

CORRECTION

Correction: Functional implications of hexameric assembly of RraA proteins from *Vibrio vulnificus*

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The images for Figs $\underline{4}$ and $\underline{5}$ are incorrectly switched. The image that appears as Fig $\underline{4}$ should be Fig $\underline{5}$, and the image that appears as Fig $\underline{5}$ should be Fig $\underline{4}$. The figure captions appear in the correct order. Please see the corrected Fig $\underline{4}$ and Fig $\underline{5}$ below.



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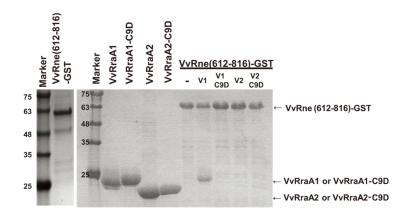


Fig 4. Interactions of VvRNase E with VvRraA proteins. Hexahistidine-tagged VvRraA1, VvRraA1-C9D, VvRraA2, VvRraA2-C9D, and the GST-fused VvRne (612–816 residues) were expressed and purified as described in the Methods section. The GST-fused VvRne protein was bound to GSH resin and incubated with VvRraA proteins and their C9D mutant proteins. Then, the proteins were eluted and the fractions were analyzed using SDS-PAGE. The protein bands were stained with Coomassie blue. Only VvRraA1 could tightly bind to VvRne.

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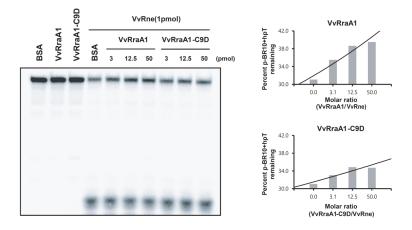


Fig 5. Inhibition of VvRraA1 and VvRraA1-C9D on the cleavage of p-BR10+hpT by VvRNase E in vitro. 0.5 pmol of 5'-end-labeled p-BR10+hpT RNA was incubated with 1 pmol of VvRne with varying concentrations of VvRraA1 and VvRraA1-C9D, 50 pmol of VvRraA1, or 50 pmol of BSA in $20~\mu$ l of $1~\times$ cleavage buffer at 37~C for 2~h for VvRne, VvRraA1 only, or BSA only controls. Samples were mixed with an equal volume of loading buffer, and then denatured at 65~C for 5~m in and loaded onto a 12% polyacrylamide gel containing 8~M urea. The percentage of uncleaved p-BR10+hpT in the gel was quantitated using a phosphorimager and OptiQuant software.

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Reference

 Song S, Hong S, Jang J, Yeom J- H, Park N, Lee J, et al. (2017) Functional implications of hexameric assembly of RraA proteins from Vibrio vulnificus. PLoS ONE 12(12): e0190064. https://doi.org/10. 1371/journal.pone.0190064 PMID: 29261778