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STUDY OF ALECTINIB VERSUS CRIZOTINIB IN  
TREATMENT-NAIVE ANAPLASTIC LYMPHOMA KINASE-  
POSITIVE ADVANCED NON-SMALL CELL LUNG CANCER

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## STATISTICAL ANALYSIS PLAN

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TREATMENT-NAIVE ANAPLASTIC LYMPHOMA  
KINASE-POSITIVE ADVANCED NON-SMALL CELL LUNG  
CANCER

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## STATISTICAL ANALYSIS PLAN AMENDMENT APPROVAL

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## LIST OF ABBREVIATIONS AND DEFINITIONS OF TERMS

Abbreviation or Term	Definition
ALK	anaplastic lymphoma kinase
CDOR	CNS duration of response
CORR	CNS objective response rate
CR	complete response
CSR	Clinical Study Report
CT	computed tomography
DOR	duration of response
ECOG	Eastern Cooperative Oncology Group
eCRF	electronic Case Report Form
EORTC	European Organization for Research and Treatment of Cancer
FDA	U.S. Food and Drug Administration
FISH	fluorescence in situ hybridization
FPP	FISH positive population
HR	hazard ratio
iDCC	independent Data Coordinating Center
iDMC	independent Data Monitoring Committee
IHC	immunohistochemistry
IRC	Independent Review Committee
ITT	intent-to-treat
IxRS	interactive voice or Web-based response system
MRI	magnetic resonance imaging
NCI CTCAE	National Cancer Institute Common Terminology Criteria for Adverse Events
NE	not evaluable
NSCLC	non-small cell lung cancer
ORR	objective response rate
OS	overall survival
PD	progressive disease
PFS	progression-free survival
PK	pharmacokinetic
PR	partial response
PRO	patient-reported outcome
PS	Performance Status
QLQ-C30	Core Quality of Life Questionnaire
QLQ-LC13	Quality of Life Questionnaire Lung Cancer Module
QoL	quality of life

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Abbreviation or Term	Definition
RANO	Response Assessment in Neuro-Oncology
RECIST	Response Evaluation Criteria in Solid Tumors
SAP	Statistical Analysis Plan
SD	stable disease
SDF	survival distribution function
TTD	time to deterioration
ULN	upper limit of normal

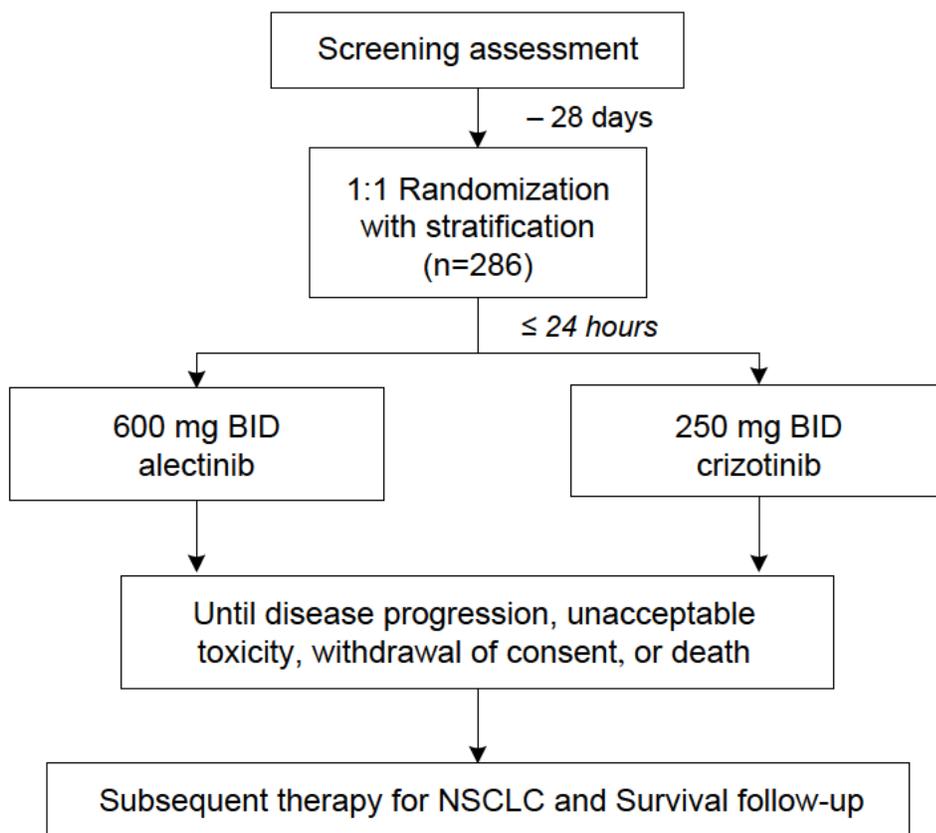
## 1. **BACKGROUND**

Alectinib is a small molecule that is highly selective and a potent inhibitor of anaplastic lymphoma kinase (ALK). Alectinib is currently being developed for the treatment of patients who have ALK-positive non-small cell lung cancer (NSCLC), as detected by a U.S. Food and Drug Administration (FDA)-approved test.

## 2. **STUDY DESIGN**

This is a randomized, active-controlled, multicenter, Phase III, open-label study in patients with treatment-naïve ALK-positive advanced NSCLC. All patients are required to provide pretreatment tumor tissue to confirm the presence of ALK rearrangement (by immunohistochemistry [IHC] test). Patients will be randomized in a 1:1 ratio into one of the two treatment arms to receive either alectinib or crizotinib (Figure 1).

**Figure 1 Summary of Study Design**



BID=twice daily; n=number of patients; NSCLC=non-small cell lung cancer.

### 2.1 **PROTOCOL SYNOPSIS**

The Protocol Synopsis is in [Appendix 1](#). The outcome measures are listed there.

## 2.2 DETERMINATION OF SAMPLE SIZE

The focus of this clinical trial is hypothesis testing, and the primary endpoint of progression-free survival (PFS) was used to determine the sample size of the study.

The Phase III PROFILE 1014 study of crizotinib versus standard pemetrexed–platinum-based chemotherapy in previously untreated patients with ALK-positive non-squamous NSCLC reported a median PFS of 10.9 months for crizotinib (Solomon et al. 2014). A hazard ratio (HR) of 0.65 for alectinib versus crizotinib (i.e., an increase in median PFS from 10.9 months to 16.8 months) will be targeted.

In this study, 286 patients will be enrolled in a 1:1 randomization allocation. Enrollment will take approximately 24 months on the basis of an assumption of non-linear recruitment as follows:

- Month 1: 1 patient per month
- Month 2: 2 patients per month
- Month 3: 4 patients per month
- Month 4: 6 patients per month
- Month 5: 8 patients per month
- Month 6: 10 patients per month
- Month 7: 12 patients per month
- Months 8–12: 13 patients per month
- Months 13–14: 14 patients per month
- Month 15 onwards: 15 patients per month

Approximately 170 PFS events are required to achieve 80% power of the log-rank test at a two-sided alpha level of 5%. This number of events is estimated to occur approximately 33 months after the first patient has been enrolled.

To illustrate sensitivity, if only 160 PFS events are observed by the clinical cutoff, the study has 78.1% power; if only 165 PFS events are observed, the study has 79.3% power with the assumed HR of 0.65 and an alpha of 0.05.

No interim analysis for efficacy or futility is planned.

An analysis of overall survival (OS) will be performed at the time of the final analysis of the primary endpoint of PFS. A survival follow-up analysis will be performed once approximately 50% of patients (i.e., 143 patients) have died. The median OS in the crizotinib arm is assumed to be 24 months and the expected median OS in the alectinib treatment arm is 30 months, equating to an HR of 0.83. On the basis of the sample size (n=286), the trial will not be powered to demonstrate a statistically significant difference

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in OS of this magnitude. At the time of the final analysis of the primary endpoint of PFS, on the basis of the above assumptions, 106 OS events are expected to have occurred. The events required for the survival follow-up analysis are expected to occur approximately 42 months after the first patient has been enrolled.

### **2.3 ANALYSIS TIMING**

This study is event driven with a recruitment period of approximately 24 months. The required number of events for the primary analysis of the primary endpoint (investigator-assessed PFS) is expected approximately 33 months after the first patient has been enrolled. If the alectinib median PFS by investigator is not estimable at the time of the primary analysis due to an insufficient number of events in this treatment arm, an exploratory update of PFS analysis will be performed when approximately 50% patients in the alectinib arm will have experienced a PFS event. This analysis will be conducted purely for the purpose of obtaining a median estimate of PFS in the alectinib treatment arm, no formal treatment comparisons will be performed for this endpoint after the primary analysis.

A survival follow-up analysis will be performed once approximately 50% of patients have died, which is estimated to occur approximately 42 months after the first patient has been enrolled.

The study will formally end once the survival follow-up analysis is complete.

## **3. STUDY CONDUCT**

### **3.1 RANDOMIZATION ISSUES**

Patients will be randomly assigned in a 1:1 allocation ratio to the two treatment arms via a block stratified randomization procedure.

Randomization will guard against systematic selection bias and should ensure the comparability of treatment groups. To assist balance in important prognostic factors, randomization will be stratified by Eastern Cooperative Oncology Group Performance Status (ECOG PS; 0/1 vs. 2), race (Asian vs. non-Asian), and CNS metastases at baseline (yes vs. no).

The randomization is checked three times during the study conduct by the independent Data Coordinating Center (iDCC).

### **3.2 INDEPENDENT REVIEW FACILITY**

An independent imaging group will be used to evaluate tumor assessments for determination of progression and response rate according to Response Evaluation Criteria in Solid Tumors (RECIST), Version 1.1 for the primary analysis. Imaging studies (computed tomography [CT]/magnetic resonance imaging [MRI]/bone scans) will be acquired according to a standard protocol and will be forwarded to the independent

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reviewers. In addition, relevant cytologic and medical data (e.g., relevant cytology reports documenting malignant pleural effusions, bone marrow aspirations, cerebral spinal fluid, CNS symptoms, corticosteroid use, etc.) may be forwarded, if available, to the independent reviewers to aid with assessment of progressive disease (PD) and response. Investigator tumor assessments will not be reconciled with the Independent Review Committee (IRC) tumor assessments. Further details are included in the IRC Charter. Details of imaging handling procedures are also described in a separate laboratory manual.

For evaluation of the CNS endpoints, the IRC will perform an assessment of scans based on RECIST v1.1 and Response Assessment in Neuro-Oncology (RANO) criteria. More details are given in the IRC charter.

### **3.3 DATA MONITORING**

Although the study is open label, the Sponsor will not analyze aggregated data by treatment group before analysis of the primary endpoint of PFS.

An independent Data Monitoring Committee (iDMC) has been incorporated into the trial design to objectively and carefully review the accumulating safety data for alectinib. Regular safety interim analyses to be reviewed by the iDMC have been planned to evaluate safety information from the study. Outputs for these reviews are prepared by an external iDCC.

## **4. STATISTICAL METHODS**

The analyses outlined in this Statistical Analysis Plan (SAP) supersede those specified in the protocol for the purpose of a regulatory filing.

### **4.1 ANALYSIS POPULATIONS**

#### **4.1.1 Intent-To-Treat Population**

The primary analysis population for efficacy is the intent-to-treat (ITT) population, defined as all randomized patients. Patients will be assigned to the treatment group to which they were randomized.

#### **4.1.2 Safety Population**

The primary analysis population for safety is the Safety Population, defined as all patients who received at least one dose of study medication. Patients will be assigned to treatment groups as treated, and all patients who received any dose of alectinib will be included in the alectinib treatment arm.

#### **4.1.3 FISH Positive Population**

A secondary analysis population for efficacy is the fluorescence in situ hybridization (FISH) Positive Population (FPP), defined as all patients in the ITT population who were ALK-positive as assessed using the Vysis FISH assay. Patients will be assigned to the

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treatment group to which they were randomized. A supportive efficacy analysis will be performed based on the FISH assay rather than the IHC.

An analysis of the primary endpoint of investigator-assessed PFS based on the FPP will be performed at the time of the primary analysis. T-tests to compare baseline characteristics by treatment group will be performed to assess any imbalance between treatment arms in the FPP. In the event that the randomization is not maintained in the FPP (t-test  $p < 0.01$ ), log-rank analyses of investigator-assessed PFS will be stratified by randomization strata and the significant baseline characteristics will be included in the Cox model. A major discrepancy between the Ventana IHC assay and the Vysis FISH assay may necessitate a follow-up analysis of the primary endpoint based on approximately 170 PFS events observed in the FPP to ensure 80% power of the log-rank test under the assumption of an HR of 0.65 in this secondary population.

#### **4.1.4 Pharmacokinetic-Evaluable Population**

The Pharmacokinetic-Evaluable Population is defined as all patients who received any dose of study medication and who have at least one post-baseline pharmacokinetic (PK) sample available.

## **4.2 ANALYSIS OF STUDY CONDUCT**

Study enrollment, patient disposition, reasons for discontinuation from the study treatment, and reason for study termination will be summarized for all patients in the ITT population.

Protocol deviations, including violations of inclusion and/or exclusion criteria and deviations during study conduct will be reported and summarized.

## **4.3 ANALYSIS OF TREATMENT GROUP COMPARABILITY**

Demographic, baseline disease characteristics, and lung cancer history, including prior brain radiation, will be compared descriptively between both treatment arms for the ITT population. Descriptive baseline summaries of continuous data will present the group mean, standard deviation, median, minimum, and maximum. Descriptive baseline summaries of discrete data will present the category counts as frequencies and percentages.

A summary of concordance of stratification factors determined by electronic Case Report Form (eCRF) versus interactive voice or Web-based response system (IxRS) will also be reported.

The baseline value of any variable will be defined as the last available value recorded prior to the first administration of study medication.

Previous and concomitant cancer therapy will also be summarized, including radiotherapy and surgery, as well as subsequent anti-cancer therapy. Previous and concurrent diseases and medications will also be summarized.

The Abbott Vysis FISH test will be used as an exploratory assay after patients have been enrolled in this study. A descriptive summary of baseline characteristics, including randomization strata by treatment arm, will be produced for the FPP to assess treatment group comparability.

#### **4.4 EFFICACY ANALYSIS**

##### **4.4.1 Primary Efficacy Endpoint**

PFS is defined as the time from date of randomization to the date of first documented disease progression or death, whichever occurs first. The primary endpoint of PFS will be determined on the basis of investigator assessment of progression using RECIST v1.1. Patients who have not experienced disease progression or death at the time of analysis will be censored at the last tumor assessment date either during study treatment or during follow-up. Patients with no post-baseline tumor assessment will be censored at the date of randomization.

Patients who discontinue treatment prior to disease progression (e.g., due to toxicity) will continue in the study and will be followed until disease progression and for OS regardless of whether they subsequently receive anti-cancer therapy.

The treatment comparison of PFS will be based on a stratified log-rank test at the 5% level of significance (two-sided). The randomization stratification factors are ECOG PS (0/1 vs. 2) and race (Asian vs. non-Asian), as recorded on the eCRF, and CNS metastases at baseline (yes vs. no). These factors will be included in the stratified log-rank analysis as long as an individual stratum includes > 10% of the ITT population. For analysis purposes, stratification according to CNS metastases at baseline will be performed on the basis of the IRC assessment rather than the investigator assessment, given that the independent assessment by neuroradiologists is deemed to be the most reliable and will correspond to the populations used to assess the CNS efficacy endpoints. Results from an unstratified log-rank test will also be presented as a supportive analysis.

Because patients were stratified on the basis of CNS metastases at baseline by investigator assessment, baseline characteristics grouped by CNS metastases at baseline by IRC will be summarized by treatment arm. Concordance of CNS metastases at baseline between investigator and IRC will also be reported.

Additional supportive analyses include Kaplan-Meier and Cox modelling approaches. The Kaplan-Meier method will be used to estimate the median PFS for each treatment arm with 95% confidence limits, and a Kaplan-Meier curve will be constructed to provide a visual description of the difference between the treatment arms. A stratified Cox

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proportional hazard regression model will be used including treatment in order to provide an estimate of the treatment effect expressed as an HR (alectinib vs. crizotinib), as well as a 95% CI. The proportional hazards assumption will be assessed both graphically from the Kaplan-Meier plot as well as by adding a treatment by time interaction term to the Cox regression model. If the proportional hazards assumption is not met, alternative appropriate methods will be used.

The difference between the two treatment groups for the primary objective will be assessed and tested for the following hypothesis: the survival distribution function (SDF) of the alectinib treatment group is the same as for the crizotinib treatment group versus the alternative that the two distributions are different:

$$H_0: \text{SDF (alectinib)} = \text{SDF (crizotinib)}$$

versus

$$H_A: \text{SDF (alectinib)} \neq \text{SDF (crizotinib)},$$

where SDF denotes the survival distribution function of the parameter PFS.

#### **4.4.2 Secondary Efficacy Endpoints**

All secondary efficacy endpoints will be analyzed in the ITT population unless otherwise specified.

If the primary endpoint of investigator-assessed PFS is statistically significant at a two-sided 5% significance level based on the stratified log-rank test, the following secondary endpoints will be tested in the following sequential order (O'Neill 1997), each at a two-sided 5% significance level:

- PFS by IRC
- Time to CNS progression by IRC RECIST criteria
- Objective response rate (ORR) by investigator assessment
- OS

All tests in the sequence will be based on a stratified log-rank test at the 5% level of significance (two-sided). The stratification factors included and the analysis population will be the same as for the primary hypothesis test. Results from unstratified tests will also be presented as supportive analyses.

##### **4.4.2.1 PFS by IRC**

An analysis of PFS on the basis of the IRC assessments will be performed using the same methodology as specified for PFS on the basis of investigator assessment.

A concordance analysis between the IRC-determined and the investigator-determined PD status (yes vs. no) will be provided by treatment arm together with a summary of agreement between the IRC-determined and the investigator-determined PD dates. The early discrepancy rate (investigator assessed prior to IRC) and the late discrepancy rate

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(investigator assessed after IRC) will be calculated for each treatment arm and the differential discordance calculated as the rate on the alectinib arm minus the rate on the crizotinib arm (Amit et al. 2011). In addition, a table will be created displaying the number of visits with investigator tumor assessment but missing IRC assessment.

#### **4.4.2.2 Time to CNS Progression by IRC RECIST Criteria**

Time to CNS progression is defined as the time from randomization until first radiographic evidence of CNS progression by independent review. An independent central radiological review will be performed for all patients, and the analysis of CNS progression will be based on the data from the independent review. All patients in the ITT population will be included in the analysis regardless of their baseline status of CNS metastases. CNS progression is defined as progression due to newly developed CNS lesions and/or progression of preexisting baseline CNS lesions. On the basis of RECIST v1.1, this is defined as a new post-baseline CNS/brain lesion(s) and/or an increase of  $\geq 20\%$  in the sum of longest diameters of the measurable baseline CNS lesions compared with nadir and/or unequivocal progression of non-measurable baseline CNS lesions.

In order to account for the competing risks inherent in the comparison of CNS progression between alectinib and crizotinib, a stratified two-sided log-rank test will be computed on the basis of a cause-specific hazard function. Results from unstratified tests will also be presented as supportive analyses.

The probability of CNS progression, non-CNS progression, and death by treatment group with 95% CIs will each be estimated using cumulative incidence functions. Gray's test to compare the risk of CNS progression between alectinib and crizotinib will also be performed as a supportive analysis.

Similar analyses of CNS progression based on IRC RANO criteria will be performed.

#### **4.4.2.3 Objective Response Rate**

ORR, on the basis of investigator assessment, is defined as the percentage of patients in the ITT population with measurable disease at baseline who attain a complete response (CR) or partial response (PR). Per RECIST v1.1, confirmation of objective response is not required for this secondary endpoint. Patients without a post-baseline tumor assessment will be considered non-responders, as will patients with a best overall response of stable disease (SD), PD, or not evaluable (NE).

An estimate of ORR and its two-sided 95% CI will be calculated using the Clopper-Pearson method for each treatment arm. Response rates in the treatment groups will be compared using a stratified Mantel-Haenszel test, including the randomization stratification factors. Results from unstratified tests will also be presented as supportive analyses. The difference in ORR between the two treatment arms will be

presented together with a two-sided 95% CI on the basis of a normal approximation to the binomial distribution.

ORR by IRC will be analyzed similarly as a supportive analysis in the population of patients identified to have measurable disease at baseline according to the IRC.

#### **4.4.2.4 Duration of Response**

For patients with measurable disease at baseline who have experienced an objective response (CR or PR) during the study as assessed by the investigator, duration of response (DOR) is defined as the duration from the first tumor assessment that supports the patient's objective response (CR or PR, whichever is first recorded) to first documented disease progression or death due to any causes, whichever occurred first. Patients who have not progressed or died at the time of analysis will be censored at the last tumor assessment date. Because the determination of DOR is based on a non-randomized subset of patients, formal hypothesis testing will not be performed. DOR will be estimated using Kaplan-Meier methodology and an HR and its 95% CI on the basis of a Cox proportional regression model with stratification factors will be calculated. The proportional hazards assumption will be assessed graphically.

#### **4.4.2.5 Overall Survival**

OS is defined as the time from the date of randomization to the date of death due to any cause. Patients who are not reported as having died at the time of analysis will be censored at the date when they were last known to be alive. Patients who do not have post-baseline information will be censored at the date of randomization. OS will be analyzed using the same methodology as specified for the primary endpoint. A survival follow-up analysis will be performed based on more mature data.

#### **4.4.2.6 CNS Objective Response Rate according to RECIST v1.1 and RANO Criteria by IRC**

CNS objective response rate (CORR) will be summarized by treatment group in the subgroup of patients in the ITT population who have measurable CNS lesions at baseline as determined by the IRC. These summaries will be provided for both RECIST v1.1 and RANO. CNS responses according to RECIST v1.1 do not have to be confirmed, whereas confirmation is incorporated into the RANO criteria.

CORR will be summarized by treatment group in the subgroup of patients in the ITT population who have measurable and non-measurable CNS lesions at baseline as determined by the IRC. For this analysis, patients with non-measurable CNS disease at baseline by IRC who experience a CR in CNS lesions will be included as responders.

In addition, a subgroup analysis of CORR will be performed according to prior brain radiation status (yes vs. no) for the two analysis populations above.

#### **4.4.2.7 CNS Duration of Response according to RECIST v1.1 and RANO Criteria by IRC**

This analysis will be performed including all patients in the ITT population who have measurable CNS lesions at baseline as determined by the IRC who had a CNS response (CR or PR). CNS duration of response (CDOR) is the time from CNS response to CNS PD by IRC.

#### **4.4.3 Exploratory Efficacy Endpoints**

The Abbott Vysis FISH test will be used as an exploratory assay after patients have been enrolled in this study. As part of the rationale for the biomarker assessment, results from these analyses will be used to quantify the degree of correlation between FISH and IHC for the detection of ALK-positive NSCLC. The proportion of patients in the ITT population who have ALK-positive NSCLC by the FISH assay will be summarized. The FPP will be a secondary analysis population for efficacy. This analysis population will be used to perform a supportive analysis of the study data based on the FISH assay rather than the IHC.

#### **4.4.4 Sensitivity Analyses**

A sensitivity analysis will be performed on the primary endpoint of PFS with the following changes from the primary analysis:

- Censor patients at the last adequate tumor assessment prior to the start of non-protocol-specified anti-cancer therapy received prior to observing progression.
- Censor patients for whom documentation of disease progression or death occurs after  $\geq 2$  missed tumor assessments. These patients will be censored at the last tumor assessment prior to the missed assessments.
- Censor patients who discontinue study treatment (due to personal preference or toxicity) and/or withdraw or are lost to follow-up prior to observing progression.

Two additional sensitivity analyses for PFS include:

- The effect of missing tumor assessments will be assessed if the number of missing assessments in either arm is  $> 5\%$ . For patients with progression determined following one or more missing tumor assessments, the progression will be backdated to the first missing tumor assessment.
- The effect of loss to follow-up will be assessed depending on the number of patients who are lost to follow-up. If  $> 5\%$  of patients are lost to follow-up for PFS in either treatment arm, a “worst-case” analysis will be performed in which patients who are lost to follow-up will be considered to have progressed at the last date they were known to be progression-free.

Summary statistics on timing of scans (regardless of outcome) will be provided by treatment arm. Nelson-Aalen methods may be used to estimate the cumulative hazard rate.

A sensitivity analysis for PFS and OS will be performed based on the stratification factors entered at randomization in the IxRS system.

#### **4.4.5 Subgroup Analyses**

PFS by investigator and IRC assessments will be presented separately for important subgroups including age (<65, ≥65), sex, race (Asian, non-Asian), and smoking status and baseline prognostic characteristics including baseline ECOG PS, CNS metastases at baseline as determined by IRC, and prior brain radiation (in patients with CNS metastases at baseline). The HR including a 95% CI will be presented separately for each level of the categorical variables.

#### **4.5 PHARMACOKINETIC AND PHARMACODYNAMIC ANALYSES**

Standard non-compartmental analysis may be conducted for PK data collected from patients participating in serial and/or intensive PK collections for relevant analytes, as data allow, as appropriate, and if needed. PK parameters, including but not limited to area under the concentration-time curve, maximum concentration, and time to maximum concentration, will be calculated on the basis of the available data as appropriate and where data allow. Additional PK parameters may be calculated as deemed appropriate.

Individual and mean plasma concentrations at each sampling timepoint and/or PK parameters for alectinib and metabolite(s) will be listed, as appropriate.

Summary statistics (e.g., means, standard deviation, coefficient of variation percent, geometric means, medians, and ranges) for plasma concentrations and/or PK parameters for alectinib and metabolite(s) will be presented by nominal collection times (plasma concentrations only), as appropriate. Additional plots or summary statistics may be constructed or calculated, as appropriate.

Results of PK and/or any PK/pharmacodynamic analyses may be reported outside the Clinical Study Report (CSR).

Non-linear mixed-effects modeling (with software NONMEM) will be used to analyze the sparse and/or serial/intensive plasma concentration-time data for alectinib. The PK data from this study may be pooled with data from other studies. Population and individual PK parameters will be estimated and the influence of various covariates (such as age, sex, and body weight) on these parameters will be investigated.

Exploratory analyses will be conducted to investigate the relationship between alectinib PK exposure and efficacy/safety parameters.

Results of the mixed-effects modeling and exploratory analyses will be reported in a document separate from the CSR.

## **4.6 SAFETY ANALYSES**

All safety analyses will be performed on the Safety Population; that is, all patients who receive any dose of study medication (see Section 4.1.2) by treatment arm with all patients who received any dose of alectinib will be included in the alectinib treatment arm.

### **4.6.1 Exposure of Study Medication**

Study treatment (alectinib and crizotinib) exposure, including treatment duration, dose intensity, number of doses, percentage of planned dose, and total cumulative dose will be summarized with descriptive statistics. The number of missed doses will also be displayed.

### **4.6.2 Adverse Events**

Verbatim description of adverse events will be mapped to MedDRA thesaurus terms and graded according to the National Cancer Institute Common Terminology Criteria for Adverse Events, Version 4.0 (NCI CTCAE v4.0). All adverse events occurring during or after the first study treatment will be listed and summarized by mapped term, appropriate thesaurus level and NCI CTCAE grade.

Summary tables of the following will be provided:

- All adverse events
- Serious adverse events
- Severe adverse events (Grade 3 or higher)
- Adverse events leading to study treatment discontinuation
- Adverse events leading to dose reduction
- Adverse events leading to dose interruption
- Treatment-related adverse events
- Adverse events leading to death
- Adverse events by highest NCI CTCAE grade
- Selected adverse events relating to ALK inhibitors and/or the tyrosine kinase inhibitor class and alectinib data

A summary table of common adverse events, that is, those occurring in at least 10% of patients, will be provided.

Selected adverse events are defined as follows:

- Hepatotoxicity
- Interstitial lung disease
- Vision disorders
- Skin disorders (e.g., photosensitivity, rash)
- Anemia
- Gastrointestinal disorders (e.g., nausea, vomiting, diarrhea)
- Abnormal renal function (e.g., serum creatinine increase, renal impairment, renal failure)
- Severe myalgia and CPK elevations
- Edema
- Bradycardia

Rates of each selected adverse event group will be summarized including incidence of Grade 3 or higher events, serious adverse events, and treatment-related events.

Multiple occurrences of the same event will be counted once at the maximum severity. All deaths and causes of death will be summarized.

Adverse events by sex, age (<65 vs. ≥65 years), race (non-Asian vs. Asian), and CNS metastases at baseline (yes vs. no) will be presented.

#### **4.6.3            Laboratory Data**

Laboratory data will be classified according to NCI CTCAE v4.0 and will be summarized descriptively over time, including change from baseline. Highest NCI CTCAE grade after baseline will also be reported and shift tables from baseline to worst value during the study post-baseline will be presented. An additional shift table for serum creatinine will include only elevations above the upper limit of normal (ULN) (i.e., elevations above baseline, which are included in the full NCI CTCAE v4.0 definition, will be excluded from this shift table).

A Hy's law analysis will be provided. The potential Hy's law quadrant is defined as ALT or AST increases above 3-fold the ULN with concomitant total bilirubin increases above 2-fold the ULN.

In order to evaluate and compare potential hypogonadism in the two treatment arms, total and free testosterone, sex hormone-binding globulin (if available), follicle-stimulating hormone, and luteinizing hormone levels in the blood will be summarized for males by treatment arm over time.

#### **4.6.4 Vital Signs and ECOG Performance Status**

Vital signs will be summarized descriptively by treatment group over time, including change from baseline. ECOG PS will also be tabulated over time by treatment group.

#### **4.6.5 Electrocardiograms**

The following parameters are captured on the eCRF and will be listed and summarized by treatment group over time: heart rate; RR (time from one R wave to next R wave); PQ; QRS and QT duration; and QT interval corrected using Fridericia's formula.

### **4.7 PATIENT-REPORTED OUTCOME ANALYSES**

Through the use of the European Organization for Research and Treatment of Cancer (EORTC) Core Quality of Life Questionnaire (QLQ-C30) and EORTC Quality of Life Questionnaire Lung Cancer Module (QLQ-LC13), lung cancer symptoms, symptoms commonly associated with cancer treatments, and disease and treatment's impact on patients' functioning and health-related quality of life will be collected at every study visit (Week 4, Week 8, and then after every 4 weeks) until disease progression and during post-progression treatment in case of isolated, asymptomatic CNS progression; at post-treatment visit (4 weeks after permanent treatment discontinuation); and at subsequent 8-weekly survival follow-up visits for 6 months. For patients who discontinue treatment for reasons other than disease progression and who progress within the first 6 months of survival follow-up, patient-reported outcome (PRO) measures will be administered every 4 weeks until disease progression and then decreased to every 8 weeks until 6 months post-treatment. For patients who discontinue treatment for reasons other than disease progression and have not yet progressed at 6 months post-treatment, PRO measures will be administered every 4 weeks until disease progression and will no longer be required thereafter.

The baseline PRO assessment will be the first PRO assessment completed within 7 days of randomization. For the following PRO analyses, there may be multiple PRO assessments completed within the same calendar day; the earliest assessment will be used for analysis. In the cases that there are multiple PRO assessments completed within the same time window scheduled for PRO assessments, the PRO assessment closest to the visit will be chosen for analysis.

The EORTC QLQ-C30 and EORTC QLQ-LC13 will be scored according to the EORTC scoring manual 3rd edition ([Fayers et al. 2001](#)). The QLQ-C30 and QLQ-LC13 are composed of both multi-item scales and single-item measures including functional scales, symptom scales, and a global health status/quality of life (QoL) scale.

For multi-item subscales, if  $\leq 50\%$  of items within the multi-item subscale are missing at a given timepoint, the multi-item score will be calculated on the basis of the non-missing items. If  $> 50\%$  of items are missing or if a single-item measure is missing, the subscale is missing.

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All of the scales and single-item measures will be linearly transformed so that each score will range from 0 to 100. A high score for a functional scale represents a high/healthy level of functioning, a high score for the global health status/QoL represents a high QoL; however, a high score for a symptom scale/item represents a high level of symptomatology/problems.

The EuroQoL 5 Dimension questionnaire analyses are out of scope of this SAP.

#### **4.7.1 Time to Deterioration of Patient-Reported Lung Cancer Symptoms, Global Health Status, and Cognitive Function**

Time to deterioration (TTD) analyses will be performed on lung cancer symptom scores, global health status scores, and cognitive function scale scores in the ITT population and will include all data collected through disease progression and survival follow-up.

TTD is defined as the time from randomization until the first confirmed clinically meaningful deterioration. Confirmed clinically meaningful deterioration in lung cancer symptoms is defined as a  $\geq 10$ -point increase from baseline in a symptom score that must be held for at least two consecutive assessments or an initial  $\geq 10$ -point increase above baseline followed by death within 5 weeks from the last assessment. Conversely, confirmed clinically meaningful deterioration in global health status or function is defined as a  $\geq 10$ -point decrease from baseline in a symptom score that must be held for at least two consecutive assessments or an initial  $\geq 10$ -point decrease from baseline followed by death within 5 weeks from the last assessment. A  $\geq 10$ -point change in the score is perceived by patients as being clinically significant (Osoba et al. 1998).

TTD will be documented for each of the following symptom, global health status, or cognitive function scale scores:

- Cough (Question 31 on the EORTC QLQ-LC13)
- Dyspnea single item (Question 8 on the QLQ-C30)
- Dyspnea multi-item subscale (Questions 33-35 on the QLQ-LC13)
- Chest pain (Question 40 on the QLQ-LC13)
- Arm and/or shoulder pain (Question 41 on the QLQ-LC13)
- Fatigue multi-item subscale (Questions 10, 12, and 18 on the QLQ-C30)
- Global Health Status scale (Questions 29 and 30 on the QLQ-C30)
- Cognitive Function Scale (Questions 20 and 25 on the QLQ-C30)

In addition to those single symptom endpoints, a 3-symptom composite endpoint will also be assessed. In this instance, symptom deterioration will be determined as a  $\geq 10$ -point increase above baseline in any of the following lung cancer symptom scores, whichever occurs first (cough, chest pain, and dyspnea multi-item subscale). In the same manner as the single symptom endpoints, symptom deterioration will need to be confirmed for the original symptom; a  $\geq 10$ -point increase from baseline in a symptom

score (i.e., cough) that must be held for at least two consecutive assessments (i.e., cough for two assessments) or an initial  $\geq 10$ -point increase above baseline followed by death within 5 weeks from the last assessment.

Patients without baseline or post-baseline EORTC symptom or scale scores will be censored at the date of randomization. Patients without deterioration at the time of analysis will be censored at the last time they were known to have not deteriorated.

There will be no imputation for missing data for the TTD analysis. If there is a substantial amount of missingness, patient baseline and disease characteristics will be compared between patients with and without missing baseline assessments using descriptive statistics.

TTD of the prespecified symptoms will be summarized using the Kaplan-Meier method. The estimated Kaplan-Meier plots will be provided for each score previously described (including the 3-symptom composite score). A stratified log-rank test will be the primary method to compare the time to first deterioration between the treatment groups. The median time and two-sided 95% CI for the median will also be provided. Estimates of the treatment effect will be expressed as HRs with use of a stratified Cox model, including 95% CIs for the ITT population.

A sensitivity analysis will be performed to account for multiple PRO assessments completed within 7 days of randomization, the same day, or multiple assessments completed within the same time window scheduled for PRO assessments. The sensitivity analyses will use the following data:

- In the case that there are multiple baseline PRO assessments completed within 7 days of randomization, the assessment with the best score (lowest score for symptoms, highest score for global health status or function scales) for any of the pre-specified scores will be selected.
- In case of multiple assessment within the same calendar day, the worst score (highest score for symptoms, lowest score for global health status or function scales) for any of the prespecified scores will be selected.
- In the case of multiple assessments within a single time window scheduled for PRO assessments, the assessment with the worst score (highest score for symptoms, lowest score for global health status or function scales) for any of the prespecified scores will be selected.

There is risk that PFS endpoints (or death) may confound differences in CNS events and in PRO cognitive function, therefore, a competing risks analysis may be done on the TTD for Cognitive Function.

#### **4.7.2 Additional Patient-Reported Outcomes Analyses**

Only patients in ITT population with a baseline assessment and at least one post-baseline assessment (PRO-evaluable subset) will be included in the following

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analyses. Missing post-baseline items and subscales will not be imputed and will be summarized as observed.

Compliance rates will be summarized by listing the number and proportion of patients in the PRO-evaluable subset who completed the PRO assessments at each timepoint by treatment arm. Reasons for non-completion will be summarized, if available, in the CRF.

All of the following analyses will be performed on all timepoints during the treatment period as well as at the time of disease progression per RECIST v1.1 (PRO assessment completed within  $\pm 7$  days of date of radiographic pharmacodynamic) at the last dose of treatment received before treatment discontinuation of any cause, at the 4-week post-treatment discontinuation visit, and at the 8-weekly survival follow-up visits through 6 months.

- EORTC scores and change from baseline will be descriptively analyzed using means, standard deviations, medians, and range, by treatment arm at baseline and each of the timepoints previously defined.
- The number and proportion of patients at each of the 4-response category reflecting no symptom/impairment to severe symptom/impairment will be documented at baseline.
- Graphs of the mean changes and standard errors over time from the baseline assessment of items and subscales will be provided for each treatment arm.
- The number and proportion of patients who improved, worsened, or remained stable compared with baseline score will be summarized for all item-score and subscales score of the EORTC QLQ-C30 questionnaire and the QLQ-LC13 by treatment arm at baseline and each of the timepoints previously defined. This analysis should be performed for the PRO-evaluable population as well as the subgroups of the patients with/without CNS metastases at baseline. A patient will be deemed improved at a timepoint if there is a  $\geq 10$ -point decrease from baseline for any symptom or a  $\geq 10$ -point increase from baseline for any functioning scales or global health status scale. A patient will be deemed worsened at that timepoint if they have a  $\geq 10$ -point increase from baseline for any symptom or a  $\geq 10$ -point decrease from baseline for any functioning scales or global health status. If patients do not fit the improved or worsened criteria, they will be defined as stable for that specific item/scale at that timepoint.

Additional analyses will be conducted specifically for the cognitive function scale per the EORTC QLQ-C30:

- The number and proportion (%) of patients with  $\geq 10$ -point decrease from baseline in the Cognitive Function Scale score per the EORTC QLQ-C30 in each treatment arm at (1) any point during the treatment or (2) at date of CNS Progression (PRO assessment completed within  $\pm 7$  days of date of CNS PD). This analysis will not include PRO data from the 4-week post-treatment discontinuation visit or the PRO data collected at the 8-weekly survival follow-up visits through 6 months.

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- A histogram of worst change scores from baseline (decline of 16.66 vs. 33.33 vs. 50 vs. 66.66 vs. 83.33 vs. 100 points from baseline) in the Cognitive Function Scale score in each treatment arm at any point on treatment. This analysis will not include PRO data from the 4-week post-treatment discontinuation visit or the PRO data collected at the 8-weekly survival follow-up visits through 6 months.

#### **4.8 EXPLORATORY ANALYSES**

Exploratory outcome measures defined in the protocol other than the Abbot Vysis FISH subgroup analysis (see Section 4.4.3) and onset of hypogonadism in adult men are outside of the scope of this SAP and will not be included in the CSR.

The objective to determine the correlation between ALK status as assessed by plasma ALK polymerase chain reaction and/or plasma ALK sequencing tests with ALK status obtained using the Ventana ALK IHC and FISH Vysis ALK Break Apart FISH Probe Kit (Abbott) is out of scope of this SAP and will be conducted outside of the CSR analyses.

#### **4.9 INTERIM ANALYSES**

No interim analysis for efficacy or futility is planned.

## 5. REFERENCES

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