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## Tuning the Innate Immune Response to Cyclic Dinucleotides Using Atomic Mutagenesis

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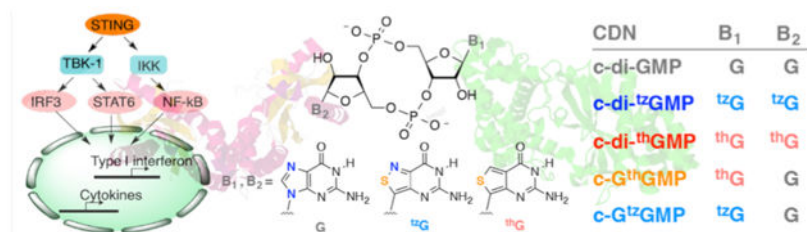
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### Abstract

Cyclic dinucleotides (CDNs) trigger the innate immune response in eukaryotic cells through the STING (stimulator of interferon genes) signalling pathway. To decipher this complex cellular process, a better correlation between structure and downstream function is required. Here we report the design and immunostimulatory effect of a novel group of c-di-GMP analogues. By employing an “atomic mutagenesis” strategy, changing one atom at a time, a class of gradually modified CDNs was prepared. These c-di-GMP analogues induce type-I interferon (IFN) production, with some being more potent than c-di-GMP, their native archetype. To the best of our knowledge, this study demonstrates for the first time that CDN analogues bearing modified nucleobases can tune the innate immune response in eukaryotic cells.

### Graphical Abstract



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Cyclic dinucleotides (CDNs) play critical regulatory roles in bacteria and trigger the innate immune response in eukaryotic cells. Here we illustrate that a systematic modification of the nucleobases, rather than the phosphate or sugar moieties, can generate STING agonists that demonstrate strong immunostimulatory effects.

## Keywords

cyclic dinucleotide; innate immune response; modified nucleosides; STING agonist

The innate immune system of eukaryotes possesses diverse mechanisms for detection of invading pathogens.<sup>[1]</sup> One of the most fundamental processes relies on cell surface or intracellular receptors that recognize molecular patterns unique to microorganisms.<sup>[2]</sup> Such pattern recognition receptors (PRRs) are capable of distinguishing pathogen-associated molecular patterns (PAMPs) from the molecular repertoire of endogenous host “self” patterns.<sup>[3]</sup> PRR engagement of PAMPs can then trigger the innate immune response to enhance antimicrobial activity, and further modulate the adaptive immune response.<sup>[2–3, 4]</sup> Cyclic dinucleotides (CDNs), which play critical roles in the cGAS-STING and RECON innate immune signaling pathways, have been recognized as PAMPs in recent years.<sup>[5]</sup> STING was the first mammalian receptor identified that directly binds CDNs,<sup>[6]</sup> most notably 2',3'-cGAMP, the product of the cytosolic DNA sensor cGAS.<sup>[7]</sup> A conformational change upon ligand binding recruits and activates the kinase TBK1 (Figure 1),<sup>[8]</sup> and phosphorylation of STING by TBK1 facilitates recruitment of transcription factor IRF3. When IRF3 itself gets phosphorylated by TBK1, it forms an activated homo-dimer that induces expression of type-I interferon (IFN  $\alpha/\beta$ ) and other cytokines within the nucleus (Figure 1).<sup>[9]</sup> In addition to the TBK1–IRF3 pathway, STING can activate other signaling pathways, including NF- $\kappa$ B and STAT6.<sup>[9–10]</sup>

Activation of STING by CDN analogues has shown pharmacological promise for improving the efficacy of cancer immunotherapies, including PD1 and CTLA-4 targeted drugs and CAR-T cell therapy.<sup>[11]</sup> Consequently, medicinal chemistry efforts have sought to develop hydrolysis-resistant CDNs with longer cellular residency time by altering the ribose and/or the phosphate moieties.<sup>[12]</sup> Chemical and chemo-enzymatic approaches have been taken for the preparation of CDN analogues bearing either backbone or nucleobase modification.<sup>[13]</sup> Analyses of the biological activities of such CDN analogues have provided valuable information on their binding properties to downstream sensors and augmented our knowledge regarding structure–activity relationships.<sup>[13–14]</sup>

In a recent publication, two fluorescent guanosine analogues developed in our laboratory have been used to prepare novel emissive CDNs, where atomic mutagenesis replaces the nucleobase's imidazole ring with a thiophene or an isothiazole moiety (<sup>th</sup>G or <sup>tz</sup>G, respectively, Figure 1).<sup>[15]</sup> While this contribution primarily focused on the photophysical properties of the emissive CDNs,<sup>[15c]</sup> these compounds could also provide insight into CDNs and their biological recognition, as the G analogues differ by one atom, and along with native guanosine, thus present a gradually altered purine molecular architecture. To deepen our molecular level understanding of CDN signaling, we analyzed here the

immunostimulatory effects of these systematically modified CDNs. We demonstrate that certain analogues can induce type-I IFN production more potently than their native archetype, highlighting potential new approaches to studying and manipulating the eukaryotic innate immune response.

The dimeric and mixed CDN analogues shown in Figure 1 were made from GTP, <sup>th</sup>GTP and <sup>tz</sup>GTP using DncV, a promiscuous dinucleotide cyclase from *Vibrio cholerae* (see Experimental Section and Figure S1 for data).<sup>[13b]</sup> The substrates and enzyme were incubated at 37 °C for 2–5 h, after which the reaction mixture was heat-inactivated and filtered before subjecting it to reverse-phase HPLC separation and purification. Pure fractions were collected, combined and lyophilized.<sup>[15c]</sup> The CDNs were re-dissolved in water for downstream experiments. To preliminarily determine whether the synthetic c-di-GMP analogues could activate the IFN response in eukaryotic cells, THP-1 cells were treated with 5 μM of c-di-GMP, c-G<sup>th</sup>GMP and c-di-<sup>th</sup>GMP. After 4 h incubation, induction of type-I IFN was measured with HEK-Blue IFN α/β reporter cells (see SI for experimental details). c-G<sup>th</sup>GMP induced type-I IFN production with comparable efficiency to c-di-GMP, while c-di-<sup>th</sup>GMP showed no activity (Figure 2).

To analyze the immunostimulatory effects of all synthetic CDNs in greater detail, RAW 264.7 cells were treated with various concentrations of c-di-GMP, c-di-<sup>tz</sup>GMP, c-di-<sup>th</sup>GMP, c-G<sup>th</sup>GMP and c-G<sup>tz</sup>GMP and the phosphorylation of IRF3 to pIRF3 was evaluated. CDNs were thus transfected into RAW 264.7 murine cells with digitonin as described in previous studies.<sup>[7b, 16]</sup> Cells were then lysed with NP-40 buffer 2 h after transfection, and total protein collected for immunoblotting against phosphorylated IRF3 (pIRF3) and β-actin. No pIRF3 was observed for untreated cells (UT) nor digitonin permeabilized cells (DG) (Figure 3). Low concentrations (1 μM) of c-di-GMP did not induce obvious IRF3 activation, while 5 and 10 μM displayed comparable efficiency in inducing IRF3 phosphorylation. Increasing amounts of phosphorylated IRF3 were observed when cells were treated with higher concentrations of c-di-<sup>tz</sup>GMP and c-G<sup>tz</sup>GMP, while no clear dose-response was observed for c-G<sup>th</sup>GMP (Figure 3a–b). The least isomorphic analogue, c-di-<sup>th</sup>GMP, did not trigger observable IRF3 activation at any of the concentrations tested. Two other biological replicates produced similar trends (Figure S2).

Since most synthetic c-di-GMP analogues activated IRF3, we analyzed their dose- and time-dependency for inducing IFN-β mRNA production using RT-qPCR. RAW 264.7 cells were transfected with 1, 5 and 10 μM of CDNs as described above and incubated for 2, 4 and 6 h. Total RNA was isolated and used for RT-qPCR (see Experimental section in the SI). As shown in Figure 4a and 4b, c-di-GMP induced the most IFN-β mRNA production 4 h post transfection, while the highest response was observed after 2 h for c-di-<sup>tz</sup>GMP, c-G<sup>th</sup>GMP, and c-G<sup>tz</sup>GMP. The same trend was observed for all three concentrations of CDNs tested (Figure 4a, Figure S3a–b). The IFN response to c-di-<sup>th</sup>GMP was minimal, but c-G<sup>th</sup>GMP showed the highest potency in inducing IFN-β mRNA production (Figure 4a–c, Figure S3a–d) among all CDNs tested. After 2 h incubation, 5 μM of c-G<sup>th</sup>GMP induced 10-fold higher IFN-β mRNA production than c-di-GMP, the native messenger. The differences in activity displayed by the analogues and their dependency on the specific assay used are discussed below.

Apparent STING activation by c-di-GMP analogues that contain unnatural isomorphous nucleobase was assessed here by three methods: type I IFN production measured by a reporter cell line, IRF3 phosphorylation measured by western blotting, and IFN- $\beta$  mRNA production measured by RT-qPCR. The initial analysis was performed in THP-1, a human cell line, while more detailed analyses were performed in RAW 264.7, a murine cell line. Our results show that all analogues except for c-di-<sup>th</sup>GMP stimulated the STING pathway in RAW 264.7 cells. The other three analogues appear to stimulate IRF phosphorylation at comparable or higher levels than the parent c-di-GMP 2 h post transfection.

To quantitatively analyze activation of the STING pathway, CDN-induced, IFN- $\beta$  production was measured by RT-qPCR in time- and dose-dependent manners. As seen in Figure 4, IFN- $\beta$  induction drops in the order: c-G<sup>th</sup>GMP > c-di-<sup>tz</sup>GMP > c-G<sup>tz</sup>GMP > c-di-GMP > c-di-<sup>th</sup>GMP, although it is apparent the cellular processes show complex concentration/time dependency. The effect of CDN concentrations above 5  $\mu$ M plateaued except for c-G<sup>tz</sup>GMP. Importantly, however, peak IFN- $\beta$  responses occurred at different times for different analogues, with the synthetic analogues c-di-<sup>tz</sup>GMP, c-G<sup>tz</sup>GMP and c-G<sup>th</sup>GMP inducing earlier and stronger maximum IFN- $\beta$  response compared to the native c-di-GMP (Figure 4, Figure S3 a–b). This pattern might result from negative feedback mechanisms of the CDNs-activated STING pathway and type-I IFN signalling.<sup>[17]</sup> We speculate that rapid and potent IFN induction might concomitantly activate early negative feedback responses, which ultimately result in down tuning IFN- $\beta$  production.

Among the synthetic CDNs tested in RAW 264.7 cells, the two analogues containing <sup>th</sup>G, the least isomorphous G surrogate that lacks the basic N7 in the native purine scaffold, displayed dramatically different potency in activating the STING pathway, with c-di-<sup>th</sup>GMP appearing essentially inactive, while c-G<sup>th</sup>GMP exerting the strongest effect on IFN- $\beta$  induction of all analogues tested. This stark difference was also observed in the THP-1 human cell line, although c-G<sup>th</sup>GMP showed comparable potency to c-di-GMP (Figure 2). Our findings could reflect differences in the assay themselves. RT-qPCR detects IFN- $\beta$  mRNA levels and not necessarily the translated active protein levels, whereas the reporter assay detects secreted type I interferons, including both IFN  $\alpha$  and  $\beta$ . Additionally, the difference between cell lines could be rationalized by the existence of multiple STING alleles in human cells compared to murine cells, which possess different sensitivity to CDNs.<sup>[7c, 18]</sup>

The observed intensity and duration of the cellular signalling response reflect both the affinity of the ligand to STING, as well as its resistance to intracellular degradation processes (assuming negligible differences in transfection efficiencies). It is perhaps not surprising that c-di-<sup>th</sup>GMP does not serve as a potent STING agonist, since it is the least isomorphous CDN analogue, with two altered purine cores. However, retaining one native G residue, as in c-G<sup>th</sup>GMP, restores STING activation. While speculative, this result is consistent with observations made for other asymmetric CDNs, 3',3'-cGAMP and 2',3'-cGAMP, that also induce more potent STING activation than c-di-GMP.<sup>[8a, 12a, 19]</sup> Either through enhanced binding affinity to STING or potentially increased resistance to hydrolytic degradation, c-G<sup>th</sup>GMP induces a faster and greater innate immune response relative to the native signal c-di-GMP in murine cells. It would be worthwhile to further investigate the

binding affinity of these CDN analogues to different STING variants, to further build correlations with their biological activity. In this context, the intrinsic fluorescence of our modified CDNs could potentially provide an effective tool to facilitate such studies.

Modifying the phosphate and sugar moieties of CDNs has been explored as a strategy to alter the pharmacological potency of STING agonists. Most of the non-cognate base-modified CDNs have not been tested in immune response assays.<sup>[14f]</sup> Here we illustrate that a systematic modification of the nucleobases, rather than the phosphate or sugar moieties, can generate STING agonists that are more potent than c-di-GMP. Particularly intriguing is the high potency of the mixed analogues c-G<sup>h</sup>GMP and c-G<sup>l</sup>GMP, where only one of the native guanosine residues is replaced by an unnatural synthetic C-nucleoside. Recognizing the complexity and intricacies of such cellular pathways, these observations put forth new approaches for the implementation of novel CDN analogues with altered recognition features, where the potency and duration of the triggered cellular immune response can potentially be tuned.

## Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

## Acknowledgements

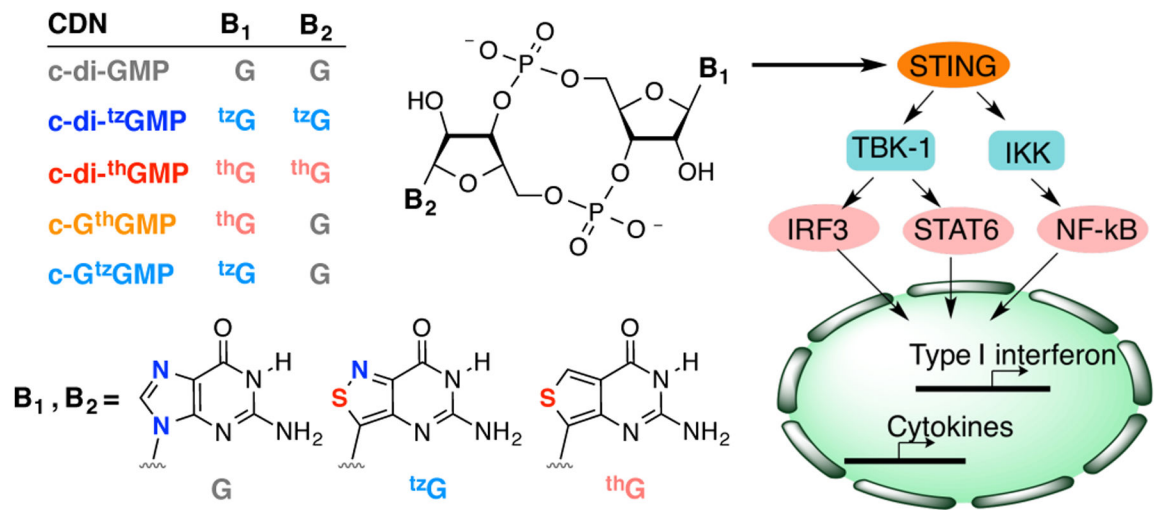
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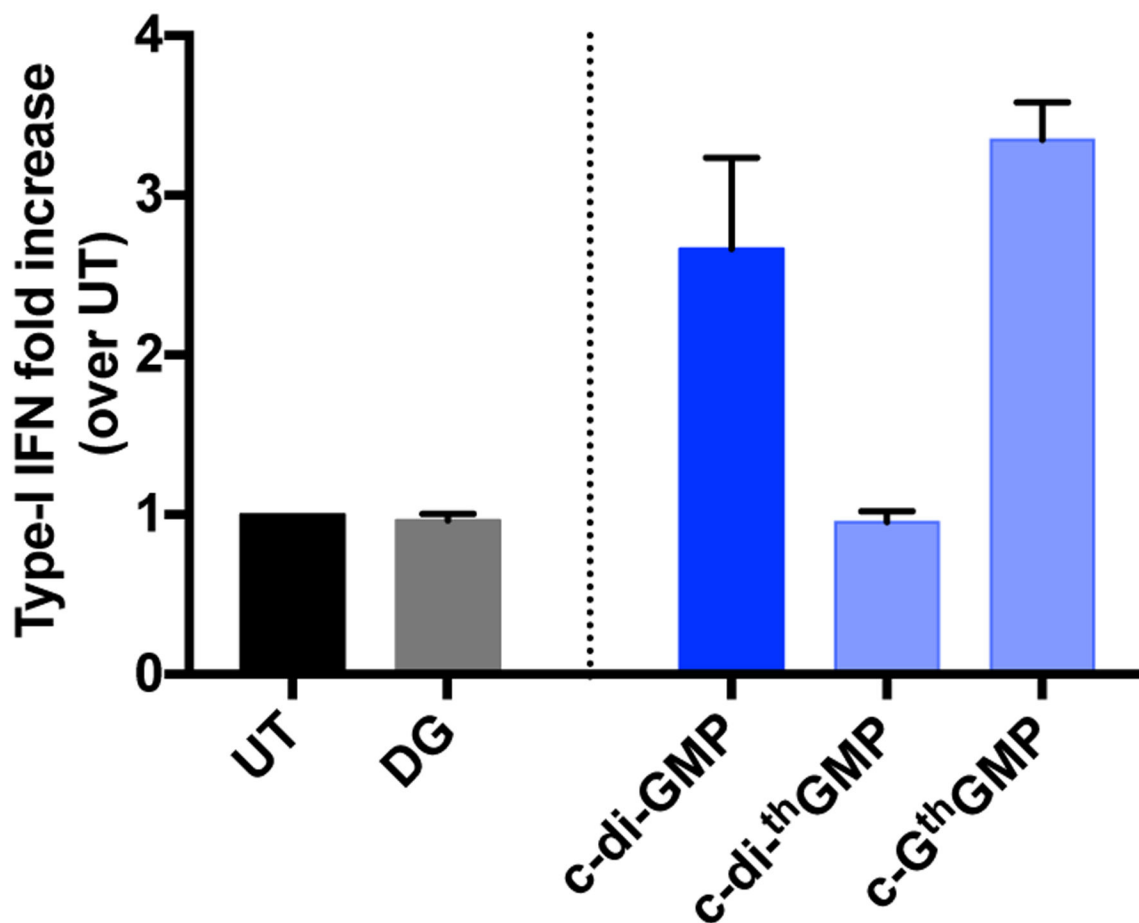
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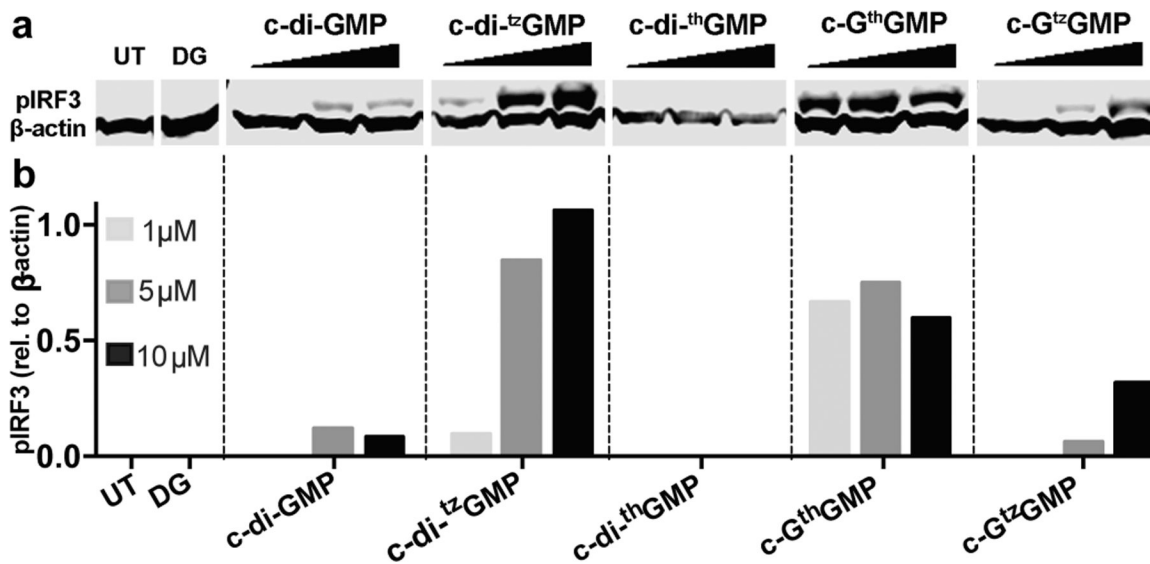
**Figure 1.**  
CDN analogues and their immunostimulatory effects.



**Figure 2.**

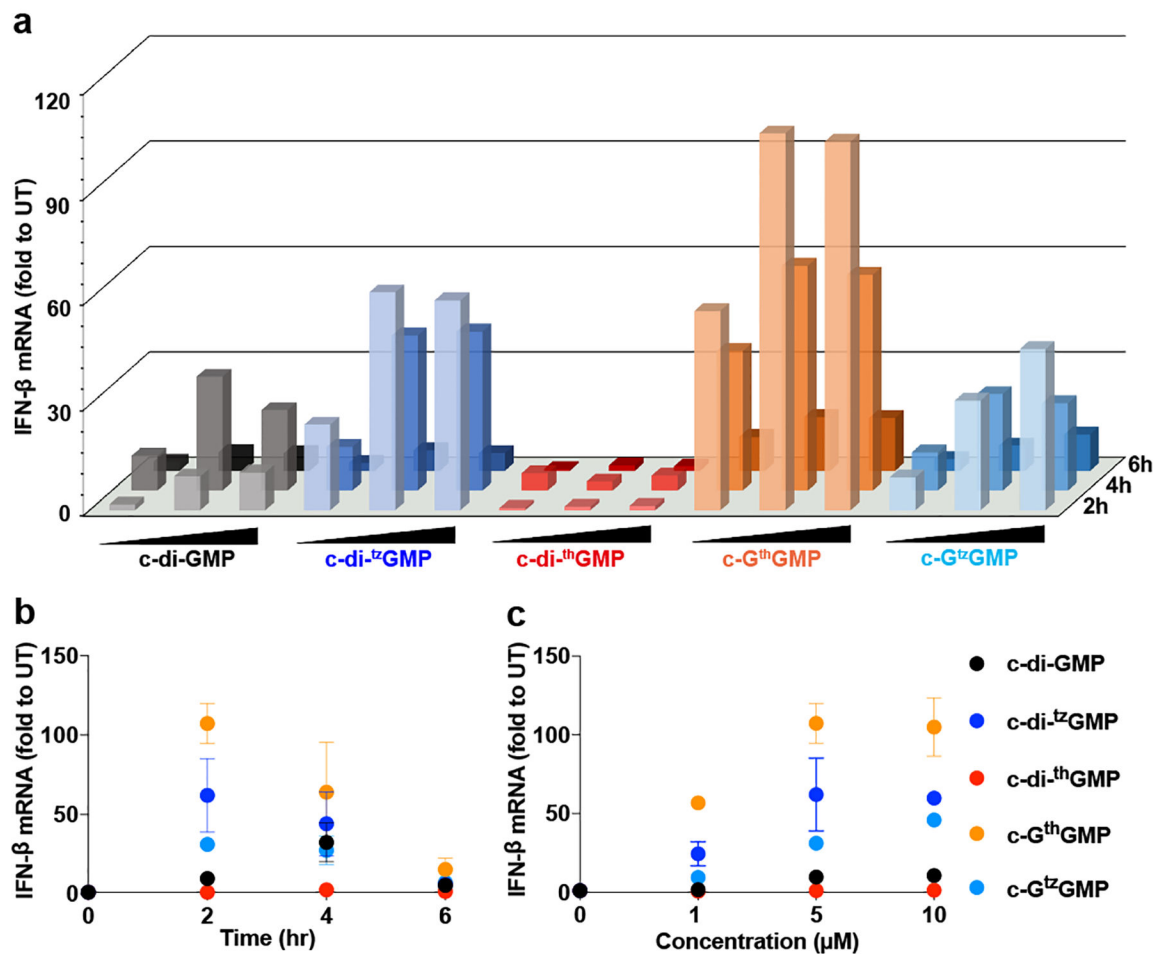
Type-I IFN induced by CDNs in THP-1 cells. THP-1 cells were seeded at a density of 100,000 cells/well in a 96-well cell culture plate and differentiated with 25 nM of PMA for approximately 20 h prior to treatment with CDNs. Cells were transfected with 5  $\mu$ M of CDNs in a permeabilization buffer containing 5  $\mu$ g/mL of digitonin, then washed and incubated in RPMI medium with 2% FBS at 37°C for 4 h. 50  $\mu$ L of cell culture supernatant per well was transferred to 150  $\mu$ L of HEK-Blue IFN  $\alpha/\beta$  reporter cells seeded at 50,000 cells/well in a 96-well cell culture plate and incubated at 37 overnight. The reporter cells were spin down the next day, and 50  $\mu$ L of cell culture supernatant per well was transferred to a 96-well plate and added with 150  $\mu$ L of QUANTI-Blue™ SEAP detection medium (InvivoGen). The samples were then incubated at 37°C for 1 h 20 min before absorption was measured at 640 nm. The absorption signal of each sample was normalized to untreated samples. Two independent assays were done in duplicates or triplicates. Error bars indicate SD.





**Figure 3.**

IRF3 phosphorylation induced by c-di-GMP and its analogues. (a) IRF3 phosphorylation induced by c-di-GMP analogues. 1, 5 and 10  $\mu$ M of each CDN was used to transfect RAW 264.7 cells. Cells were lysed with NP-40 lysis buffer 2 h post transfection, 20  $\mu$ g of total protein was loaded on SDS-polyacrylamide gel. Proteins were transferred to PVDF membrane after gel electrophoresis, and immunoblotted against pIRF3 and  $\beta$ -actin. (b) quantification of western blot. Y-axis indicates relative intensity of pIRF3 compare to  $\beta$ -actin.



**Figure 4.**

IFN- $\beta$  mRNA induction triggered by c-di-GMP and its analogues. (a) IFN production induced by c-di-GMP and its analogues. RAW 264.7 cells were transfected with 1, 5, 10  $\mu$ M of c-di-GMP, c-di-<sup>tz</sup>GMP, c-G<sup>th</sup>GMP, c-di-G<sup>th</sup>GMP and c-G<sup>tz</sup>GMP respectively, and incubated for 2, 4, 6 h respectively before lysed by TRIzol. RNA purification and RT-qPCR were conducted following the protocol described in the Methods section. (b) IFN response after 2, 4, 6 h incubation with 5  $\mu$ M of CDNs. (c) IFN response to 1  $\mu$ M, 5  $\mu$ M and 10  $\mu$ M of CDNs after 2 h incubation. Two independent assays were performed in triplicates (n=2). Error bars indicate SD.