Supplemental Table. The Thyroid hormone Receptor Interacting Proteins family and their functional roles are presented except for TRIP12, which is the topic of this review.

	Uniprot Human	Alternative Short Name	Alternative Name	Function
TRIP1	P62195	PRS8 P45/SUG1	26S proteasome regulatory subunit 8, 26S proteasome AAA-ATPase subunit RPT6, Proteasome 26S subunit ATPase 5. Proteasome subunit p45	Transcriptional activator [1–4] p45 ATPase subunit of PA700, the regulatory complex of the 26S proteasome [5,6].
TRIP2	Q15648	MED1	Mediator of RNA polymerase II transcription subunit 1	Component of the Mediator complex, a coactivator involved in the regulated transcription of nearly all RNA polymerase II-dependent genes [7].
		TRAP220	Thyroid hormone receptor-associated protein complex 220 kDa component	Interaction with nuclear receptors in a ligand-dependent manner, global activator for the nuclear receptor superfamily [8–10].
		DRIP205	Vitamin D receptor-interacting protein complex component	
		ARC205	Activator-recruited cofactor 205 kDa component	
		CRSP200	Cofactor Required for Sp1	Subunit p200 of the transcriptional factor CRSP is identical to TRIP2 [11].
		PBP, PPAR-binding protein	Peroxisome proliferator-activated receptor-binding protein	Interaction with peroxisome proliferator-activated receptor [12].
		RB18A	Recognized by PAb1801 antibody	Identification of common antigenic and functional properties with P53 [13].
TRIP3	Q15649	ZNHIT3	ZNHI3 Zinc finger HIT domain-containing protein 3	The heterodimer NUFIP1 (nuclear fragile X mental retardation-interacting protein 1)-ZNHIT3 is implicated in the assembly of the box C/D small nucleolar ribonucleoprotein (snoRNP) [14] and in starvation-induced ribophagy [15].
		HNF-4a coactivator	Hepatocyte Nuclear Factor 4a coactivator	[16]
TRIP4	Q15650	ASC-1	Activating signal cointegrator 1	Transcription coactivator that associates with nuclear receptors, transcriptional coactivators and basal transcription factors [17]. Component of the human RQC-trigger (hRQT) complex that facilitates ribosome-associated quality control [18].
TRIP5	P52732	KIF11 HKSP Eg5	Kinesin-like protein KIF11 Kinesin-like spindle protein	Motor protein required for establishing a bipolar spindle during mitosis [19,20]. Required in non-mitotic cells for transport of secretory proteins from the Golgi complex to the cell surface [21]. Modulates neuronal migration [21,22].
TRIP6	Q15654	ZRP-1 OIP-1	Zyxin-related protein 1 Opa-interacting protein 1	Relays signals from the cell surface to the nucleus to weaken adherent junction and promote actin cytoskeleton reorganization and

				cell invasiveness, transcriptional coactivator for NF-kappa-B and JUN [23].
TRIP7	Q15651	HMGN3	High mobility group nucleosome-binding domain-containing protein 3	Binds to nucleosomes, regulating chromatin structure [24,25].
TRIP8	Q15652	JMJD1C	Probable JmjC domain-containing histone demethylation protein 2C	Probable histone demethylase that specifically demethylates Lysine 9 residue of histone H3 [26].
TRIP9	Q15653	NF-Kappa-BIB	NF-kappa-B inhibitor beta	Inhibits NF-kappa-B by complexing with and trapping it in the cytoplasm [27].
TRIP10	Q15642	CIP4 STP	Cdc42-interacting protein 4 Salt tolerant protein Protein Felic	Required for translocation of GLUT4 to the plasma membrane [28]. Required for membrane tubulation during endocytosis, the formation of podosomes. Promotes membrane invagination and the formation of tubules. Promotes CDC42-induced actin polymerization [29].
TRIP11	Q15643	GMAP-210 CEV14	Golgi-associated microtubule-binding protein 210 Clonal evolution-related gene on chromosome 14 protein	Is a membrane tether required for vesicle tethering to Golgi, essential role in the maintenance of Golgi structure and function [30].
TRIP12	Q14669			
TRIP13	Q15645	PCH2 16E1-BP	Pachytene checkpoint protein 2 homolog Human papillomavirus type 16 E1 protein-binding protein	Plays a key role in chromosome recombination and chromosome structure development during meiosis [31]. Promotes early steps of the DNA double-strand breaks (DSBs) repair process [32]. Plays a role in mitotic spindle assembly checkpoint activation [33].
TRIP14	Q15646	OASL 2′-5′-OAS-RP p59OASL	2′-5′-oligoadenylate synthase-like protein 2′-5′-OAS-related protein p59OASL 59 kDa 2′-5′-oligoadenylate synthase-like protein	Can bind double-stranded RNA. Displays antiviral activity against encephalomyocarditis and hepatitis C virus [34].
TRIP15	P61201	SGN2	COP9 signalosome complex subunit 2 Alien homolog JAB1-containing signalosome subunit 2	Essential component of the COP9 signalosome complex (CSN) [35].

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