TransAM™ T-bet

(version A)

Catalog Nos. 51396 & 51896

Active Motif North America

1914 Palomar Oaks Way, Suite 150 Carlsbad, California 92008, USA Toll free: 877 222 9543

Telephone: 760 431 1263 Fax: 760 431 1351

Active Motif Europe

104 Avenue Franklin Roosevelt B-1330 Rixensart, Belgium UK Free Phone: 0800 169 31 47

France Free Phone: 0800 90 99 79

Germany Free Phone: 0800 181 99 10

Telephone: +32 (0)2 653 0001

Fax: +32 (0)2 653 0050

Active Motif Japan

Azuma Bldg, 7th Floor 2-21 Ageba-Cho, Shinjuku-Ku Tokyo, 162-0824, Japan

Telephone: +81 3 5225 3638 Fax: +81 3 5261 8733

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Overview

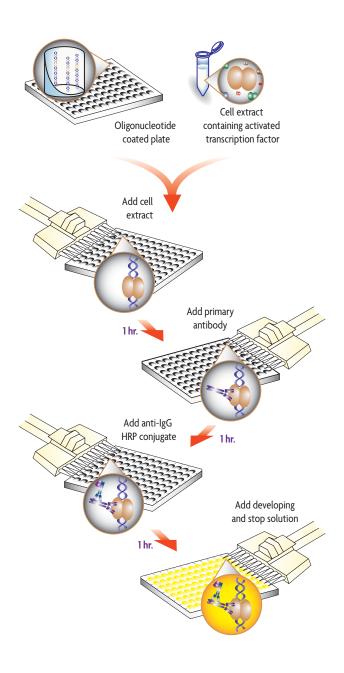
Members of the T-box gene family of transcription factors are defined by their homology within a 200-aa DNA binding domain called the T-box. These proteins are important regulators of several early developmental processes. Seventeen different T-box genes with diverse regulatory functions in development and diseases have been identified in mammals. Two T-box genes are expressed in cells of the immune system: Eomesodermin and Tbx-21, which is also known as T-bet for T-box expressed in T-cells. T-bet is a 535 amino acids transcription factor essential to T helper 1 (Th1) cell generation and effector function. Recently, the expression of T-bet has also been described in B, natural killer (NK), CD8+ T and dentritic cells. T-bet regulates expression of numerous immune system-associated genes, including cytokines, cytokine receptors, chemokines, chemokine receptors and cytotoxicity.

With its patented TransAM™ method, Active Motif introduced the first ELISA-based kits to detect and quantify transcription factor activation. TransAM Kits combine a fast, user-friendly format with a sensitive, specific assay. TransAM T-bet Kits are designed specifically to detect and quantify T-bet activation. They contain a 96-well plate to which oligonucleotide containing the T-bet consensus binding site has been immobilized. T-bet contained in nuclear extract bind specifically to this oligonucleotide and is detected through use of an antibody directed against T-bet. Addition of a secondary antibody conjugated to horseradish peroxidase (HRP) provides a sensitive colorimetric readout that is easily quantified by spectrophotometry. The 96-well plate with individual strips of 8-wells is suitable for manual use or for high-throughput screening applications.

product	format	catalog no.
TransAM™ T-bet	1 x 96 rxns	51396
TransAM™ T-bet	5 x 96 rxns	51896



^{*} Technology covered by AAT-filed patents and licensed to Active Motif.



Introduction

T-bet Transcription Factor

T-bet (T-box expressed in T-Cells; Tbx21) is a member of the T-box transcription factor family and plays a critical role in the generation of T helper 1 (Th1) cells¹. Th1 cells secrete IFN- γ and Tumor Necrosis Factor β (TNF- β) and are central to cellular immunity against intracellular pathogens. Th1 cells lead to susceptibility to infectious diseases, particularly those of parasites, bacteria, and fungi for which phagocytosis and intracellular killing are important². T-bet has been shown to promote Th1 development, and at least part of its role in this process is due to its ability to directly regulate key lineage determinant genes such as IFN- γ and CXCR3. T-bet has been shown to bind directly to the promoter regions of these genes, and the expression of T-bet is required and sufficient to induce transcription. The level of T-bet expression is increased by signals mediated by the T cell receptor (TCR). T-bet expression has also been shown to correlate with permissive histone acetylation and methylation, as well as the induction of DNase hypersensitivity at the IFN- γ locus in Th1 cells³.

Transcription Factor Assays

To date, three methods are widely used to measure T-bet activation, either directly or indirectly:

- T-bet activation can be determined by Western blot using antibodies specific for T-bet proteins. This method is time consuming (up to 2 days once the nuclear extracts are prepared), and is not suitable for processing large numbers of samples.
- 2. The DNA-binding capacity of T-bet can be assayed by gel retardation, also called electrophoretic mobility shift assay (EMSA). In this method, nuclear extracts are incubated with a radioactive double-stranded oligonucleotide probe containing the consensus sequence for T-bet binding. If T-bet is active in the nuclear extract, it will bind to the probe. Samples are then resolved by electrophoresis on a native polyacrylamide gel, followed by autoradiography. This method is sensitive, but like the previous procedure, it is time consuming (multiple days of gel exposure may be required to achieve sufficient sensitivity) and it cannot be applied to high-throughput screening. Gelshift assays also require special precautions and equipment for handling radioactivity.
- 3. Another method used to assay T-bet activation is based on reporter genes, typically luciferase or β -galactosidase, placed under the control of a promoter containing the T-box recognition sequence. The promoter can be artificial and made of several consensus-binding elements. The procedure is limited by the following issues: (i) reporter gene assays have to be repeated several times to obtain statistically reliable data; and (ii) reporter gene assays are sensitive to confounding factors that may influence the expression level of the reporter gene. Therefore, assays have to be carefully standardized. This method is sensitive and easy to perform with a large number of samples but requires efficient cell transfection with the reporter plasmid.

TransAM T-bet

The TransAM Kit combines a fast and user-friendly ELISA format with a sensitive and specific assay for transcription factors. TransAM T-bet Kit contains a 96-well plate on which an oligonucleotide containing the T-box consensus binding site (5 ´-AC ACC TAG GTG T-3 ´) has been immobilized. The active form of T-bet contained in nuclear extract specifically binds to this oligonucleotide. The primary antibody used to detect T-bet recognizes an epitope on the T-bet protein upon DNA binding. Addition of an HRP-conjugated secondary antibody provides a sensitive colorimetric readout easily quantified by spectrophotometry. Once the nuclear extracts are prepared, this assay is completed in less than 3.5 hours. As this assay is performed in 96-well plates, a large number of samples can be handled simultaneously, enabling high-throughput automation. This assay is specific for T-bet activation and has been shown to be 5-fold more sensitive and 20-fold faster than the gel-retardation technique. With the 3.5-hour procedure of TransAM, we could detect T-bet activation using as little as 1 µg of nuclear extract.

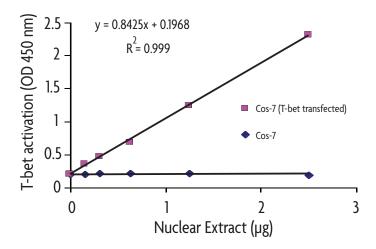
Kit Performance and Benefits

Detection limit: < 1 µg nuclear extract/well.

Range of detection: TransAM provides quantitative results from 0.125 to 2.5 µg of nuclear extract per well.

Cross-reactivity: TransAM T-bet specifically detects T-bet from human origin.

Assay time: 3.5 hours.



Monitoring T-bet activation with the TransAM T-bet Kit:

0.125 to 2.5 µg of Cos-7 (T-bet transfected and treated with PMA/ionomycin) nuclear extract (rectangle-shape) and untransfected Cos-7 nuclear extracts (diamond-shape) were assayed per well. Data shown are the results from wells assayed in duplicate.

Kit Components and Storage

TransAM T-bet Kits are for research use only. Not for use in diagnostic procedures. Except for the nuclear extract that must be kept at -80°C, kit components can be stored at -20°C prior to first use. Then, we recommend storing each component at the temperature indicated in the table below. Store the T-bet antibody at 4°C after it has been thawed for use. All components are guaranteed stable for 6 months from date of receipt when stored properly.

Reagents	Quantity 1 plate / 5 plates	Storage
T-bet antibody	5 μl / 25 μl	4°C
anti-mouse HRP-conjugated antibody	10 μl / 50 μl	4°C
Wild-type oligonucleotide AM31	100 μl / 500 μl (10 pmol/μl)	-20°C
Mutated oligonucleotide AM31	100 μl / 500 μl (10 pmol/μl)	-20°C
Cos-7	20 μl / 100 μl	
nuclear extract (T-bet transfected) PMA/ionomycin-treated	(2.5 μg/μl)	-80°C
Dithiothreitol (DTT) (1 M)	100 μl / 500 μl	-20°C
Protease Inhibitor Cocktail	100 μl / 500 μl	-20°C
poly d[I-C]	100 µl / 500 µl (1 µg/µl)	-20°C
Lysis Buffer AM1	10 ml / 50 ml	4°C
Binding Buffer AM5	10 ml / 50 ml	4°C
10X Wash Buffer AM2	22 ml / 110 ml	4°C
10X Antibody Binding Buffer AM3	2.2 ml / 11 ml	4°C
Developing Solution	11 ml / 55 ml	4°C
Stop Solution	11 ml / 55 ml	4°C
96-well T-bet assay plate	1/5	
Plate sealer	1/5	

Additional materials required

- Multi-channel pipettor
- Multi-channel pipettor reservoirs
- Rocking platform
- Microplate spectrophotometer capable of reading at 450 nm (655 nm as optional reference wavelength)

Protocols

Buffer Preparation and Recommendations

Preparation of Complete Lysis Buffer

We provide an excess of Lysis Buffer AMI in order to perform the assay AND to prepare customized nuclear extracts. Please refer to the Appendix Section A for a protocol to prepare a nuclear extract. Our Nuclear Extract Kit can also be purchased separately (Cat. Nos. 40010 & 40410). Lysis Buffer AMI contains phosphatase inhibitors to prevent dephosphorylation of transcription factors during the extract preparation and the assay. The presence of these inhibitors gives a yellow coloration to Lysis Buffer AMI. Prepare the amount of Complete Lysis Buffer required for the assay by adding 1 μ l of 1 M DTT and 10 μ l Protease Inhibitor Cocktail per ml of Lysis Buffer AMI (see the Quick Chart for Preparing Buffers in this section). Some of the protease inhibitors lose their activity after 24 hours once diluted. Therefore, we recommend using the Complete Lysis Buffer immediately for cell lysis. The remaining amount should be discarded if not used in the same day.

Preparation of Complete Binding Buffer

Prepare the amount of Complete Binding Buffer AM5 required for the assay by adding 1 μ l of DTT and 10 μ l of 17 μ g/ml poly[d(l-C)] per ml of Binding Buffer (see the Quick Chart for Preparing Buffers in this section). After use, discard remaining Complete Binding Buffer.

Preparation of 1X Wash Buffer

Prepare the amount of 1X Wash Buffer required for the assay as follows: For every 100 ml of 1X Wash Buffer required, dilute 10 ml 10X Wash Buffer AM2 with 90 ml distilled water (see the Quick Chart for Preparing Buffers in this section). Mix gently to avoid foaming. The 1X Wash Buffer may be stored at 4°C for one week. The Tween 20 contained in the 10X Wash Buffer AM2 may form clumps, therefore homogenize the buffer by incubating at 50°C for 2 minutes and mixing prior to use.

Preparation of 1X Antibody Binding Buffer

Prepare the amount of 1X Antibody Binding Buffer required for the assay as follows: For every 10 ml of 1X Antibody Binding Buffer required, dilute 1 ml 10X Antibody Binding Buffer AM3 with 9 ml distilled water (see the Quick Chart for Preparing Buffers in this section)*. Mix gently to avoid foaming. Discard remaining 1X Antibody Binding Buffer after use. The BSA contained in the 10X Antibody Binding Buffer AM3 may form clumps, therefore homogenize the buffer by warming to room temperature and vortexing for 1 minute prior to use. Dilute the primary antibody with the 1X Antibody Binding Buffer to 1:2000 for the T-bet and the HRP-conjugated secondary antibody to 1:1000. Depending on the particular assay, the signal:noise ratio may be optimized by using higher dilutions of both antibodies. This may decrease the sensitivity of the assay.

* Volumes listed refer to the preparation of buffer for diluting both the primary & secondary antibodies.



Developing Solution

The Developing Solution should be warmed to room temperature before use. The Developing Solution is light sensitive, therefore, we recommend avoiding direct exposure to intense light during storage. The Developing Solution may develop a yellow hue over time. This does not affect product performance. A blue color present in the Developing Solution indicates that it has been contaminated and must be discarded. Prior to use, place the Developing Solution at room temperature for at least 1 hour. Transfer the amount of Developing Solution required for the assay into a secondary container before aliquoting into the wells (see the Quick Chart for Preparing Buffers in this section). After use, discard remaining Developing Solution.

Stop Solution

Prior to use, transfer the amount of Stop Solution required for the assay into a secondary container (see the Quick Chart for Preparing Buffers in this section). After use, discard remaining Stop Solution

WARNING: The Stop Solution is corrosive. Wear personal protective equipment when handling, i.e. safety glasses, gloves and labcoat.

Cos-7 nuclear extract (T-bet transfected) PMA/ionomycin-treated

The Cos-7 (T-bet transfected) PMA/ionomycin-treated nuclear extract is provided as a positive control for T-bet activation. Sufficient extract is supplied for 25 reactions if using 1.25 μ g per well. This extract is optimized to give a strong signal when used at 1.25 μ g/well. We recommend aliquoting the extract in 5 μ l fractions and storing at -80°C. Avoid multiple freeze/thaw cycles of the extract.

Wild-type and mutated consensus oligonucleotides

The wild-type consensus oligonucleotide is provided as a competitor for T-bet binding in order to monitor the specificity of the assay. Used at 40 pmol/well, the oligonucleotide will prevent T-bet binding to the probe immobilized on the plate. Conversely, the mutated consensus oligonucleotide should have no inhibitory effect on T-bet binding. Prepare the required amount of wild-type and/or mutated consensus oligonucleotide by adding 4 μ l of appropriate oligonucleotide to 36 μ l of Complete Binding Buffer per well being used (see the Quick Chart for Preparing Buffers in this section). To allow for optimum competition, add the oligonucleotide to the well prior to addition of the nuclear extract.



Quick Chart for Preparing Buffers

Reagents to prepare	Components	1 well	1 strip (8 wells) (6 strips 48 wells) (12 strips 96 wells)
Complete Lysis Buffer	DTT	0.01 µl	0.1 μl	0.6 μl	1.2 µl
	Protease Inhibitor Cocktail	0.12 µl	0.9 μl	5.4 μl	10.8 µl
	Lysis Buffer	11.12 µl	89.0 μl	534.0 μl	1.07 ml
	Total Required	11.25 µl	90.0 μl	540.0 μl	1.08 ml
Complete Binding Buffer	DTT	0.04 μl	0.3 μl	2.16 µl	4.3 μl
	poly d[I-C]	0.45 μl	3.6 μl	21.6 µl	43.2 μl
	Binding Buffer	44.5 μl	356.1 μl	2.14 ml	4.27 ml
	Total Required	45 μl	360 μl	2.16 ml	4.32 ml
Binding Buffer with	wt or mut oligont	4 μl	32 μl	192 µl	N/A
T-bet wt or	Complete Binding Buffer	41 μl	328 μl	1.97 ml	N/A
mut oligont	Total Required	45.0 μl	360.0 μl	2.16 ml	N/A
1X Washing Buffer	Distilled Water	2.025 n	nl 16.2 ml	97.2 ml	194.4 ml
	10X Washing Buffer	225.0 µl	1.8 ml	10.8 ml	21.6 ml
	Total Required	2.25 m	l 18.0 ml	108.0 ml	216.0 ml
1X Antibody Binding Buffer*	Distilled Water 10X Antibody Binding Buffer Total Required	202.5 μl 22.5 μl 225.0 μl	1.62 m 180.0 µl 1.8 ml	l 9.72 ml 1.08 ml 10.8 ml	19.44 ml 2.16 ml 21.6 ml
Developing Solution	Total Required	112.5 µl	900.0 μl	5.4 ml	10.8 ml
Stop Solution	Total Required	112.5 µl	900.0 μl	5.4 ml	10.8 ml

^{*} Volumes listed refer to the preparation of buffer for diluting both the primary & secondary antibodies.

T-bet Transcription Factor Assay

Determine the appropriate number of microwell strips required for testing samples, controls and blanks in duplicate. If less than 8 wells in a strip need to be used, cover the unused wells with a portion of the plate sealer while you perform the assay. The content of these wells is stable at room temperature if kept dry and, therefore, can be used later for a separate assay. Store the unused strips in the aluminum pouch at 4°C. Use the strip holder for the assay.

Prepare the Complete Lysis Buffer, Complete Binding Buffer, 1X Wash Buffer and 1X Antibody Binding Buffer as described above in the section Buffer Preparation and Recommendations. Multichannel pipettor reservoirs may be used for dispensing the Complete Binding Buffer, Wash Buffer, Antibody Binding Buffer, Developing Solution and Stop Solution into the wells being used.

Step 1: Binding of T-bet to its consensus sequence

- Add 40 µl Complete Binding Buffer to each well to be used. If you wish to perform competitive binding experiments, add 40 µl Complete Binding Buffer that contains 40 pmol (4 µl) of the wild-type or mutated oligonucleotide (see the Buffer Preparation section above for a description of competitive binding).
- Sample wells: Add 10 µl of sample diluted in Complete Lysis Buffer per well. We recommend
 using 2-10 µg of nuclear extract diluted in Complete Lysis Buffer per well. A protocol for
 preparing nuclear extracts is provided on page 11.
 - Positive control wells: Add 1.25 μ g of the provided nuclear extract diluted in 10 μ l of Complete Lysis Buffer per well (0.5 μ l of nuclear extract in 9.5 μ l of Complete Lysis Buffer per well).
 - Blank wells: Add 10 µl Complete Lysis Buffer only per well.
- 3. Use the provided adhesive cover to seal the plate. Incubate for 1 hour at room temperature with mild agitation (100 rpm on a rocking platform).
- 4. Wash each well 3 times with 200 μ l 1X Wash Buffer. For each wash, flick the plate over a sink to empty the wells, then tap the inverted plate 3 times on absorbent paper towels.

Step 2: Binding of primary antibody

- Add 100 μl diluted T-bet antibody (1:2000 dilution in 1X Antibody Binding Buffer) to each well being used.
- 2. Cover the plate and incubate for 1 hour at room temperature without agitation.
- 3. Wash the wells 3 times with 200 µl 1X Wash Buffer (as described in Step 1, No. 4).

Step 3: Binding of secondary antibody

- 1. Add 100 μ l of diluted HRP-conjugated antibody (1:1000 dilution in 1X Antibody Binding Buffer) to all wells being used.
- 2. Cover the plate and incubate for 1 hour at room temperature without agitation.
- 3. During this incubation, place the Developing Solution at room temperature.
- 4. Wash the wells 4 times with 200 µl 1X Wash Buffer (as described in Step 1, No. 4).

Step 4: Colorimetric reaction

- 1. Add 100 µl Developing Solution to all wells being used.
- Incubate 2-15 minutes at room temperature protected from direct light. Monitor the blue color development in the sample and positive control wells until it turns medium to dark blue. Do not overdevelop.
- 3. Add 100 µl Stop Solution. In presence of the acid, the blue color turns yellow.
- 4. Read absorbance on a spectrophotometer within 5 minutes at 450 nm with a reference wavelength of 655 nm. Blank the plate reader according to the manufacturer's instructions using the blank wells.

References

- 1. Beima, K.M., et al. (2006) J Biol. Chem, 281: 11992-12000.
- 2. Tong, Y., et al.. (2005) PNAS, 102: 2034-2039.
- 3. Lewis, M.D., et al. (2007) Mol Cell Biol, 27:8510-8521.

Appendix

Section A. Preparation of Nuclear Extract

For your convenience, Active Motif offers a Nuclear Extract Kit (Cat. Nos. 40010 & 40410). This kit contains buffers optimized for use in TransAM Kits, which serves to reduce inconsistencies in the assay that may arise from using homemade or other buffers. If you prefer to make your own buffers, please refer to the following protocol.

This procedure can be used for a confluent cell layer of 75 cm 2 (100-mm dish). The yield is approximately 0.15 mg of nuclear proteins for 9 x 10 6 cells.

- 1. Wash cells with 10 ml ice-cold PBS/PIB. Discard PBS/PIB.
- 2. Add 10 ml ice-cold PBS/PIB and scrape the cells off the dish with a cell lifter. Transfer cells into a pre-chilled 15 ml tube and spin at 300 x g for 5 minutes at 4°C.
- 3. Resuspend the pellet in 1 ml ice-cold HB buffer by gentle pipetting and transfer the cells into a pre-chilled 1.5 ml tube.
- 4. Allow the cells to swell on ice for 15 minutes.
- 5. Add 50 μ l 10% Nonidet P-40 (0.5 % final) and vortex the tube vigorously for 10 seconds.
- Centrifuge the homogenate for 30 seconds at 4°C in a microcentrifuge. Remove the supernatant (cytoplasmic fraction) and, if you wish to save this for other uses, transfer it into a pre-chilled microcentrifuge tube. (Store the cytoplasmic fraction at –80°C.)
- 7. Resuspend the nuclear pellet in 50 μl Complete Lysis Buffer and rock the tube gently on ice for 30 minutes on a shaking platform.
- 8. Centrifuge for 10 minutes at 14,000 x g at 4°C and save the supernatant (nuclear extract). Aliquot and store at -80°C. Avoid freeze/thaw cycles.
- 9. Determine the protein concentration of the extract by using a Bradford-based assay.

Preparation of Buffers for Nuclear Extract

10X PBS For 25	0 ml, mix:
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0.1 M phosphate buffer, pH 7.5 3.55 g Na, HPO₄ + 0.61 g KH, PO₄

1.5 M NaCl 21.9 g 27 mM KCl 0.5 g

Adjust to 250 ml with distilled water. Prepare a 1X PBS solution by adding 10 ml 10X PBS to 90 ml distilled water. Sterilize the 1X PBS by filtering through a 0.2 μ m filter. The 1X PBS is at pH 7.5. Store the filter-sterilized 1X PBS solution at 4°C.

PIB (Phosphatase Inhibitor Buffer) For 10 ml, mi	ix:
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125 mM NaF 52 mg
250 mM β-glycerophosphate 0.55 g
250 mM p-nitrophenyl phosphate (PNPP) 1.15 g
25 mM NaVO₃ 31 mg

Adjust to 10 ml with distilled water. Mix the chemicals by vortexing. Incubate the solution at 50° C for 5 minutes. Mix again. Store at -20° C.

PBS/PIB

Prior to use, add 0.5 ml PIB to 10 ml 1X PBS.

HB (Hypotonic Buffer)	For 50 ml, mix:
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20 mM Hepes, pH 7.5 0.24 g 5 mM NaF 12 mg

10 μ M Na₂MoO₄ 5 μ l of a 0.1 M solution
0.1 mM EDTA 10 μ l of a 0.5 M solution

Adjust pH to 7.5 with 1 N NaOH. Adjust volume to 50 ml with distilled water. Sterilize by filtering through a $0.2 \mu m$ filter. Store the filter-sterilized solution at $4^{\circ}C$.

Section B: Troubleshooting Guide

Problem/question	Possible cause	Recommendation
No signal or weak signal	Omission of key reagent	Check that all reagents have been added in all wells in the correct order
	Substrate or conjugate is no longer active	Test conjugate and substrate for activity
	Enzyme inhibitor present	Sodium azide will inhibit the peroxidase reaction, follow our recommendations to prepare buffers
	Plate reader settings not optimal	Verify the wavelength and filter settings in the plate reader
	Incorrect assay tem- perature	Bring substrate to room temperature
	Inadequate volume of Developing Solution	Check to make sure that correct volume is delivered by pipette
High background in all wells	Developing time too long	Stop enzymatic reaction as soon as the positive wells turn medium-dark blue
	Concentration of anti- bodies is too high	Increase antibody dilutions
	Inadequate washing	Ensure all wells are filled with Wash Buffer and follow washing recommendations
Uneven color develop- ment	Incomplete washing of wells	Ensure all wells are filled with Wash Buffer and follow washing recommendations
	Well cross-contami- nation	Follow washing recommendations
High background in sample wells	Too much nuclear extract per well	Decrease amount of nuclear extract down to 1-2 μg/well
	Concentration of anti- bodies is too high	Perform antibody titration to determine optimal working concentration. Start using 1:2000 for primary antibody and 1:5000 for the secondary antibody. The sensitivity of the assay will be decreased
No signal or weak signal in sample wells	Not enough nuclear extract per well	Increase amount of nuclear extract to 50 µg/well
	T-bet is poorly activated or inactivated in nuclear fractions	Perform a time course for T-bet activation in the studied cell line
	Nuclear extracts are not from correct species	

Section C. Related Products

Transcription Factor ELISAs	Format	Catalog No.
TransAM™ AML-1/Runx1	1 x 96-well plate	47396
TransAM™ AML-3/Runx2	1 x 96-well plate	44496
TransAM™ AP-1 Family	2 x 96-well plates	44296
TransAM™ AP-1 c-Fos	1 x 96-well plate	44096
TransAM™ AP-1 c-lun	1 x 96-well plate	46096
TransAM™ AP-1 FosB	1 x 96-well plate	45096
TransAM™ AP-1 JunD	1 x 96-well plate	43496
TransAM™ ATF-2	1 x 96-well plate	42396
TransAM™ c-Myc	1 x 96-well plate	43396
TransAM™ C/EBP α/β	1 x 96-well plate	44196
TransAM™ CREB	1 x 96-well plate	42096
TransAM™ pCREB	1 x 96-well plate	43096
TransAM™ Elk-1	1 x 96-well plate	44396
TransAM™ ER	1 x 96-well plate	41396
TransAM™ FKHR (FOXO1/4)	1 x 96-well plate	46396
TransAM™ GATA Family	2 x 96-well plates	48296
TransAM™ GATA-4	1 x 96-well plate	46496
TransAM™ GR	1 x 96-well plate	45496
TransAM™ HIF-1	1 x 96-well plate	47096
TransAM™ HNF Family	2 x 96-well plates	46296
TransAM™ HNF-1	1 x 96-well plate	46196
TransAM™ IRF-3 (Human)	1 x 96-well plate	48396
TransAM™ IRF-3 (Mouse)	1 x 96-well plate	48496
TransAM™ IRF-7	1 x 96-well plate	50196
TransAM™ MAPK Family	2 x 96-well plates	47296
TransAM™ MEF2	1 x 96-well plate	43196
TransAM™ MyoD	1 x 96-well plate	47196
TransAM™ NF-YA	1 x 96-well plate	40396
TransAM™ NFATc1	1 x 96-well plate	40296
TransAM™ NFKB Family	2 x 96-well plates	43296
TransAM™ Flexi NFKB Family	2 x 96-well plates	43298
TransAM™ NFκB p50	1 x 96-well plate	41096
TransAM™ Flexi NFκB p50	1 x 96-well plate	41098
TransAM™ NFκB p50 Chemi	1 x 96-well plate	41097
TransAM™ NFκB p52	1 x 96-well plate	48196
TransAM™ NFκB p52 Chemi	1 x 96-well plate	48197
•	•	40096
TransAM™ NFκB p65 TransAM™ Flexi NFκB p65	1 x 96-well plate 1 x 96-well plate	40098
TransAM™ NFKB p65 Chemi	1 x 96-well plate	40098
TransAM™ Oct-4	1 x 96-well plate	41196
TransAM™ p53	1 x 96-well plate	42496
TransAM™ PPARγ	1 x 96-well plate	40196
•	•	41296
TransAM™ Sp1	1 x 96-well plate	
TransAM™ Sp1/Sp3 TransAM™ STAT Family	1 x 96-well plate	40496 42206
TransAM™ STAT Family TransAM™ STAT3	2 x 96-well plates	42296
Halisaivi SIAIS	1 x 96-well plate	45196

Chromatin Assembly	Format	Catalog No.
Chromatin Assembly Kit	10 rxns	53500
Histone Acetyltransferase Activity	Format	Catalog No.
HAT Assay Kit (Fluorescent)	1 x 96 rxns	56100
Histone Purification	Format	Catalog No.
Histone Purification Kit	10 rxns	40025
Co-Immunoprecipitation	Format	Catalog No.
Nuclear Complex Co-IP Kit	50 rxns	54001
Universal Magnetic Co-IP Kit	25 rxns	54002
DNA Methylation	Format	Catalog No.
MethylDetector™	50 rxns	55001
MethylCollector™	25 rxns	55002
SUMOylation	Format	Catalog No.
SUMOlink™ SUMO-1 Kit	20 rxns	40120
SUMOlink™ SUMO-2/3 Kit	20 rxns	40220

Technical Services

If you need assistance at any time, please call Active Motif Technical Service at one of the numbers listed below.

Active Motif North America

1914 Palomar Oaks Way, Suite 150

Carlsbad, CA 92008

USA

Toll Free: 877 222 9543
Telephone: 760 431 1263
Fax: 760 431 1351

E-mail: tech service@activemotif.com

Active Motif Europe

104 Avenue Franklin Roosevelt

B-1330 Rixensart, Belgium

UK Free Phone: 0800 169 31 47
France Free Phone: 0800 90 99 79
Germany Free Phone: 0800 181 99 10
Telephone: +32 (0)2 653 0001
Fax: +32 (0)2 653 0050

E-mail: eurotech@activemotif.com

Active Motif Japan

 $Azuma\ Bldg, 7th\ Floor$

2-21 Ageba-Cho, Shinjuku-Ku

Tokyo, 162-0824, Japan

Telephone: +81 3 5225 3638 Fax: +81 3 5261 8733

E-mail: japantech@activemotif.com

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