PureQuant™ CD3+ T Cell Assay

Catalog Number A47197

Pub. No. MAN0019205 Rev. C.0



WARNING! Read the Safety Data Sheets (SDSs) and follow the handling instructions. Wear appropriate protective eyewear, clothing, and gloves. Safety Data Sheets (SDSs) are available from thermofisher.com/support.

Product description

The Applied Biosystems[™] PureQuant[™] CD3+ T Cell Assay is a TaqMan[™] qPCR-based assay that measures genomic DNA demethylation at unique sites. This assay is used to measure the percentage of CD3+ T cells in a heterogenous population of cells such as peripheral blood mononuclear cells (PBMC) or a cell therapy product. Genomic DNA isolated from cells is first subjected to bisulfite conversion, followed by qPCR using methylation specific primers. The assay utilizes Standards for estimating copy number. Calibrator and Reference serve as controls. Each 96-well plate (0.1 mL block) can accommodate from 1 to 7 test samples. Unlike other PureQuant[™] assays available, the PureQuant[™] CD3+ T Cell Assay uses PureQuant[™] qPCR Master Mix II rather than PureQuant[™] qPCR Master Mix.

Contents and storage

Table 1 PureQuant™ CD3+ T Cell Assay, (Cat. No. A47197)[1]

Contents	Amount	Storage
Dynabeads™ SILANE Genomic DNA Kit	1 Box	4°C
Lysis Binding Buffer		
 Dynabeads[™] magnetic beads 		
Wash Buffer 1		
Wash Buffer 2		
Elution Buffer		
PureQuant™ qPCR Master Mix II	1 × 5 mL	-20°C
PureQuant™ qPCR Bisulfite Conversion Reagents	1 Box	RT
 PureQuant™ Ammonium Bisulfite 	4 × 2.5 mL	
PureQuant™ THFA	1 × 3.5 mL	
PureQuant™ CD3+ T Cell Module	1 Box	-20°C
 PureQuant™ High Copy Standard B 	1 mL	
 PureQuant™ Reference Genomic DNA 	6 µg	
 PureQuant™ Calibrator B 	350 μL	
Lambda DNA	1.7 mL	
 TaqMan™ GAPDH Assay 	200 μL	
TaqMan™ CD3 Assay	200 µL	

PureQuant[™] CD3+ T Cell Assay is only available as a complete kit. Individual components are not sold separately.



Required material not supplied

Unless otherwise indicated, all materials are available through **thermofisher.com**. "MLS" indicates that the material is available from **fisherscientific.com** or another major laboratory supplier.

Catalog numbers that appear as links open the web pages for those products.

Item	Source		
Reagents			
PureLink™ Genomic DNA Mini Kit	K1820-01		
2-Propanol	MLS		
Ethanol	MLS		
TE, pH 8.0	MLS		
Instruments and equipment			
HulaMixer™ Sample Mixer	15920D		
DynaMag [™] -2 Magnet	12321D		
Eppendorf™ ThermoMixer™ C and SmartBlock™ 2 mL	Eppendorf™ 5382000023 and 5362000035		
Eppendorf™ Safe-Lock™ Tubes, 2.0 mL	Eppendorf™ 022363344		
Real Time-PCR Machine	MLS		

IMPORTANT! The performance of this assay has been validated using a Eppendorf ThermoMixer C temperature control device. A dry heat block can be used as an alternative. However, use of the Eppendorf ThermoMixer C is recommended for most consistent results.

Assay overview



Guidelines to isolate genomic DNA and prepare samples

- Use the 2 mL Safe-Lock[™] tubes for sample preparation.
- Use the Elution Buffer supplied with the Dynabeads[™] SILANE Genomic DNA Kit.
- Do not add water into the Eppendorf[™] ThermoMixer[™] C with SmartBlock[™] wells or heat block.

Isolate genomic DNA and prepare sample

Isolate genomic DNA from 1-2 × 10⁶ cells using PureLink[™]
Genomic DNA Mini Kit, then check the purity of gDNA using a spectrophotometer or NanoDrop[™] spectrophotometer.

The $\mbox{OD}_{260/280}$ and $\mbox{OD}_{260/230}$ of your gDNA should fall within the specified ranges:

(OD_{260/280}:1.7-2.0 and OD_{260/230}:1.5-2.4)

2. Prepare Sample, Calibrator, and Reference in 2 mL tubes according to the following table:

Item	Amount/volume of DNA	Make up the volume w/ Elution Buffer to		
Sample	400-1200 ng	142 µL		
PureQuant™ Calibrator B	75 μL	142 µL		
PureQuant™ Reference Genomic DNA ^[1]	1000–1200 ng	142 µL		

^[1] Take 1 µL PureQuant™ Reference Genomic DNA and estimate the concentration using TE (pH 8.0) as blank. Expected concentration range between 150–250 ng/µL).

Incubate at 56°C for 5 min with gentle shaking (900 rpm) using an Eppendorf[™] ThermoMixer[™] C with 2 mL SmartBlock[™] (or dry bath with 2 mL block).

- 4. Briefly spin down the samples.
- Adjust the Eppendorf[™] ThermoMixer [™] C (or dry heat block) temperature to 80°C for the bisulfite conversion.

Guidelines for bisulfite conversion

- During bisulfite conversion, follow the recommended incubation times as over-incubation or under-incubation will impact the assay result.
- Bring Lysis Binding Buffer and Dynabeads[™] magnetic beads to room temperature before use.
- Make a homogeneous suspension of Dynabeads[™] magnetic beads before use.

Perform bisulfite conversion

1. Add 270 μ L ammonium bisulfite and 90 μ L THFA to the Samples, Calibrator, and Reference.

Vortex thoroughly and briefly spin down the samples.

2. Eppendorf™ ThermoMixer™ C: Incubate at 80°C for 45 minutes with gentle shaking (900 rpm). Briefly centrifuge to spin down the samples and allow them to cool down to room temperature before proceeding to the next step.

OR

Heat block: Incubate at 80°C for 45 minutes with intermittent vortexing for 1–2 seconds at 4.5-minute intervals. Minimize any delay that may occur during vortexing. Briefly spin down the samples at the end of the 45 minute incubation. Allow samples to cool down to room temperature before proceeding to the next step.

- 3. Add 870 µL Lysis Binding Buffer and 105 µL Dynabeads[™] magnetic beads from the (Dynabeads[™] SILANE Genomic DNA Kit) to each reaction.
- 4. Mix thoroughly by vortexing and briefly spin down the tubes.
- 5. Add 570 µL of 2-propanol and vortex thoroughly.
- 6. Incubate at room temperature for 7 minutes on a rotating mixer or HulaMixer[™] Sample Mixer under constant vertical rotation (50 rpm).
- Briefly spin down the tubes and place them in the DynaMag[™]-2 Magnet for 5 minutes.
- 8. While in the magnetic rack carefully pipette off the supernatant without transferring beads.

Note: The beads contain the DNA.

Guidelines to purify DNA

- Bring Wash Buffers and Elution Buffer from Dynabeads[™] SILANE Genomic DNA Kit to room temperature before use.
- Vortex sufficiently to ensure that beads are resuspended completely in each wash.
- Remove any residual Wash Buffer 2 before drying the beads at 65°C.

 Add ethanol and isopropanol to the wash buffers as recommended on the bottles.

Purify DNA

- 1. Remove the tubes from the magnetic rack and add 900 μL of Wash Buffer 1.
- 2. Vortex at maximum setting to make sure that beads are resuspended completely.
 - Briefly spin down the tubes.
- 3. Place the samples in the magnet for 3 minutes.
- 4. While the tubes are in the magnet, remove the supernatant without transferring beads.
- 5. Repeat the washing (steps 1–3) with Wash Buffer 1.
- 6. Add 900 µL of Wash Buffer 2.
- Resuspend the beads completely by vortexing. Briefly spin down the tubes.
- 8. Place the samples in the magnet for 3 minutes.
- 9. While the tubes are in the magnet, remove the supernatant without transferring beads.
- 10. Add 400 µL Wash Buffer 2.
- Resuspend the beads completely by vortexing.
 Briefly spin down the tubes.
- 12. Place the samples in the magnet for 3 minutes.
- While the tubes are in the magnet, remove the supernatant without transferring beads.
 - Briefly spin down the tubes.
- 14. Place the samples in the magnet for 3 minutes.
- **15.** Remove any residual Wash Buffer and remove the tubes from the magnetic rack.
- 16. Dry the beads with the lid open at 65°C for 15 minutes in a Eppendorf[™] ThermoMixer[™] C (or a dry-bath).
- 17. Add 60 µL Elution Buffer.
- **18.** Incubate at room temperature for 7 min under constant shaking (1400 rpm) in the Eppendorf[™] ThermoMixer[™] C, or use a vortex with a foam adapter at moderate speed.
- 19. Place the tubes in the magnet for 2 minutes.
- Carefully transfer the eluate (55 μL) to a fresh tube. The eluate contains bisulfite-converted DNA that is used in subsequent qPCR.

Guidelines for setting up qPCR

- The PureQuant[™] CD3+ T Cell Assay requires the use of PureQuant[™] qPCR Master Mix II.
- Use the same six standard dilutions for both CD3 and GAPDH Standards.
- All Samples, Standards and Controls are run in triplicate.
- Refer to instrument user manual for instructions programming the qPCR run with a Standard Curve.

IMPORTANT! Assign all the standard dilutions as "Standard." Assign Samples, Calibrator and Reference as "Unknown"; and no template control as "NTC or N" in triplicate.

IMPORTANT! While assigning wells for each standard, provide copy numbers that are specified in the Final Copy Number column (see "Prepare Standard dilutions").

Prepare Standard dilutions

Prepare a full panel of six standard dilutions by serially diluting the PureQuant™ High Copy Standard B starting at 31250 copies/3 µL according to the following table. The same standard dilutions are used for both CD3 and GAPDH in qPCR. Store the standard dilutions at −20°C.

Standard Dilutions	Copy Number STD volume	Diluent DNA [1]	Final Copy Number per 3 µL
STD#1	1000 μL	_	31250
STD#2	mix 200 μL STD#1	800 µL	6250
STD#3	mix 200 μL STD#2	800 µL	1250
STD#4	mix 200 μL STD#3	800 µL	250
STD#5	mix 200 μL STD#4	800 µL	50
STD#6 ^[2]	mix 30 µL STD#3	1200 µL	30

 $^{^{[1]}~}$ 10 ng/µL Lambda DNA in TE (10 mM Tris, 1 mM EDTA, pH 8.0)

^[2] Use STD#3 to prepare STD#6.

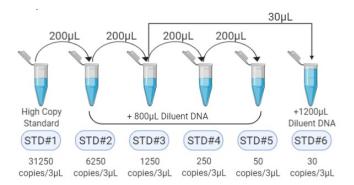


Figure 1 Serial dilutions

Prepare PCR plates

 Prepare the qPCR Master Mix II Cocktail in two separate tubes, one each for CD3 and GAPDH, excluding Template DNA according to the following table:

Reagent	Amount
Template DNA	3 µL
Lambda DNA (50 ng/µL in TE, pH8.0)	1 µL
TaqMan™ assay	0.5 μL
Water, Nuclease-free	0.5 μL
PureQuant™ qPCR Master Mix II	5 μL
TOTAL	10 μL

Load the templates first and then add 7 μL of the qPCR Master Mix II Cocktail.

	CD3			GAPDH								
	1	2	3	4	5	6	7	8	9	10	11	12
Α		STD-1_CD	3	Sample#1		STD-1_GAPDH		Sample#1				
В		STD-2_CD	3		Sample#2		STD-2_GAPDH		Sample#2			
С		STD-3_CD	3	Sample#3		STD-3_GAPDH		Sample#3				
D		STD-4_CD	3	Sample#4		STD-4_GAPDH		Sample#4				
E		STD-5_CD	3	Sample#5		STD-5_GAPDH		Sample#5				
F		STD-6_CD	3	Sample#6		STD-6_GAPDH		Sample#6				
G		Reference	1	Sample#7		Reference		Sample#7				
Н		Calibrator		NTC		Calibrator		NTC				

Figure 2 A representative 96-well plate layout illustrating CD3 assay set up. A 384-well plate can also be used for qPCR using the same volumes for a 96-well plate.

Note: All Samples, Standards and Controls are run in triplicate.

- 3. Seal the plate with film and briefly centrifuge before placing into the qPCR instrument.
- 4. Select FAM or equivalent as dye and Non Fluorescent Quencher (NFQ) or equivalent as quencher.

Table 2 qPCR cycle setup parameters

Step	Time	Temp	Cycles	
Pre-incubation	35 min	95°C	1X	
A :G +:	15 sec	95°C	50X	
Amplification	1 min	61°C		
Cooldown	5 sec	42°C	1X	

Run qPCR using the cycle set up parameters listed in Table 2.

IMPORTANT! See "Guidelines for setting up qPCR" for instructions on setting up the qPCR standard curve and other controls.

 After the qPCR run is complete, execute analysis on the qPCR instrument to calculate: Ct Average, Ct Standard Deviation, and Copy Number. Export data in .txt or .xlsx format. Analyze data using PureQuant[™] CD3+ T Cell Assay Microsoft[™] Excel[™] Analysis Template. Download Analysis Template from thermofisher.com/order/catalog/product/ A47197.

The analysis template has three tabs: Tab#1 contains instructions and guidance on how to perform data analysis, Tab#2 is the Analysis Template, and Tab#3 contains an Example.

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Revision history: Pub. No. MAN0019205

Revision	Date	Description
C.0	1 December 2021	The storage temperature for the PureQuant™ qPCR Master Mix II was corrected from 4°C to -20°C.
B.0		Updated the product description and guidelines for setting up qPCR to include the requirement of PureQuant™ qPCR Master Mix II.
A.0	29 May 2020	Baseline for this revision history.

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