

A

Uniprot NO.	Ordered Locus Names	Domains	AA	Protein MW (KD)
Q9KRD2	VC1710	EAL	818	92
Q9KYR6	VC0137	EAL	432	51
Q9KN64	VCA0101	EAL	518	59
Q9KSP5	VC1211	EAL	704	79
Q9KRP4	VC1592	EAL	284	32
Q9KQZ6	VC1851	EAL HDOD	404	46
Q9KMV8	VCA0210	HD-GYP	522	60
Q9KPM7	VC2340	HD-GYP	384	44
Q9KSB1	VC1348	HD-GYP	441	49
Q9KU35	VC0694	GAF	558	61
Q9KSG1	VC1295	HD-GYP	492	55
Q9KLI8	VCA0931	HD-GYP	460	51
Q9KPT8	VC2497	HD-GYP	417	47
Q9KSV4	VC1152	HDOD	311	35
Q9KU70	VC0653	EAL	684	78
Q9KM53	VCA0536	EAL	288	32
Q9KMO0	VCA1083	EAL	235	27
Q9KUW1	VC0398	EAL	673	78
Q9KUK3	VC0515	EAL HDOD	410	47
Q9KLS3	VCA0895	HD-GYP	901	112
Q9KLE4	VCA0802	Phosphodiester	657	74
Q9KNY0	VC2600	Phosphodiester	622	72
Q9KM78	VCA0510	Phosphodiester	968	110
Q9KRF5	VC1687	DHH	301	33
Q9KT24	VC1081	HD-GYP	380	43
Q9KT19	VC1086	EAL	536	61
H9L4Q6	VC1652	EAL	584	65
Q9KU65	VC0658	EAL	685	77
Q9KJ9	VC1641	EAL	257	29
Q9KLR1	VCA0681	HD-GYP	431	49
Q9KYR7	VC0072	EAL	768	88
Q9KQR4	VC1934	EAL	655	76
Q9KN01	VCA0165	GAF	710	81
Q9KN95	VCA0080	EAL	607	70
Q9KN18	VC2750	EAL	849	95
Q9KLF9	VCA0785	EAL GAF	627	71

B

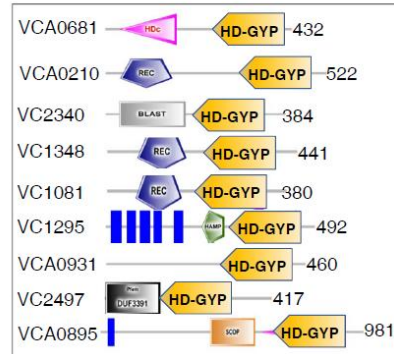


Figure S2 Identification of cGAMP PDEs of *V. cholerae* (V-cGAPs) that inhibit DncV-induced IRF3 activation.

(A) Summary of PDE candidates tested in this study and indicated detailed information. (B) Illustration of all HD-GYP domain containing proteins in *V. cholerae*.