

CORRECTION

# Correction: Exome Sequencing of Phenotypic Extremes Identifies *CAV2* and *TMC6* as Interacting Modifiers of Chronic *Pseudomonas aeruginosa* Infection in Cystic Fibrosis

Mary J. Emond, Tin Louie, Julia Emerson, Jessica X. Chong, Rasika A. Mathias, Michael R. Knowles, Mark J. Rieder, Holly K. Tabor, Debbie A. Nickerson, Kathleen C. Barnes, NHLBI GO Exome Sequencing Project, Lung GO, Ronald L. Gibson, Michael J. Bamshad

In the Funding section, two of the grant numbers from the funder Cystic Fibrosis Foundation are listed incorrectly: OBSERV04KO should be OBSERV13K0, and BAMSHA14P should be BAMSHA14P0.

## Reference

1. Emond MJ, Louie T, Emerson J, Chong JX, Mathias RA, Knowles MR, et al. (2015) Exome Sequencing of Phenotypic Extremes Identifies *CAV2* and *TMC6* as Interacting Modifiers of Chronic *Pseudomonas aeruginosa* Infection in Cystic Fibrosis. *PLoS Genet* 11(6): e1005273. doi: [10.1371/journal.pgen.1005273](https://doi.org/10.1371/journal.pgen.1005273) PMID: [26047157](https://pubmed.ncbi.nlm.nih.gov/26047157/)



## OPEN ACCESS

**Citation:** Emond MJ, Louie T, Emerson J, Chong JX, Mathias RA, Knowles MR, et al. (2015) Correction: Exome Sequencing of Phenotypic Extremes Identifies *CAV2* and *TMC6* as Interacting Modifiers of Chronic *Pseudomonas aeruginosa* Infection in Cystic Fibrosis. *PLoS Genet* 11(8): e1005424. doi:10.1371/journal.pgen.1005424

**Published:** August 18, 2015

**Copyright:** © 2015 Emond et al. This is an open access article distributed under the terms of the [Creative Commons Attribution License](https://creativecommons.org/licenses/by/4.0/), which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.