Conventional PCR Genotyping – Accustart[™] II Genotyping Kit

Completely reagent-based system enables reliable PCR genotyping with minimal pipetting skill.



4.1 Two mouse tail snips (2.5 mm) were extracted according to the recommended conditions for each kit. The volume of each extract was brought to 300 µl and diluted 1/20 with TE buffer. 5 µl of diluted extract was used in a 25 µl PCR reactions.

SNP Genotyping (Real-Time qPCR) – AccuStart Genotyping ToughMix

Genotyping ToughMix enables probe-based genetic analysis directly from crude extracts, DBS punches, plant tissue, and clinical specimens.





Influence of PCR inhibitor

4.1 Comparison to conventional master mixes AccuStart Genotyping ToughMix stands up to the challenge where other genotyping master mixes fall apart. ToughMix can be used with clean templates where it generates higher fluorescent signal and tighter clusters than the leading competitors.

4.2 Comparison to conventional master mixes. In the presence of a common PCR inhibitor, humic acid (50 ng/µl), the competitors system is completely shut down while ToughMix delivers robust, accurate results.

microRNA Expression Profiling:

Highly regarded 3-component system enables detection and guantification of miRNA and other small non-coding RNA with superior sensitivity.



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Featured Product Guide



cDNA synthesis kits: qScript cDNA SuperMix / XLT cDNA SuperMix

Single-tube Supermix minimizes pipetting steps, simplifies reaction assembly, and improves overall accuracy. XLT Supermix delivers superior sensitivity and yield.

FEATURES AND BENEFITS:

- 5x concentrated SuperMix maximizes input volume with dilute samples of total RNA
- Superior results across a wide dynamic range of input RNA (10⁸ fold)
- Pre-blended with ribonuclease inhibitor protein (RIP) and an optimized blend of random hexamer and oligo(dT) primers to ensure accurate representation of 5' and 3' sequences
- Stable through 20 freeze/thaw cycles



qPCR

PerfeCra qPCR reagents combine a stringent, ultrapure antibody hotstart with performance engineered DNA polymerase in stabilized 1-tube formulations optimized for the specific performance needs of real-time quantitative PCR.

PerfeCта qPCR ToughMix®

Advanced single-component reagent technology ensures reliable and sensitive qPCR detection even in the presence of common PCR inhibitors.

PerfeCтa MultiPlex qPCR ToughMix

Advanced 1-tube Supermixes supporting robust multiplexing while withstanding a broad spectrum of PCR inhibitors.

PerfeCта FastMix II

Convenient 1-tube reagent solution supports robust and reliable probe-based DNA detection with fast or standard thermal cycling conditions.

PerfeCта SYBR Green FastMix / SuperMix

Sensitive and precise DNA amplification using SYBR® Green detection chemistry.

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(right) that introduce numerous opportunities for error.

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Tough-tested Toughmix reagents deliver superior performance in the presence of PCR inhibitors

Inhibitor	Common sources	Reagent performance	
		Competitor	PerfeC⊤a ToughMix
Polyphenols	Plant extracts	_	1
Humic acids	Soil Plant tissues	-	1
Hematin	Dried bloods Blood spots	_	1
Hemoglobin	Blood	1	1
Polysaccharides	Feces Plant tissues	-	1
Melanin	Hair Skin	_	1

One-Step RT-qPCR and RT-P(

One-Step RT-qPCR and RT-PCR

1-step kits incorporate both the reverse transcription and PCR steps in a convenient single reaction tube.

	Conventional One-Step RT-PCR	Tough-Tested MultiPlex One-Step RT-qPCR	Tough-Tested One-Step Probe-based RT-qPCR	One-Step SYBR-based RT-qPCR
Kit	qScript XLT One-Step RT-PCR Kit	UltraPlex One-Step ToughMix	qScript XLT One-Step RT- qPCR ToughMix	qScript One-Step SYBR Green RT-qRT PCR
Detection Chemistry	N/A	Hydrolysis Probes	Hydrolysis Probes	SYBR Green I dye
Sensitivity	+++	++++	+++	++
Multiplex Compatibility	N/A	>4	<4	No
Reagent Components	2	1	1	2
RNA Input (Linear Range)	1 pg – 1 µg	-	-	-
Amplicon Length	4+ kb	<1 kb	<1 kb	<1 kb



31 One-Step SYBR Green RT-qPCR with broad dynamic range, high sensitivity and high specificity. A 202 bp fragment of glyceraldehyde-3-phosphate dehydrogenase (GAPD) mRNA was amplified from log-fold serial dilutions of HeLa cell total RNA (100 ng to 0.1 pg). Eight replicate reactions for each RNA quantity, and the no template control (NTC) were carried out in 25 µl volumes with the qScript One-Step SYBR Green RT-qPCR Kit and 200 nM each GAPD specific primers (PrimerBank ID 7669492a2, Wang, X. and Seed, **B** (2003) NAR 31(24): e154; pp.1-8). Reactions were assembled on ice, transferred to a MyiQ[™] real-time detection system (Bio-Rad Laboratories), and incubated for 5 min at 50°C followed by 2 min at 95°C. PCR cycling was for 40 cycles of 3 s, 95°C; 30 s, 60°C. Immediately following PCR cycling the block temperature was ramped from 60°C to 95°C and melt curve data was collected. Panel A) Amplification plots and standard curve regression analysis. Panel B) Dissociation results (melt curve) for NTC, 0.1 pg and 1 pg reactions.

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