Combining pMINFLUX with graphene energy transfer for precise 3D localizations. a Top: Schematic of a DNA origami structure with a single dye positioned at a height of 16 nm above a graphene-on-glass coverslip. Bottom: Fluorescence intensity trace of the total fluorescence intensity of a single dye molecule in a single DNA origami structure. b Fluorescence decays for each of the four pulsed interleaved vortex-shaped beams which are focused on the sample arranged in a triangular pattern with the fourth beam placed at the center of the triangular structure. The star indicates the xy position of the dye molecule. c xy localization histogram of time bins. d Distribution of fluorescence lifetimes obtained from the time bins. e Distribution of the distances to graphene z calculated from the fluorescence lifetimes of d). f 3D localizations of the full fluorescence intensity trace using the 2D information of pMINFLUX and the z distances from the fluorescence lifetimes. The individual localizations are shown
Super-resolution microscopy methods are essential for uncovering the structures of cells and the dynamics of molecules. Since researchers overcame the resolution limit of around 250 nanometers (while winning the 2014 Nobel Prize in Chemistry for their efforts), which had long been considered absolute, the methods of microscopy have progressed rapidly.

Now a team led by LMU chemist Prof. Philip Tinnefeld has made a further advance through the combination of various methods, achieving the highest resolution in three-dimensional space and paving the way for a fundamentally new approach for faster imaging of dense molecular structures. The new method permits axial resolution of under 0.3 nanometers.

The researchers combined the so-called pMINFLUX method developed by Tinnefeld's team with an approach that utilizes special properties of graphene as an energy acceptor. pMINFLUX is based on the measurement of the fluorescence intensity of molecules excited by laser pulses. The method makes it possible to distinguish their lateral distances with a resolution of just 1 nanometer.

Graphene absorbs the energy of a fluorescent molecule that is no more than 40 nanometers distant from its surface. The fluorescence intensity of the molecule therefore depends on its distance from graphene and can be used for axial distance measurement.
a, pMINFLUX interrogates the position of a fluorophore with multiple spatially displaced doughnut beams and yields 2D fluorescence lifetime images with nanometer precision. b, Graphene provides a measure for the axial distance to graphene. The fluorescence lifetime shortens, the closer a fluorophore is to graphene. c, Combining the lateral information of pMINFLUX with the axial graphene distance information yields 3D localizations. GET-pMINFLUX yields photon efficient localizations with nanometer precision. This enables L-PAINT. The schematic of the DNA origami structure has a DNA-pointer protruding. The fluorophore modified DNA-pointer can transiently to one of three binding sites spaced with 6 nm. Within 2 s this dense structure is with nanometer precision localized in 3D by combining L-PAINT and GET-pMINFLUX. Credit: by Jonas Zähringer, Fiona Cole, Johann Bohlen Florian Steiner, Izabela Kamińska, Philip Tinnefeld

**DNA-PAINT increases the speed**
Consequently, the combination of pMINFLUX with this so-called graphene energy transfer (GET) furnishes information about molecular distances in all three dimensions—and does this in the highest resolution achievable to date of under 0.3 nanometers. "The high precision of GET-pMINFLUX opens the door to new approaches for improving super-resolution microscopy," says Jonas Zähringer, lead author of the paper.

The researchers also used this to further increase the speed of super-resolution microscopy. To this end, they drew on DNA nanotechnology to develop the so-called L-PAINT approach. In contrast to DNA-PAINT, a technique that enables super-resolution through the binding and unbinding of a DNA strand labeled with a fluorescent dye, the DNA strand in L-PAINT has two binding sequences.

In addition, the researchers designed a binding hierarchy, such that the L-PAINT DNA strand binds longer on one side. This allows the other end of the strand to locally scan the molecule positions at a rapid rate.

"As well as increasing the speed, this permits the scanning of dense clusters faster than the distortions arising from thermal drift," says Tinnefeld. "Our combination of GET-pMINFLUX and L-PAINT enables us to investigate structures and dynamics at the molecular level that are fundamental to our understanding of biomolecular reactions in cells."

The findings are published in the journal Light: Science & Applications.

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