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Identification of Human Lineage-Specific Transcriptional Coregulators Enabled by a Glossary of Binding Modules and Tunable Genomic Backgrounds

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Due to a Production error, an incorrect version of Table S4 was originally published online with this article. The AUROC values within the table were correct, but some were attributed to the wrong ChIP-seq dataset as several headings had been erroneously swapped. The correct version of Table S4 is now available online. We regret this error and apologize for any confusion that it has caused.

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