

CORRESPONDENCE

Reply to letter comments on: Molecular targeted maintenance therapy versus standard of care in advanced biliary cancer: an international, randomised, controlled, open-label, phase III umbrella trial (SAFIR-ABC10-Precision Medicine)



Ruan and colleagues cite the example of the ClarIDHy trial¹ to highlight the risk of dilution, in the event of crossover upon disease progression, of any overall survival (OS) benefit of molecular targeted therapies (MTT) over first-line standard of care (1L-SoC; chemoimmunotherapy) in the SAFIR ABC10 trial.² In fact, in the ClarIDHy trial, 70% of patients in the placebo arm switched to the experimental treatment [the isocitrate dehydrogenase 1 (IDH1) inhibitor ivosidenib], so that the numerical OS benefit [a secondary endpoint, progression-free survival (PFS) having logically been chosen as the primary endpoint] did not reach statistical significance [median 10.3 versus 7.5 months, hazard ratio (HR) 0.79, 95% confidence interval (CI) 0.56-1.12, $P = 0.09$]. The context, however, is different for SAFIR ABC10. Although a crossover may be ethically justified, it is now unavoidable given the widespread availability of MTT, which all patients in the trial will potentially receive at some point. Therefore, we opted for a crossover within the trial rather than outside it, to ensure maximum standardization. Indeed, OS, as opposed to PFS, is sensitive to the confounding effects of post-study treatment, generally at the investigators' discretion. In addition to the intention-to-treat (ITT) analysis, we will adjust for the impact of crossover on OS using the rank-preserving structural failure time (RPSFT) approach.³ This approach is based on a common assumption: the treatment effect is the same for all patients, regardless of when treatment is received. Although not intended to provide a formal proof-of-treatment effect, the RPSFT method compares treatment groups as randomized, with results that have the same significance level as those of the ITT analysis. In ClarIDHy, The RPSFT-adjusted median OS was 5.1 months with placebo (HR 0.49, 95% CI 0.34-0.70, $P < 0.001$).¹ Correcting for crossover has been employed for decades in other phase III studies, and the RPSFT model has been recognized by health technology assessment bodies, e.g. the National Institute of Clinical Excellence in the UK.⁴

Second, we agree that the umbrella design aggregates a heterogeneous portfolio of MTTs, with reported objective response rates ranging from 2% with ivosidenib to up to 50% with fibroblast growth factor receptor (FGFR), human epidermal growth factor receptor 2 (HER2) and v-Raf murine sarcoma viral oncogene homolog B/mitogen-activated protein kinase kinase inhibitors. Our statistical analysis plan

includes subgroup analyses stratified by alteration and Bayesian hierarchical models to account for heterogeneity.

Lastly, Ruan and colleagues emphasize the value of circulating tumour DNA (ctDNA) analysis for elucidating the primary or secondary mechanisms of resistance to chemoimmunotherapy and MTT. In fact, we have planned ctDNA analysis at study entry, to assist in tumour molecular profiling, and to compare tissue versus 'liquid' molecular profiling (all patients); after four cycles of 1L-SoC (all patients); after three cycles of treatment in randomized patients (MTT, including after crossover, or continuation of 1L-SoC); and at first disease progression in randomized patients (including after crossover) and non-randomized patients (including early dropouts, before the fourth cycle of 1L-SoC). Of note, randomization is based on baseline alterations, and will occur before progression develops, thus limiting the risk of variation of resistance mechanisms, although this is unlikely with 1L-SoC chemoimmunotherapy.⁵

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