

Correction

# Correction: Expression and Replication Studies to Identify New Candidate Genes Involved in Normal Hearing Function

The PLOS ONE Staff

Table 4 has been corrected for orientation and readability. Please see the corrected Table 4 here:

**Table 4.** Results of the replication association study for the candidate SNPs of expressed genes.

Gene	Top SNP	Eff. All	Meta-analysis Beta±SE*	Repl. Beta±SE	Repl. p-value	Maf	Trait	Str	SNP type	Info Score	Conc	HWE p-value	Repl. Model
CSMD1	rs10091102	C	0.08 ± 0.03	0.04 ± 0.02	0.022	0.25	PC1	+	imp	0.98106	96.2	-	add
ANK2	rs17045059	G	0.25 ± 0.06	0.08 ± 0.03	0.010	0.30	4 kHz	+	imp	0.92767	97.2	-	add
CDH13	rs17195859	A	0.27 ± 0.06	0.07 ± 0.03	0.024	0.15	6 kHz	+	gen	-	95.9	0.37	dom
DCLK1	rs9574464	G	-0.17 ± 0.03	-0.06 ± 0.02	0.008	0.32	0.25 kHz	+	imp	0.95981	97.3	-	dom

doi:10.1371/journal.pone.0085352.t004

**Citation:** The PLOS ONE Staff (2014) Correction: Expression and Replication Studies to Identify New Candidate Genes Involved in Normal Hearing Function. PLoS ONE 9(3): e91446. doi:10.1371/journal.pone.0091446

**Published** March 7, 2014

**Copyright:** © 2014 The PLOS ONE Staff. This is an open-access article distributed under the terms of the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.