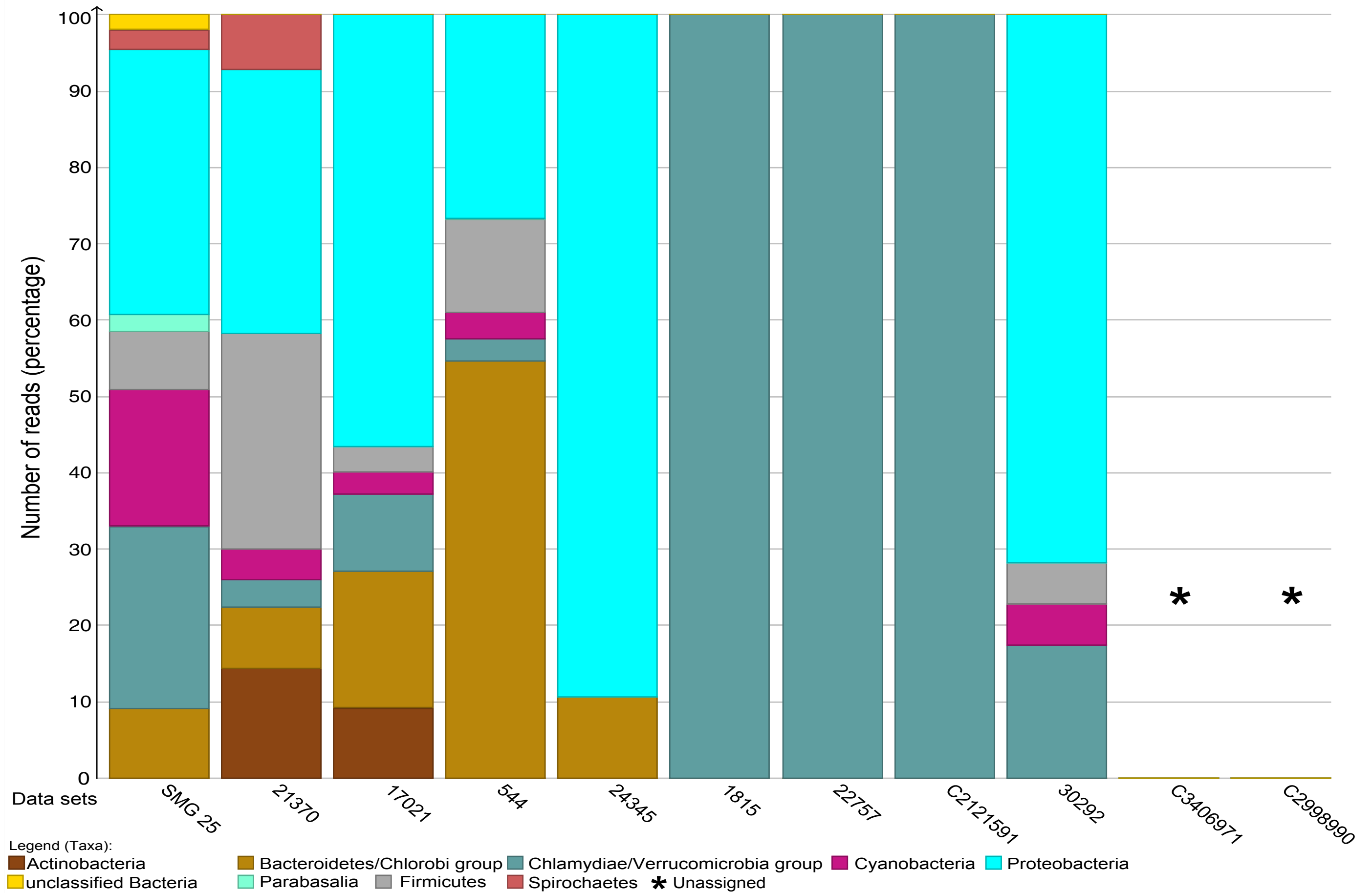


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<b>Author(s)</b>	Culligan, Eamonn P.; Sleator, Roy D.; Marchesi, Julian R.; Hill, Colin
<b>Publication date</b>	2013
<b>Original citation</b>	Culligan EP, Sleator RD, Marchesi JR, Hill C (2013) Functional Environmental Screening of a Metagenomic Library Identifies stlA; A Unique Salt Tolerance Locus from the Human Gut Microbiome . PLoS ONE 8(12): e82985. doi:10.1371/journal.pone.0082985
<b>Type of publication</b>	Article (peer-reviewed)
<b>Link to publisher's version</b>	<a href="http://dx.doi.org/10.1371/journal.pone.0082985">http://dx.doi.org/10.1371/journal.pone.0082985</a> Access to the full text of the published version may require a subscription.
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**Figure S3. Taxonomic assignment of scaffold sequences from Human Microbiome Project on which an *stIA* homologue was found.**



**Figure S3. Taxonomic assignment of scaffold sequences from Human Microbiome Project on which an *stIA* homologue was found.**

Scaffold sequences were analysed using BLASTX. The BLASTX results were then downloaded and imported in MEGAN 4 software program which performed taxonomic assignment of each scaffold based on BLAST reads. Two of the shorter scaffolds indicated with an asterisk (\*) could not be assigned any taxonomic classification.