

DATA NOTE

A guide to selecting high-performing antibodies for TAF15

(UniProt ID: Q92804) for use in western blot,

immunoprecipitation, and immunofluorescence

[version 1; peer review: 2 approved]

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Abstract

TAF15 (TATA-box binding protein-associated factor 15) is a member of the FET protein family, known for their roles in transcriptional regulation and RNA metabolism. Here we have characterized five TAF15 commercial antibodies for western blot, immunoprecipitation, and immunofluorescence using a standardized experimental protocol based on comparing read-outs in knockout cell lines and isogenic parental controls. These studies are part of a larger, collaborative initiative seeking to address antibody reproducibility issues by characterizing commercially available antibodies for human proteins and publishing the results openly as a resource for the scientific community. While use of antibodies and protocols vary between laboratories, we encourage readers to use this report as a guide to select the most appropriate antibodies for their specific needs.

Keywords

Q92804, TAF15, TATA-binding protein-associated factor 2N, TATA-box binding protein associated factor 15, antibody characterization, antibody validation, western blot, immunoprecipitation, immunofluorescence





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Any reports and responses or comments on the article can be found at the end of the article.

This article is included in the YCharOS (Antibody

Characterization through Open Science)

gateway.

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Competing interests: For this project, the authors developed partnerships with high-quality antibody manufacturers and KO cell line providers. The partners provide antibodies and KO cell lines at no cost. These partners include: - Abcam-ABCD antibodies- ABclonal-Aviva Systems Biology -Bio Techne -Cell Signalling Technology -Developmental Studies Hybridoma Bank -GeneTex – Horizon Discovery – Proteintech – Synaptic Systems –Thermo Fisher Scientific.

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Introduction

The FET family of RNA binding proteins includes TATA-binding protein-associated factor 2N (TAF15), Fused in Sarcoma (FUS)¹ and RNA binding protein EWS (EWS). TAF15 functions primarily as a transcriptional co-activator by interacting with components of the transcription machinery, including RNA polymerase II and other transcription factors, to regulate gene expression.² TAF15 contains RNA-binding motifs, enabling it to influence RNA splicing, transport, and stability.³ Dysregulation or mutations in TAF15 have been implicated in certain cancers and neurodegenerative diseases, highlighting its critical role in cellular homeostasis.⁴

This research is part of a broader collaborative initiative in which academics, funders and commercial antibody manufacturers are working together to address antibody reproducibility issues by characterizing commercial antibodies for human proteins using standardized protocols, and openly sharing the data.^{5–7} Here we evaluated the performance of five commercial antibodies for TAF15 for use in western blot, immunoprecipitation, and immunofluorescence, enabling biochemical and cellular assessment of TAF15 properties and function. The platform for antibody characterization used to carry out this study was endorsed by a committee of industry and academic representatives.⁸ It consists of identifying human cell lines with adequate target protein expression and the development/contribution of equivalent knockout (KO) cell lines, followed by antibody characterization procedures using most commercially available antibodies against the corresponding protein. The standardized consensus antibody characterization protocols are openly available on Protocol Exchange, a preprint server (DOI: 10.21203/rs.3.pex-2607/v1).⁹

The authors do not engage in result analysis or offer explicit antibody recommendations. Our primary aim is to deliver top-tier data to the scientific community, grounded in Open Science principles. This empowers experts to interpret the characterization data independently, enabling them to make informed choices regarding the most suitable antibodies for their specific experimental needs. Guidelines on how to interpret antibody characterization data found in this study are featured on the YCharOS gateway.¹⁰

Results and discussion

Our standard protocol involves comparing readouts from wild type (WT) and KO cells.^{11,12} The first step was to identify a cell line(s) that expresses sufficient levels of a given protein to generate a measurable signal using antibodies. To this end, we examined the DepMap (Cancer Dependency Map Portal, RRID:SCR_017655) transcriptomics database to identify all cell lines that express the target at levels greater than $2.5 \log_2$ (transcripts per million "TPM" + 1), which we have found to be a suitable cut-off.⁵ The HAP1 expresses the TAF15 transcript at 7.9, which is above the average range of cancer cells analyzed. *TAF15* KO in HAP1 cells were obtained from Horizon Discovery (Table 1).

To screen all antibodies by western blot, WT and *TAF15* KO protein lysates were ran on SDS-PAGE, transferred onto nitrocellulose membranes, and then probed with the five TAF15 antibodies in parallel (Figure 1).

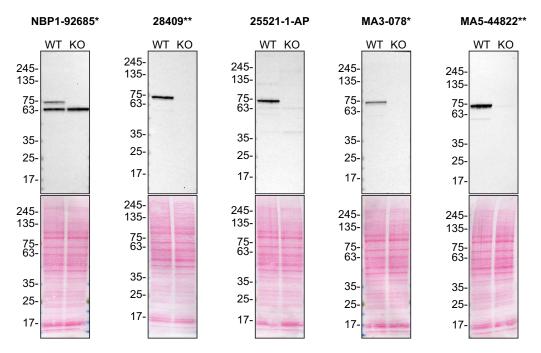
We then assessed the capability of all five antibodies to capture TAF15 from HAP1 protein extracts using immunoprecipitation techniques, followed by western blot analysis. For the immunoblot step, a specific TAF15 antibody identified previously (refer to Figure 1) was selected. Equal amounts of the starting material (SM) and the unbound fractions (UB), as well as the whole immunoprecipitate (IP) eluates were separated by SDS-PAGE (Figure 2).

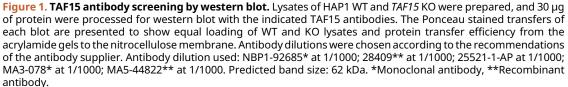
For immunofluorescence, the five antibodies were screened using a mosaic strategy. First, HAP1 WT and *TAF15* KO cells were labelled with different fluorescent dyes in order to distinguish the two cell lines, and the TAF15 antibodies were evaluated. Both WT and KO lines imaged in the same field of view to reduce staining, imaging and image analysis bias (Figure 3). Quantification of immunofluorescence intensity in hundreds of WT and KO cells was performed for each antibody tested, and the images presented in Figure 3 are representative of this analysis.⁹

In conclusion, we have screened five TAF15 commercial antibodies by western blot, immunoprecipitation, and immunofluorescence by comparing the signal produced by the antibodies in human HAP1 WT and *TAF15* KO cells. To assist users in interpreting antibody performance, Table 3 outlines various scenarios in which antibodies may

Institution	Catalog number	RRID (Cellosaurus)	Cell line	Genotype
Horizon Discovery	C631	CVCL_Y019	HAP1	WT
Horizon Discovery	HZGHC004653c001	CVCL_TR46	HAP1	TAF15 KO

Table 1. Summary of the cell lines used.





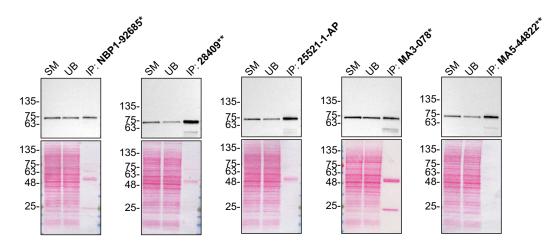


Figure 2. TAF15 antibody screening by immunoprecipitation. HAP1 lysates were prepared, and immunoprecipitation was performed using 1 mg of lysate and 2.0 µg of the indicated TAF15 antibodies pre-coupled to Dynabeads protein A or protein G. Samples were washed and processed for western blot with the indicated TAF15 antibody. For western blot, 28409** was used at 1/1000. The Ponceau stained transfers of each blot are shown. SM=4% starting material; UB=4% unbound fraction; IP=immunoprecipitate. *Monoclonal antibody, **Recombinant antibody.

perform in all three applications.⁵ Several high-quality and renewable antibodies that successfully detect TAF15 were identified in all applications. Researchers who wish to study TAF15 in a different species are encouraged to select high-quality antibodies, based on the results of this study, and investigate the predicted species reactivity of the manufacturer before extending their research.

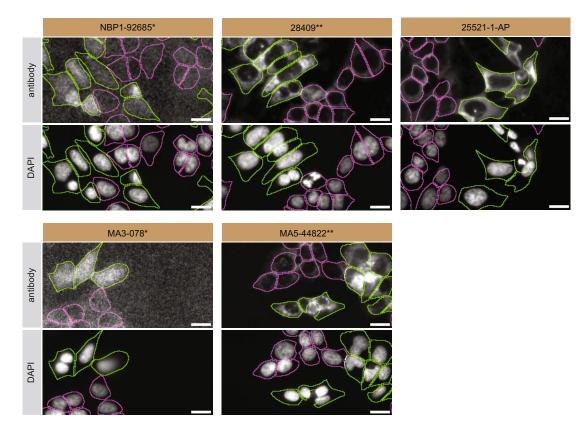


Figure 3. TAF15 antibody screening by immunofluorescence. HAP1 WT and *TAF15* KO cells were labelled with a green or a far-red fluorescent dye, respectively. WT and KO cells were mixed and plated to a 1:1 ratio on coverslips. Cells were stained with the indicatedTAF15 antibodies and with the corresponding Alexa-fluor 555 coupled secondary antibody including DAPI. Acquisition of the blue (nucleus-DAPI), green (WT), red (antibody staining) and far-red (KO) channels was performed. Representative images of the merged blue and red (grayscale) channels are shown. WT and KO cells are outlined with green and magenta dashed line, respectively. When an antibody was recommended for immunofluorescence by the supplier, we tested it at the recommended dilution. The rest of the antibodies were tested at 1 and 2 µg/ml and the final concentration was selected based on the detection range of the microscope used and a quantitative analysis not shown here. Antibody dilution used: NBP1-92685* at 1/1000; 28409** at 1/150; 25521-1-AP at 1/50; MA3-078* at 1/500; MA5-44822** at 1/1000. Bars = 10 µm. *Monoclonal antibody, **Recombinant antibody.

The underlying data listed below for this study can be found on Zenodo, an open-access repository for which YCharOS has its own collection of antibody characterization reports.

Limitations

Inherent limitations are associated with the antibody characterization platform used in this study. Firstly, the YCharOS project focuses on renewable (recombinant and monoclonal) antibodies and does not test all commercially available TAF15 antibodies. YCharOS partners provide approximately 80% of all renewable antibodies, but some top-cited polyclonal antibodies may not be available through these partners.

Secondly, the YCharOS effort employs a non-biased approach that is agnostic to the protein for which antibodies have been characterized. The aim is to provide objective data on antibody performance without preconceived notions about how antibodies should perform or the molecular weight that should be observed in western blot. As the authors are not experts in TAF15 only a brief overview of the protein's function and its relevance in disease is provided. TAF15 experts are invited to analyze and interpret observed banding patterns in western blots and subcellular localization in immunofluorescence.

Thirdly, YCharOS experiments are not performed in replicates primarily due to the use of multiple antibodies targeting various epitopes. Once a specific antibody is identified, it validates the protein expression of the intended target in the selected cell line, confirms the lack of protein expression in the KO cell line and supports conclusions regarding the

specificity of the other antibodies. All experiments are performed using master mixes, and meticulous attention is paid to sample preparation and experimental execution. In IF, the use of two different concentrations serves to evaluate antibody specificity and can aid in assessing assay reliability. In instances where antibodies yield no signal, a repeat experiment is conducted following titration. Additionally, our independent data is performed subsequently to the antibody manufacturers internal validation process, therefore making our characterization process a repeat.

Lastly, as comprehensive and standardized procedures are respected, any conclusions remain confined to the experimental conditions and cell line used for this study. The use of a single cell type for evaluating antibody performance poses as a limitation, as factors such as target protein abundance significantly impact results.⁹ Additionally, the use of cancer cell lines containing gene mutations poses a potential challenge, as these mutations may be within the epitope coding sequence or other regions of the gene responsible for the intended target. Such alterations can impact the binding affinity of antibodies. This represents an inherent limitation of any approach that employs cancer cell lines.

Method

The standardized protocols used to carry out this KO cell line-based antibody characterization platform was established and approved by a collaborative group of academics, industry researchers and antibody manufacturers.⁸ The detailed materials and step-by-step protocols used to characterize antibodies in western blot, immunoprecipitation and immuno-fluorescence are openly available on Protocol Exchange, a preprint server (DOI: 10.21203/rs.3.pex-2607/v1).⁹ Brief descriptions of the experimental setup used to carry out this study can be found below.

Cell lines and antibodies

Cell lines used and primary antibodies tested in this study are listed in Tables 1 and 2, respectively. To ensure that the cell lines and antibodies are cited properly and can be easily identified, we have included their corresponding Research Resource Identifiers, or RRID.^{13,14}

Peroxidase-conjugated goat anti-rabbit and anti-mouse antibodies are (Thermo Fisher Scientific, cat. number 65-6120 and 62-6520). Alexa-555-conjugated goat anti-rabbit and anti-mouse secondary antibodies (Thermo Fisher Scientific, cat. number A-21429 and A-21424). Peroxidase-conjugated Protein A for IP detection is from MilliporeSigma, cat. number P8651.

Antibody screening by western blot

HAP1 WT and *TAF15* KO cells were collected in RIPA buffer (25 mM Tris-HCl pH 7.6, 150 mM NaCl, 1% NP-40, 1% sodium deoxycholate, 0.1% SDS) (Thermo Fisher Scientific, cat. number 89901) supplemented with $1 \times$ protease inhibitor cocktail mix (MilliporeSigma, cat. number P8340). Lysates were sonicated briefly and incubated 30 min on ice. Lysates were spun at ~110,000 × g for 15 min at 4°C and equal protein aliquots of the supernatants were analyzed by SDS-PAGE and western blot. BLUelf prestained protein ladder (GeneDireX, cat. number PM008-0500) was used.

Western blots were performed with precast midi 4-20% Tris-Glycine polyacrylamide gels (Thermo Fisher Scientific, cat. number WXP42012BOX) ran with Tris/Glycine/SDS buffer (Bio-Rad, cat. number 1610772), loaded in Laemmli loading sample buffer (Thermo Fisher Scientific, cat. number AAJ61337AD) and transferred on nitrocellulose membranes. Proteins on the blots were visualized with Ponceau S staining (Thermo Fisher Scientific, cat. number BP103-10) which is scanned to show together with individual western blot. Blots were blocked with 5% milk for 1 hr, and antibodies were incubated O/N at 4°C with 5% milk in TBS with 0.1% Tween 20 (TBST) (Cell Signalling Technology, cat. number 9997). Following three washes with TBST, the peroxidase conjugated secondary antibody was incubated at a dilution of ~0.2 μ g/ml in TBST with 5% milk for 1 hr at room temperature followed by three washes with TBST. Membranes were incubated with Pierce ECL (Thermo Fisher Scientific, cat. number 32106) prior to detection with the iBrightTM CL1500 Imaging System (Thermo Fisher Scientific, cat. number A44240).

Antibody screening by immunoprecipitation

Antibody-bead conjugates were prepared by adding 2 μ g of antibody to 500 μ l of Pierce IP Lysis Buffer from Thermo Fisher Scientific (cat. number 87788) in a microcentrifuge tube, together with 30 μ l of Dynabeads protein A- (for rabbit antibodies) or protein G- (for mouse antibodies) (Thermo Fisher Scientific, cat. number 10002D and 10004D, respectively). 10 μ l of antibodies 28409** and MA3-078* were used in the IP. Tubes were rocked for ~1 hr at 4°C followed by two washes to remove unbound antibodies.

HAP1 WT were collected in Pierce IP buffer (25 mM Tris-HCl pH 7.4, 150 mM NaCl, 1 mM EDTA, 1% NP-40 and 5% glycerol) supplemented with protease inhibitor. Lysates were rocked 30 min at 4°C and spun at 110,000 × g for 15 min at 4°C. 0.5 ml aliquots at 2 mg/ml of lysate were incubated with an antibody-bead conjugate for ~1 h at 4°C. The unbound

Company	Catalog number	Lot number	RRID (Antibody Registry)	Clonality	Clone ID	Host	Concentration (µg/µL)	Vendors recommended applications
Bio-Techne	NBP1-92685*	102313	AB_11037253	monoclonal	4D71	mouse	1	Wb, IF
Cell Signaling Technology	28409 **	1	AB_2798957	recombinant mono	D8V6Q	rabbit	0.146	Wb, IP
Proteintech	25521-1-AP	41674	AB_2880116	polyclonal		rabbit	2.66	Wb
Thermo Fisher Scientific	MA3-078*	WL343399	AB_2633323	monoclonal	8TA-2B10	mouse	n/a	Wb, IP, IF
Thermo Fisher Scientific	MA5-44822 **	YE3913388B	AB_2931279	recombinant mono	JE61-92	rabbit	1	Wb, IF
Wb=western blot; IF=immunofluorescence; IP=immunoprecipitation.	escence; IP=immunopr	ecipitation.						

Table 2. Summary of the TAF15 antibodies tested.

Wb=western ມາບ, າ. ... *Monoclonal antibody. **Recombinant antibody, n/a=not available.

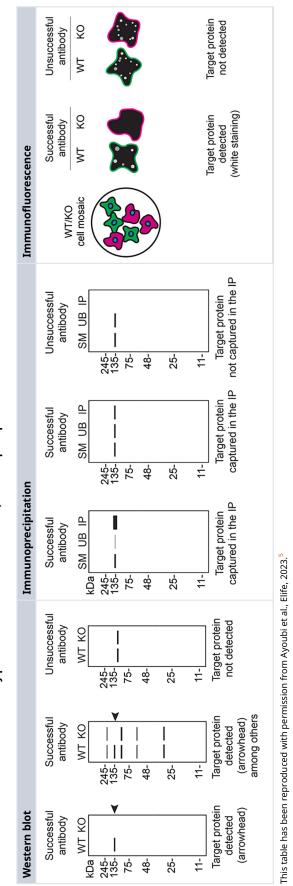


Table 3. Illustrations to assess antibody performance in all western blot, immunoprecipitation and immunofluorescence.

fractions were collected, and beads were subsequently washed three times with 1.0 ml of IP buffer and processed for SDS-PAGE and western blot on precast midi 4-20% Tris-Glycine polyacrylamide gels. Protein A:HRP was used as a secondary detection system at a concentration of $0.3 \mu g/ml$.

Antibody screening by immunofluorescence

HAP1 WT and *TAF15* KO cells were labelled with a green and a far-red fluorescence dye, respectively (Thermo Fisher Scientific, cat. number C2925 and C34565). The nuclei were labelled with DAPI (Thermo Fisher Scientific, cat. Number D3571) fluorescent stain. WT and KO cells were plated on 96-well plate with optically clear flat-bottom (Perkin Elmer, cat. number 6055300) as a mosaic and incubated for 24 hrs in a cell culture incubator at 37° C, 5% CO₂. Cells were fixed in 4% paraformaldehyde (PFA) (VWR, cat. number 100503-917) in phosphate buffered saline (PBS) (Wisent, cat. number 311-010-CL). Cells were permeabilized in PBS with 0,1% Triton X-100 (Thermo Fisher Scientific, cat. number 16210-064) and 0.01% Triton X-100 for 30 min at room temperature. Cells were incubated with IF buffer (PBS, 5% BSA, 0.01% Triton X-100) containing the primary TAF15 antibodies overnight at 4°C. Cells were then washed 3×10 min with IF buffer and incubated with corresponding Alexa Fluor 555-conjugated secondary antibodies in IF buffer at a dilution of 1.0 µg/ml for 1 hr at room temperature with DAPI. Cells were washed 3×10 min with IF buffer and once with PBS.

Images were acquired on an ImageXpress micro confocal high-content microscopy system (Molecular Devices), using a 20x NA 0.95 water immersion objective and scientific CMOS cameras, equipped with 395, 475, 555 and 635 nm solid state LED lights (lumencor Aura III light engine) and bandpass filters to excite DAPI, Cellmask Green, Alexa-555 and Cellmask Red, respectively. Images had pixel sizes of 0.68 x 0.68 microns, and a z-interval of 4 microns. For analysis and visualization, shading correction (shade only) was carried out for all images. Then, maximum intensity projections were generated using 3 z-slices. Segmentation was carried out separately on maximum intensity projections of Cellmask channels using CellPose 1.0, and masks were used to generate outlines and for intensity quantification.¹⁵ Figures were assembled with Adobe Illustrator.

Data availability

Underlying data

Zenodo: Antibody Characterization Report for TAF15, https://doi.org/10.5281/zenodo.10085319.16

Zenodo: Dataset for the TAF15 antibody screening study, https://doi.org/10.5281/zenodo.1453680517

Data are available under the terms of the Creative Commons Attribution 4.0 International license (CC-BY 4.0).

Acknowledgment

We would like to thank the NeuroSGC/YCharOS/EDDU collaborative group for their important contribution to the creation of an open scientific ecosystem of antibody manufacturers and KO cell line suppliers, for the development of community-agreed protocols, and for their shared ideas, resources, and collaboration. Members of the group can be found below. We would also like to thank the Advanced BioImaging Facility (ABIF) consortium for their image analysis pipeline development and conduction (RRID:SCR_017697). Members of each group can be found below.

NeuroSGC/YCharOS/EDDU collaborative group: Thomas M. Durcan, Aled M. Edwards, Peter S. McPherson, Chetan Raina and Wolfgang Reintsch

ABIF consortium: Claire M. Brown and Joel Ryan

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Sian E. Evans

University of Leicester, Leicester, England, UK **Carolyn Jones** University of Leicester, Leicester, England, UK

This article offers a clear and straightforward demonstration of the specificity of several antibodies targeting the protein TAF15. It utilises three techniques—Western Blotting (WB), Immunoprecipitation (IP), and Immunofluorescence (IF)—with all protocols readily accessible for researchers to replicate. Table 3 presents example results, which are incredibly helpful for interpreting the data. Finally, adding a limitations section provides valuable insights to inspire further reflection during the planning and execution of such experiments.

Is the rationale for creating the dataset(s) clearly described?

Yes

Are the protocols appropriate and is the work technically sound?

Yes

Are sufficient details of methods and materials provided to allow replication by others? $\ensuremath{\mathsf{Yes}}$

Are the datasets clearly presented in a useable and accessible format?

Yes

Competing Interests: No competing interests were disclosed.

Reviewer Expertise: Cell culture (primary cells and cell lines), patient-derived explants (PDE), Western blotting, Immunohistochemistry (IHC), immunofluorescence (IF), qPCR, RNA/DNA/protein purification We confirm that we have read this submission and believe that we have an appropriate level of expertise to confirm that it is of an acceptable scientific standard.

Reviewer Report 01 April 2025

https://doi.org/10.5256/f1000research.176259.r370781

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Deborah Moshinsky ២

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The authors describe Western Blot, Immunoprecipitation, and Immunofluorescence testing results for 5 commercially available antibodies against TATA-box binding protein-associated factor 15 (TAF15). Clear methods were given and referenced, and the results were clearly delineated in figures and tables. A cartoon figure was also shown to assist in result interpretation.

Brief information was given about the biology of the TAF15. The authors adequately describe that they are not experts in the biology of the target, so this is understandable. However, in light of the immunofluorescence data where some antibodies are detecting a cytoplasmic protein and others a protein in the nucleus, a mention about the expected cellular localization of the target would have been welcome, or information about the current thoughts in the field about localization.

Is the rationale for creating the dataset(s) clearly described?

Yes

Are the protocols appropriate and is the work technically sound?

Yes

Are sufficient details of methods and materials provided to allow replication by others? $\ensuremath{\mathsf{Yes}}$

Are the datasets clearly presented in a useable and accessible format?

Yes

Competing Interests: No competing interests were disclosed.

Reviewer Expertise: Antibody characterization and validation

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