Erratum for Baddal et al., Dual RNA-seq of Nontypeable *Haemophilus influenzae* and Host Cell Transcriptomes Reveals Novel Insights into Host-Pathogen Cross Talk

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Volume 6, no. 6, doi:10.1128/mBio.01765-15, 2015. The transcriptomic data sets relative to host and nontypeable *Haemophilus influenzae* (NTHi) response were omitted from the supplemental material. They have been added to the supplemental material online. Gene modulation is expressed as log$_2$ fold change at specific time points compared to the reference condition. The revised supplemental list should appear as below:

**Supplemental Material**

Supplemental material for this article may be found at [http://mbio.asm.org/lookup/suppl/doi:10.1128/mBio.01765-15/-/DCSupplemental](http://mbio.asm.org/lookup/suppl/doi:10.1128/mBio.01765-15/-/DCSupplemental).

Figure S1, PDF file, 0.8 MB.
Figure S2, PDF file, 0.5 MB.
Figure S3, PDF file, 0.4 MB.
Figure S4, PDF file, 0.4 MB.
Figure S5, PDF file, 0.4 MB.
Table S1, PDF file, 0.1 MB.
Table S2, PDF file, 0.2 MB.
Table S3, PDF file, 0.2 MB.
Data Set S1, XLSX file, 0.2 MB.
Data Set S2, XLSX file, 0.2 MB.

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