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24th November 2018

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3 Prof. Rajiv Jalan
4 Editor-in-Chief
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6 Journal of Hepatology
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10 Dear Prof. Jalan,
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14 On behalf of all co-authors, we would like to submit a reply letter entitled “Reply to
15 Correspondence Concerning Models For Prediction Of Early Recurrence of HCC After Surgical
16 Resection Of Hepatocellular Carcinoma” to respond *four* Letters to the Editors commenting our
17 recent paper concerning early recurrence after resection of Hepatocellular Carcinoma (HCC) (J
18 Hepatol. 2018 Dec;69(6):1284-1293. doi: 10.1016/j.jhep.2018.08.027). Those four letters include
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- 21 1. “Time to Recurrence, But Not Recurrence-free Survival, Should Be the Endpoint of
22 Predicting Early Recurrence after HCC Resection” (JHEPAT-D-18-01944) by Wen-Tao
23 Yan et al.
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- 25 2. “Predicting early hepatocellular carcinoma recurrence after resection: a comment for moving
26 forward” (JHEPAT-D-18-01803R1) by Xiao-Ying Zhang et al.
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- 28 3. “Toward the universal scoring system in treatment for patients with hepatocellular
29 carcinoma” (JHEPAT-D-18-01939) by Jiro Kusakabe et al.
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- 31 4. “Factors Predicting Early Recurrence After Surgical Resection of Hepatocellular Carcinoma”
32 (JHEPAT-D-18-01989) by Yao-Ming Zhang et al.
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41 Thank you for your attention.
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44 Yours sincerely,
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48 Philip J Johnson, MD, FRCP
49 Professor in Translational Oncology
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**Reply to Correspondence Concerning Models For Prediction Of Early Recurrence of
HCC After Surgical Resection Of Hepatocellular Carcinoma**

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1 We are grateful to your correspondents [1-4] for their interest in, and comment on, our recent
2 paper concerning early recurrence after resection of Hepatocellular Carcinoma (HCC).[5]
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7 As Royston et al., have pointed out ‘Surprisingly, there is no widely agreed approach to
8 building a multivariable prognostic model from a set of candidate predictors’.[6] In building
9 our prognostic models we have two guiding principles. Firstly, those variables involved
10 should be readily available in all centers treating patients with HCC. Secondly, again as
11 emphasized by Royston et al., the models should be ‘transportable’ ‘i.e. broadly, if not
12 universally, applicable’. To achieve these criteria, we need large cohorts from multiple
13 centers, worldwide. With these two principles in mind we had to be pragmatic both in the
14 variables used to develop the models and the datasets available to us for analysis. Thus, the
15 price to pay, in most cases, for the large numbers is that we have to accept those cohorts that
16 have already been collected. We are not in a position to specify exactly what variables are in
17 the cohort, and we acknowledge that this not ideal. We are pursuing prospective studies to
18 validate our models and collect other relevant variables not included in the original study.
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38 *The variables entering the final models*

39 Like Zhang X.Y. et al., and Zhang Y.M. et al.,[3, 4] we found it surprising that macrovascular
40 invasion (MaVI), a well-described risk factor for early recurrence after resection, was not in
41 our final models. Had we considered MaVI in isolation (i.e. just in a univariate analysis),
42 MaVI was a risk factor (HR 3.306, P<0.001). However, it was not an *independent* predictor in
43 the multivariable analyses, a finding in agreement with other authors.[7, 8] When all
44 identified risk factors are considered in a multivariable model, the presence or absence of
45 MaVI does not add to the predictive utility of our model. Indeed, if we rebuild the model but
46 exclude any one of the significant parameters (male gender, multiple tumors, tumor size or
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AFP, except ALBI) then MaVI *does* enter the model.

Zhang Y.M. et al. questioned the high figure for microvascular invasion (MiVI) in the US series.[4] In our study we assumed that all patients with MaVI also had MiVI. We have confirmed the figures in the US series. The rate of MiVI in resected specimen was 44.6% (similar to that in Italy, 49.5%) and 28.6% for MaVI, giving the figure reported in Table 1 of 73.1%.

With regards to the inclusion of gender into our models,[4] we can only report what the analysis showed and recognize that multivariable models only relate to statistical associations and offer no insight into the underlying mechanism of the factors identified. In fact, Shim et al. also demonstrated that male gender was an independent risk factor associated with early recurrence in their cohort of 1085 patients.[8] Although it is unclear why male patients are prone to early recurrence, they may have more aggressive HCCs than female patients.[9]

Kusakabe et al. expressed concern that certain variables were excluded.[1] We have examined the effect of excluding the variables ALT, ALP, INR and positive margin on the performance of the ERASL-pre and ERASL-post models. Although the excluded variables modestly improved the fit of the model (LR test, $P=0.0121$ and $P=0.0102$), there was no further improvement in the survival prediction, as measured by Harrell's C index ($P=0.064$ and $P=0.060$). There was also no added enhancement in survival prediction when both models were rebuilt with the excluded variables added back in ($P=0.107$ and $P=0.161$).

All the variables in our study refer to the pre-operative assessment apart from MiVI which was only available after histological examination of the resected tumor and is only involved

1 in the post-operative model. We acknowledge that more accurate information about tumor
2 size, number and MaVI might be forthcoming after visual inspection of the liver at the time of
3 operation, but doubt that any changes will have a major impact on the model performance.
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7 8 9 *Issues surrounding the ALBI score*

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11 The ALBI score is an evidence-based and extensively validated refinement/simplification of
12 the Child-Pugh score (CPS).[10, 11] Its performance is at least as good or, more often, better
13 than the CPS.[12] Yan et al.,[2] expressed surprise that ALBI, a measure of liver function,
14 entered our early recurrence models since liver function is usually considered to be a factor
15 influencing *long term survival* after HCC resection and indeed we have previously confirmed
16 this in relation to ALBI.[13] Hsu et al., also reported, in a study of over 1900 patients
17 undergoing resection, that ALBI was an independent factor associated with early
18 recurrence.[14] Why this should be so must remain speculative, but one possibility is that a
19 fraction of ‘early recurrences’ may actually be ‘de novo’ tumor. Such cases might be expected
20 to occur more frequently in patients with poor liver function in more advanced underlying
21 chronic liver disease.
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41 We acknowledge that, in principle, it is preferable to consider continuous variables in their
42 continuous form rather than after categorization. In the event however, replacing the ALBI
43 grade with the ALBI score did not improve the survival prediction of the model: ERASL-
44 pre,(P=0.716 and P=0.422 respectively) and ERASL-post,(P=0.644 and P=0.913 respectively).
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53 Again, a case for applying albumin and bilirubin as separate parameters in developing our
54 models can be made. However, such an approach risks losing the conceptual basis of the
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1 score as a measure of liver function and the same issue would arise in applying any composite
2 score (e.g. CPS or tumor stage) in a multivariable model.
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6 7 ***Issues concerning early deaths*** 8

9 A small percentage of patients (<10%) died before their recurrence status was known. In our
10 study, that aimed to predict recurrence within the first two years of resection, we censored all
11 deaths within this period. We believe this was the correct procedure but acknowledge, as Yan
12 et al.,[2] point out, that this approach would be more appropriately labelled as ‘time to
13 recurrence’ rather than ‘recurrence free survival’.
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24 ***Might the models suggest which sub-group of HCC patients would benefit from*** 25 ***transplantation rather than resection?*** 26 27

28 As noted by Kusakabe et al.,[1] it is likely that at least those in the high risk ERASL group,
29 would do better with transplantation although it will be difficult to design a study to formally
30 predict the optimal potentially curative treatment. We have already shown that the ‘cure
31 fraction’ after transplantation for HCC is 74.1% compared to only 24.1% in the cases
32 undergoing resection.[15] In the near future we hope to be able to add a figure for ‘likelihood
33 of cure’ at the individual patient level, to the ERASL model.
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46 **Conclusion** 47

48 It is inevitable that different researchers using different datasets and different statistical
49 methodologies will arrive at different prognostic models. Like Kusakabe et al.,[1] we are
50 aiming to develop an overarching model for HCC. To this end we have collected a database
51 of around 150,000 cases with detailed clinical data and are starting to apply machine learning
52 and artificial intelligence in addition to conventional statistical methodology. The only way to
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1 arrive at a comprehensive and consistent model will be through large collaborative projects
2 involving high quality datasets, preferably prospectively-collected. We hope the present study,
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4 and the lessons learnt therefrom, will be a significant step in the right direction.
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10 11 **References**

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