 9. Progression of brain metastasis or nervous function decline due to brain metastasis.
- 10. Serious heart disease.
- 11. Left ventricular ejection fraction (LVEF) of < 50%.
- 12. History of stroke or acute myocardial infarction during 4 mo before enrollment.
- 13. History of acute gastrointestinal bleeding during 4 wk before enrollment.
- 14. Administration of other investigational drugs during 4 wk before enrollment.
- 15. Patients assessed by the investigator as unwilling to participate in this study due to other clinically important medical conditions, or who have difficulty in complying with the trial requirements, as assessed by the investigator.
- 16. Patients who may be pregnant, pregnant women or lactating women.
- 17. An inappropriate decision made by the investigator.

which there is no standard treatment. Target cancers include bile duct, urothelial, uterine, ovarian, and other solid cancers. Breast, gastric and colorectal cancers are excluded from this study because these indications have been approved or clinical trials are ongoing.

Inclusion and exclusion criteria are shown in Table 1. Patients meeting the eligibility criteria will be accrued in this trial. Patients are treated with intravenous trastuzumab (8 mg/kg loading dose followed by 6 mg/kg maintenance dose) and pertuzumab (840 mg loading dose followed by 420 mg maintenance dose) every 3 weeks until disease progression, experience of unacceptable adverse events (AEs), death, or withdrawal of informed consent (Fig. 1). No interim analyses are planned in this trial. The trial schedule is shown (Supplemental Digital Content Table S2, http://links.lww.com/MD/E657). The study will be conducted between March 2019 and March 2021, and the data cutoff date will be

September 2020. No compensation was provided for participation. In case of adverse events, care is provided without additional financial burden for the participants.

2.3. Drug supply

The monoclonal antibodies trastuzumab and pertuzumab are provided by Chugai Pharmaceutical Co., Ltd., Tokyo, Japan, and stored at each study site.

2.4. Interventions

Dose and regimen of trastuzumab and pertuzumab are in accordance with the Japanese package inserts for each product. [21,22] The loading dose in the first cycle and the following maintenance dose after cycle 2 of trastuzumab are 8 mg/kg and

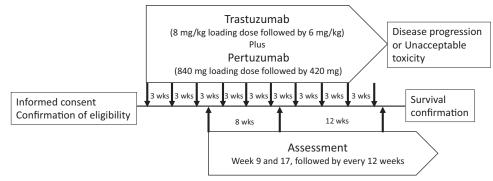


Figure 1. Study design. After informed consent and eligibility confirmation, trastuzumab and pertuzumab are administered every 3 wk until disease progression or unacceptable toxicity. Response assessment is performed at weeks 9 and 17, followed by every 12 wk.

6 mg/kg, respectively. The loading dose in the first cycle and the following maintenance dose after cycle 2 of pertuzumab are 840 mg and 420 mg, respectively. Both drugs are administered intravenously and the administration sequence is not specified. The investigator should confirm that no infusion-related reactions occur during administration and 60 minutes after administration (30 minutes observation after cycle 1). The administration cycle will be repeated every 21 days (3 weeks) until disease progression, the patient experiences unacceptable AEs, death, or withdrawal of informed consent. If the previous administration interval is 6 weeks or longer, administration of loading doses of trastuzumab and pertuzumab is re-started, followed by maintenance doses in the following cycles. If an administered dose is <50% of the loading dose due to infusion-related reactions or other events, if possible, the remaining dose is administered within 1 week of starting administration, and the maintenance dose is administered at day 1 of the next cycle (day 22 from the first cycle). If an administered dose is $\geq 50\%$ to < 75% of the loading dose, if possible the remaining dose is administered within 2 weeks of starting administration. If an administered dose is $\geq 75\%$ of the loading dose, administration of the remaining dose is not necessary. If reduction of left ventricular ejection fraction or AEs for which administration is considered inappropriate are observed, administration is suspended until resolved. Dose reduction is not allowed when re-administration is started. Treatment can be continued after data cutoff according to a separate protocol. Prohibited concomitant medications and therapies are as follows:

- (1) other anti-neoplastic agent,
- (2) other investigational agent,
- (3) radiation, except for palliative radiation for non-targeted lesion.
- (4) Immune suppressive agents (high-dose steroid, tumor necrosis factor-alpha inhibitor) (Supplemental Digital Content Fig. S1, http://links.lww.com/MD/E655).

The investigator can provide appropriate supportive care such as antihistamines and/or steroids for the treatment of infusion reactions; antidiarrheal agents; palliative surgery; palliative radiotherapy for bone metastases; and central venous catheterization, preferably 7 days before initiation of treatment. There is no limitation to subsequent treatment after termination of this study, but it should be started after all required examinations are completed.

2.5. Efficacy endpoints

The primary endpoint for efficacy is the objective response rate (ORR), the proportion of patients who achieve complete response or partial response. Responses are assessed by independent central radiology imaging review (ICR) according to the Response Evaluation Criteria in Solid Tumors (RECIST) version 1.1. Secondary efficacy endpoints include ORR assessed by the investigator; progression free survival (PFS), defined as the time from the day of initiation of treatment to radiologic progression assessed by ICR or death by any cause; overall survival (OS), defined as the time from the day of initiation of treatment to death from any cause; and duration of response, defined as the time of first documentation of an objective response to the date of tumor progression/death. Exploratory efficacy endpoints are the ratio of PFS in the latest treatment to PFS in this trial, and consistency of accuracy and sensitivity of HER2 amplification using FoundationOne CDx and other genetic tests.

2.6. Efficacy assessment

The first time point for the primary assessment is week 8, and the second analysis is performed in week 16 after initiation of treatment. If duration of treatment is ≥16 weeks, assessment is performed every 12 weeks until treatment termination. Tumor shrinkage and progression are assessed using RECIST version 1.1 by ICR. The investigators evaluate efficacy outcomes by themselves, according to the standardized efficacy assessment method, and submit all image data captured at screening and during treatment to ICR.

2.7. Safety assessment

Safety is assessed according to AEs, laboratory tests (hematology and biochemistry), vital signs, body weight, 12-lead electrocardiogram, and echocardiogam. AEs are graded using the common terminology criteria for adverse events v 5.0 (common terminology criteria for adverse events v5.0-JCOG). All AEs are to be documented and the investigator will assess intensity, severity, and relatedness of an AE. All serious AEs is reported according to a standardized serious AEs report form.

2.8. Efficacy and safety assessment committee

The independent committee consists of 3 physicians (medical oncologists). They recommend continuation, changing or discontinuation of the trial to the investigators, based on evaluation of safety information and efficacy information, as necessary.

2.9. Statistical analysis

2.9.1. Sample size. Assuming that the threshold response rate is 5% and the expected response rate is 20%, the sample size was calculated to be 38 at a one-sided level of significance of 2.5%, with power of 80%. The threshold response rate of 5% was set based on the response rate of trastuzumab which was administered to patients without information about gene mutations, in the phase $1 \text{ trial.}^{[23]}$ In the MyPathway study, the response rate to a combination of trastuzumab and pertuzumab was 26% in patients with solid cancers with HER2 amplification, and 20.8% in those excluding colorectal cancer. Therefore, the expected response rate of 20% was set as a clinically significant response rate.

2.9.2. Statistical analysis plan. Safety analysis will be performed for subjects in the safety analysis set (SAF) and efficacy is analyzed for the full analysis set. The SAF consists of subjects who receive the study drug and have a safety measurement. The full analysis set includes the SAF subjects who have at least 1 efficacy measurement after the first treatment. The cut-off date for the primary analysis is the time when the imaging evaluation at week 3 of the last enrolled patient is completed. The primary efficacy endpoint, ORR will be calculated based on data obtained from RECIST version 1.1 by ICR, and the 95% CI will be estimated using the Clopper-Pearson method. For the secondary efficacy endpoint, ORR based on the data assessed by the investigators and the 95% CI will be calculated. The survival curves for PFS and OS will be estimated using Kaplan-Meier method. The median PFS and OS with their 95% CIs will be calculated. Duration of response will be estimated using the Kaplan-Meier method for the subjects' response to the treatment, and the median and the 95% CI will then be estimated.

Subgroup analyses will be performed for primary and secondary endpoints for each cancer type. The response rate in each cancer type will also be estimated using the Bayesian information-borrowing approach. Safety will be assessed for all patients who receive at least 1 treatment with the combination of trastuzumab and pertuzumab. All analyses will be conducted with SAS version 9.4 (SAS Institute, Inc., Cary, NA).

2.10. Data collection and monitoring

All data are collected via a case report form using an electronic data capture system. Monitoring is conducted independently by a contract research organization according to good clinical practice guidelines. Monitoring will be conducted periodically during the trial to confirm the trial is conducted in accordance with the study protocol. Data will not be publicly available.

2.11. Auditing

The contract research organization will perform a planned audit independently from monitoring. Regulatory agencies may also conduct a regulatory inspection of this trial. These audit/inspections can occur at any time during or after completion of the study. The investigator and study site agree to allow the auditor/inspector direct access to all relevant documents. The auditor/inspector can ask the investigator and study site for access to the storage room containing the study products, the pharmacy, and the clinical trial facility.

2.12. Dissemination policy

Trial result will be published as a press-release, or in publication. Also, the result will be presented in academic meetings.

3. Discussion

JUPITER trial is a Japanese phase 2, multicenter, basket trial to evaluate the efficacy and safety of combination therapy with trastuzumab and pertuzumab in patients with locally advanced or metastatic solid cancers harboring *HER2* amplification.

Quartino et al^[24] reported that no evidence of a drug-drug interaction effect was observed between trastuzumab and pertuzumab in the neoadjuvant setting of early breast cancer, and there was no association between the pertuzumab serum concentration and pathological complete response within the dose range of pertuzumab (20–100 µg/mL), suggesting no dose adjustments is needed for patients with lower exposure. Furthermore, the same doses and regimens used in breast are applied in patients with solid cancers other than breast and gastric cancers in the MyPathway study. ^[20] The doses and regimens of trastuzumab and pertuzumab approved for use in breast cancer in Japan were used in this study.

The development of trastuzumab and pertuzumab in breast and gastric cancers was conducted using an organ-specific cancer approach in these cancers according to conventional randomized control trials. Instead of focusing on single cancer type, our trial design is a basket study focusing on *HER2* amplification, regardless of the site or origin of the cancer. Efficacy and safety is evaluated in 1 protocol in a variety of cancers harboring *HER2* amplification such as bile duct, urothelial, uterine, ovarian, and other solid cancers. In our study, we use *HER2* gene amplification defined by next-generation sequencing technology as a biomarker, instead of protein overexpression and fluorescent

in situ hybridization because there are reports that selection of patients using genomic biomarkers revealed better clinical outcomes including higher median ORR, and prolonged median PFS and OS than use of a protein biomarker. [25,26] The incidence of *HER2* amplification in our target cancers is reported to be less than 5%. [4] Although considerable therapeutic potential is expected in these rare cancers, a conventional development strategy requires a randomized controlled trial to obtain appropriate indications. Conversely, our basket study can identify a rare cancer that responds to anti-HER2 agents. This is a paradigm shift to genomic biomarker-driven personalized medicine (precision medicine) in cancer treatment, as pembrolizumab has been approved based on the presence of a biomarker rather than the cancer location, without evidence from a randomized controlled trial for each cancer types. [27]

Several limitations is considered in relation to our trial. First, molecular diversity is known to exist within tumor sites of individual patients, resulting in underestimation of the genomic complexity of solid cancers. [28] Furthermore, *HER2* amplification appears to be an actionable target in some tumor types, but not all. Liquid biopsy might be extremely helpful in understanding or characterizing specific cancer information in real-time fashion, using a simple blood draw, but still is under development for routine clinical use. [29] Second, the planned number of patients enrolled in this trial is 38, which may be adequate to explore efficacy in *HER2*-amplified solid tumors. However, this number is too small for evaluation of the efficacy in each type of cancer, particularly where no response is seen. Therefore, there is a risk to miss potential efficacy due to the small number of patients.

The results of our trial will advance clinical and scientific knowledge about the treatment of rare, locally advanced *HER2*-amplified solid cancers using the combination of trastuzumab and pertuzumab.

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Author contributions

Akihiro Hirakawa: study design, statistical analysis plan, reviewing manuscript.

Eri Ishibashi: concept generation, protocol development, writing manuscript, reviewing manuscript.

Hirotoshi Dosaka-Akita, Chikashi Ishioka, Hisahiro Matsubara, Hiroshi Nishihara, Naoko Aragane, Shinichi Toyooka, Manabu Muto: site PI, accrual of patients, reviewing manuscript.

Kenta Takahashi: concept generation, protocol development, accrual of patients, reviewing manuscript.

Sadakatsu Ikeda: study design, protocol development, accrual of patients, writing manuscript, reviewing manuscript.Satoshi Miyake: overall supervise, reviewing manuscript.

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- Toshio Kubo, Yohei Harada, Hideyuki Hayashi, Masayuki Kano, Yasuhi Shimizu, Hidekazu Shirota, Yukiko Mori: site manager, accrual of patients, reviewing manuscript.
- **Ukihide Tateishi:** protocol development for imaging review and response criteria, reviewing manuscript.

References

- Carr TH, McEwen R, Dougherty B, et al. Defining actionable mutations for oncology therapeutic development. Nat Rev Cancer 2016;16:319– 29
- [2] Morash M, Mitchell H, Beltran H, et al. The role of next-generation sequencing in precision medicine: a review of outcomes in oncology. J Pers Med 2018;8:E30.
- [3] Colombo I, Kurnit KC, Westin SN, et al. Moving from mutation to actionability. Am Soc Clin Oncol Educ Book 2018;23:495–503.
- [4] Zehir A, Benayed R, Shah RH, et al. Mutational landscape of metastatic cancer revealed from prospective clinical sequencing of 10,000 patients. Nat Med 2017;23:703–13.
- [5] Schmidt KT, Chau CH, Price DK, et al. Precision oncology medicine: the clinical relevance of patient-specific biomarkers used to optimize cancer treatment. J Clin Pharmacol 2016;56:1484–99.
- [6] Turner BM, Hicks DG. Pathologic diagnosis of breast cancer patients: evolution of the traditional clinical-pathologic paradigm toward "precision" cancer therapy. Biotech Histochem 2017;92:175–200.
- King CR, Kraus MH, Aaronson SA. Amplification of a novel v-erbB-related gene in a human mammary carcinoma. Science 1985;229:974–6.
 Albagoush SA1, Limaiem F2. HER2. StatPearls [Internet]. Treasure Island (FL): StatPearls Publishing; 2018-.2018 Dec 29.
- [8] Tokunaga E, Oki E, Nishida K, et al. Trastuzumab and breast cancer: developments and current status. Int J Clin Oncol 2006;11:199–208.
- [9] Swain SM, Baselga J, Kim SB, et al. CLEOPATRA Study Group-Pertuzumab, trastuzumab, and docetaxel in HER2-positive metastatic breast cancer. N Engl J Med 2015;372:724–34.
- [10] Richard S, Selle F, Lotz JP, et al. Pertuzumab and trastuzumab: the rationale way to synergy. An Acad Bras Cienc 2016;88(Suppl 1):565–77.
- [11] Hierro C, Alsina M, Sánchez M, et al. Targeting the fibroblast growth factor receptor 2 in gastric cancer: promise or pitfall? Ann Oncol 2017;28:1207–16.
- [12] Meric-Bernstam F, Johnson AM, Dumbrava EEI, et al. Advances in HER2-targeted therapy: novel agents and opportunities beyond breast and gastric cancer. Clin Cancer Res 2018;25:2033–41.
- [13] Saito M, Shiraishi K, Kunitoh H, et al. Gene aberrations for precision medicine against lung adenocarcinoma. Cancer Sci 2016;107:713–20.
- [14] Salem ME, Weinberg BA, Xiu J, et al. Comparative molecular analyses of left-sided colon, right-sided colon, and rectal cancers. Oncotarget 2017;8:86356–68.

- [15] Kiss B, Wyatt AW, Douglas J, et al. Her2 alterations in muscle-invasive bladder cancer: patient selection beyond protein expression for targeted therapy. Sci Rep 2017;7:42713.
- [16] Galdy S, Lamarca A, McNamara MG, et al. HER2/HER3 pathway in biliary tract malignancies; systematic review and meta-analysis: a potential therapeutic target? Cancer Metastasis Rev 2017;36:141–57.
- [17] Aumayr K, Klatte T, Neudert B, et al. HER2 and TOP2A gene amplification and protein expression in upper tract urothelial carcinomas. Pathol Oncol Res 2018;24:575–81.
- [18] Teplinsky E, Muggia F. Targeting HER2 in ovarian and uterine cancers: challenges and future directions. Gynecol Oncol 2014;135:364–70.
- [19] Birkeland AC, Yanik M, Tillman BN, et al. Identification of targetable ERBB2 aberrations in head and neck squamous cell carcinoma. JAMA Otolaryngol Head Neck Surg 2016;142:559–67.
- [20] Hainsworth JD, Meric-Bernstam F, Swanton C, et al. Targeted therapy for advanced solid tumors on the basis of molecular profiles: results from MyPathway, an open-label, phase IIa multiple basket study. J Clin Oncol 2018;36:536–42.
- [21] Chugai Pharmaceutical Co., Ltd. Package insert for Herceptin (trastuzumab) (in Japanese). Available at: https://chugai-pharm.jp/hc/ ss/pr/drug/her_via0060_01/shiori/PDF/en/her_s_en.pdf [access March 13, 2019].
- [22] Chugai Pharmaceutical Co., Ltd. Package insert for Perjeta I.V. Infusion 420 mg/14 mL (Pertuzumab) (in Japanese). Available at: https://chugaipharm.jp/hc/ss/pr/drug/per_via0420/shiori/PDF/en/per_s_en.pdf [access March 13, 2019].
- [23] Tsimberidou AM, Wen S, Hong DS, et al. Personalized medicine for patients with advanced cancer in the phase I program at MD Anderson: validation and landmark analyses. Clin Cancer Res 2014;20:4827–36.
- [24] Quartino AL, Li H, Jin JY, et al. Pharmacokinetic and exposure-response analyses of pertuzumab in combination with trastuzumab and docetaxel during neoadjuvant treatment of HER2+ early breast cancer. Cancer Chemother Pharmacol 2017;79:353–61.
- [25] Schwaederle M, Zhao M, Lee JJ, et al. Impact of precision medicine in diverse cancers: a meta-analysis of phase II clinical trials. J Clin Oncol 2015;33:3817–25.
- [26] Schwaederle M, Zhao M, Lee JJ, et al. Association of biomarker-based treatment strategies with response rates and progression-free survival in refractory malignant neoplasms: a meta-analysis. JAMA Oncol 2016;2:1452–9.
- [27] Goldberg KB, Blumenthal GM, McKee AE, et al. The FDA oncology center of excellence and precision medicine. Exp Biol Med (Maywood) 2018;243:308–12.
- [28] Bashashati A, Ha G, Tone A, et al. Distinct evolutionary trajectories of primary high-grade serous ovarian cancers revealed through spatial mutational profiling. J Pathol 2013;231:21–34.
- [29] Palmirotta R, Lovero D, Cafforio P, et al. Liquid biopsy of cancer: a multimodal diagnostic tool in clinical oncology. Ther Adv Med Oncol 2018;10:1758835918794630.