

# Visualizing endogenous Nanog behavior using a novel nanobody-based toolkit

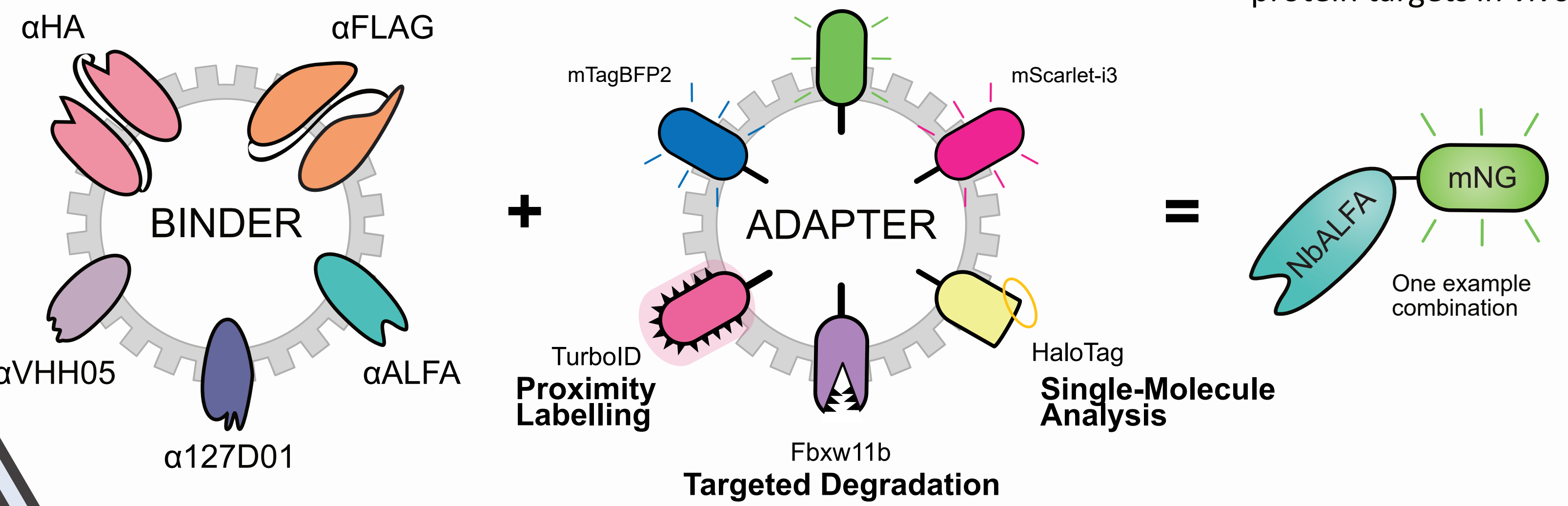
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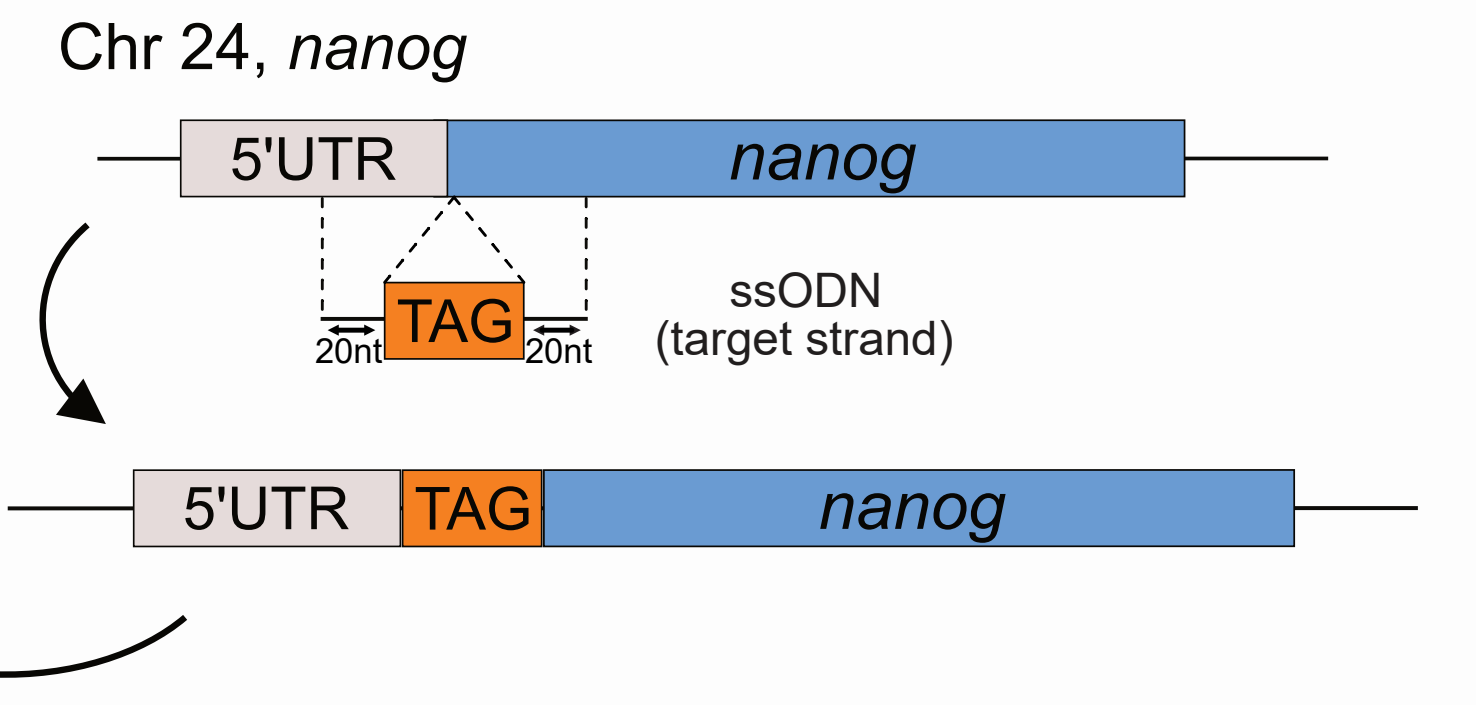


## 1. Genetically encoded affinity reagents (GEARs): A toolkit for visualizing and manipulating endogenous protein function *in vivo*.

Probing endogenous protein localization and function *in vivo* remains challenging due to laborious gene targeting and monofunctional alleles. To address this, we developed a multifunctional toolkit based on genetically encoded affinity reagents (GEARs). GEARs use nanobodies (Nb)<sup>1-3</sup> and single chain variable fragments (scFv)<sup>4-5</sup>, which recognize small epitopes (<15aa), enabling visualization and selective degradation of protein targets *in vivo*.



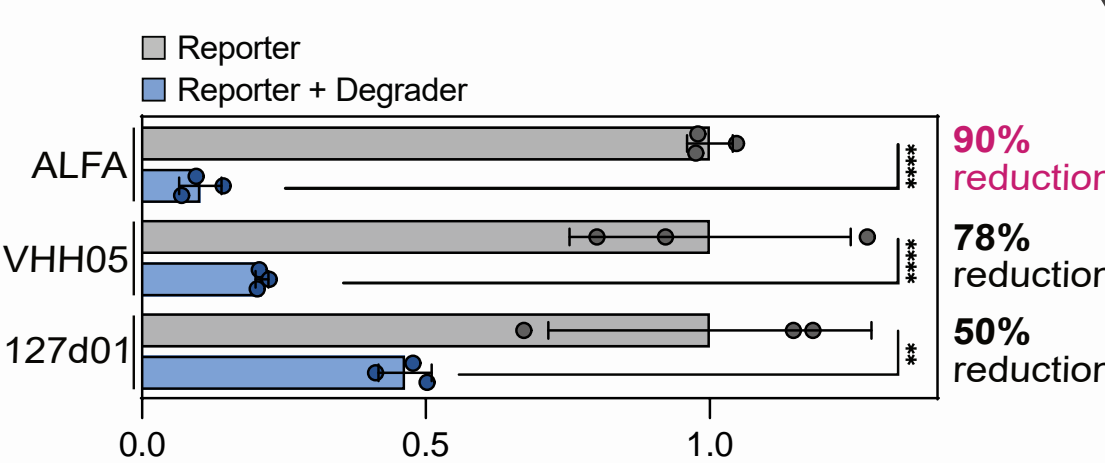
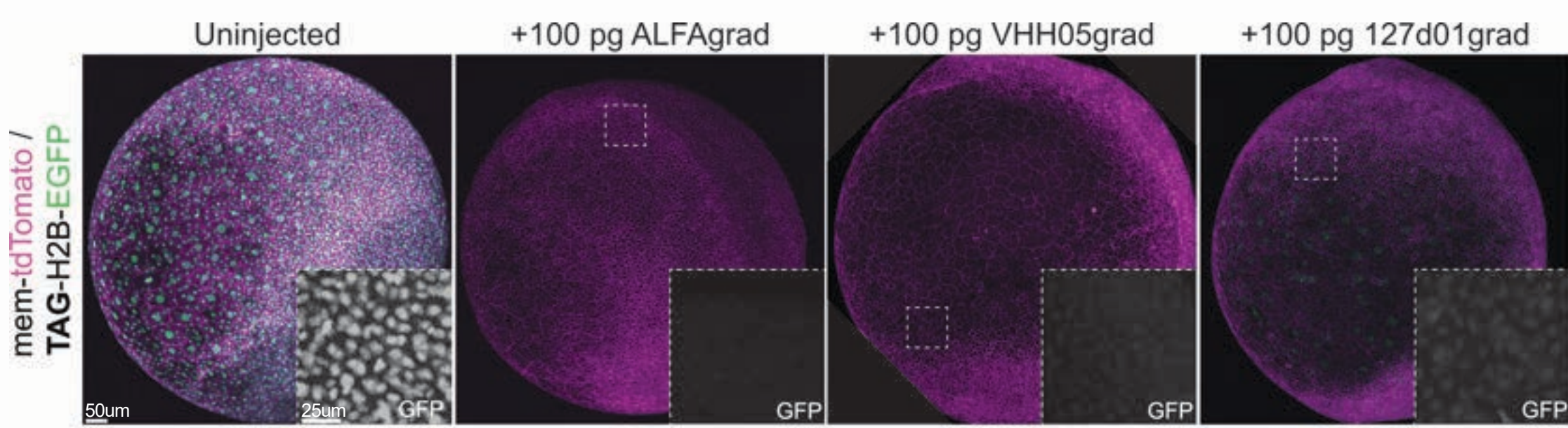
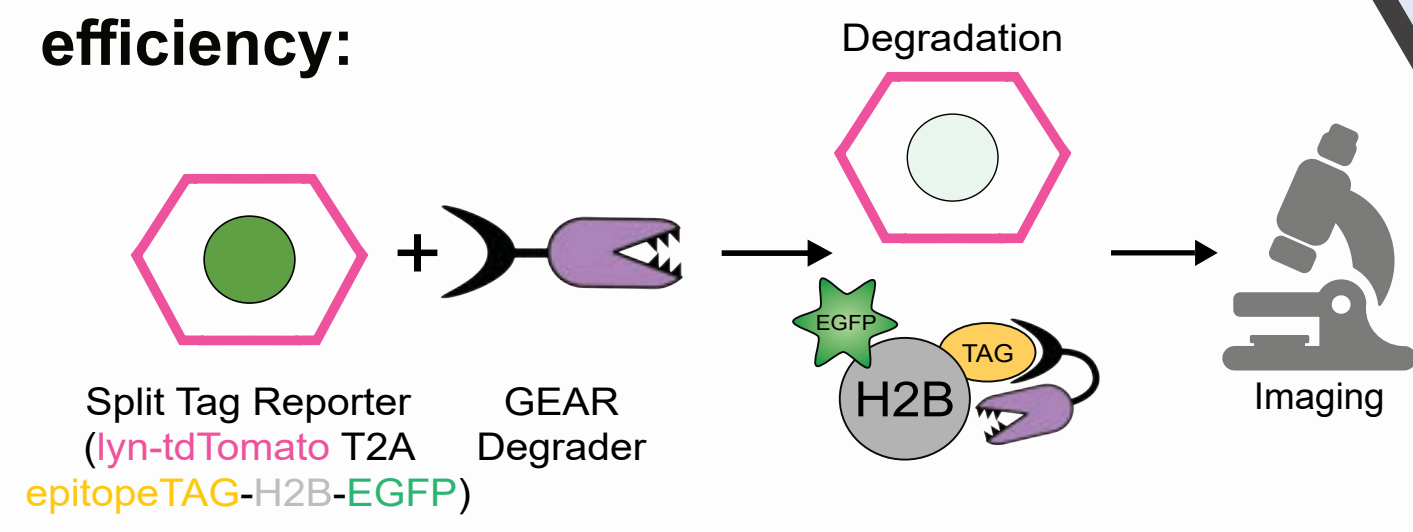
We also established a purely synthetic CRISPR/Cas9-based epitope tagging pipeline to generate endogenous knock-in alleles with high efficiency.



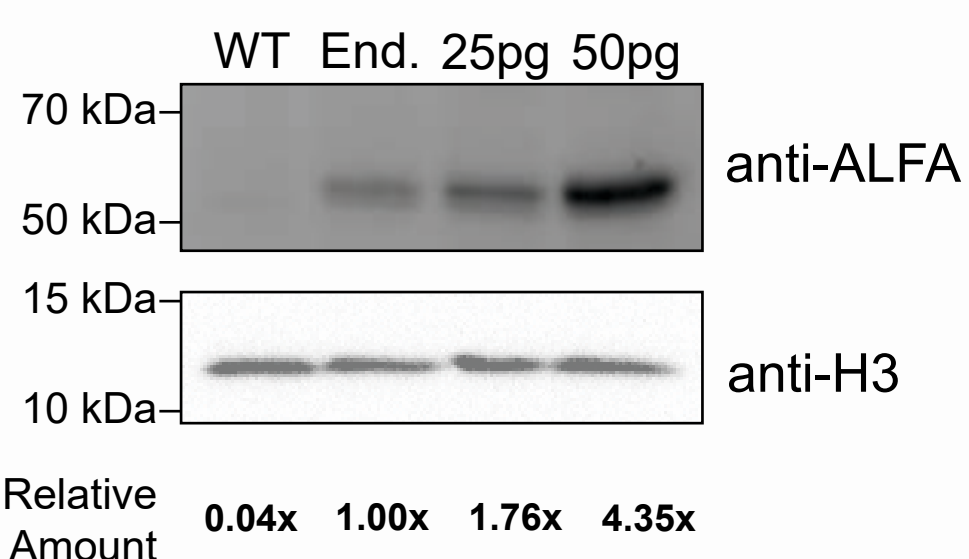
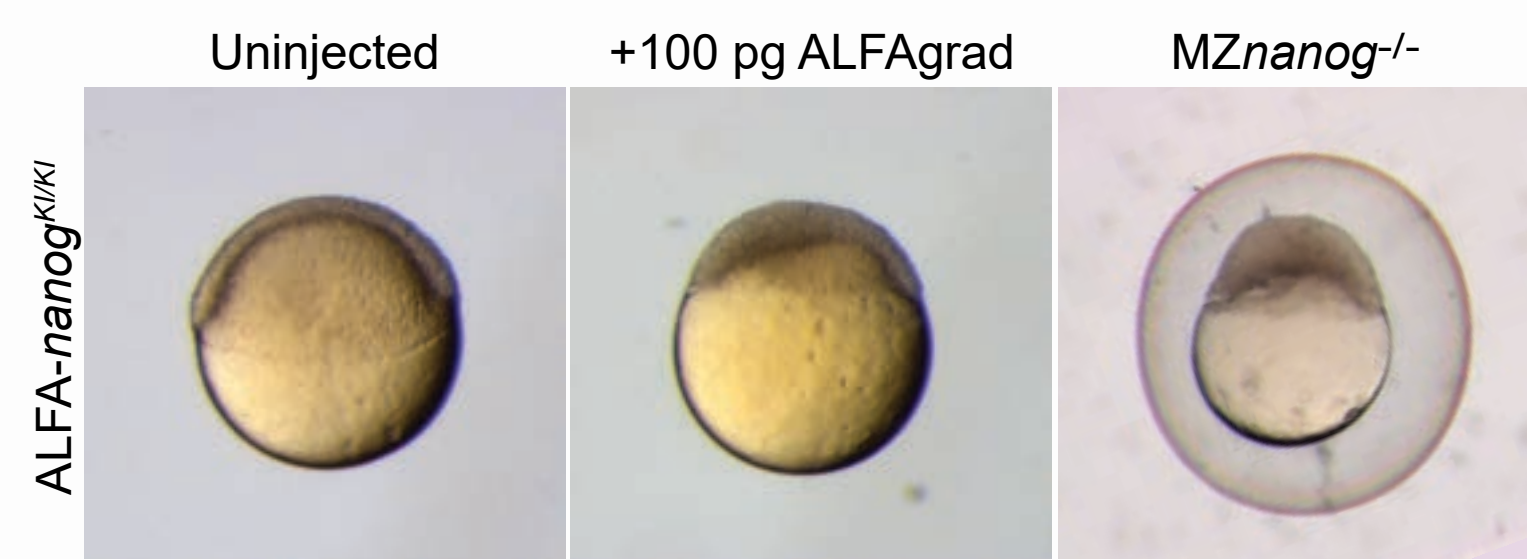
## 2. GEARs can bind and degrade protein targets with high efficiency and mimic loss-of-function mutation

Nanobody-based GEARs were fused to zebrafish Fbxw11b (based on zGrad system<sup>6</sup>) to generate degraders (GEARgrads), and tested for their performance at degrading fluorescent reporters, which were injected into 1-cell staged zebrafish embryos. ALFAgrad performed best with **90% clearance**.

A bicystronic reporter was used to quantify GEARgrad efficiency:



ALFAgrad injection into embryos homozygous for endogenously tagged ALFA-Nanog induced a **maternal-zygotic nanog loss-of-function phenotype<sup>7</sup>** in 100% of the embryos.

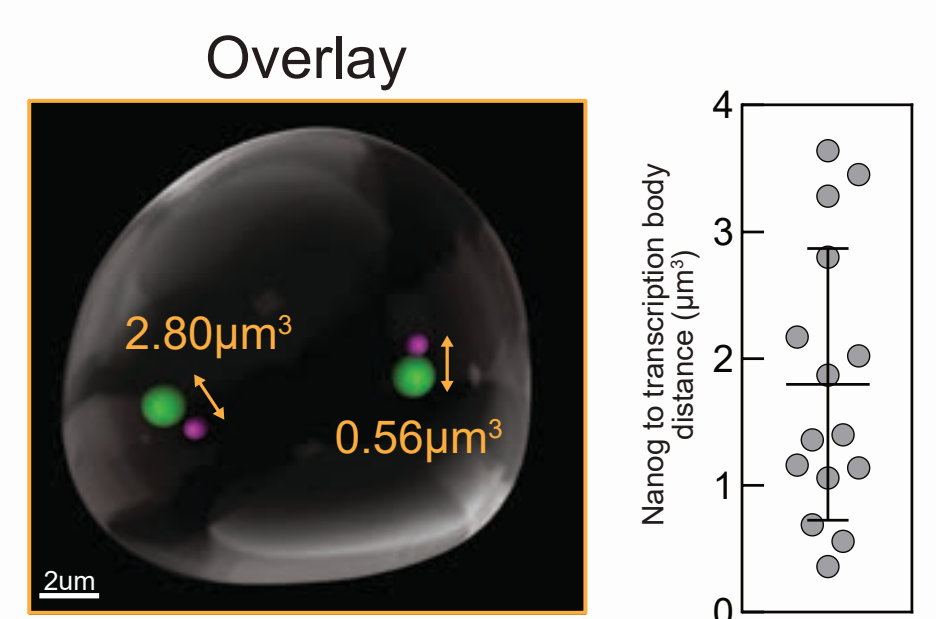
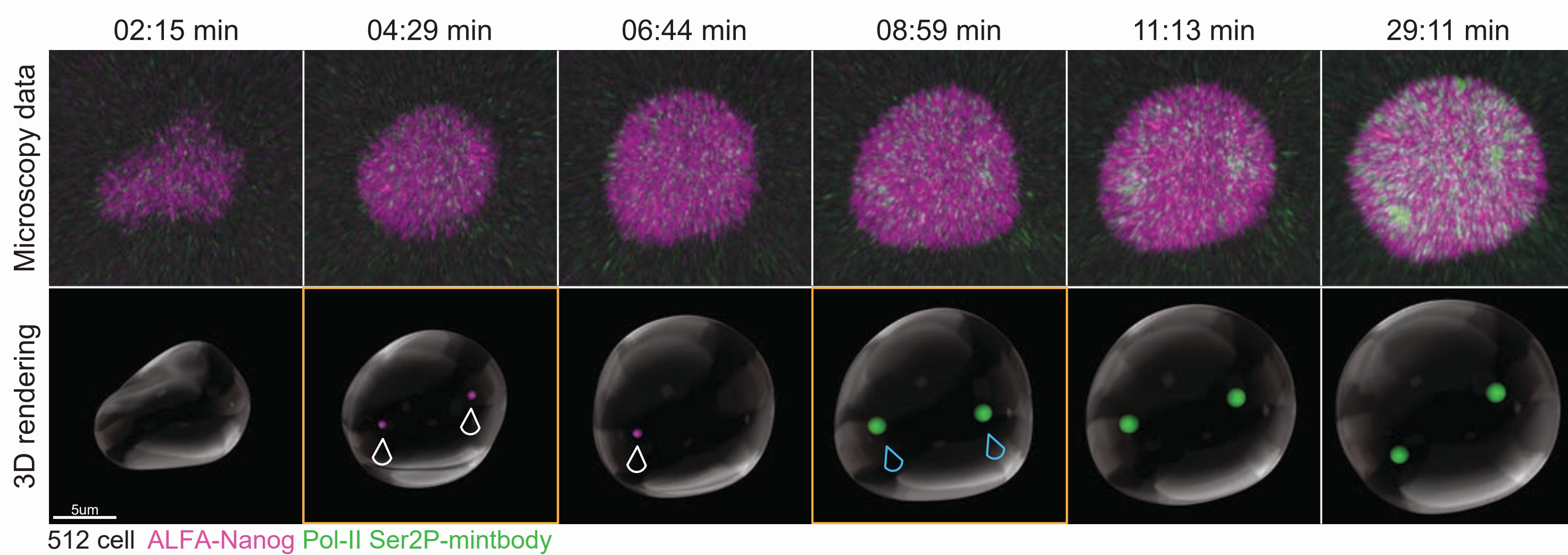
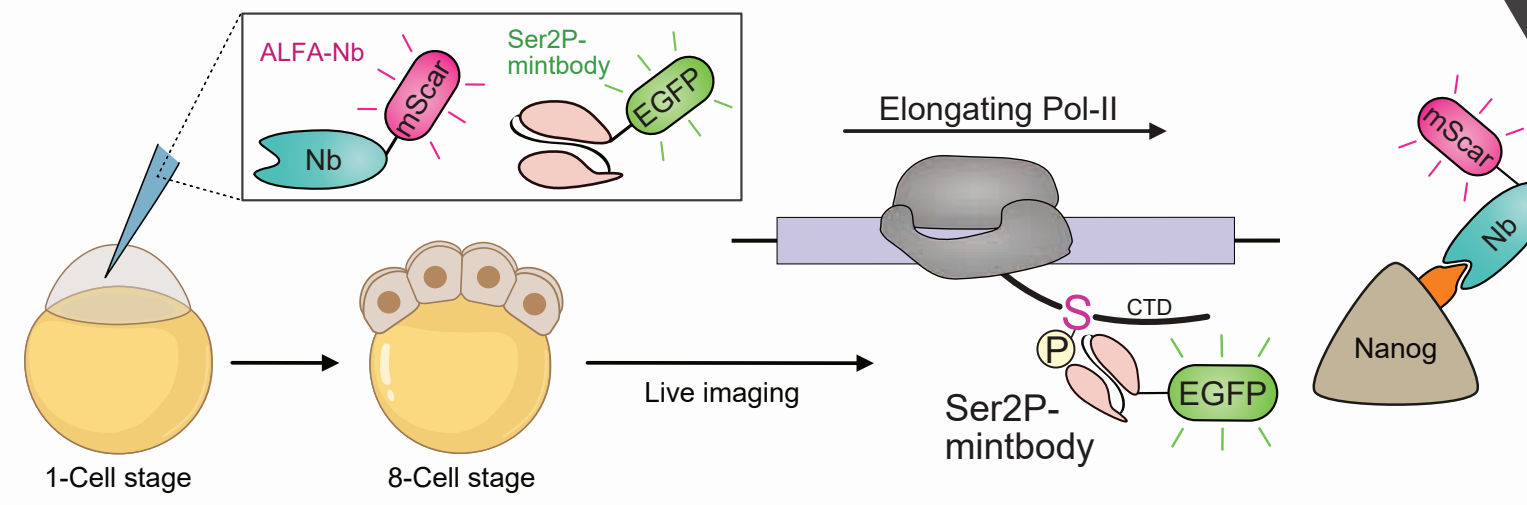


A western blot using anti ALFA antibody showed that the endogenous amount of ALFA-tagged Nanog was nearly half the amount relative to exogenous Nanog.

## 4. Endogenous Nanog foci prime the formation of large transcription bodies

Foci formation of both TFs and active transcription, in the form of transcription bodies, have been observed in a number of systems<sup>8-12</sup>. In zebrafish, the first transcription bodies are seeded at the miR430 locus<sup>8-10</sup>.

Live imaging setup:



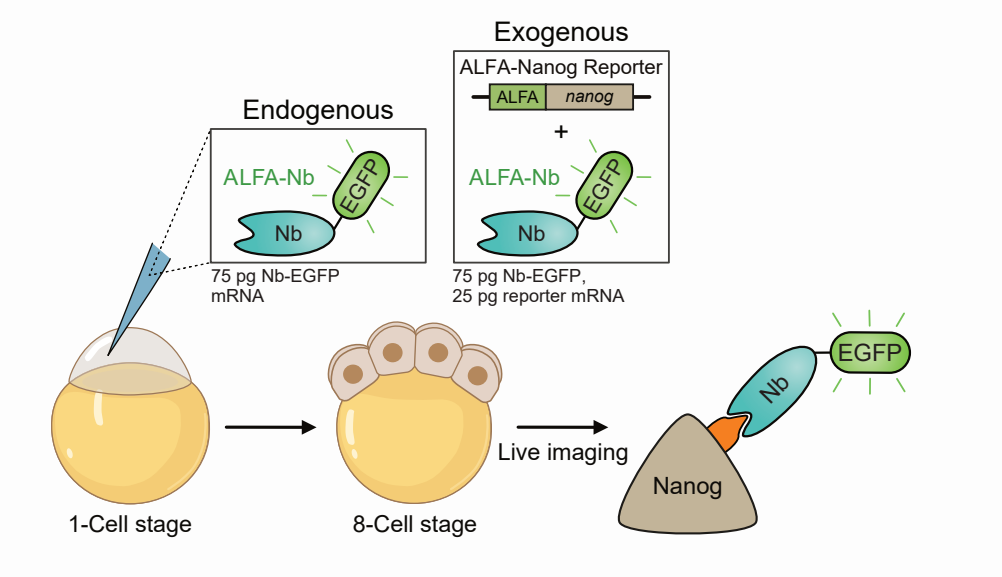
We used an mScarlet-13-ALFA GEAR to detect Nanog protein, as well as a Pol-II-Ser2P<sup>6</sup> EGFP reporter (Mintbody detecting RNA Pol II phosphorylated on Serine 2)<sup>13</sup> to visualize transcriptional elongation.

Like exogenous studies, we observed that Nanog foci precede transcription bodies in close proximity but are rapidly dissolved while transcription bodies are long lived

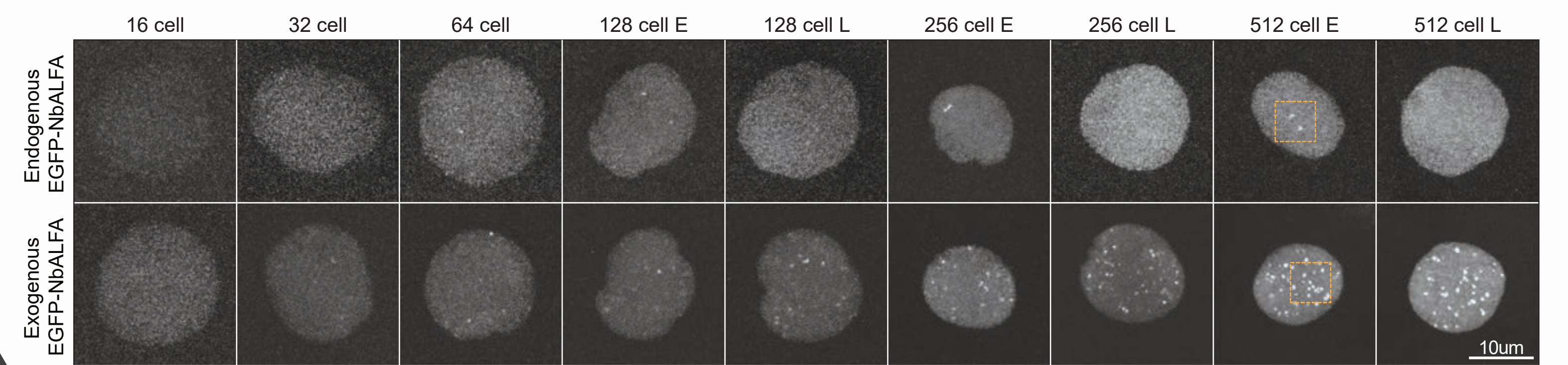
## 3. GEARs visualize the native behavior of Nanog during the earliest transcriptional events in zebrafish embryos

Using GEARs we investigated the spatiotemporal dynamics of endogenous Nanog protein. Nanog is a pioneer transcription factor that has intrinsically disordered regions that engage in concentration dependent interactions and exogenous fluorescently tagged Nanog localizes to subnuclear puncta during the maternal-to-zygotic transition<sup>8-9</sup>. While live imaging of exogenous Nanog has revealed important molecular behaviors<sup>8-9</sup>, the localization of endogenous Nanog has not been studied due to the lack of appropriate tools.

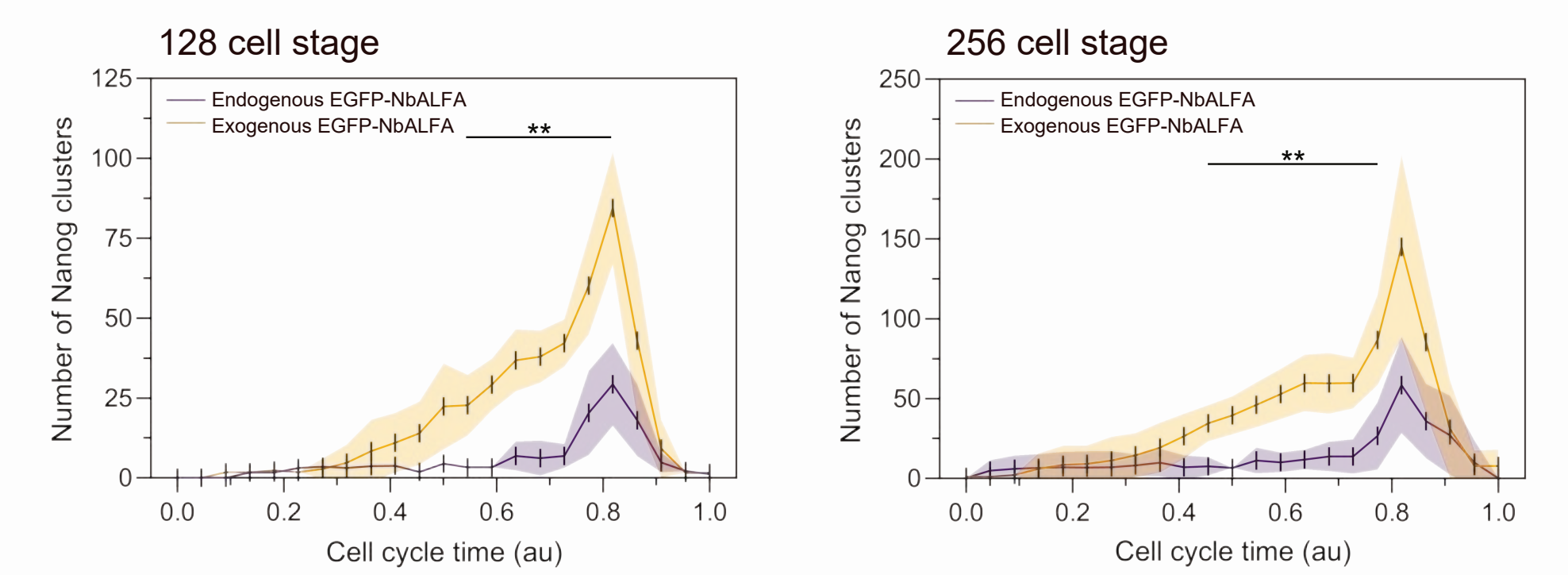
Live imaging setup:



### Endogenous vs exogenous Nanog visualization



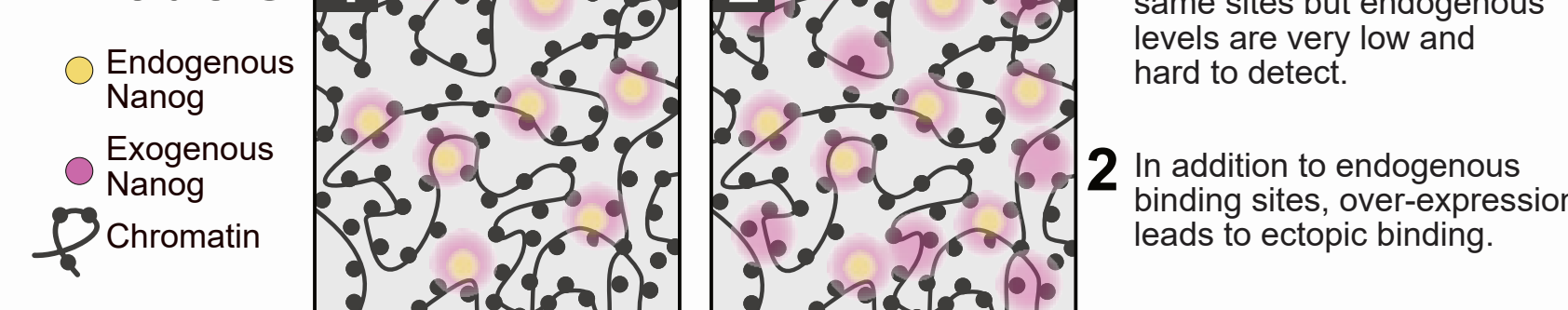
Endogenous Nanog protein forms significantly fewer foci than overexpressed Nanog.



## 5. Conclusions

- GEARs are powerful and versatile tools to interrogate endogenous proteins. The short epitope tags required can be introduced into the genome with high efficiency.
- GEARs can be used to efficiently degrade proteins and furthermore target the pool of maternal proteins, enabling robust clearance of proteins that establish the earliest events of embryogenesis.
- ALFA-nanog knock-in fish report the physiological concentration of endogenously produced Nanog protein, which is found at much lower levels than previously appreciated.
- Large difference in Nanog foci number suggests that overexpressed Nanog can either seed new foci or enlarge existing foci, most likely aided by its IDRs.

### Models



## A little bit extra: GEARs function also in other organisms



ALFA-tag reporter mRNA was injected into one-cell stage mouse zygotes and the ALFAgrad in one of the two cells at the 2 cell stage, such that the uninjected cell acts as no degradation control. We saw robust clearance of nuclear EGFP signal in the degraon injected side, with a clearance efficiency of 96%.

The GEARs toolkit provides an easy plug-and-play approach allowing users to generate endogenously tagged alleles with limitless versatility and custom control over protein function *in vivo*.

### References

- Götzke et al., 2019 Nat Comms.
- Ling et al., 2019 Mol Immunol.
- Bradley et al., 2015 Mol Pharmacol.
- Zhao et al., 2019 Nat Comms.
- Liu et al., 2021 Genes Cells.
- Yamaguchi et al., 2019 Elife.
- Gagnon et al., 2018 Dev.
- Kuznetsova et al., 2023 Curr Biol.
- Pownall et al., 2023 Science.
- Hadzhiev et al., 2023 Dev. Cell.
- Cissé et al., 2013 Science.
- Mir et al., 2017 Genes Dev.
- Shibuta et al., 2021 Commun Biol.

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This work is supported by



Live imaging movies