

**Alternative sigma factor RpoX is a part of RpoE regulon and plays distinct roles  
in stress response, motility, biofilm formation and hemolytic activities in the  
marine pathogen *Vibrio alginolyticus***

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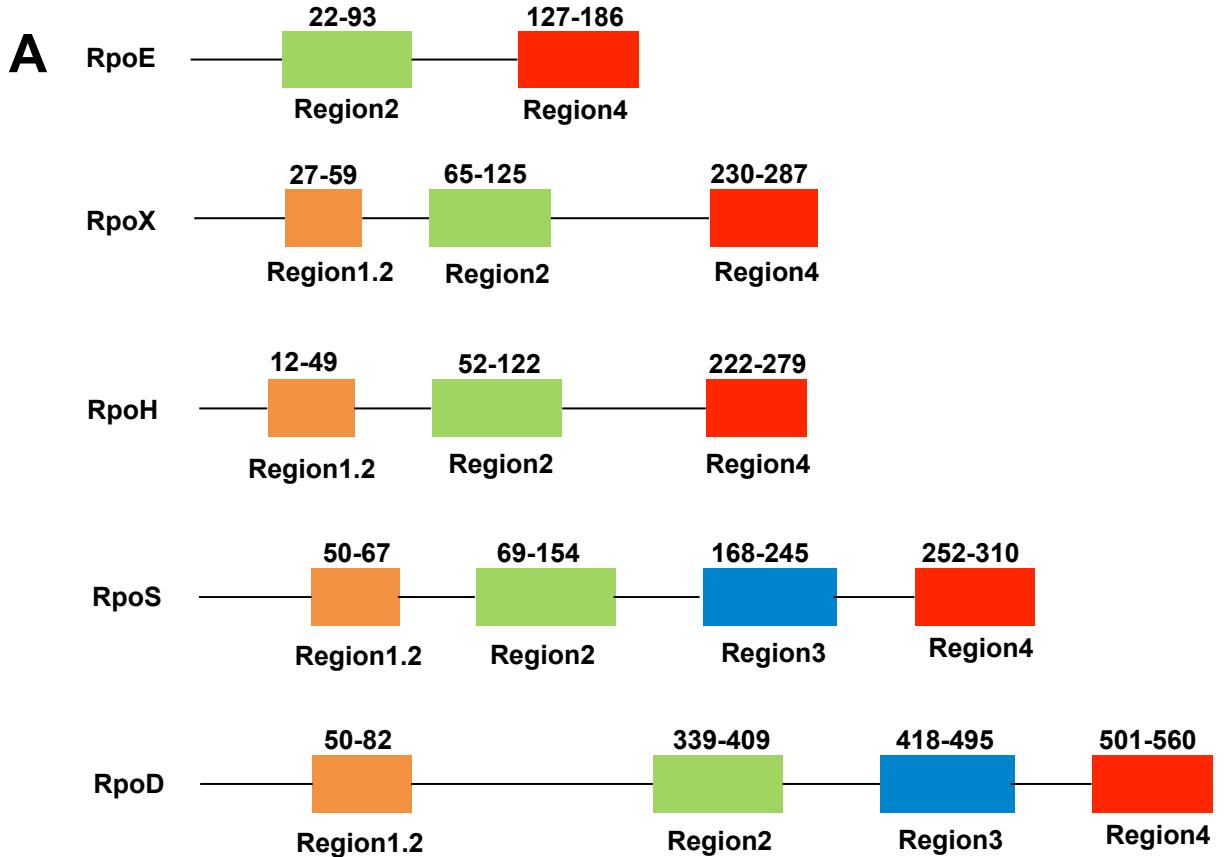
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**Running title:** RpoX regulates stress adaptation and pathogenesis

## Legends for supplemental figure

**FIGURE S1.** Genetic analysis of *rpoX* in vibrios. **(A)** Comparison of the conserved domains between RpoD and the alternative sigma factors RpoE, RpoX, RpoH, and RpoS. The numbers indicate the related amino acid residues. **(B)** Alignment of the amino acid sequence of *V. alginolyticus* RpoX (VARpoX) with the protein sequences of RpoX from *V. parahaemolyticus* (VPRpoX), *V. harveyi* (VHRpoX), and *V. splendidus* (VSRpoX). The identical and highly conserved amino acids in the sequences are highlighted in black and gray, respectively.



**B**

VARpoX :	HHNNNECFAERERAVTEDELSMKESLIVV-E-KDPYAHYLKDIVGIDLLSREEDELHYAKLNRA	GCIDEKARE-LIE8NLRLVVVKIA	NCY : 80
VPRpoX :	HHNNNECFAERERAVTEDELSMKESLIVV-E-KDPYAHYLKDIVGIDLLSREEDELHYAKLNRA	GCIDEKARE-LIE8NLRLVVVKIA	NCY : 80
VHRpoX :	-----MKESLIVV-E-KDPYAHYLKDIVGIDLLSREEDELHYAKLNRA	GCIDEKARE-SMIE8NLRLVVVKIA	NCY : 63
VSRpoX :	HHNNNDSEAAEQGETDDTPIKESFPTT-AE-DYAHYLKDIVGIDLLSREDEVYYAKLNRA	GCIDEKARE-SMIE8NLRLVVVKIA	NCY : 80
	64e s g kdpYahYLKdIVgidiLLJrE E6 Ya4Ln4aGD AR 6IE8NLRLVVK6AK Y		
VARpoX :	LLKRPQHNFTLLDLIEEGNIGLIKA	DVKDFP-PGYRFSTYAVWWIRENIEAALMNN	GRTVRLPAHVCKEINSLALKRHDASK : 161
VPRpoX :	LLKRPQHNFTLLDLIEEGNIGLIKA	DVKDFP-PGYRFSTYAVWWIRENIEAALMNN	GRTVRLPAHVCKEINSLALKRHDASK : 161
VHRpoX :	LLKRPQHNFTLLDLIEEGNIGLIKAIDKYDPE	DVKDFP-PGYRFSTYAVWWIRENIEAALMNN	GRTVRLPAHVCKEINSLALKRHDASK : 144
VSRpoX :	LLKRPQHNFTLLDLIEEGNIGLIKAIDKYDPE	DVKDFP-PGYRFSTYAVWWIRENIEAALMNN	GRTVRLPAHVCKEINSLALKRHDASK : 161
	1KR gnift6LD6IEEGN6GLIKA6DK51PepGYRFSTYAVWWIRENIEaAlmN	gRTVRLPaH6cKE6N3la kk d SK	
VARpoX :	SEKKEISIISSEL8RCGVRABRIDELVAL-I	-SGEIDV8ITIADHRVVEVLELCSESEYTD	PPODCENOFFTQAINEIIKTLA : 241
VPRpoX :	SEKKEISIISSEL8RCGVRABRIDELVAL-I	-SGEIDV8ITIADHRVVEVLELCSESEYTD	PPODCENOFFTQAINEIIKTLA : 241
VHRpoX :	SEKKEISIISSEL8RCGVRABRIDELVAL-I	-SGEIDV8ITIADHRVVEVLELCSESEYTD	PPODCENOFFTQAINEIIKTLA : 224
VSRpoX :	KMSEKKEISIISSEL8DSGVRABRIDELVAL-I	-SGEIDV8ITIADHRVVEVLELCSESEYTD	PPODCENOFFTQAINEIIKTLA : 241
	84e S1se6S sgvk 46d 66al sgf6dv 3t	P vle cesey dPq dce ftqag6e 6 3lp	
VARpoX :	KLQVVLIIHRPGFLPQDVWVKTLSIGLU--	-INVSNERVRQMCIEAVWRVQ---	KRLRFDGKV----- : 296
VPRpoX :	KLQVVLIIHRPGFLPQDVWVKTLSIGLU--	-INVSNERVRQMCIEAVWRVQ---	KRLRFDGKV----- : 296
VHRpoX :	KLQVVLIIHRPGFLPQDVWVKTLSIGLU--	-INVSNERVRQMCIEAVWRVQ---	KRLRFDGKV----- : 279
VSRpoX :	KLQVVLIIHRPGFLPQDVWVKTLSIGLU--	-INVSNERVRQMCIEAVWRVQ---	KRLRFDGKV----- : 298
	41q 66lhR5G1F VKTL e6G	6168nERVRQ6Q Ea6e46q	K41 fDgw6

Fig. S1