

Reagents for Molecular Biology Research

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2020 Product Catalogue

Reagents for Molecular Biology Research

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PCR

■ High-Fidelity PCR

| Product Name | Size | Cat. No.# |
|--|-------------------------|---------------|
| Phanta Max Super-Fidelity DNA Polymerase | 100 U / 500 U / 1,000 U | P505-d1/d2/d3 |
| (HOT) 2 Phanta Max Master Mix | 1 ml / 5 ml / 15 ml | P515-01/02/03 |
| New 2 Phanta Max Master Mix (Dye Plus) | 1 ml / 5 ml / 15 ml | P525-01/02/03 |

■ Conventional PCR

| | Product Name | Size | Cat. No.# |
|-----|---|------------------------------|---------------|
| | Taq DNA Polymerase (Mg ² plus Buffer) | 1,000 U / 5,000 U / 10,000 U | P101-01/02/03 |
| | Taq DNA Polymerase (Mg ² free Buffer) | 1,000 U / 5,000 U / 10,000 U | P102-01/02/03 |
| | Taq DNA Polymerase (Mg² plus Buffer, with dNTP) | 1,000 U / 5,000 U / 10,000 U | P101-d1/d2/d3 |
| | Taq DNA Polymerase (Mg ² free Buffer, with dNTP) | 1,000 U / 5,000 U / 10,000 U | P102-d1/d2/d3 |
| | 2 Taq Master Mix | 5 ml / 15 ml / 50 ml | P111-01/02/03 |
| HOT | 2 Taq Master Mix (Dye Plus) | 5 ml / 15 ml / 50 ml | P112-01/02/03 |
| | Green Taq Mix | 5 ml / 15 ml / 50 ml | P131-01/02/03 |

■ High-Yield PCR

| Product Name | Size | Cat. No.# |
|-------------------------------------|---------------------------|---------------|
| Taq Plus DNA Polymerase | 250 U / 1,000 U / 3,000 U | P201-01/02/03 |
| Taq Plus DNA Polymerase (with dNTP) | 250 U / 1,000 U / 3,000 U | P201-d1/d2/d3 |
| 2 Taq Plus Master Mix | 5 ml / 15 ml / 50 ml | P211-01/02/03 |
| 2 Taq Plus Master Mix II (Dye Plus) | 5 ml / 15 ml / 50 ml | P213-01/02/03 |

■ Long-Fragment PCR

| Product Name | | Cat. No.# |
|---|---------------------|---------------|
| Vazyme LAmp DNA Polymerase (Mg ² plus buffer) | 125U/500U | P301-01/02 |
| Vazyme LAmp DNA Polymerase (Mg ² plus buffer, with dNTP) | 125U/500U | P301-d1/d2 |
| Vazyme LAmp DNA Polymerase (Mg² free buffer, with dNTP) | 125U/500U | P302-d1/d2 |
| 2 Vazyme LAmp Master Mix | 1 ml / 5 ml / 15 ml | P311-01/02/03 |
| 2 Vazyme LAmp Master Mix (Dye Plus) | 1 ml / 5 ml / 15 ml | P312-01/02/03 |

■ Direct PCR

| | Product Name | | Cat. No.# |
|-----|-------------------------------------|------------------|-------------|
| HOT | ne Step Mouse Genotyping it | 200 rxn | PD101-01 |
| | ne Step U Probe Mouse Genotyping it | 200 rxn | PD104-01 |
| | Blood Direct PCR it V2 | 50 rxn / 200 rxn | PD103-01/02 |

Rapid PCR

| | Product Name | | Cat. No.# |
|------------------------|-------------------|------------|-----------|
| 2 Rapid Taq Master Mix | 5 ml / 15 ml | P222-01/02 | |
| | 50 ml (50 x 1 ml) | P222-03 | |
| | 50 ml (10 x 5 ml) | P222-04 | |



■ Hot-Start PCR

| Product Name | Size | Cat. No.# |
|------------------------------------|---------------------------|---------------|
| HOT AceTaq DNA Polymerase | 250 U / 1,000 U / 3,000 U | P401-d1/d2/d3 |
| 2 AceTaq Master Mix | 1 ml / 5 ml /15 ml | P411-01/02/03 |
| 2 AceTaq Master Mix (Dye Plus) | 1 ml / 5 ml / 15 ml | P412-01/02/03 |
| Champagne Taq antibody | 500 U | P121-01 |
| (HOT) Champagne Taq DNA Polymerase | 500 U (2.5 / 5 / 10 U/ I) | P122-d1/d2/d3 |

■ Multiplex PCR

| Product Name | Size | Cat. No.# |
|------------------|------------------------------|----------------|
| Multiplex PCR it | 50 rxn / 200 rxn / 1,000 rxn | PM101-01/02/03 |

■ Isothermal Amplification

| Product Name | | Cat. No.# |
|-----------------------------------|------------|------------|
| Bst DNA Polymerase Large Fragment | 00U/ .000U | P 01-01/02 |

PCR-Related

| Product Name | Size | Cat. No.# |
|------------------------|-----------------|------------|
| PCR Enhancer | 500 I | P021-01 |
| dNTP Mix (10 mM each) | 1 ml / 5 ml | P031-01/02 |
| dNTP Mix (2.5 mM each) | 1 ml / 5 ml | P032-01/02 |
| eat-labile UDG | 100 U / 500 U | P051-01/02 |
| E.coli UDG | 500 U / 5,000 U | P061-01/02 |

Cloning/Mutagenesis

Fast Cloning

| | | | Cat. No.# |
|-----|---------------------------------------|-----------------|------------|
| | ClonExpress II ne Step Cloning it | 25 rxn / 50 rxn | C112-01/02 |
| | ClonExpress MultiS ne Step Cloning it | 10 rxn / 25 rxn | C113-01/02 |
| HOT | ClonExpress Ultra ne Step Cloning it | 25 rxn / 50 rxn | C115-01/02 |

Fast Mutagenesis

| Product Name | | |
|---|-----------------|------------|
| Mut Express II Fast Mutagenesis it V2 | 10 rxn / 25 rxn | C214-01/02 |
| Mut Express MultiS Fast Mutagenesis it V2 | 10 rxn / 25 rxn | C215-01/02 |

■ TA Cloning

| Product Name | | |
|-----------------------------|------------------|------------|
| T4 DNA Ligase | 40,000 U | C301-01 |
| Smin Universal Ligation Mix | 50 rxn / 100 rxn | C311-01/02 |

■ TOPO Cloning

| Product Name | | |
|-----------------------------------|-----------------|------------|
| HOT 5min TA/Blunt- ero Cloning it | 25 rxn / 50 rxn | C601-01/02 |



Nucleic Acid Electrophores is

GelRed Nucleic Acid Stain

| | Product Name | | Cat. No.# |
|-----|--|-----------------------|----------------|
| HOT | Ultra GelRed Nucleic Acid Stain (10000) | 0.5 ml / 5 ml / 50 ml | GR501-01/02/03 |

DNA Marker

| Product Name | Size | Cat. No.# |
|------------------------|---------------|-------------|
| DL2000 Plus DNA Marker | 250 I / 500 I | MD101-01/02 |
| DL5000 DNA Marker | 250 I / 500 I | MD102-01/02 |
| DL15000 DNA Marker | 250 1 / 500 | MD103-01/02 |
| 100 bp DNA Ladder | 250 1 / 500 | MD104-01/02 |

ReverseTranscription

■ Conventional RT-PCR

| | Product Name | | Si | ze Cat. No.# |
|-----|---------------------------------------|------------------|-----------------------------|-----------------|
| | iScript III Reverse Transcriptase | | 10,000 | U R302-01 |
| | iScript II 1st Strand cDNA Synthesis | it | 50 rxn / 100 rxn (20 I / rx | kn) R211-01/02 |
| HOT | iScript III 1st Strand cDNA Synthesis | it (gDNA wiper) | 50 rxn / 100 rxn (20 I / rx | (n) R312-01/02 |
| | M-MLV(-) Reverse Transcriptase | | 10,000 | U R021-01 |
| | Murine RNase inhibitor | | 2,000 U / 10,000 U / 20,000 | U R301-01/02/03 |

RT-qPCR SuperMix

| | Product Nar | ne | | Size | Cat. No.# |
|-----|---------------|---|-------------|----------|-----------|
| | iScript II | RT SuperMix for qPCR | 100 rxn (20 | I / rxn) | R222-01 |
| HOT | iScript III F | RT SuperMix for qPCR (gDNA wiper) | 100 rxn (20 | I / rxn) | R323-01 |
| | iScript II | Select RT SuperMix for qPCR | 100 rxn (20 | I / rxn) | R232-01 |
| | iScript II | Select RT SuperMix for qPCR (gDNA wiper) | 100 rxn (20 | I / rxn) | R233-01 |

■ One-Step RT-PCR

| iScript II | ne Step RT-PCR | it | 50 rxn (50 | I / rxn) | P611-01 |
|------------|----------------|---------------|------------|----------|---------|
| iScript II | ne Step RT-PCR | it (Dye Plus) | 50 rxn (50 | I / rxn) | P612-01 |

■ Single Cell Sequence Amplification

| Product Name | | |
|--|---------|---------|
| Single Cell Sequence Speci c Ampli cation it | 200 rxn | P621-01 |

miRNA

■ miRNA Reverse Transcription

| Product Name | | Cat. No.# |
|---|------------------------------|-------------|
| miRNA 1st Strand cDNA Synthesis it (by stem-loop) | 50 rxn / 100 rxn (20 1/ rxn) | MR101-01/02 |

miRNA qPCR

| Product Name | | Cat. No.# |
|--------------------------------------|--------------------------------|-------------|
| miRNA Universal S BR qPCR Master Mix | 125 rxn / 500 rxn (20 1 / rxn) | M 101-01/02 |



qPCR

qPCR Master Mix (SYBR)

| | Produc | | | Cat. No.# |
|-----|--------|--------------------------------|----------------------------------|-----------|
| HOT | Cham | Universal S BR qPCR Master Mix | 500 rxn / 2,500 rxn (20 l / rxn) | 11-02/03 |
| | Ace | Universal S BR qPCR Master Mix | 500 rxn / 2,500 rxn (20 l / rxn) | 511-02/03 |
| | Ace | qPCR S BR Green Master Mix | 500 rxn / 2,500 rxn (20 l / rxn) | 111-02/03 |

qPCR Master Mix (Probe)

| | | | Cat. No.# |
|-----|-------------------------------------|----------------------------------|-----------|
| | Ace qPCR Probe Master Mix | 500 rxn / 2,500 rxn (20 I / rxn) | 112-02/03 |
| HOT | Ace Universal U Probe Master Mix V2 | 500 rxn / 2,500 rxn (20 I / rxn) | 513-02/03 |
| HOT | Cham Geno-SNP Probe Master Mix | 500 rxn / 2,500 rxn (20 I / rxn) | 11-02/03 |

One-Step qRT-PCR Mix

| Product Name | | Cat. No.# |
|--|----------------------|-----------|
| iScript II ne Step qRT-PCR S BR Green it | 250 rxn (20 I / rxn) | 221-01 |
| iScript II ne Step qRT-PCR Probe it | 250 rxn (20 I / rxn) | 222-01 |
| iScript II U ne Step qRT-PCR Probe it | 250 rxn (20 l / rxn) | 223-01 |

GenomeEditing

| Product Name | Size | Cat. No.# |
|---------------------------------|--------------------|-------------|
| Cas Nuclease | 50 pmol / 250 pmol | EN301-01/02 |
| T Endonuclease I | 50 pmol / 250 pmol | EN303-01/02 |
| In Vitro Transcription | | |
| Product Name | | |
| T inh ield RNA Transcription it | 50 rvn / 100 rvn | TR101-01/02 |

25 rxn / 50 rxn

TR102-01/02

NucleicAcidIsolation

■ Rapid Sample Treatment

HOT) T RNAi Transcription it

| | Product Name | | Size | Cat. No.# |
|-----|-----------------------|----|---|---------------|
| New | RoomTemp Sample Lysis | it | 250 rxn (5 ml) / 1000 rxn (20 ml) / 5000 rxn (100 ml) | P0 3-01/02/03 |

■ RNA Isolation (Column)

| Product Name | Size | Cat. No.# |
|--|--------|-----------|
| FastPure Cell / Tissue Total RNA Isolation Mini it | 50 rxn | RC101 |
| FastPure Plant Total RNA Isolation it (Polysaccharides / Polyphenolics-Rich) | 50 rxn | RC401 |



■ DNA Isolation (Column)

| Product Name | Size | Cat. No.# |
|--|------------------|-------------|
| FastPure Blood DNA Isolation Mini it V2 | 50 rxn / 200 rxn | DC111-01/02 |
| FastPure Cell/Tissue DNA Isolation Mini it | 100 rxn | DC102 |
| FastPure Bacteria DNA Isolation Mini it | 100 rxn | DC103 |
| FastPure Plant DNA Isolation Mini it | 50 rxn | DC104 |
| FastPure FFPE DNA Isolation it | 50 rxn | DC105 |
| Lysozyme | 200 mg | DE103 |

■ Tissue Stabilizer

| Product Name | Size | Cat. No.# | |
|-----------------------------|--------|-----------|--|
| RNA eeper Tissue Stabilizer | 100 ml | R501-01 | |

■ Exosome Isolation

| Product Name | Size | Cat. No.# |
|--|-------|-----------|
| VE Exosome Isolation Reagent (from cell culture media) | 50 ml | R601 |
| VE Exosome Isolation Reagent (from serum) | 10 ml | R602 |
| VE Exosome Isolation Reagent (from plasma) | 10 ml | R603 |

CellBiology/ProteinResearch

CellCounting

| | Product Name | | Cat. No.# |
|-----|-----------------------|---------------------|------------|
| HOT | CC - Cell Counting it | 500 rxn / 1,000 rxn | A311-01/02 |

CellTransfection

| Product Name | | |
|----------------------------------|----------------------|---------------|
| ExFect 2000 Transfection Reagent | 0.5 ml / 1 ml / 5 ml | T202-01/02/03 |

DualLuciferaseReporterAssay

| | Product Name | | Cat. No.# | |
|-----|-----------------------------------|---------|-----------|--|
| HOT | Dual Luciferase Reporter Assay it | 100 rxn | DL101-01 | |

Mycoplasma

| | Product Name | | | |
|-----|------------------------------|---------------------|---------------|--|
| HOT | MycoBlue Mycoplasma Detector | 20 rxn / 50 rxn | D101-01/02 | |
| | Myco- ff Mycoplasma Cleaner | 100 1/500 1/1.000 1 | D103-01/02/03 | |



PCR

Selection Guide

| Applications | Products (Cat.#) | Features | Applicable for |
|----------------------|--|---|--|
| Conventional PCR | 2 Taq Master Mix (P111) 2 Taq Master Mix (Dye Plus) (P112) Green Taq Mix (P131) | No 3' 5' exonuclease activity. Excellent compatibility. Products contain A at 3'-end. | Colony PCR Large-scale gene identi cation TA Cloning for small fragments. |
| igh- ield PCR | Taq Plus Master Mix (P211) Taq Plus Master Mix II (Dye Plus) (P213) | With delity 6-fold higher than Taq. Mixed products with 3'-end blunt or containing A. | PCR that requires some delity. |
| Rapid PCR | 2 Rapid Taq Master Mix (P222) | Ampli cation speed up to 15 sec / kb. | Colony PCR. |
| Long-Fragment PCR | Vazyme LAmp Master Mix (P311) Vazyme LAmp Master Mix (Dye Plus) (P312) | Ef ciently amplify fragments 20 kb. | Long-fragment ampli cation. |
| ot-Start PCR | AceTaq Master Mix (P411) AceTaq Master Mix (Dye Plus) (P412) Champagne Taq Antibody (P121) Champagne Taq DNA Polymerase (P122) | Excellent speci city. Excellent sensitivity. | Ampli cation that requires higher sensitivity and speci city Ampli cation of genes with low copy or qPCR assay from complex templates (genomic DNA, cDNA). |
| Multiplex PCR | Multiplex PCR it (PM101) | 1 -plex PCR in one single reaction. | Detection or typing of pathogens. |
| Direct PCR | ne Step Mouse Genotyping it (PD101) Blood Direct PCR it V2 (PD103) | Easy and fast, without DNA puri cation. | ne step mouse genotyping Direct PCR from plant tissues Direct PCR from blood. |
| igh-Fidelity PCR | Phanta Max Super-Fidelity DNA Polymerase (P505) 2 Phanta Max Master Mix (P515) 2 Phanta Max Master Mix (Dye Plus) (P525) | With super delity 53-fold higher than Taq igh resistance to PCR inhibitors. | igh- delity PCR. Ampli cation of templates with high GC-content Long-fragment (up to 40 kb) ampli cation. |



High-Fidelity PCR



→ 2x Phanta Max Master Mix (#P515)

→ 2x Phanta Max Master Mix (Dye Plus) (#P525)



Super Fidelity 53-fold higher than Taq DNA Polymerase.

Long Fragment amplify fragments up to 40 kb.

Suitable for templates with high GC-content.

Suitable for Direct-PCR using crude materials as templates .

Validated crude materials bacteria, fungi, whole blood, cultured cells, plant or animal tissue lysate, food lysates, etc.



Selected Product Citations

hao , et al. Metabolic coupling of two small-molecule thiols programs the biosynthesis of lincomycin A. *Nature*, 2015, 51 (53)115- .

Tian , et al. An enzymatic 4 2 cyclization cascade creates the pentacyclic core of pyrroindomycins. *Nature Chemical Biology*, 2015, 11(4) 25 -65.

an , et al. Mapping the Mouse Cell Atlas by Microwell-Seq. Cell, 201 , 1 2(5) 10 1-10 .

Cheng , et al. Pacer Mediates the Function of Class III PI3 and PS Complexes in Autophagosome Maturation by Engaging Stx1 . *Molecular Cell*, 201 , 65(6) 102 -43.

Lv M, et al. Characterization of a C3 Deoxygenation Pathway Reveals a ey Branch Point in Aminoglycoside Biosynthesis. Journal of the American Chemical Society, 2016, 13 (20) 642 -35.



High-Yield PCR



→ 2x Taq Plus Master Mix II (Dye Plus) (#P213)

Features

- * Robust performance for high-yield PCR in most primer-template systems.
- * Ready-to-use master mix with no need for operations on ice.

 PCR products can be directly loaded for electrophoresis with no need for loading buffer.

Validation Data







2x Taq Plus Master Mix II (Vazyme, #P213) demonstrated excellent template compatibility. Fragments (0.5 kb to 15 kb) were amplified from genomic DNA (mouse, human, wheat, rice), bacterial culture, and λ DNA, respectively. A specific corresponding band was observed in each PCR.

₩,

Selected Product Citations

hang , et al. (2014) Complementary sequence-mediated exon circularization. Cell, 15 (1) 134-4 .

uan , et al. Gyrl-like proteins catalyze cyclopropanoid hydrolysis to confer cellular protection. *Nature Communications*, 201 , (1).14 5.



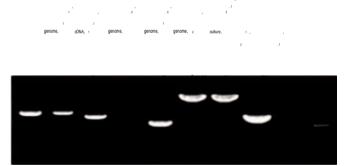
Rapid PCR



Features

- * Rapid ampli cation speed is 15 sec / kb, with an extreme speed of 1 sec / kb for fragments within 1 kb.
- * Ready-to-use master mix with no need for operations on ice. PCR products can be directly loaded for electrophoresis with no need for loading buffer.
- * Excellent stability: remains stable after 50 freeze-thaw cycles.

Validation Data



Fragments (1 kb - 2 kb) was amplified from genomic DNA (human, mouse, wheat, rice), cDNA (human), bacterial culture, colony, and λ DNA, respectively. The extension time was set as 1 sec / kb. Ten μl of PCR product was loaded for agarose gel electrophoresis. Specific bands were observed.

Selected Product Citations

hang B, et al. Enzyme-catalysed 6 4 cycloadditions in the biosynthesis of natural products. Nature, 201, 56 (50) 122-6. Wang S, et al. Molecular Basis for the Final xidative RearrangementSteps in Chartreusin Biosynthesis. J Am Chem Soc, 201 , 140(34) 10 0 -14.



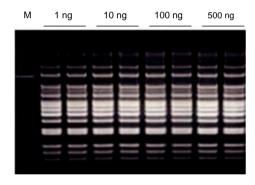
Multiplex PCR



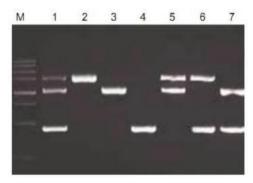
Features

- Multiplex 19-plex PCR or even higher.
 Excellent target-to-target ampli cation uniformity and extremely low target preference.
- * Highly sensitive ampli cation from trace amount of genomic DNA (1 ng).

Validation Data



Uniformamplificationcoverageofdifferentregions.Human genomic DNA was used as template for 19-plex PCR. The size of the amplicons ranged from 70 bp to 916 bp. The result indicated that Multiplex PCR Kit (Vazyme, #PM101) has a uniform amplification coverage of different regions for 1 ng-500 ng of template.



The Multiplex PCR Kit showed excellent compatibility with fragment length. Mouse genomic DNA was used as template for amplification of 1.55 kb, 1.07 kb, and 0.45 kb fragments, respectively. The result indicated that Multiplex PCR Kit (Vazyme, #PM101) is compatible with amplicons of various lengths in one single reaction system.

1: 3-plex PCR 2-4: 1-plex PCR 5-7: 2-plex PCR M: DL5000 DNA Marker



Cloning / Mutagenesis

Selection Guide

| Applications | Products (Cat.#) | | Applicable for |
|------------------|--|---|--|
| Fast Cloning | ClonExpress Ultra ne Step Cloning it (C115) ClonExpress II ne Step Cloning it (C112) ClonExpress MultiS ne Step Cloning it (C113) | Easy, fast, and ef cient. No need to consider the restriction enzyme cutting sites on the inserts. Ligase-independent. Positive Clone Rate 5. Ef cient cloning of fragments of 50 bp - 10 kb. | Cloning or assembly of 1-5 fragments. |
| Fast Mutagenesis | Mut Express II Fast Mutagenesis it V2 (C214) Mut Express MultiS Fast Mutagenesis it V2 (C215) | Ef cient ampli cation of any plasmids within 20 kb. Site-directed mutations of 1-5 discontinuous sites in one reaction. | 1-5 separate site-directed mutagenesis on one plasmid. |
| T P Cloning | 5min TA/Blunt- ero Cloning it (C601) | Cloning within 5 min. Positive Clone Rate 5 | TA cloning. cloning with blunt ends. |

TOPO Cloning



→ 5min TA/Blunt- ero Cloning it (#C601)

Features

Ready-to-use master mix.

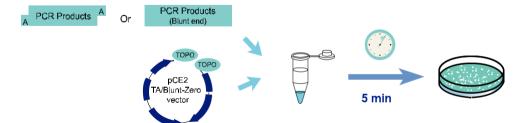
Suitable for both TA cloning and blunt-end cloning.

Rapid cloning within 5 min.

igh cloning ef ciency with Positive Clone Rate

Ampicillin and ana dual resistance vector.

ork ow



5.



Fast Cloning



→ ClonExpress

Itra One Step Cloning

it (#C115)

Features

Cloning within 5 min.

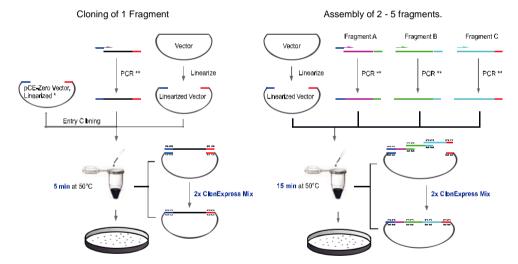
Ready-to-use super mix in one tube.

Ef cient cloning of fragments of 50 bp - 10 kb with Positive Clone Rate 5.

Suitable for cloning of 1 fragment, assembly of 2 - 5 fragments, and entry cloning.

Independent of DNA ligase, signi cantly reducing the self-ligated colonies.

Mechanism



^{*} pCE- ero Vector, Linearized, is supplied with ClonExpress Ultra ne Step Cloning Kit (Vazyme, #C115).

^{**} It is highly recommended to use Vazyme s APP - CE Design - for easy primer design.



Selected Product Citations of ClonExpress

Wu N, et al. TB 6 null variants and a common hypomorphic allele in congenital scoliosis. New England Journal of Medicine, 2015, 3 2(4) 341-50.

Ge , et al. Architecture of the mammalian mechanosensitive Piezo1 channel. Nature, 2015, 52 (5 6) 64- .



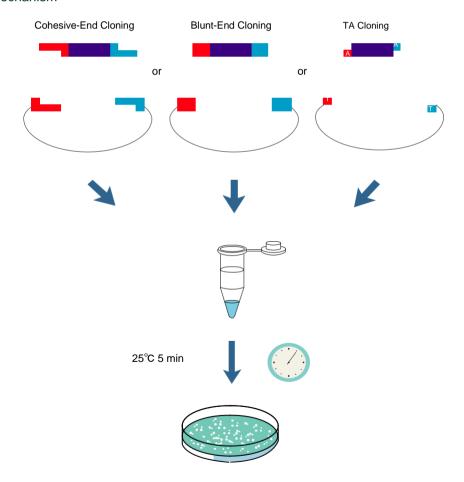


→ 5min niversal Ligation Mix (#C311)

Features

- * Versatile Suitable for TA cloning, blunt-end cloning, cohesive-end cloning, and ligation of linkers or adapters. Fast Cloning within 5 min at 25°C.
- * Ef cient Positive Clone Rate

Mechanism



GelRed

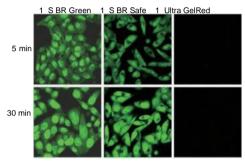




Itra GelRed Nucleic Acid Stain (10000x) (#GR501)

Perfect substitute for ethidium bromide (EB)

No toxicity



Ultra GelRed is unable to cross cell membranes.

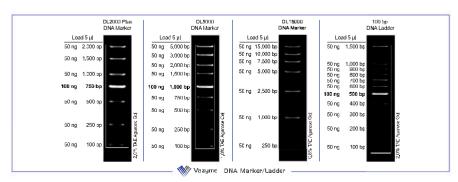
DNA Marker/Ladder



→ DNA Markers / Ladders

Stable

Clear Bands





Reverse Transcription

| Selection | Guide | | | | | | | | |
|-----------------------------------|----------------------|--|-------------|----------------------------|-----------------|--|---------------------|--|--------------|
| | nestate in the state | Harder II to State of the State | HEEREN ICHT | PERSONAL MERCHANIST STREET | HERE'N CERECHIE | Here and the state of the state | Hacide III de Steen | A THE STATE OF THE | or all color |
| Applications | | | | | | | | | ĺ |
| RT-qPCR | | | • | - | - | | | | |
| RT-PCR | | | | | | | | | |
| Features | | | | | | | | | |
| SuperMix | | | | | | | | | |
| Long-fragment cDNA | | | | | | | | | |
| Rapid removal of Genomic DNA | | | | | | | - | | |
| Primers | | | | | | | | | 1 |
| ligo dT ₂₃ VN / N6 Mix | | | | | | | | | |
| ptional | | | | | | | | | |

| | M-MLV (H-) (#R021) | HiScript II Reverse Transcriptase (#R201) | HiScript III Reverse Transcriptase (#R302) |
|-----------------------|-----------------------|---|--|
| Reaction temperature | 3 °C-42°C | 42°C - 55°C | 3 °C-50°C |
| Thermal stability | *** | **** | *** |
| RNase activity | No | No | No |
| cDNA length | 2 kb-3 kb | Up to 20 kb | Up to 20 kb |
| Template adaptability | *** | *** | **** |



RT-qPCR SuperMix

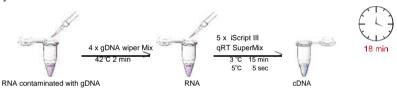


Features

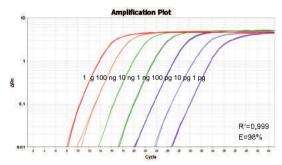
- * Ready-to-use SuperMix reverse transcription within 20 min by only adding template RNA.
- * Excellent ef ciency for low-input RNA or degraded RNA.
- * Excellent tolerance for impurities (i.e. ethanol, isopropanol, phenol water, guanidine thiocyanate, humic acid). Lower CT value and higher ef ciency than most other commercially available reverse transcription reagents.

Validation Data

1. Easy Fast



2. Excellent Sensitivity



RNA of HeLa cells was serially diluted and reverse transcribed using HiScript III RT SuperMix for qPCR (gDNA wiper) (Vazyme, #R323), followed by qPCR detection of gene ACT. The results show an excellent linear relationship across a wide range of RNA concentrations. The target gene (ACT) was detected in 1 pg of RNA.



qPCR

Selection Guide

| Applications | Products (Cat.#) |
|------------------------|--|
| S BR | Cham Universal S BR PCR Master Mix (11) |
| Probe | Ace Universal U Probe Master Mix V2 (513) |
| SNP (TaqMan MGB Probe) | Cham Geno-SNP Probe Master Mix (11) |

qPCR Master Mix (SYBR)



niversal SYBR qPCR Master Mix (# 711)

Best Combination of Specificity Sensitivity

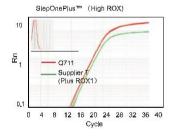
ot-Start Tag Unique specificity-promoting Factors

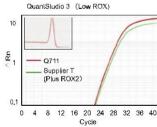
ptimal Concentrations of Mg² and Dye
 Universal

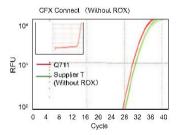
Validation Data

Unique

Applicable for almost all qPCR instruments.







Selected Product Citations

u L, et al. The transcription factor TCF-1 initiates the differentiation of TF *Immunology*, 2015, 4 (3) 53 -51.

cells during acute viral infection. Nature

Guo C, et al. Cholesterol omeostatic Regulator SCAP-SREBP2 Integrates NLRP3 Inflammasome Activation and Cholesterol Biosynthetic Signaling in Macrophages. *Immunity*, 201, 4 (5) 42-56.



qPCR Master Mix (Probe)



→ Ace

niversal

Probe Master Mix V2 (# 513)

Features

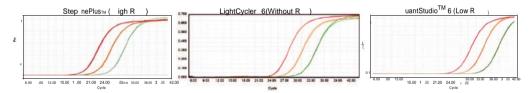
* Excellent sensitivity: ot-start AceTaq and optimal buffer ensure high sensitivity and effectively inhibit nonspeci c ampli cation.

Excellent linear relationship over a large range of input amount of template. Suitable for the detection of single-copy templates.

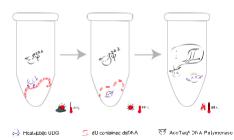
- * Anti-contamination: the dUTP/UDG system eliminates possible contaminations and ensures reliable results.
- * niversal: applicable for almost all qPCR instruments.

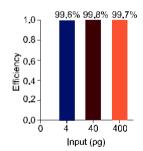
Validation Data

1. Applicable for almost all qPCR instruments.



2. d TP/ DG system.





For Vazyme # 513, the removal rate of the contaminated template is as high as 99.6, effectively ensuring the accuracy of experimental results. U-containing templates (4 pg, 40 pg, 400 pg) were added respectively to the reaction system to evaluate the removal efficiency of the contaminated template by Vazyme # 513.



qPCR Master Mix (Probe)





Geno-SNP Probe Master Mix (# 811)

Advantages

Compatible with 1 ng - 10 ng of input genomic DNA.

Accurate genotyping of SNP sites with GC-content of 25 - 3.

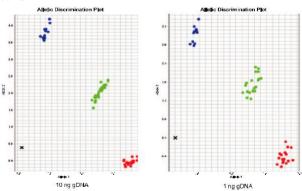
Excellent stability stable signal and accurate genotyping results can be obtained both 2 hr pre-PCR and 2 hr post-PCR.

72 hr pre-PCR PCR reaction solutions were prepared and left in darkness (at room temperature) for 2 hr before PCR 72 hr post-PCR after PCR, the samples were left in darkness (at room temperature) for 2 hr.

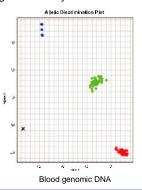
* Blood lysate can be directly used as a template for SNP genotyping, with no need for blood genomic DNA extraction.

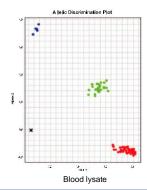
Validation Data

1. Flexible input amounts.



2. Direct genotyping with blood lysate.







Nucleic Acid Isolation

Selection Guide

| Category | Series | Sample / Application | Products | Cat.# |
|-------------------------------|---|---|--|-------|
| DNA Isolation | Rapid Sample Treatment | Blood | RoomTemp Sample Lysis it | P0 3 |
| | Extraction DNA (Column) | Blood | FastPure Blood DNA Isolation Mini it V2 | DC111 |
| | | Cell/tissue | FastPure Cell/Tissue DNA Isolation Mini it | DC102 |
| | | FastPure Bacteria DNA Isolation Mini it | | DC103 |
| Purification | | Plant | FastPure Plant DNA Isolation Mini it | DC104 |
| | | FFPE | FastPure FFPE DNA Isolation it | DC105 |
| | | Lysozyme | Lysozyme | DE103 |
| | RNA tissue eeper RNA eeper for fresh tissue | | RNA eeper Tissue Stabilizer | |
| RNA Isolation Purification | Column RNA Extraction | Cell/tissue total RNA | FastPure Cell/Tissue Total RNA Isolation Mini it | RC101 |
| | | Polysaccharide & Polyphenol-rich Plant total RNA | FastPure Plant Total RNA Isolation it (Polysaccharides & Polyphenolics-rich) | RC401 |
| Exosome Isolation | Cell supernatant | | VE Exosome Isolation Reagent (from cell culture media) | |
| | Serum | | VE Exosome Isolation Reagent (from serum) | R602 |
| | Plasma | | VE Exosome Isolation Reagent (from plasma) | |

Plant RNA and DNA Isolation





FastPure Plant Total RNA Isolation it (Polysaccharides Polyphenolics-rich) (#RC401)

Features

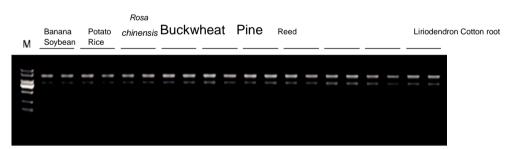
High purity.

Rapid extraction of total RNA from plant tissues, especially from those rich in polysaccharide & polyphenol. Low genomic DNA residue.

Validated Samples

Pine needles, *Eriobotrya aponica* leaves, potato tubers, grape fruits, apples, pears, tobacco leaves, mature leaves and roots of wheat, peach fruit, lotus, chrysanthemum rhizome, bananas, *Rosa chinensis*, buckwheat leaves and seeds, poplar, *Catharanthus roseus* leaves, liriodendron, reed, rice plant, roots and leaves of cotton, strawberry leaf, *Phoebe neurantha* leaves, ginkgo (root, leaf, ower and fruits), Arabidopsis seeds, corn seeds, fungal hyphae, etc.

Validation Data



Total RNA was extracted using Vazyme #RC401 from 50 mg of banana fruit, potato tubers, rose petals, pine needles, reed leaves, Liriodendron leaves, cotton roots, soybean leaves, rice leaves, or 20 mg of buckwheat seed, respectively. The RNA products were loaded for agarose gel electrophoresis. Vazyme #RC401 showed great compatibility to above plants, especially to those that were rich in polysaccharide polyphenol, and the RNA extracted using Vazyme #RC401 was with good integrity and high yield.

M: DL2000 Plus DNA Marker (Vazyme, #MD101). The elution volume was 100 μl and the loading amount was 4 μl-10 μl for agarose gel electrophoresis.



Rapid Sample Treatment



Features

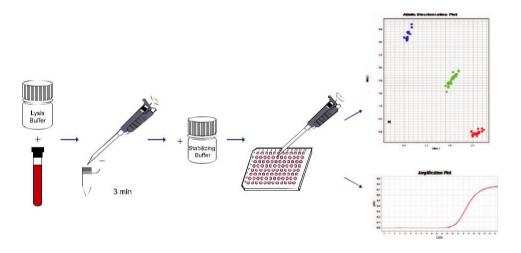
Replace cumbersome template extraction protocols with a simple one-step cell lysis procedure.

Lyse samples in just $3 \, \text{min}$ at room temperature. qPCR reagent is compatible with the fast program, total operation time is less than 1 hr.

Lyse different anticoagulant blood, FTA card, buccal swab and other samples.

The lysis reagent is consistent with the traditional kit for extracting the genome.

ork ow



Work ow of RoomTemp Sample Lysis



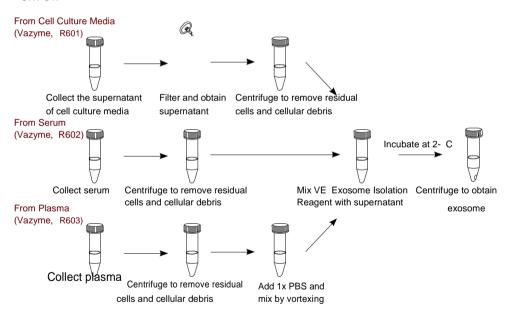
Exosome Isolation



Features

Easy isolation of exosomes by one-step precipitation, avoiding time-consuming ultra-centrifugation. Intact exosomes with high yield obtained by low-speed centrifugation.

ork ow



Cell Counting



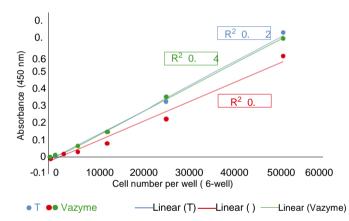
Features

Ready-to-use solution.

igh sensitivity, with excellent linear correlation and repeatability.

Low cytotoxicity.

Validation Data



HEK293 suspension cells were serially diluted and inoculated to a 96-well plate. The cell density in each group (n 3) is: 0, 400, 00, 1600, 3200, 6400, 12 00, 25600, 51200 cells per well. CCK-reagents from Vazyme (#A311, green), Supplier T (blue), and Supplier (red) were used for cell counting, respectively. The R value of Vazyme #A311 is 0.99.

Selected Product Citations

heng, et al. Thiopeptide antibiotics exhibit a dual mode of action against intracellular pathogens by affecting both host and microbe. *Chemistry & Biology*, 2015, 22() 1002-.

Liu , et al. Adiponectin reduces ER stress-induced apoptosis through PPAR transcriptional regulation of ATF2 in mouse adipose. *Cell Death & Disease*, 2016, (11) e24.

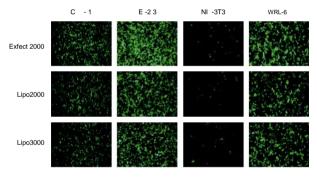
Cell Transfection



Features

- * High transfection of ciency in a variety of cell lines.
- * Low cytotoxicity to avoid damaging the normal physiological state of cells.
- * Add directly to cells in culture medium, in the presence or absence of serum.
- * Applicable for co-transfection with multiple plasmids.

Validation Data



ExFect 2000 exhibits higher transfection efficiency than that of Lipo2000 and Lipo3000. CH K1, HEK293, NIH3T3 and RL6 cells were transfected using various reagents in a 24-well plate, respectively. The expression of FP was analyzed after transfection for 24 h.

Selected Product Citations

Sun, et al. Usp regulates ippo pathway through deubiquitinating the transcriptional coactivator orkie. *Nature Communications*, 201, 10(1) 411.

Liu , et al. 1-L-MT, an ID inhibitor, prevented colitis-associated cancer by inducing CDC20 inhibition-mediated mitotic death of colon cancer cells. *International Journal of Cancer*, 201 , 143(6) 1516-2 .

Luciferase Assay



→ Dual Luciferase Reporter Assay it (#DL101)

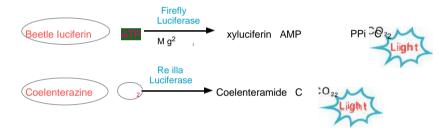
Features

Robust luminescent signals applicable for analysis of weak promoters and other genetic regulatory elements.

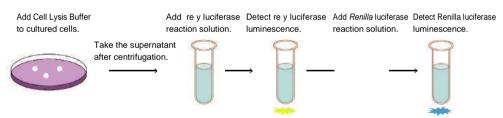
Detection linear range covers up to orders of magnitude (R₂ 0.).

Detection sensitivity of 10-1 mole.

Mechanism



ork ow





Selected Product Citations

Liu, et al. Circular RNA hsa circ 001 3 regulates breast cancer progression via sponging miR-200c-3p. Cell Death & Disease, 201, 10 55

Wu, et al. Ubiquitination is essential for avibirnavirus replication by supporting VP1 Polymerase activity. Journal of Virology, 201, 3(3) e01-1.

Wu, et al. SUM 1 Modification Facilitates Avibirnavirus Replication by Stabilizing Polymerase VP1. Journal of Virology, 201, VI. 0222 -1.

Mycoplasma Detection



→ Myco-Blue Mycoplasma Detector (#D101)

Features

Cell culture supernatant can be used directly for detection.

Results are obtained after incubation at 60°C for 1 hr and can be determined by visual observation.

Accuracy is higher than PCR method, and comparable to gPCR method.

Suitable for detection of all kinds of mycoplasma that are commonly found in cell culture.

Validated Cell Lines

Validated cells and media serum include (but are not limited to)

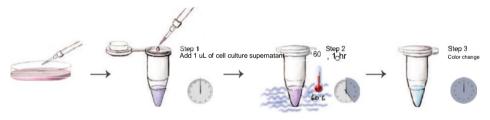
Suspension cells C , NS0, 2 3F, mouse hybridoma, Sf, B 21, etc.

Adherent cells Vero, MDC, SP2/0, 23T, epG2, eLa, A54, MB-MDA231, L 2, MEF, etc.

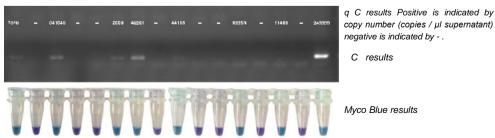
Medium CD FortiC , CDM4, Expi 2 3 Medium, CD ybridoma, Grace, DMEM, 1640, F12, etc.

Serum fetal calf / calf serum horse serum Gibco SR serum replacement, etc.

ork ow



Validated Data



Randomly selected 16 cell cultures, and mycoplasma were detected by three methods.



In Vitro Transcription



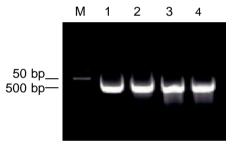
T7 RNAi Transcription it (#TR102)

Features

- * High yield yields up to 80 g of dsRNA in a single reaction.
- * Magnetic bead puri cation: recovery ef ciency up to 0.
- * Able to transcribe both siRNA (21 bp) and dsRNA (long fragment).

Validation Data

1. Excellent transcription ef ciency.



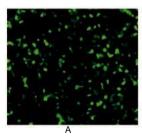
Agarose gel electrophoresis (2) of 500 bp dsRNA.

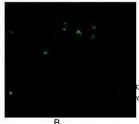
M: DL2000 Plus DNA Maker.

and : products before and after enzymatic hydrolysis of dsRNA, respectively

and : products before and after enzymatic hydrolysis of dsRNA, respectively.

2. nock-down of GFP expression by transcribed siRNA.





3T cells were co-transfected for 24 hrs with both FP plasmid ontrol FP siRNA (A) or positive FP siRNA ().



High-Fidelity PCR

hao , Wang M, u D, et al. Metabolic coupling of two small-molecule thiols programs the biosynthesis of lincomycin A. Nature. 2015 Feb5 51 (53) 115-. IF: 42.351

hang B, Wang B, Wang W, et al. Enzyme-catalysed 6 4 cycloadditions in the biosynthesis of natural products . *Nature.* 201 Apr 56 (50) 122-126. IF: 41.577

Ma , hu L, Song T, et al. A paralogous decoy protects Phytophthora sojae apoplastic effector Ps EG1 from a host inhibitor . *Science*. 201 Feb 1 355(6326) 10-14. IF: 34.661

an , Wang R, hou , et al. Mapping the MouseCell Atlas by Microwell-Seq . Cell. 201 May1 1 3(5) 130 . IF: 31.398

hang B, Li , ang , et al. Crystal Structures of Membrane Transporter MmpL3, an Anti-TB Drug Target . Cell. 201 an 24 1 6(3) 636-64 .e13. IF: 31.398

Wang S, hang B, hu , et al. Molecular Basis for the Final xidative Rearrangement Steps in Chartreusin Biosynthesis . Am Chem Soc. 201 Aug 2 140(34) 10 0 -10 14. IF: 14.357

Cheng , Ma , Ding , et al. Pacer Mediates the Function of Class III Pl3 and PS Complexes in Autophagosome Maturation by Engaging Stx1 . *Mol Cell.* 201 Mar 16 65(6) 102 -1043.e5. IF: 14.248

Wu , in F, Luo , et al. Unusual Processing Generates SPA LncRNAs that Sequester Multiple RNA Binding Proteins . Mol Cell. 2016 Nov 3 64(3) 534-54 . IF: 13.958

Tian , Sun P, an , et al. An enzymatic 4 2 cyclization cascade creates the pentacyclic core of pyrroindomycins . Nat Chem iol. 2015 Apr 11(4) 25 -65. IF: 13.217

hang M, hou C, Wei , et al. uman cleaving embryos enable robust homozygotic nucleotide substitutions by base editors . enome iol. 201 May 22 20(1) 101. IF: 13.214

Wang M, hao , hang , et al. Differences in PLP-Dependent Cysteinyl Processing Lead to Diverse S-Functionalization of Lincosamide Antibiotics . Am Chem Soc. 2016 May 25 13 (20) 634 -51. IF: 13.038

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Sun , Ding , han M, et al. Usp regulates ippo pathway through deubiquitinating the transcriptional coactivator orkie . Nature Communications. 201 an 24 10(1) 411. IF: 12.353

Duan GF, e , u S, et al. Signal peptide represses Glu 1 surface and synaptic traf cking through binding to amino-terminal domain . *Nature Communications*. 201 Nov 1 (1) 4 . IF: 12.353

Chen T, iang F, hu S, et al. ADAR1 is required for differentiation and neural induction by regulating microRNA processing in a catalytically independent manner. . Cell Res. 2015 Apr 25(4) 45 - 6. IF: 12.413

Ding W, i W, Wu , et al. Biosynthesis of the nosiheptide indole side ring centers on a cryptic carrier protein Nos . *Nature Communications*. 201 Sep 5 (1) 43 . IF: 12.124

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i , Li , ie L, et al. Expanding Radical SAM Chemistry by Using Radical Addition Reactions and SAM Analogues . Angew Chem Int Ed Engl. 2016 Sep 1 55(3) 11 45-. IF: 11.71

i , Li , Ding W, et al. Substrate-Tuned Catalysis of the Radical S-Adenosyl-L-Methionine Enzyme NosL Involved in Nosiheptide Biosynthesis . Angew Chem Int Ed Engl. 2015 ul 2 54(31) 021-4. IF: 11.261

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Autophagy. 201 ul 15() 1234-125 . IF: 11.1

u D, Shan B, Sun , et al. USP14 regulates autophagy by suppressing 63 ubiquitination of Beclin1 . enes Dev. 2016 Aug 1 30(15) 1 1 -30. IF: 10.042



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hang B, Wang B, Wang W, et al. Enzyme-catalysed 6 4 cycloadditions in the biosynthesis of natural products . Nature. 201 Apr 56 (50) 122-126. IF: 41.577

hang , Wang B, hang , et al. Complementary sequence-mediated exon circularization . *Cell.* 2014 Sep 25 15 (1) 134-14 . IF: 33.116

Wang S, hang B, hu, et al. Molecular Basis for the Final xidative Rearrangement Steps in Chartreusin Biosynthesis. Am Chem Soc. 201 Aug 2 140(34) 10 0 -10 14. IF: 14.357

Sun , Liu , heng , et al. Distinct chemokine signaling regulates integrin ligand speci city to dictate tissue-speci c lymphocyte homing . Dev Cell. 2014 ul 14 30(1) 61- 0. IF: 12.86

uan , hang , Cai , et al. Gyrl-like proteins catalyze cyclopropanoid hydrolysis to confer Cellular protection . *Nature Communications*. 201 Nov 14 (1) 14 5. IF: 12.124

hang , Wang T T, u L, et al. Genome Mining and Comparative Biosynthesis of Meroterpenoids from Two Phylogenetically Distinct Fungi . *Angew Chem Int Ed Engl.* 201 ul 2 5 (2) 1 4-1 . IF: 12.102

Chen C, hai S, hang L, et al. Uhrf1 regulates germinal center B Cell expansion and af nity maturation to control viral infection . Exp Med. 201 May 215(5) 143-144 . IF: 11.991

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Li , Wang , et al. Base editing with a Cpf1-cytidine deaminase fusion . *Nat iotechnol.* 201 Apr 36(4) 324-32 . IF: 41.667 in S, ong , et al. Cytosine, but not adenine, base editors induce genome-wide off-target mutations in rice . *Science.* 201 Apr 1 364(643)2 2-2 5. IF: 41.058

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ong , Song , Li C, et al. Ef cient C-to-T base editing in plants using a fusion of nCas and human AP BEC3A . *Nat iotechnol.* 201 ct.1. IF: 35.724

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Li , Li , ang S, et al. CRISPR-Cas -mediated base-editing screening in mice identi es DND1 amino acids that are critical for primordial germ Cell development . Nat Cell iol. 201 Nov 20(11) 1315-1325. IF: 19.064

Wang L, ue W, an L, et al. Enhanced base editing by co-expression of free uracil DNA glycosylase inhibitor . Cell Res. 201 ct 2 (10) 12 -12 2. IF: 15.606

hao , Wu , Geng , et al. Ion Permeation and Mechanotransduction Mechanisms of Mechanosensitive Piezo Channels . *Neuron.* 2016 Mar 16 (6) 124 -1263. IF: 15.054

Mo F, huang , Liu , et al. Acetylation of Aurora B by TIP60 ensures accurate chromosomal segregation . Nat Chem iol. 2016 Apr 12(4) 226-32. IF: 14.273

- Cheng , Ma , Ding , et al. Pacer Mediates the Function of Class III Pl3 and PS Complexes in Autophagosome Maturation by Engaging Stx1 . *Mol Cell.* 201 Mar 16 65(6) 102 -1043.e5. IF: 14.248
- ing , Pan C, Shao T, et al. Mixed Lineage inase Domain-like Protein ML L Breaks Down Myelin following Nerve Injury . Mol Cell. 201 Nov 1 2(3) 45 -46 .e5. IF: 14.248
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- i , Si , hang , et al. Conferring DNA virus resistance with high speci city in plants using virus-inducible genome-editing system . enome iol. 201 Nov 15 1 (1) 1 . IF: 13.214
- uang , Gu L, hang , et al. An oomycete plant pathogen reprograms host pre-mRNA splicing to subvert Immunity .

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- uang C, ang F, hang , et al. Mrg15 stimulates Ash1 3 36 methyltransferase activity and facilitates Ash1 Trithorax group protein function in Drosophila . *Nature Communications*. 201 Nov21 (1) 164 . IF:12.124
- in , Liu , Luo D, et al. DELLA proteins are common components of symbiotic rhizobial and mycorrhizal signalling pathways . *Nature Communications*. 2016 Aug 12 12433. IF: 11.33
- e M, u , Chen , et al. MoSnt2-dependent deacetylation of histone 3 mediates MoTor-dependent autophagy and plant infection by the rice blast fungus Magnaporthe oryzae . *Autophagy*. 201 14() 1543-1561. IF: 11.1
- Tan L M, hang C , ou M, et al. The PEAT protein complexes are required for histone deacetylation and heterochromatin silencing . *EM* . 201 ct 1 3 (1). pii e 0. IF: 10.557



Fast Mutagenesis

- ing , ao R W, hang , et al. SLERT Regulates DD 21 Rings Associated with Pol I Transcription . Cell. 201 May 4 16 (4) 664-6 .e16. IF: 30.409
- Li , Liu C , ue W, et al. Coordinated circRNA Biogenesis and Function with NF 0/NF110 in Viral Infection . *Mol Cell.* 201 ul 20 6 (2) 214-22 .e . IF: 14.713
- Mo F, huang , Liu , et al. Acetylation of Aurora B by TIP60 ensures accurate chromosomal segregation . *Nat Chem iol.* 2016 Apr 12(4) 226-32. IF: 14.273
- u D, hang T, iao , et al. Modi cation of BECN1 by ISG15 plays a crucial role in autophagy regulation by type IFN/interferon . *Autophagy*. 2015 Apr 3 11(4) 61 -2 . IF: 11.753
- uang W , Liu , McCormick S, et al. Tomato Pistil Factor STIG1 Promotes in Vivo Pollen Tube Growth by Binding to Phosphatidylinositol 3-Phosphate and the ExtraCellular Domain of the Pollen Receptor inase LePR 2 . *Plant Cell*. 2014 un 26(6) 2505-2523, IF: 10.125



Traditional Total RNA Isolation

Chen B, ou W, u, et al. Ef cient labeling and imaging of protein-coding genes in living cells using CRISPR-Tag. Nature Communications, 201, (1) 5065. IF: 12.353



RNA Tissue eeper

ang L,Li,Gong R, et al.The Long Non-coding RNA- RLNC1 Regulates Bone Mass by Directing Mesenchymal Stem Cell Fate .201, Mol Ther, 2 (2) 3 4-410. IF: 7.008



miRNA

Wang M, Wu W, Li L, et al. Analysis of the miRNA Expression Pro les in the earalenone-Exposed TM3 Leydig Cell Line . *International ournal of molecular sciences*, 201, 20(3) 635. IF: 3.687



Reverse Transcription

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Meng , Wang , Brunetti T, et al. The DGCR5 long noncoding RNA may regulate expression of several schizophrenia-related genes . *Sci Transl Med.* 201 Dec 1 10(4 2). pii eaat6 12. IF: 16.71

in S, Tian S, Luo M, et al. Tetherin Suppresses Type I Interferon Signaling by Targeting MAVS for NDP52-Mediated Selective Autophagic Degradation in uman Cells . *Mol Cell*. 201 ct 1 6 (2) 30 -322.e4. IF: 14.248

Guo M, Li C, Lei , et al . Role of the adipose PPAR -adiponectin axis in susceptibility to stress and depression/anxiety-related behaviors . *Mol Psychiatry*. 201 ul 22() 1056-106 . IF: 13.3

ang L, Wang W , iu W L, et al. A single-Cell transcriptomic analysis reveals precise pathways and regulatory mechanisms underlying hepatoblast differentiation . *Hepatology.* 201 Nov 66(5) 13 -1401. IF: 13.246

Chen B, ou W, u , et al. Ef cient labeling and imaging of protein-coding genes in living Cells using CRISPR-Tag .

Nature Communications. 201 Nov 2 (1) 5065. IF: 12.353

Liu, in ,Wu C, et al. Downregulated NDR1 protein kinase inhibits innate immune response by initiating an miR146a-STAT1 feedback loop. . Nature Communications. 201 ul 1 (1) 2 . IF: 12.353

Sun , Ding , han M, et al. Usp regulates ippo pathway through deubiquitinating the transcriptional coactivator orkie . Nature Communications. 201 an 24 10(1) 411. IF: 12.353

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Liu M, Shi , hang , et al. Inducible overexpression of Ideal Plant Architecture1 improves both yield and disease resistance in rice . *Nat Plants*. 201 Apr 5(4) 3 -400. IF: 11.471

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