



Reagents for Molecular Biology Research

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Overview of Vazyme

Vazyme: InoVation in Enzyme Technology

With the faith of "InnoVation in Enzyme Technology", Vazyme Biotech Co., Ltd. has passionately focused on developing enzyme and antibody technologies and products for years. The Vazyme Biotech is now staffed by more than 1,000 employees. The headquarter is located in Nanjing of China with a R&D / manufacturing base that covers 25,000 m² and a GMP workshop of 4,000 m². Vazyme has developed a powerful sales network in China and is expanding into international markets.



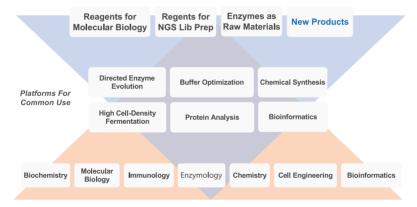




Vazyme Technologies and Products

With years of experience, Vazyme has developed six technology platforms that can be commonly used for R&D and manufacturing, including (1) directed enzyme evolution, (2) buffer optimization, (3) chemical synthesis, (4) high cell-density fermentation, (5) protein analysis, and (6) bioinformatics.

Based on the above platforms, Vazyme now provides a variety of products, solutions, and services, which fall into three major product lines, including (1) solutions for molecular biology research, (2) solutions for Next-Generation Sequencing (NGS) library preparation, and (3) enzymes as raw materials for industrial use.



Developing Technologies to Improve Human Health

Fascinated by the enzyme and antibody technologies, we regard enzymes and antibodies as the key factor of the biotechnology industry. Vazyme's vision is to develop technology to improve human health.

Experts for Experts

2020 Vazyme Product Catalogue

Reagents for Molecular Biology Research

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PCR

High-Fidelity PCR

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

Conventional PCR

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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 (Mg ²⁺ 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
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 x Tag Plus Master Mix II (Dve Plus)	5 ml / 15 ml / 50 ml	P213-01/02/03

Long-Fragment PCR

5 5		
Product Name		Cat. No.#
Vazyme LAmp DNA Polymerase (Mg ²⁺ plus buffer)	125 U / 500 U	P301-01/02
Vazyme LAmp DNA Polymerase (Mg2+ plus buffer, with dNTP)	125 U / 500 U	P301-d1/d2
Vazyme LAmp DNA Polymerase (Mg ²⁺ free buffer, with dNTP)	125 U / 500 U	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	One Step Mouse Genotyping Kit	200 rxn	PD101-01
	One Step U* Probe Mouse Genotyping Kit	200 rxn	PD104-01
	Blood Direct PCR Kit V2	50 rxn / 200 rxn	PD103-01/02

Rapid PCR

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

Multiplex PCR

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



■ Hot-Start PCR

	Cat. No.#
250 U / 1,000 U / 3,000 U	P401-d1/d2/d3
1 ml / 5 ml /15 ml	P411-01/02/03
1 ml / 5 ml / 15 ml	P412-01/02/03
500 U	P121-01
500 U (2.5 / 5 / 10 U/μI)	P122-d1/d2/d3
	250 U / 1,000 U / 3,000 U 1 ml / 5 ml /15 ml 1 ml / 5 ml / 15 ml 500 U

Product Name	Size	Cat. No.#
Taq Pro HS DNA Polymerase	250 U / 1,000 U / 5,000 U	PN101-01/02/03
Taq Pro HS Master Mix	500 rxn / 1,500 rxn (20 μl/rxn)	PN111-01/02
Taq Pro HS U⁺ Master Mix	500 rxn / 1,500 rxn (20 μl/rxn)	PN112-01/02

■ Isothermal Amplification

Product Name		Cat. No.#
Bst DNA Polymerase Large Fragment	800 U / 8,000 U	P701-01/02

PCR-Related

Product Name	Size	Cat. No.#
PCR Enhancer	500 μl	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
Heat-labile UDG	100 U / 500 U	P051-01/02
E.coli UDG	500 U / 5,000 U	P061-01/02

Cloning / Mutagenesis

■ Fast Cloning

Product Name		
ClonExpress II One Step Cloning Kit	25 rxn / 50 rxn	C112-01/02
ClonExpress MultiS One Step Cloning Kit	10 rxn / 25 rxn	C113-01/02
(HOT) ClonExpress Ultra One Step Cloning Kit	25 rxn / 50 rxn	C115-01/02

■ Fast Mutagenesis

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

■ TA Cloning

	T4 DNA Ligase	40,000 U	C301-01
ew	5min Universal Ligation Mix	50 rxn / 100 rxn	C311-01/02

■ TOPO Cloning

Product Name		
101 5min TA/Blunt-Zero Cloning Kit	25 rxn / 50 rxn	C601-01/02



Nucleic Acid Electrophoresis

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		Cat. No.#
DL2000 Plus DNA Marker	250 µl / 500 µl	MD101-01/02
DL5000 DNA Marker	250 µl / 500 µl	MD102-01/02
DL15000 DNA Marker	250 µl / 500 µl	MD103-01/02
100 bp DNA Ladder	250 µl / 500 µl	MD104-01/02

Reverse Transcription

■ Conventional RT-PCR

Product Name		Cat. No.#
HiScript II Reverse Transcriptase	2,000 U / 10,000 U	R201-01/02
HiScript III Reverse Transcriptase	10,000 U	R302-01
HiScript II 1st Strand cDNA Synthesis Kit	50 rxn / 100 rxn (20 μl / rxn)	R211-01/02
HiScript III 1st Strand cDNA Synthesis Kit (+gDNA wiper)	50 rxn / 100 rxn (20 μl / rxn)	R312-01/02
M-MLV(H-) 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

Pr	oduct Name		Cat. No.#
His	Script II Q RT SuperMix for qPCR	100 rxn (20 µl / rxn)	R222-01
His	Script II Q RT SuperMix for qPCR (+gDNA wiper)	100 rxn (20 µl / rxn)	R223-01
His	Script II Q Select RT SuperMix for qPCR	100 rxn (20 µl / rxn)	R232-01
His	Script II Q Select RT SuperMix for qPCR (+gDNA wiper)	100 rxn (20 μl / rxn)	R233-01
His	Script III RT SuperMix for qPCR (+gDNA wiper)	100 rxn (20 µl / rxn)	R323-01
His	Script III All-in-one RT SuperMix Perfect for qPCR	100 rxn (20 µl / rxn)	R333-01

■ One-Step RT-PCR

Product Name		Cat. No.#
HiScript II One Step RT-PCR Kit	50 rxn (50 μl / rxn)	P611-01
HiScript II One Step RT-PCR Kit (Dve Plus)	50 rxn (50 ul / rxn)	P612-01

■ Single Cell Sequence Amplification

Product Name		Cat. No.#
Single Cell Sequence Specific Amplification Kit	200 rxn	P621-01

miRNA Reverse Transcription

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

HOT



qPCR

qPCR Master Mix (SYBR-Green)

	Product Name	Size	Cat. No.#
HOT	ChamQ Universal SYBR® qPCR Master Mix	500 rxn / 2,500 rxn (20 μl / rxn)	Q711-02/03
	AceQ Universal SYBR® qPCR Master Mix	500 rxn / 2,500 rxn (20 μl / rxn)	Q511-02/03
	AceQ qPCR SYBR® Green Master Mix	500 rxn / 2,500 rxn (20 μl / rxn)	Q111-02/03

qPCR Master Mix (Probe)

	Product Name	Size	Cat. No.#
	AceQ qPCR Probe Master Mix	500 rxn / 2,500 rxn (20 µl / rxn)	Q112-02/03
HOT	AceQ Universal U* Probe Master Mix V2	500 rxn / 2,500 rxn (20 µl / rxn)	Q513-02/03
HOT	ChamQ Geno-SNP Probe Master Mix	500 rxn / 2,500 rxn (20 µl / rxn)	Q811-02/03
	Animal Detection U* Probe Master Mix	5 x 1 ml / 1 x 10 ml	QV110-01/02
HOT	Taq Pro HS Probe Master Mix	500 rxn / 2,500 rxn (20 µl / rxn)	QN111-01/02
HOT	Taq Pro HS U⁺ Probe Master Mix	500 rxn / 2,500 rxn (20 μl / rxn)	QN112-01/02
HOT	Taq Pro HS Universal Probe Master Mix	500 rxn / 2,500 rxn (20 μl / rxn)	QN113-01/02
HOT	Taq Pro HS Universal U* Probe Master Mix	500 rxn / 2,500 rxn (20 μl / rxn)	QN114-01/02

■ One-Step qRT-PCR Mix

Product Name		Cat. No.#
HiScript II One Step qRT-PCR SYBR® Green Kit	250 rxn (20 µl / rxn)	Q221-01
HiScript II One Step qRT-PCR Probe Kit	250 rxn (20 µl / rxn)	Q222-01
HiScript II U+ One Step qRT-PCR Probe Kit	250 rxn (20 µl / rxn)	Q223-01
HiScript II U One Step gRT-PCR Probe Kit	5000 rxn (30 μl / rxn)	Q222-CN

miRNA qPCR

Product Name	Size	Cat. No.#
miRNA Universal SYBR® qPCR Master Mix	125 rxn / 500 rxn (20 ul / rxn)	MQ101-01/02

Genome Editing

Product Name		Cat. No.#
Cas9 Nuclease	50 pmol / 250 pmol	EN301-01/02
T7 Endonuclease I	50 pmol / 250 pmol	EN303-01/02

In Vitro Transcription

	Product Name			
	T7 High Yield RNA Transcription Kit	50 rxn / 100 rxn	TR101-01/02	
нот	T7 RNAi Transcription Kit	25 rxn / 50 rxn	TR102-01/02	



Nucleic Acid Isolation

RNA Isolation (Column)

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

■ DNA Isolation (Column)

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

■ Tissue Stabilizer

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

■ Exosome Isolation

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

Cell Biology / Protein Research

■ Cell Counting

	Product Name	Size	Cat. No.#
HOT	CCK-8 Cell Counting Kit	500 rxn / 1,000 rxn	A311-01/02

■ Dual Luciferase Reporter Assay

	Product Name	Size	Cat. No.#	ı
HOT	Dual Luciferase Reporter Assay Kit	100 rxn	DL101-01	
	Duo-Lite Luciferase Assay System	10 ml / 100 ml	DD1205-01/02	

Mycoplasma

	Product Name		Cat. No.#
HOT	MycoBlue Mycoplasma Detector	20 rxn / 50 rxn	D101-01/02

Protein Marker

	Product Name		Cat. No.#
HOT	180 kDa Prestained Protein Marker	2 × 250 μl / 10 × 250 μl (5 μl/rxn)	MP102-01/02



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 identification; TA Cloning for small fragments.
High-Yield PCR	2× Taq Plus Master Mix (#P211) 2× Taq Plus Master Mix II (Dye Plus) (#P213)	With fidelity 6-fold higher than Taq. Mixed products with 3"-end blunt or containing A.	PCR that requires some fidelity.
Rapid PCR	2× Rapid Taq Master Mix (#P222)	Amplification speed: up to 15 sec / kb.	Colony PCR.
Long-Fragment PCR	2× Vazyme LAmp Master Mix (#P311) 2× Vazyme LAmp Master Mix (Dye Plus) (#P312)	Efficiently amplify fragments > 20 kb.	Long-fragment amplification.
Hot-Start PCR	2× AceTaq Master Mix (#P411) 2× AceTaq Master Mix (Dye Plus) (#P412) Champagne Taq Antibody (#P121) Champagne Taq DNA Polymerase (#P122) Taq Pro HS DNA Polymerase (#PN101) Taq Pro HS Master Mix (#PN111) Taq Pro HS U+ Master Mix (#PN112)	Excellent specificity. Excellent sensitivity.	Amplification that requires higher sensitivity and specificity; Amplification of genes with low copy or qPCR assay from complex templates (genomic DNA, cDNA).
Multiplex PCR	Multiplex PCR Kit (#PM101)	19-plex PCR in one single reaction.	Detection or typing of pathogens.
Direct PCR	One Step Mouse Genotyping Kit (#PD101) Blood Direct PCR Kit V2 (#PD103)	Easy and fast, without DNA purification.	One step mouse genotyping; Direct PCR from plant tissues; Direct PCR from blood.
High-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 fidelity 53-fold higher than Taq; High resistance to PCR inhibitors.	High-fidelity PCR. Amplification of templates with high GC-content; Long-fragment (up to 40 kb) amplification.



High-Fidelity PCR



→ 2× Phanta Max Master Mix (#P515)

→ 2× 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

Zhao Q, et al. Metabolic coupling of two small-molecule thiols programs the biosynthesis of lincomycin A. *Nature*, 2015, 518(7537):115-9.

Tian Z, et al. An enzymatic [4+2] cyclization cascade creates the pentacyclic core of pyrroindomycins. *Nature Chemical Biology*, 2015, 11(4):259-65.

Han X, et al. Mapping the Mouse Cell Atlas by Microwell-Seq. Cell, 2018, 172(5):1091-107.

Cheng X, et al. Pacer Mediates the Function of Class III PI3K and HOPS Complexes in Autophagosome Maturation by Engaging Stx17. *Molecular Cell*, 2017, 65(6):1029-43.

Lv M, et al. Characterization of a C3 Deoxygenation Pathway Reveals a Key Branch Point in Aminoglycoside Biosynthesis. *Journal of the American Chemical Society*, 2016, 138(20):6427-35.



High-Yield PCR

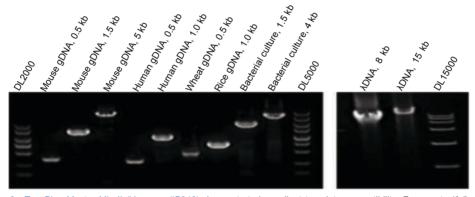


→ 2× Taq Plus Master Mix II (Dye Plus) (#P213)

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



2× 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.

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Selected Product Citations

Zhang X, et al. (2014) Complementary sequence-mediated exon circularization. Cell, 159(1):134-47.

Yuan H, et al. Gyrl-like proteins catalyze cyclopropanoid hydrolysis to confer cellular protection. *Nature Communications*, 2017, 8(1).1485.



Rapid PCR

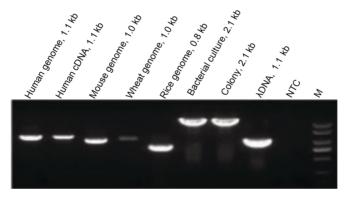


→ 2× Rapid Tag Master Mix (#P222)

Features

- * Rapid: amplification 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

Zhang B, et al. Enzyme-catalysed [6+4] cycloadditions in the biosynthesis of natural products. Nature, 2019, 568(7750):122-6.

Wang YS, et al. Molecular Basis for the Final Oxidative RearrangementSteps in Chartreusin Biosynthesis. J Am Chem Soc, 2018, 140(34):10909-14.



Multiplex PCR

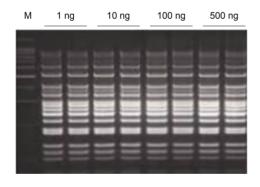


→ Multiplex PCR Kit (#PM101)

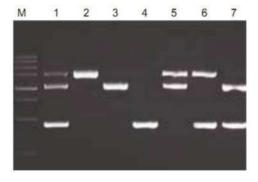
Features

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

Validation Data



Uniformamplification coverage of different regions. 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			
Fast Cloning	ClonExpress Ultra One Step Cloning Kit (#C115) ClonExpress II One Step Cloning Kit (#C112) ClonExpress MultiS One Step Cloning Kit (#C113)	Easy, fast, and efficient. No need to consider the restriction enzyme cutting sites on the inserts. Ligase-independent. Positive Clone Rate > 95%. Efficient cloning of fragments of 50 bp - 10 kb.	Cloning or assembly of 1-5 fragments.
Fast Mutagenesis	Mut Express II Fast Mutagenesis Kit V2 (#C214) Mut Express MultiS Fast Mutagenesis Kit V2 (#C215)	Efficient amplification 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.
		Cloning within 5 min. Positive Clone Rate > 95%	TA cloning. cloning with blunt ends.

TOPO Cloning

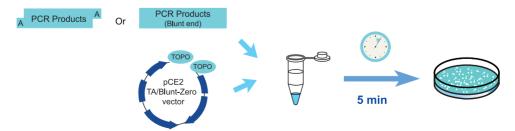




Features

- * Ready-to-use master mix.
- * Suitable for both TA cloning and blunt-end cloning.
- * Rapid cloning within 5 min.
- * High cloning efficiency with Positive Clone Rate > 95%.
- * Ampicillin and Kana dual resistance vector.

Workflow





Fast Cloning



→ ClonExpress Ultra One Step Cloning Kit (#C115)

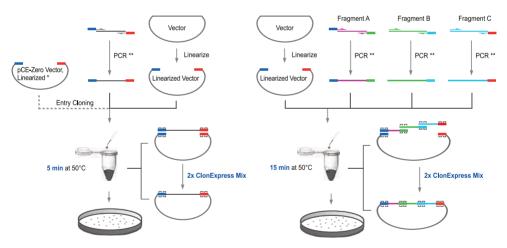
Features

- * Cloning within 5 min.
- * Ready-to-use super mix in one tube.
- * Efficient cloning of fragments of 50 bp 10 kb with Positive Clone Rate > 95%.
- * Suitable for cloning of 1 fragment, assembly of 2 5 fragments, and entry cloning.
- * Independent of DNA ligase, significantly reducing the self-ligated colonies.

Mechanism

Cloning of 1 Fragment

Assembly of 2 - 5 fragments.



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

Selected Product Citations of ClonExpress

Wu N, et al. TBX6 null variants and a common hypomorphic allele in congenital scoliosis. New England Journal of Medicine, 2015, 372(4):341-50.

Ge J, et al. Architecture of the mammalian mechanosensitive Piezo1 channel. Nature, 2015, 527(7576):64-9.

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





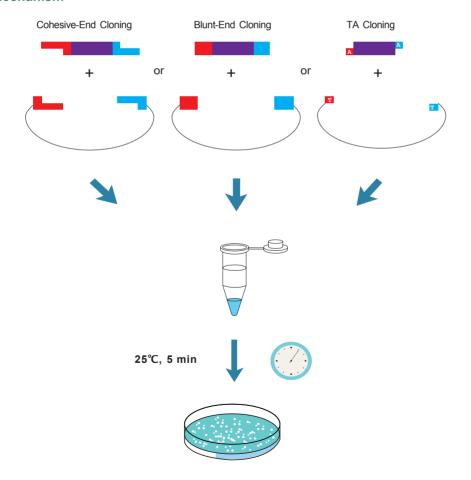


→ 5min Universal Ligation Mix (#C311)

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.
- * Efficient: Positive Clone Rate > 95%.

Mechanism

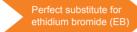




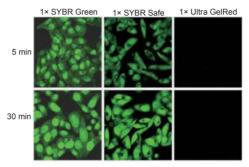
GelRed



→ Ultra GelRed Nucleic Acid Stain (10000×) (#GR501)



No toxicity



Ultra GelRed is unable to cross cell membranes.

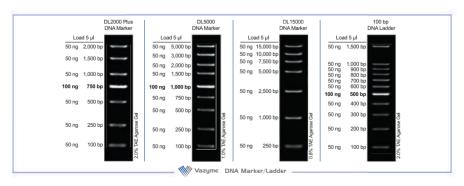
DNA Marker/ Ladder



→ DNA Markers / Ladders

Stable

Clear Bands



Reverse Transcription

Selection Guide

		JHA .	contact i	nklik st	Mark for apock	Superfult for	Jupe Mix for	EDWA MX for
	HEGITA II SH	HEERDE II SEEDE	HERITORY LOWE	History of Super	Herder NO Select AT	HEERE HAD ENGLISHED F	HEERT HEERT HEERT	HEER HALE SORHWAY
Applications								
RT-qPCR								
RT-PCR								
Features								
SuperMix								
Long-fragment cDNA								
Rapid removal of Genomic DNA								
Primers								
Oligo dT ₂₃ VN / N6 Mix								
Optional								

	M-MLV (H-) (#R021)	HiScript II Reverse Transcriptase (#R201)	HiScript III Reverse Transcriptase (#R302)
Reaction temperature	37℃ - 42℃	42℃ - 55℃	37℃ - 50℃
Thermal stability	_ራ አ አ	***	ታ ታ ታ ታ
RNase H activity	No	No	No
cDNA length	2 kb - 3 kb	Up to 20 kb	Up to 20 kb
Template adaptability	☆☆☆	***	***
Crude material adaptability	☆☆☆	***	***

RT-qPCR SuperMix



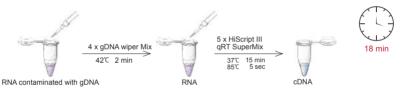
→ HiScript III RT SuperMix for qPCR (+gDNA wiper) (#R323)

Features

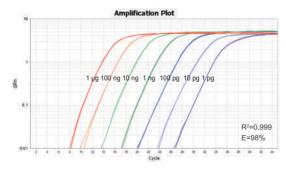
- * Ready-to-use SuperMix: reverse transcription within 20 min by only adding template RNA.
- * Excellent efficiency for low-input RNA or degraded RNA.
- * Excellent tolerance for impurities (i.e. ethanol, isopropanol, phenol water, guanidine thiocyanate, humic acid).
- * Lower C_T value and higher efficiency 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 ACTB. The results show an excellent linear relationship across a wide range of RNA concentrations. The target gene (ACTB) was detected in 1 pg of RNA.



qPCR

Selection Guide

Applications	Products (Cat.#)	
SYBR	ChamQ Universal SYBR QPCR Master Mix (#Q711)	
Probe	AceQ Universal U+ Probe Master Mix V2 (#Q513)	
SNP (TaqMan MGB Probe)	ChamQ Geno-SNP Probe Master Mix (#Q811)	

qPCR Master Mix (SYBR)



→ ChamQ Universal SYBR® qPCR Master Mix (#Q711)

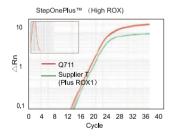
Best Combination of Specificity + Sensitivity

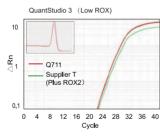
Unique Hot-Start Tag

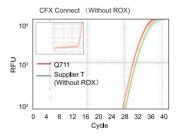
- Unique specificity-promoting Factors
- Optimal Concentrations of Mg²⁺ and Dye
- Universal

Validation Data

Applicable for almost all qPCR instruments.







~W

Selected Product Citations

Xu L, et al. The transcription factor TCF-1 initiates the differentiation of TFH cells during acute viral infection. *Nature Immunology*, 2015, 47(3):538-51.

Guo C, et al. Cholesterol Homeostatic Regulator SCAP-SREBP2 Integrates NLRP3 Inflammasome Activation and Cholesterol Biosynthetic Signaling in Macrophages. *Immunity*, 2018, 49(5): 842-56.



qPCR Master Mix (Probe)



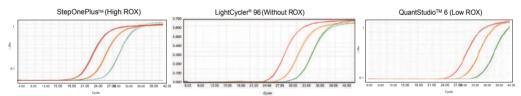
AceQ Universal U+ Probe Master Mix V2 (#Q513)

Features

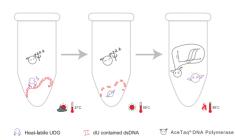
- * Excellent sensitivity: Hot-start AceTaq and optimal buffer ensure high sensitivity and effectively inhibit non-specific amplification.
- * Excellent linear relationship over a large range of input amount of template. Suitable for the detection of single-copy templates.
- * Anti-contamination: the dUTP/UDG system eliminates possible contaminations and ensures reliable results.
- * Universal: applicable for almost all qPCR instruments.

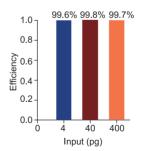
Validation Data

1. Applicable for almost all qPCR instruments.



2. dUTP/UDG system.





For Vazyme #Q513, 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) were added respectively to the reaction system to evaluate the removal efficiency of the contaminated template by Vazyme #Q513.



qPCR Master Mix (Probe)



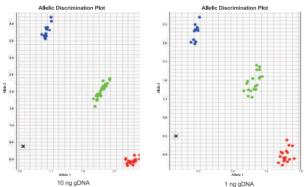
→ ChamQ Geno-SNP Probe Master Mix (#Q811)

Advantages

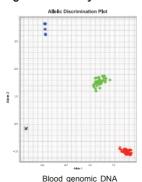
- * Compatible with 1 ng 10 ng of input genomic DNA.
- * Accurate genotyping of SNP sites with GC-content of 25% 73%.
- * Excellent stability: stable signal and accurate genotyping results can be obtained both 72 hr pre-PCR and 72 hr post-PCR.
 - * 72 hr pre-PCR: PCR reaction solutions were prepared and left in darkness (at room temperature) for 72 hr before PCR;
 - * 72 hr post-PCR: after PCR, the samples were left in darkness (at room temperature) for 72 hr.
- * Blood lysate can be directly used as a template for SNP genotyping, with no need for blood genomic DNA extraction.

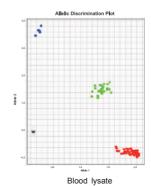
Validation Data

1. Flexible input amounts.



2. Direct genotyping with blood lysate.







Nucleic Acid Isolation

Selection Guide

Category	Series	Sample / Application	Products	Cat.#
		Blood	FastPure Blood DNA Isolation Mini Kit V2	DC111
		Cell / Tissue	FastPure Cell/Tissue DNA Isolation Mini Kit	DC102
DNA Isolation	DNA Extraction (Column)	Bacterial FastPure Bacteria DNA Isolation Mini Kit		DC103
& Purification	(Column)	Plant	FastPure Plant DNA Isolation Mini Kit	DC104
		FFPE	FastPure FFPE DNA Isolation Kit	DC105
		Lysozyme	Lysozyme	DE103
	RNA Tissue Keeper	RNA Keeper for fresh tissue	RNA Keeper Tissue Stabilizer	R501
RNA Isolation	O-IIII	Cell / tissue total RNA	FastPure Cell/Tissue Total RNA Isolation Mini Kit	RC101
		Polysaccharide & Polyphenol-ric Plant total RNA		FastPure Plant Total RNA Isolation Kit (Polysaccharides & Polyphenolics-rich)
	Cell supernatant		VEX Exosome Isolation Reagent (from cell culture media)	
Exosome Isolation	Serum		VEX Exosome Isolation Reagent (from serum)	R602
	Plasma		VEX Exosome Isolation Reagent (from plasma)	R603

Plant RNA and DNA Isolation





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

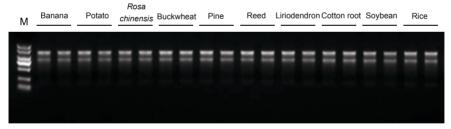
Features

- * 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 japonica 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, flower 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.

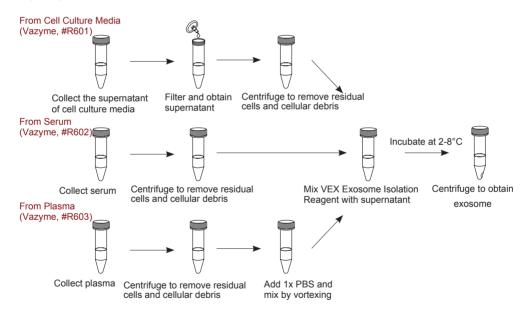
Exosome Isolation



Features

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

Workflow



Cell Counting

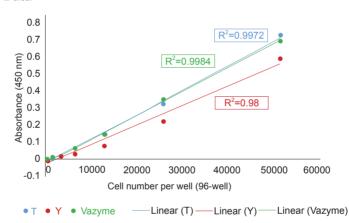


CCK-8 Cell Counting Kit (#A311)

Features

- * Ready-to-use solution.
- * High 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, 800, 1600, 3200, 6400, 12800, 25600, 51200 cells per well. CCK-8 reagents from Vazyme (#A311, green), Supplier T (blue), and Supplier Y (red) were used for cell counting, respectively. The \mathbb{R}^2 value of Vazyme #A311 is > 0.99.

Selected Product Citations

Zheng Q, et al. Thiopeptide antibiotics exhibit a dual mode of action against intracellular pathogens by affecting both host and microbe. *Chemistry & Biology*, 2015, 22(8):1002-7.

Liu Z, et al. Adiponectin reduces ER stress-induced apoptosis through PPAR α transcriptional regulation of ATF2 in mouse adipose. *Cell Death & Disease*, 2016, 7(11):e2487.

Luciferase Assay

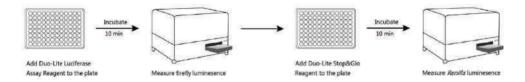


→ Duo-Lite Luciferase Assay System (#DD1205)

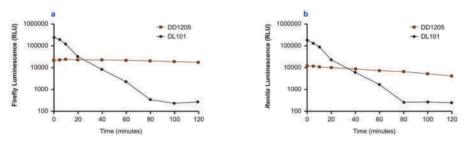
Features

- * Easy to operate: The experiment could be completed by two steps: adding sample and reading plate. There is no need for cell lysis.
- * Stable signal: Glow-type kit with 2h half-life of fluorescence. Suitable for high-throughput operation.
- * High accuracy: The system contains Renilla luciferase, which could correct the errors that caused by differences among cell number, transfection efficiency and cell growth state.

Workflow



Validation Data



Sample: HEK293 cells co-transfected with firefly + plasmid (96 well plate incubate)

Experimental design: Detect the dynamic change of fluorescent values of glow-type kit (Vazyme #DD1205) and flash-type kit (Vazyme #DL101) within 120 min simultaneously.

Conclusion: Compared with flash-type kit (Vazyme #DL101), glow-type kit (Vazyme #DD1205) shows higher light stability. The half-life of fluorescence is up to 2h.

Luciferase Assay

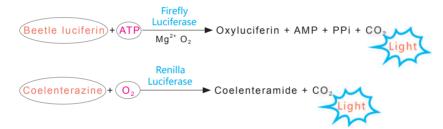


→ Dual Luciferase Reporter Assay Kit (#DL101)

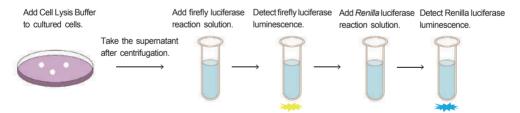
Features

- * Robust luminescent signals: applicable for analysis of weak promoters and other genetic regulatory elements.
- * Detection linear range covers up to 8 orders of magnitude (R² > 0.99).
- * Detection sensitivity of 10⁻¹⁸ mole.

Mechanism



Workflow



Selected Product Citations

Liu Z, et al. Circular RNA hsa_circ_001783 regulates breast cancer progression via sponging miR-200c-3p. *Cell Death & Disease*, 2019, 10:55

Wu H, et al. Ubiquitination is essential for avibirnavirus replication by supporting VP1 Polymerase activity. *Journal* of *Virology*, 2019, 93(3): e01899-18.

Wu H, et al. SUMO1 Modification Facilitates Avibirnavirus Replication by Stabilizing Polymerase VP1. *Journal of Virology*, 2019, JVI. 02227-18.

Mycoplasma Detection



→ Myco-Blue Mycoplasma Detector (#D101)

Features

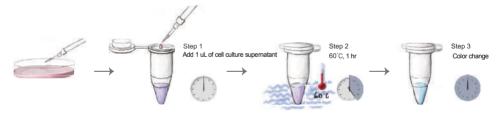
- * Cell culture supernatant can be used directly for detection.
- * Results are obtained after incubation at 60°C for 1 hr and can be determined by visual observation.
- * Accuracy is higher than PCR method, and comparable to gPCR method.
- * Suitable for detection of all kinds of mycoplasma that are commonly found in cell culture.

Validated Cell Lines

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

- * Suspension cells: CHO, NS0, 293F, mouse hybridoma, Sf9, BHK21, etc.;
- * Adherent cells: Vero, MDCK, SP2/0, 293T, HepG2, HeLa, A549, MB-MDA231, L929, MEF, etc.;
- * Medium: CD FortiCHO, CDM4, Expi 293 Medium, CD Hybridoma, Grace, DMEM, 1640, F12, etc.;
- * Serum: fetal calf / calf serum; horse serum; Gibco KSR serum replacement, etc.

Workflow



Validated Data



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



In Vitro Transcription

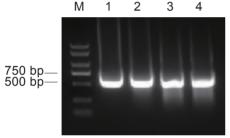


Features

- * High vield: vields up to 80 ug of dsRNA in a single reaction.
- * Magnetic bead purification: recovery efficiency up to 80%.
- * Able to transcribe both siRNA (21 bp) and dsRNA (long fragment).

Validation Data

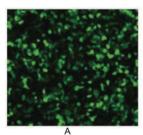
1. Excellent transcription efficiency.

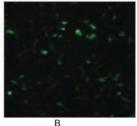


Agarose gel electrophoresis (2%) of 500 bp dsRNA.

- M: DL2000 Plus DNA Maker.
- 1 and 3: products before and after enzymatic hydrolysis of dsRNA, respectively;
- 2 and 4: products before and after enzymatic hydrolysis of dsRNA, respectively.

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





293T cells were co-transfected for 24 hrs with both GFP plasmid and negative control GFP siRNA (A) or positive GFP siRNA (B).



High-Fidelity PCR

Zhao Q, Wang M, Xu D, et al. Metabolic coupling of two small-molecule thiols programs the biosynthesis of lincomycin A[J]. Nature. 2015 Feb5;518(7537):115-9. IF: 42.351

Zhang B, Wang K B, Wang W, et al. Enzyme-catalysed [6+4] cycloadditions in the biosynthesis of natural products[J]. Nature. 2019 Apr;568(7750):122-126. IF: 41.577

Ma Z, Zhu L, Song T, et al. A paralogous decoy protects Phytophthora sojae apoplastic effector PsXEG1 from a host inhibitor[J]. Science. 2017 Feb 17:355(6326):710-714. IF: 34.661

Han X, Wang R, Zhou Y, et al. Mapping the MouseCell Atlas by Microwell-Seq[J]. Cell. 2018 May17;173(5):1307. IF: 31.398

Zhang B, Li J, Yang X, et al. Crystal Structures of Membrane Transporter MmpL3, an Anti-TB Drug Target[J]. Cell. 2019 Jan 24;176(3):636-648.e13. IF: 31.398

Wang YS, Zhang B, Zhu J, et al. Molecular Basis for the Final Oxidative Rearrangement Steps in Chartreusin Biosynthesis[J]. J Am Chem Soc. 2018 Aug 29:140(34):10909-10914. IF: 14.357

Cheng X, Ma X, Ding X, et al. Pacer Mediates the Function of Class III Pl3K and HOPS Complexes in Autophagosome Maturation by Engaging Stx17[J]. *Mol Cell.* 2017 Mar 16;65(6):1029-1043.e5. IF: 14.248

Wu H, Yin QF, Luo Z, et al. Unusual Processing Generates SPA LncRNAs that Sequester Multiple RNA Binding Proteins[J]. Mol Cell. 2016 Nov 3;64(3):534-548. IF: 13.958

Tian Z, Sun P, Yan Y, et al. An enzymatic [4+2] cyclization cascade creates the pentacyclic core of pyrroindomycins[J]. *Nat Chem Biol.* 2015 Apr;11(4):259-65. **IF: 13.217**

Zhang M, Zhou C, Wei Y, et al. Human cleaving embryos enable robust homozygotic nucleotide substitutions by base editors[J]. Genome Biol. 2019 May 22;20(1):101. IF: 13.214

Wang M, Zhao Q, Zhang Q, et al. Differences in PLP-Dependent Cysteinyl Processing Lead to Diverse S-Functionalization of Lincosamide Antibiotics[J]. *J Am Chem Soc.* 2016 May 25;138(20):6348-51. **IF:** 13.038

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Sun X, Ding Y, Zhan M, et al. Usp7 regulates Hippo pathway through deubiquitinating the transcriptional coactivator Yorkie[J]. Nature Communications. 2019 Jan 24;10(1):411. IF: 12.353



Conventional PCR

Zhang B, Wang K B, Wang W, et al. Enzyme-catalysed [6+4] cycloadditions in the biosynthesis of natural products[J]. Nature. 2019 Apr;568(7750):122-126. IF: 41.577

Zhang XO, Wang HB, Zhang Y, et al. Complementary sequence-mediated exon circularization[J]. Cell. 2014 Sep 25;159(1):134-147. IF: 33.116

Wang Y S, Zhang B, Zhu J, et al. Molecular Basis for the Final Oxidative Rearrangement Steps in Chartreusin Biosynthesis[J]. J Am Chem Soc. 2018 Aug 29;140(34):10909-10914. IF: 14.357

Sun H, Liu J, Zheng Y, et al. Distinct chemokine signaling regulates integrin ligand specificity to dictate tissue-specific lymphocyte homina[J]. Dev Cell. 2014 Jul 14:30(1):61-70. IF: 12.86

Yuan H, Zhang J, Cai Y, et al. Gyrl-like proteins catalyze cyclopropanoid hydrolysis to confer Cellular protection[J]. *Nature Communications*. 2017 Nov 14;8(1):1485. IF: 12.124

Zhang X, Wang T T, Xu Q L, et al. Genome Mining and Comparative Biosynthesis of Meroterpenoids from Two Phylogenetically Distinct Fungi[J]. *Angew Chem Int Ed Engl.* 2018 Jul 2;57(27):8184-8188. IF: 12.102

Chen C, Zhai S, Zhang L, et al. Uhrf1 regulates germinal center B Cell expansion and affinity maturation to control viral infection[J]. J Exp Med. 2018 May 7;215(5):1437-1448. IF: 11.991

Bai D, Zhang J, Li T, et al. The ATPase hClNAP regulates 18S rRNA processing and is essential for embryogenesis and tumour growth[J]. Nature Communications. 2016 Aug 1;7:12310. IF: 11.33

Zhang X O, Dong R, Zhang Y, et al. Diverse alternative back-splicing and alternative splicing landscape of circular RNAs[J]. Genome Res. 2016 Sep;26(9):1277-87. IF: 11.351



Fast Cloning

Wu N, Ming X, et al. TBX6 null variants and a common hypomorphic allele in congenital scoliosis[J]. N Engl J Med. 2015 Jan 22;372(4):341-50. IF: 54.42

Ge J, Li W, et al. Architecture of the mammalian mechanosensitive Piezo1 channel[J]. *Nature*. 2015 Nov 5;527(7576):64-9. IF: 42.351

Li X, Wang Y, et al. Base editing with a Cpf1-cytidine deaminase fusion[J]. Nat Biotechnol. 2018 Apr 36(4):324-327. IF: 41.667

Jin S, Zong Y, et al. Cytosine, but not adenine, base editors induce genome-wide off-target mutations in rice[J]. Science. 2019 Apr 19;364(6437):292-295. IF: 41.058

Wang X, Li J, Wang Y, et al. Efficient base editing in methylated regions with a human APOBEC3A-Cas9 fusion[J]. *Nat Biotechnol.* 2018 Nov;36(10):946-949. **IF:** 35.724

Zong Y, Song Q, Li C, et al. Efficient C-to-T base editing in plants using a fusion of nCas9 and human APOBEC3A[J]. Nat Biotechnol. 2018 Oct 1. IF: 35.724

Li T, Yang X, et al. Domestication of wild tomato is accelerated by genome editing[J]. Nat Biotechnol. 2018 Oct 1. IF: 35.724

Zhang Y, Li W, et al. Structural damage in the C. elegans epidermis causes release of STA-2 and induction of an innate immune response[J]. *Immunity*. 2015 Feb 17;42(2):309-320. IF: 19.748

Li Q, Li Y, Yang S, et al. CRISPR-Cas9-mediated base-editing screening in mice identifies DND1 amino acids that are critical for primordial germ Cell development[J]. Nat Cell Biol. 2018 Nov;20(11):1315-1325. IF: 19.064

Wang L, Xue W, Yan L, et al. Enhanced base editing by co-expression of free uracil DNA glycosylase inhibitor[J]. Cell Res. 2017 Oct;27(10):1289-1292. IF: 15.606



Fast Mutagenesis

Xing Y H, Yao R W, Zhang Y, et al. SLERT Regulates DDX21 Rings Associated with Pol I Transcription[J]. Cell. 2017 May 4:169(4):664-678.e16. IF: 30.409

Li X, Liu C X, Xue W, et al. Coordinated circRNA Biogenesis and Function with NF90/NF110 in Viral Infection[J]. *Mol Cell*. 2017 Jul 20;67(2):214-227.e7. IF: 14,713

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Xu D, Zhang T, Xiao J, et al. Modification of BECN1 by ISG15 plays a crucial role in autophagy regulation by type I IFN/interferon[J]. Autophagy. 2015 Apr 3;11(4):617-28. IF: 11.753

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Traditional Total RNA Isolation

Chen B, Zou W, Xu H, et al. Efficient labeling and imaging of protein-coding genes in living cells using CRISPR-Tag[J]. Nature Communications, 2018, 9(1): 5065. IF: 12.353



RNA Tissue Keeper

Yang L,Li Y,Gong R, et al.The Long Non-coding RNA-ORLNC1 Regulates Bone Mass by Directing Mesenchymal Stem Cell Fate[J].2019, Mol Ther, 27(2):394-410. IF: 7.008



miRNA

Wang M, Wu W, Li L, et al. Analysis of the miRNA Expression Profiles in the Zearalenone-Exposed TM3 Leydig Cell Line[J]. International journal of molecular sciences, 2019, 20(3): 635. IF: 3.687



Reverse Transcription

Zhou H, Liu J, Zhou C, et al. In vivo simultaneous transcriptional activation of multiple genes in the brain using CRISPR-dCas9-activator transgenic mice[J]. *Nat Neurosci.* 2018 Mar;21(3):440-446. **IF:** 17.839

Meng Q, Wang K, Brunetti T, et al. The DGCR5 long noncoding RNA may regulate expression of several schizophrenia-related genes[J]. Sci Transl Med. 2018 Dec 19;10(472). pii: eaat6912. IF: 16.71

Jin S, Tian S, Luo M, et al. Tetherin Suppresses Type I Interferon Signaling by Targeting MAVS for NDP52-Mediated Selective Autophagic Degradation in Human Cells[J]. *Mol Cell*. 2017 Oct 19;68(2):308-322.e4. IF: 14.248

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Yang L, Wang W H, Qiu W L, et al. A single-Cell transcriptomic analysis reveals precise pathways and regulatory mechanisms underlying hepatoblast differentiation[J]. *Hepatology*. 2017 Nov:66(5):1387-1401. IF: 13.246

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Liu Z, Qin Q, Wu C, et al. Downregulated NDR1 protein kinase inhibits innate immune response by initiating an miR146a-STAT1 feedback loop[J]. *Nature Communications*. 2018 Jul 17;9(1):2789. IF: 12.353

Sun X, Ding Y, Zhan M, et al. Usp7 regulates Hippo pathway through deubiquitinating the transcriptional coactivator Yorkie[J]. *Nature Communications*. 2019 Jan 24;10(1):411. **IF:** 12,353

Xian H, Xie W, Yang S, et al. Stratified ubiquitination of RIG-I creates robust immune response and induces selective gene expression[J]. Sci Adv. 2017 Sep 22;3(9):e1701764. IF: 11.511

Liu M, Shi Z, Zhang X, et al. Inducible overexpression of Ideal Plant Architecture1 improves both yield and disease resistance in rice[J]. Nat Plants. 2019 Apr;5(4):389-400. IF: 11.471

Yin Z, Chen C, et al. Histone acetyltransferase MoHat1 acetylates autophagy-related proteins MoAtg3 and MoAtg9 to orchestrate functional appressorium formation and pathogenicity in Magnaporthe oryzae[J]. Autophagy. 2019 Jul 15(7):1234-1257. [F: 11.1

Liu X, Wei W, Li X, et al. BMI1 and MEL18 Promote Colitis-Associated Cancer in Mice via REG3B and STAT3[J]. Gastroenterology. 2017 Dec;153(6):1607-1620. IF: 18.392

Cheng X, Ma X, Zhu Q, et al. Pacer Is a Mediator of mTORC1 and GSK3-TIP60 Signaling in Regulation of Autophagosome Maturation and Lipid Metabolism[J]. *Mol Cell.* 2019 Feb 21;73(4):788-802.e7. IF: 14.248

Ju J, Chen A, Deng Y, et al. NatD promotes lung cancer progression by preventing histone H4 serine phosphorylation to activate Slug expression[J]. *Nature Communications*. 2017 Oct 13;8(1):928. **IF**: 12.124

Han X, Yu H, Huang D, et al. A molecular roadmap for induced multi-lineage trans-differentiation of fibroblasts by chemical combinations[J]. Cell Res. 2017 Jun;27(6):843. IF: 15.393



QPCR

Guo C, Chi Z, et al. Cholesterol Homeostatic Regulator SCAP-SREBP2 Integrates NLRP3 Inflammasome Activation and Cholesterol Biosynthetic Signaling in Macrophages[J]. *Immunity*. 2018 Nov 20;49(5):842-856.e7. IF: 19,734

Zhou H, Liu J, Zhou C, et al. In vivo simultaneous transcriptional activation of multiple genes in the brain using CRISPR-dCas9-activator transgenic mice[J]. *Nat Neurosci.* 2018 Mar;21(3):440-446. **IF:** 17.839

Meng Q, Wang K, Brunetti T, et al. The DGCR5 long noncoding RNA may regulate expression of several schizophrenia-related genes[J]. Sci Transl Med. 2018 Dec 19:10(472). pii: eaat6912. IF: 16.71

Huang, Li T, Wang L, et al. HepatoCellular carcinoma redirects to ketolysis for progression under nutrition deprivation stress[J]. Cell Res. 2016 Oct;26(10):1112-1130. IF: 14.812

Cheng X, Ma X, Zhu Q, et al. Pacer Is a Mediator of mTORC1 and GSK3-TIP60 Signaling in Regulation of Autophagosome Maturation and Lipid Metabolism[J]. *Mol Cell*. 2019 Feb 21;73(4):788-802.e7. IF: 14.248

Liu Y, Fu Y, Wang Q, et al. Proteomic profiling of HIV-1 infection of human CD4(+) T Cells identifies PSGL-1 as an HIV restriction factor[J]. Nat Microbiol. 2019 May;4(5):813-825. IF: 14.174

Yang L, Wang W H, Qiu W L, et al. A single-Cell transcriptomic analysis reveals precise pathways and regulatory mechanisms underlying hepatoblast differentiation[J]. *Hepatology*. 2017 Nov;66(5):1387-1401. IF: 13.246

Han X, Chen H, Huang D, et al. Mapping human pluripotent stem Cell differentiation pathways using high throughput single-Cell RNA-sequencing[J]. *Genome Biol.* 2018 Apr 5;19(1):47. **IF: 13.214**

Chen B, Zou W, Xu H, et al. Efficient labeling and imaging of protein-coding genes in living Cells using CRISPR-Tag[J]. Nature Communications. 2018 Nov 29;9(1):5065. IF: 12.353

Liu Z, Qin Q, Wu C, et al. Downregulated NDR1 protein kinase inhibits innate immune response by initiating an miR146a-STAT1 feedback loop[J]. *Nature Communications*. 2018 Jul 17:9(1):2789. IF: 12.353

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