

Phase 2 Study of Single Agent Navitoclax (ABT-263) and Biomarker Correlates in Patients with Relapsed Small Cell Lung Cancer

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TRANSLATIONAL RELEVANCE

Bcl-2 is a key apoptotic resistance gene, and is upregulated in the majority of small cell lung cancers (SCLC). Navitoclax, or ABT-263, is an orally bioavailable small molecule with selective and potent inhibition of Bcl-2, resulting in single agent activity in preclinical SCLC models. This paper describes the first phase II evaluation of navitoclax in patients with recurrent and metastatic SCLC patients. Navitoclax administration at this dose and schedule was feasible, but the observed response rate in this context was low. This result is in contrast to results using SCLC cell line xenograft models, but consistent with prior observations in primary tumor xenograft SCLC models. Finally, the study validates a number of correlative biomarkers previously associated with small cell lung cancer prognosis, or implicated as possible predictive correlates of benefit from Bcl-2-directed therapy.

ABSTRACT

Purpose: Bcl-2 is a critical regulator of apoptosis that is overexpressed in the majority of small cell lung cancers (SCLC). Navitoclax (ABT-263) is a potent and selective inhibitor of Bcl-2 and Bcl-x_L. The primary objectives of this Phase 2a study included safety at the recommended Phase 2 dose and preliminary, exploratory efficacy assessment in patients with recurrent and progressive SCLC after at least one prior therapy.

Experimental Design: 39 patients received navitoclax 325mg daily, following an initial lead-in of 150mg daily × 7 days. Study endpoints included safety and toxicity assessment, response rate, progression-free and overall survival (PFS and OS), as well as exploratory pharmacodynamic correlates.

Results: The most common toxicity associated with navitoclax was thrombocytopenia, which reached grade 3-4 in 41% of patients. Partial response was observed in one (2.6%) patient and stable disease in 9 (23%). Median PFS was 1.5 months, and median OS 3.2 months. A strong association between plasma pro-gastrin releasing peptide (pro-GRP) level and tumor Bcl-2 copy number (R=0.93) was confirmed. Exploratory analyses revealed baseline levels of cytokeratin 19 fragment antigen 21-1, neuron-specific enolase, pro-GRP, and circulating tumor cell number as correlates of clinical benefit.

Conclusion: Bcl-2 targeting by navitoclax demonstrates limited single agent activity against advanced and recurrent SCLC. Correlative analyses suggest several putative biomarkers of clinical benefit. Preclinical models support that navitoclax may

enhance sensitivity of SCLC and other solid tumors to standard cytotoxics. Future studies will focus on combination therapies.

INTRODUCTION

Recurrent metastatic small cell lung cancer (SCLC) has a dismal prognosis and few therapeutic options (1). The only therapy for recurrent SCLC currently approved for use in the United States is topotecan, based on a study that demonstrated improvement in adverse symptoms and similar outcome relative to combination chemotherapy.(2) The clinical benefit of topotecan is generally limited to patients with objective response or stable disease persistent for at least 3 months following completion of first line chemotherapy. For patients with refractory SCLC, defined as progressive disease during or within 3 months after first line therapy, there are no effective therapies, and topotecan demonstrates a response rate of only 2 – 6% (2, 3). There are no FDA-approved therapies for disease progression following topotecan.

In contrast to the marked heterogeneity of non-small cell lung cancer, SCLC is characterized by mutations or aberrant expression of several key oncogenes and tumor suppressors (*TP53*, *RB*, *BCL2*) in a large majority of cases (4). Bcl-2 is a central apoptotic inhibitor, overexpression of which is associated with both malignant transformation and chemotherapeutic resistance (5). Overexpression of Bcl-2 has been reported in up to 80% of SCLC, associated with gene amplification of the *BCL2* locus on chromosome 18q21 (6-9).

Navitoclax is a selective high-affinity small molecule inhibitor of Bcl-2 and the related apoptotic inhibitor Bcl-x_L (10, 11). Navitoclax binds to and inhibits Bcl-2 in a protein domain similar to that of naturally occurring Bcl-2 inhibitors such as Bax or Bak. Treatment of mice bearing any of several SCLC cell line xenograft tumors with single

agent navitoclax, or the closely related parent molecule ABT-737, resulted in dramatic tumor responses (11-14). The primary toxicity of ABT-737 and of navitoclax in preclinical models was a dose-dependent rapid thrombocytopenia, an on-target effect due to inhibition of Bcl-x_L in platelets (15, 16).

We initiated a phase 1/2 study of single agent navitoclax, with the phase 1 portion of this study open to patients with all solid tumors. Key observations from the phase 1 portion of the study included confirmation of rapid and dose-dependent thrombocytopenia, a plasma half-life of approximately 15 hours, and preliminary evidence of anti-cancer activity including a durable partial response in a patient with SCLC (17). The gene encoding Pro-GRP is in close proximity to *BCL2* on 18q21, and data from the phase 1 study suggested a correlation between *BCL2* gene copy number and plasma pro-GRP levels, as well as between the percent change in pro-GRP over the first 35 days of continuous treatment and reduction in tumor size (17). Exploration of doses and schedules in phase I suggested a recommended phase 2 dose of 325 mg daily, and demonstrated that the severity of initial thrombocytopenia could be attenuated by a 7-day lead-in of 150 mg daily. This regimen was further explored in a phase 2 expansion, limited to patients with relapsed or refractory SCLC. Here we summarize the clinical outcome data in this cohort of SCLC patients, present confirmatory data regarding exploratory correlates from the phase 1, and extend these data to include evaluation of potential prognostic markers previously reported in SCLC, including cytokeratin 19 fragment antigen 21-1 (CYFRA 21-1), neuron-specific enolase (NSE), and circulating tumor cell (CTC) enumeration (18-25).

METHODS

Patient Population

Eligible candidates for the phase 2 portion of this study (NCT00445198) were adults with histologically or cytologically confirmed SCLC, with progressive disease after at least one prior chemotherapy regimen. Any number of prior therapies were allowed. Eligible patients had an Eastern Cooperative Oncology Group (ECOG) performance status of ≤ 1 , and had adequate bone marrow, renal, and hepatic function. Patients were excluded if they had underlying or predisposing condition of bleeding (history of non-chemotherapy induced thrombocytopenia with bleeding within 1 year, active peptic ulcer or hemorrhagic esophagitis/gastritis, active immune thrombocytopenic purpura, etc.).

This study was conducted according to the Declaration of Helsinki and with approval from Institutional Review Boards of all participating study sites. All participants provided written informed consent before participating.

Study Design

The phase 2 component of this clinical trial was an open-label, single arm study of patients with recurrent and progressive SCLC after at least one prior therapy. Based on the dose and schedule determined in phase 1, patients were treated with navitoclax 150 mg daily for 1 week, and 325 mg daily thereafter. Cycle duration was defined as 21 days. Subjects could remain on therapy indefinitely, until disease progression or intolerable toxicity.

Assessments

Safety

Safety assessments included history and physical examinations, vital signs, ECOG performance status, AEs, blood chemistry and complete blood counts with differential. Safety assessments were performed at screening, weekly during cycle 1, and at the start of subsequent cycles. Adverse event severity was graded according to the National Cancer Institute Common Terminology Criteria for Adverse Events (NCI CTCAE), version 3.0 (26). Relationships of adverse events to navitoclax (definitely, probably, possibly, unlikely, or unrelated) were assessed by the Principal Investigator at each site.

Efficacy

Tumor response was assessed using RECIST after every 2 cycles of therapy (27). Additional efficacy variables included PFS and OS.

Pharmacodynamic Correlates

Blood specimens for analyzing circulating tumor cells (CTC) were collected at screening, on day 14 of cycles 1 and 2, and at final study visit for US patients only. Circulating tumor cell detection was performed as previously described (28), using the CellSearch system (Veridex, Raritan, NJ). Samples enriched for CTC were removed from the Veridex cartridge after imaging and enumeration, washed in PBS, pelleted, resuspended in 100 μ L PBS, dropped and then dried on a positively charged slide

(Biogenex, San Ramon, CA). Slides were rinsed in water prior to imaging using the Bioview imaging system (Tel Aviv, Israel) to identify and record loci of CTC. Fluorescence in-situ hybridization (FISH) was performed as previously described (29) using a Vysis LSI Bcl-2® (orange) probe and chromosome 18 probe (green) developed by Abbott Molecular (Des Plaines, IL). The slides were returned to the Bioview imaging system and the previously identified CTC assessed for DNA copy number.

Serum and plasma samples collected at the same intervals were stored at -70°C or colder until analyzed for quantitative assessment of tumor markers. Plasma CYFRA 21-1 and Pro-GRP were measured using automated ARCHITECT chemiluminescent microparticle immunoassay (Abbott Diagnostics, Abbott Park, IL) and NSE was measured in plasma using automated electrochemoluminescent assays on Elecsys 2010 (Roche Diagnostics, Germany) in the laboratory of Petra Stieber (Institute of Clinical Chemistry, University Hospital Munich, Germany). Serum samples were analyzed for cleaved and full length cytokeratin 18 fragment (M30 and M65) enzyme-linked immunosorbent assays (ELISAs; Peviva, Stockholm, Sweden) using previously described assays validated to good clinical laboratory practice in the laboratory of author C.D. (30, 31). These determinants have been previously explored as diagnostic, prognostic, or predictive biomarkers in cancer patients (18-21).

Statistical Analysis

All subjects enrolled were included in clinical outcome analyses. Descriptive statistics were used to summarize demographic variables. Progression free survival and overall survival were computed using Kaplan-Meier methodology and 95% confidence

intervals were provided. Progression free survival was defined as the number of days from the date the subject started study drug to the date the subject experienced disease progression or death. Overall survival was defined as the number of days from the date the subject started study drug to the date of death. A sample size of 40 subjects was chosen to provide approximately 16 patients with chemosensitive disease and 16 with chemoresistant disease. This sample size provided 90% confidence that the true response rate was within 20-25% of the observed response rate.

Correlations between median *BCL2* copy number and pro-GRP level were performed by the Pearson correlation. M30 and change in CTC number from baseline comparisons among Pro-GRP groups were performed by one-way ANOVA using JMP 8.0 statistical software. Optimized thresholds for the tumor markers were obtained using BATTing (Bootstrapping and Aggregating Thresholds from Trees) method (32, 33). BATTing uses tree-based model for threshold estimation. However, a single tree may be unstable and not robust enough against small perturbations in the distribution of the data. In addition, single tree-based models are known to be prone to overfitting and have poor prediction power. Those issues are addressed in BATTing by aggregating the thresholds from multiple trees to get a more robust estimate. Each tree is built using a bootstrap random sample drawn from the original population and provides its own cutoff. The final estimate of the threshold is calculated as the median from the distribution of cutoffs generated from the multiple trees. This procedure was implemented using the R statistical software.

RESULTS

Patient Characteristics and Study Drug Dosing

A total of 39 patients participated in the phase 2 study. The majority of patients had performance status 1. The median number of prior therapeutic regimens was 2 (range 1 – 6). A summary of patient demographics is shown in **Table 1**. All patients were started at the planned dose and schedule, including dose escalation to 325 mg in week 2. The primary reasons for study discontinuation included disease progression (N = 26, 67%) and AEs (N=11, 28%). The median number of treatment cycles was 2 (range, 1 – 11) and the median treatment duration was 1.3 months (range, 0.2 - 7.3).

Safety and Tolerability

Navitoclax-related AEs are summarized in **Table 2**. The most common AE was thrombocytopenia, however this was not associated with bleeding sequelae. A total of 17 of 39 patients (46.3%) required dose interruption and 7 (17.9%) required dose reduction on study. The most common reasons for dose interruption were thrombocytopenia (12.8%), fatigue (7.7%) and nausea (7.7%), all of which reversed with drug discontinuation. Eleven (28%) subjects discontinued for AE's. Of these, 4 (10%) were considered possibly or probably related to ABT-263.

Efficacy

A waterfall plot representing best % change from baseline in identified target lesions is shown in **Figure 1**. Only one confirmed partial response was observed

(2.6%). Nine patients (23.1%) experienced stable disease as best response. Sixteen patients (41%) had disease progression, and another 13 (33.3%) were not evaluable for response. These 13 did not complete 2 cycles of therapy, and did not have post-treatment tumor assessments on protocol. The majority of these patients discontinued therapy due to disease progression or decline in performance status. Median progression-free survival was 1.5 months (95% CI 1.4 – 1.7) and median overall survival 3.2 months (95% CI 2.3 – 8.1) (**Table 3**).

Pharmacodynamic Correlates

Several exploratory correlative biomarkers were included in this study, including both CTC enumeration and plasma protein markers associated with small cell lung cancer. The biomarker analysis included all phase II patients; to increase the sample size, patients with SCLC dosed with ≥ 325 mg navitoclax on the phase I portion of this study were also included in the analysis. Biomarkers correlating with outcome that are assessable prior to treatment are of particular potential utility, consequently we determined optimized thresholds prognostic for patient outcome using BATTing for several tumor markers including CYFRA 21.1, NSE, and CTC at baseline and on cycle 1 day 14. A summary of pretreatment biomarkers evaluated are presented in **Table 4**. Interestingly, baseline levels of CTC, and of both plasma biomarkers, appear to be associated with both PFS and OS in patients treated with navitoclax. CTC levels at cycle 1 day 14 were also significantly associated with outcome (data not shown).

By grouping patients above these thresholds for both NSE and CYFRA in the analysis, we identified a population of patients with poor prognosis (median PFS: 41

days [n=16] vs. 55 days [n=23], **Figure 2A**, $p=0.0006$; median OS: 61 days [n=15] vs. 242 days [n=23], **Figure 2B**, $p=0.0009$).

Pro-GRP was of particular interest with regard to navitoclax, as the *GRP* gene is in close chromosomal proximity to *BCL2*, the gene encoding the primary target for this drug. *BCL2* copy number correlates with relative sensitivity to navitoclax in SCLC cell lines. Plasma pro-GRP levels may correlate with tumor *BCL2* copy number, and were implicated in our phase I study as a relatively non-invasive means of assessing tumor *BCL2* gene amplification.(17) To further evaluate this association, we assessed plasma Pro-GRP, and *BCL2* copy number by FISH in CTC from patients on this study (n=10). A strong correlation was confirmed (Pearson correlation 0.93, $p < 0.0001$). We identified an optimal Pro-GRP threshold linked with amplification of Bcl-2 at 600 pg/mL and compared the activity of navitoclax between patients above and below this threshold using two measurements of activity.

First, we examined the early activation of the apoptotic pathway in patient serum, with the M30 ELISA assay. This assay measures the release of caspase-cleaved cytokeratin 18 following apoptotic cell death. M30 concentrations were significantly increased in patients in the high Pro-GRP group (n=19) when compared to those with low Pro-GRP (n=13) (212% vs 109%, **Figure 2C**, $p=0.0178$). Second, we examined the changes in CTC numbers and found that the median CTC number increased more in the low pro-GRP group (n=8) by 518% of baseline, than in the high pro-GRP group (n=12) (186% of baseline, **Figure 2D**, $p=0.0433$).

DISCUSSION

A primary objective of this Phase 2a study was to assess safety at recommended Phase 2 dose. As anticipated from the Phase 1 study, the most common grade 3-4 toxicity was thrombocytopenia (17). However, this was not associated with clinically significant bleeding events and was reversible with interruption of study drug. Thrombocytopenia represents an expected and on-target toxicity of this drug, due to potent inhibition of Bcl-x_L, a close functional homolog of Bcl-2 that is expressed in platelets (15). Bcl-x_L inhibition has been demonstrated to result in platelet apoptosis, with a bias toward apoptosis of older platelets in circulation (16).

The level of single agent activity of navitoclax in recurrent SCLC was disappointing, given its remarkable preclinical *in vivo* activity as a single agent in multiple SCLC cell line xenograft models (12). This activity was in keeping with our earlier report using the closely related parental drug, ABT-737, in both *in vitro* and *in vivo* testing of multiple cell lines (13). In sharp contrast to these results, evaluation of ABT-737 activity in primary xenograft SCLC tumor models, derived from direct transfer of human tumors into recipient mice, demonstrated low level single agent activity, varying from no effect relative to control to inhibition of growth trajectory without evident tumor reduction (34). Gene expression patterns in primary SCLC xenografts more closely mimic the human tumors of origin than do related cancer cell lines (35). In retrospect, the clinical data presented here support the predictive value of the primary xenograft model as a tool in cancer drug development, and more broadly underline the importance of evaluating potential therapeutics across a spectrum of relevant preclinical disease models.

Multiple preclinical models, including primary xenografts, in fact do support that ABT-737 and navitoclax, through lowering of the tumor apoptotic threshold, enhance the efficacy of standard cytotoxic agents against SCLC and other solid tumors (36-42). Thus combinatorial regimens involving navitoclax with chemotherapy or radiation may be of particular interest. This strategy is being actively tested in a variety of solid tumors.

The correlative biomarkers evaluated in this study are currently exploratory in nature. The relationship between tumor size and these markers has not been explored for first line treatments in SCLC, and the independent prognostic role of the markers is not clear.. We hypothesize that some of these endpoints, including CTC enumeration and tumor marker expression, may represent prognostic biomarkers, associated with poor outcome across a variety of therapeutic strategies and in a variety of clinical contexts (reviewed in (43)). NSE, CYRFA 21-1, CTC, and M30 have been reported as poor prognostic factors in newly diagnosed SCLC (18, 22-24). Plasma pro-GRP, in contrast, we hypothesize may reflect tumor dependence on Bcl-2, and may be more closely linked to activity of potent and specific Bcl-2 inhibitors. If pro-GRP is confirmed as a predictive factor for sensitivity to Bcl-2-targeted inhibition, this could represent a strategy for selective treatment of patients for future clinical trials of navitoclax. In contrast, hypothesized prognostic markers (CYFRA-21-1, NSE and CTC), if confirmed in other settings, could be considered as stratification factors for future clinical research in SCLC. Evaluation of these putative biomarkers in future clinical trials in SCLC will better define their utility as prognostic markers for SCLC or predictive markers for treatment with navitoclax.

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TABLES

Table 1. Patient demographics

Demographic category	N = 39	
Median age (range)	64 (45 – 78)	
Female, n (%)	21 (54)	
Refractory disease n (%)	21 (54)	
Sensitive disease n (%)	17 (44)	
ECOG PS, n (%)	0	13 (33)
	1	25 (64)
	(missing)	1 (3)
Prior therapeutic regimens	1 – 2	28 (72)
	≥ 3	11 (28)

Table 2. Treatment-related adverse events occurring in ≥ 10% of patients

Adverse event	Any grade N (%)	Grade 3/4 N (%)
Thrombocytopenia	24 (62)	16 (41)
Diarrhea	19 (49)	2 (5)
Nausea	18 (46)	1 (3)
Fatigue	14 (36)	2 (5)
Decreased appetite	7 (18)	-
AST elevation	7 (18)	3 (8)
Neutropenia	6 (15)	4 (10)
ALT elevation	6 (15)	3 (8)
Dehydration	4 (10)	2 (5.1)
Dyspepsia	4 (10)	-
Dysgeusia	4 (10)	-

AE possibly, probably, or definitely related are included.

AST: aspartate aminotransferase; ALT: alanine aminotransferase.

Table 3. Efficacy data summary

Efficacy endpoint	Patients (N = 39)
Median PFS, mo (95% CI)	1.5 (1.4, 1.7)
Median OS, mo (95% CI)	3.2 (2.3, 8.1)
ORR, % (95% CI)	2.6 (0.1, 13.5)
Best response, N (%)	
PR	1 (2.6)
SD	9 (23.1)
PD	16 (41.0)
inevaluable*	13 (33.3)

PFS=progression-free survival; OS=overall survival; mo= months; CI=confidence interval;
 ORR=overall response rate.
 *Baseline tumor data only.

Table 4. Baseline biomarker summary

Baseline biomarker	Threshold	Median PFS			Median OS		
		Above threshold D (N)	Below threshold D (N)	p-value	Above threshold D (N)	Below threshold D (N)	p-value
CYFRA 21.1	2.3 ng/ml	44 (18)	57 (14)	0.0016	72(18)	242 (14)	0.007
NSE	15 ng/ml	44 (20)	68 (12)	0.0044	90 (20)	242 (12)	0.0088
CTC	12 /7.5ml	37 (16)	59 (12)	0.0183	67 (16)	NR (12)	0.0057

D=days; N=number of patients; CYFRA=cytokeratin-19 fragment; NSE=neuron-specific enolase; CTC= circulating tumor cells;
 NR= not reached.

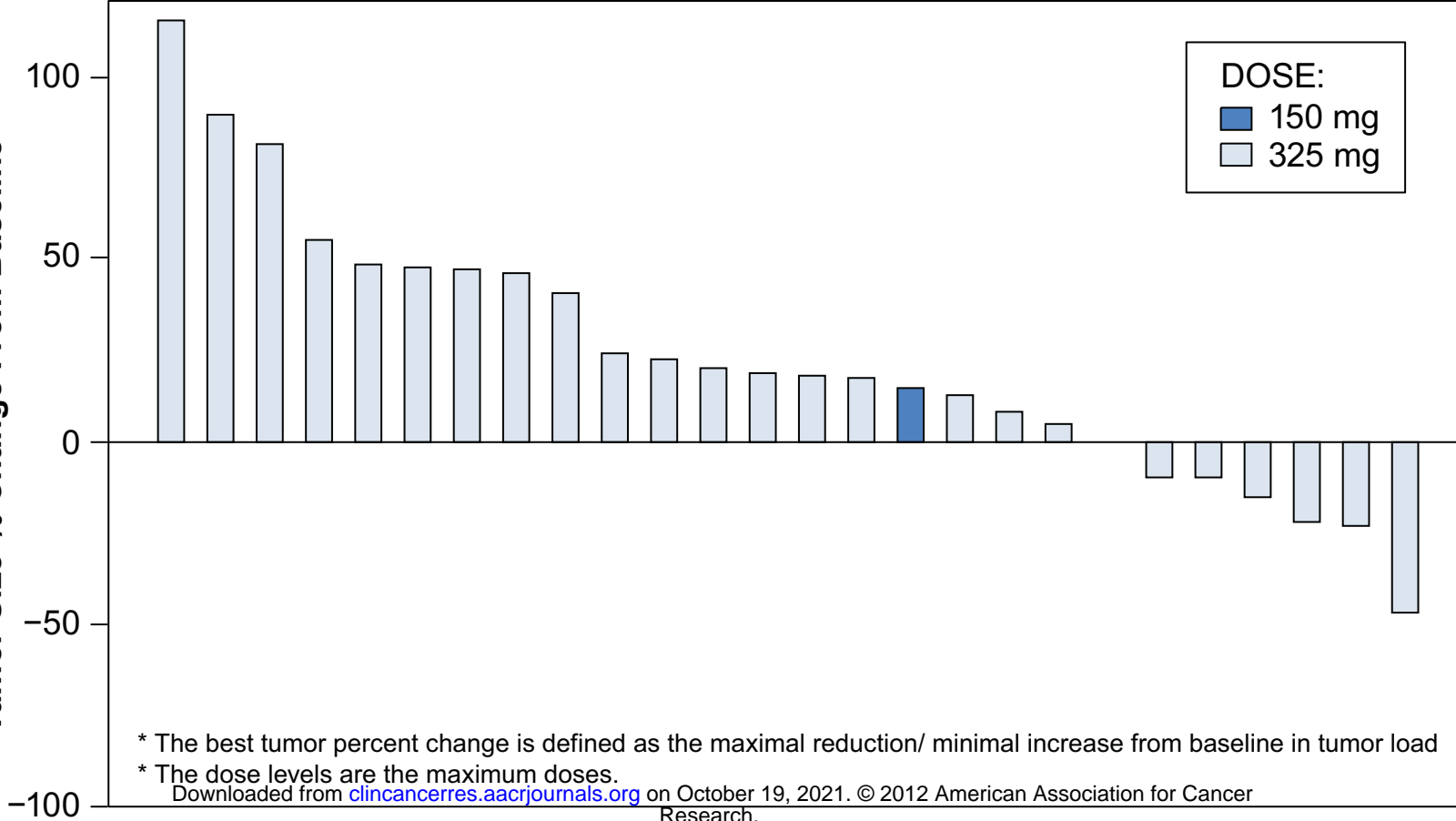
FIGURE LEGENDS

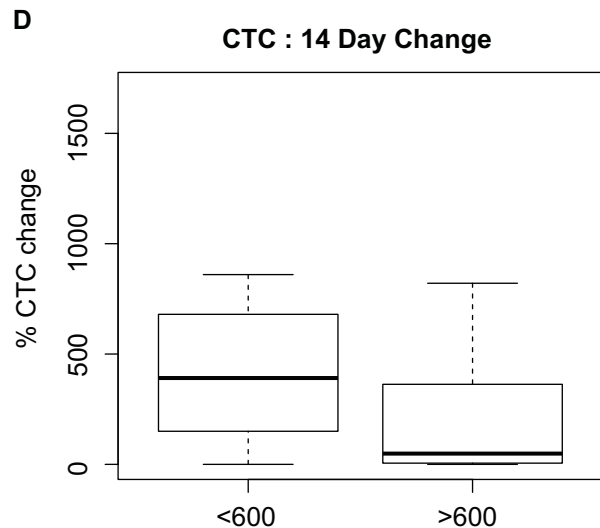
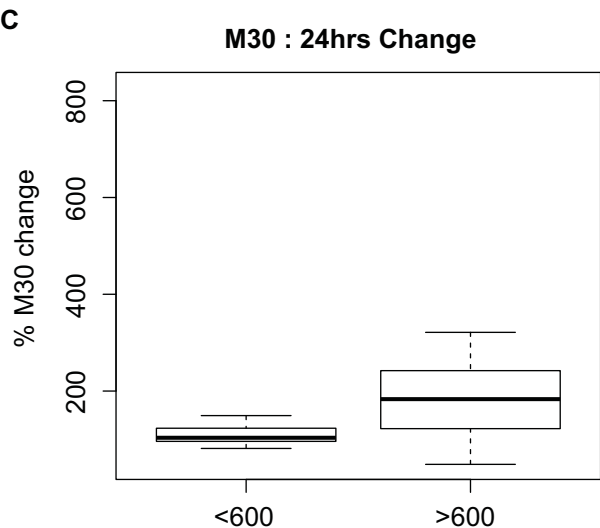
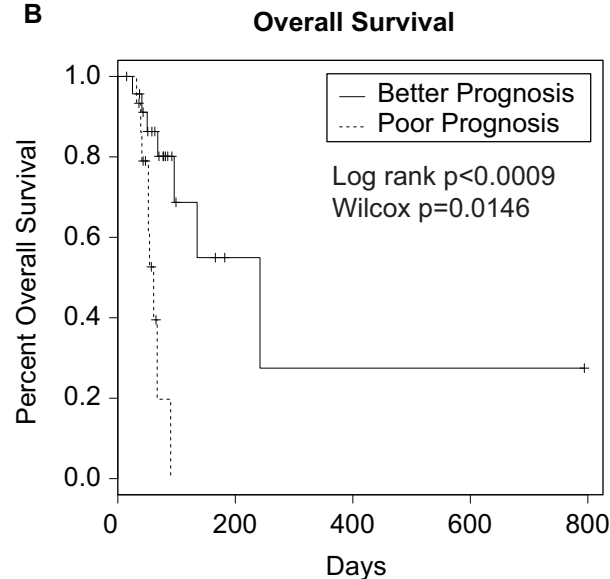
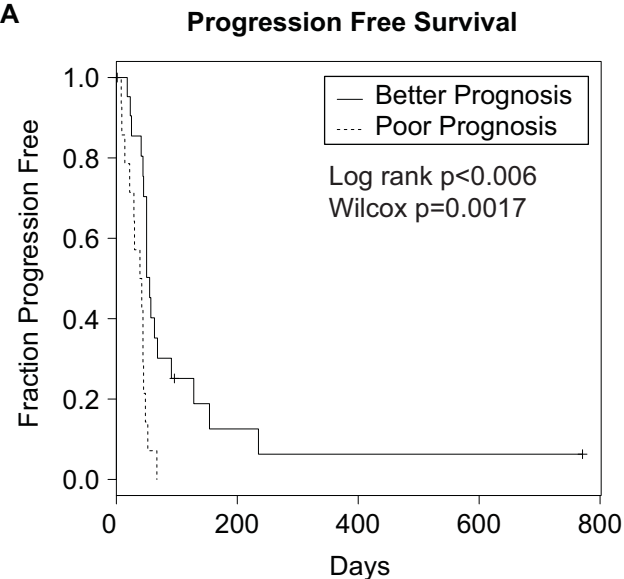
Figure 1. Waterfall plot of best fractional change in tumor size relative to baseline.

The best tumor size % change from baseline is defined as the maximal reduction or minimal increase in sum of longest dimensions of target lesions relative to pretreatment assessment.

Figure 2. Pharmacodynamic biomarker assessment. NSE and CYFRA thresholds define a patient population with markedly poorer prognosis as assessed by progression-free survival (**A**) and overall survival (**B**). Patients with high plasma Pro-GRP (correlating with *BCL2* gene amplification) demonstrated increased apoptosis as assessed by the M30 ELISA assay (**C**), and did not demonstrate an increase in CTC from baseline to day 14, in contrast to patients with low plasma Pro-GRP (**D**).

Tumor Size % Change From Baseline





Clinical Cancer Research

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Charles M. Rudin, Christine L Hann, Edward B Garon, et al.

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