

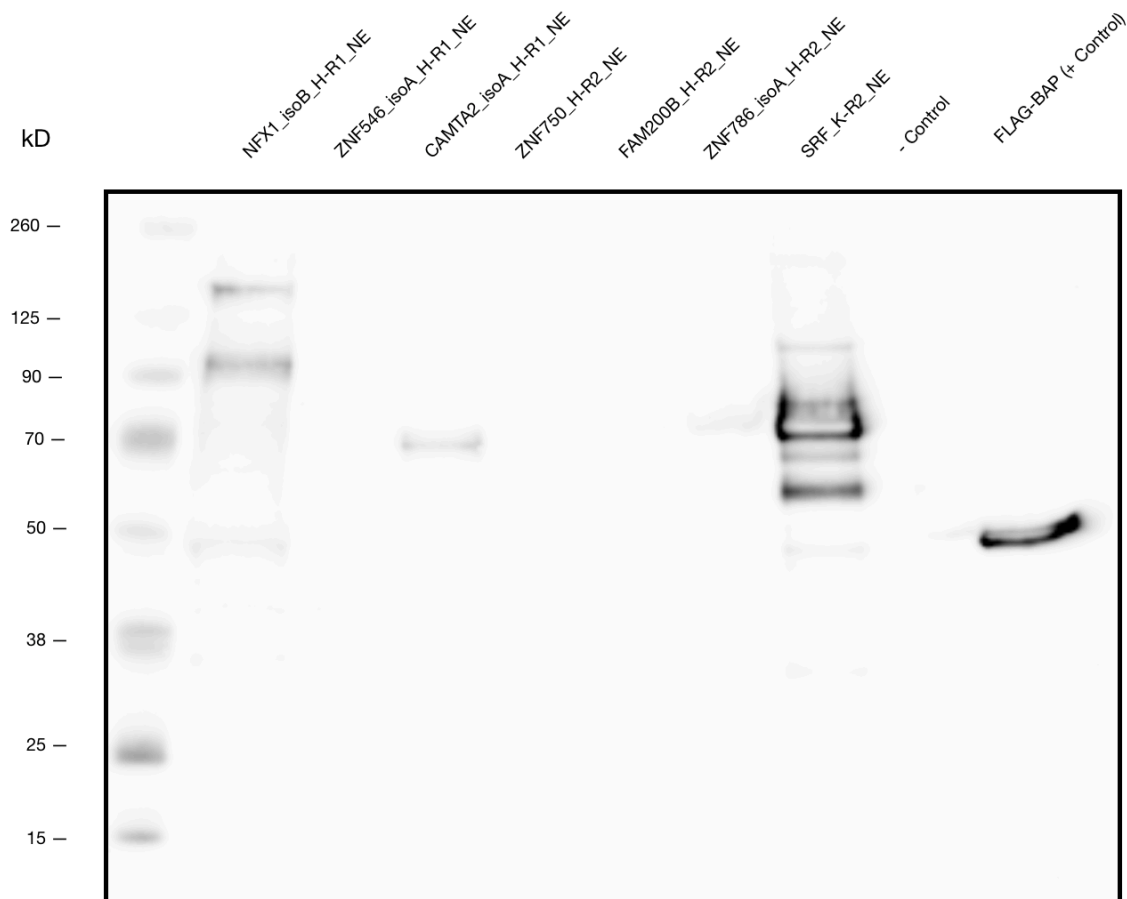
NFX1 (*Homo sapiens*), ZNF546 (*Homo sapiens*), CAMTA2 (*Homo sapiens*), ZNF750 (*Homo sapiens*), FAM200B (*Homo sapiens*), ZNF786 (*Homo sapiens*), and SRF (*Homo sapiens*)

Method:

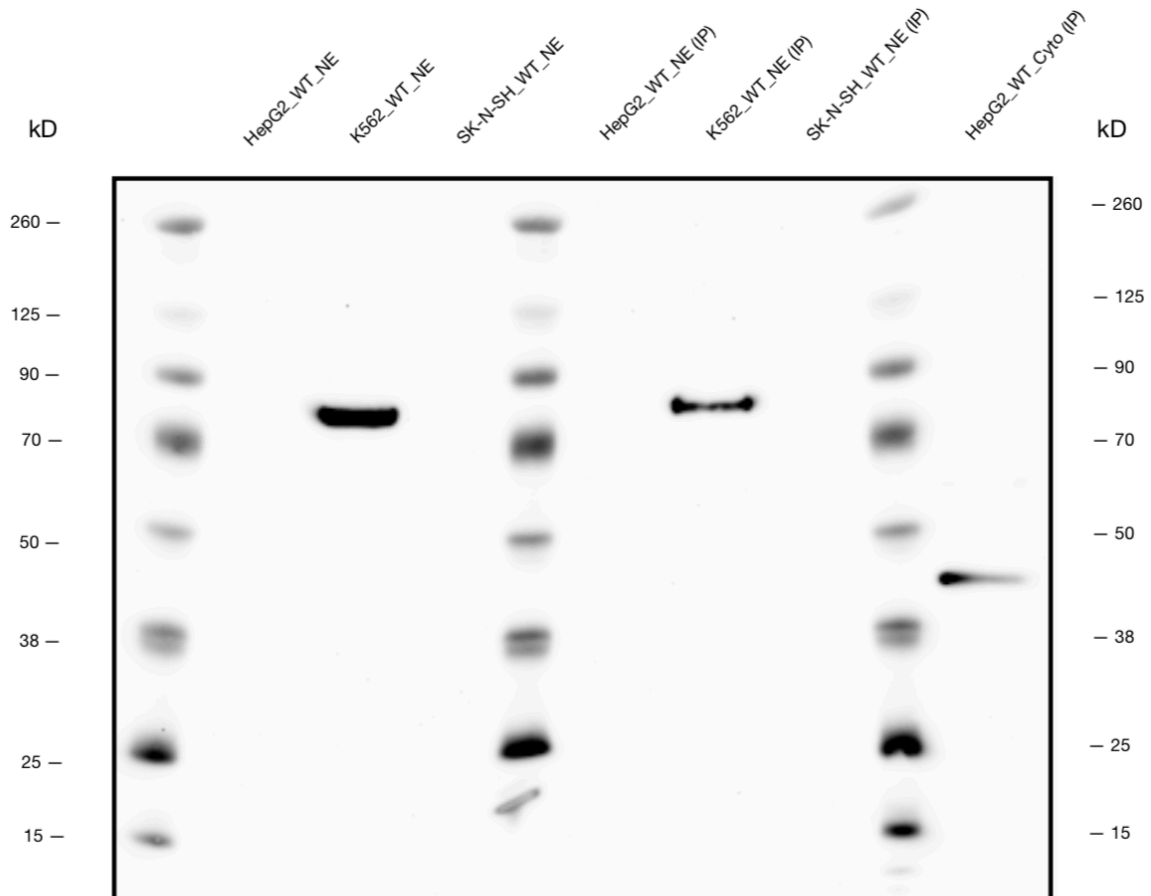
Western Blot Validation

Caption:

Each FLAG-tagged sample was immunoprecipitated from its corresponding nuclear protein isolate (500 uL) using the FLAG Immunoprecipitation Kit (Sigma-Aldrich; cat# FLAGIPT1). The final elution step was performed by suspending the sample-bound resin in NuPage Sample Reducing Agent 10X and NuPage LDS Sample Buffer 4X (Thermo Fisher Scientific) and heating for 3 minutes at 90C. Followed by cooling on ice, the protein samples were loaded onto a NuPage 4-12% Bis-Tris gel (Thermo Fisher Scientific) and separated using a PowerEase 90W system (Thermo Fisher Scientific) running at 150 V for 1 hour. A blank IP was included as a negative control, and an immunoprecipitated FLAG-BAP fusion protein provided in the kit as a positive control. The protein bands were transferred to a nitrocellulose membrane using the Invitrogen iBlot 2 System (Thermo Fisher Scientific), and blocked overnight at 4C in 5% milk solution with gentle rocking. The membrane was treated with a 1:5000 dilution of monoclonal M2-Peroxidase-conjugated ANTI-FLAG antibody (diluted in 5% BSA solution) (Sigma-Aldrich; cat# A8592) for 1 hour. Following four 5-minute washes with 1X TBST, visualization was attained with the Super Signal West Femto solution kit (Thermo Fisher Scientific) and a MyECL Imager (Thermo Fisher Scientific). The second western blot image depicts negative control IPs prepared with HepG2 nuclear lysate (Lane 6) and K562 nuclear lysate (Lane 7).



Lane	Loaded Sample	Expected Band Size (kDa)	Comments
1	Ladder	N/A	N/A
2	FLAG-NFX1_isoB_HepG2 rep 1 (nuclear extract)	127	Distinct band around 100 kDa. Found comparable western image with banding around 120 kDa, within 20% of the observed band: https://www.proteinatlas.org/ENSG00000086102-NFX1/antibody#western_blot . PTMs: Phosphorylation and Ubl conjugation
3	FLAG-ZNF546_isoA_HepG2 rep 1 (nuclear extract)	101	No visible banding. PTMs: Acetylation, Isopeptide bonding and Ubl conjugation
4	FLAG-CAMTA2_isoA_HepG2 rep 1 (nuclear extract)	135	Single distinct band further than 20% from the expected size
5	FLAG-ZNF750_HepG2 rep 2 (nuclear extract)	80	No visible banding
6	FLAG-FAM200B_HepG2 rep 2 (nuclear extract)	79	No visible banding
7	FLAG-ZNF786_isoA_HepG2 rep 2 (nuclear extract)	93	Single faint band around 70 kDa. Found comparable western image with banding around 72 kDa, within 20% of the observed band: https://www.proteinatlas.org/ENSG00000197362-ZNF786/antibody#western_blot
8	FLAG-SRF_K562 rep 2 (nuclear extract)	55	Distinct band near the expected size. PTMs: Glycosylation and Phosphorylation
9	FLAG-BAP (Positive Control)	49	Dark band around 50 kDa
10	Negative Control (Blank IP)	None	No visible banding



Lane	Loaded Sample	Expected Band Size (kDa)	Comments
1	Ladder	N/A	N/A
2	HepG2 Wild-Type (nuclear extract)	None	No visible banding
3	K562 Wild-Type (nuclear extract)	None	Single non-distinct band at around 80 kDa
4	SK-N-SH Wild-Type (nuclear extract)	None	No visible banding
5	Ladder	N/A	N/A
6	HepG2 Wild-Type (nuclear extract IP)	None	No visible banding
7	K562 Wild-Type (nuclear extract IP)	None	Single non-distinct band at around 80 kDa
8	SK-N-SH Wild-Type (nuclear extract)	None	No visible banding
9	Ladder	N/A	N/A
10	HepG2 Wild-Type (cytoplasmic extract IP)	None	Single non-distinct band at around 45 kDa

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Grant:

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