

PURITY IN PRACTICE SAFETY IN SOLUTION

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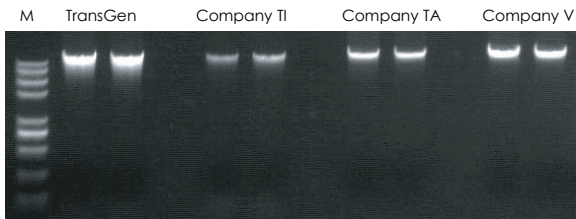
EasyPure[®] Universal Plant Genomic DNA Kit (EE112)

- Wide range of applications: applicable to various plant tissues, especially those rich in polysaccharides, polyphenols or starch.
- Fast operation: high-quality genomic DNA can be extracted in as little as 52 minutes.
- Safe and low toxicity: no toxic organic reagents such as phenol and chloroform.
- High purity: the unique technology can efficiently remove impurities such as pigments, polyphenols and polysaccharides in the sample, low inhibition of PCR amplification.

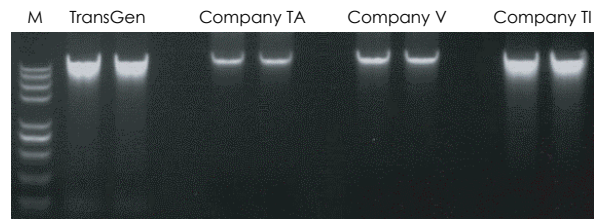
Data

Common plants

Tobacco leaves

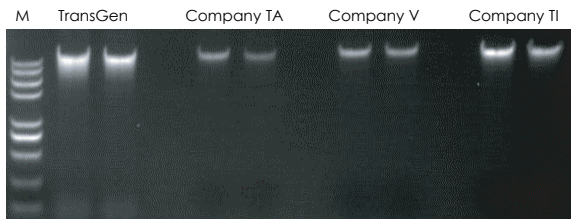


Corn leaves

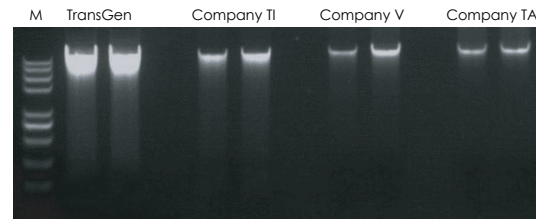


Polysaccharide-rich plants

Lychee leaves

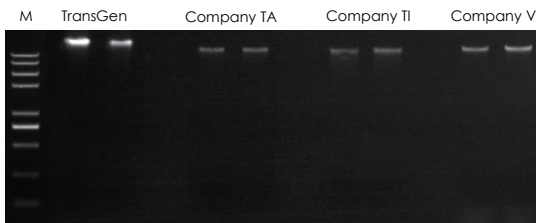


Sugarcane leaves

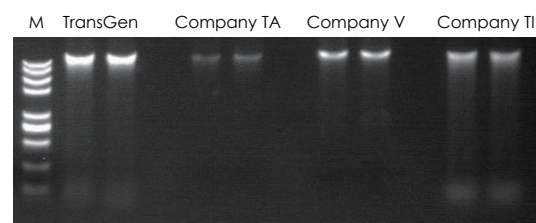


Polyphenol-rich plants

Pine needles



Corn roots

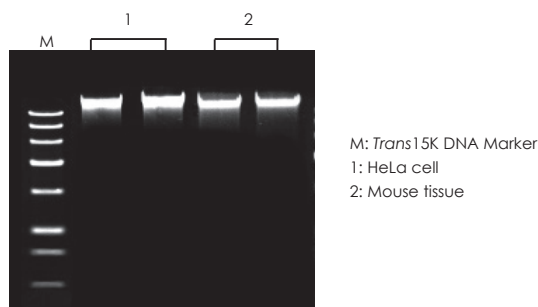


M: Trans2K[®] Plus II DNA Marker

EasyPure[®] Genomic DNA Kit (EE101)

- Suitable for the extraction of various types of materials (animal cells, animal tissues, mouse tails, E. coli, yeast).
- Fast operation, high yield (up to 15 µg).
- High integrity of cDNA ensured by mild lysis conditions without physical disruption, thus reducing damage to gDNA during cell lysis process.
- Safety: no toxic organic reagents such as phenol and chloroform.
- High purity enabled by spin column which can efficiently and specifically bind to DNA and remove protein, salts, lipids or other contaminants.

Data



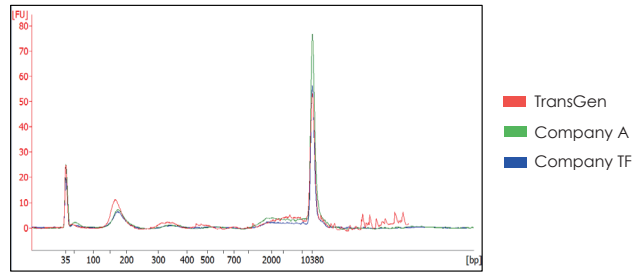
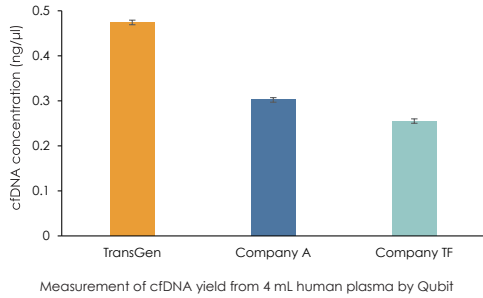
Tissue samples that have been successfully tested

Mouse	Testicle, epididymis, uterus, ovary, kidney, liver, lung, heart, brain, spleen, mesenteric lymph nodes, inguinal lymph nodes, skin, sciatic nerve, bone marrow, spinal cord, blood
Cynomolgus monkey	Testicle, epididymis, uterus, ovary, kidney, liver, lung, heart, brain, spleen, mesenteric lymph nodes, inguinal lymph nodes, skin, sciatic nerve, bone marrow, salivary gland, lacrimal gland

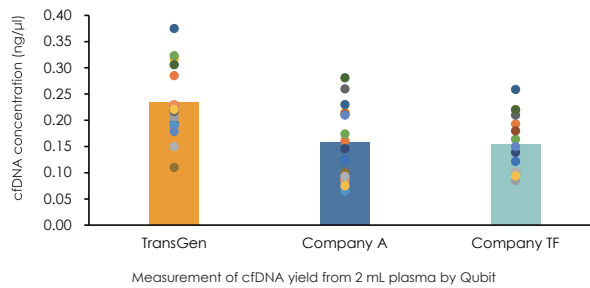
MagicPure[®] Cell-Free DNA Kit (EC201)

- Suitable for 0.5 mL-10 mL serum or plasma.
- Simple and fast, no centrifugation required.
- High yield and purity.

Data



Human plasma cfDNA was extracted using reagents from TransGen, Company A, and Company TF. The results showed that the cfDNA extracted using TransGen's product had a higher concentration, with the main fragment concentration being greater than that of competitors. Additionally, it demonstrated superior coverage for fragments in the 150-200 bp range compared to competing products.



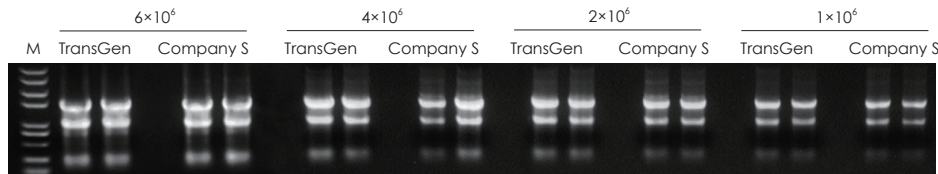
cfDNA was extracted from 16 human plasma samples using reagents from TransGen, Company A, and Company TF, respectively. The results showed that the TransGen product yielded higher cfDNA concentrations than competing products.

EasyPure[®] Fast Cell RNA Kit (ER111)

- Simple operation: operate at room temperature during the whole process, no low-temperature centrifugation required.
- Fast: extraction completes in 7 minutes.
- Safety: no need for organic reagents such as phenol and chloroform.
- High purity: efficient removal of impurity contamination.

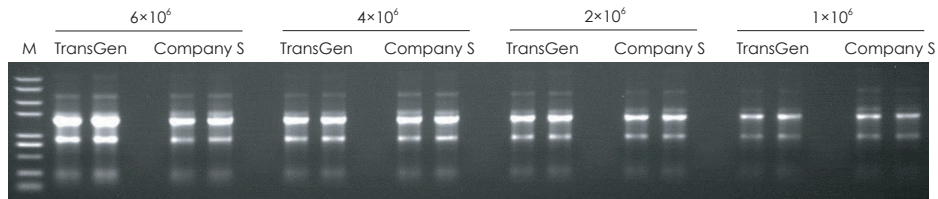
Data

Jurkat cells



M: Trans2K[®] Plus II DNA Marker

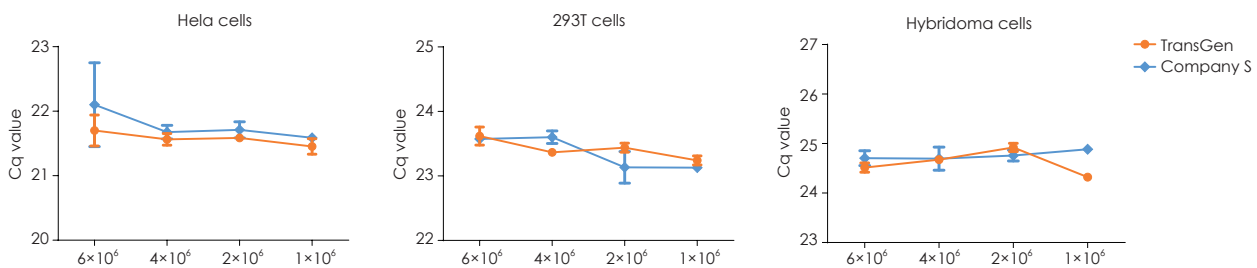
HeLa cells



M: Trans2K[®] Plus II DNA Marker

RNA was extracted from Jurkat cells and HeLa cells using reagents from TransGen and Company S respectively. The extracted RNA was analyzed by agarose gel electrophoresis.

Sample type	RNA yield (µg)			
	1×10 ⁶ cells	2×10 ⁶ cells	4×10 ⁶ cells	6×10 ⁶ cells
HeLa cells	4	7	10	13
293T cells	3	6	10	12
HEK293 cells	3	5	8	10
A549 cells	8	14	25	30
Jurkat cells	10	15	23	26
CHO cells	4	7	15	20
Hybridoma cells	7	15	20	30



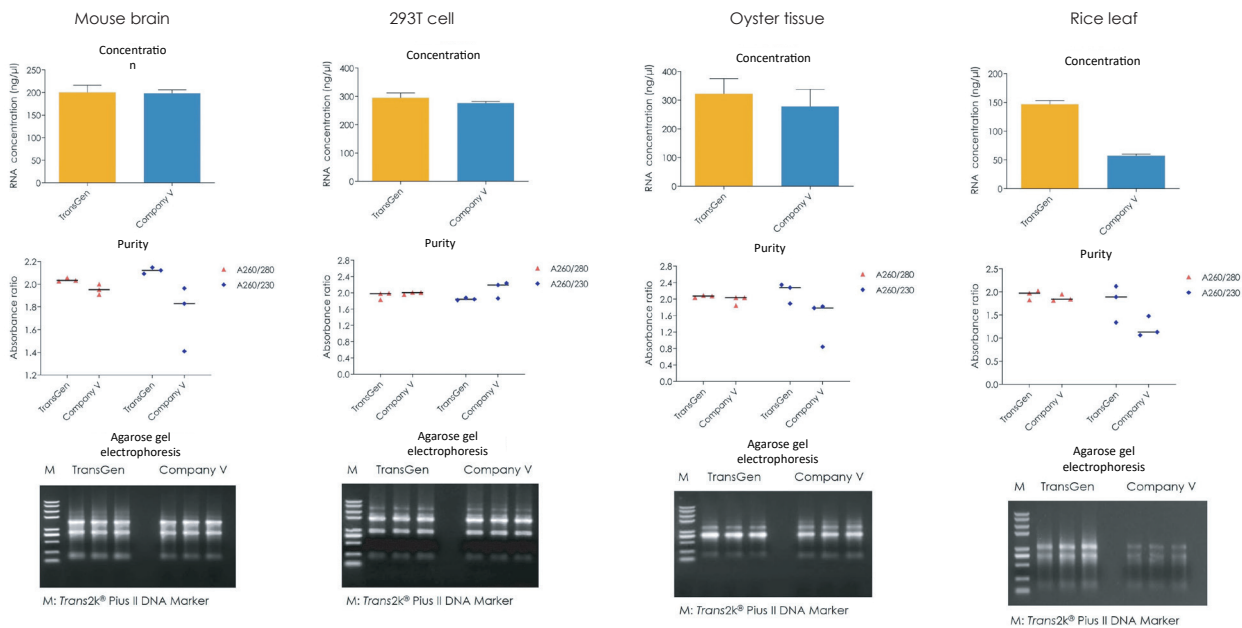
RNA was extracted from HeLa cells, 293T cells and Hybridoma cells using reagents from TransGen and Company S respectively. 200 ng of extracted RNA was used as a template for RT and qPCR. The results showed that extracted RNA by TransGen reagent had no inhibition for downstream qPCR.

TransZol Simple RNA Kit (ER511)

- Simple operation: Centrifugation at room temperature, no centrifugation-based phase separation with easier supernatant collection.
- High safety: Free from toxic reagents such as chloroform.
- Fast extraction speed: Completed in 8.5 minutes.
- High extraction purity: Effectively removes impurities such as gDNA and proteins.
- Wide application: Suitable for a wide range of samples, including animal cells/tissues, plant tissues, virus and bacteria samples.

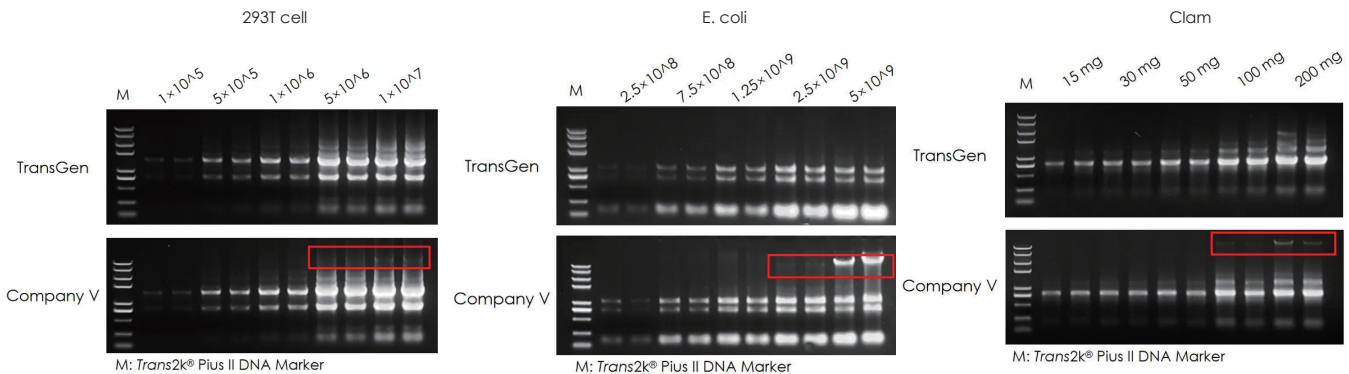
Data

Better extraction result



Extract RNA from various samples, including animal cells/tissues, plant tissues and microorganisms using TransGen (ER511) and Company V products and then test the RNA concentration, purity, run agarose gel electrophoresis. The results show that the RNA extracted with ER511 has higher quality and ER511 is compatible with a variety of sample types.

Wider range of sample size



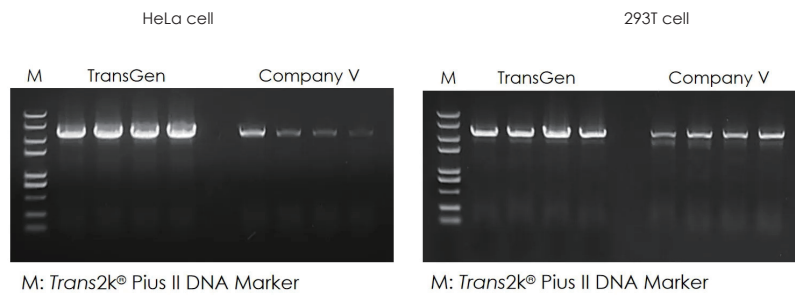
Extract RNA from 293T cells, Escherichia coli (E. coli), and clam (Solenidae) using TransGen (ER511) and Company V products and then the RNA quality is assessed by agarose gel electrophoresis. The results show that the RNA extracted with ER511 has higher quality and ER511 demonstrates good compatibility with different sample input amounts. Compared to Company V, ER511 shows no residual genomic DNA, ensuring the accuracy of downstream experimental results.

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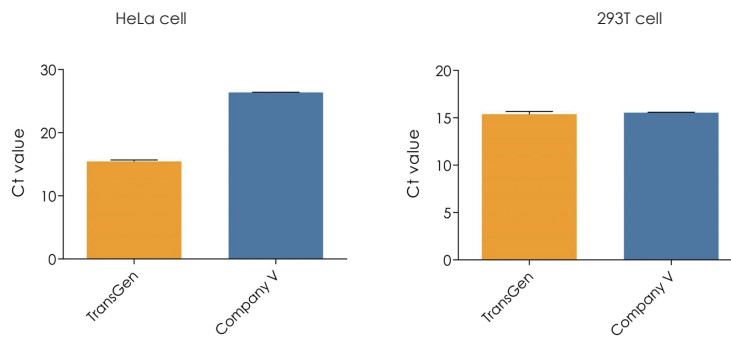
No inhibition in downstream experiments

1. PCR and qPCR experiments

PCR experiment

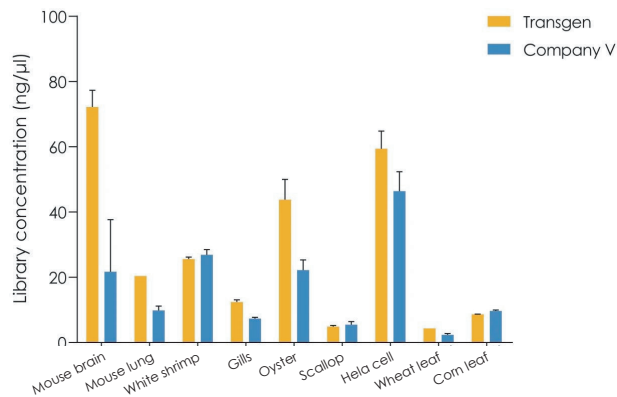


qPCR experiment



Extract RNA from HeLa and 293T cells using TransGen (ER511) and Company V products and the extracted RNA is used as template for PCR and qRT-PCR experiments. The results show that the RNA extracted with TransGen product (ER511) has higher quality and show no inhibition in both PCR and qRT-PCR reactions, while the RNA extracted with Company V exhibits stronger inhibition in downstream experiments.

2. Library preparation



Extract RNA from multiple species using TransGen (ER511) and Company V products and the extracted RNA is used as template for library preparation using an automated library preparation system with KP701. The results show that RNA extracted with TransGen product (ER511) successfully undergo RNA library preparation, yielding higher library output.

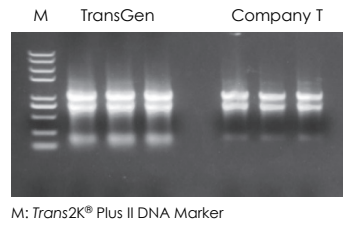
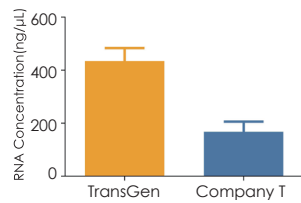
EasyPure[®] Universal Plant Total RNA Kit (ER302)

- Wide range of applications: applicable to various plant tissues, especially those rich in polysaccharides, polyphenols or starch.
- Fast operation: high-quality genomic RNA can be extracted in less than 30 minutes.
- Safe and low toxicity: no toxic organic reagents such as phenol and chloroform.
- High purity: the unique technology can efficiently remove impurities such as pigments, polyphenols and polysaccharides in the sample.

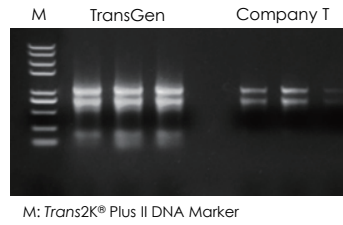
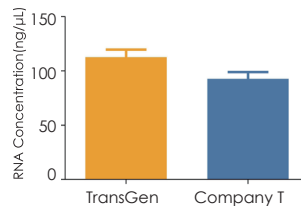
Data

Wide range of applications

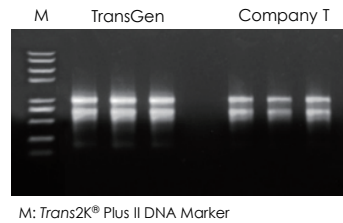
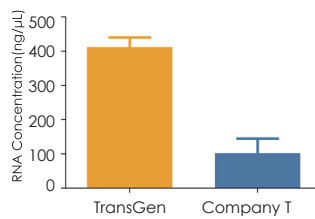
Tea tree leaves



Corn root



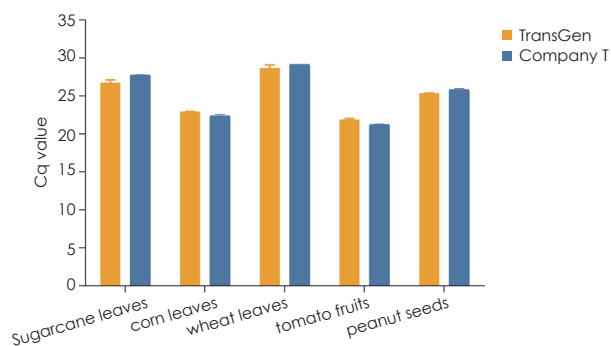
Soybean seeds



RNA was extracted from 100 mg of different plant samples using reagents from TransGen and Company T respectively. The concentration of extracted RNA was detected. And the extracted RNA was analyzed by agarose gel electrophoresis.

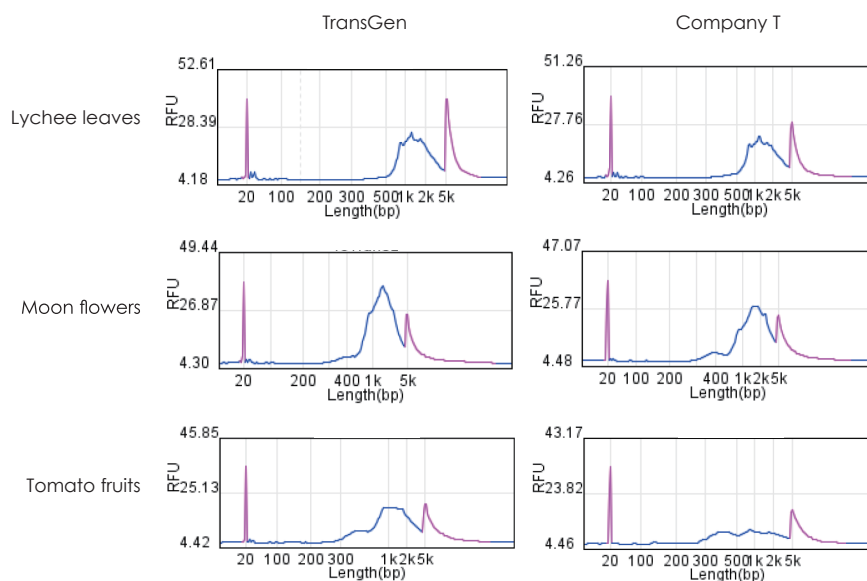
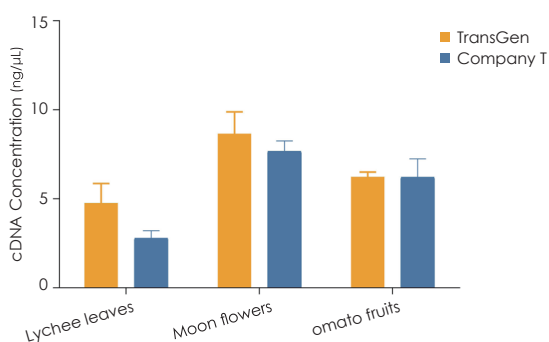
Suitable for a wide range of downstream experiments

qRT-PCR



RNA was extracted from different plant samples using reagents from TransGen and Company T respectively. The extracted RNA was used as a template for qRT-PCR (reagent: AQ311). The results showed that extracted RNA by TransGen reagent had no inhibition for downstream qPCR.

Full-length cDNA synthesis and amplification



RNA was extracted from different plant samples using reagents from TransGen and Company T respectively. The extracted RNA was used as a template for full-length cDNA synthesis (reagent: KC901).

GelStain (GS101)

- **Non-Toxic Safety:** GelStain's unique oily and large molecular structure prevents it from penetrating cell membranes. Ames test results confirm it is far less mutagenic than Ethidium Bromide (EB).
- **High Sensitivity:** Ideal for staining nucleic acid fragments of all sizes, with less impact on nucleic acid mobility than SYBR Green I.
- **High Stability:** Compatible with microwave or other heating methods for agarose gel preparation. Remains highly stable under both acidic and alkaline conditions at room temperature and exhibits strong light resistance.
- **Superior Signal-to-Noise Ratio:** Delivers intense sample fluorescence with very low background signal for clear, sharp images.
- **Simple & Convenient:** Like EB, the dye does not degrade during gel casting or electrophoresis. Post-electrophoresis staining requires only 30 minutes with no destaining or washing steps, allowing for direct visualization under a standard UV transilluminator.
- **Broad Compatibility:** Offers the flexibility of both pre-cast (in-gel) and post-electrophoresis (soaking) staining methods. Compatible with agarose and polyacrylamide gel electrophoresis. Suitable for staining dsDNA, ssDNA, and RNA.
- **No Equipment Changes Needed:** Compatible with standard EB or SYBR filter sets. Can be visualized using the same standard UV transilluminator as for EB, with optimal excitation around 300 nm.

GelStain Blue (GS102)

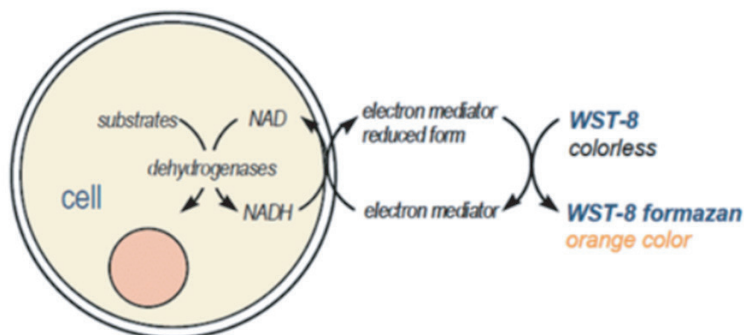
- **Non-Toxic:** Its unique, large oily molecular structure prevents it from penetrating cell membranes. Ames test results confirm significantly lower mutagenicity compared to Ethidium Bromide (EB). It is readily biodegradable and non-carcinogenic.
- **Exceptional Sensitivity:** Ideal for staining nucleic acid fragments of all sizes. It affects nucleic acid mobility less than SYBR Green I and exhibits no band-shifting, representing the highest sensitivity available in gel nucleic acid stains.
- **High Stability:** Compatible with microwave or other heating methods for agarose gel preparation. Remains highly stable under both acidic and alkaline conditions at room temperature and exhibits strong light resistance.
- **Superior Signal-to-Noise Ratio:** Delivers intense sample fluorescence with very low background signal for clear, sharp images.
- **Simple & Convenient:** Just like EB, it does not degrade during gel casting or electrophoresis. For post-electrophoresis staining, the process requires only 30 minutes with no need for a destaining step, and gels can be immediately visualized using a standard visible light gel transilluminator.
- **Broad Compatibility:** Offers the flexibility of both pre-cast (in-gel) and post-electrophoresis (soaking) staining methods. Compatible with agarose and polyacrylamide gel electrophoresis. Suitable for staining dsDNA, ssDNA, and RNA.
- **Visualization Method:** GelStain Blue is optimally excited at a wavelength near 474 nm, and observation with a blue light transilluminator is recommended. It can also be visualized using the standard UV transilluminator commonly used for EB.

TransDetect[®] Cell Counting Kit (CCK) (FC101)

Designed for cell proliferation assays as well as cytotoxicity assays by utilizing a water-soluble tetrazolium salt. The faster the cell proliferation, the lower cytotoxicity and the more cells, the darker the color. The color development has a good linear relationship with the number of cells.

- Fast detection and high sensitivity.
- Low cytotoxicity.
- Broad linear range.
- Good stability and high repeatability.

Schematic diagram of CCK

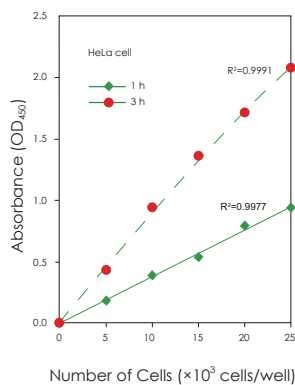


High Sensitivity

HeLa cells were seeded in a 96-well plate at densities of 0, 5,000, 10,000, 15,000, 20,000, and 25,000 cells per well and incubated at 37°C for 3 hours. After removing the cell culture supernatant, 110 µl of culture medium pre-mixed with CCK reagent (100 µl of DMEM with 10% FBS + 10 µl CCK reagent) was added to each well. The cells were then returned to the 37°C incubator for further culture. At 1 hour and 3 hours after adding CCK, the plate was placed in a microplate reader to measure the absorbance at a wavelength of 450 nm. A standard curve was plotted. The results showed a strong linear relationship between the absorbance and the cell number, with $R^2 > 0.99$.

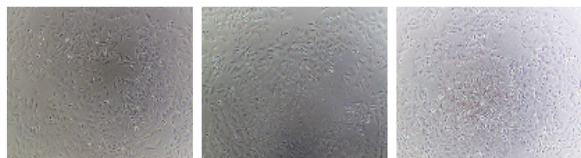
Data

CCK sensitivity test



Low Cytotoxicity

Based on the above experiment, after the 3-hour measurement, the culture medium in the wells seeded with 10,000 cells was replaced with fresh DMEM medium containing 10% FBS. Within 24 hours after adding CCK, the cell morphology in the culture wells was observed. As shown in the figure, the cells were in good condition, indicating low cytotoxicity of FC101 and that the cells remained viable for subsequent experiments after the assay.



3 hours incubation 6 hours incubation 24 hours incubation

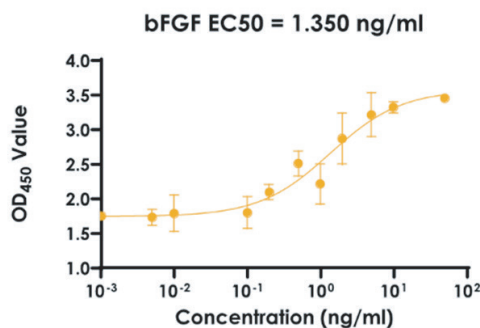
Good Stability

Experimental data demonstrate that the key performance parameters of this CCK product exhibited no significant alterations after three freeze-thaw cycles at -20°C, thereby meeting all predefined quality specifications.

Freeze-Thaw Cycles	R ² Value (Indicator of Linearity)	CV% @ 5000 cells (Indicator of Precision)
0 (Fresh)	0.995	4.5%
1 Cycle	0.993	5.1%
2 Cycles	0.990	6.2%
3 Cycles	0.985	7.8%

Application Example: bFGF Cytokine Activity Assay

NIH-3T3 cells were treated with varying concentrations of the growth factor bFGF (0.001, 0.005, 0.01, 0.1, 0.2, 0.5, 1, 2, 5, 10, 50 ng/ml) for 48 hours, followed by detection with the CCK reagent. The EC₅₀ of bFGF was determined to be 1.350 ng/ml.



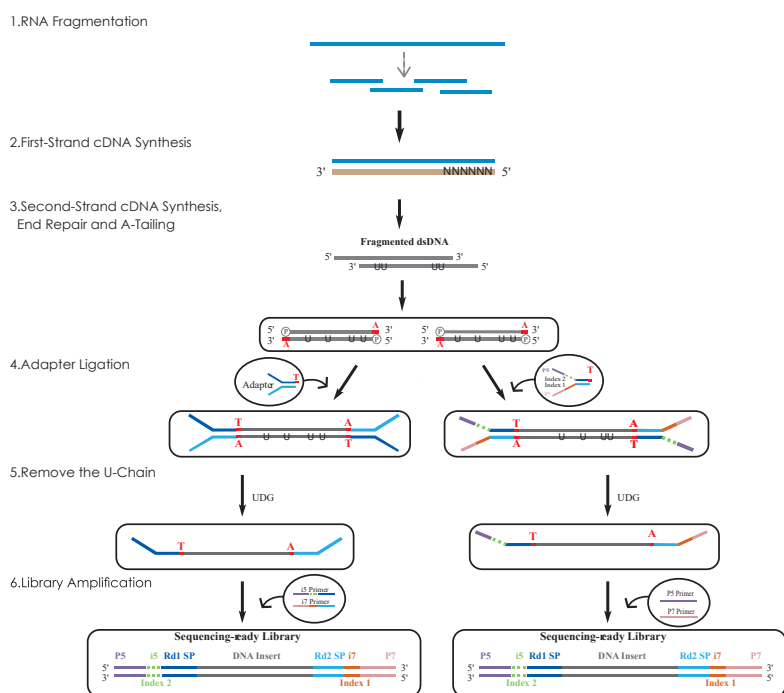
TransNGS[®] Fast RNA-Seq Library Prep Kit for Illumina[®] (KP701)

- High library yield
- High library conversion efficiency: Robust performance even with partially degraded RNA (RIN ≥ 3).
- Safe handling: Strand-specific reverse transcription components are actinomycin D-free, requiring no light protection. Low toxicity and odorless.
- Flexible fragmentation conditions: Achieve consistent results across varying fragmentation parameters.
- Broad species compatibility: Suitable for RNA isolated from animals, plants, and microorganisms.
- Versatile upstream compatibility: Compatible with multiple RNA preprocessing methods (e.g., mRNA capture or rRNA depletion).

Applications

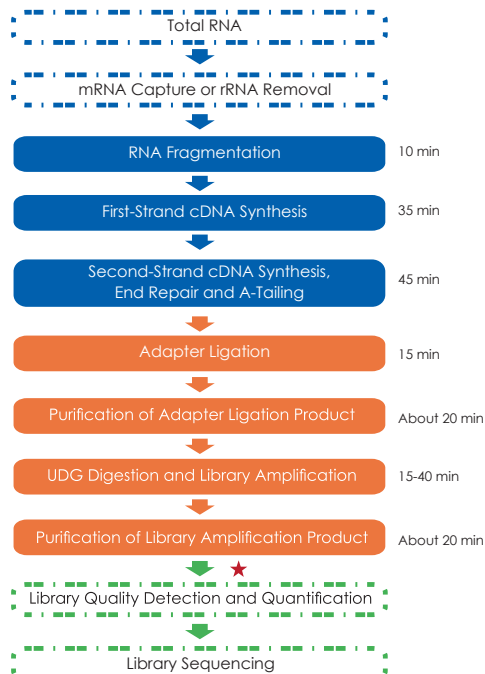
- Whole transcriptome sequencing
- Gene expression analysis
- Single nucleotide variation analysis
- Alternative Splicing Detection
- Fusion Gene Detection
- Analysis of non-coding RNA and RNA precursor

Schematic Diagram of Library Preparation



Principle chart of strand-specific library construction
(No dUTP incorporation and UDG digestion steps in second-strand synthesis of non-strand-specific library)

Workflow Diagram



[-] The dotted lines are not included in this kit flow

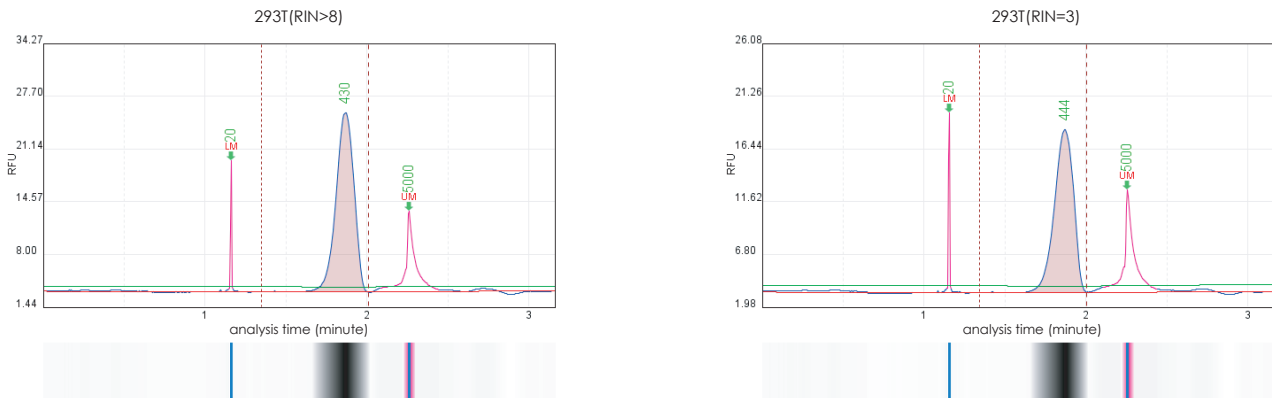
★ Steps where fragments size selection can be performed

Flow chart of strand-specific library construction
(No dUTP incorporation and UDG digestion steps in second-strand synthesis of non-strand-specific library)

Data

Library Peak Profile

Using 293T RNA samples with varying RIN values (high vs. low integrity), we perform mRNA capture and library construction. TransGen libraries exhibit normal peak profiles (no aberrant size distribution).



Using rRNA-depleted *Staphylococcus aureus* RNA, we construct both strand-specific and non-strand-specific libraries with TransGen's kits. The sequencing data shows high quality.

Library Prep Solution	Clean data (%)	rRNA (%)	Q30 (%)	GC (%)	Total mapped (%)	Duplicate (%)	Multiple mapped (%)	Exon (%)	Gene
Strand-specific Library	91.95	0.86	95.15	34.7	94.95	27.59	0.24	92.04	2639
Non-strand-specific Library	85.23	0.90	94.99	34.85	94.62	23.91	0.24	91.15	2657

We construct both strand-specific and non-strand-specific libraries with TransGen's technology using mouse fecal microbial RNA post rRNA depletion. The results show highly concordant taxonomic profiles between the two library types.

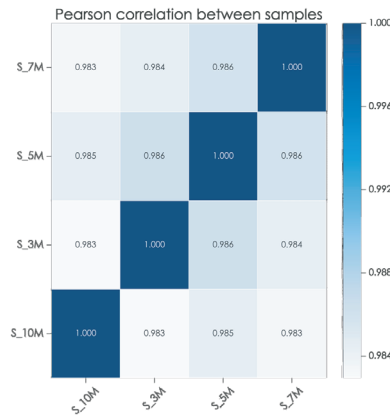
Taxonomic Annotation of rRNA-Depleted Mouse Fecal RNA Libraries

Classification	Non-strand-specific Library	Strand-specific Library
s__Ligilactobacillus_murinus	0.14395	0.1334
s__Duncaniella_dubosii	0.03768	0.03752
s__Bacteroides_sp._KGM10229	0.03804	0.037
s__Staphylococcus_aureus	0.0167	0.03449
s__Bacteroides_uniformis	0.02367	0.02409
s__Muribaculum_gordoncarteri	0.02314	0.02315
s__Pusillimonas_faecalis	0.01763	0.01746
s__Muribaculum_intestinale	0.01649	0.01675
s__Roseburia_intestinalis	0.01675	0.01551
s__Duncaniella_sp._B8	0.01537	0.01623
s__Flavonifractor_plautii	0.01546	0.01574
s__Murine_astrovirus	0.01527	0.01521
s__Pseudomonas_aeruginosa	0.01508	0.01537
s__Vescimonas_coprocola	0.01421	0.01376
s__Plasmodium_knowlesi	0.01494	0.01265
s__Dysosmobacter_sp._Marseille-Q4140	0.01306	0.01333
s__Oscillibacter_sp._NSJ-62	0.01314	0.01324
s__Lachnoclostridium_sp._YL32	0.01244	0.01275
s__Roseburia_hominis	0.01023	0.0094
s__Acutalibacter_muris	0.00956	0.00964

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Compatible with Variable Fragmentation Durations

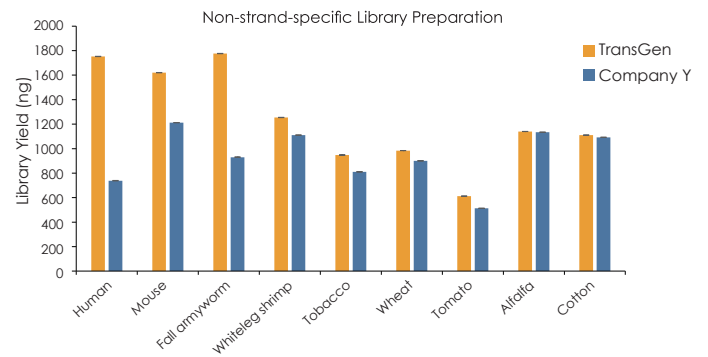
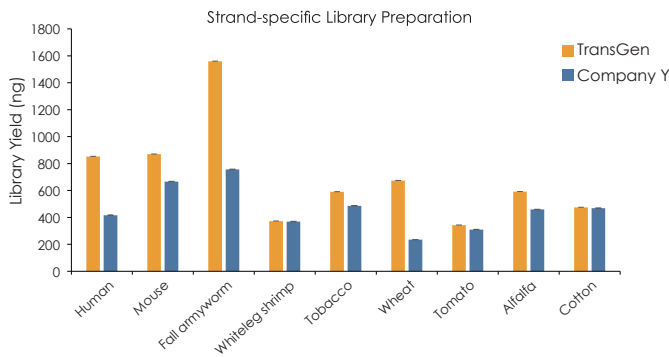
Using 500 ng of 293T total RNA, we test fragmentation time of 3, 5, 7, and 10 minutes, followed by mRNA capture and library construction. Correlation analysis demonstrate Pearson correlation >98% between groups.



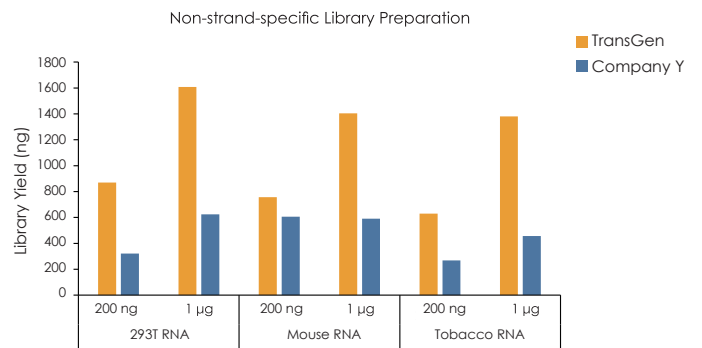
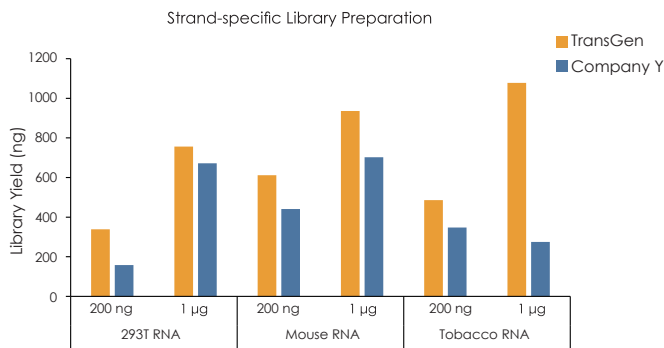
Comparison with Competitors

Library Yield

Using high-quality RNA samples (RIN > 8) from nine different species, we perform mRNA capture and subsequent library preparation. The results demonstrate that TransGen's strand-specific and non-strand-specific library preparation kits consistently generated higher library yields compared to those of Company Y.



Using severely degraded RNA (RIN = 3) from human (293T cells), mouse, and tobacco, we perform mRNA capture followed by library construction with high (1 µg) and low (200 ng) input amounts. The results demonstrate that the TransGen's kit (both strand-specific and non-strand-specific) consistently outperforms Company Y in library yield across all species and input amounts.



Sequencing Data Quality

Comparative analysis of sequencing data across multiple species reveal that RNA libraries using TransGen kit significantly outperform Company Y's in percentage of clean data, number of reliably detected genes.

Strand-specific Library Preparation (after mRNA capture)

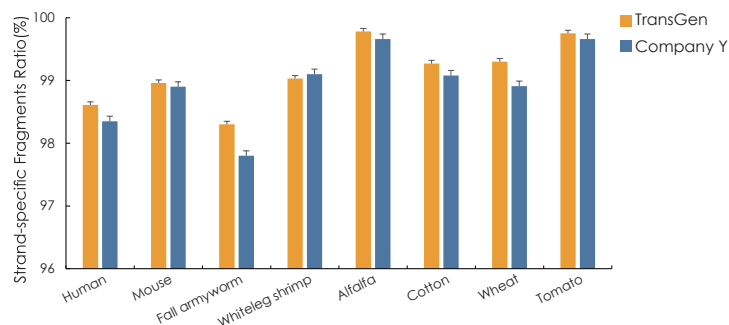
Sample (500 ng total RNA)	Library Prep Solution	Clean data (%)	rRNA (%)	Q30 (%)	GC (%)	Total mapped (%)	Duplicate (%)	Multiple mapped (%)	Exon (%)	Gene
Human	TransGen	91.7	0.21	95.29	48.66	98.24	23.64	7.17	85.88	15421
	Company Y	82.22	0.47	95.7	49.46	98.34	30.56	7.76	84.70	14410
Mouse	TransGen	93.10	0.55	95.73	48.76	97.47	19.62	7.60	89.83	12817
	Company Y	83.19	0.31	96.03	49.48	97.83	20.64	7.73	88.13	12706
Fall armyworm	TransGen	87.52	0.85	94.06	45.29	86.26	24.47	4.15	82.64	7607
	Company Y	84.88	0.58	94.6	47.55	84.81	32.95	4.05	83.67	7503
Whiteleg shrimp	TransGen	83.65	2.76	94.24	47.98	90.77	24.15	14.61	80.25	11858
	Company Y	82.76	3.30	94.65	49.52	90.67	27.88	15.92	78.69	11405
Alfalfa	TransGen	84.66	0.25	95.98	43.65	61.49	28.77	3.17	69.46	15387
	Company Y	81.98	0.21	94.37	43.19	59.25	28.42	2.92	68.50	15231
Cotton	TransGen	87.94	0.98	94.65	43.88	98.23	20.30	4.84	90.65	45809
	Company Y	85.63	0.41	94.63	43.77	98.35	26.50	5.05	88.37	44705
Wheat	TransGen	85.62	1.31	95.58	52.3	95.39	21.53	5.66	87.22	46235
	Company Y	56.42	1.40	94.78	57.74	94.85	28.15	6.66	87.24	47038
Tomato	TransGen	85.6	1.33	93.48	44.34	97.75	25.32	1.56	94.56	17704
	Company Y	83.8	1.34	94.68	44.16	97.76	35.41	1.67	93.71	17563

Non-strand-specific Library Preparation (after mRNA capture)

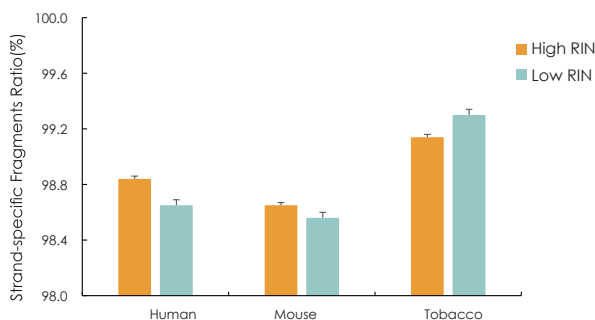
Sample (500 ng total RNA)	Library Prep Solution	Clean data (%)	rRNA (%)	Q30 (%)	GC (%)	Total mapped (%)	Duplicate (%)	Multiple mapped (%)	Exon (%)	Gene
Human	TransGen	93.20	0.71	95.88	47.11	98.35	20.79	7.52	87.15	15037
	Company Y	91.97	0.80	95.71	48.55	98.00	29.62	7.93	85.58	14426
Mouse	TransGen	93.35	0.59	95.79	48.39	96.13	23.79	7.77	93.41	12777
	Company Y	91.24	0.47	95.84	49.35	97.36	25.35	8.14	93.56	12530
Fall armyworm	TransGen	87.22	1.07	93.82	44.88	81.45	22.89	3.39	82.12	7771
	Company Y	83.97	0.52	94.5	45.4	82.45	32.45	3.96	80.11	7668
Whiteleg shrimp	TransGen	83.89	2.61	94.13	46.83	90.2	23.95	19.42	82.17	11783
	Company Y	80.49	3.55	94.83	48.22	90.84	26.69	17.77	83.90	11474
Alfalfa	TransGen	86.11	0.28	93.49	42.3	59.06	21.45	4.33	69.21	15535
	Company Y	80.77	0.22	94.71	42.5	58.21	24.83	4.78	68.35	15639
Cotton	TransGen	85.46	1.07	93.89	43.00	96.91	20.76	4.99	89.40	45096
	Company Y	80.90	1.26	94.6	43.23	98.23	26.93	5.39	87.46	44908
Wheat	TransGen	86.11	1.17	93.98	52.08	87.81	19.59	5.37	84.61	46979
	Company Y	85.39	0.82	94.94	54.73	95.22	30.35	6.44	85.88	47522
Tomato	TransGen	82.82	0.98	93.82	43.73	91.67	27.27	1.58	94.91	17639
	Company Y	79.06	0.96	94.52	43.55	97.64	34.09	1.56	93.73	17412

Higher strand-specific Fragments Ratio

Using high-integrity RNA samples (RIN > 8) from eight diverse species, we perform mRNA capture and library preparation. The results demonstrate that TransGen's libraries consistently achieve higher strand-specific fragments ratio compared to Company Y's across all species tested.



Using high-quality (RIN > 6) and degraded (RIN = 3) RNA samples from human, mouse, and tobacco, we perform mRNA capture and library preparation. TransGen strand-specific libraries maintain high strand specificity regardless of RNA degradation level.



Category	Product Name	Cat.No.
DNA Extraction	EasyPure® Universal Plant Genomic DNA Kit	EE112
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	MagicPure® Cell-Free DNA Kit	EC201
RNA Extraction	EasyPure® Fast Cell RNA Kit	ER111
	TransZol Up Simple RNA Kit	ER511
	EasyPure® Universal Plant Total RNA Kit	ER302
Nucleic Acid Gel Stain	GelStain	GS101
	GelStain Blue	GS102
CCK	TransDefect® Cell Counting Kit (CCK)	FC101
RNA Library Prep	TransNGS® Fast RNA-Seq Library Prep Kit for Illumina®	KP701

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Cat#	Description	Cond.
NB-45-00042-100	Super NI-NTA Agarose Resin	100ml
NB-45-00042-25	Super NI-NTA Agarose Resin	25ml
NB-45-00058-4	Proteus 1 -step Batch Mini Spin Column Pack	40pc
NB-12-6001-3	NeoLine pipette 2-20 µl	1unit
NB-12-0023C	Mini Centrifuge N500C @10,000rpm (Including 6x1.5/2.0ml angle rotor)	1pcs
NB-03-0160	Proteinase K (Powder)	100mg
NB-60-0001	NeoPrep mini	50columns
NB-12-8001-19	Combs for NeoPRO4 mini (1.5mm, 15 wells)	5pieces
NB-12-8001-20	Spacer glasses flat for NeoPRO4 mini (0.75mm, 100*83mm)	5pieces
NB-12-8001-04	Short glasses flat for NeoPRO4 mini (1.0mm, 100*73mm)	10pieces

“Remember, if you ever get lost or have questions, don’t hesitate to reach out to our customer support team. They’re here to help you address any concerns you may have.”

clinisciences

Austria

Company: CliniSciences GmbH
Address: Sternwartestrasse 76, A-1180
Wien - Austria
Telephone: +43 720 115 580
Fax: +43 720 115 577
Email: oesterreich@clinisciences.com
Web: <https://www.clinisciences.com>



Belgium

Company: CliniSciences S.R.L
Address: Avenue Stalingrad 52, 1000
Brussels - Belgium
Telephone: +32 2 31 50 800
Fax: +32 2 31 50 801
Email: belgium@clinisciences.com
Web: <https://www.clinisciences.com>



Denmark

Company: CliniSciences ApS
Address: Oesterbrogade 226, st. 1,
Copenhagen, 2100 - Denmark
Telephone: +45 89 888 349
Fax: +45 89 884 064
Email: denmark@clinisciences.com
Web: <https://www.clinisciences.com>



Finland

Company: CliniSciences ApS
Address: Oesterbrogade 226, st. 1,
Copenhagen, 2100 - Denmark
Telephone: +45 89 888 349
Fax: +45 89 884 064
Email: suomi@clinisciences.com
Web: <https://www.clinisciences.com>



France

Company: CliniSciences S.A.S
Address: 74 Rue des Suisses, 92000
Nanterre- France
Telephone: +33 9 77 40 09 09
Fax: +33 9 77 40 10 11
Email: info@clinisciences.com
Web: <https://www.clinisciences.com>



Germany

Company: Clinisciences GmbH
Address: Wilhelm-Mausier-Str. 41-43,
50827 Köln - Germany
Telephone: +49 221 9498 320
Fax: +49 221 9498 325
Email: deutschland@clinisciences.com
Web: <https://www.clinisciences.com>



Iceland

Company: CliniSciences ApS
Address: Oesterbrogade 226, st. 1,
Copenhagen, 2100 - Denmark
Telephone: +45 89 888 349
Fax: +45 89 884 064
Email: island@clinisciences.com
Web: <https://www.clinisciences.com>



Ireland

Company: CliniSciences Limited
Address: Ground Floor, 71 lower Baggot street
Dublin D02 P593 - Ireland
Telephone: +353 1 6971 146
Fax: +353 1 6971 147
Email: ireland@clinisciences.com
Web: <https://www.clinisciences.com>



Italy

Company: CliniSciences S.r.l
Address: Via Maremmana inferiore 378
Roma 00012 Guidonia Montecelio - Italy
Telephone: +39 06 94 80 56 71
Fax: +39 06 94 80 00 21
Email: italia@clinisciences.com
Web: <https://www.clinisciences.com>



Netherlands

Company: CliniSciences B.V.
Address: Kraaijenhoffstraat 137A,
1018RG Amsterdam, - Netherlands
Telephone: +31 85 2082 351
Fax: +31 85 2082 353
Email: nederland@clinisciences.com
Web: <https://www.clinisciences.com>



Norway

Company: CliniSciences AS
Address: c/o MerVerdi Munkerudtunet 10
1164 Oslo - Norway
Telephone: +47 21 988 882
Email: norge@clinisciences.com
Web: <https://www.clinisciences.com>



Poland

Company: CliniSciences sp.Z.o.o.
Address: ul. Rotmistrza Witolda Pileckiego 67
lok. 200 - 02-781 Warszawa -Poland
Telephone: +48 22 307 0535
Fax: +48 22 307 0532
Email: polska@clinisciences.com
Web: <https://www.clinisciences.com>



Portugal

Company: Quimigen Unipessoal LDA
Address: Rua Almada Negreiros, Lote 5, Loja 14,
2615-275 Alverca Do Ribatejo - Portugal
Telephone: +351 30 8808 050
Fax: +351 30 8808 052
Email: info@quimigen.com
Web: <https://www.quimigen.pt>



Spain

Company: CliniSciences Lab Solutions
Address: C/ Hermanos del Moral 13
(Bajo E), 28019, Madrid - Spain
Telephone: +34 916 750 700
Fax: +34 91 269 40 74
Email: espana@clinisciences.com
Web: <https://www.clinisciences.com>



Sweden

Company: CliniSciences ApS
Address: Oesterbrogade 226, st. 1,
Copenhagen, 2100 - Denmark
Telephone: +45 89 888 349
Fax: +45 89 884 064
Email: sverige@clinisciences.com
Web: <https://www.clinisciences.com>



Switzerland

Company: CliniSciences AG
Address: Fracht Ost Flughafen Kloten
CH-8058 Zürich - Switzerland
Telephone: +41 (044) 805 76 81
Fax: +41 (044) 805 76 75
Email: switzerland@clinisciences.com
Web: <https://www.clinisciences.com>



UK

Company: CliniSciences Limited
Address: 11 Progress Business center, Whittle
Parkway, SL1 6DQ Slough- United Kingdom
Telephone: +44 (0)1753 866 511
or +44 (0) 330 684 0982
Fax: +44 (0)1753 208 899
Email: uk@clinisciences.com
Web: <https://www.clinisciences.com>



USA

Company: CliniSciences LLC
Address: c/o Carr Riggs Ingram,
500 Grand Boulevard, Suite 210 Miramar
Beach, FL 32550- USA
Telephone: +1 850 650 7790
Fax: +1 850 650 4383
Email: usa@clinisciences.com
Web: <https://www.clinisciences.com>

