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Harnessing stress granule formation by small molecules to inhibit the cellular replication of SARS-CoV-2†

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We identified small-molecule enhancers of cellular stress granules by observing molecular crowding of proteins and RNAs in a time-dependent manner. Hit molecules sensitized the IRF3-mediated antiviral mechanism in the presence of poly(I:C) and inhibited the replication of SARS-CoV-2 by inducing stress granule formation. Thus, modulating multimolecular crowding can be a promising strategy against SARS-CoV-2.

The coronavirus disease 2019 (COVID-19) pandemic, caused by severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2), has caused an unprecedented public health and economic crisis around the world, resulting in over 218 million reported cases of COVID-19 and 4.5 million deaths worldwide as of August 2021.^{1,2} Additionally, there are now more than 1000 confirmed variants of SARS-CoV-2 deposited in the viral genome sequence databases, and several variants, such as B.1.1.7. (alpha) and B.1.617.2. (delta), identified in the UK and India, respectively, have greater infectivity and severity to human beings.^{3,4} New variants may even escape the immunity elicited by vaccination and weaken the antiviral effects of conventional drugs.^{4,5} Despite this situation, there are only a few vaccines and SARS-CoV-2-targeted antiviral reagents available to date. Therefore, effective therapeutics are in high demand to end this pandemic, and a host-targeted antiviral strategy can be one of the most efficient options.

Stress granules are cytosolic membraneless organelles mainly composed of RNAs and proteins *via* multimolecular crowding.⁶ The formation of stress granules is induced by cellular stress, such as viral infection, to reduce the burden of translation, one of the most energy-consuming mechanisms in eukaryotic cells.⁷ Upon recognizing various types of stress, four

kinds of eIF2 α kinases can induce the phosphorylation of eIF2 α , resulting in the disruption of initiation codon recognition by the 48S pre-initiation complex, thereby inhibiting cellular translation and minimizing cellular damage.⁸ Concurrently, Ras GTPase-activating protein-binding protein 1 (G3BP1) captures polysome-released mRNAs and reversibly nucleates hundreds of proteins and RNA molecules through its RNA-binding domain and intrinsically disordered regions, which mediates multiple macromolecular interactions.^{9,10} Under viral infection, cellular translation is arrested by the phosphorylation of protein kinase R (PKR) *via* its recognition of viral double-stranded RNA (dsRNA) produced in viral replication cycles (Fig. 1).¹¹ Particularly, temporal suspension of cellular translation in the host can directly inhibit the viral replication because many viruses including coronaviruses utilize host translational machinery to proliferate themselves within the cell.^{12,13}

Stress granules are also the central hub for innate antiviral immune responses. In the presence of viral stress, host antiviral immune regulators, such as RIG-I and MDA5, are recruited and assembled together with mRNAs and translation machinery proteins to form the multimolecular condensate. This antiviral stress granule can serve as a signaling platform to activate various antiviral signaling pathways that converge to phosphorylate IFN-regulatory



Fig. 1 The schematic figure of stress granule formation and its antiviral function.

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factor 3 (IRF3), resulting in the induction of type I interferons and interferon-stimulated genes (ISGs) (Fig. 1).^{12,14,15} Most viruses including SARS-CoV-2 are known to alter multiple signaling pathways related to stress granule formation and subsequent type I interferon and ISG responses,^{16–18} indicating the importance of stress granules in antiviral responses. Considering that type I interferons have been used clinically as antiviral drugs for years and have beneficial effects on COVID-19 patients,¹⁹ enhancing the formation of stress granules using small molecules, thereby activating type I interferons, could be a novel strategy to treat viral diseases including COVID-19.

Given that stress granules are reversible and dynamic organelles, a time-dependent observation of stress granules in live cells could be informative. Thus, we utilized U2OS human osteosarcoma epithelial cells stably expressing the green fluorescent protein (GFP) tagged G3BP1, the marker protein of stress granules, for screening a drug-like small-molecule library. By simply counting cytoplasmic G3BP1 puncta without fixation in live cells, hit compounds at different time points can be discovered in a high-throughput manner. To confirm that the assay system can monitor the dynamics of stress granules in a 96-well plate format, we monitored GFP-G3BP1-expressed U2OS cells upon treatment with two well-known stress granule enhancers, sodium arsenite²⁰ and thapsigargin,²¹ at various concentrations and clearly observed a dose-dependent induction of stress granules (Fig. S1, ESI[†]). We observed that the number of stress granules was maximized at 2 h after treatment of both enhancers and gradually reduced afterward. In sodium arsenite-treated cells, the time for maximal stress granules was postponed as its concentration decreases, which is consistent with previous reports.²⁰

Since candidate compounds should be able to induce stress granule formation in virus-infected cells, we used polyinosinic:polycytidylic acid (poly(I:C)), a synthetic dsRNA analogue, to mimic virus-infected cells. Poly(I:C) is known to induce stress granule formation and activate the innate antiviral response upon recognition of viral dsRNA by PKR.²² Indeed, the transfection of poly(I:C) to human airway epithelial cells, Calu-3, activates the PKR-eIF2 α -IRF3 pathway and arrests global translation, confirmed by immunoblot analysis (Fig. S2a, ESI[†]). Immunofluorescence detection of G3BP1 also revealed that poly(I:C) transfection can time-dependently increase the number of stress granules in both Calu-3- and G3BP1-GFP-expressing U2OS cells, indicating that poly(I:C) is an appropriate viral dsRNA mimic (Fig. S2b and S3, ESI[†]).

Subsequently, we applied the time-course monitoring system of cellular stress granules to discover small-molecule enhancers of the stress granule formation in virus-infected cells from a 464 member representative set of privileged substructure-based diversity-oriented synthesis (pDOS) library.²³ Briefly, we transfected U2OS reporter cells with poly(I:C), treated them with individual compounds, and performed image-based monitoring of changes in cellular stress granules at 5 different time points (Fig. 2a). Compounds that enhanced the total count of stress granules per cell by >3-fold of standard deviations over negative controls were selected as



Fig. 2 (a) Screening result of the time-course stress granule monitoring system. After transfection of G3BP1-GFP-expressed U2OS cells with 1.25 $\mu\text{g mL}^{-1}$ poly(I:C), 10 μM compounds were treated. (b) Core structure of hit candidates and the chemical structure of optimized hit. (c) Immunofluorescence assay upon treatment of **C01** in Calu-3 cells. (d) Poly(I:C)-dependent induction of stress granule formation of **C01** in Calu-3 cell. 2.5 $\mu\text{g mL}^{-1}$ poly(I:C) was treated. Data represent mean \pm SD from two independent biological replicates.

hit compounds. Using our image-based time-course assay, we identified two classes of compounds as stress granule enhancers: benzopyranylpiperazine-cored compounds (Fig. 2b) and a tamoxifen derivative (Fig. S4, ESI[†]). Particularly, we observed different onset times for maximal stress granules in the case of several hit compounds, showing significant stress granule enhancement only after 3 h, which may not be detected without our time-course monitoring approach.

To improve the potency of initial hits, we performed a structure-activity relationship (SAR) study of benzopyranylpiperazine-cored compounds having novel core structures with better activities compared to tamoxifen derivatives. Through a systematic combination of substituted benzopyranyl β -keto aldehydes with various arylhydrazines, we synthesized over 100 analogues in a parallel process (Scheme S1, ESI[†])²⁴ and performed an image-based time-course evaluation of these analogues in a 96-well plate format. Several compounds consistently showed the induction of stress granule formation after 3 h, which is consistent with those from our initial screening. As shown in Fig. S5 (ESI[†]), the benzopyranylpiperazines containing *para*-CF₃-phenyl substituents were the key elements for the enhancement of cellular stress granules in the presence of poly(I:C), and analogues (**C01–C11**) containing various substituents with amide, urea, thiourea, and sulfonamide linkages at the R₃ position generally showed good to moderate stress granule enhancement activities. Most analogues with different substituents at the R₁ position did not show any activities at 5 μM concentration, and analogues containing the trifluoromethyl group at either the



ortho or the *meta* position did not induce the stress granule formation (Fig. S6, ESI[†], **NI**, **NJ**, and **NK12**). The modification of the secondary amine group of piperazine at the R₂ position failed to enhance the stress granules and significantly decreased their solubility (Fig. 2b and Fig. S6, ESI[†], **NI01–NI11**). Based on our SAR studies, we concluded that **C01–C04** effectively enhance the cellular stress granules in the presence of poly(I:C), which led us to conduct further evaluation. To examine whether compounds induce stress granule formation in lung epithelial cells where SARS-CoV-2 causes atypical respiratory failure,² we performed an immune-fluorescence assay for G3BP1 in Calu-3 human epithelial cells. **C01**, **C02** and **C04** increased the total number of stress granules, whereas **C03** weakly induced stress granule formation (Fig. 2c and d and Fig. S7, ESI[†]). In a tetrazolium-based viability assay with Calu-3 cells (Fig. S8, ESI[†]), **C04** showed a more cytotoxic effect (CC₅₀ = 37.5 μM) compared to **C01–C03** (CC₅₀ > 60 μM). **C01** and **C02** also induced stress granule assembly in Vero kidney epithelial cells from the African green monkey, especially in poly(I:C)-transfected cells, commonly used for COVID-19 antiviral experiments (Fig. S9a, ESI[†]). In particular, **C01** showed more specific induction of stress granule formation than **C02** in both Calu-3 and Vero cells (Fig. S9b, ESI[†]). Based on these observations, we selected compound **C01** for subsequent biological studies.

Having found that **C01** can facilitate the formation of cellular stress granules under poly(I:C)-treated conditions, we then investigated the compound effects on the innate antiviral responses. The **C01**-only treatment showed a marginal impact on the PKR-eIF2α-IRF3 signaling pathway in Calu-3 cells, but **C01** significantly activated this antiviral pathway in a dose-dependent manner and induced translation arrest, confirmed by the puromycin experiment, in the presence of poly(I:C), which mimics the viral infection (Fig. 3a). However, the negative analogue **NI02**, without enhancing the stress granule formation, did not activate antiviral responses. We then examined whether the antiviral efficacies of **C01** were influenced by the quantity of treated dsRNA, and observed that **C01**-based antiviral responses were positively correlated with the concentration of treated poly(I:C) confirmed by western blotting of the PKR-eIF2α-IRF3 signaling pathway, indicating that **C01** specifically upregulates innate antiviral responses in the presence of external dsRNA (Fig. 3b). Next, we performed a quantitative real-time PCR (qRT-PCR) to examine whether **C01** treatment induced the cellular production of type I interferons and ISG56. As shown in Fig. 3c and d, **C01** increased the cellular levels of IFNβ1 and ISG56 in poly(I:C)-treated Calu-3 cells; however, both mRNA levels were not significantly changed in the absence of poly(I:C) or upon treatment of **NI02**, which is consistent with our previous image-based observations of stress granule formation. Therefore, we concluded that **C01** inhibits viral replication by enhancing the formation of cellular stress granules and subsequent innate antiviral immune responses.

Next, we investigated the antiviral effects of **C01–C04** against SARS-CoV-2. Briefly, Calu-3 and Vero cells were infected with SARS-CoV-2 at multiplicity of infection (MOI) values of 0.2 and 0.0125, respectively. Then, the cells were treated with various concentrations of compounds. At 24 h post viral infection,

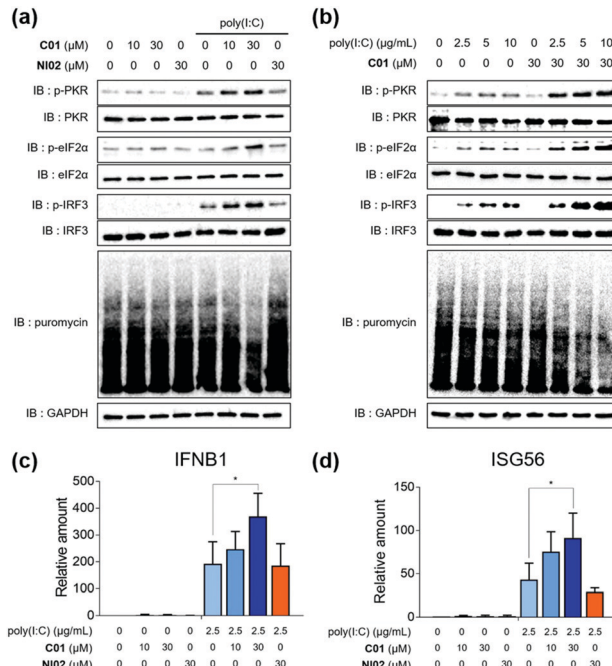


Fig. 3 (a) Poly(I:C)-specific dose-dependent activation of the antiviral response. 2.5 μg mL⁻¹ poly(I:C) was treated. (b) Poly(I:C)-dependent activation of the antiviral response. qPCR quantification of (c) IFNβ1 and (d) ISG56. Calu-3 cells were used. Data represent mean ± SD from at least two independent biological replicates. **p* < 0.05 (Student's *t* test).

immunofluorescence detection of the SARS-CoV-2 N protein and host cell nucleus was performed to quantitatively measure the inhibition of viral infection as well as cell viability. We observed that **C01–C04** showed dose-dependent inhibition of SARS-CoV-2 replication (Fig. 4a and Fig. S10, ESI[†]). In particular, **C01** showed the highest selective index (S.I.), the ratio of CC₅₀ to IC₅₀, in Calu-3 cells (Table S1, ESI[†]). On the other hand, the negative analogue **NI02** failed to inhibit the replication of SARS-CoV-2 in both Calu-3 and Vero cells. To elucidate the inhibitory mechanism of the compounds, we performed an immunofluorescence assay to G3BP1 for SARS-CoV-2-infected cells. SARS-CoV-2 infection triggered weak stress granule formation in Vero cells, and sodium arsenite-induced stress granule formation was inhibited by SARS-CoV-2 (Fig. S11, ESI[†]). In contrast, the **C01** and **C02** treatments, but not the **NI02** treatment, induce stress granule formation in the virus-infected cells, indicating that the stress granules induced by **C01** and **C02** activate innate immune responses against foreign viral dsRNA and block the viral translation (Fig. 4b and Fig. S12, ESI[†]). To specify the effective window of treatment concentration, which is limited by its cytotoxicity at high concentrations, we conducted a drug combination assay. In the presence of lopinavir (**LPV**), an antiviral reagent targeting viral proteases,¹⁹ **C01** showed significantly enhanced antiviral activity (IC₅₀ from 7.64 to < 0.78 μM) and selective index (from 2.09 to > 17.91) against SARS-CoV-2 in Vero cells, whereas **NI02** did not (Fig. 4c and Fig. S13, ESI[†]). These results suggested that the drug combination of our stress granule



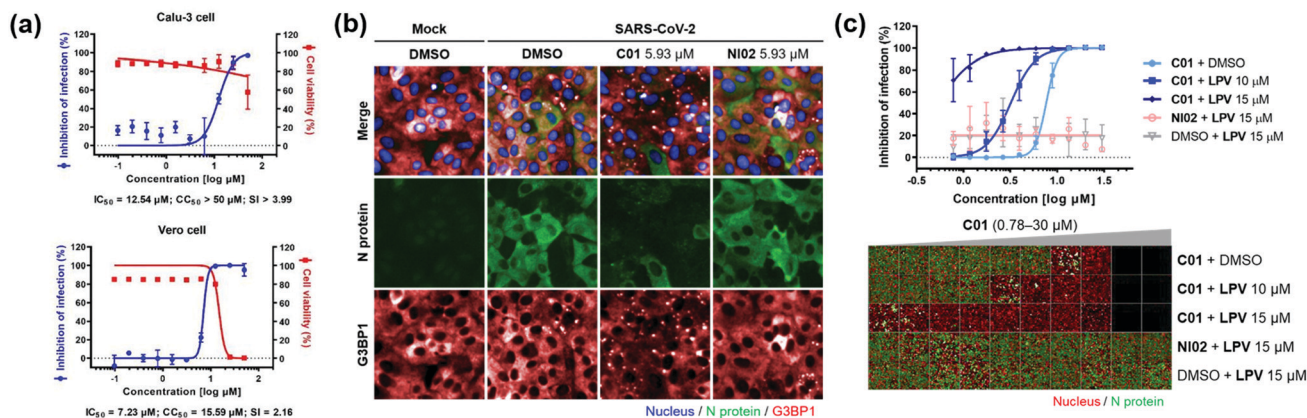


Fig. 4 (a) Dose–response curves of **C01** against SARS-CoV-2 in Calu-3 and Vero cells. (b) Co-immunofluorescence assay of **C01** and **NI02** (negative control) using viral N protein and G3BP1 antibodies in Vero cells. (c) Drug combination assay via co-treatment of **C01** with lopinavir in Vero cells. See the ESI† for details. Data represent mean \pm SD from two independent biological replicates.

enhancers and other antiviral reagents with different modes of action can be a novel strategy for treating viral diseases including COVID-19.

In summary, we discovered small-molecule stress granule enhancers by monitoring the time-course formation of cellular stress granules, cytoplasmic aggregates by multimolecular crowding of proteins and RNAs. To the best of our knowledge, this is the first dynamic screening of small molecules for stress granule regulators only in the presence of dsRNA and its potential application for antiviral disease treatment. We confirmed that our benzopyranopyrazole-based hit compounds activated innate antiviral immune responses and increased the expression of type I interferons and ISGs in a dsRNA-dependent manner in various epithelial cells. Hit compounds also increased the quantity of stress granules in SARS-CoV-2-infected cells and inhibited the viral replication of SARS-CoV-2. A drug combination strategy with the FDA-approved drug, lopinavir, enhanced the antiviral effects of our hit compounds against SARS-CoV-2. Our host immune-targeted strategy harnessing the role of antiviral stress granules might overcome the evolution and mutation of SARS-CoV-2 variants, which render the existing vaccines and virus-targeting drugs ineffective.

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Conflicts of interest

There are no conflicts to declare.

Notes and references

- World Health Organization, Coronavirus (COVID-19) Dashboard, <https://covid19.who.int/>, accessed August 31, 2021.
- A. G. Harrison, T. Lin and P. Wang, *Trends Immunol.*, 2020, **41**, 1100.
- B. Giles, P. Meredith, S. Robson, G. Smith and A. Chauhan, *Lancet Infect. Dis.*, 2021, **21**, 1213.
- C. del Rio, P. N. Malani and S. B. Omer, *JAMA*, 2021, **326**, 1001.
- W. T. Harvey, A. M. Carabelli, B. Jackson, R. K. Gupta, E. C. Thomson, E. M. Harrison, C. Ludden, R. Reeve, A. Rambaut, S. J. Peacock and D. L. Robertson, *Nat. Rev. Microbiol.*, 2021, **19**, 409.
- P. Anderson and N. Kedersha, *Nat. Rev. Mol. Cell Biol.*, 2009, **10**, 430.
- P. Anderson and N. Kedersha, *Trends Biochem. Sci.*, 2008, **33**, 141.
- A. Marcelo, R. Koppenol, L. P. de Almeida, C. A. Matos and C. Nóbrega, *Cell Death Dis.*, 2021, **12**, 592.
- P. Yang, C. Mathieu, R. M. Kolaitis, P. Zhang, J. Messing, U. Yurtsever, Z. Yang, J. Wu, Y. Li, Q. Pan, J. Yu, E. W. Martin, T. Mittag, H. J. Kim and J. P. Taylor, *Cell*, 2020, **181**, 325.
- T. Matheny, B. V. Treeck, T. N. Huynh and R. Parker, *RNA*, 2021, **27**, 174.
- S. J. DeWitte-Orr and K. L. Mossman, *Future Virol.*, 2010, **5**, 325.
- C. McCormick and D. A. Khapersky, *Nat. Rev. Immunol.*, 2017, **17**, 647.
- P. D. Slaine, M. Kleer, N. K. Smith, D. A. Khapersky and C. McCormick, *Viruses*, 2017, **9**, 388.
- L. C. Reineke and R. E. Lloyd, *J. Virol.*, 2015, **89**, 2575.
- N. Eiermann, K. Haneke, Z. Sun, G. Stoeklin and A. Ruggieri, *Viruses*, 2020, **12**, 984.
- D. Campos-Melo, Z. C. E. Hawley, C. A. Droppelmann and M. J. Strong, *Front. Cell Dev. Biol.*, 2021, **9**, 621779.
- H. H. Rabouw, M. A. Langereis, R. C. Knaap, T. J. Dalebout, J. Canton, I. Sola, L. Enjuanes, P. J. Bredenbeek, M. Kikkert, R. J. de Groot and F. J. van Kuppeveld, *PLoS Pathog.*, 2016, **12**, e1005982.
- B. Gao, X. Gong, S. Fang, W. Weng, H. Wang, H. Chu, Y. Sun, C. Meng, L. Tan, C. Song, X. Qiu, W. Liu, M. Forlenza, C. Ding and Y. Liao, *PLoS Pathog.*, 2021, **17**, e1008690.
- S. Alhumaid, A. A. Mutair, Z. A. Alawi, N. Alhmeed, A. R. Z. Zaidi and M. Tobaiqy, *Trop. Med. Infect. Dis.*, 2020, **5**, 180.
- A. P. Sfakianos, L. E. Mellor, Y. F. Pang, P. Kritsiligkou, H. Needs, H. Abou-Hamdan, L. Desaubry, G. B. Poulin, M. P. Ashe and A. J. Whitmarsh, *Cell Death Differ.*, 2018, **25**, 1766.
- M. G. Thomas, L. J. Martinez Tosar, M. A. Desbats, C. C. Leishman and G. L. Boccaccio, *J. Cell Sci.*, 2009, **122**, 563.
- G. Clavarino, N. Claudio, T. Couderc, A. Dalet, D. Judith, V. Camosseto, E. K. Schmidt, T. Wenger, M. Lecuit, E. Gatti and P. Pierre, *PLoS Pathog.*, 2012, **8**, e1002708.
- J. Kim, H. Kim and S. B. Park, *J. Am. Chem. Soc.*, 2014, **136**, 14629.
- S. O. Park, J. Kim, M. Koh and S. B. Park, *J. Comb. Chem.*, 2009, **11**, 315.

