TECHNICAL DATA SHEET

THUNDER™ Human GRANZYME B TR-FRET Biomarker Assay Kit



CATALOG NUMBERS KIT-GRB-100 (100 tests)

KIT-GRB-100 (100 tests)
KIT-GRB-500 (500 tests)
KIT-GRB-2500 (2500 tests)
KIT-GRB-5000 (5000 tests)
KIT-GRB-10000 (10000 tests)

Store at -80°C.
For research use only.
Not for use in diagnostic procedures.



PRODUCT DESCRIPTION

This assay kit is designed for the quantitative determination of **human Granzyme B** in cell culture supernatants using a simple, rapid and sensitive immunoassay based on the homogeneous (no wash) THUNDER™ TR-FRET technology. The kit <u>does not</u> include a recombinant **human Granzyme B** standard, which must be purchased separately from Sino Biological (catalogue number 10345-H08H).

SPECIFICITY

This assay kit contains two specific and selective antibodies that recognize both natural and recombinant **human Granzyme B** in cell culture supernatant samples.

SPECIES REACTIVITY

Human (Swiss-Prot Acc. P10144; Entrez Gene Id 3002).

TR-FRET ASSAY PRINCIPLE

The THUNDER™ **human Granzyme B** assay kit is a homogeneous time-resolved Förster resonance energy transfer (TR-FRET) sandwich immunoassay (Figure 1). The assay workflow consists of a one-step reagent addition (Figure 2). Human Granzyme B in the cell supernatant sample is detected with a pair of fluorophorelabeled antibodies in a simple "add-incubate-measure" format (no wash steps). One antibody is labeled with a donor fluorophore (Europium chelate; Eu-Ab1) and the second with a far-red acceptor fluorophore (FR-Ab2). The binding of the two labeled antibodies to distinct epitopes on the target analyte takes place in solution and brings the two dyes into close proximity. Excitation of the donor Europium chelate molecules with a flash lamp (320 or 340 nm) or a laser (337 nm) triggers a FRET from the donor to the acceptor molecules, which in turn emit a TR-FRET signal at 665 nm. Residual energy from the Eu chelate generates light at 615 nm. The signal at 665 nm is proportional to the concentration of human Granzyme B in the cell supernatant. Data can be expressed as either the signal at 665 nm or the 665 nm/615 nm ratio. A distinct advantage of the THUNDER™ TR FRET technology is that it does not require addition of potassium fluoride to ensure signal stability. The non-toxic nature of THUNDER™ makes it an ideal candidate for multiplexing with other assay technologies using a sequential assay protocol.

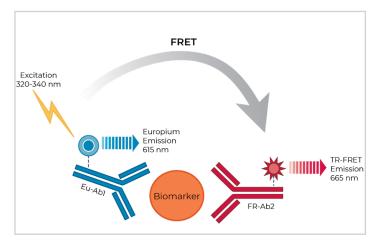


Figure 1 Schematic representation of the TR-FRET assay principle.



Figure 2 THUNDER™ human Granzyme B assay workflow.

THUNDER™ ASSAY PROTOCOL

This protocol must be read in its entirety prior to beginning the assay.

- · We cannot guarantee the performance of the product outside the conditions detailed in this Assay Protocol.
- · Bring all reagents to room temperature prior to running the assay.
- · Centrifuge all tubes before use to improve recovery of content (2000x g, 10-15 sec).
- \cdot Use ultrapure water (Milli-Q[®] grade water; 18 M Ω cm) to dilute Assay Buffer.
- · It is recommended to test all standards in triplicate and samples at least in duplicate.

KIT COMPONENTS	100 points*	500 points*
Eu-labeled anti-Granzyme B antibody (Eu-Ab1)	5 μL (1 clear tube, red cap)	25 µL (1 clear tube, red cap)
Acceptor-labeled anti-Granzyme B antibody (FR-Ab2)	20 µL (1 brown tube, blue cap)	100 µL (1 brown tube, blue cap)
Assay Buffer 1 (5X)**	1 mL (1 tube, yellow cap)	3 mL (3 tubes, yellow cap)

^{*} The number of assay points is based on an assay volume of 20 μL in lowvolume 384-well assay plates using the kit components at the recommended concentrations.

ADDITIONAL MATERIALS REQUIRED	Recommended source	Catalog No.
Recombinant human Granzyme B	Sino Biological	10345-H08H
Ultrapure laboratory grade water	Many options available	NA
Low-volume 384-well microplate, white	PerkinElmer Greiner Corning	6007290 784075 4513
Multi- and single-channel pipettes	Many options available	NA
Adhesive sealing film for plates	Many options available	NA
A plate reader equipped with the TR-FRET option	Many options available	NA

REAGENT PREPARATION

- The instructions described below are for testing the entire number of assay points in each kit. Adjust volumes accordingly for testing of fewer assay points.
- · Prepare only as much reagent as is needed on the day of the experiment.

STEP 1 PREPARATION OF 1X ASSAY BUFFER 1

- · Mix end-over-end the 5X Assay Buffer 1 before use.
- The thawed 5X Assay Buffer 1 can be stored at 4°C for 1-2 weeks. For longer periods of time, buffer should be stored at -80°C.
- · Unused 1X Assay Buffer 1 can be stored at 4°C for 2 days.

For the 100-point kit

Add 1 mL of 5X Assay Buffer 1 to 4 mL of water.

For the 500-point kit

Add 3 mL of 5X Assay Buffer 1 to 12 mL of water.

STEP 2

PREPARATION OF hGranzyme B STANDARD SERIAL DILUTIONS

IMPORTANT NOTE

The analyte human Granzyme B is not included in the kit. The kit has been validated using human Granzyme B recombinant protein from Sino Biological (Catalog number 10345-H08H). Other commercial recombinant human Granzyme B may also work but have not been tested with this kit.

- · Prepare working standards just before use and use within one hour. Do not store working standards.
- Each well requires 15 µL of working standard.
- · Change tip between each standard dilution.
- · Mix each tube thoroughly before the next transfer.
- · The following section describes the preparation of a standard curve for triplicate measurements (recommended).

NOTE

The Working Stock Solution should be used within one hour or aliquoted into screw-capped polypropylene vials and stored at -80°C for long term storage.



^{**} Extra Assay Buffer can be ordered separately (cat # TRFRET-AB1).

REAGENT PREPARATION (CONTINUED)

- A. Follow the manufacturer's instructions to reconstitute the analyte (typically at 100-200 µg/mL) to create a Stock Solution.
- B. Create a 1 µg/mL Working Stock Solution by diluting the Stock Solution in either 1X Assay Buffer 1 if the standard curve is done in Assay buffer or in culture medium with 10% fetal bovine serum (FBS) if the standard curve is done in culture medium. Use this Working Standard Solution to prepare a standard dilution series.
- C. Label 12 tubes, one for each working standard. Alternatively, serial dilutions can be conducted using a polypropylene 96-well plate.
- D. Prepare serial dilutions for the standard curve as indicated in the table below using either 1X Assay Buffer 1 or culture medium with 10% FBS (for testing of culture supernatant samples):

Tube	Volume of Granzyme B	Vol. of 1X Assay Buffer or culture medium +10% FBS (μL)	[Granzyme B] in standard curve (g/mL in 15 µL)
1	30 µL diluted hGranzyme B (1E-6 g/mL)	70	3.0E-7
2	30 μL of tube 1	70	1.0E-7
3	30 µL of tube 2	60	3.0E-8
4	30 μL of tube 3	70	1.0E-8
5	30 µL of tube 4	60	3.0E-9
6	30 μL of tube 5	70	1.0E-9
7	30 μL of tube 6	60	3.0E-10
8	30 μL of tube 7	70	1.0E-10
9	30 µL of tube 8	60	3.0E-11
10	30 μL of tube 9	70	1.0E-11
11	30 µL of tube 10	60	3.0E-12
12	30 µL of tube 11	70	1.0E-12
13*	0	60	0

^{*} In order to calculate the Limit of Detection (LOD), three additional zero standard tubes (13-16) are assayed in triplicate (12 wells in the 384-well plate).

STEP 3 PREPARATION OF 4X ANTIBODY DETECTION MIX

- · Prepare and mix just before use.
- · Each well requires 5 µL of 4X Antibody Detection Mix
- The unused 4X Antibody Detection Mix may be stored at 4°C for 2 days or aliquoted and stored at -80°C. Avoid repeated freeze/thaw cycles.

For the 100-point kit

NOTE

Due to the low reagent volumes in the 100-point kit, the antibodies are diluted with 1X Assay Buffer 1 directly in the vial.

- A. Add 255 µL of 1X Assay Buffer 1 into the vial containing 5 μL of Eu Ab1 stock solution.
- B. Add 240 µL of 1X Assay Buffer 1 into the vial containing 20 µL of FR Ab2 stock solution.
- C. Mix gently 260 µL of pre-diluted Eu Ab1 with 260 µL of pre-diluted FR-Ab2.

For the 500-point kit

- A. Mix gently 1,275 µL of 1X Assay Buffer 1 with the 25 µL of Eu Abl stock solution.
- B. Mix gently 1,200 µL of 1X Assay Buffer 1 with the 100 µL
- C. Mix gently 1,300 µL of pre-diluted Eu Ab1 with 1,300 µL of pre-diluted FR Ab2.



SAMPLE PREPARATION

- · Each well requires 15 µL of sample (cell supernatant).
- Just after their collection, put the samples at +4°C and assay immediately or aliquot and store -80°C. Avoid repeated freeze/thaw cycles.
- · Do not use a water bath to thaw samples. Thaw samples at room temperature.
- · If the analyte concentration in the sample possibly exceeds the highest point of the standard curve, prepare one or more 10-fold dilutions of the sample. Prepare serial dilutions using the same medium used to culture the cells.

ASSAY PROCEDURE

- · Samples and standards must be assayed at least in duplicate each time the assay is performed.
- · When loading reagents in the low-volume 384-well microplate, change tips between each standard or sample addition and after each set of reagents.
- · If using a multichannel pipettor, always use a new disposable reagent reservoir.
- · When reagents are added to the microplate, make sure the liquids are at the bottom of the well.
 - **A.** Add 15 μ L of each working standard or 15 μ L of cell supernatant sample.
 - B. Add 5 µL of 4X Antibody Detection Mix (Eu-Ab1 + FR-Ab2) to each of the assay wells.
 - C. Cover the plate with a plate sealer and incubate 2 hours at room temperature.
 - D. Gently remove the adhesive plate sealer. Read plate on a TR-FRET compatible microplate reader.

NOTE The same plate can be read several times without any negative effect on the assay performance.

TR-FRET PLATE READER SETTINGS

For optimal assay performance, we recommend that you read the TR FRET assays at two wavelengths, detecting both the emission from the Europium chelate donor fluorophore at 615 nm, and the acceptor fluorophore at 665 nm. The following instrument settings are provided as guidelines.

	TR-FRET Compatible Plate Reader*			
Parameter	Flash lamp excitation Laser excitation			
Excitation filter	320 nm (or 340 nm)	Not applicable		
Emission filter	615 nm (or 620 nm)	615 nm (or 620 nm)		
Delay time	90 µs	50 µs		
Flash energy level	100% or High	100%		
Number of flashes	100-200	20		
Window (integration time)	300 μs	100 µs		

^{*}These settings are provided as guidelines only. Settings should be optimized for each reader. For optimal settings, visit bioauxilium.com/resources.

DATA ANALYSIS

- TR-FRET data are typically calculated and presented ratiometrically using the following formula: [(665 nm/615 nm) x 1,000]
- · Alternatively, the signals at 665 nm can be used directly to process your data.
- · Calculate the TR-FRET ratio for each well.
- · Since TR-FRET signal is read in a time-resolved mode, background substraction is usually not necessary.
- · Create a standard curve by analyzing data according to a nonlinear regression using the 4 parameter logistic equation (sigmoidal doseresponse curve with variable slope) and a $1/Y^2$ data weighting. For correct analysis, the highest standard point should be removed from the curve if it is present after the hook point (i.e., if it shows lower counts compared to the next standard).
- · Do not use a value of zero for the zero standard. Use instead the actual background values obtained with the zero standard.
- · If samples have been diluted, the concentration read from the standard curve must be multiplied by the dilution factor.



TYPICAL STANDARD CURVE

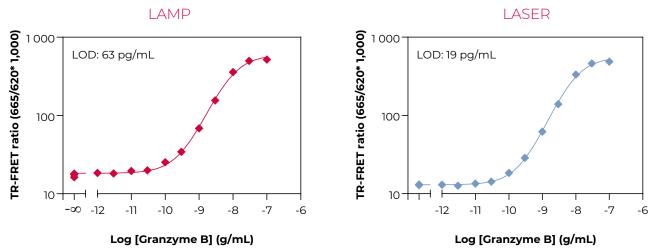


Figure 3 Typical nonlinear human Granzyme B standard curves in Assay Buffer 1. Left: lamp excitation, Right: laser excitation. Data represent the mean ± standard deviation of triplicate measurements (3 wells) for each standard. The data was generated using a low-volume 384-well white plate read on the PHERAstar® FSX equipped with TR-FRET option.

VALIDATION DATA

The THUNDER™ human Granzyme B kit has been validated using human Granzyme B protein from Sino Biological (Catalog number 10345-H08H; NOT included in the kit).

HUMAN Granzyme B THUNDER™ ASSAY SUMMARY					
Assay Type	Homogeneous sand	Homogeneous sandwich immunoassay			
Format	384-we	II plate			
Hands-on time	15 mir	nutes			
Incubation time	2 ho	urs			
Sample type (volume)	Cell culture supernatant (15 μL)				
Specificity	Natural and recombinant human Granzyme B				
	Laser excitation Lamp excitation				
Limit of Detection (LOD)	19 pg/mL 63 pg/mL				
Lower Limit of Quantification (LLOQ)	82 pg/mL 268 pg/mL				
EC _{so}	9.2 ng/mL 8.8 ng/mL				
Dynamic Range	19 – 100,000 pg/mL	63 – 100,000 pg/mL			

SENSITIVITY

The Limit of Detection (LOD) was calculated by adding two standard deviations to the mean background counts of 12 zero standard replicates and calculating the corresponding concentration on the standard curve.

The Lowest Level of Quantitation (LLOQ) was calculated by adding 10 standard deviations to the mean background counts of 12 zero standard replicates and calculating the corresponding concentration on the standard curve.

	ASSAY BUFFER 1		DMEM +10% FBS		RPMI +10% FBS	
	Laser	Lamp	Laser	Lamp	Laser	Lamp
# of standard curves	10	10	3	3	3	3
LOD (pg/mL)	19	63	14	72	43	123
LLOQ (pg/mL)	82	268	100	324	198	627



PRECISION

INTRA-ASSAY PRECISION

Three samples of known concentration (low, medium, high) were tested twenty-one times each on the same plate to assess intra-assay precision. The assay was run in a 384-well plate with standards and samples prepared in Assay Buffer 1. The resulting low %CV (<10%) for each sample indicates good reproducibility within an assay.

Sample concentration	Low	Med	High
# of samples	21	21	21
Mean (pg/mL)	921	2 730	12 229
SD (pg/mL)	40	178	931
CV (%) Intra-Assay	4.4	6.5	7.6

INTER-ASSAY PRECISION

Three samples of known concentration were tested in separate assays to assess inter-assay precision. A total of eight independent experiments were performed by two operators using two different kit lots: The assay was run in 384-well plates with standards and samples prepared in Assay Buffer 1 and tested in triplicate. The resulting low %CV (<10%) for each sample indicates good reproducibility between assays.

Sample concentration	Low	Med	High
# of samples	24	24	24
Mean (pg/mL)	917	2 652	10 372
SD (pg/mL)	52	150	969
CV (%) Intra-Assay	5.7	5.6	9.3

RECOVERY

To assess the accuracy of the assay, spike-to-recovery experiments were conducted. Each assay consisted of one standard curve and three spike concentrations (low, medium and high), assayed in triplicate in Assay Buffer 1, DMEM and RPMI (supplemented with 10% FBS). The average recovery was calculated from the average of 3 wells and compared to a standard curve prepared in the corresponding diluent. Data show recovery values within the range of 80–120% demonstrating assay accuracy.

	Spike level	Expected (pg/mL)	Observed (pg/mL)	Average % Recovery
Assay Buffer 1	High	10 000	10 277	103
	Medium	3 000	2 927	98
	Low	1 000	938	94
DMEM + 10% FBS	High	10 000	8 351	84
	Medium	3 000	2 405	80
	Low	1 000	810	81
RPMI + 10% FBS	High	10 000	9 145	91
	Medium	3 000	3 238	108
	Low	1 000	1 111	111



LINEARITY

To assess the linearity of the assay, samples were spiked with a high concentration of analyte in culture medium supplemented with 10% FBS and diluted with the same culture medium with 10% FBS. The assay was run in a 384-well plate with standards prepared in culture media with 10% FBS. Each sample was measured in triplicate. Data show recovery values within 80-120% confirming assay linearity.

	Expected (pg/mL)	Observed (pg/mL)	Average % Recovery
DMEM + 10% FBS	15 625	15 131	97
	7 813	6 815	87
	3 906	3 174	81
	1 953	1 554	80
	977	794	81
	488	525	108
	244	212	87
	122	102	84
	61	53	87
RPMI + 10% FBS	15 625	13 281	85
	7 813	6 599	84
	3 906	3 366	86
	1 953	1 605	82
	977	870	89
	488	423	87
	244	199	81
	122	126	103
	61	59	97
	31	27	87

