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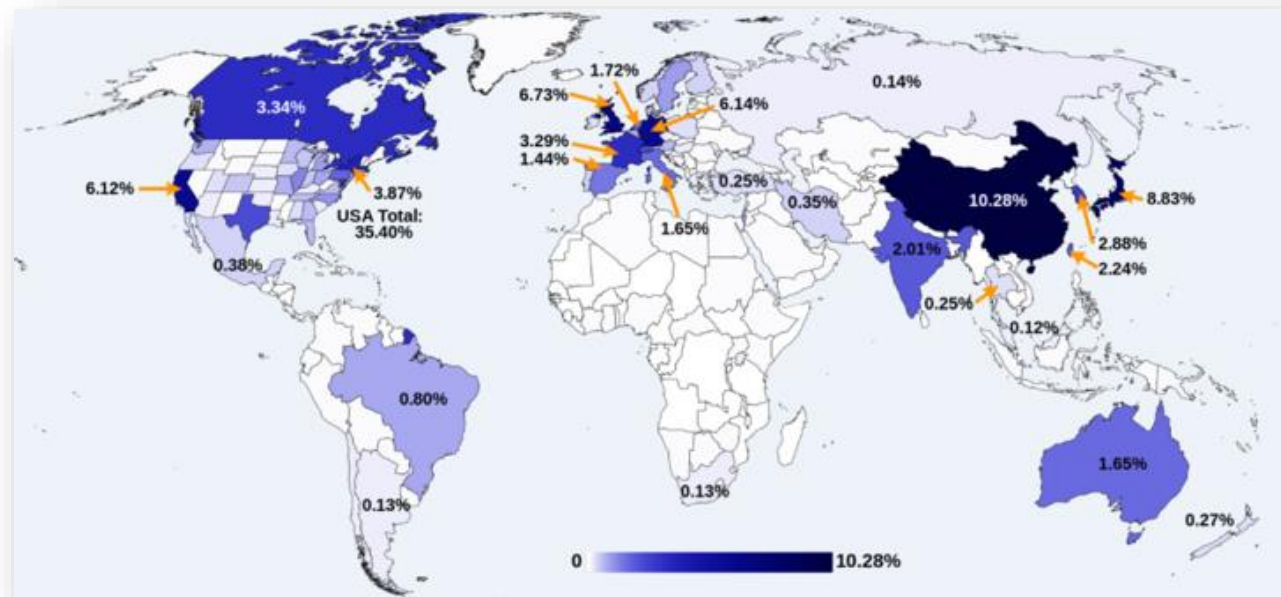
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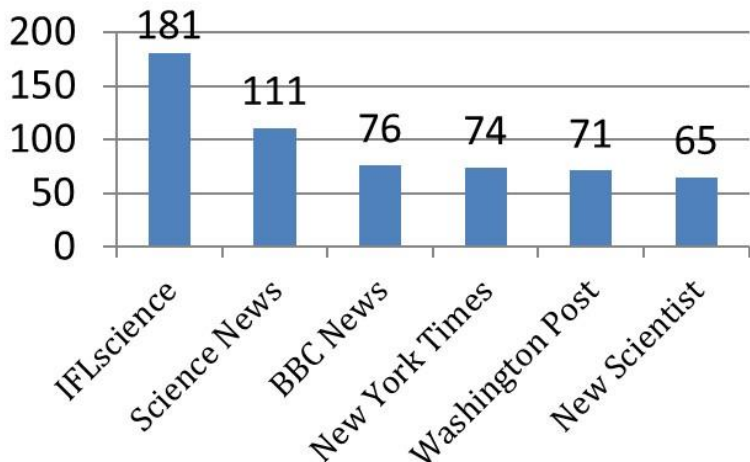


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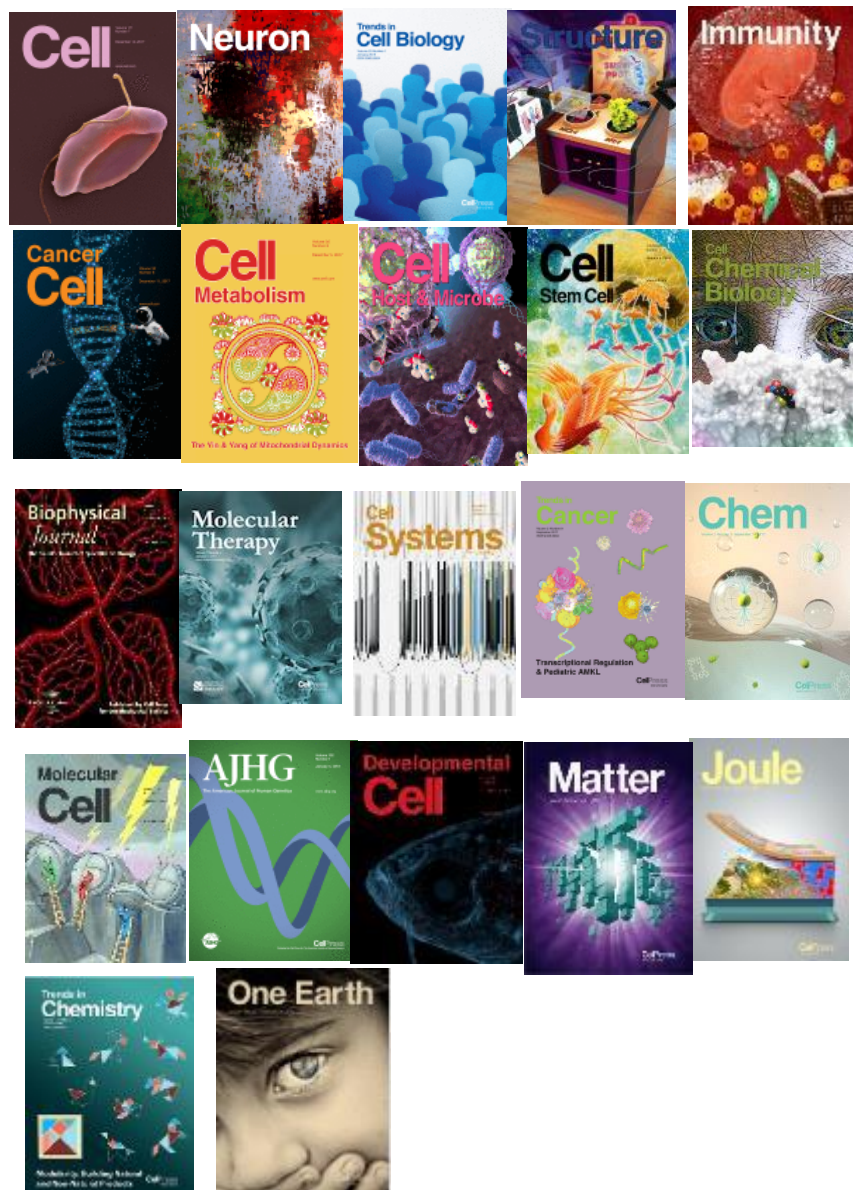
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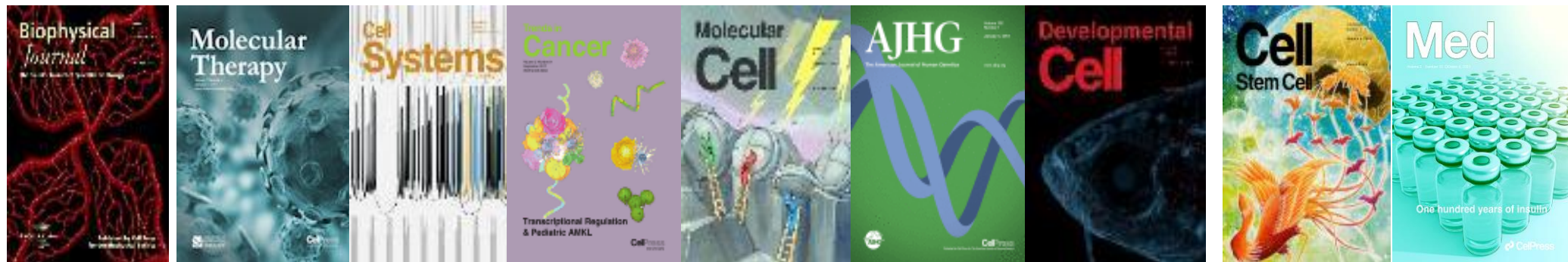






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## One Earth




Available online 3 March 2021

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## PM<sub>2.5</sub> reductions in Chinese cities from 2013 to 2019 remain significant despite the inflating effects of meteorological conditions

Qirui Zhong<sup>1,2</sup>, Shu Tao<sup>1,2,3,6,8,✉</sup>, Jianmin Ma<sup>2</sup>, Junfeng Liu<sup>2</sup>, Huizhong Shen<sup>1</sup>, Guofeng Shen<sup>2</sup>, Dabo Guan<sup>4,5</sup>, Xiao Yun<sup>2</sup>, Wenjun Meng<sup>2</sup>, Xinyuan Yu<sup>2</sup>, Hefa Cheng<sup>2</sup>, Dongqiang Zhu<sup>2</sup>, Yi Wan<sup>2</sup>, Jianying Hu<sup>2</sup>

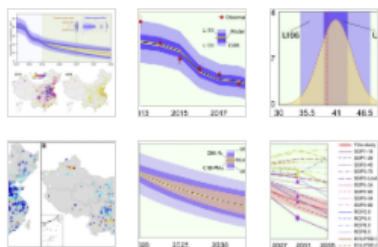
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## Highlights


- National mean PM<sub>2.5</sub> decreased by 34% from 2013 to 2019 through emission mitigation
- National PM<sub>2.5</sub> reductions were overestimated because of meteorological effects
- Mitigation efforts for individual cities are related to initial pollution level and GDP
- National mean PM<sub>2.5</sub> will decrease to 24.2 μg/m<sup>3</sup> by 2035 under continuous action

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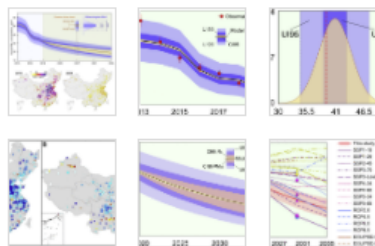
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## PM<sub>2.5</sub> reductions in Chinese cities from 2013 to 2019 remain significant despite the inflating effects of meteorological conditions


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Volume 171, Issue 6, 30 November 2017, Pages 1229-1231



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## How Cryo-EM Became so Hot

Yifan Cheng<sup>1</sup>, Robert M. Glaeser<sup>2</sup>, Eva Nogales<sup>2,3</sup>

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The Royal Swedish Academy of Sciences awarded the 2017 Nobel Prize for Chemistry to Jacques Dubochet, Joachim Frank, and Richard Henderson for “developing **cryoelectron microscopy** for the high-resolution structure determination of biomolecules in solution.” Achieving this goal, which required

## Cryo-Electron Microscopy

Cryoelectron microscopy showed that the flexible dimerization domain gets ordered when Ase1/PRC1 crosslinks two antiparallel microtubules [78], leading to the rod domains projecting from the microtubule-binding domains at a fixed angle [74,78].

From: *Trends in Cell Biology*, 2013

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## Cryo-EM, Part C: Analyses, Interpretation, and Case studies

Friedrich Förster, Elizabeth Villa, in *Methods in Enzymology*, 2010

## Abstract

Cryoelectron microscopy (cryo-EM) is an increasingly popular method to elucidate the structures of macromolecular complexes. However, in many applications the resolution of cryo-EM densities is limited to the low or intermediate resolution regime, that is,  $(10 \text{ \AA})^{-1}$  or worse. Therefore, unambiguous molecular interpretation of cryo-EM densities requires efficient use of additional information, such as

## Recent Advances in Electron Cryomicroscopy, Part B

Michael G. Rossmann, ... Pavel Plevka, in *Advances in Protein Chemistry and Structural Biology*, 2011

## Abstract

**Cryo-electron microscopy** (cryo-EM) in combination with **single-particle analysis** has begun to complement crystallography in the study of large **macromolecules** at near-atomic resolution. Furthermore, advances in **cryo-electron tomography** have made possible the study of macromolecules within their cellular

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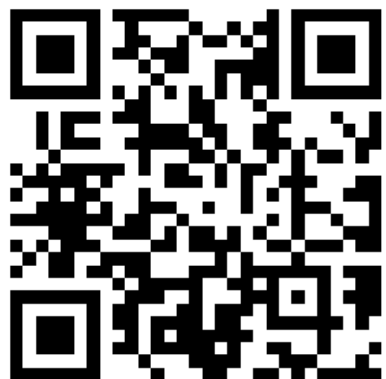
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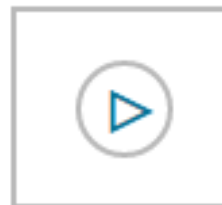
Using Alerts 

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Reviewing documents 

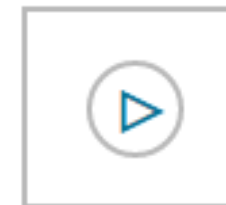
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Browsing journals



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