



CORRECTION

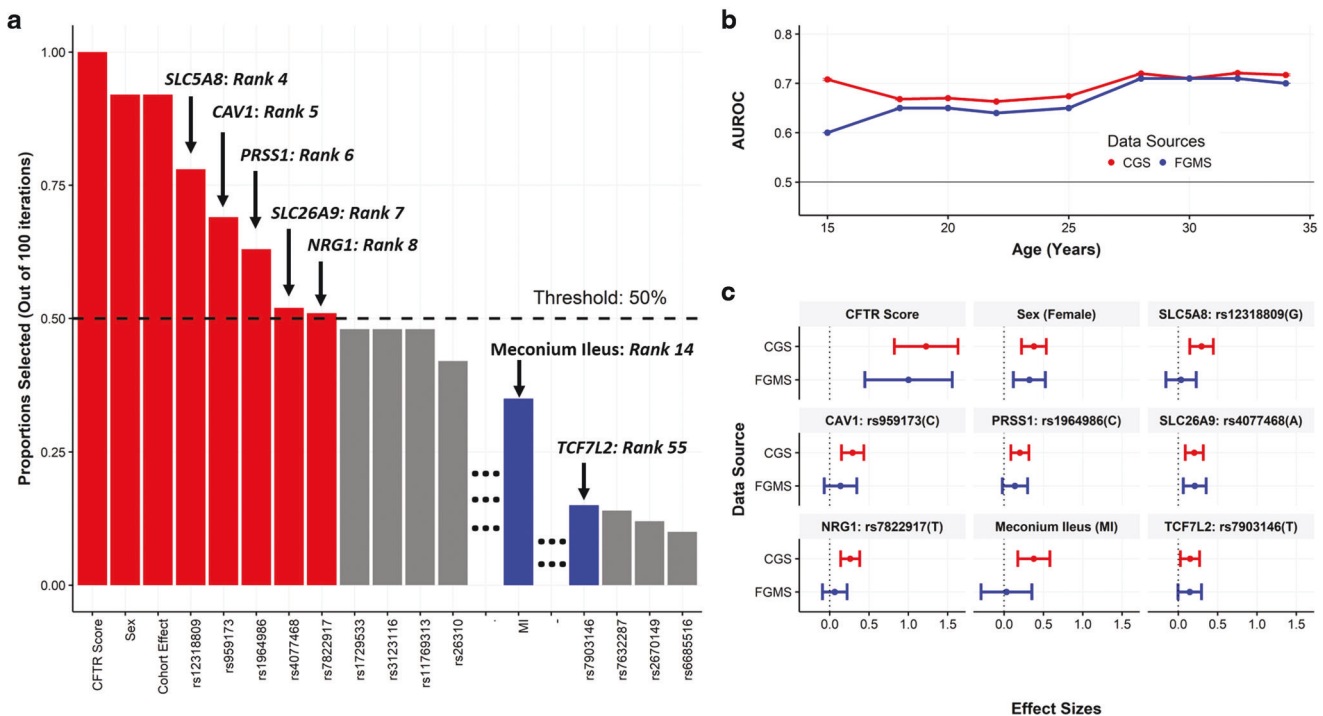
Correction to: Cystic fibrosis–related diabetes onset can be predicted using biomarkers measured at birth

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The risk alleles of rs1964986 (PRSS1) and rs959173 (CAV1) should be the C allele for both variants instead of the A and T alleles listed in the paper. The changes have been reflected in both Figure 1 and Table 2 shown below.



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The original article has been corrected.

Table 2. Effect sizes (hazard ratios) and the 95% confidence intervals (CIs) fitted using a multivariate Cox PH model in the CGS. Risk allele/risk group noted in parentheses after the listed predictor.

Gene annotation	Predictor	Hazard ratio	95% CI
<i>CFTR</i>	<i>CFTR</i> mutation score	3.02	(2.01, 4.54)
—	Sex (female)	1.48	(1.26, 1.74)
<i>SLC5A8</i>	rs12318809 (G)	1.35	(1.16, 1.57)
<i>CAV1</i>	rs959173 (C)	1.27	(1.10, 1.47)
<i>PRSS1</i>	rs1964986 (C)	1.23	(1.09, 1.38)
<i>SLC26A9</i>	rs4077468 (A)	1.20	(1.07, 1.34)
<i>NRG1</i>	rs7822917 (T)	1.31	(1.16, 1.48)
—	Meconium ileus (MI)	1.29	(1.05, 1.59)
<i>TCF7L2</i>	rs7903146 (T)	1.18	(1.05, 1.34)



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