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Article

# All-RNA-mediated targeted gene integration in mammalian cells with rationally engineered R2 retrotransposons

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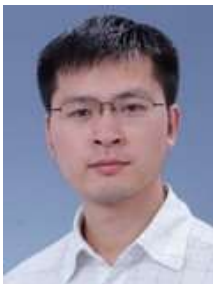
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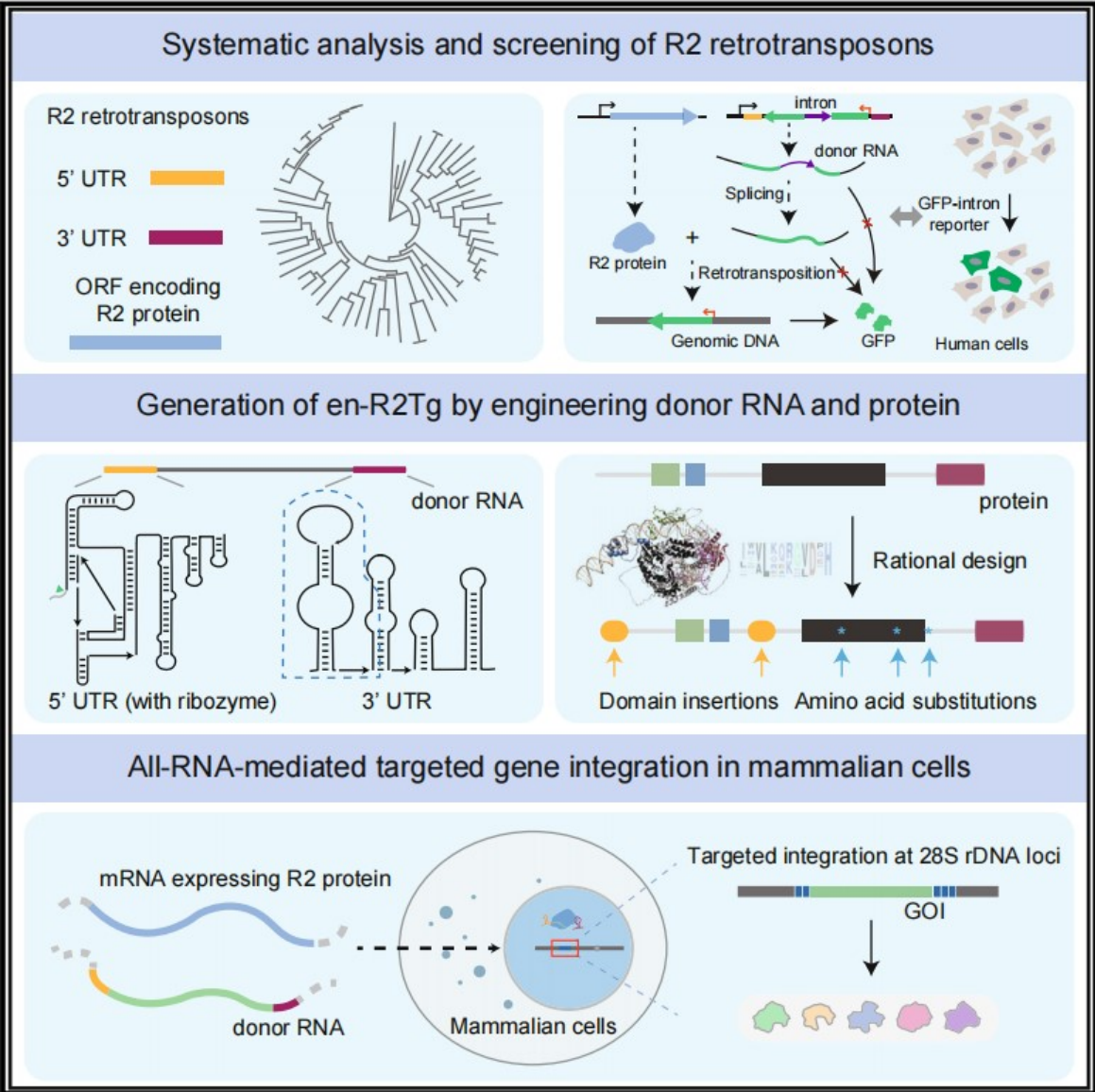
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- 干细胞与再生医学
- 干细胞应用研究与转化



筛选

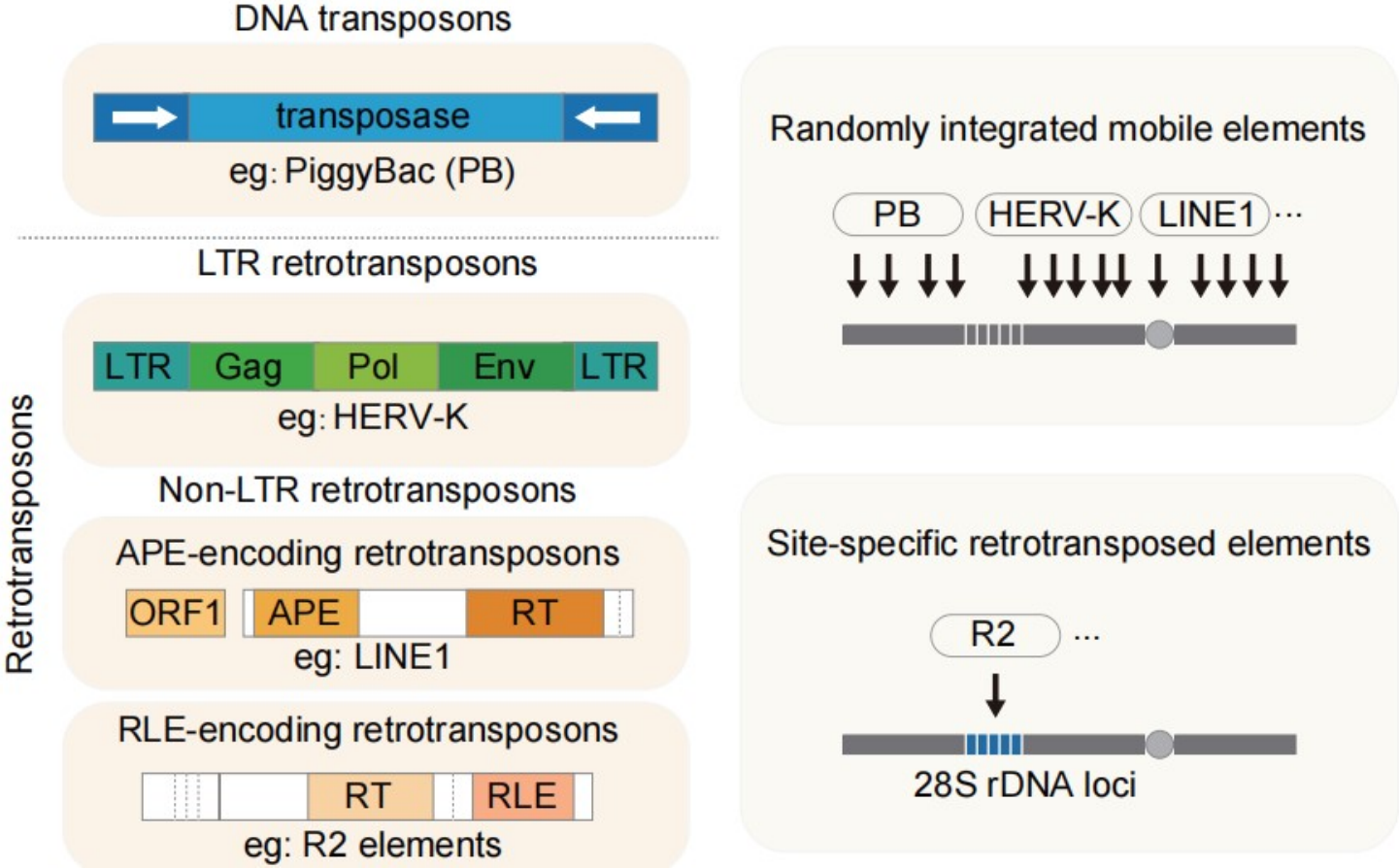


改造



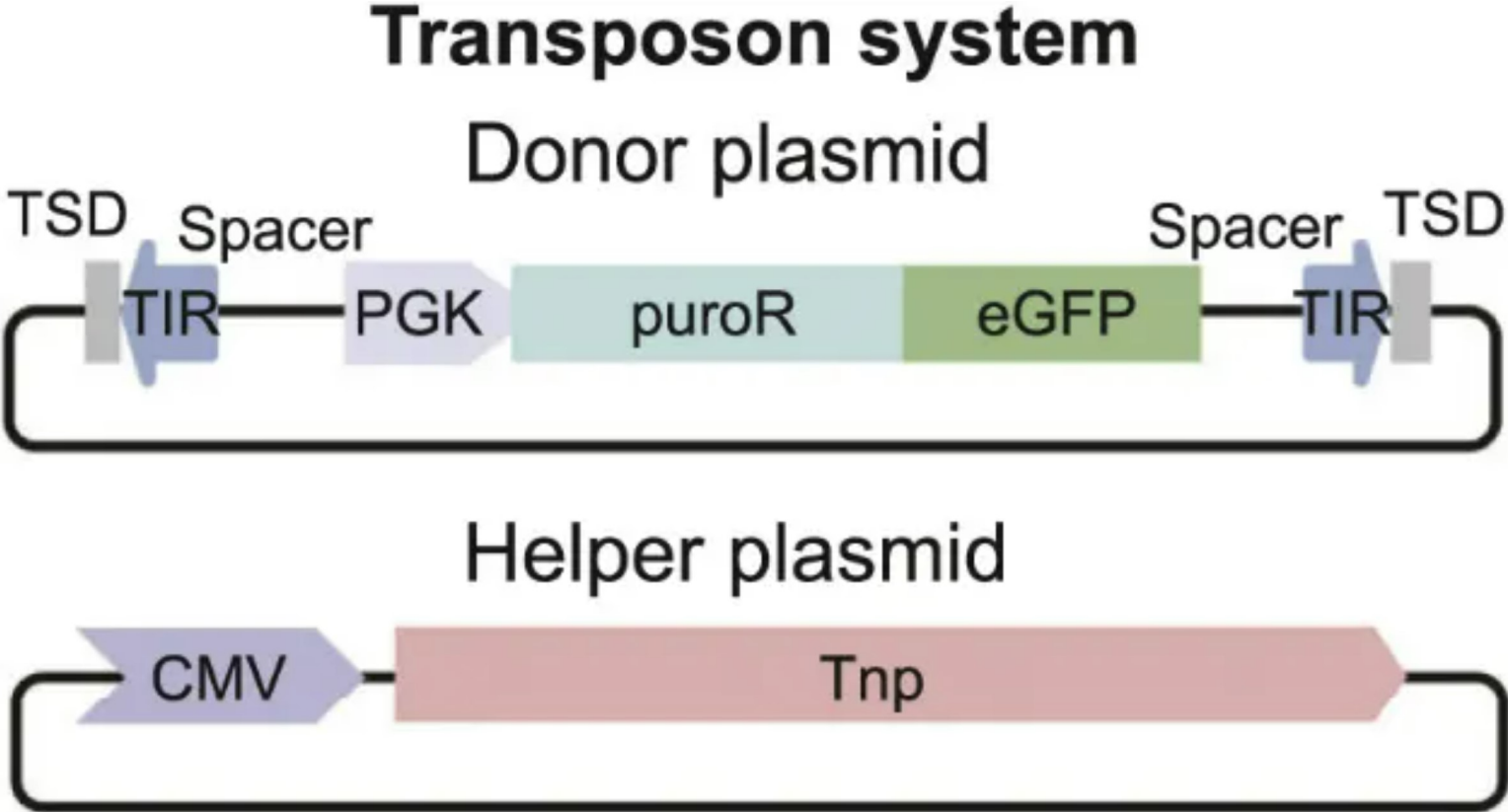
功能验证

# Overview of Transposons



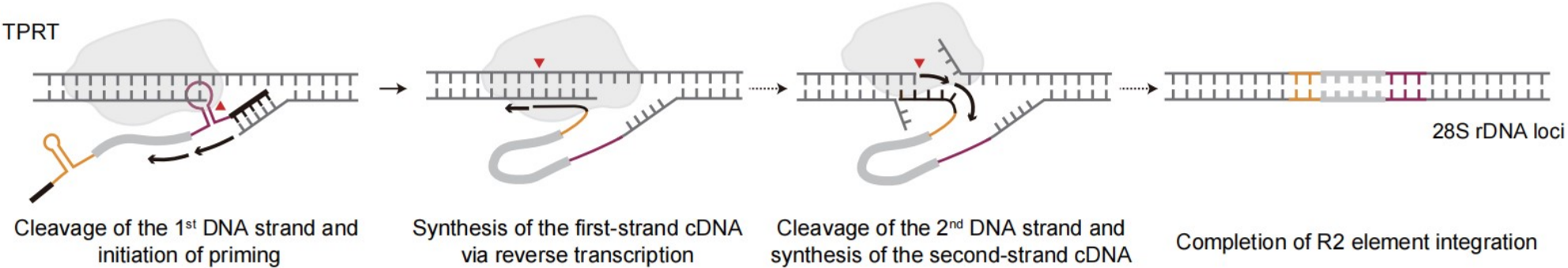
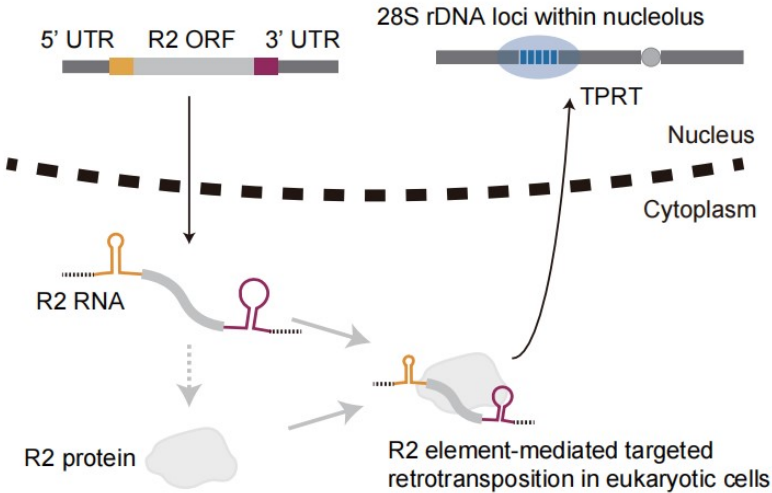
R2 is a site-specific retrotransposon targeted only 28S rDNA.

Typical gene integration system designed from DNA transposons.



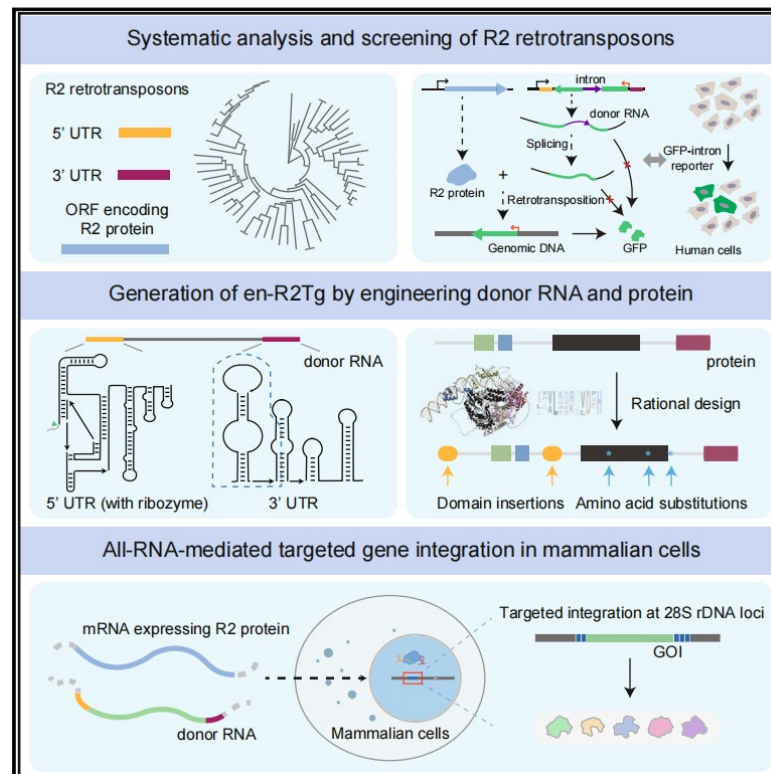
# Advantages of R2 to target gene integration

- Mature and efficient ssRNA delivery technology.
- Safety (target 28S rDNA, does not disturb functional mRNAs)

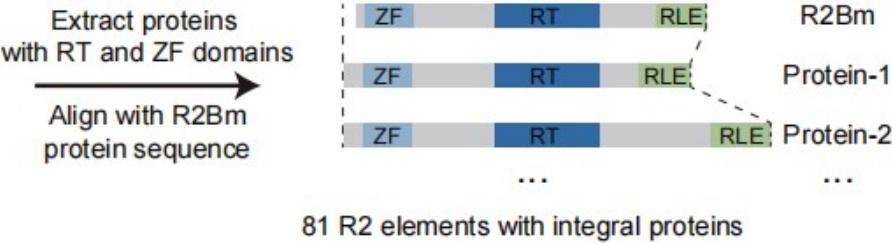
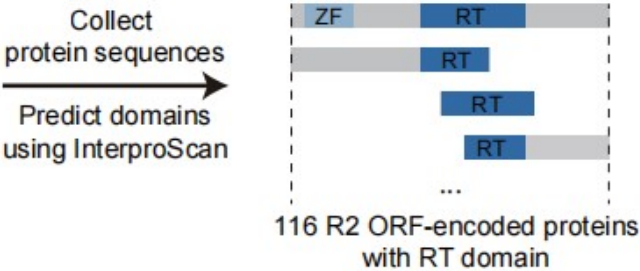
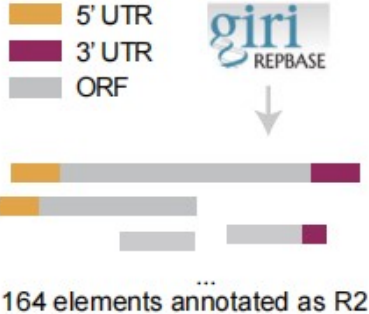


## Study Objectives

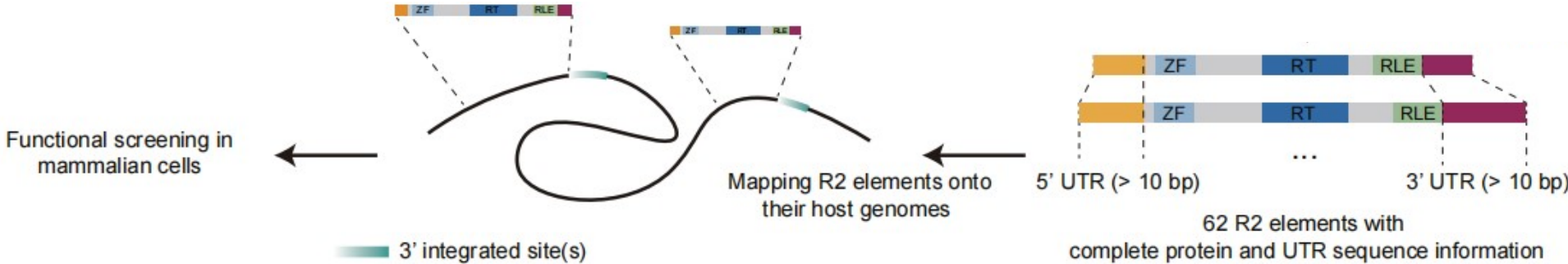
- Systematic R2 retrotransposons analysis
- Engineering R2 elements for enhanced integration
- Demonstrating all-RNA-mediated targeted integration in mammalian cells



Bioinformatics screened 62 complete R2 (integral for transposition)

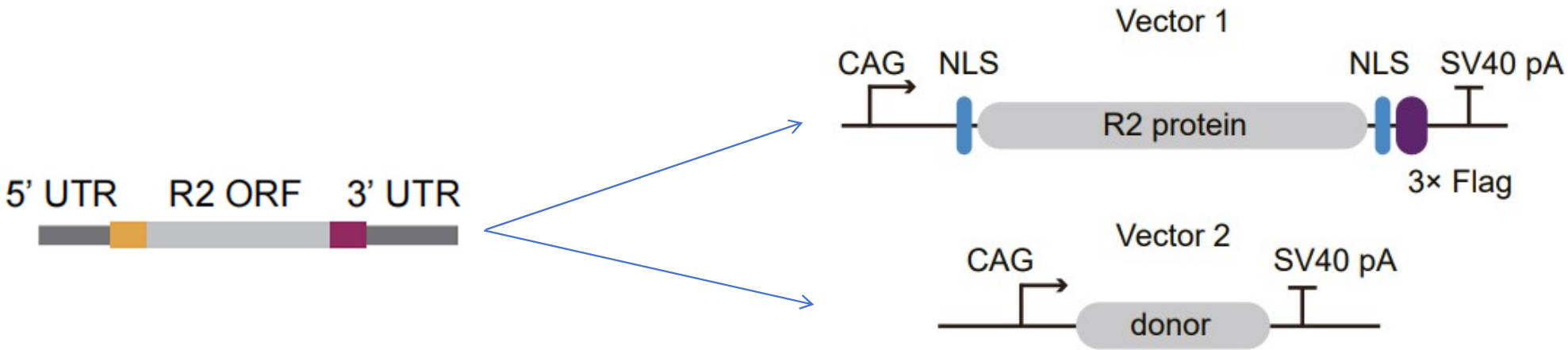


Extract R2 elements with 5' UTR and 3' UTR information

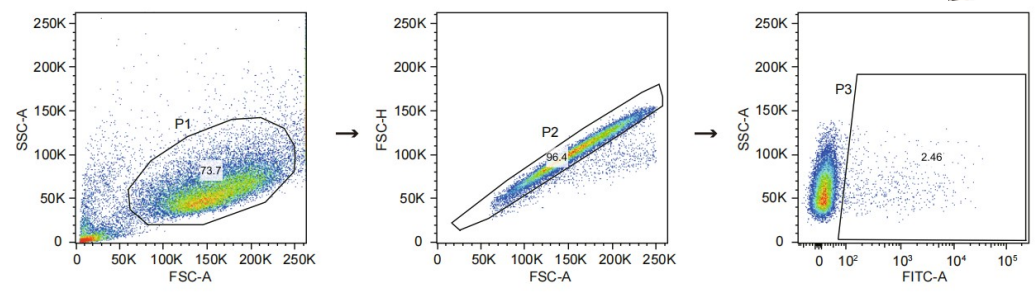
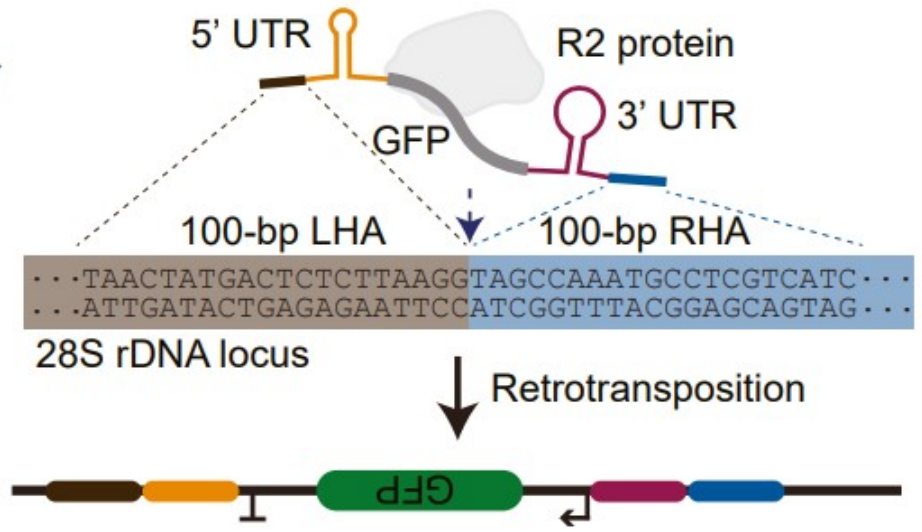
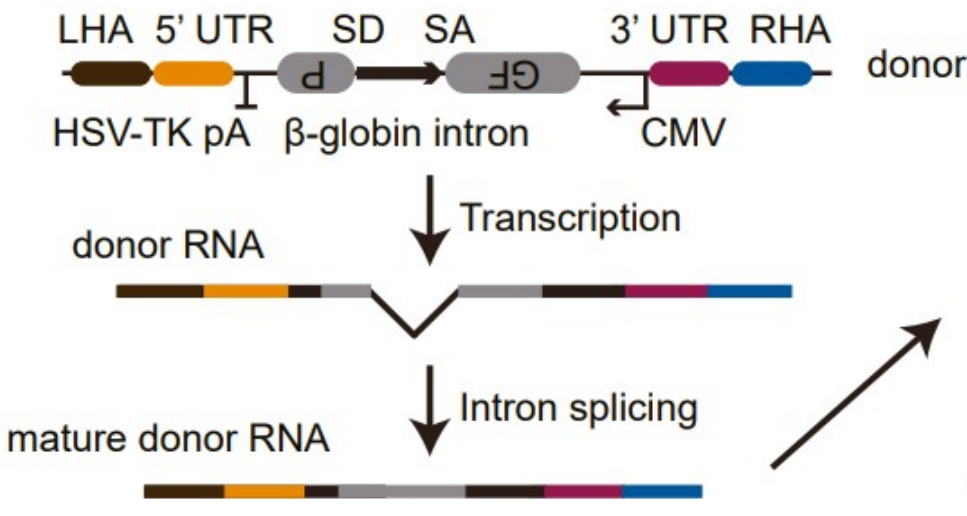




Modular expression system

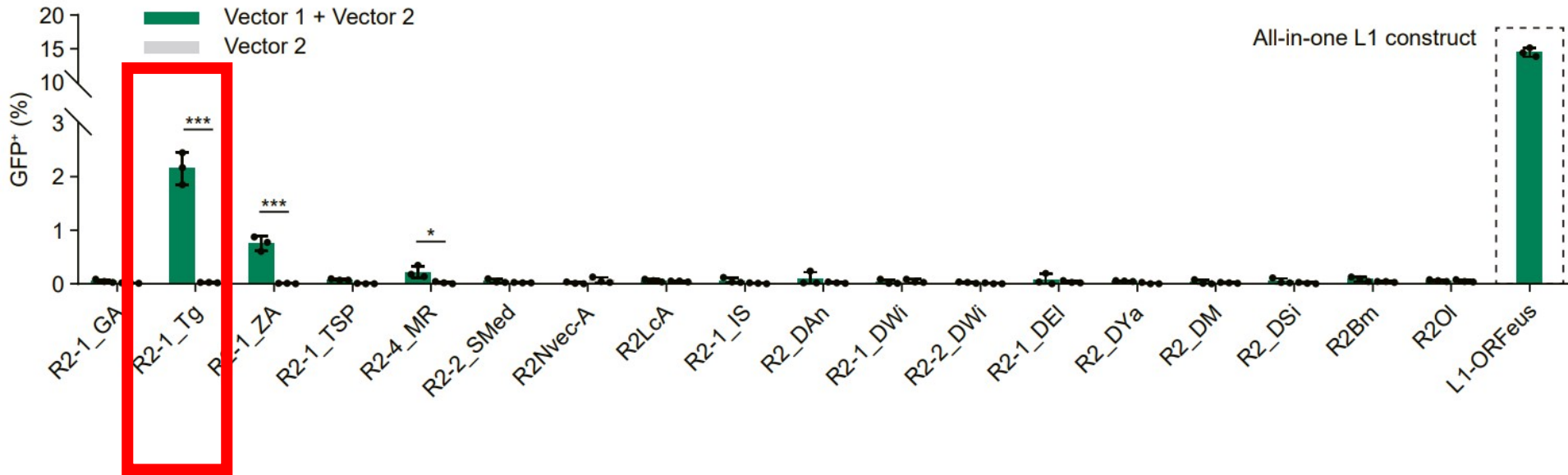


# GFP-intron reporter system



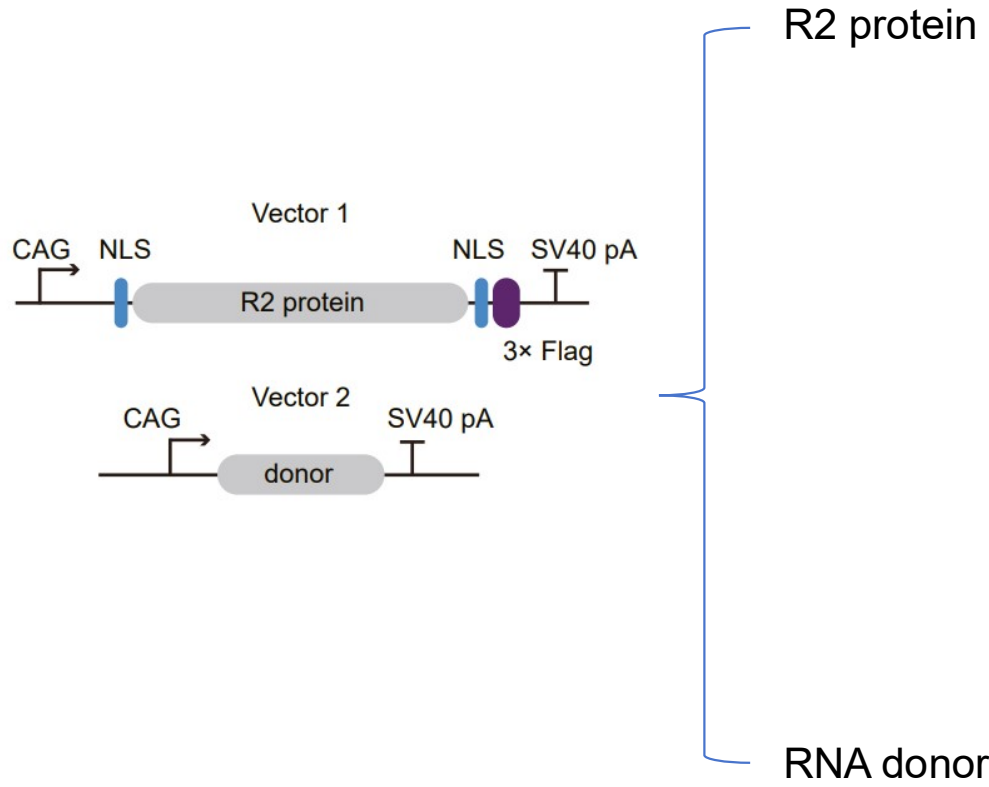
Calculate GFP+ (%) rate

# R2Tg has optimal retrotransposition activity

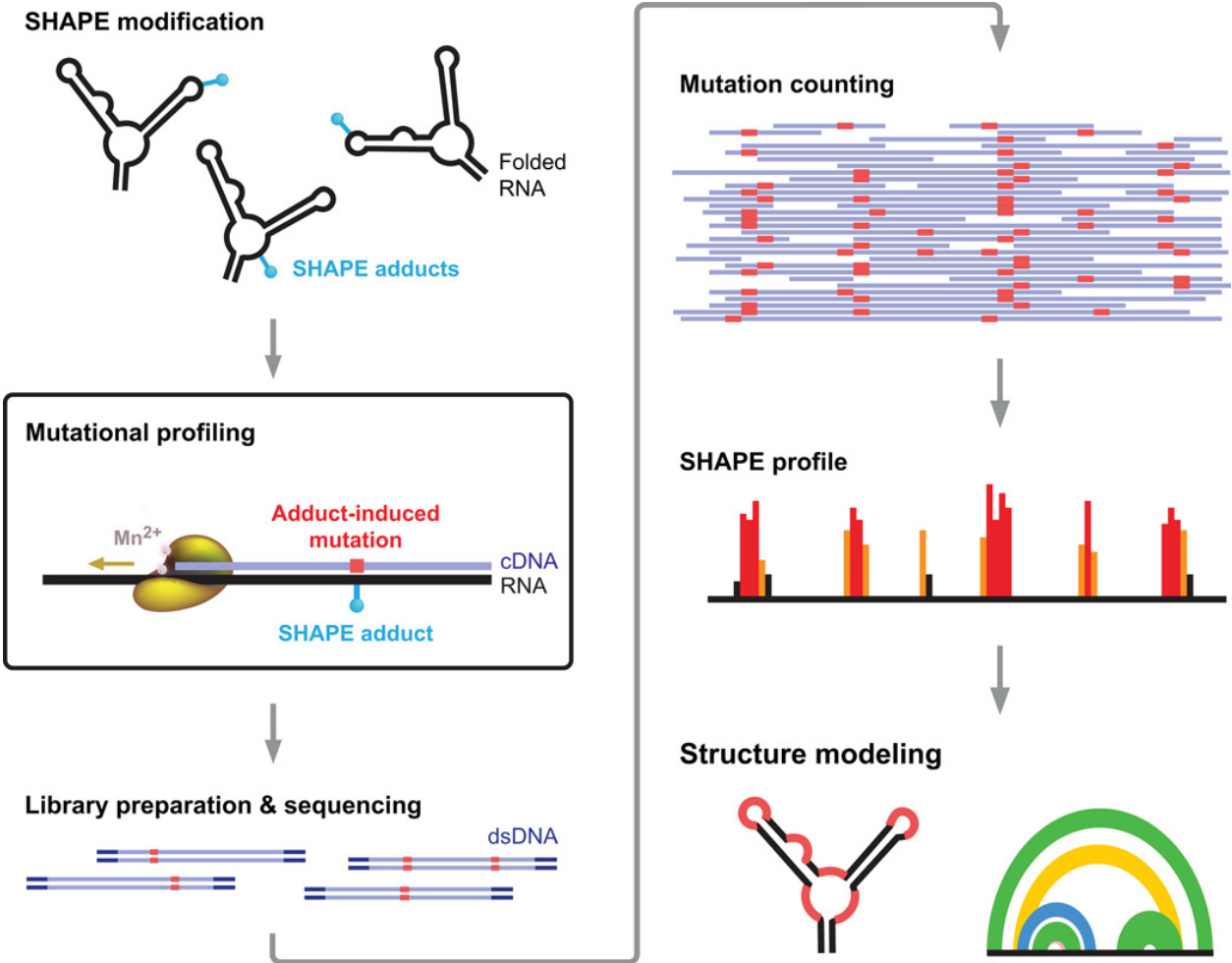


An, W. et al. Characterization of a synthetic human LINE-1 retrotransposon ORFeus-Hs. Mobile DNA, 2011.

# Engineering strategy

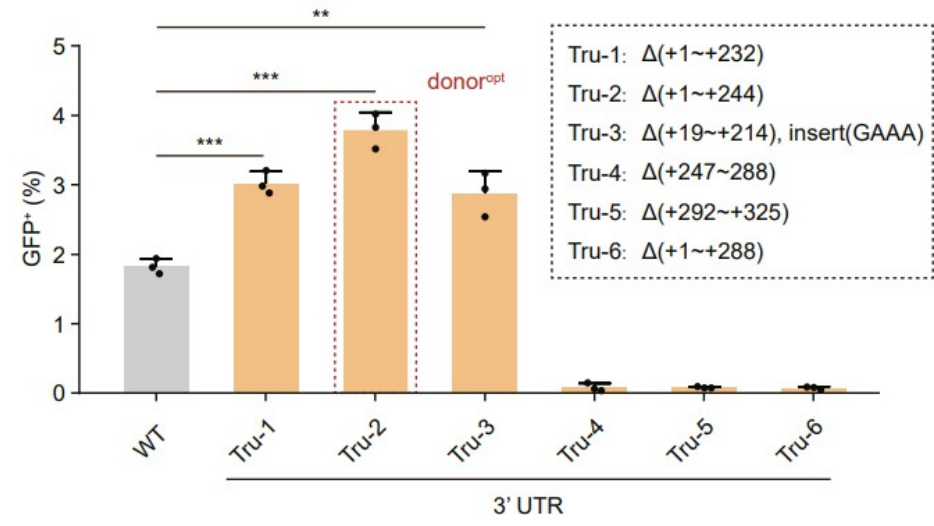
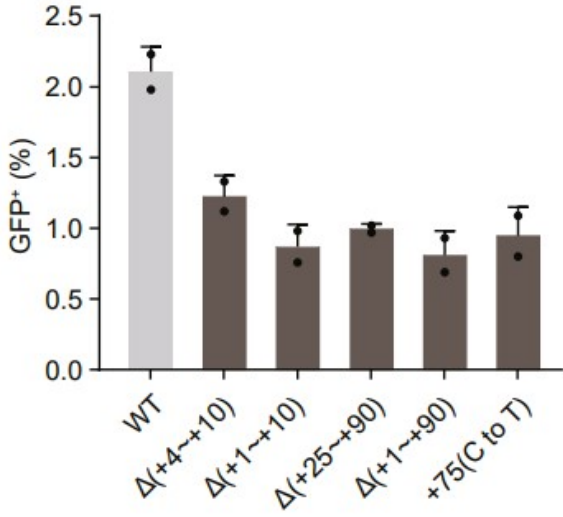
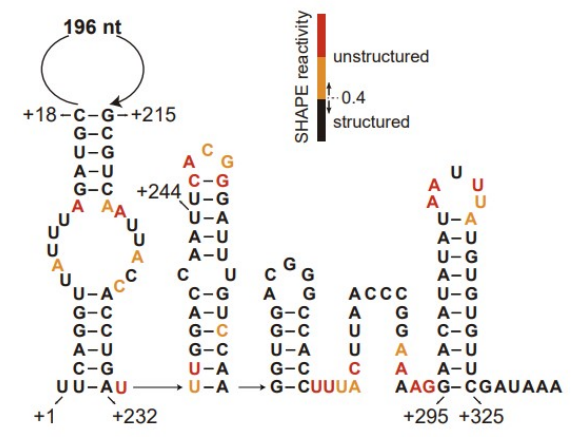
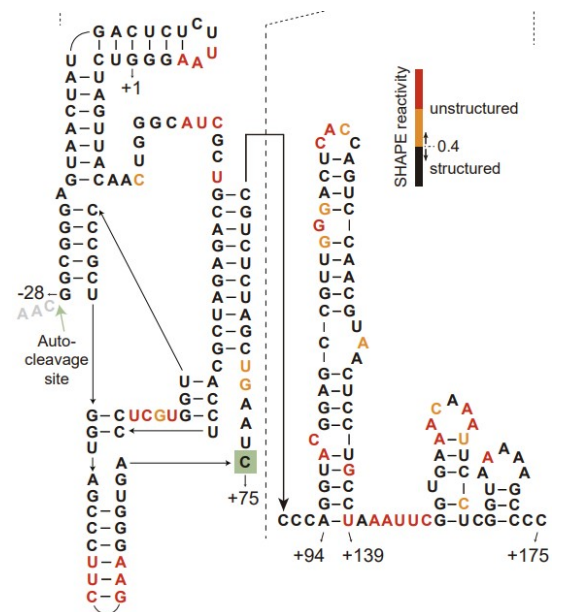


# RNA structure determined by SHAPE-Map

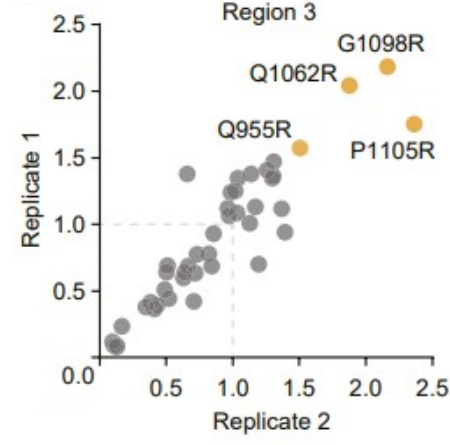
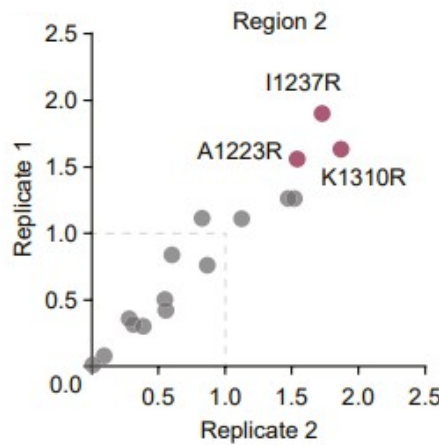
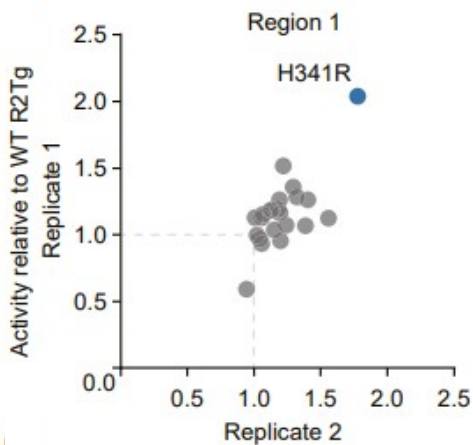
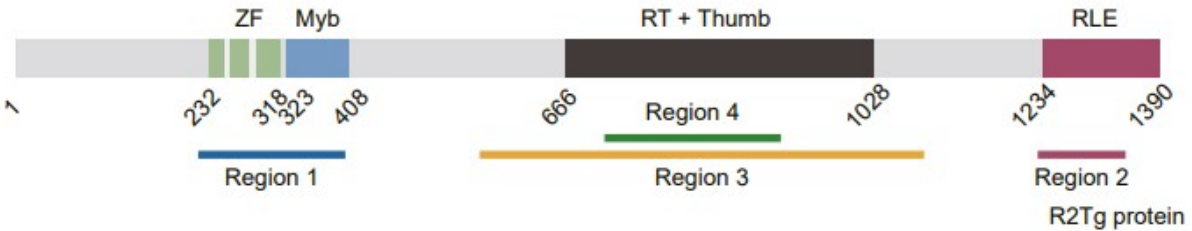


Siegfried NA et al., "RNA motif discovery by SHAPE and mutational profiling (SHAPE-MaP)," Nat Methods, 2014.

# Truncation of 3'UTR (Tru-2, donor<sup>opt</sup>) enhances integration.

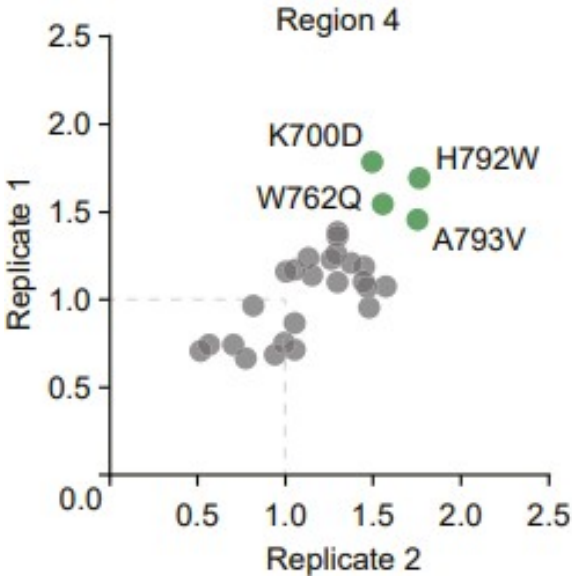
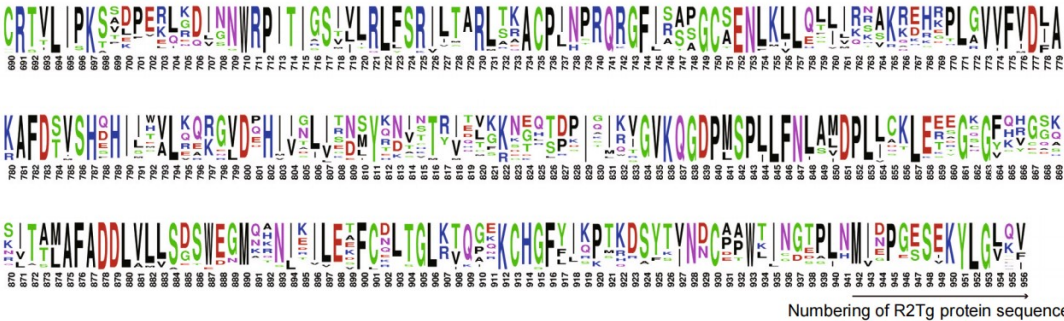
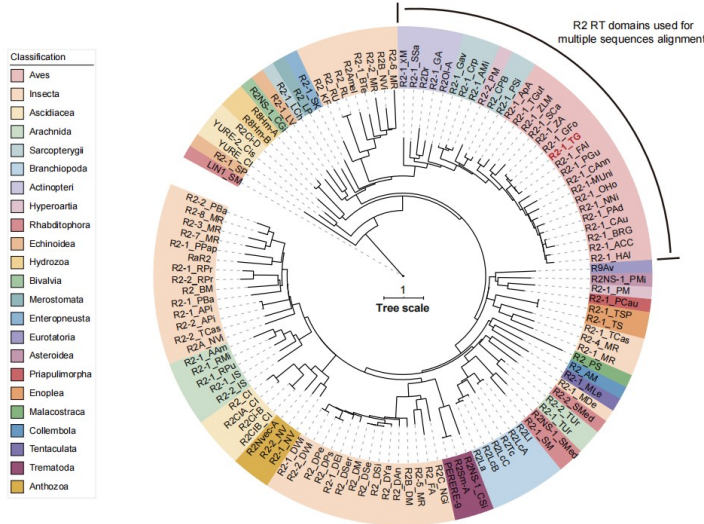
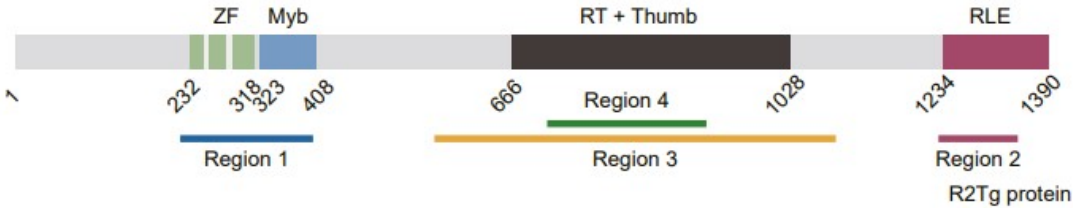


# Point mutation in R2 protein



Strategy 1. Arginine with positive charge enhances DNA binding ability.

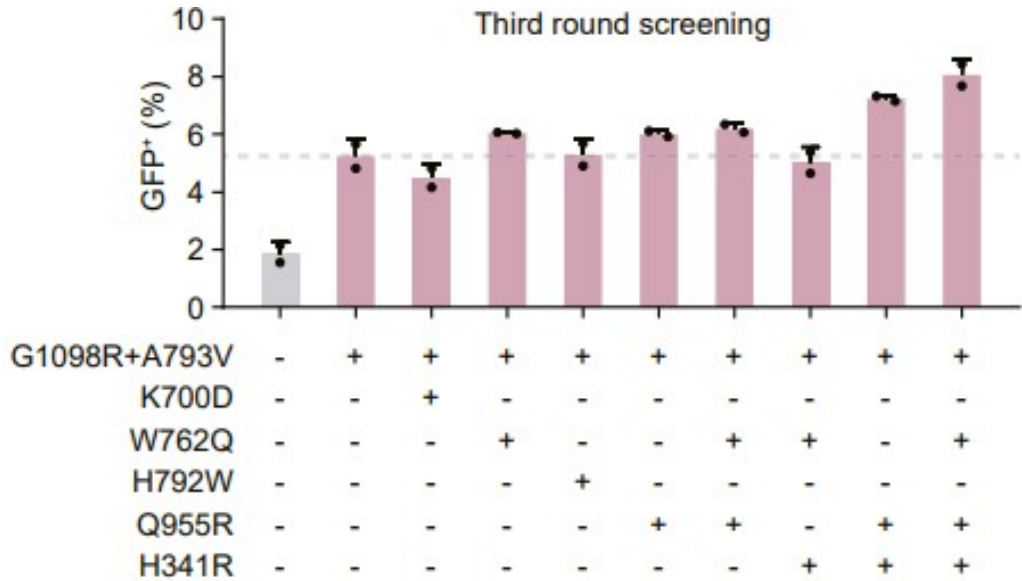
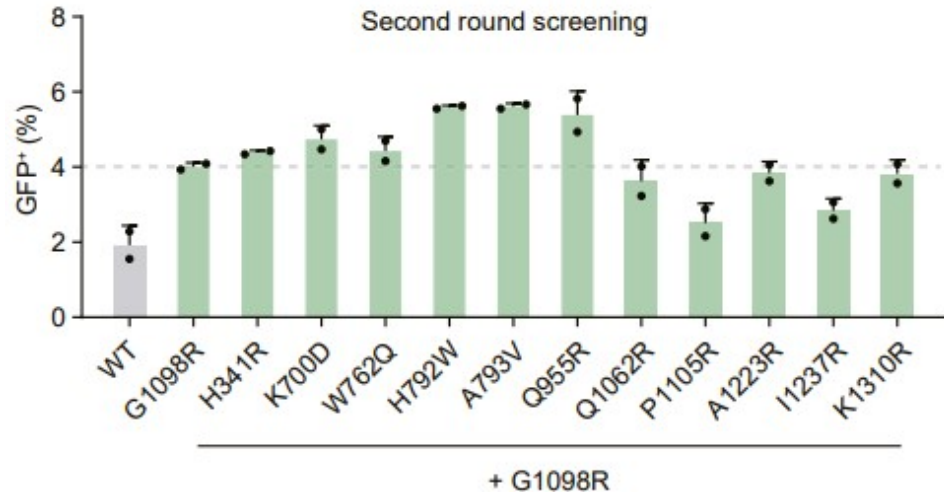
# Point mutation in R2 protein



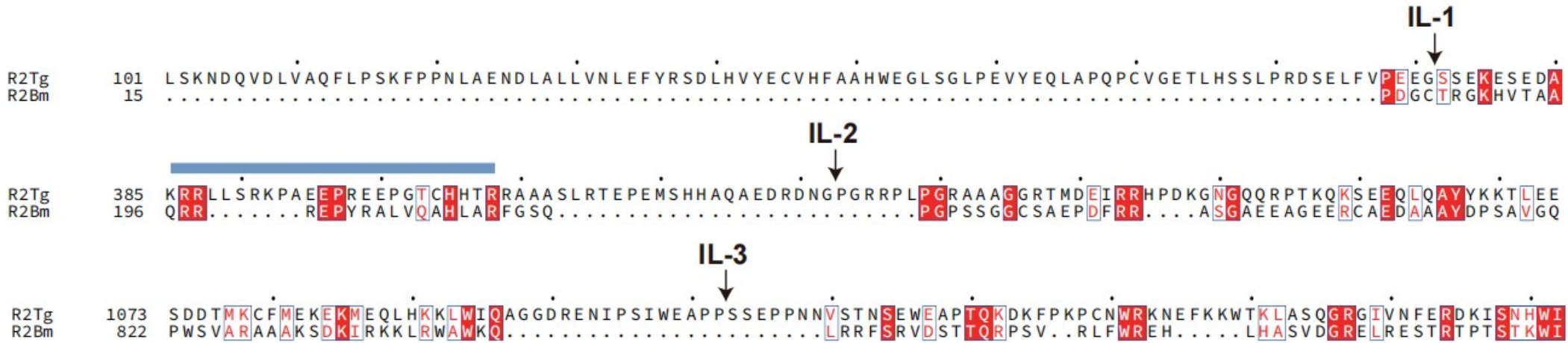
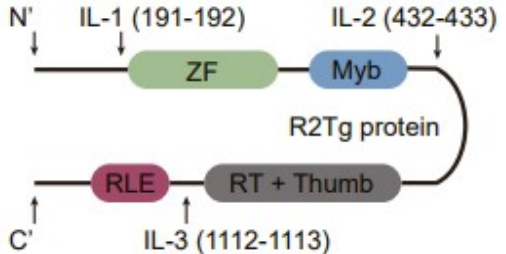
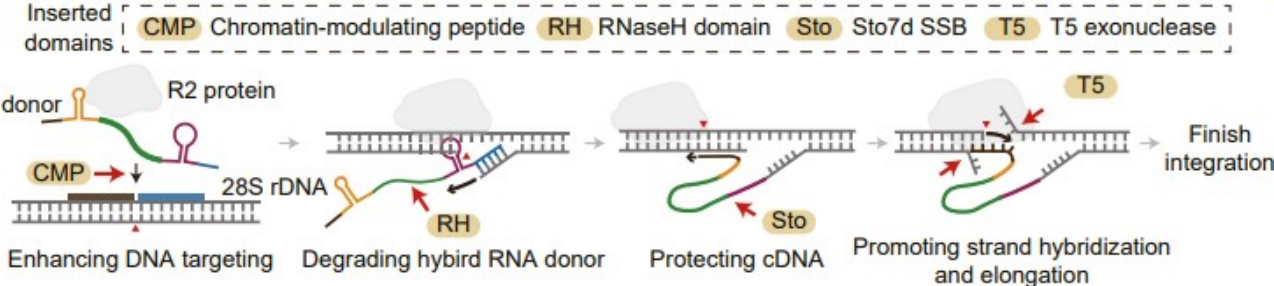
Strategy 2. Integrate consensus amino acids of RT domain.



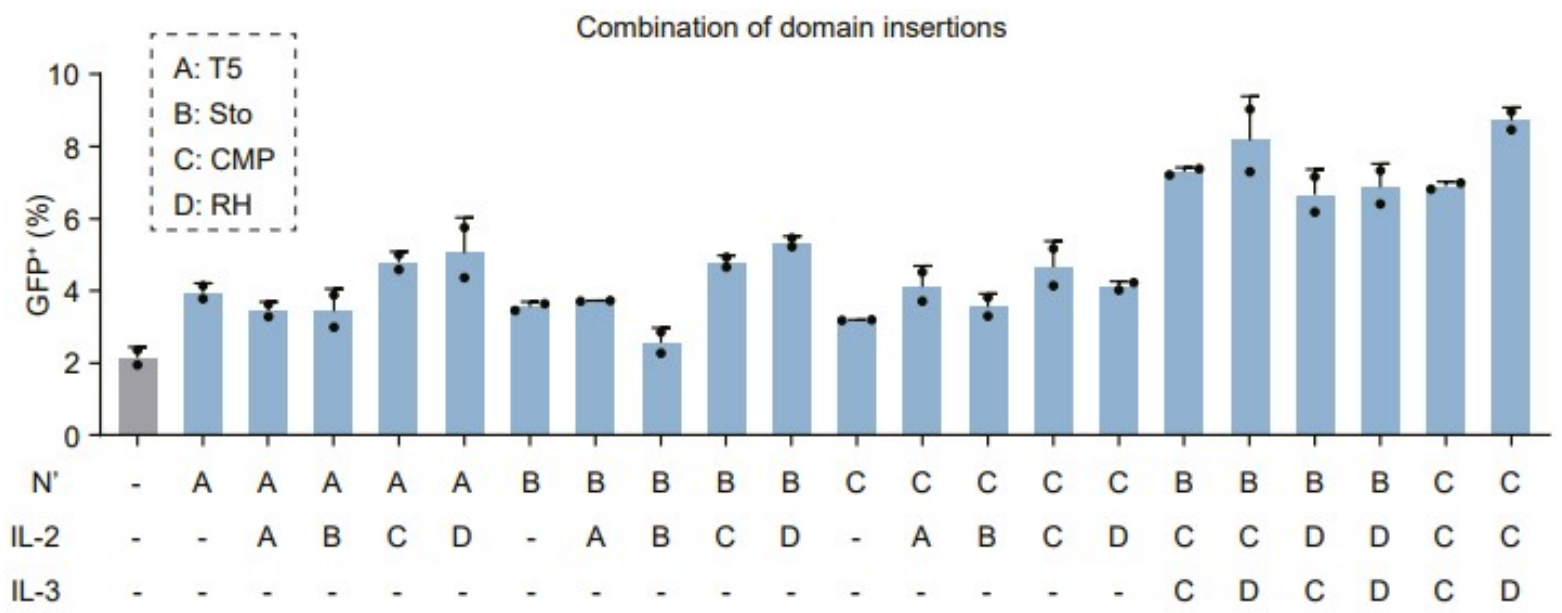
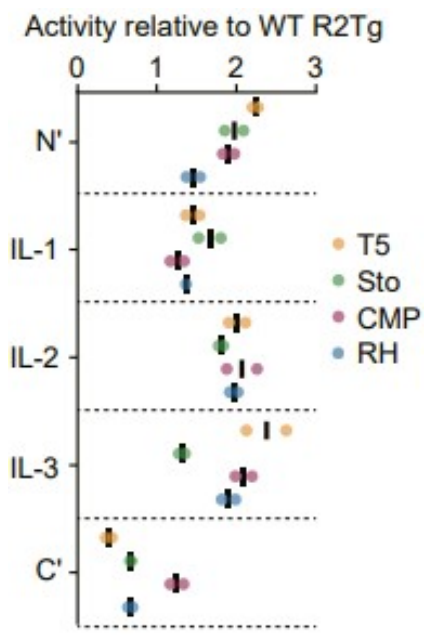
Point mutation in R2 protein



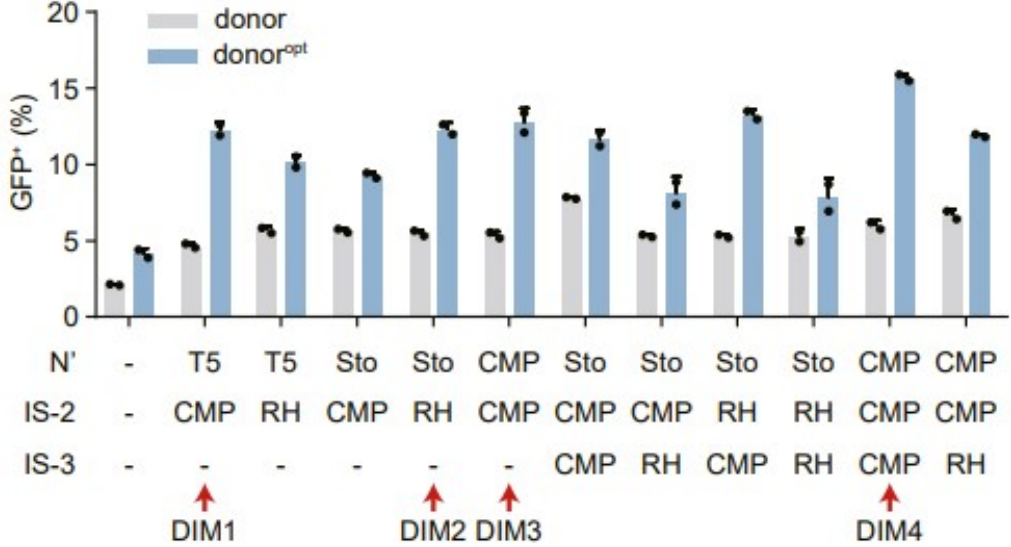
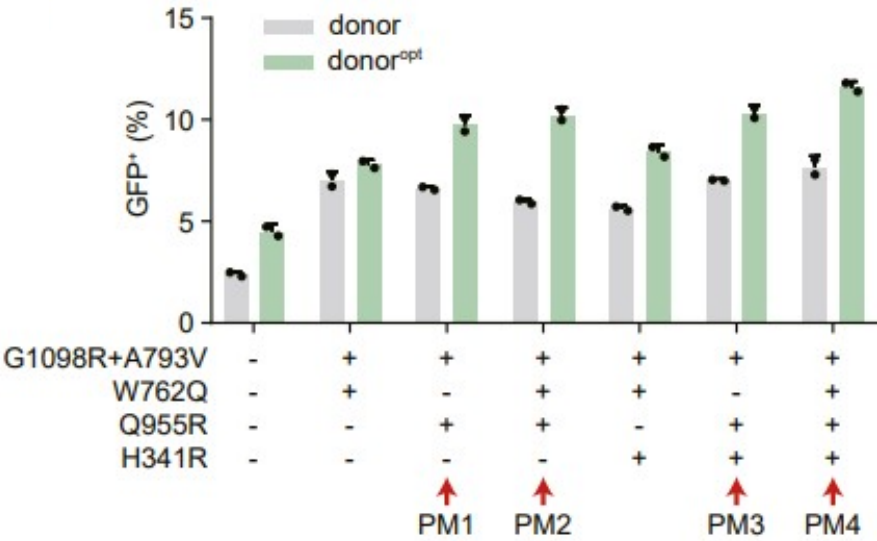
# Domain insertion in R2 protein



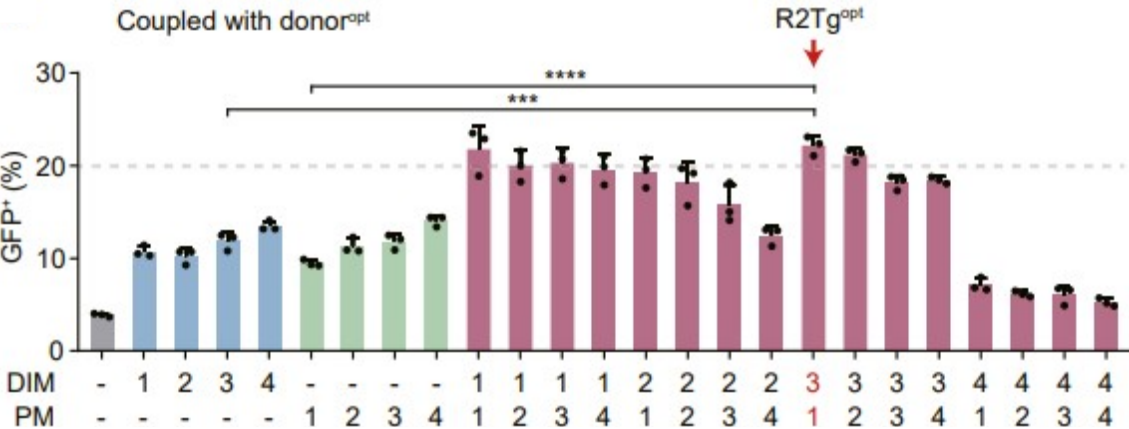
# Domain insertion in R2 protein



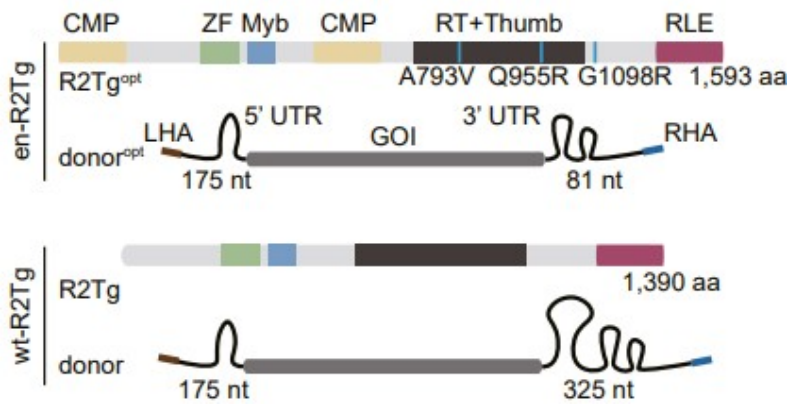
Engineered RNA donor and R2 protein enhances integration synergistically.



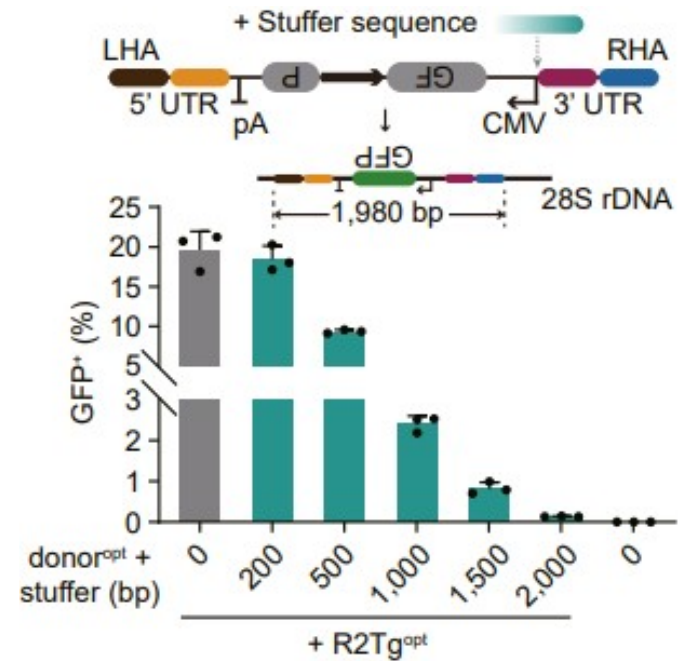
# Combination of strategies generated en-R2Tg



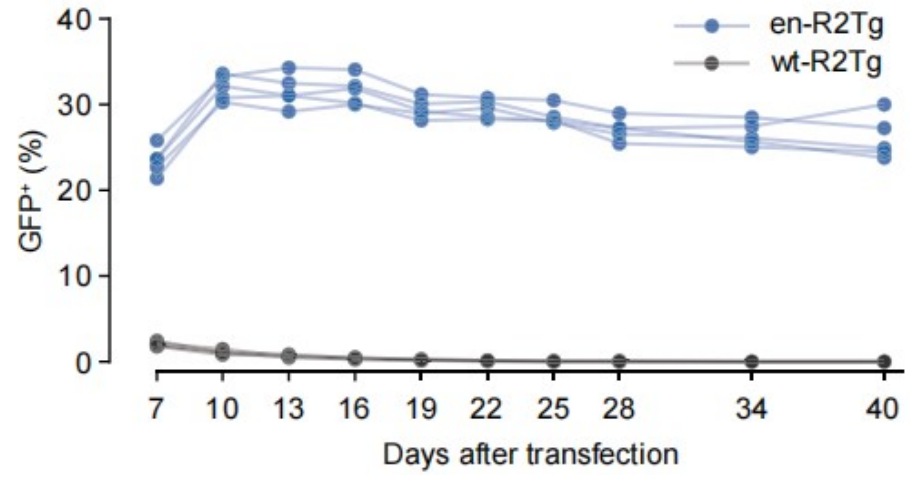
# CMP(N)-CMP(IL-2)-R2Tg(A793V/Q955R/G1098R)



Bigger than 2.5kb insertion capacity.

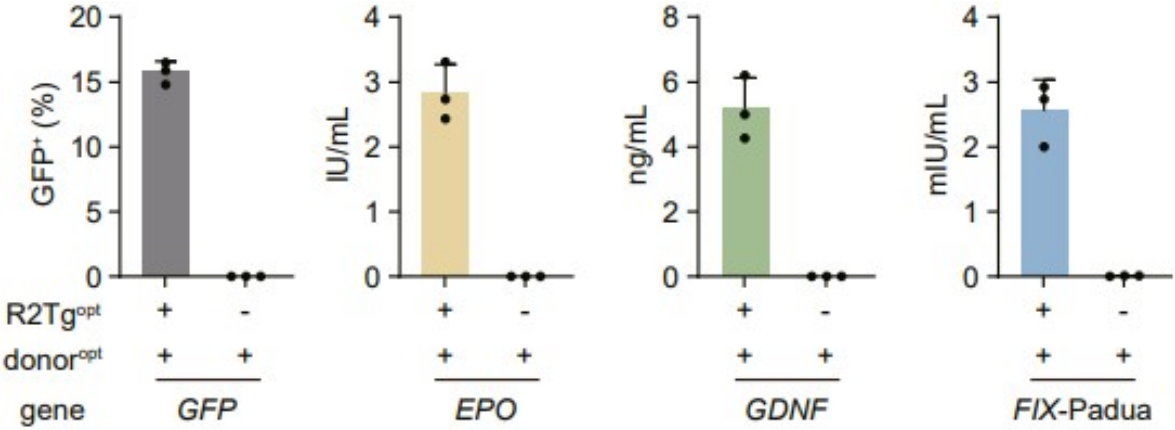


Stable expression over 1 month.



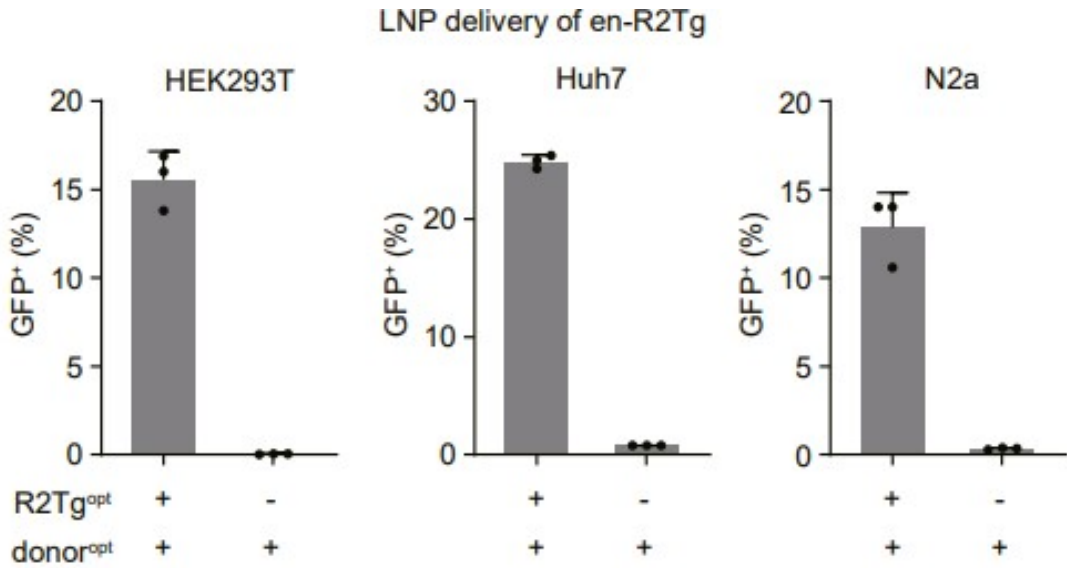
(in HEK293T cells by extended cell culture)

Integrate different genes in HEK293T.

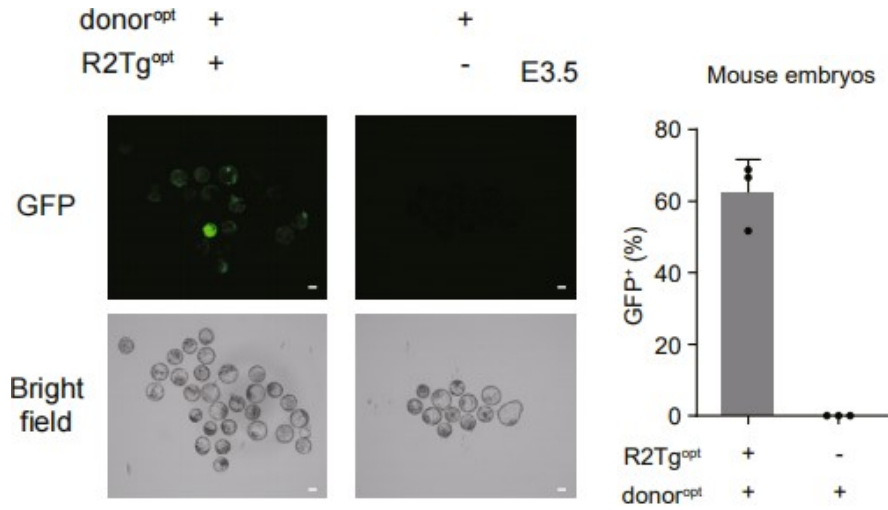
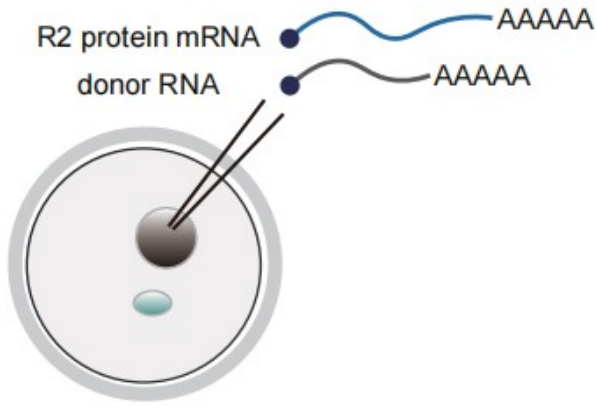


*In-vitro* transcribed RNA, delivered by Lipo3000.

Examples in multiple cell lines and zygote cells.

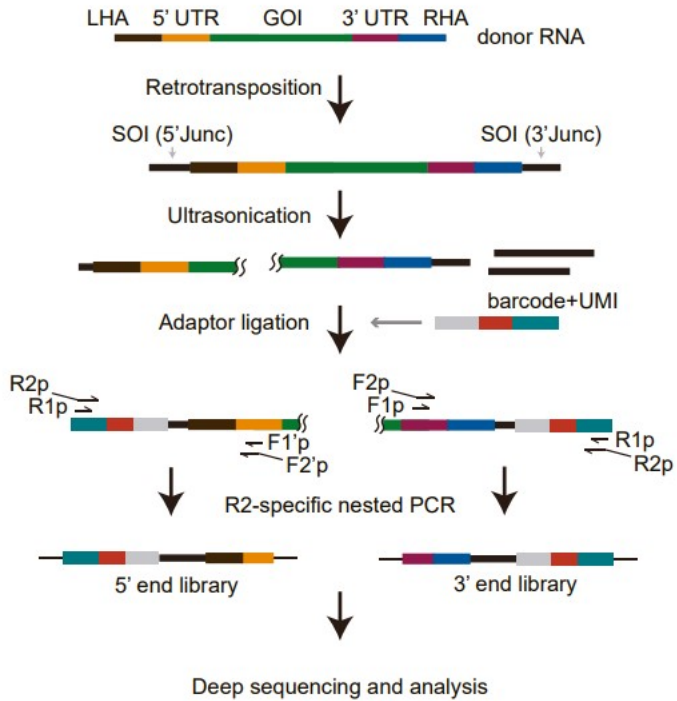


Microinjection of RNAs into mouse zygotes

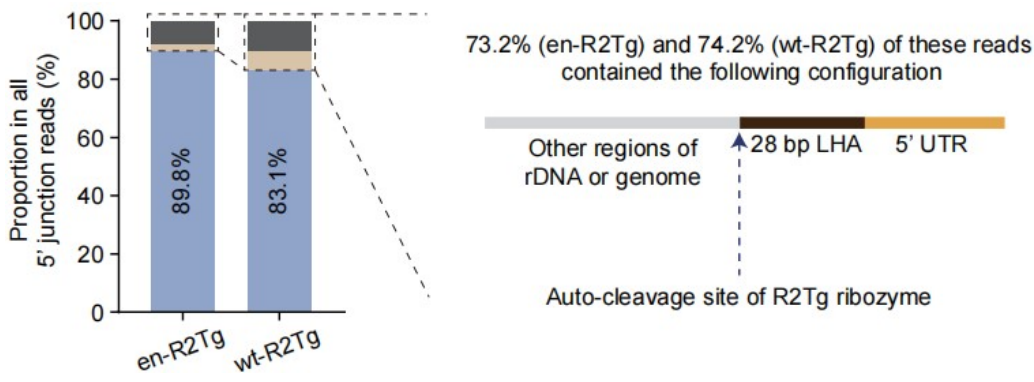




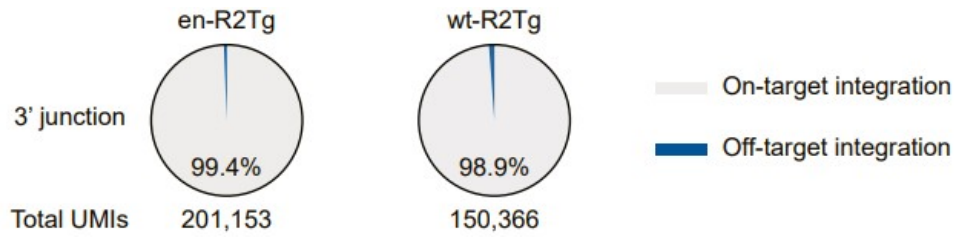
# Precise integration specificity at 28S rDNA.



## 89.8% of 5' junction reads have exact integration sites

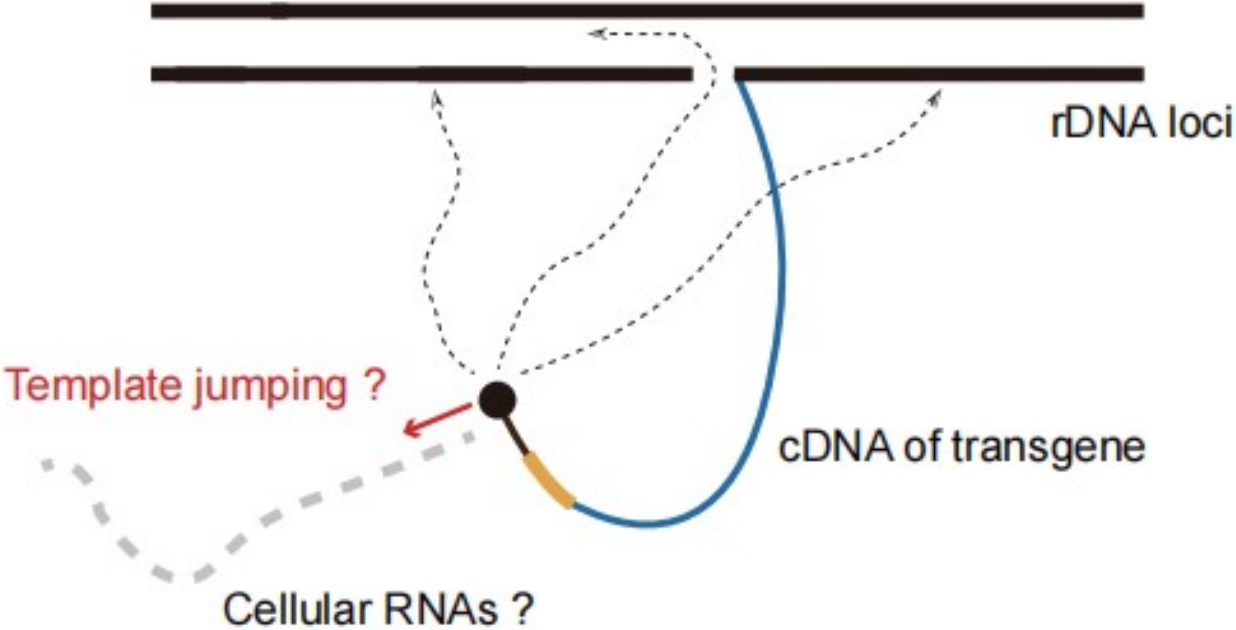


## 99.4% of 3' junction reads have exact integration sites



# Capturing retrotransposon-integrated transgene sequencing (CRIT-seq)

Models explain the differences between 5' and 3' junction precision



## Summary

- Systematic analysis and screening of naturally occurring R2 retrotransposons.
- Engineering of donor RNA and protein enhances R2-mediated gene insertion activity.
- All-RNA-delivered en-R2Tg enables effective gene integration in mammalian cells.
- Optimized en-R2Tg system has high on-target integration specificity in genome.