

Etra A, Gergoudis S, Morales G, et al. Assessment of systemic and gastrointestinal tissue damage biomarkers for GVHD risk stratification. *Blood Adv.* 2022;6(12):3707-3715.

Page 3710: Due to formatting errors, Table 2 does not correctly delineate the individual biomarker models from each other. "Single Biomarker Model" should show 5 individual biomarker models, not one 5-biomarker grouping. "Two Biomarker Model" should show six 2-biomarker models, not one 12-biomarker grouping. "Three Biomarker Model" should show four 3-biomarker models, not one 12-biomarker grouping. Also, "REG3" should read "REG3 $\alpha$ ." The corrected Table 2 is shown below:

**Table 2. Training cohort biomarker algorithms and their respective *p* values**

	<i>P</i> value
<b>Single Biomarker Model (threshold <i>p</i> &lt; .3)</b>	
<b>TNFR1</b>	<b>&lt;.001</b>
<b>TIM3</b>	<b>.250</b>
IL6	.630
<b>ST2</b>	<b>&lt;.001</b>
<b>REG3<math>\alpha</math></b>	<b>&lt;.001</b>
<b>Two Biomarker Model (threshold <i>p</i> &lt; .15)</b>	
<b>TNFR1</b>	<b>&lt;.001</b>
<b>TIM3</b>	<b>.140</b>
<b>TNFR1</b>	<b>.017</b>
<b>ST2</b>	<b>.007</b>
TIM3	.940
ST2	<.001
<b>TNFR1</b>	<b>.014</b>
<b>REG3<math>\alpha</math></b>	<b>&lt;.001</b>
TIM3	.910
REG3 $\alpha$	<.001
<b>ST2</b>	<b>.003</b>
<b>REG3<math>\alpha</math></b>	<b>&lt;.001</b>
<b>Three Biomarker Model (threshold <i>p</i> &lt; .05)</b>	
TIM3	.530
ST2	.003
REG3 $\alpha$	<.001
TNFR1	.004
TIM3	.11
ST2	.005
TNFR1	.2
ST2	.029
REG3 $\alpha$	<.001
TNFR1	.005
TIM3	.2
REG3 $\alpha$	<.001
<b>Four Biomarker Model (threshold <i>p</i> &lt; .05)</b>	
TNFR1	.078
TIM3	.13
ST2	.021
REG3 $\alpha$	<.001

Biomarkers meeting the statistical criteria for further study are shown in bold.

The errors have been corrected in the published article.

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