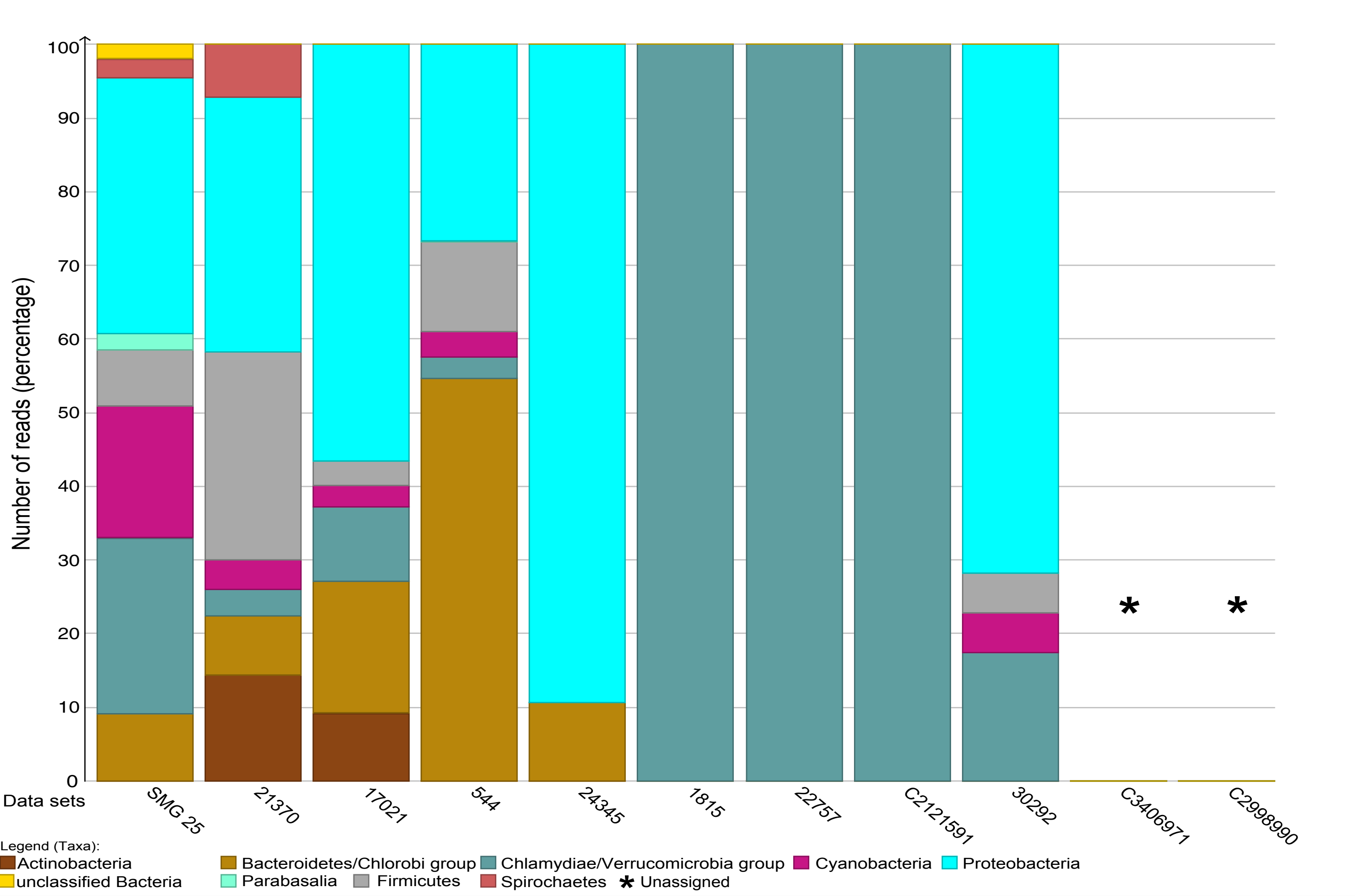


Title	Functional environmental screening of a metagenomic library identifies stlA; a unique salt tolerance locus from the human gut microbiome
Authors	Culligan, Eamonn P.;Sleator, Roy D.;Marchesi, Julian R.;Hill, Colin
Publication date	2013
Original Citation	Culligan EP, Sleator RD, Marchesi JR, Hill 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
Type of publication	Article (peer-reviewed)
Link to publisher's version	10.1371/journal.pone.0082985
Rights	© 2015 Culligan et al. This is an open access article distributed under the terms of the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited - <a href="http://creativecommons.org/licenses/by/4.0/">http://creativecommons.org/licenses/by/4.0/</a>
Download date	2024-03-29 15:28:02
Item downloaded from	<a href="https://hdl.handle.net/10468/2350">https://hdl.handle.net/10468/2350</a>

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



**Figure S3. Taxonomic assignment of scaffold sequences from Human Microbiome Project on which an *stlA* 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.