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 "REG3α." The corrected Table 2 is shown below:

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

	P value
Single Biomarker Model (threshold p < .3)	
TNFR1	**<.001**
TIM3	**.250**
IL6	.630
ST2	**<.001**
REG3α	**<.001**
Two Biomarker Model (threshold p < .15)	
TNFR1	**<.001**
TIM3	**.140**
TNFR1	**.017**
ST2	**.007**
TIM3	.940
ST2	<.001
TNFR1	**.014**
REG3α	**<.001**
TIM3	.910
REG3α	<.001
ST2	**.003**
REG3α	**<.001**
Three Biomarker Model (threshold p < .05)	
TIM3	.530
ST2	.003
REG3α	<.001
TNFR1	.004
TIM3	.11
ST2	.005
TNFR1	.2
ST2	.029
REG3α	<.001
TNFR1	.005
TIM3	.2
REG3α	<.001
Four Biomarker Model (threshold p < .05)	
TNFR1	.078
TIM3	.13
ST2	.021
REG3α	<.001

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

The errors have been corrected in the published article.