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## Article:

Pettit, L.J., Browne, H.P., Yu, L. et al. (10 more authors) (2014) Functional genomics reveals that Clostridium difficile SpoOA coordinates sporulation, virulence and metabolism. BMC Genomics, 15. 160-160. ISSN 1471-2164
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Additional file 1: Identification of differentially expressed genes in C. difficile 630 spo0A mutant by transcriptional profiling. Scatter plot of the log2 fold changes against the normalised mean read abundance per gene (calculated at the base level). Red dots represent genes considered to be significantly differentially expressed ( $\mathrm{P}=\leq 0.01$ ). Black dots signify genes not deemed to be significantly differentially expressed according to these criteria.

