Dosing *de novo* combinations of two targeted drugs: Towards a customized precision medicine approach to advanced cancers

Sariah Liu^{1,*}, Mina Nikanjam^{2,*}, Razelle Kurzrock³

¹Department of Hematology-Oncology, Kaiser Permanente San Diego Medical Center, San Diego, CA, USA

²Division of Hematology-Oncology, University of California Los Angeles, Los Angeles, CA, USA

³Center for Personalized Cancer Therapy and Division of Hematology and Oncology, UC San Diego Moores Cancer Center, San Diego, CA, USA

^{*}These authors have contributed equally to this work

Correspondence to: Mina Nikanjam, e-mail: mnikanjam@mednet.ucla.edu

Keywords: oncology, targeted therapy, maximum tolerated dose, recommended phase 2 dose, precision medicine *Received:* October 05, 2015 *Accepted:* January 15, 2016 *Published:* January 25, 2016

ABSTRACT

Metastatic cancers harbor complex genomic alterations. Thus, monotherapies are often suboptimal. Individualized combinations are needed in order to attenuate resistance. To help inform selection of safe starting doses for novel, two-agent, targeted drug combinations, we identified clinical trials in adult oncology patients who received targeted drug doublets (PubMed, January 1, 2010 through December 31, 2013). The dose percentage was calculated for each drug: (safe dose in combination divided by single agent full dose) X 100. Additive dose percentage represented the sum of the dose percentage for each drug. A total of 144 studies (N = 8568 patients; 95 combinations) were analyzed. In 51% of trials, each of the two drugs could be administered at 100% of their full dose. The lowest safe additive dose percentage was 60% if targets and/or class of drugs overlapped, or in the presence of mTor inhibitors, which sometimes compromised the combination dose. If neither class nor target overlapped and if mTor inhibitors were absent, the lowest safe additive dose percentage was 143%. The current observations contribute to the knowledge base that informs safe starting doses for new combinations of targeted drugs in the context of clinical trials or practice, hence facilitating customized combination therapies.

INTRODUCTION

A rapidly growing body of knowledge in cancer genomics has unveiled a complicated and heterogeneous molecular landscape in metastatic cancers. Indeed, it has been recently reported that patients with advanced tumors interrogated by next generation sequencing often have unique and complex genomic profiles [1]. For instance, in 57 patients with metastatic breast cancer, 216 somatic aberrations in 70 different genes, including 131 distinct aberrations were observed. Furthermore, no two patients shared the same molecular portfolio [2]. Molecular heterogeneity exists between histologies as well as within the same diagnostic group, and even within individual patients [3]. This diversified genomic landscape speaks to the need for customized combination treatments based on the genetic signature associated with each tumor [4, 5].

Combination therapies with targeted agents are frequently adopted to overcome resistance and maximize efficacy. This is of particular importance given that patients with advanced cancer frequently carry multiple genomic aberrations simultaneously. In a patientcentric approach, combined therapies would be highly individualized, which poses challenges as to how to ensure safe delivery of *de novo* combined therapies. Phase I oncology trials are traditionally designed to address concerns about drug safety. However, with at least 300 anti-cancer drugs approved or in advanced clinical trials, there are about 45,000 two-drug combinations and approximately 4,500,000 three-drug combinations, with even higher numbers of combinations if all permutations of drug dosing are considered. Testing each combination therefore poses a herculean challenge. Furthermore, the most reasonable starting doses for clinical trials with two targeted agents remains unclear.

Outside the cancer field, drugs are combined routinely and safely, based on established algorithms, for patients with multiple comorbidities. Indeed, the average patient suffering from cancer is often on many therapeutic agents, often designated "polypharmacy," for conditions as diverse as depression, heart disease, pain, and other illnesses. The safety of these drugs in combination has rarely if ever been formally tested in phase I studies. Yet physicians routinely prescribe a median of eight medications for patients with cancer, based on an understanding of drug-drug interactions and other factors [6]. Therefore, the prohibition against de novo combinations of drugs, and the demand for formal phase I testing of new combinations, often with slow and conservative dose escalation schemes, seems to be unique to the oncology sphere, and is perhaps a legacy from the era of cytotoxic drugs, which are toxic and have narrow therapeutic windows, especially compared to targeted agents that are generally better tolerated [7]. Importantly, within the context of oncology clinical trials, there is often considerable uncertainty as to what the initial dose levels should be, in the quest to balance safety, efficacy, and efficiency. In order to explore the correlation between dosing and toxicity for de novo combinations of targeted

agents, we conducted an analysis of previously published clinical trials. The goal of this study was to use literature review to establish a process that would help determine safe initial dosing for novel combinations of two-drug combinations of targeted agents, in order to inform both clinical trials and practice.

RESULTS

During the four-year period of publications evaluated, the total number of trials of two targeted agents that met the inclusion criteria was 144 (8568 patients; 95 drug combinations) (**Supplemental Table 1**, **Figure 1, Figure 2, Table 1**). A dose percentage was calculated to compare the dose of drug used in each combination to the single agent recommended dose (prioritizing the Food and Drug Administration-(FDA-) approved dose or, if not FDA approved, the recommended phase 2 dose (RP2D) or maximum tolerated dose (MTD), respectively) and the sum of the dose percentages for the combination was referred to as the additive dose percentage (see Methods, paragraph on Calculating "dose percentage").



Figure 1: Consort Diagram. Articles were identified by PubMed search and screened to identify two targeted drug combinations excluding studies of immunotherapy, hormonal therapy, radiation, or special populations (organ dysfunction, pediatric, or elderly patients).

First drug at 100% dose percentage of the FDA-approved dose/RP2D/MTD

Between January 1, 2010 and December 31, 2013, 121 trials (including 78 drug combinations) were published (N = 7748 patients) where both agents were targeted and at least one was given at full (100%) dose [8–18] (Supplemental references 1 to 110) (Table 2). These included 47 phase I trials (N = 1449 patients), 63 Phase II or III trials (N = 5742 patients), and 11 phase I/II combined trials (N = 557 patients). (In the phase I trials, only a subset of patients were treated at the RP2D/MTD)

The median dose percentage for the second agent was at 100% of the FDA-approved dose/RP2D/MTD (range, 25% to 100%). The median (range) for the additive dose percentage was 200% (125% to 200%) of the additive FDA-approved dose/RP2D/MTD. The lowest safe additive dose percentage was 125%; this lower dose was needed in some studies where both drugs were of the same class, i.e., small molecule inhibitors.

In total, 75 trials (51% of the 144 trials of the trials) (N = 48 drug combinations) administered each targeted agent at 100% dose percentage (N = 5229 patients received each drug at 100% dose percentage). These trials (wherein additive dose percentage was 200%) included the following types of combinations: 10 combinations that involved an antibody and small molecule tyrosine kinase inhibitor (TKI); 7 combinations, two antibodies; 17 combinations, antibody and small molecule non-TKI; 6 combinations, two small molecule TKIs; 6 combinations,

a small molecule non-TKI and small molecule TKI; and 2 combinations, involved two small molecule non-TKIs. In 3 of the 48 combinations where each drug was given at 100%, dose percentage, the target of the two molecules overlapped. These included the following: gefitinib and nimotuzumab (both targeting epidermal growth factor receptor (EGFR)) [9]; bevacizumab and ABT-510 (targeting vascular endothelial growth factor (VEGF)) [8]; and trastuzumab and pertuzumab (targeting HER2) [10, 11].

Subset analyses with two antibodies, two small molecule inhibitors, and an antibody with a small molecule inhibitor were performed. For all 9 studies of two antibodies given in combination, each drug was given at 100% of the FDA-approved dose/RP2D/MTD dose. When two small molecules were administered, each drug could be administered at 100% of the dose in 25 of the 68 total trials (37%) (Table 3). When a small molecule and antibody were administered in combination, each drug could be administered at 100% in 40 out of 55 total trials (73%) (Table 3). Limitations of this analysis are due to the small number of antibody-antibody combinations (N = 9).

First drug at >50% but < 100% dose percentage of the FDA-approved dose/RP2D/MTD

There were 13 trials (N = 13 drug combinations) where the first drug was administered at >50% but < 100% of the FDA-approved dose/RP2D/MTD due to toxicity of higher doses [19–21] (Supplemental references 111-120).





Table 1: Two targeted drug combinations reported over four years (Phase I, II, III studies on PubMed January 1,2010 to December 21, 2013)

	Two targeted agents ^a
Number of trials	144
Number of drug combinations	95
Number of patients	8568
Median (range) additive dose percentage	200% (60%-200%)
Number (percent) of trials where \geq one drug dose percentage was 100%	121 (84%)
Number of drug combinations where \geq one drug dose percentage was 100%	78
Number of patients where \geq one drug dose percentage was 100%	7748
Median (range) percentile for second drug when one drug dose percentage was 100%	100% (25%-100%)
Number (percent) of trials where each drug's dose percentage was 100% (e.g. additive dose percentage ^b = 200%)	74 (51%)
Number of drug combinations where each drug's dose percentage was 100% (e.g. additive dose percentage = 200%)	48
Number of patients where each drug's dose percentage was 100% (e.g. additive dose percentage = 200%)	5229
Number (percent) of trials where additive dose percentage was $>100\%$, with no single drug given at 100%	13 (9%)
In trials where additive dose percentage was $>100\%$, with no single drug given at 100%, median (range) of additive dose percentage	127% (104%-148%)
Number (percent) of trials where additive dose percentage was $\leq 100\%$ and safe dose was found	5 (4%)
In trials where additive dose percentage was $\leq 100\%$ and safe dose was found, median (range) of additive dose percentage	75% (60%-100%)
Number (percent) of trials where additive dose percentage was $\leq 100\%$ and safe dose was not found ^c	3 (2%)
In trials where additive dose percentage was $\leq 100\%$ and safe dose was not found, median (range) of additive dose percentage studied	75% (65%-100%)
Number (percent) of trials that were aborted early with additive dose percentage $>100\%$	2 (1%)
In trials that were aborted early with additive dose percentage >100%, median (range) of additive dose percentage	200%

^aExcludes hormonal modulators and immunotherapy

^bAdditive dose percentage = (dose of drug A in combination/standard dose of drug A as a single agent) X 100 + (dose of drug B in combination/standard dose of drug B as a single agent) X 100.

^cSee Results for details. In the case of two trials in which safe dose could not be defined, other trials of the same combination did define a safe dose (additive dose percentage = 100% or 75%); one trial had fatigue as a dose-limiting toxicity, but no acute or irreversible toxicities.

The lowest safe additive dose of the combination was 104% and the latter was required for sunitinb (75% of dose) combined with everolimus (29% of dose) [20]. The lowest safe additive dose percentage was 143% [19] (rapamycin and bevacizumab) when the drugs did not overlap in either class or target (**Table 2**)

First drug at less than or equal to 50% dose percentage of the FDA-approved dose/RP2D/MTD

Five trials (bevacizumab and sorafenib [22]; sorafenib and temsirolimus [23]; sorafenib and sirolimus [24]; bevacizumab and telatinib (VEGFR inhibitor) [25];

Table 2: Summary of Two Targeted Drugs in Combination^a

	Second drug at 100% dose percentage of FDA-approved dose/ RP2D/MTD	Lowest safe dose percentage of second drug if both are of the same class and/or have overlapping targets	Lowest additive dose percentage of the combination
First drug at 100% dose percentage of FDA-approved dose/RP2D/MTD ^b	61% of trials (74/121) (Note: 74 of the 144 total trials (51%) administered each drug at 100% dose)	25% of FDA/RP2D/MTD ^c	125%
First drug at 51-99% dose percentage of the FDA- approved dose/RP2D/MTD	Not applicable (13 total trials)	29% of FDA/RP2D/MTD ^d	104% ^d 143% (for non-overlapping targets and different classes) ^e
First drug at ≤ 50% dose percentage of the FDA- approved dose/RP2D/MTD	Not applicable (5 total trials)	10% of FDA/RP2D/MTD ^f	60% (overlapping targets in each case) ^f 90% (for non-overlapping targets, but both same class (small molecule inhibitors) ^g

^aThe five studies where no safe dose was found or study was aborted early due to unacceptable toxicity were excluded from this table and include: bevacizumab and sorafenib [28, 29], pazopanib and temsirolimus [27], bevacizumab and everolimus [31], and bevacizumab and temsirolimus [32]

^bFirst drug had the dose percentage closest to the FDA-approved/RP2D/MTD dose

^cIn these cases, the combinations were same class (small molecule inhibitors) with non-overlapping targets (sorafenib at 100% with everolimus at 25%) and imatinib at 100% with everolimus at 25%) [12–14].

^dSunitinib was at 75% and everolimus at 29% [20]

^eRapamycin was at 93% and bevacizumab was at 50% [19]

^fBevacizumab with vatalinib [26] and bevacizumab with telatinib [25] each included an anti-VEGF antibody and a small molecule VEGFR inhibitor (both at 10% and 50%, respectively)

^gSorafenib was at 50% and temsirolimus at 40% [23]; however the combination of pazopanib and temsirolimus was above the FDA-approved/RP2D/MTD at an additive dose percentage of 65% (albeit with no acute or irreversible side effects and with the nonspecific side effect of fatigue as dose limiting in one patient).

Abbreviations: MTD = maximum tolerated dose; RP2D = recommended phase II dose

bevacizumab and vatalinib (VEGFR inhibitor) [26] were published, where both agents were targeted and the first drug was administered at $\leq 50\%$ dose percentage of the FDA-approved dose/RP2D/MTD due to toxicity of higher doses, thus the additive dose percentage was $\leq 100\%$ of the additive FDA-approved dose/RP2D/MTD. The dose percentage for bevacizumab and sorafenib was 50% and 25% of the RP2D of each drug, respectively; for sorafenib and temsirolimus, 50% and 40%, respectively; for sorafenib and sirolimus, 50% and 50%, respectively; for bevacizumab and telatinib, 10% and 50%, respectively; for bevacizumab and vatalinib, 10% and 50%, respectively (additive dose percentage = 75%, 90%, 100%, 60% and 60%). Of note, the lowest additive dose percentages (75%, 60% and 60%, respectively) applied to bevacizumab and sorafenib, bevacizumab and telatinib, and bevacizumab and vatalinib, which, in each case, overlap in their targeting angiogenesis. In addition, the combination of pazopanib and temsirolimus was given at 65% additive dose percentage and was considered above the MTD, but the toxicity was fatigue, which is often hard to quantify [27] **(Table 2).**

mTor inhibitor-based combinations

Combinations with mTor inhibitors such as everolimus or temsirolimus often required compromised doses: (i) the combination of sorafenib (100% of dose) with everolimus necessitated dosing the latter at 25% [12]; (ii) the everolimus dose was reduced to 25% in combination with imatinib, when used at 100% [13, 14]; (iii) when sunitinib (75% of dose) was combined with everolimus, only 29% of the dose of the latter could be given [20]; and (iv) as mentioned above, the dose of pazopanib (25%) and temsirolimus (40%) resulted in dose limiting toxicity of fatigue [27].

Table 3: Summary of Subset Analysis for Combination of Two Small Molecule Inhibitors, as well as Small Molecule Inhibitor and Antibody Combinations^a

Combination of Two Small Molecule Inhibitors (N = 68 Studies)				
	Second drug at 100% dose percentage of FDA-approved dose/RP2D/MTD	Lowest additive dose percentage of the combination		
First drug at 100% dose percentage of FDA-approved dose/RP2D/MTD	44% of trials (25/57) (Note: 25 of the 68 total trials (37%) administered each drug at 100% dose)	125% ^b		
First drug at 51-99% dose percentage of the FDA-approved dose/RP2D/MTD	Not applicable (9 total trials)	104% ^c		
First drug at \leq 50% dose percentage of the FDA-approved dose/RP2D/MTD	Not applicable (2 total trials)	90% ^d		
Combination of Small Molecule Inhibitor and Antibody Combinations (N = 62 Studies)				
First drug at 100% dose percentage of FDA-approved dose/RP2D/MTD	73% of trials (40/55) (Note: 40 of the 62 total trials (65%) administered each drug at 100% dose)	150%e		
First drug at 51-99% dose percentage of the FDA-approved dose/RP2D/MTD	Not applicable (4 total trials)	117% ^f 143% (for non-overlapping targets) ^g		
First drug at \leq 50% dose percentage of the FDA-approved dose/RP2D/MTD	Not applicable (3 total trials)	60% (overlapping targets in each case) ^h		

^aAll combinations of two antibodies had each drug given at 100% of the FDA-approved/RP2D/MTD dose

^bIn these cases, the combinations had non-overlapping targets (sorafenib at 100% with everolimus at 25%), and imatinib at 100% with everolimus at 25%) [12–14].

^cSunitinib was at 75% and everolimus at 29% [20]

^dSorafenib was at 50% and temsirolimus at 40% [23].

^eBevacizumab and erlotinib were each given at 50% and 100%, respectively [15–17] while panobinostat and bevacizumab were given at 50% and 100%, respectively [18].

^fVandetanib was at 67% and bevacizumab was at 50% [21].

^gRapamycin was at 93% and bevacizumab was at 50% [19].

^hBevacizumab with vatalinib [26] and bevacizumab with telatinib [25] each included an anti-VEGF antibody and a small molecule VEGFR inhibitor (both at 10% and 50%, respectively).

Two targeted agents where the additive dose percentage was $\leq 100\%$ and safety was unacceptable

Three trials (bevacizumab and sorafenib (2 trials giving 50% and 50%; 50% and 25%) [28, 29]; one trial, pazopanib and temsirolimus (25% and 40% dose, respectively) [27]) were published where the lowest dose level did not have an acceptable safety profile. Of interest, for the bevacizumab and sorafenib combination, other studies have found acceptable safety at 50% and 25% of the dose [22] or for 50% and 50% of the dose [30]. Therefore the only combination with an undefined safety dosing in this group (and no alternative studies demonstrating safety) was the study of pazopanib and

temsirolimus mentioned above [27]. As mentioned, doselimiting toxicity in this study included fatigue, and did not include acute irreversible events.

Two targeted agents where the study was aborted early or safety defined as unacceptable and the additive dose percentages investigated were > 100%

There were two trials published where the additive dose percentage was $\geq 100\%$ and the studies did not find an acceptable dose: bevacizumab and everolimus (both drugs at 100%) [31]; bevacizumab and temsirolimus (both at 100% of each) [32]. These trials did not attempt to lower the dose.

DISCUSSION

Targeted agents matched to advanced tumors bearing cognate alterations are often given as monotherapy. While significant salutary effects can be achieved [5, 33, 34] responses generally last only a few months. This is perhaps not surprising since metastatic malignancies mostly harbor multiple genomic alterations [35–38], strongly suggesting that individualized combination treatment will be need to be deployed to further improve outcomes.

When two targeted agents are combined, safety considerations may include whether both belong to the same class of drugs (e.g., both are small molecule inhibitors) or if there are overlapping targets (e.g. both target angiogenesis) [7]. In the current study, we have reviewed phase I-III clinical trials of targeted therapeutics over a four-year span (N = 8568 patients) to determine safe starting doses for novel two-drug combinations of targeted agents.

In over half of the studies, both drugs could be given at 100% of the individually defined optimum dose (i.e., the FDA-approved dose or RP2D or MTD) (Table 2). When giving full doses was not possible, studies were able to define safe starting doses by lowering the additive dose percentage of the combination. When one drug was given at 100% of the full dose, the lowest safe dose for the second drug was 25%. Lowering the dose was needed in the presence of two drugs of the same class when both were small molecule inhibitors or two drugs with overlapping targets. In some cases, neither drug was administered at 100% of the full dose. The lowest additive dose percentage was 60% and was relevant to bevacizumab and telatinib, and bevacizumab and vatalinib, which overlapped in their target of angiogenesis [25, 26]. However, other studies were able to administer 100% of each agent (200% additive dose percentage) despite overlapping targets: gefitinib and nimotuzumab (targeting EGFR) [9]; bevacizumab and ABT-510 (targeting VEGF) [8]; and trastuzumab and pertuzumab (targeting HER2) [10, 11]. Thus, the presence of overlapping targets needs to be considered for starting doses, but, in many cases, will not limit the ability to administer full doses of agents. Combinations that included mTor inhibitors such as everolimus or temsirolimus also resulted in compromised doses. At times, these combinations could not be given at more than 65 to 100% of the additive dose percentage. Finally, the lowest safe additive dose percentage for drugs with non-overlapping targets or class was 143%.

In implementing novel drug combinations tailored to the genomic aberration of each individual cancer, considerations include efficacy of the combination and toxicity. The effect of administration of less than 100% of the single agent MTD for combination therapies was addressed in two separate studies of phase I data. Jain et al demonstrated, in a single institution study of 24 clinical trials, that patients who received lower drug doses did not fare worse than those on higher doses, and suggested that targeted agents may have different dose response relationships than cytotoxic chemotherapies [39]. A separate study of 55 clinical trials sponsored by a single entity with multiple sites suggested that patients on higher doses had better response rates and overall survival [40]. Thus, it is unclear if dose reductions to allow for the administration of multiple agents will alter efficacy.

This study has several limitations. First, the publications reviewed for the current analysis were limited to two-drug combinations (targeted agents) in adult patients without organ dysfunction. The results are likely not applicable to patients with renal or hepatic impairment, or children, who may require dosing modifications depending on metabolism of the therapeutic or maturation, respectively, and were excluded from the current analysis as often occurs in clinical trials. In addition, immunotherapies, hormonal modulators, and cytotoxics were not included, which may alter the additive dose percentages seen in the study as hormonal modulators and immunotherapies may be better tolerated in combination therapy while cytotoxics may increase toxicity. It is also plausible that some trials with two targeted agents that showed significant toxicity were never published, and so the data presented herein, while derived from a large number of patients, does not guarantee safety with all possible drug combinations. Drug-drug interactions and effects on metabolic proteins such as CYP enzymes, which can lead to changes in the pharmacokinetic profile of therapeutics, may have resulted in lower safe dose levels for combination therapy as compared to single agent dosing. While this was not addressed in the study, we still observed that, in 51% of trials, both drugs could still be administered at 100% of their FDA-approved dose/RP2D/ MTD. The study also did not address target engagement for therapy where optimal doses may be lower than the FDA-approved dose/RP2D/MTD. Still it is of interest that for some drugs such as everolimus, administering 5 mg (which is 50% of the approved dose) offers 100% target engagement [41]. Another limitation of the study was that it included different types of trials (phase I, phase II, phase III trials) with different primary objectives. Finally, there are myriad of possible dosing schemes and some investigators or practitioners may want to hold one of the drugs at a preconceived dosing level. While our study did define that, if one drug is held at 100%, the lowest safe additive dosing percentage was 125%, it was not possible to define all permutations because they do not exist in the literature reviewed.

In conclusion, classical cytotoxic chemotherapy dosing was previously limited by significant toxicity, and thus the administration of two or more drugs in combination often necessitated very conservative initial dosing, even within the controlled environment of a clinical trial. Although targeted therapies can have fewer side effects than traditional cytotoxic chemotherapy, a process to calculate initial safe doses, either within a clinical trial or in practice, for combinations of two targeted agents, remains a matter of debate. The molecular heterogeneity of cancer indicates that prosecuting malignancies with an optimized personalized/ precision medicine strategy will require combination therapy matched to individual molecular profiles. Yet, with an increasing number of targeted agents deployed in the clinic, there are thousands of drug combinations possible, and there is an increasingly urgent need for more knowledge that can inform safely combining them. Outside of oncology, for patients with multiple comorbidities, drugs are routinely given together based on established algorithms; indeed, the average cancer patient is on 5 to 10 drugs for their other health problems, often before starting treatment for their malignancy. The current study documented the following in adults with intact organ function treated with two targeted agents: (i) compromised dosing most often was needed for overlap of drug class (e.g., two small molecule inhibitors (but not two antibodies)) and/or targets (especially angiogenesis) or in the presence of mTor inhibitors; (ii) without overlap of class or target and in the absence of mTor inhibitors, the lowest safe additive dose percentage was 143%; (iii) in the presence of overlapping class and/or targets and/or mTor inhibitors, the lowest safe additive dose percentage was about 60%; and (iv) dose escalation to full dose was possible with most two targeted drug combinations, since over half of these combinations were administered safely at 100% dose percentage of the FDA-approved/ RP2D/MTD of each drug (additive dose percentage = 200%). Therefore starting doses of two targeted drugs in combination in a clinical trial or practice could be about 70% of each drug if there is no overlap in targets or class and no mTor inhibitor in the regimen, and about 30% of each drug with overlap of class and/or target or inclusion of an mTor inhibitor. If one drug was held at 100% of full dose, the lowest safe starting dose of the second drug was 25%, and this was required in the case of overlapping drug class/target. Since over half of the combinations could be given with both drugs at full dose, in the absence of significant toxicity, intra-patient dose escalation can occur, to allow for improved efficacy if needed. Of course, further adjustments may be needed, depending on co-morbidities, patient age, organ function, other concomitant medications, and consideration of absorption, distribution, metabolism and excretion (ADME) of individual drugs. Despite these limitations, our current observations can help inform the safe starting dose of de novo two targeted agent combinations, both in clinical trials and practice, as a step toward customization of therapy to the complex molecular landscape seen in patients with cancer.

MATERIALS AND METHODS

To identify research articles for the analysis, we first conducted a search of PubMed for studies published between January 1, 2010 and December 31, 2013, using the search terms "cancer, phase, combination." We then manually screened the resulting articles and included studies that meet the inclusion criteria: (i) phase I-III clinical trials; (ii) solid tumors or hematology malignancy; and (iii) two-drug combination therapy where both drugs were targeted agents. Targeted agents are generally cytostatic and broadly include antibodies that have a specific protein as their target or small molecule inhibitors with low nM IC50s (concentration that results in 50% inhibition of enzyme function) for the specific protein target. Exclusion criteria were as follows: (i) the dose of any drug in the combination was greater than 100% of the standard dose as a single agent; (ii) the dose of any drug in the combination was chosen to be low due to reasons other than toxicity, such as optimized activity at lower doses due to different biological impact; (iii) the study was performed on selected patient populations such as pediatric, elderly, or patients with organ dysfunction; and (iv) the study treatment regimen included radiation. Hormonal agents and immonotherapeutics were excluded.

Data

Clinical data were manually extracted from each clinical trial. Data included drug names, targets of action (the target of small molecule inhibitors was felt to be relevant if it was impacted at an IC50 <250 nM [42]), drug type, status of Food and Drug Administration (FDA) approval, number of drugs in the combination, disease, number of participants, dose of each drug in the combination, recommended phase 2 dose (RP2D) or maximal tolerated dose (MTD) achieved in the study, dose limiting toxicities (DLTs), and grade \geq 3 toxicities.

Calculating "dose percentage"

Based on the extracted data, a "dose percentage" was calculated, which was defined as the dose of one drug in the combination, divided by the standard dose of the same drug used as a single agent, multiplied by 100 ((dose percentage = dose of drug in the combination/ standard dose of drug as a single agent) X 100). Single agent dose was defined as the FDA approved dose, or for drugs that were not FDA approved, the RP2D or MTD dose from phase I studies. FDA approved dose was always prioritized as the reference full dose and RP2D was prioritized over MTD. For drugs where the standard single agent dose as the accepted dose. "Additive dose

percentage" for combinations of two targeted agents was calculated by adding the dose percentage of each drug in a given combination. Hence, in combination therapy, if the maximum safe dose percentage of drug A was 50% of the FDA-approved dose/RP2D/MTD of drug A as a single agent, and the maximum safe dose percentage of drug B was 25% of the FDA-approved dose/RP2D/MTD of drug B as a single agent, the "additive dose percentage" of the combination was 75%. The maximum "additive dose percentage" for any two-drug combination is 200% (i.e., 100% of each drug). The "first drug" was defined as the drug with the highest dose percentage of the combination (i.e., the drug with the dose that was closest to the FDA-approved dose/RP2D/MTD).

ACKNOWLEDGMENTS

This study was funded in part by the Joan and Irwin Jacobs philanthropic fund.

CONFLICTS OF INTEREST

Dr. Kurzrock receives consultant fees from Sequenom Inc. as well as research funds from Genentech, Pfizer, Foundation Medicine, Guardant, Sequenom and Merck Serono, and has an ownership interest in RScueRx. Dr. Liu and Dr. Nikanjam do not have any conflicts of interest to report.

FUNDING

This study was funded in part by the Joan and Irwin Jacobs philanthropic fund (Dr. Kurzrock).

REFERENCES

- 1. Kurzrock R, Giles FJ. Precision oncology for patients with advanced cancer: the challenges of malignant snowflakes. Cell Cycle. 2015; 14: 2219–2221.
- Wheler JJ, Parker BA, Lee JJ, Atkins JT, Janku F, Tsimberidou AM, Zinner R, Subbiah V, Fu S, Schwab R, Moulder S, Valero V, Schwaederle M, et al. Unique molecular signatures as a hallmark of patients with metastatic breast cancer: implications for current treatment paradigms. Oncotarget. 2014; 5: 2349–2354. doi: 10.18632/oncotarget.1946.
- Gerlinger M, Rowan AJ, Horswell S, Larkin J, Endesfelder D, Gronroos E, Martinez P, Matthews N, Stewart A, Tarpey P, Varela I, Phillimore B, Begum S, et al. Intratumor heterogeneity and branched evolution revealed by multiregion sequencing. N Engl J Med. 2012; 366: 883–892.
- 4. Wheler J, Lee JJ, Kurzrock R. Unique molecular landscapes in cancer: implications for individualized, curated drug combinations. Cancer Res. 2014; 74: 7181–7184.
- 5. Tsimberidou AM, Iskander NG, Hong DS, Wheler JJ, Falchook GS, Fu S, Piha-Paul S, Naing A, Janku F, Luthra

R, Ye Y, Wen S, Berry D, et al. Personalized medicine in a phase I clinical trials program: the MD Anderson Cancer Center initiative. Clin Cancer Res. 2012; 18: 6373–6383.

- Borad MJ, Curtis KK, Babiker HM, Benjamin M, Tibes R, Ramanathan RK, Wright K, Dueck AC, Jameson G, Von Hoff DD. The impact of concomitant medication use on patient eligibility for phase I cancer clinical trials. J Cancer. 2012; 3: 345–353.
- Liu S and Kurzrock R. Toxicity of targeted therapy: Implications for response and impact of genetic polymorphisms. Cancer Treat Rev. 2014; 40: 883–891.
- Uronis HE, Cushman SM, Bendell JC, Blobe GC, Morse MA, Nixon AB, Dellinger A, Starr MD, Li H, Meadows K, Gockerman J, Pang H, Hurwitz HI. A phase I study of ABT-510 plus bevacizumab in advanced solid tumors. Cancer Med. 2013; 2: 316–324.
- 9. Kim SH, Shim HS, Cho J, Jeong JH, Kim SM, Hong YK, Sung JH, Ha SJ, Kim HR, Chang H, Kim JH, Tania C, Cho BC. A phase I trial of gefitinib and nimotuzumab in patients with advanced non-small cell lung cancer (NSCLC). Lung Cancer. 2013; 79: 270–275.
- Gianni L, Pienkowski T, Im YH, Roman L, Tseng LM, Liu MC, Lluch A, Staroslawska E, de la Haba-Rodriguez J, Im SA, Pedrini JL, Poirier B, Morandi P, et al. Efficacy and safety of neoadjuvant pertuzumab and trastuzumab in women with locally advanced, inflammatory, or early HER2-positive breast cancer (NeoSphere): a randomised multicentre, open-label, phase 2 trial. Lancet Oncol. 2012; 13: 25–32.
- Baselga J, Gelmon KA, Verma S, Wardley A, Conte P, Miles D, Bianchi G, Cortes J, McNally VA, Ross GA, Fumoleau P, Gianni L. Phase II trial of pertuzumab and trastuzumab in patients with human epidermal growth factor receptor 2-positive metastatic breast cancer that progressed during prior trastuzumab therapy. J Clin Oncol. 2010; 28: 1138–1144.
- Finn RS, Poon RT, Yau T, Klumpen HJ, Chen LT, Kang YK, Kim TY, Gomez-Martin C, Rodriguez-Lope C, Kunz T, Paquet T, Brandt U, Sellami D, et al. Phase I study investigating everolimus combined with sorafenib in patients with advanced hepatocellular carcinoma. J Hepatol. 2013; 59: 1271–1277.
- Ryan CW, Vuky J, Chan JS, Chen Z, Beer TM, Nauman D. A phase II study of everolimus in combination with imatinib for previously treated advanced renal carcinoma. Invest New Drugs. 2011; 29: 374–379.
- Schoffski P, Reichardt P, Blay JY, Dumez H, Morgan JA, Ray-Coquard I, Hollaender N, Jappe A, Demetri GD. A phase I-II study of everolimus (RAD001) in combination with imatinib in patients with imatinib-resistant gastrointestinal stromal tumors. Ann Oncol. 2010; 21: 1990–1998.
- 15. Lubner SJ, Mahoney MR, Kolesar JL, Loconte NK, Kim GP, Pitot HC, Philip PA, Picus J, Yong WP, Horvath L, Van Hazel G, Erlichman CE, Holen KD. Report of a multicenter phase II trial testing a combination of biweekly

bevacizumab and daily erlotinib in patients with unresectable biliary cancer: a phase II Consortium study. J Clin Oncol. 2010; 28: 3491–3497.

- Hsu CH, Kang YK, Yang TS, Shun CT, Shao YY, Su WC, Sandoval-Tan J, Chiou TJ, Jin K, Hsu C, Cheng AL. Bevacizumab with erlotinib as first-line therapy in Asian patients with advanced hepatocellular carcinoma: a multicenter phase II study. Oncology. 2013; 85: 44–52.
- 17. Johnsson A, Hagman H, Frodin JE, Berglund A, Keldsen N, Fernebro E, Sundberg J, De Pont Christensen R, Garm Spindler KL, Bergstrom D, Jakobsen A. A randomized phase III trial on maintenance treatment with bevacizumab alone or in combination with erlotinib after chemotherapy and bevacizumab in metastatic colorectal cancer: the Nordic ACT Trial. Ann Oncol. 2013; 24: 2335–2341.
- Drappatz J, Lee EQ, Hammond S, Grimm SA, Norden AD, Beroukhim R, Gerard M, Schiff D, Chi AS, Batchelor TT, Doherty LM, Ciampa AS, Lafrankie DC, et al. Phase I study of panobinostat in combination with bevacizumab for recurrent high-grade glioma. J Neurooncol. 2012; 107: 133–138.
- Choo SP, Chowbay B, Ng QS, Thng CH, Lim C, Hartono S, Koh TS, Huynh H, Poon D, Ang MK, Chang S, Toh HC. A Phase 1 dose-finding and pharmacodynamic study of rapamycin in combination with bevacizumab in patients with unresectable hepatocellular carcinoma. Eur J Cancer. 2013; 49: 999–1008.
- Molina AM, Feldman DR, Voss MH, Ginsberg MS, Baum MS, Brocks DR, Fischer PM, Trinos MJ, Patil S, Motzer RJ. Phase 1 trial of everolimus plus sunitinib in patients with metastatic renal cell carcinoma. Cancer. 2012; 118: 1868–1876.
- 21. Kummar S, Gutierrez ME, Chen A, Turkbey IB, Allen D, Horneffer YR, Juwara L, Cao L, Yu Y, Kim YS, Trepel J, Chen H, Choyke P, et al. Phase I trial of vandetanib and bevacizumab evaluating the VEGF and EGF signal transduction pathways in adults with solid tumours and lymphomas. Eur J Cancer. 2011; 47: 997–1005.
- 22. Galanis E, Anderson SK, Lafky JM, Uhm JH, Giannini C, Kumar SK, Kimlinger TK, Northfelt DW, Flynn PJ, Jaeckle KA, Kaufmann TJ, Buckner JC. Phase II study of bevacizumab in combination with sorafenib in recurrent glioblastoma (N0776): a north central cancer treatment group trial. Clin Cancer Res. 2013; 19: 4816–4823.
- Kelley RK, Nimeiri HS, Munster PN, Vergo MT, Huang Y, Li CM, Hwang J, Mulcahy MF, Yeh BM, Kuhn P, Luttgen MS, Grabowsky JA, Stucky-Marshall L, et al. Temsirolimus combined with sorafenib in hepatocellular carcinoma: a phase I dose-finding trial with pharmacokinetic and biomarker correlates. Ann Oncol. 2013; 24: 1900–1907.
- 24. Desar IM, Timmer-Bonte JN, Burger DM, van der Graaf WT, van Herpen CM. A phase I dose-escalation study to evaluate safety and tolerability of sorafenib combined with sirolimus in patients with advanced solid cancer. Br J Cancer. 2010; 103: 1637–1643.

- 25. Langenberg MH, Witteveen PO, Roodhart J, Lolkema MP, Verheul HM, Mergui-Roelvink M, Brendel E, Kratzschmar J, Loembe B, Nol-Boekel A, Christensen O, Schellens JH, Voest EE. Phase I evaluation of telatinib, a VEGF receptor tyrosine kinase inhibitor, in combination with bevacizumab in subjects with advanced solid tumors. Ann Oncol. 2011; 22: 2508–2515.
- Jones SF, Spigel DR, Yardley DA, Thompson DF, Burris HA, 3rd. A phase I trial of vatalanib (PTK/ZK) in combination with bevacizumab in patients with refractory and/or advanced malignancies. Clin Adv Hematol Oncol. 2011; 9: 845–852.
- Semrad TJ, Eddings C, Dutia MP, Christensen S, Lara PN, Jr. Phase I study of the combination of temsirolimus and pazopanib in advanced solid tumors. Anticancer Drugs. 2013; 24: 636–640.
- Castellano D, Capdevila J, Sastre J, Alonso V, Llanos M, Garcia-Carbonero R, Manzano Mozo JL, Sevilla I, Duran I, Salazar R. Sorafenib and bevacizumab combination targeted therapy in advanced neuroendocrine tumour: a phase II study of Spanish Neuroendocrine Tumour Group (GETNE0801). Eur J Cancer. 2013; 49: 3780–3787.
- Mina LA, Yu M, Johnson C, Burkhardt C, Miller KD, Zon R. A phase II study of combined VEGF inhibitor (bevacizumab+sorafenib) in patients with metastatic breast cancer: Hoosier Oncology Group Study BRE06-109. Invest New Drugs. 2013; 31: 1307–1310.
- 30. Falchook GS, Wheler JJ, Naing A, Piha-Paul SA, Fu S, Tsimberidou AM, Hong DS, Janku F, Zinner R, Jiang Y, Huang M, Lin Q, Parkhurst K, et al. Dual antiangiogenic inhibition: a phase I dose escalation and expansion trial targeting VEGF-A and VEGFR in patients with advanced solid tumors. Invest New Drugs. 2015; 33: 215–224.
- Harshman LC, Barbeau S, McMillian A, Srinivas S. A phase II study of bevacizumab and everolimus as treatment for refractory metastatic renal cell carcinoma. Clin Genitourin Cancer. 2013; 11: 100–106.
- 32. Alvarez EA, Brady WE, Walker JL, Rotmensch J, Zhou XC, Kendrick JE, Yamada SD, Schilder JM, Cohn DE, Harrison CR, Moore KN, Aghajanian C. Phase II trial of combination bevacizumab and temsirolimus in the treatment of recurrent or persistent endometrial carcinoma: a Gynecologic Oncology Group study. Gynecol Oncol. 2013; 129: 22–27.
- 33. Shaw AT, Kim DW, Nakagawa K, Seto T, Crino L, Ahn MJ, De Pas T, Besse B, Solomon BJ, Blackhall F, Wu YL, Thomas M, O'Byrne KJ, et al. Crizotinib versus chemotherapy in advanced ALK-positive lung cancer. N Engl J Med. 2013; 368: 2385–2394.
- 34. Sosman JA, Kim KB, Schuchter L, Gonzalez R, Pavlick AC, Weber JS, McArthur GA, Hutson TE, Moschos SJ, Flaherty KT, Hersey P, Kefford R, Lawrence D, et al. Survival in BRAF V600-mutant advanced melanoma treated with vemurafenib. N Engl J Med. 2012; 366: 707–714.

- 35. Bean J, Brennan C, Shih JY, Riely G, Viale A, Wang L, Chitale D, Motoi N, Szoke J, Broderick S, Balak M, Chang WC, Yu CJ, et al. MET amplification occurs with or without T790M mutations in EGFR mutant lung tumors with acquired resistance to gefitinib or erlotinib. Proc Natl Acad Sci U S A. 2007; 104: 20932–20937.
- 36. Flaherty KT, Infante JR, Daud A, Gonzalez R, Kefford RF, Sosman J, Hamid O, Schuchter L, Cebon J, Ibrahim N, Kudchadkar R, Burris HA, 3rd, Falchook G, et al. Combined BRAF and MEK inhibition in melanoma with BRAF V600 mutations. N Engl J Med. 2012; 367: 1694–1703.
- 37. Pao W, Miller VA, Politi KA, Riely GJ, Somwar R, Zakowski MF, Kris MG, Varmus H. Acquired resistance of lung adenocarcinomas to gefitinib or erlotinib is associated with a second mutation in the EGFR kinase domain. PLoS Med. 2005; 2: e73.
- Yap TA, Omlin A, de Bono JS. Development of therapeutic combinations targeting major cancer signaling pathways. J Clin Oncol. 2013; 31: 1592–1605.

- 39. Jain RK, Lee JJ, Hong D, Markman M, Gong J, Naing A, Wheler J, Kurzrock R. Phase I oncology studies: evidence that in the era of targeted therapies patients on lower doses do not fare worse. Clin Cancer Res. 2010; 16: 1289–1297.
- Gupta S, Hunsberger S, Boerner SA, Rubinstein L, Royds R, Ivy P, LoRusso P. Meta-analysis of the relationship between dose and benefit in phase I targeted agent trials. J Natl Cancer Inst. 2012; 104: 1860–1866.
- 41. Tabernero J, Rojo F, Calvo E, Burris H, Judson I, Hazell K, Martinelli E, Ramon y Cajal S, Jones S, Vidal L, Shand N, Macarulla T, Ramos FJ, et al. Dose- and schedule-dependent inhibition of the mammalian target of rapamycin pathway with everolimus: a phase I tumor pharmacodynamic study in patients with advanced solid tumors. J Clin Oncol. 2008; 26: 1603–1610.
- Selleckchem.com: Inhibitor Expert. Selleck Chemicals; 2015. http://www.selleckchem.com. Accessed February 24, 2015.