Operating characteristics of tumor growth inhibitionoverall survival models to support early Phase Ib decisions:

An evaluation in first-line metastatic non-small cell lung cancer patients treated with atezolizumab plus chemotherapy based on the Phase III study IMpower150



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## Progress and Opportunities to Advance Clinical Cancer Therapeutics Using Tumor Dynamic Models

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# TGI-OS models predict atezolizumab vs. control HR in atezolizumab Phase II and III clinical studies



Model predictions (dots) and 95% prediction interval (1000 replicates, bars) with observed (squares)

Chan et al, CPT Pharmacometrics Syst Pharmacol. 2021;00:1–12

#### Background

- Typical tumor dynamic profiles following single agent or combination with a hypothetical agent that would reduce growth rate (KG) by 20%–40% and associated expected OS HR in randomized studies of the combination versus single agent
- We hypothesized that effect size in TGI metrics (relative to control) predicts for OS hazard ratio



Bruno et al, Clin Cancer Res, 2020;26,1785-1795

#### Atezolizumab IMpower150 study in 1<sup>st</sup>-line NSCLC

- Atezolizumab in combination with bevacizumab plus carboplatin-paclitaxel (Arm B: ABCP) significantly improved PFS and OS compared to the control treatment (Arm C: BCP) (Socinski, N Engl J Med 2018;378, 2288-301)
- Typical tumor dynamic profiles showed a clear separation too and estimated KG predicted study outcome (OS distributions and hazard ratio) (Yoshida, ACoP 2019)



In black: tumor dynamic endpoints that may support early decisions

#### Methods

- □ Resampled tumor data from IMpower150
  - N=40 patients, 6 months recruitment, 24 weeks follow up with control
- Assessed probability to detect the effect if true
  - **Resampled in Arm B (ABCP) and Arm C (control, BCP)**
  - Estimated effect size based on based on TGI metrics (biexponential TGI model, Claret, CCR 2018)
    - Declared Go if significant difference (two-sided Wilcoxon test)
  - Predicted OS HR using integrated NSCLC TGI-OS model (Chan, CPT-PSP 2021)
    Estimated effect size (HR)
    - □ Simulated a Phase III design (400 patients/arm resampled among the 40 patients)
    - Declared Go if significant difference (log-rank test)
  - **Correct go decision (power or sensitivity)** = % replicates with significant difference
- Assess probability to detect an effect if absent (false positive)
  - Resampled in Arm C vs. Arm C
  - Incorrect go decision (Type I error or specificity)
- □ Assessed **PFS** the same way

| Metric                   | Median<br>Effect size (%) | 95%CI          | Percent of success (%) <sup>1</sup> |
|--------------------------|---------------------------|----------------|-------------------------------------|
| KG (week <sup>-1</sup> ) | -23.0                     | [-52.1 – 34.9] | 73.8                                |
| KS (week <sup>-1</sup> ) | 6.7                       | [-37.0 – 75.0] | 36.4                                |
| TR24                     | -19.3                     | [-38.1 – 3.49] | 64.4                                |
| TTG (week)               | 18.8                      | [-24.9 - 87.2] | 69.4                                |

Power to detect a difference is fair except with KS, best with KG

| Metric                   | Median<br>Effect size (%) | 95%CI          | Percent of success (%) <sup>1</sup> |  |  |
|--------------------------|---------------------------|----------------|-------------------------------------|--|--|
| KG (week <sup>-1</sup> ) | 0.3                       | [-13.1 - 17.8] | 4.2                                 |  |  |
| KS (week <sup>-1</sup> ) | -0.4                      | [-22.7 - 27.8] | 4.8                                 |  |  |
| TR24                     | 0.1                       | [-17.4 - 21.5] | 4.4                                 |  |  |
| TTG (week)               | -0.4                      | [-15.5 - 14.9] | 3.0                                 |  |  |

Type I error is acceptable

<sup>1</sup>% replicates with significant difference, two-sided Wilcoxon at p<0.05



#### **OS** simulations



HR estimate quite uncertain

Power to detect a difference is fair in the Phase III simulations

<sup>1</sup>% replicates with significant difference, two-sided log-rank



#### Conclusions so far...

- Data selection process based on IMpower150 mimics a Phase Ib design with control
  - N=40 patients, 6 months recruitment, 24 weeks follow up with control
- "Success" has been defined when TGI metrics were significantly different in experimental vs. control arm
  - As expected KG was the most sensitive metric to predict "success" but TTG and TR24 have good sensitivity too and could offer interesting, easier to estimate alternatives to KG
  - Type I error was close to 5% specified by the test
- OS simulations predicted a HR of 0.82, Phase III probability of success was fair for a 400 patient per arm trial
- The observed PFS analysis based on the same resampled and truncated data had a 28.0% power to show a difference across Arms (2-sided log rank test at p=0.05)
- This evaluation suggests that model-based TGI metrics may be useful exploratory endpoints to inform early clinical decisions
- Alternative designs are being investigated
  - Less patients (N=20, 30)
  - Shorter follow up (3 months)
  - Single arm study (options to generate virtual control are explored, Marchand, ACoP 2017)
- Expansion of this work is ongoing in other settings

### Backups

#### IMpower150 study design





#### Selection of data

- Select only Arm B (Atezo + CP+Bev) and Arm C (CP+Bev) data
  - Select TGI evaluable patients *i.e.* at least baseline and one post-treatment
    SLD assessment: 91 % of treated patients
- Arm B:
  - Rank the patients by date of first dose in each arm
  - Randomly sample one patient in Arm B
  - Define the 10-month period after first dose
  - Sample 39 patients with replacement
- Arm C:
  - Sample with replacement 40 patients in the same 6-month period
- Check the date of the SLD assessments for the last patient (selected in arm B) #40 and select the last assessment before 24w+1w weeks after first dose
  - Cut the dataset for SLD assessment visit after this date for the whole dataset (Arm B and Arm C)





Tumor profile of the last patient selected (SLD up to ~24w)



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#### N 500 replicates

| TRT     | N replicates | Median FU<br>(weeks) | Min FU<br>(weeks) | Max FU<br>(weeks) | Median<br>Scan | Min Scan | Max Scan |
|---------|--------------|----------------------|-------------------|-------------------|----------------|----------|----------|
| Control | 500          | 23.7                 | 3.9               | 49.4              | 5              | 2        | 9        |
| Atezo   | 500          | 24.8                 | -3.9              | 50.6              | 5              | 1        | 10       |

In the analysis datasets (N=500), the median follow-up is 24 weeks in both atezo and control arms, with a maximum of 50 weeks (~1 years).

The patients had a median of 5 scans for the tumor assessment.



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- The N datasets are used to estimate TGI metrics using Stein TGI model (biexponential model).
- The individual TGI metrics (KG, KS, TTG, and TR24) are summarized by treatment arms and replicates.
- An effect size for each metrics and each replicate is derived as follow:
  - Metric Atezo Metric Control Metric Control \* 100
- A two-sided Wilcoxon test ( $\alpha$ =5%) is performed on the TGI metrics of the two arms
- P-values of the test are used to derive the percent of success of each replicate if p-value < 0.05 (consider doing < 0.10)
- The percent of success is summarized for each metric
- To investigate the type I error, the same approach is used with randomly selected patients from Arm C compared to patients in Arm C.



## TGI model estimates on one virtual analysis analysis dataset

| Parameter                | Estimate | SE      | RSE  | shrinkage |              |
|--------------------------|----------|---------|------|-----------|--------------|
| KG[Atezo]                | 0.00740  | 0.0016  | 21.9 | -         | Atezo+CP+Bev |
| KS[Atezo]                | 0.0704   | 0.0068  | 9.72 | -         |              |
| KG[control]              | 0.0137   | 0.0019  | 13.6 | -         | CP+Bev       |
| KS[control]              | 0.0675   | 0.0099  | 14.7 | -         |              |
| TS[0]                    | 67       | 4.9048  | 7.32 | -         |              |
| sigma^2                  | 42.1     | 11.1456 | 26.5 | -         |              |
| omega[KG, atezo]         | 0.611    | 0.2459  | 40.3 | 17.5      |              |
| omega[KS, atezo]         | 0.156    | 0.0469  | 30.1 | 23.4      |              |
| omega[KG, control]       | 0.261    | 0.1253  | 48   | 19.9      |              |
| omega[KS, control]       | 0.428    | 0.15    | 35   | 18.2      |              |
| omega[TS0]               | 0.41     | 0.0576  | 14.1 | 2.85      |              |
| omega[corrKGKS, atezo]   | -0.0706  | 0.0831  | 118  | -         |              |
| omega[corrKGKS, control] | -0.0443  | 0.1093  | 247  | -         |              |
| Obj                      | 2620     | 0       | 0    | -         |              |
|                          |          |         |      |           |              |

#### Derivation of the % change ArmB versus ArmC on the individual TGI metrics values

| Ν  | parameter   | ARMB    | ARMC   | %change |
|----|-------------|---------|--------|---------|
| 40 | median.KG   | 0.00864 | 0.0142 | -39.2   |
| 40 | median.KS   | 0.0729  | 0.0710 | 2.7     |
| 40 | median.TR24 | 0.44    | 0.651  | -32.4   |
| 40 | median.TTG  | 25.2    | 18.1   | 39.2    |

#### Derivation of the % change ArmB versus ArmC on the typical TGI metrics values

| parameter | ARMB    | ARMC   | %change |
|-----------|---------|--------|---------|
| KG        | 0.00740 | 0.0137 | -46.0   |
| KS        | 0.0704  | 0.0675 | 4.3     |



#### Methods for OS simulations

- Integrated NSCLC TGI-OS model (Chan, CPT-PSP 2021) has been reestimated without IMpower150 data (no big change, see next slide)
- This model was used to simulate OS as follows:
  - 500 replicates of 40 patients by arm (Arm B vs. Arm C or Arm C vs. Arm C) with individual covariates and individual estimated KG
- For each of these scenario and across the 500 replicates, the followings analyses were preformed:
  - KM of OS distribution by arm
  - ➢ HR 95%PI
  - Percent of success using log-rang test at p-values of 0.05, 0.1 and 0.15





| Pooled      |           |            |       |           | without IMpower150 |          |            |       |           |
|-------------|-----------|------------|-------|-----------|--------------------|----------|------------|-------|-----------|
|             | Value     | Std. Error | z     | р         |                    | Value    | Std. Error | z     | р         |
| (Intercept) | 3.47      | 0.173      | 20.1  | 8.41E-90  | (Intercept)        | 3.41     | 0.201      | 17    | 6.44E-65  |
| logKG       | -0.616    | 0.0224     | -27.4 | 1.35E-165 | logKG              | -0.621   | 0.0256     | -24.3 | 4.46E-130 |
| BCRP        | -0.00385  | 0.000348   | -11.1 | 1.75E-28  | BCRP               | -0.00374 | 0.000403   | -9.29 | 1.60E-20  |
| BECOG       | -0.233    | 0.0298     | -7.81 | 5.61E-15  | BECOG              | -0.179   | 0.0348     | -5.14 | 2.74E-07  |
| nsite5      | -0.0764   | 0.0139     | -5.51 | 3.53E-08  | nsite5             | -0.0723  | 0.0153     | -4.73 | 2.21E-06  |
| asian       | 0.244     | 0.0443     | 5.52  | 3.42E-08  | asian              | 0.194    | 0.0508     | 3.82  | 0.000135  |
| BALBUM      | 0.0135    | 0.00304    | 4.43  | 9.48E-06  | BALBUM             | 0.0135   | 0.0036     | 3.74  | 0.000184  |
| ICTC        | 0.119     | 0.0286     | 4.16  | 3.20E-05  | ICTC               | 0.131    | 0.0331     | 3.96  | 7.64E-05  |
| BLDH        | -0.000141 | 4.00E-05   | -3.53 | 0.00041   | BLDH               | -0.00012 | 4.17E-05   | -2.8  | 0.00515   |
| BNLR        | -0.009    | 0.00262    | -3.44 | 0.000582  | BNLR               | -0.0165  | 0.00379    | -4.36 | 1.27E-05  |
| line        | -0.109    | 0.0341     | -3.2  | 0.00138   | line               | -0.103   | 0.0357     | -2.88 | 0.00403   |
| mliver      | -0.118    | 0.0401     | -2.94 | 0.00332   | mliver             | -0.121   | 0.0452     | -2.68 | 0.00734   |
| Log(scale)  | -0.264    | 0.0161     | -16.3 | 4.95E-60  | Log(scale)         | -0.276   | 0.0185     | -14.9 | 2.82E-50  |

