

Supplementary Methods

Table S1. Sequence of the primers employed for PCR amplification in this study. The table includes oligonucleotide sequences employed for cloning (lower case letters indicate the SalI and NotI restriction sites, respectively) and RT-qPCR.

	gene symbol	primers sequence 5'→3'	
1	<i>Bd</i> GUCD1 (for cloning)	F	ATTCCCGGgtcgcacTATGGTGCTCAGGACGCTAG
		R	AGTCACGATgcggccgcTTACAGAGAGCTGTCTGG AAGTTTGC
2	<i>Bd</i> PepR2 (for RT-qPCR)	F	CGTGTACAGCTACGGAGTCG
		R	GCAGAATCCACAGGCATCTT
probe #128 (cat. no. 04693647001)			
3	<i>Bd</i> GUCD1 (for RT-qPCR)	F	GACATCGATAGAGTGGACGAACT
		R	GCACTGATGGATCTGCATTG
probe #151 (cat. no. 04694376001)			
4	<i>Bd</i> ACT1 (for RT-qPCR)	F	TGCTCCTCCTGAAAGGAAGT
		R	ACTCAGCCTTTGCAATCCAC
probe #120 (cat. no. 04693540001)			

Table S2. The qPCR efficiency values for the genes analyzed.

	Gene	Efficiency
1	<i>Bd</i> PepR2	1,983
2	<i>Bd</i> GUCD1	1,977
3	<i>Bd</i> ACT1	2,000

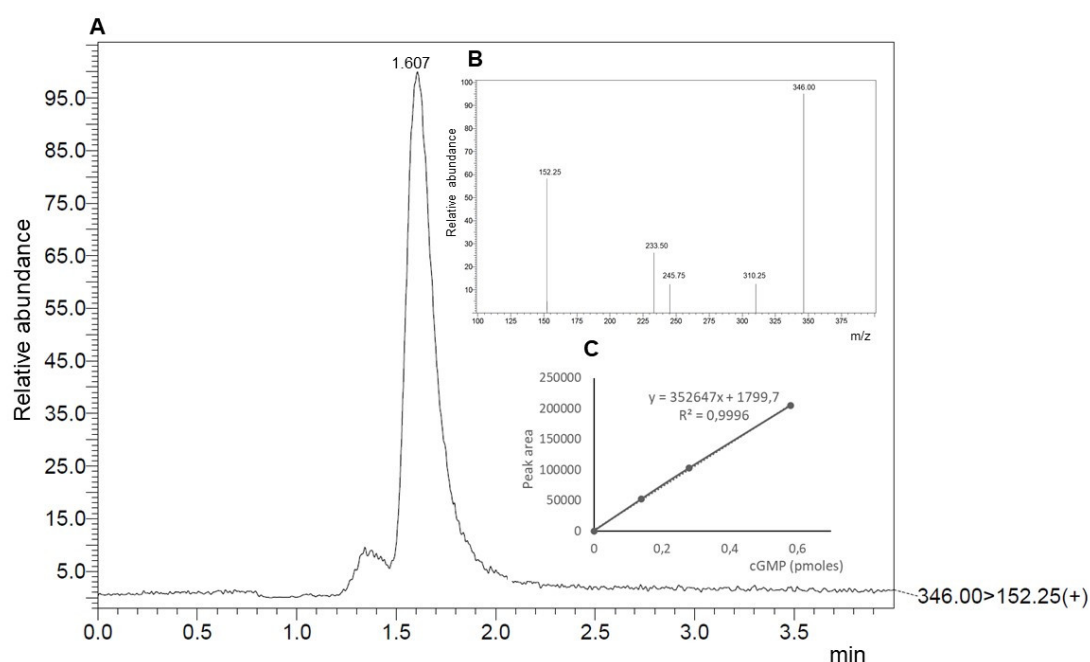


Figure S1. Detection of cGMP generated by BdGUCD1 by LC-MS/MS. (A) Determination of GC level by LC-MS/MS. Ion chromatogram of cGMP was generated from a reaction mixture containing 5 μ g of purified protein (B) Inset showing parent cGMP ion at m/z 346.00 [M + H]⁺ and corresponding fragmented daughter ion at m/z 125.25 [M + H]⁺. Fragmented product ion was used for quantitation. (C) Inset showing the cGMP calibration curve performed with 0–0.58 pmoles of pure cGMP on the column.

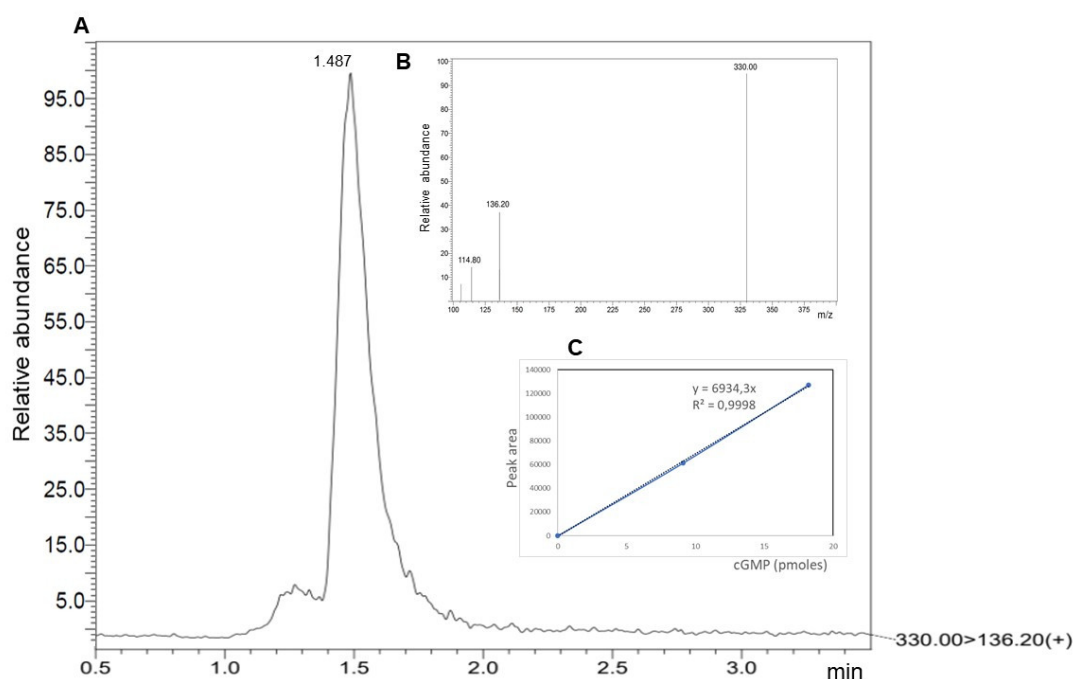


Figure S2. Detection of cAMP generated by BdGUCD1 by LC-MS/MS. (A) Determination of AC level by LC-MS/MS. Ion chromatogram of cAMP was generated from a reaction mixture containing 5 μ g of purified protein or (B) Inset showing parent cAMP ion at m/z 330.00 [M + H]⁺ and corresponding fragmented daughter ion at m/z 136.20 [M + H]⁺. Fragmented product ion was used for quantitation. (C) Inset showing the cAMP calibration curve performed with 0–18.22 pmoles of pure cAMP on the column.