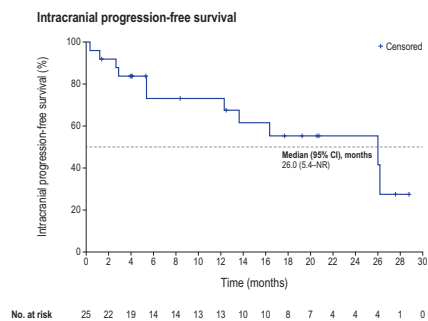


## Erratum

We have discovered several errors in our paper. First, there is an error in the published version in Fig 3 of the supplementary data. We have included the correct figure here. The number of EGFR driver mutation detection at screening should be 63, which consistent with the description of the article content. There is also a minor correction to Figure 1C which is also presented in the erratum.



**Figure 1.**