Two-Dimensional Nonlinear Structured Illumination Microscopy with rsEGFP2

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Abstract: Superresolution microscopy enables imaging of subcellular structures and dynamics 9 with nanoscale detail. Among the various superresolution techniques, structured illumination 10 microscopy (SIM) stands out for its compatibility with live-cell imaging. Linear SIM is 11 restricted to a resolution improvement of a factor of two, improving the resolution to about 12 100 nm. Nonlinear SIM (NSIM) utilizes reversibly switchable fluorescent proteins to generate 13 a nonlinear response, allowing for the collection of higher spatial frequency information and 14 theoretically extending the resolution without limit. By employing rsEGFP2 and patterned 15 depletion illumination (PD) to generate the desired nonlinearity in the fluorescent response, we 16 have successfully achieved 2D PD-NSIM imaging of actin in live U2OS cells with sub-80 nm 17 resolution. 18

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20 1. Introduction

Structured Illumination Microscopy (SIM), a popular superresolution method for live cell imaging, 21 offers a unique approach to surpassing the diffraction limit from the perspective of the frequency 22 domain [1,2]. In SIM, the sample is illuminated with a sinusoidal pattern that modulates the 23 emitted fluorescence intensity to induce Moiré fringes, lower-frequency patterns resulting from 24 the mixing of the illumination pattern and the structure of the sample. These Moiré fringes are 25 observable with the microscope even when the sample structure is not resolvable. Because the 26 illumination pattern is known, the Moiré fringes can be used to reconstruct the sample structure 27 beyond the resolution limit of the microscope. 28

One of the biggest advantages of SIM over other superresolution methods is the imaging speed. 29 For live cell imaging, temporal resolution is important, and SIM has been demonstrated imaging 30 whole live cells [3,4]. Most of the current superresolution methods achieve remarkable spatial 31 resolution often at the expense of diminished temporal resolution and/or diminished field of view. 32 Since its invention [1,2], SIM has been integrated with various imaging techniques including 33 total internal reflection microscopy (TIRF) [3,5,6], light sheet microscopy [7,8], and multifocal 34 imaging [9–11]. However, the resolution enhancement of linear SIM is capped at a 2-fold 35 improvement because the resolution improvement is proportional to the frequency of the 36 structured illumination pattern. Consequently, achieving superresolution below 100 nm for *in* 37 vivo live cell imaging with SIM remains challenging. This can be achieved with very high NA 38 objectives that can use toxic immersion media [6]. 39

The resolution of SIM can be extended by harnessing the nonlinear response of fluorophores. When the fluorescence emission is a nonlinear function of the excitation intensity, higher harmonics of the structured illumination pattern are generated, resulting in higher-frequency information being shifted into the passband of the microscope. As the nonlinearity increases, more high frequency harmonics emerge, enabling higher spatial frequencies to be mixed down. Theoretically, the extent of the resolution improvement is unlimited since it is dictated by the number of high-frequency harmonic terms [12]. In reality, however, the final resolution is affected
by the noise in the system, and only the frequency terms above the noise level contribute to the
resolution of the final image [13]. Extending the resolution of SIM with the nonlinear response
of the fluorophores is referred to as Nonlinear SIM (NSIM).

There are different ways of generating the desired nonlinearity. NSIM was first demonstrated 50 using excitation saturation as the nonlinearity [12]. But excitation saturation requires very 51 high excitation intensities and is not compatible with live-cell imaging. Reversibly switchable 52 fluorescent proteins (rsFPs) that can toggle between a fluorescent and a non-fluorescent state can 53 generate a non-linear fluorescent response at much lower excitation intensities and have been used 54 for most NSIM realizations. rsFPs typically use two wavelengths. Negatively photoswitchable 55 rsFPs (np-rsFPs) use one wavelength for activation and a longer wavelength for both excitation and 56 deactivation. Positively photoswitchable rsFPs use one wavelength for activation and excitation 57 and another wavelength for deactivation. NSIM with rsFPs was first demonstrated with the 58 np-rsFP Dronpa [14]. NSIM has since been developed using several different np-rsFPs. NSIM 59 has also been used with the positively photoswitchable rsFP (pp-rsFP) Kohinoor [15]. 60

In principle the activation, deactivation, and excitation light can all be structured. However, 61 for a np-rsFP, the activation and deactivation/excitation beams are at different wavelengths and 62 care must be taken to match the patterns accounting for chromatic aberrations. To avoid this 63 problem, patterned depletion NSIM (PD-NSIM) can be used. In this approach, for an np-rsFP, 64 all fluorophores are activated with uniform illumination at the activation wavelength. Then the 65 sample is illuminated with a depletion pattern that turns off the rsFPs, and, finally, the sample 66 is illuminated with an excitation pattern that is π out of phase with the depletion pattern. This 67 approach requires 3 separate illuminations per raw image. 68

To reduce the number of required illuminations, patterned activation NSIM (PA-NSIM) has been used [6]. In PA-NSIM, the sample is illuminated with a sinusoidal activation pattern and then illuminated with a sinusoidal excitation pattern. So only two exposures are needed rather than three. However, care must be taken to ensure that the activation and excitation patterns are matched since they are at different wavelengths and chromatic aberrations may affect their relative magnification. As the nonlinearity is increased in PA-NSIM, the background increases because the DC term increases with increasing nonlinearity.

So far, the rsFPs that have been used for NSIM include Dronpa [14], Skylan-NS [6, 16], 76 rsEGFP2 [16], and Kohinoor [17]. Rego et al. succesfully demonstrated a lateral resolution of 77 40 nm using PD-NSIM with Dronpa to image purified microtubules, and nuclear pores in the 78 intact nuclear membrane, capturing 63 raw readout images (7 phases and 9 orientations) [14]. 79 They also imaged actin in intact fixed cells at 60 nm resolution. The switching rate of Dronpa is 80 slow, requiring about 0.4 seconds of exposure to turn off with approximately 20 W/cm² at 488 81 nm. This limits the imaging speed. Another significant limitation with Dronpa was its inability 82 to go beyond 15 switching cycles before substantial bleaching set in. Achieving even this limited 83 performance necessitated the use of toxic anti-bleaching chemicals, which limits its application 84 in live-cell NSIM imaging. Li and and colleagues utilized Skylan-NS to perform PA-NSIM, 85 imaging intracellular dynamics in living cells. They claimed a spatial resolution of approximately 86 60 nm and a time resolution of around 40 frames per second. Compared with other rsFPs, the 87 contrast ratio of Skylan-NS reduces as the number of switching cycles increases, reducing the 88 nonlinearity [16]. All resolution values reported by Li are stated as "theoretical", calculated based 89 on the number of orders used in NSIM, and the resolution is not independently evaluated [18]. 90 Zhang et al. further compared Skylan-NS with rsEGFP2 and Dronpa using PA-NSIM and 91 claimed that Skylan-NS outperforms other rsFPs for PA-NSIM [16]. While Skylan-NS offers 92 higher photon yields and contrast ratios, rsEGFP's faster switching kinetics, longer fluoresence 93 decay, and low light intensity requirements also give it advantages. Although the studies [16, 19] 94 show that rsEGFP2 is not ideal for PA-NSIM, its potential for PD-NSIM hasn't been well studied. 95

Here, we present the first demonstration of live 2D PD-NSIM using rsEGFP2. We demonstrate
 imaging of actin filaments in U2OS cells, achieving sub-80 nm resolution in live-cell imaging.
 We examine the images in both in both real-space and Fourier-space to support our claim of
 increased resolution.

100 2. Methods

101 2.1. Principle of PD-NSIM



Fig. 1. (a) Imaging formation of NSIM. (b) Frequency-space representation of object and effective illumination pattern. (c) Frequency content of first higher-order harmonic order NSIM within 5 circular regions. (d) Effective OTF of first higher-order harmonic order NSIM after rotating 5 angles.

In fluorescence microscopy, each image is the product of sample structure and the excitation 102 pattern convolved with the microscope point spread function (PSF), as shown in Fig. 1(a). If the 103 excitation pattern is sinusoidal, it will create additional copies of the sample structure that are 104 shifted in spatial frequency. In frequency-space additional band-limited copies of the sample 105 information are generated as shown in Fig. 1(b). With SIM, the excitation pattern cannot have 106 a sample frequency greater than $\frac{2NA}{\lambda}$, and the copies cannot be shifted further than the green 107 circles shown in Fig. 1(b). Thus a frequency outside that region will still be unresolvable. In 108 NSIM, the nonlinearity distorts the fluorescence emission pattern generating higher harmonics 109 which create additional copies of the sample information shifted by larger amounts as shown 110 in Fig. 1(c). The pattern is rotated to cover an isotropic region of frequency-space, Fig. 1(d). 111 Excellent explanations of linear SIM can be found in [20]. 112

In patterned depletion NSIM, the sample is illuminated three times for every exposure. First 113 a uniform beam at the activation wavelength, 405 nm, activates the sample. Then the sample 114 is illuminated with a patterned depletion beam at the depletion/excitation wavelength, 488 nm. 115 This beam serves to deactivate the fluorophores. The longer the beam is left on, the more the 116 fluorophores are deactivated. However, at the intensity zeros of the pattern, the fluorophores will 117 not be deactivated. So, the longer the depletion is turned on, the narrower the peaks of activated 118 fluorophores as shown in Fig. 2. The fluorescence is then excited with a third beam that is π out 119 of phase with the depletion beam so that the intensity maxima are now centered on the remaining 120 activated fluorophores. The excitation is then more sharply peaked than a sinusoid resulting in 121 higher harmonics that provide higher resolution information. 122



Fig. 2. Patterned depletion Illumination for NSIM

123 2.2. Characterization of rsEGFP2

rsEGFP2 is an rsFP derived from the enhanced green fluorescent protein (EGFP) by replacing 124 threenine 65 by alanine [21]. Fig. S1(a) shows the 3D structure of the protein. It operates in a 125 negative switching mode, with a 480 nm depletion/excitation wavelength, and a 405 nm activation 126 wavelength. rsEGFP2 is significantly faster than rsEGFP. At a light intensity of 5.5 kW/cm^2 127 rsEGFP2 switches off about 6.5 times faster than rsEGFP. With a light intensity of 100 W/cm². 128 rsEGFP2 can achieve an off-rate greater than 90% with a 4ms exposure time at 480 nm. Similarly, 129 rsEGFP2 achieves greater than 90% activation with a 5ms exposure to 250 W/cm² at 405 nm. 130 rsEGFP2 can undergo over 2,100 switching cycles without a significant reduction in fluorescence 131 intensity, making it highly durable for PD-NSIM which requires more exposures than PA-NSIM. 132 The relatively low light intensities required for switching between states help reduce phototoxicity 133 and photobleaching making rsEGFP2 well suited for long-term live-cell imaging. Its successful 134 application in RESOLFT further demonstrates its utility in high-resolution imaging [21]. 135

Compared to Skylan-NS, the switching time of rsEGFP2 is 4 ms compared to 10 ms for Skylan-NS [16]. rsEGFP2 is a factor of 4 less bright than Skylan-NS. rsEGFP2 is more photostable than Skylan-NS, losing 20% of intensity over 1000 switching cycles and having a stable contrast ratio of 20. Whereas, Skylan-NS loses over 50% of intensity and the contrast ratio degrades from 37 to 17.

141 2.3. Optical Setup

Fig. 3 shows our experimental setup for 2D PD-NSIM. The system is built on an Olympus IX71 142 inverted microsope with a Prior Proscan XY Stage and a Prior Nanoscan SP200 Z-stage. The 143 system includes a 405 nm activation laser (Coherent OBIS 405 nm LX 100 mW) and a 488 nm 144 depletion/excitation laser (Coherent OBIS 488 nm LX 150 mW). The depletion beam is reflected 145 by a polarizing beam splitter (PBS, 10FC16PB.3 Newport), through an achromatic half-wave plate 146 (AHWP10M-600, Thorlabs) onto a spatial light modulator (SLM, Forth Dimension QXGA-3DM). 147 The combination of PBS, HWP, and SLM generates binary phase-patterns as described in [3]. 148 The activation beam is reflected by the PBS into the excitation path without reflecting off of 149 the SLM, providing a uniform illumination to activate all the rsFPs without patterning. The 150 activation and depletion beams are sent through a series of relay lenses to the microscope body 151 which contains the tube lens and objective (UPLAPO60XOHR TIRF Objective, 60x, 1.50 NA. 152 Olympus). The total magnification from the SLM to the sample is 240×. The relay lenses allow 153 for the alignment of the illumination beams onto the sample with mirrors M5 and M7 which 154 are close to the sample and pupil conjugate planes. Relay lenses L7 and L8 are used to create 155 an image of the pupil at mirror M9 which can be replaced by a deformable mirror for adaptive 156 optics correction. For this work, M9 is a dielectric mirror. 157

The fluorescence emission is directed to the sCMOS camera (Andor SONA 4BV6U, 6.5μ m pixel size) through the "image flat" long-pass dichroic mirror (Di03-R488-t3-25x36, Semrock). The total magnification from the sample to the camera is 180×, and the effective pixel size is 36.1 nm. An emission filter (Semrock BrightLine quad band bandpass filter, FF01-446/523/600/677-25) and a notch filter (Semrock StopLine quad-notch filter, NF03-405/488/561/635E-25) are put before the camera to block unwanted light, assuring a low background noise level.

To maximize the modulation strength of the sinusoidal pattern, the polarization state of the 164 two interference beams must be normal to the direction of the pattern wave vector (s-polarized) 165 to achieve maximum contrast of the pattern. Therefore, the polarization state of the incident 166 beam needs to be rotated along with the pattern orientation. A half-wave plate (HWP2 in Fig. 3) 167 mounted in a fast-motorized rotation stage (8MRU, Altechna) is used to control the polarization. 168 All but the ± 1 diffraction orders are filtered out by the mask after mirror M4 (custom design, 169 Chrome mask on Soda Lime substrate, Photosciences, Inc.). The activation beam is tilted to 170 align with one of the first-order diffracted beams of the depletion beam to ensure that it will pass 171



Fig. 3. Experimental setup of 2D-NSIM. HWP: half-wave plate. PBS: polarized beam splitter. PH: pinhole. M1-M4: flat mirror. L1-L11: lens. $f_1 = 50$ mm, $f_2 = 400$ mm, $f_3 = 400$ mm, $f_4 = 125$ mm, $f_5 = 125$ mm, $f_6 = 100$ mm, $f_7 = 250$ mm, $f_8 = 250$ mm, $f_9 = 300$ mm, $f_1 0 = 50$ mm, $f_1 1 = 200$ mm. The blue represents the illumination path (488nm) and the green path is the detection path.

172 through the mask.

173 2.4. Data Acquisition

The data acquisition process for 2D PD-NSIM begins by activating all fluorophores using uniform illumination of ~ 10 W/cm² for 5 ms at 405 nm. Then the patterned depletion illumination of ~ 5 W/cm² for 10 ms at 488 nm is applied. Finally, the phase of the illumination pattern is shifted by π , and the 488 nm excitation beam is turned on for 6 ms at ~ 5 W/cm².

The pattern at the SLM has a period of 15 pixels split 7/8 between positive and negative phase pixels. The nonlinear pattern generation is repeated with 5 phase shifts and 6 angles of rotation. Patterns at angles 0°, 30°, 60°, 90°, 120°, and 150° were used. A total of 30 raw images is required to reconstruct a single 2D PD-NSIM image. To increase SNR while maximizing the nonlinear response of rsEGFP2, we repeat the acquisition process 5 times for each phase, rather than using a single long exposure. The images are then averaged to create a final image stack for reconstruction.

185 2.5. Cell Culture and Transfection

The U2OS cells used in this paper were cultured in Dulbecco's Modified Eagle Medium (DMEM) supplemented with 10% (vol/vol) fetal bovine serum (FBS) and 1% (vol/vol) penicillinstreptomycin (Pen-Strep) and maintained at 37°C and 5% (vol/vol) CO₂ in a humidified incubator. Cells were passaged upon reaching apporximately 80-90% confluence by washing with phosphatebuffered saline (PBS), detaching with 500 μ L of trypsin-EDTA solution, neutralizing with four volumes of complete medium and centrifuging at 1500 rpm for 5 minutes. The pellet was resuspended in fresh growth medium and seeded at an appropriate density for subsequent use.

To visualize actin structures, the Actin-chromobody plasmid (a gift from Testa's lab) was obtained. We performed standard transformation into E. coli strain α 5, followed by inoculation into a large flask (50 mL culture volume) and overnight culture at 37° C. The next day, after centrifugation, the bacterial pellet was collected, and the plasmid was purified using a Qiagen midiprep kit. The purified plasmid, dissolved in ddH₂O at a concentration of $1 - 2\mu g/\mu L$, was used for subsequent transfection into U2OS cells.

The transfection was performed using PEI reagent. The imaged U2OS cells were plated on 199 35 mm ibidi dishes pre-coated with 0.1% PBS-diluted fibronectin human plasma and incubated 200 overnight to achieve 90-95% confluence. For transfection, $3\mu g$ of rsEGFP2 plasmid DNA was 201 diluted in 50 μ L of Opti-MEM medium, while 6 μ L of PEI reagent was diluted in another 50 μ L 202 of Opti-MEM. The diluted PEI was added to the DNA solution, mixed thoroughly, and incubated 203 at room temperature for 10 minutes to form complexes. The transfection mixture was added 204 dropwise to the cells and incubated at 37° C and 5% CO₂ for 24 hours. The cells were then 205 imaged within 6 hours. 206

207 3. Results

208 3.1. Modulation Strength

Fig. S2 shows simulation results demonstrating how depletion and excitation exposures affect 209 modulation strength. The modulation strength is measured as the OTF overlap magnitude. As 210 expected, with excitation exposure time fixed at 10 ms, an increase in depletion exposure time 211 enhances modulation strength for both linear and nonlinear orders. Conversely, with depletion 212 exposure fixed at 10 ms, longer excitation times diminish modulation strength for both harmonic 213 order. However, the simulation results only show the effects of the depletion and excitation 214 illumination exposures and do not account for specific characterists of rsEGFP2 such as the 215 switching-off time, activation time, or photobleaching. We did measurements using rsEGFP2 216 labeled actin in U2OS cells; the results are shown in Fig. 4. We measure the modulation contrast 217 for both the first and second order signals as a function of the excitation exposure or depletion 218 exposure time. Each measurement was repeated on five different cells, and we plot the error bars 219 from the different measurements. 220

For excitation exposure, similar to the simulation results, increasing exposure time led to 221 higher modulation strength for both harmonic orders. However, in the case of depletion, the 222 modulation contrast for the first harmonic initially increased with longer depletion times, leveling 223 off at 0.25 at 6 ms. In contrast, for the second harmonic, increasing depletion exposure resulted 224 in a reduction in modulation strength. This may be due to a lower than desired modulation 225 contrast of the excitation pattern. This will result in the fluorophores being turned off everywhere 226 if the depletion beam is kept on too long. Therefore, to balance the linear and nonlinear response, 227 we chose 10 ms for the depletion exposure and 6 ms for the excitation exposure. 228

To increase the SNR of the collected data, we repeated each imaging cycle multiple times and averaged the images. To determine the number of times to repeat each image, the illumination cycle was repeated with the setting of 10 ms depletion exposure and 6 ms excitation exposure for up to 50 cycles. The results, Fig. 4(c), show the modulation strength increases rapidly initially



Fig. 4. (a) Experimental results showing the effects of depletion and excitation pattern exposure on the nonlinear response. Data was collected from rsEGFP2-labeled U2OS cells with a coarse pattern spacing of $0.5NA/\lambda$. For each plot, five cells were imaged, and the data represents the mean values with error bars indicating the standard deviation. Top inset: Relationship between excitation exposure time and modulation depth, with depletion exposure time fixed at 10 ms. Bottom inset: Relationship between depletion exposure time and modulation depth, with excitation exposure time fixed at 10 ms. (b) Experimental result of repeating test. Data was collected from same U2OS cell with an illumination pattern spacing of $0.5NA/\lambda$. (c) Synchronization trigger sequence of all components during 2D PD-NSIM data acquisition for a single imaging cycle at one angle.

²³³ but only very slowly after 5 cycles. The increase in OTF overlap indicates an increase in the
 ²³⁴ SNR. Therefore, we chose to repeat each raw image cycle 5 times. Fig. 4(d) shows the timing
 ²³⁵ sequence of the lasers, polarization rotation, SLM, and camera.

236 3.2. NSIM

The performance of 2D PD-NSIM was demonstrated by imaging actin labeled live U2OS cells 237 transfected with rsEGFP2. Results are shown in Fig. 5, Fig. S4, and Fig. S5. Fig. 5(a) shows live 238 cell imaging of the U2OS cells with a pattern frequency of 1.66NA/ λ . The effective OTF after 239 the NSIM reconstruction has an extended bandwidth of $5.32NA/\lambda$, as shown Fig. 5(b), 2.66× the 240 extent of the widefield OTF. This corresponds to a resolution of ~ 70 nm. A clear improvement 241 in resolution can be observed between WF, SIM, and PD-NSIM images as shown in the insets 242 in Fig. 5(b) and (c). The Fourier Transform (FFT) of the spatial images also reveals strong 243 features in frequency space, demonstrating the extended frequency space captured by PD-NSIM. 244 Fig. 5(d) shows a different cell imaged with the same pattern spacing. A line profile through 245 an actin fiber is shown in Fig. 5(e). The measured Full-Width at Half-Maximum (FWHM) of 246 the fiber measured from the NSIM image is measured to be 80 nm, three times smaller than the 247 measurement from the WF image. This improvement is greater than the increase in the OTF 248 extent, because the Wiener filtering provides a further increase in the measured resolution. 249



Fig. 5. Live-cell imaging of 2D PD-NSIM with rsEGFP2. (a) A live U2OS cell transfected with rsEGFP2 imaged with widefield (WF), Linear SIM, and 2D PD-NSIM. (b) Bottom inset: zoomed-in view of the pink boxed region in (a). Top inset: corresponding fast Fourier transform (FFT) image. (c) Top inset: zoomed-in view of the orange boxed region in (a). Bottom inset: corresponding fast Fourier transform (FFT) image. (d) A live U2OS cell transfected with rsEGFP2 imaged with WF, Linear SIM, and 2D PD- NSIM. Right inset: Zoomed-in view of the blue boxed region in the left construction image. (e) Intensity plot of the yellow line marked in yellow boxed region from (d). The FFT image is displayed with logarithmic scaling for clarity with the green, red, and yellow circles represent effective OTF for WF, Linear SIM and the NSIM reconstructions in frequency domain.

Theoretically, for a pattern spacing of $1.66 \text{NA} / \lambda$ (NA=1.5 and $\lambda = 503 \text{ nm}$), the WF resolution 250 is 167.6 nm. From simulations, the resolution after SIM and NSIM reconstruction with a Wiener 251 filter is 92 nm and 68 nm respectively. To better demonstrate the repeatability of our approach 252 and the real resolution improvement, we quantified the FWHM of 19 actin filaments from 7 253 different U2OS cells. We measured the filaments about 1μ m from the edge of the actin fiber to 254 increase the chance that we are measuring a single actin fiber, and the FWHM represents the 255 system resolution. Fig. S3 shows the resolution distribution of the measured actin fibers, giving 256 a mean FWHM of 208 nm for WF, 97 nm for SIM, and 70 nm for PD-NSIM. The resolution 257 improvement of PD-NSIM is 3×. 258

259 4. Conclusion

We successfully achieved live-cell superresolution imaging using 2D PD-NSIM with rsEGFP2.
 We optimized the data acquisition based on the characteristics of rsEGFP2 to enhance both the
 nonlinearity and modulation contrast. Live-cell imaging was successfully demonstrated with
 sub-80 nm resolution. In the future we plan to use PD-NSIM to perform time-lapse imaging of
 actin dynamics.

Several promising developments could further enhance the utility and performance of 2D
 PD-NSIM. One promising direction is to combine PD-NSIM with TIRF or light-sheet illumination
 which could improve the SNR and modulation contrast, and thereby increase the strength of

- the nonlinear harmonic orders [14, 19]. Another potential direction is extending 2D PD-NSIM 268
- into three-dimensions. By implementing 3D structured illumination patterns and performing 269
- axial scanning superresolution volumetric live-cell imaging could be achieved. Additionally, 270
- integrating Adaptive Optics into PD-NSIM could significantly improve image quality, and may 271
- even be essential for 3D PD-NSIM, particularly for deep tissue imaging, where optical aberrations 272
- pose substantial challenges [22, 23]. 273

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Disclosures 278

The authors declare that there are no conflicts of interest related to this article. 279

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