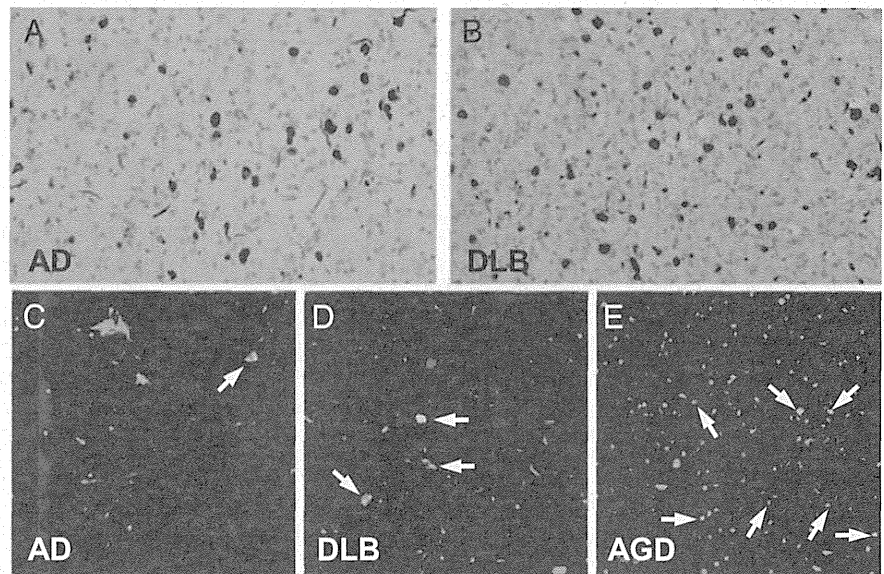


Fig. 3 Trans-activation response (TAR) DNA-binding protein of M_r 43 kDa (TDP-43)-positive structures in other neurodegenerative disorders. Neuronal cytoplasmic inclusions (NCIs) and short dystrophic neurites (DNs) stained with the pTDP-43-specific antibody (pS403/404) in the temporal cortex of the Alzheimer's disease (AD) case (A) and the dementia with Lewy bodies (DLB) case (B). (C–E) Double-label immunofluorescence histochemistry of the temporal cortex of AD (C) and DLB (D) and of the amygdala of argyrophilic grain disease (AGD) (E). The green fluorescence reveals the immunoreactivity for phosphorylated tau (AT8) in C and E, and that for phosphorylated alpha-synuclein in D, while the red fluorescence represents the immunopositivity for pS403/404 in C–E. Arrows indicate the colocalization of tau and pTDP-43 in C and E, and that of alpha-synuclein and pTDP-43 in D.



still unclear. A higher Braak NFT stage in the TDP-43 positive patients than in the TDP-43 negative ones was found in DLB+AD cases⁹¹ and in our study of AD cases.⁹⁵ We also reported parallel distribution of TDP-43 positive structures and tau positive grains and higher AGD stages in cases with TDP-43 immunoreactivity than in those without TDP-43 immunoreactivity in AGD.⁹⁷ Double-label immunofluorescence microscopy reveals partial colocalization of tau and TDP-43 in AD, DLB, AGD, Guamanian PDC and CBD^{83,84,86,87,90,91,97} or of α -synuclein and TDP-43 in DLB.^{87,91,95} These findings suggest that there may be common factors or mechanisms that affect the conformation or modification of these proteins, leading to their intracellular accumulation.

Regarding the typing of TDP-43 pathology, neocortical TDP-43 pathology in AD and DLB corresponded to type A.^{90,95,99} Immunoblot analyses of the sarkosyl insoluble fraction from AD and DLB cases with neocortical TDP-43 pathology also showed that the band pattern of these CTFs in AD and DLB corresponded to that of type A⁵⁸. These results suggest that the morphological and biochemical features of TDP-43 pathology are common between AD or DLB and a specific subtype of FTLD-TDP. Since all FTLD-TDP cases with *GRN* mutations show type A pathology,⁴⁵ there may be genetic factors, such as mutations or genetic variants of *GRN* underlying the co-occurrence of abnormal deposition of TDP-43, tau and α -synuclein. Indeed, recently, *GRN* loss-of-function mutation has been confirmed in patients clinically diagnosed with AD^{100–105} and Parkinson's disease.¹⁰⁶ The association between rs5848 variant in the 3' untranslated region of *GRN* and risk of AD has been reported in a Taiwanese population,¹⁰⁷ suggesting that homozygous TT genotype accentuates the risk

of AD. These findings suggest that PGRN reduction may induce both TDP-43 pathology and AD pathology.

Cell death and TDP-43 pathology

A report using a cell culture system showed that intracellular aggregate formation of TDP-43 induced cell death.⁷⁷ In brains of FTLD and ALS cases, basically, the occurrence of TDP-43-positive neuronal structures is related to degenerative changes.³⁶ However, the issue of the relation between the formation of TDP-43-positive inclusions and cell death may not be straightforward, since neuronal loss was not evident in the hippocampal granule cells³⁶ and the neostriatum¹⁰⁸ where TDP-43-positive structures were present in ALS cases. The reason for such a discrepancy between the results of the cell culture experiments and the findings of diseased brains should be discussed as a future issue.

CONCLUSION

The relevance of the pathological classification of TDP-43 proteinopathy is supported by clinical, biochemical and genetic correlations, although there is still highly significant heterogeneity in cases with type A pathology (Table 2). The results of the biochemical analyses of the diseased brains and the cellular models suggest that different strains of TDP-43 with different conformations may determine the clinicopathological phenotypes of TDP-43 proteinopathy, like prion disease. Detecting each TDP-43 strain in biological fluids may be useful for the differential diagnosis of TDP-43 proteinopathy. Furthermore, elucidating the mechanism of the conformational changes leading to the formation of multiple TDP-43 strains may be important for developing disease-modifying therapy for these diseases.

Table 2 Clinical, pathological and genetic associations of TDP-43 proteinopathy and other neurodegenerative disorders

Pathological subtype	Associated genes	Clinical phenotypes				Other diseases	
		bvFTD	PNFA	SD			MND
				Semantic aphasia	Prosopagnosia		
A	GRN, C9ORF72	+	+			PLS, ALS	
B	C9ORF72	+	+			ALS	
C				+			
D	VCP	+			+	PLS ALS	

ALS, amyotrophic lateral sclerosis; AD, Alzheimer's disease; bvFTD, behavioral variant frontotemporal dementia; DLB, dementia with Lewy bodies; MND, motor neuron disease; PLS, primary lateral sclerosis; PNFA, progressive non-fluent aphasia; SD, semantic dementia.

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Scientific correspondence

***C9ORF72* repeat-associated non-ATG-translated polypeptides are distributed independently of TDP-43 in a Japanese patient with c9ALS**

Hexanucleotide (GGGGCC) repeat expansion in a noncoding region of *C9ORF72* is the major genetic cause of frontotemporal dementia and amyotrophic lateral sclerosis (c9FTD/ALS) in the Caucasian population [1], but it is very rare in the Japanese population, possibly because of the difference in genetic background [2,3]. TDP-43 pathology indistinguishable from that of non-mutational ALS/FTLD-TDP has been observed in c9FTD/ALS [4]. In addition, the presence of p62-, ubiquitin- and ubiquilin-positive, and TDP-43-negative inclusions in the cerebellar cortex and hippocampus has been reported to be a unique and consistent feature in Caucasian patients with c9FTD/ALS [5,6]. Recently, it was demonstrated that these TDP-43-negative inclusions are derived from aggregated dipeptide repeat (DPR) proteins bidirectionally translated from the expanded repeat in *C9ORF72* by repeat associated non-ATG (RAN) translation, and that such DPR protein pathology is widely distributed in the central nervous system (CNS) [7,8]. However, it still remains unknown whether these distinct neuropathological features are also reproduced in Japanese patients with c9FTD/ALS. In the present study, we performed an immunohistochemical analysis focusing especially on DPR proteins in a Japanese patient with *C9ORF72* repeat expansion (c9ALS) [2], who to our knowledge represents the only autopsy case of this genetic disease to have been reported in the Japanese population so far.

The present study was conducted with approval from the Institutional Review Board of Niigata University. The clinical and pathological findings in this case have been reported previously (case 4 [9]). Briefly, the patient had a sibling who had also been diagnosed as having ALS. At the age of 61, he noticed hand clumsiness, and progressively developed bulbar palsy and limb weakness. He died 20 months after disease onset due to respiratory failure. He had no clinical features suggestive of dementia. The presence of *C9ORF72* repeat expansion was confirmed by

repeat-primed PCR using the frozen cerebellar tissue [2]. The accurate hexanucleotide repeat length was unknown, because we could not perform Southern blot analysis due to lack of the amount of genomic DNA. Histologically, neuronal loss and gliosis were evident in the spinal anterior horns, brainstem motor nuclei and motor cortex, as well as degeneration in the anterior and lateral columns of the spinal cord. Bunina bodies and ubiquitin-positive skein-like inclusions were observed in the remaining lower motor neurones. Neuronal and glial (oligodendrocytic) cytoplasmic inclusions (NCIs and GCIs) recognized with anti-TDP-43 (polyclonal, Protein Tec Group, Chicago, IL, USA; 1:4000) were present in the lower motor nuclei and motor cortex, and much less frequently in the subcortical non-motor nuclei, such as the basal ganglia, whereas no such NCIs or GCIs were observed in the cerebellar cortex and hippocampus [9]. There were p62-positive and TDP-43-negative NCIs in the cerebellar granule cells and in the granule cells and pyramidal CA4-CA2 neurones of the hippocampus; at that time, we failed to show phosphorylated TDP-43 (pTDP-43)-positive NCIs in the lower motor nuclei and motor cortex [2].

In the present study, we confirmed that NCIs and GCIs recognized with a 'polyclonal' antibody against pTDP-43 (pS409/410; Cosmo Bio, Tokyo, Japan; 1: 1000) were present with the same distribution pattern as that of TDP-43 mentioned above (Figure 1a,c). Immunostaining with an antibody against another RNA-binding protein, RNA-binding motif 45 (RBM45) (polyclonal, Sigma-Aldrich, St. Louis, MO, USA; 1:50), which has been known to accumulate in inclusions in ALS [10], also revealed a distribution of positive inclusions strikingly similar to that of TDP-43-positive inclusions, although such NCIs and GCIs were comparatively small in number. The characteristic morphologies of NCIs and GCIs were shared by both anti-TDP-43/anti-pTDP-43 and anti-RBM45 (Figure 1a–d).

We generated polyclonal antibodies against putative DPR proteins from the GGGGCC repeat by RAN translation [11]. Although immunostaining with three antibodies against different polypeptides, poly Gly-Ala (GA), poly Gly-Pro and poly Gly-Arg, arising from RAN translation revealed similar DPR protein pathology, abundant positive

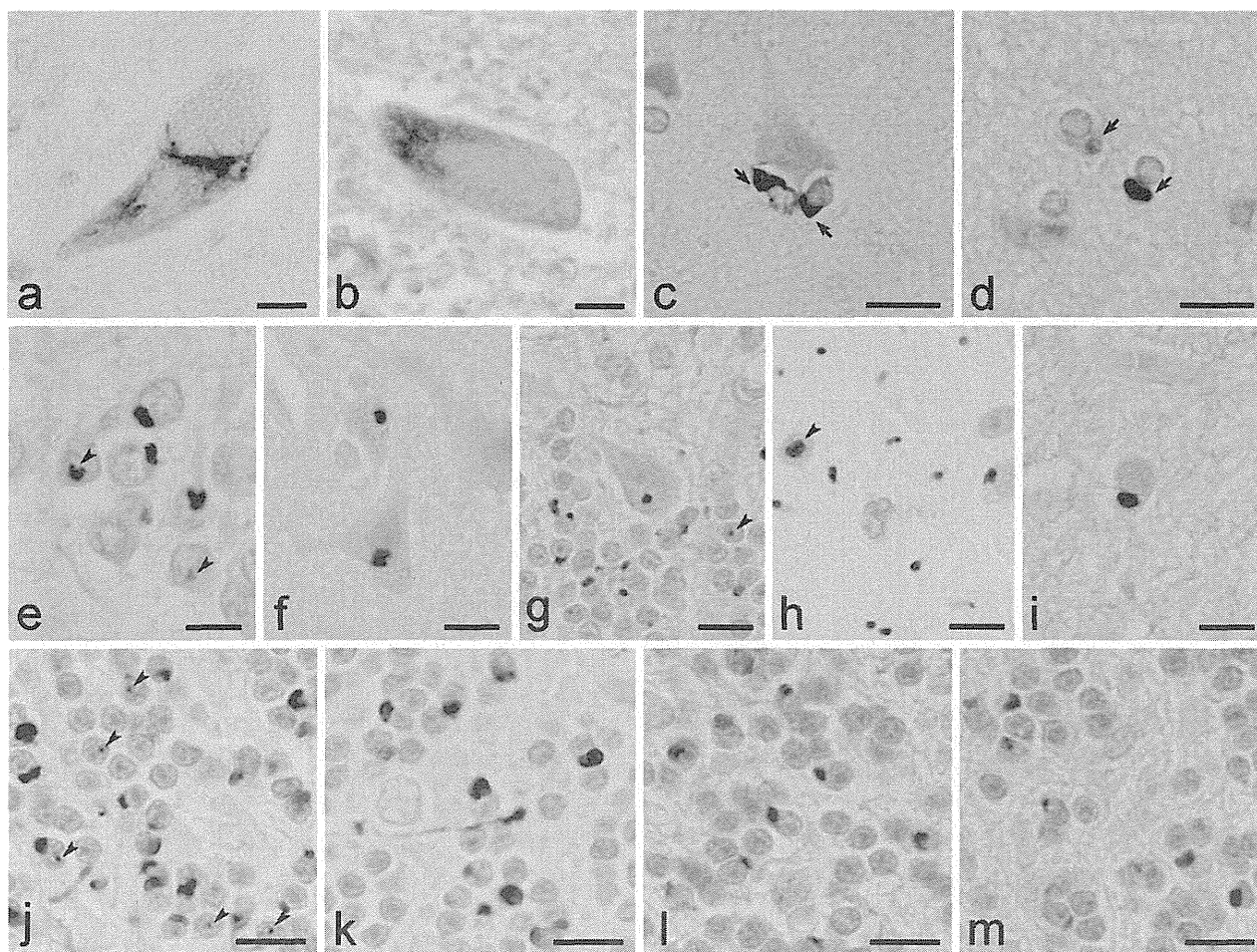


Figure 1. Immunohistochemical study. (a–d) Skein-like cytoplasmic inclusions positive for phosphorylated TDP-43 (pTDP-43) (a) and RBM45 (b) in the cervical anterior horn cells. C-like or fibrillary tangle-like glial cytoplasmic inclusions positive for pTDP-43 (c) and RBM45 (d) in the motor cortex. Note glial cytoplasmic inclusions indicated by arrows (c,d). (e–h) Numerous poly Gly-Ala (GA)-positive neuronal cytoplasmic and intranuclear inclusions (arrowheads) are observed widely beyond the regions showing TDP-43-positive inclusions [(e) hippocampal dentate gyrus, (f) temporal cortex, (g) cerebellar granular-Purkinje layer, (h) cerebellar molecular layer; also see Table 1]. A Purkinje cell also contains a positive cytoplasmic inclusion (g). Punctate or linear inclusions in the neuropil of the cerebellar molecular layer are also evident (h). (i) Here, an oligodendroglial cytoplasmic inclusion is shown in the precentral subcortical white matter. (j–m) In the cerebellar granular layer, it is evident that GA-positive neuronal cytoplasmic inclusions (j) are also positive for ubiquitin (k), ubiquitin (l) and p62 (m). Note GA-positive intranuclear inclusions indicated by arrowheads (j). Bars: 10 μ m (a–m).

inclusions were recognized most clearly with anti-poly GA (1:1000). The distribution pattern of DPR protein-positive NCIs was apparently different from that of TDP-43-positive NCIs, the latter being clearly associated with neuronal loss in the lower and upper motor neurones systems. DPR protein-positive NCIs were widely distributed in the brain (Figure 1e–h,j), with the highest frequency in the hippocampal dentate gyrus (Figure 1e) and cerebellar granular layer (Figure 1g,j), and were distributed almost evenly in the cerebral neocortex examined (Figure 1f). DPR protein-positive GCIs were comparatively

rare (Figure 1i), although TDP-43- and RBM45-positive GCIs were observed frequently (Figure 1c,d). Neuronal intranuclear inclusions were frequently associated with anti-DPR proteins (Figure 1e,g,h,j), but not with anti-TDP-43. DPR protein-positive punctate or filamentous structures were also encountered in the neuropil of the cerebellar molecular layer (Figure 1h), which were never recognized with anti-TDP-43, and only rarely with anti-ubiquitin (monoclonal, clone 5F5, Abnova, Walnut, CA, USA; 1:10 000). DPR protein-positive NCIs appeared as irregular dot-like, granular or star-like inclusions

Table 1. Summary of regional immunohistochemical findings

	Lower motor neurones*	Motor cortex	Non-motor cortex†	Dentate gyrus	Hippocampal CA4	Cerebellar granule cells
Neuronal loss	++	+	-	-	-	-
TDP-43	+	+	-	-	-	-
pTDP-43	+	+	-	-	-	-
RBM45	+	+	-	-	-	-
p62	+	++	++	++	++	++
Ubiquitin	+	+	+	+	+	++
Ubiquilin	-	+	+	++	+	++
DPR‡	-	++	++	+++	++	+++

Severity of neuronal loss are represented as: - = not noted, + = mild, ++ = moderate-severe.

Neurones containing each antibody-positive inclusions were counted per 100 neurones in high-power fields, and ratio is represented as: - = none, + = ~10%, ++ = 10-30%, +++ = >30%.

*Cervical anterior horn cells and hypoglossal nucleus.

†Frontal and temporal cortex were examined.

‡poly Gly-Ala.

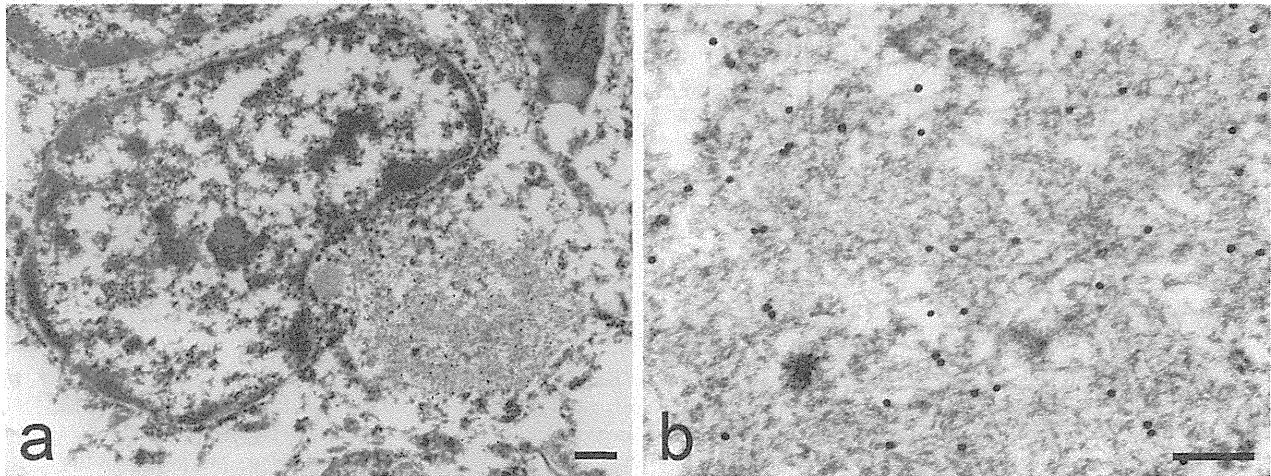


Figure 2. Immunoelectron microscopy for the presence of ubiquilin. (a) A cerebellar granule cell is seen to have a round filamentous cytoplasmic aggregate. (b) Higher-magnification view of the aggregate, showing immunogold particles on the randomly arranged, approximately 10-nm-wide filamentous structures. Bars: (a) 500 nm, (b) 200 nm.

(Figure 1e-h,j) that generally differed in morphology from TDP-43- and RBM45-positive NCIs (Figure 1a-d). Such inclusions were frequently labelled with antibodies against ubiquilin, ubiquitin (polyclonal, Dako, Glostrup, Denmark; 1:800) and p62 (monoclonal, BD biosciences, San Jose, CA, USA; 1:500) (Figure 1j-m). The immunohistochemical findings are summarized in Table 1.

Under immunogold-labelling electron microscopy for ubiquilin in the cerebellar granular layer tissue, the labelled NCIs were shown to be composed of randomly arranged filamentous structures (Figure 2a,b); their morphological features were very similar to those of ubiquitin-

positive and TDP-43-negative NCIs in the entorhinal cortex in a case of c9FTD/ALS [12]. Double-labelling immunofluorescence revealed that co-localization of TDP-43 (monoclonal, clone 2E2-D3, Abnova, Taipei, Taiwan; 1:250) and RBM45 (Figure 3a-c), but not TDP-43 (2E2-D3) and DPR proteins (Figure 3d-f), was a feature in the cytoplasmic inclusions. The majority of p62-positive NCIs were positive for DPR proteins, whereas p62-positive GCIs were negative for DPR proteins (Figure 3g-i).

In the present Japanese case of c9ALS, the histological and molecular pathology of sporadic ALS was observed: especially, the occurrence of Bunina bodies and TDP-43-

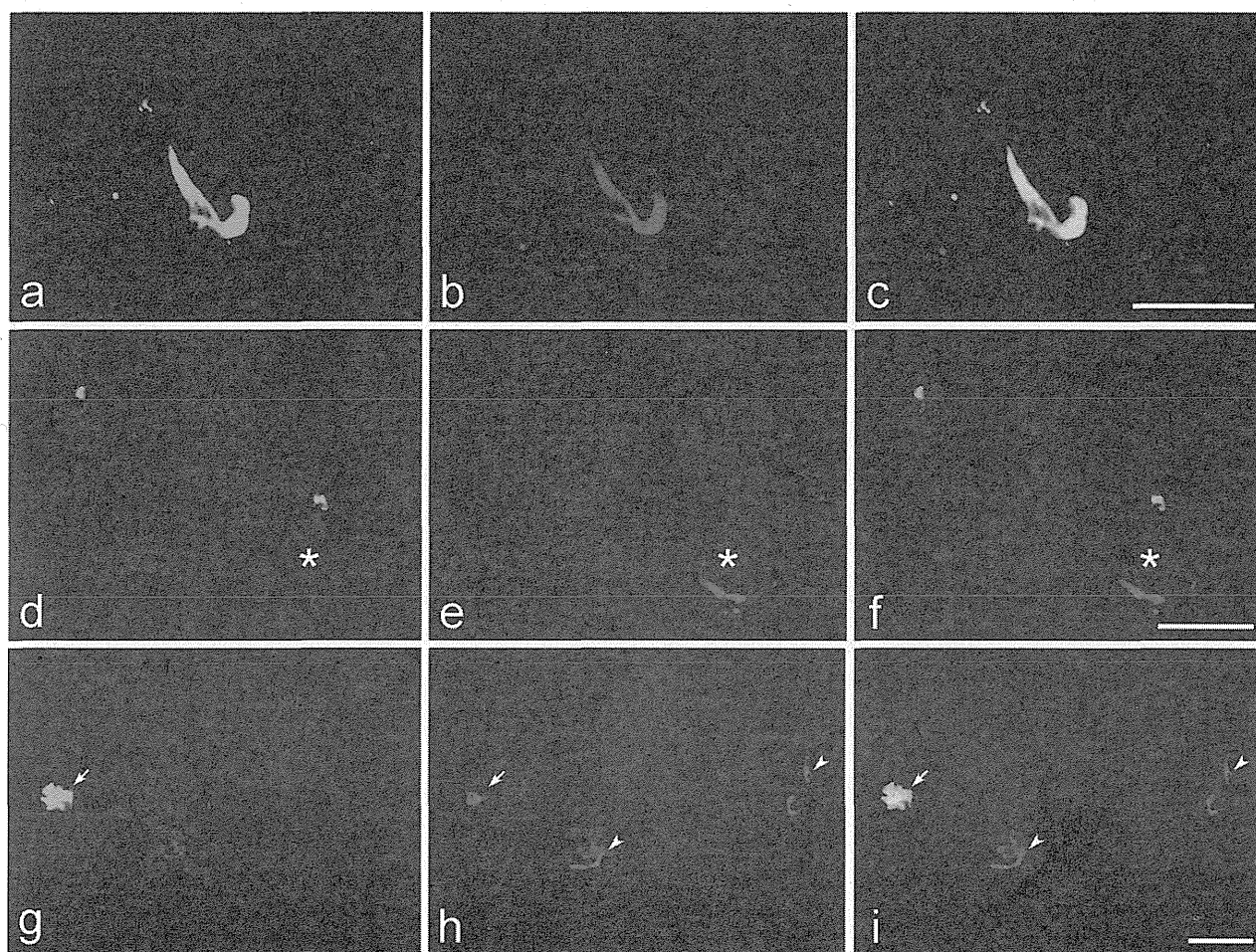


Figure 3. Double-labelling immunofluorescence for the presence of TDP-43, RBM45, DPR proteins and p62. (a–i) Inclusions observed in the motor cortex cells. (a–c) Colocalization of RBM45 (a, green) and TDP-43 (b, red) is evident in the cytoplasmic inclusions in an oligodendrocytic cell (c, merge). (d–f) Poly Gly-Ala (GA) (d, green) and TDP-43 (e, red) never co-localize in the cytoplasmic inclusions (f, merge), although co-existence of poly GA- and TDP-43-positive inclusions are rarely observed in the same cells, mostly in the neuronal cells (asterisks) (d–f). (g–i) Co-localization of poly GA (g, green, arrow) and p62 (h, red, arrow) is evident in cytoplasmic inclusions in a neuronal cell (i, merge, arrow). Poly GA-negative (g, green) and p62-positive inclusions (h, red, arrowheads) are also evident in two oligodendrocytic cells (i, merge, arrowheads). Bars: 10 μ m (a–i).

positive NCIs was confirmed. It is important to note that the TDP-43 pathology and neuronal loss were correlated with each other; this was also the case for another RNA-binding protein, RBM45, which has been reported to be increased in the cerebrospinal fluid of ALS patients, and also colocalized with TDP-43 in inclusions of patients with ALS and FTLD-TDP [10]. In addition, the occurrence of p62-positive, and TDP-43-negative granular or star-shaped NCIs in the cerebellar cortex and hippocampus was a feature. Importantly, such p62-positive and TDP-43-negative NCIs were also labelled with antibodies against DPR proteins we generated. The DPR protein

pathology was distributed independently of TDP-43 pathology: co-localization of DPR proteins and TDP-43 was extremely rare, if present, in the same inclusion, indicating that unlike TDP-43 and RBM45, there was no correlation between DPR protein pathology and neuronal loss (Table 1), or between DPR protein pathology and clinical symptoms, as pointed out previously in Caucasian cases of c9FTD/ALS [13].

It is noteworthy that most recently, the DPR proteins have been reported to be translated from not only a sense direction of the repeats but also an antisense direction (CCCCGG); in fact, two additional DPR proteins (poly