Imaging Biomarkers for Neurodegeneration in Presymptomatic Familial Frontotemporal Lobar Degeneration

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Frontotemporal lobar degeneration (FTLD) is a neurodegenerative disorder characterized by behavioral changes, language abnormality, as well as executive function deficits and motor impairment. In about 30–50% of FTLD patients, an autosomal dominant pattern of inheritance was found with major mutations in the MAPT, GRN, and the C9orf72 repeat expansion. These mutations could lead to neurodegenerative pathology years before clinical symptoms onset. With potential disease-modifying treatments that are under development, non-invasive biomarkers that help determine the early brain changes in presymptomatic FTLD patients will be critical for tracking disease progression and enrolling the right participants into the clinical trials at the right time during the disease course. In recent years, there is increasing evidence that a number of imaging biomarkers show the abnormalities during the presymptomatic stage. Imaging biomarkers of presymptomatic familial FTLD may provide insight into the underlying neurodegenerative process years before symptom onset. Structural magnetic resonance imaging (MRI) has demonstrated cortical degeneration with a mutation-specific neurodegeneration pattern years before onset of clinical symptoms in presymptomatic familial FTLD mutation carriers. In addition, diffusion tensor imaging (DTI) has shown the loss of white matter microstructural integrity in the presymptomatic stage of familial FTLD. Furthermore, proton magnetic resonance spectroscopy (1H MRS), which provides a non-invasive measurement of brain biochemistry, has identified early biochemical abnormalities in presymptomatic MAPT mutation carriers. Positron emission tomography (PET) imaging with [18F]-fluorodeoxyglucose (FDG) has demonstrated the glucose hypometabolism in the presymptomatic stage of familial FTLD. Also, a novel PET ligand, 18F-AV-1451, has been used in this group to evaluate tau deposition in the brain. Promising imaging biomarkers for presymptomatic familial FTLD have been identified and assessed for specificity and sensitivity for accurate prediction of symptom onset and tracking disease progression during the presymptomatic stage when clinical measures are not useful. Furthermore, identifying imaging biomarkers for the presymptomatic stage is important for the design of disease-modifying trials. We review the recent progress in imaging biomarkers of the presymptomatic phase of familial FTLD and discuss the imaging techniques and analysis methods, with a focus on the potential implication of these imaging techniques and their utility in specific mutation types.

Keywords: imaging biomarker, presymptomatic, familial frontotemporal lobar degeneration, MAPT, GRN, C9orf72
INTRODUCTION

Frontotemporal lobar degeneration (FTLD) is a progressive neurodegenerative disorder characterized by behavioral abnormalities, language impairment, impaired social cognition, and executive function, as well as progressive supranuclear palsy (PSP), corticobasal syndrome (CBS), or motor neuron disease (1). FTLD has a strong genetic background with an autosomal dominant pattern of inheritance in about 30–50% FTLD patients (2) with major mutations in the microtubule-associated protein tau gene (MAPT) (3), progranulin (GRN), and the repeat expansions in the chromosome 9 open reading frame 72 gene (C9orf72). Mutations in the gene encoding the microtubule-associated protein tau (MAPT) on chromosome 17 was first reported in 1998 (4) and have been found in many kindreds with familial FTLD. The vast majority of known mutations occurring in the coding region are in the repeats, causing the reduced ability of the mutant tau proteins to interact with microtubules, leading to hyperphosphorylated tau accumulation in glia and neurons causing neurodegeneration, white-matter integrity alterations, and brain atrophy years before clinical symptom onset (3, 5–8). Subtypes of MAPT mutations were associated with different types of tauopathies. These are mutations inside exon 10 (i.e., N279K, S305N, and P301L) and outside exon 10 (i.e., R406W and V337M). Mutations inside exon 10 tend to form four tandem microtubule-binding domain repeat (4R-tau) pathology rather than 3 repeat (3R-tau) pathology (9), while mutations outside exon 10 tend to form mixed 3R/4R tau pathology.

In 2006, loss-of-function mutations in the progranulin gene (GRN) were first reported to cause familial FTLD (10, 11). There are more than 70 different pathogenic GRN mutations that have been identified until this date (http://www.molgen.ua.ac.be/FTDmutations). Mutations in the GRN gene leads to a loss of progranulin levels through haploinsufficiency, and the intraneuronal aggregation of TAR DNA-binding protein (TDP)-43 protein (12). The most frequent clinical phenotypes of GRN mutation carriers were behavioral variant frontotemporal dementia (bvFTD), CBS, and primary progressive aphasia (PPA) (13, 14). GRN mutation carriers have FTLD with TDP-43 inclusions and present with a diverse clinical phenotype and a highly heterogeneous age of disease onset.

In 2011, the hexanucleotide GGGGCC (G4C2) repeat expansions of the chromosome 9 open reading frame 72 gene (C9orf72) C9orf72 gene were found as a common cause of both FTD and amyotrophic lateral sclerosis (ALS) (15, 16). The most common clinical phenotypes associated with C9orf72 expansions are bvFTD (17), ALS, or the combination of both in one person (18). Cases with the C9orf72 repeat expansion with histopathological correlation had TDP-43 depositions (18).

These mutations associated with familial FTLD could lead to brain neurodegeneration years before symptom onset. Investigation of families with the presence of MAPT, GRN, or C9orf72 provide a unique opportunity to shed light on early neurodegenerative changes and also identify biomarkers for tracking disease progression in future disease-modifying trials. In recent years, there is growing evidence that a number of imaging biomarkers show abnormalities during the presymptomatic stage. Imaging biomarkers of presymptomatic familial FTLD may provide insight into the underlying neurodegenerative process years before symptom onset.

DISCUSSION

Structural MRI

Structural magnetic resonance imaging (sMRI) has captured cortical degeneration with a mutation-specific neurodegeneration pattern years before onset of clinical symptoms in presymptomatic familial FTLD mutation carriers (6, 19–32). Studies of sMRI using different analysis methods, such as region of interest (ROI) in specific brain regions, cortical thickness analysis and voxel-based morphometry (VBM), could capture the gray matter and white matter volumes, cortical thickness, and the subcortical gray matter volume.

MAPT_sMRI

Previous cross-sectional MRI studies demonstrated atrophy in the anteromedial temporal lobe and orbitofrontal cortex in asymptomatic MAPT mutation carriers (6, 19, 20), while others found no difference between asymptomatic MAPT mutation carriers and controls (6, 7, 33). Recently, two longitudinal studies from a cohort of asymptomatic MAPT mutation carriers have reported that hippocampal volumes decline during a 2-year follow-up, but no cortical atrophy was found in longitudinal analysis with 4 years of follow-up (21, 22). During a 10-year follow-up, the rates of temporal lobe atrophy were accelerated in asymptomatic MAPT mutation carriers who were asymptomatic during follow-up, while accelerated atrophy rates in the temporal, parietal and frontal lobes were reported in MAPT mutation carriers who became symptomatic compared to non-carriers (23). These data altogether suggest the consistent finding of early involvement of the anterior–medial temporal lobe in asymptomatic MAPT mutation carriers. Taken together, the cross-sectional and longitudinal studies of lobar cortical atrophy in MAPT mutation carriers suggest a sequential pattern throughout the disease course. The cortical volume appears to decline in the temporal lobe early in the asymptomatic stage, with an acceleration of atrophy rates along with development of symptoms, followed by the frontal and parietal lobe atrophy with sparing of the occipital lobe (23) (Table 1).

Across different subtypes of MAPT mutations (IVS10+16, IVS10+3, N279K, S305N, P301L, and V337M), similar patterns of atrophy were reported in the later symptomatic phase (41). However, the atrophy pattern in the early disease phase may be varied across the different subtypes of MAPT mutations. For example, patients with MAPT N279K mutations have prominent motor symptoms early in the disease process (42–44) (Figure 1A), and therefore involvement of the primary and supplementary motor cortices may be expected in this subtype group (45–49). The N279K MAPT mutation carriers' trajectories of lobar atrophy, such as the supplemental motor cortex involvement, may be specific to the suspected 4R-tau associated neurodegeneration in N279K kindred. Further studies
| No. | Author | Year | Study design | No. of subjects | Techniques | Findings |
|-----|--------|------|--------------|----------------|------------|----------|
| 1   | Miyoshi et al. (34) | 2010 | Cross-sectional | 3 aMAPT+ vs. 9 HC | $^{[11]}$C DAA1106 PET | Gial activities were increased in the frontal cortex of aMAPT+, the occipital cortex of 2 aMAPT+, and the posterior cingulate cortex of 1 aMAPT+; $^{[11]}$C dopa PET | Low dopamine synthesis in putamen |
|     |        |      |              |                | $^{[11]}$C MP4A PET | Reduced AChE activity in the temporal, parietal cortex |
| 2   | Kantarci et al. (8) | 2010 | Cross-sectional | 14 aMAPT+ vs. 24 HC | $^1$H MRS | Elevated mI/Cr and decreased NAA/mI in PCC voxel |
| 3   | Whitwell et al. (33) | 2011 | Cross-sectional | 8 aMAPT+ vs. 8 NC | sMRI | No difference |
|     |        |      |              |                | rfMRI | Reduced connectivity in the DMN |
| 4   | Dopper et al. (7) | 2014 | Cross-sectional | 11 aMAPT+ vs. 8 NC | sMRI | No difference |
|     |        |      |              |                | DTI | Decreased FA and increased RD in bilateral uncinate fasciculi and reduced FA in the forceps minor |
| 5   | Rohrer et al. (19) | 2015 | Cross-sectional | 15 aMAPT+ vs. 8 NC | sMRI | Atrophy in the hippocampus, amygdala, temporal lobe, and insula |
| 6   | Dopper et al. (35) | 2016 | Longitudinal | 11 aMAPT+ vs. 31 NC | ASL | No difference |
| 7   | Fumagalli et al. (20) | 2018 | Cross-sectional | 24 aMAPT+ vs. 148 NC | sMRI | No difference |
| 8   | Cash et al. (6) | 2018 | Cross-sectional | 23 aMAPT+ vs. 144 NC | sMRI | No difference |
| 9   | Jiskoot et al. (36) | 2018 | Cross-sectional | 17 aMAPT+ vs. 115 NC | DTI | Reduced FA and increased diffusivity in the uncinate fasciculus and cingulum |
| 10  | Jones et al. (37) | 2018 | Cross-sectional | 3 aMAPT+ vs. 241 HC | $^{18}$F-AV-1451 tau PET | Low level of uptake in 1 asymptomatic N279K mutation carrier; little to no signal in 1 R406W mutation carrier, high uptake in 1 R406W mutation carrier |
| 11  | Panman et al. (22) | 2019 | Longitudinal | 14 aMAPT+ vs. 53 NC | sMRI | Baseline: No difference Follow-up: lower GM volume in the left temporal pole, a trend toward cortical thinning of the right inferior temporal lobe; Longitudinal: GM volume decline in the hippocampus |
| 12  | Chen et al. (38) | 2019 | Longitudinal | 14 aMAPT+ vs. 50 NC | DTI | No difference |
| 13  | Chen et al. (23) | 2019 | Longitudinal | 14 aMAPT+ (include 4 converters) vs. 23 NC | sMRI | Baseline: higher MD in entorhinal WM Longitudinal: accelerated annualized change of entorhinal MD |
| 14  | Chen et al. (39) | 2019 | Cross-sectional | 9 aMAPT+ vs. 25 NC | $^1$H MRS | Baseline: No difference |
| 15  | Chen et al. (40) | 2019 | Longitudinal | 8 aMAPT+ converters | $^1$H MRS | Baseline: No difference |

are needed to characterize the trajectories of rates of cortical atrophy across the different subtypes of MAPT mutations.

**GRN_sMRI**

In FTLD patients with GRN mutations, the pattern of brain atrophy was reported to be asymmetric and widespread, predominantly involving the frontal, inferior parietal, and posterior temporal cortices (6, 13, 50–54). In asymptomatic GRN mutation carriers, the structural MRI studies were inconsistent (24–27).

Although most cross-sectional (6, 7, 20, 32, 55–57) and longitudinal structural MRI studies (22) reported no gray matter volume differences between asymptomatic GRN mutation carriers and controls, a few recent studies found a gray matter
atrophy pattern mainly involving the frontal (25), parietal (58), and temporal lobes (27). Brain atrophy was reported in the insula early, 15 years before the expected symptom onset, followed by the temporal and parietal lobes at 10 years before the expected symptom onset, then in the striatum at 5 years before the expected symptom onset (19). In a longitudinal study of asymptomatic GRN mutation carriers with a 20-month follow-up, no gray matter volume loss was found at baseline, but left inferior and middle temporal gyri atrophy was reported 20 months later (26). Reduced gray matter density was also reported in bilateral orbitofrontal, anterior temporal, and insular cortices at baseline with greater annualized GM density changes in right orbitofrontal and left occipital cortices during follow-up (27). A recent study reported a sequential pattern of regional cortical atrophy rates with up to 10 years of follow-up, involving the frontal and parietal lobe cortices early during the asymptomatic stage, followed by the temporal lobe cortex after symptom onset in GRN mutation carriers (28) (Table 2).

These inconsistencies might be due to the age at examination, the proximity to onset of clinical symptoms, associated lobar cortical atrophy changes, the sample size of GRN mutation carriers, and the statistical analysis methodology with different sensitivity and specificity. For example, voxel-based analysis needs larger effect sizes and corrections for multiple comparisons to detect differences in cortical volumes between groups than the targeted ROI analysis. Furthermore, many studies have demonstrated the heterogeneity of clinical presentation during the early phase and variability in age of onset in GRN mutations carriers (65, 66). In addition, investigations in different time windows during the disease course may capture the variable findings in the laterality of lobar cortical atrophy across individuals with GRN mutations. It may partly explain the inconsistent results of cortical atrophy among the prior cross-sectional studies of asymptomatic GRN mutation carriers.

In GRN mutation carriers, asymmetric atrophy is a common finding with left or right asymmetry reported even within the same family (51, 54). Asymmetric cortical atrophy was found early, 5 years before expected symptom onset in asymptomatic GRN mutation carriers (19), while the underlying pathological cause for the asymmetry remains unclear. In contrast, symmetric rates of brain atrophy were also found in symptomatic GRN mutation carriers in a longitudinal study (67). This discrepancy suggests that asymmetry in cortical atrophy may be more predominant during the early stages, becoming more symmetric during the later stages of the disease (67). Also, as noted above, the specific phenotype and associated lobar cortical atrophy changes and the differences in analysis methods may be underlying these discrepancies. It is important to note that laterality occurs at various stages of neurodegenerative disease progression, which may increase the variability in atrophy rates in individual GRN mutation carriers (Figure 1B).
| No. | Author                | Year | Study design | No. of subjects | Techniques        | Findings                                                                                   |
|-----|-----------------------|------|--------------|-----------------|-------------------|--------------------------------------------------------------------------------------------|
| 1   | Borroni et al. (55)   | 2008 | Cross-sectional | 7 aGRN+ vs. 15 HC | sMRI, DTI        | No differences                                                                            |
|     |                       |      |              |                  |                   | Reduced FA in left uncinate fasciculus, left inferior occipitofrontal fasciculus           |
| 2   | Borroni et al. (56)   | 2012 | Cross-sectional | 9 aGRN+ vs. 13 NC | sMRI, rMRI       | No differences                                                                            |
|     |                       |      |              |                  |                   | Increased connectivity in SN                                                              |
| 3   | Jacova et al. (59)    | 2013 | Cross-sectional | 9 aGRN+ vs. 11 NC | 18FDG-PET        | Lower FDG uptake in right medial, ventral frontal cortex and insula                       |
| 4   | Moreno et al. (57)    | 2013 | Cross-sectional | 13 aGRN+ vs. 13 HC | sMRI,            | No difference                                                                             |
|     |                       |      |              |                  |                   | Global reserve index was inversely related to functional activation of the ventral SN in the right precentral gyrus and of the dorsal SN in the right middle temporal gyrus |
| 5   | Premi et al. (60)     | 2013 | Cross-sectional | 17 aGRN+        | rfMRI,           | No difference                                                                             |
|     |                       |      |              |                  |                   | Reduced functional connectivity in SN and DMN                                            |
| 6   | Dopper et al. (7)     | 2014 | Cross-sectional | 28 aGRN+ vs. 28 NC | sMRI, DTI,       | No difference                                                                             |
|     |                       |      |              |                  |                   | Reduced functional connectivity in SN and DMN                                            |
| 7   | Pevani et al. (25)    | 2014 | Cross-sectional | 5 aGRN+ vs. 5 NC  | sMRI, DTI,       | No difference                                                                             |
|     |                       |      |              |                  |                   | Atrophy in the right orbitofrontal, precentral gyrus, and left rostral middle frontal gyrus|
|     |                       |      |              |                  |                   | Increased AD in the right cingulum, superior longitudinal fasciculus, and corticospinal tract. |
| 8   | Premi et al. (61)     | 2014 | Cross-sectional | 17 aGRN+ vs. 38 HC | rMRI,            | No difference                                                                             |
|     |                       |      |              |                  |                   | Reduced ReHo in the left parietal region                                                  |
| 9   | Premi et al. (62)     | 2014 | Cross-sectional | 17 aGRN+ vs. 14 HC | rMRI,            | No difference                                                                             |
|     |                       |      |              |                  |                   | Increased ReHo in the bilateral mesial frontal cortex                                     |
|     |                       |      |              |                  |                   | Decreased brain connectivity within the left frontoparietal network and increased connectivity in the executive network |
|     |                       |      |              |                  |                   | The TMEM106B at-risk polymorphism (T/T) was associated with decreased connectivity within the ventral SN and the left frontoparietal network |
| 10  | Caroppo et al. (26)   | 2015 | Longitudinal  | 16 aGRN+ vs. 17 NC (baseline) | sMRI            | Baseline: no difference                                                                 |
|     |                       |      |              | 14 aGRN+ vs. 14 NC (follow-up) |                | Follow-up: atrophy in the left middle and inferior temporal gyri                          |
|     |                       |      |              |                  |                   | Baseline: hypometabolism in the left middle temporal gyrus                                |
|     |                       |      |              |                  |                   | Follow-up: hypometabolism in the frontal, temporal lobes and thalamus                      |
|     |                       |      |              |                  |                   | Atrophy in the insula (15 years before EO), then in the temporal and parietal lobes (10 years before EO) and the striatum (5 years before EO) |
| 11  | Rohrer et al. (19)    | 2015 | Cross-sectional | 45 aGRN+ vs. 93 NC | sMRI,            | Baseline: no difference                                                                 |
|     |                       |      |              |                  |                   | Follow-up: fronto-parietal hypoperfusion                                                   |
|     |                       |      |              |                  |                   | Longitudinal: stronger decrease in CBF in the frontal, temporal, parietal and subcortical regions |
|     |                       |      |              |                  |                   | No correlation of GM with plasma progranulin levels in symptomatic and asymptomatic GRN mutation carriers as a whole group |
| 12  | Dopper et al. (35)    | 2016 | Longitudinal  | 23 aGRN+ vs. 31 NC | ASL,            | Baseline: no difference                                                                   |
|     |                       |      |              |                  |                   | Follow-up: fronto-parietal hypoperfusion                                                   |
|     |                       |      |              |                  |                   | Longitudinal: stronger decrease in CBF in the frontal, temporal, parietal and subcortical regions |
|     |                       |      |              |                  |                   | No correlation of GM with plasma progranulin levels in symptomatic and asymptomatic GRN mutation carriers as a whole group |
| 13  | Galimberti et al. (63) | 2018 | Cross-sectional | 19 sym GRN+, 64 aGRN+ vs. 77 NC | sMRI,           | Lower task-evoked functional activation in the anterior and posterior ventrolateral prefrontal cortex |
|     |                       |      |              |                  |                   | No difference                                                                             |
| 14  | Alexander et al. (64) | 2018 | Cross-sectional | 5 aGRN+ vs. 11 NC | Task-based fMRI | No difference                                                                             |
| 15  | Cash et al. (6)       | 2018 | Cross-sectional | 65 aGRN+ vs. 144 NC | sMRI,           | No difference                                                                             |
| 16  | Jiskoot et al. (36)   | 2018 | Cross-sectional | 52 aGRN+ vs. 115 NC | sMRI, DTI       | Reduced FA and increased diffusivity in the internal capsule                               |
| 17  | Olm et al. (27)       | 2018 | Longitudinal  | 11 aGRN+ vs. 11 NC | sMRI,           | Baseline: reduced GM density in bilateral orbitofrontal, insular, and anterior temporal cortices |
|     |                       |      |              |                  |                   | (Continued)                                                                                 |
TABLE 2 | Continued

| No. | Author               | Year | Study design | No. of subjects | Techniques | Findings |
|-----|----------------------|------|--------------|-----------------|------------|----------|
| 18  | Gazzina et al. (58)  | 2018 | Cross-sectional | 19 aGRN+ vs. 17 NC | sMRI       | Increased cortical thickness and decreased surface area of the right parietal lobe |
| 19  | Fumagalli et al. (20) | 2018 | Cross-sectional | 66 aGRN+ vs. 148 NC | sMRI       | No difference |
| 20  | Popuri et al. (32)   | 2018 | Cross-sectional | 9 aGRN+ vs. 37 NC  | sMRI       | No difference |
| 21  | Chen et al. (28)     | 2019 | Longitudinal  | 8 aGRN+ vs. 10 NC  | sMRI       | Increased rates of atrophy in the frontal and parietal lobe cortices |
| 22  | Panman et al. (22)   | 2019 | Longitudinal  | 33 aGRN+ vs. 53 NC  | sMRI       | No difference |
|     |                      |      |              | 28 aGRN+ vs. 50 NC  | DTI        | No difference |

C9orf72_sMRI

In asymptomatic C9orf72 mutation carriers, cross-sectional structural MRI studies consistently found a diffuse pattern of atrophy including frontal, temporal, parietal, insular, and posterior cortical regions, as well as subcortical volumes of thalamus, hippocampus, and the cerebellum (6, 22, 29–32), while very few studies reported no difference between asymptomatic C9orf72 mutation carriers and healthy controls (20, 68). The atrophy of subcortical regions is estimated to occur as early as 25 years before the expected symptom onset in C9orf72 mutation carriers, and later involves temporal and frontal lobes at around 20 years before the expected symptom onset, and finally involving the cerebellum at around 10 years before the expected symptom onset (19). The early involvement of subcortical regions is a specific atrophy pattern in asymptomatic C9orf72 mutation carriers that distinguishes them from other FTLD mutation carriers.

To date, two longitudinal investigations have reported that there is no difference in the trajectory of cortical atrophy rates between asymptomatic C9orf72 mutation carriers and controls (22, 68), suggesting that longer time follow-up duration is needed to track the trajectory in asymptomatic C9orf72 mutation carriers (Table 3). Figure 1C shows serial structural MRIs from a C9orf72 mutation carrier who converted from asymptomatic phase to symptomatic phase during follow up.

A few studies on familial FTLD have combined different mutation carriers as one group and compared them with non-carriers (Table 4). Combined asymptomatic GRN and C9orf72 mutation carriers demonstrated overlapping atrophy in the insula (70), which has also been reported in a study combining asymptomatic MAPT, GRN, and C9orf72 mutation carriers (6). No difference was observed in both cross-sectional (7) and longitudinal investigations (21) in the combined cohort of asymptomatic MAPT and GRN mutation carriers. However, in a small cohort of 5 MAPT mutation and 3 GRN mutation carriers who converted from asymptomatic to symptomatic phase, extensive temporal and frontal cortical atrophy were observed, while no gray matter volume loss was observed longitudinally in asymptomatic MAPT and GRN mutation carriers who remained asymptomatic compared to non-carriers (21).

There has been a comparison of atrophy patterns resulting from mutations in different genes (22); the distinct pattern of atrophy in the early stage of disease course among the MAPT, GRN, and C9orf72 mutation carriers align well with previous studies in symptomatic mutation carriers (67).

Diffusion Tensor Imaging

Diffusion tensor imaging (DTI) is a sensitive technique to detect the white matter microstructure alterations. Damage to the white matter is a common finding in postmortem studies of sporadic and familial FTLD patients (71–73). Loss of white matter integrity on DTI was characterized by increased mean diffusivity (MD), radial diffusivity (RD), and axial diffusivity (AD), as well as reduced fractional anisotropy (FA).

MAPT_DTI

In asymptomatic MAPT mutation carriers, reduced FA and increased diffusivity were reported in bilateral uncinate fasciculus and cingulum in cross-sectional studies (7, 36), while a longitudinal study only found increased MD in entorhinal white matter with accelerated increase in MD during follow-up (38). Furthermore, in a longitudinal study, lower FA in uncinate fasciculus and other white matter tracts were reported in MAPT and GRN mutations 2 years before symptom onset, while no difference in FA was found between the combined group of asymptomatic MAPT and GRN mutation carriers who remained asymptomatic and non-carriers (21). Another recent longitudinal investigation also reported no difference in DTI measurements between asymptomatic MAPT mutation carriers and non-carriers (22).

Compared to FA, the diffusivity parameters such as MD appear to be more sensitive measures of the white matter...
**TABLE 3 |** Studies investigating asymptomatic C9orf72 mutation vs. controls.

| No. | Author          | Year | Study design | No. of subjects | Techniques | Findings                                                                 |
|-----|-----------------|------|--------------|-----------------|------------|--------------------------------------------------------------------------|
| 1   | Rohrer et al. (19) | 2015 | Cross-sectional | 18 aC9+         | sMRI       | Atrophy in subcortical areas (the thalamus, insula, and posterior cortical areas) at 25 years before EO, followed by the frontal and temporal lobes at 20 years before EO, and the cerebellum at 10 years before EO |
| 2   | Walhout et al. (29) | 2015 | Cross-sectional | 16 aC9+ vs. 23 NC | sMRI       | Cortical thinning in the temporal, parietal and occipital regions, and smaller volume in the left caudate and putamen. No difference |
| 3   | Roeter et al. (68) | 2016 | Longitudinal  | 7 aC9+ vs. 28 HC | sMRI, DTI  | Baseline: no difference Longitudinal: no difference |
| 4   | Lee et al. (30) | 2017 | Cross-sectional | 5 aC9+ vs. 23 HC | sMRI       | Lower GM intensity in the bilateral posterior mid-cingulate, left medial pulvinar thalamus, and small, scattered regions in the bilateral dorsolateral prefrontal cortex |
|     |                 |      |              | 12 aC9+ vs. 29 HC | DTI        | Reduced FA in the corpus callosum, cingulum bundles, corticospinal tracts, uncinate fasciculi and inferior longitudinal fasciculi. |
|     |                 |      |              | 13 aC9+ vs. 30 HC | rMRI       | Intrinsic connectivity deficits in DMN, sensorimotor, but most prominent in SN and medial pulvinar thalamus-seeded networks |
| 5   | Papma et al. (69) | 2017 | Cross-sectional | 18 aC9+ vs. 15 NC | sMRI       | No difference for whole group; in a subgroup >40 years, lower gray matter volume in the right inferior temporal gyrus, right cerebellum, left postcentral and precentral gyrus, the left superior parietal lobe and the left thalamus |
|     |                 |      |              |                 | DTI        | Lower FA and higher RD within the right superior corona radiata, inferior longitudinal fasciculus, uncinate fasciculus, internal and external capsule, bilateral anterior thalamic radiation and corticospinal tract |
| 6   | Bertrand et al. (31) | 2018 | Cross-sectional | 41 aC9+ vs. 39 NC | sMRI, DTI  | Atrophy in the frontal, inferior temporal and parietal cortex and bilateral thalamus |
| 7   | Cash et al. (6) | 2018 | Cross-sectional | 40 aC9+ vs. 144 NC | sMRI       | GM loss bilaterally in the thalamus, right superior posterior cerebellum, superior temporal and inferior frontal regions |
| 8   | Jiskoot et al. (36) | 2018 | Cross-sectional | 35 aC9+ vs. 115 NC | DTI        | Posteriorly located WM tracts (posterior thalamic radiation, splenium of the corpus callosum, posterior corona radiata) |
| 9   | Fumagalli et al. (20) | 2018 | Cross-sectional | 24 aC9+ vs. 148 NC | sMRI       | No difference |
| 10  | Popuri et al. (32) | 2018 | Cross-sectional | 15 aC9+ vs. 37 NC | sMRI       | Cortical thinning in the temporal, parietal and frontal regions; reduced volumes of bilateral thalamus and left caudate |
| 11  | Panman et al. (22) | 2019 | Longitudinal | 11 aC9+ vs. 53 NC | sMRI       | Baseline: GM volume loss in the cerebellum, insula, left fronto-temporal lobes; cortical thinning in the right postcentral gyrus Follow-up: GM volume loss in the thalamus, cerebellum, and several bilateral orbitofrontal and insular cortices, and the postcentral gyrus; Cortical thinning in bilateral precentral gyrus and right superior parietal lobe |
|     |                 |      |              | 12 aC9+ vs. 50 NC | DTI        | Baseline and follow-up: lower FA in frontotemporal tracts and higher MD in the entire skeleton |

Integrity alterations in asymptomatic familial FTLD (36, 38). One possible explanation is that FA is more influenced by the crossing fibers, which may limit its sensitivity to detect subtle white matter changes (74). The underlying pathologic mechanism of white matter diffusion abnormalities in MAPT mutation carriers remains unclear. It may be due to the Wallerian degeneration secondary to the cortical tau pathology or to the tau pathology observed in the white matter. Future studies are needed to correlate these findings with pathology.

**GRN_DTI**

White matter involvement from DTI studies in asymptomatic GRN mutation carriers showed reduced FA in the left uncinate fasciculus, left inferior occipitofrontal fasciculus, and the genu of the corpus callosum (55). Reduced FA and increased diffusivity in the internal capsule was also reported (36). Another DTI investigation did not find a difference in FA, MD, and RD, but reported increased AD in the right cingulum, superior longitudinal fasciculus, and corticospinal tract (25). In a longitudinal design, reduced FA was found in bilateral
superior longitudinal fasciculus, frontal corpus callosum, and left corticospinal tract in asymptomatic GRN mutation carriers, with greater annualized FA change in frontal corpus callosum and right superior longitudinal fasciculus compared to controls (27). Others did not report any abnormalities of DTI metrics in both cross-sectional (7) and longitudinal investigations (22) of asymptomatic GRN mutation carriers.

As indicated in sMRI, left–right asymmetry was also found in most white matter tracts of GRN mutation carriers. The most consistent asymmetry was reported in the uncinate fasciculus, retrolenticular part, and anterior limb of the internal capsule, as well as external capsule (36). However, the development of asymmetry over time may have different patterns in various white matter tracts. Longitudinal data with larger cohorts are needed for such investigations.

C9orf72 DTI

Investigations in asymptomatic C9orf72 carriers with DTI have shown reduced white matter integrity in the fiber tracts connecting frontal lobe (the uncinate fasciculus and inferior longitudinal fasciculus), motor function-related tracts (corticospinal tracts, corona radiata, and internal/external capsule), thalamic radiation, as well as corpus callosum and cingulum bundles and cerebellar peduncles (30, 31, 36, 69).

One cross-sectional investigation reported no DTI abnormality in asymptomatic C9orf72 mutation carriers (29). A recent longitudinal investigation reported a more diffuse white matter involvement in asymptomatic C9orf72 mutation carriers, with lower FA in the frontotemporal tracts and higher MD in the entire white matter skeleton compared to non-carriers. However, these white matter differences remained relatively stable during a 2-year follow-up period (22).

The heterogeneity in clinical phenotypes that ranged from motor neuron disease to FTD and the variation in time of symptom onset within and between families of C9orf72 mutation carriers may be responsible for the variable results from cross-sectional DTI investigations in asymptomatic C9orf72 mutation carriers. Longitudinal follow-up is essential in gaining insight into clinical and neuroimaging characteristics. Therefore, studying families with varying disease phenotypes along the ALS-FTD spectrum would be of particular interest in addressing the
issue of clinical heterogeneity in relation to specific early neuroimaging changes.

Functional MRI
Functional MRI can evaluate the alterations from functionally connected networks in spatially distinct subcortical and cortical areas in asymptomatic FTLD mutation carriers. In asymptomatic MAPT mutation carriers, resting-state fMRI demonstrated decreased connectivity in the default mode network (DMN) predominantly between lateral temporal lobe and precuneus (33). Components of the DMN, such as lateral temporal lobes and medial prefrontal cortex, were implicated in functions of semantic memory (75) and theory of mind (76), which show abnormalities in both patients with sporadic bvFTD and symptomatic MAPT mutation carriers. However, another fMRI study found no difference in functional connectivity between asymptomatic MAPT mutation carriers and controls (7). The change of functional connectivity was observed early in the disease course even without gray matter loss in asymptomatic MAPT mutation carriers (33), suggesting that functional abnormalities may precede the occurrence of atrophy in these regions.

In the asymptomatic C9orf72 mutation carriers, the reduced functional connectivity was found prominently in salience network (SN), as well as in medial pulvinar of thalamus-seeded networks, DMN, and sensorimotor network (30). The fMRI alterations can already be distinguished in persons younger than 40 years of age (30). The SN is the main functional network associated with FTLD (77), which plays a role in behavioral functioning and emotion processing (78). Thus, reduced connectivity across central nodes of the SN may lead to some of the clinical features of FTLD.

Increasing evidence from the resting-state fMRI studies demonstrate widely affected networks in asymptomatic GRN mutation carriers, including both increased (56) and reduced connectivity in the SN (7), as well as reduced connectivity in the DMN (7). Functional activation of the SN was inversely related to global reserve index in asymptomatic GRN mutation carriers (60). Furthermore, hypoconnectivity of the left fronto-parietal network (FPN) and a hyper-connectivity within the executive network have been demonstrated (62). The functional impairment of parietal lobes had been thought to be the earliest feature of FTD with GRN mutations (61), while one study reported no difference in functional connectivity between asymptomatic GRN mutation carriers and non-carriers (25). When performing a relational reasoning task, asymptomatic GRN mutation carriers presented lower task-evoked functional activation in anterior and posterior ventrolateral prefrontal cortex compared to controls (64).

As suggested by theoretical concept of molecular neopathies (79), a specific proteinopathy affects specific networks, while other brain networks could be dynamically involved during the course of the disease. This may possibly explain the heterogeneity in the fMRI findings in asymptomatic FTLD mutation carriers. However, other environmental or genetic factors can also affect the involved networks in a complex way. For example, TMEM106B genetic variations had a modulating effect on the functional connectivity in asymptomatic GRN mutation carriers. For example, the TT genotype of TMEM106B may cause additional neurodegeneration by reduced connectivity in the left fronto-parietal network and ventral SN (62). More studies with longitudinal follow-up are needed to further validate the use of functional connectivity as a potential biomarker for clinical trials.

Proton Magnetic Resonance Spectroscopy
Proton magnetic resonance spectroscopy (1H MRS), which provides non-invasive measurements of brain biochemistry, is a potential imaging marker for early detection of neurodegenerative disease progression in familial FTLD. 1H MRS metabolite measurements have been sensitive biomarkers of early neurodegenerative pathology in FTLD, AD, and Lewy body dementia (80–85). N-acetylaspartate/Cre(tine (NAA/Cr) is considered a biomarker of neuronal integrity (86), neuronal viability (87, 88), and synaptic integrity (89), while elevated mI may be associated with astrocytic and microglial activation, as well as glial proliferation (90, 91).

In both asymptomatic and symptomatic MAPT mutation carriers, neurochemical alterations from the posterior cingulate voxel were found on single-voxel 1H MRS (92). An elevation in myo-inositol (mI) or mI to creatine (mI/Cr) and a decrease in the neuronal integrity marker NAA or NAA/Cr have been found in symptomatic patients with FTLD (83), while only elevated mI/Cr in the posterior cingulate gyrus has been reported in asymptomatic MAPT mutation carriers (8). During longitudinal follow-up in MAPT mutation carriers who converted from the asymptomatic to symptomatic disease, serial 1H MRS from the posterior cingulate voxel demonstrated that metabolite ratio changes, characterized by increasing mI/Cr and decreasing NAA/mI ratios, begin to accelerate ~2 years prior to symptom onset (40).

However, 1H MRS from the medial frontal lobe voxel was more sensitive than 1H MRS from the PCC voxel in the asymptomatic MAPT mutation carriers, characterized by decreased NAA/Cr and NAA/mI ratios (39), suggesting specific regional involvement in neurodegenerative diseases. Regional differences that are apparent in the early stage of disease course may be lost as neurodegenerative pathology spreads to most of the brain regions during the later stages in MAPT mutation carriers.

Potentially, anterior temporal lobe 1H MRS measurements would be the most sensitive to early changes in MAPT mutation carriers. However, this may be offset by a reduction in the quality of the 1H MRS data, since anterior temporal lobes are proximal to the magnetic susceptibility artifacts from the skull base that impact the quality of the 1H MR spectra.

To date, no MRS study was reported in asymptomatic GRN and C9orf72 mutation carriers. In the future, whole-brain and multi-voxel MRS could provide metabolite changes in more brain regions than single-voxel MRS (93). By utilizing more advanced acquisition methods, it would be possible to quantify metabolites, such as glutamine, glutathione, and Scyllo-Inosital (94). For example, an advanced 1H-MRS protocol composed of semi-localization by adiabatic selective refocusing (sLASER) localization and FAST(EST)MAP shimming detected
lower glutamate concentration in patients with amnestic mild cognitive impairment than clinically normal controls indicating early Alzheimer's disease pathophysiology (95).

**Positron Emission Tomography**

Positron emission tomography (PET) imaging is often suggested as a useful biomarker for the earliest stage of FTD and had demonstrated differences in metabolism by different tracers in the presymptomatic stage of familial FTLD.

**MAPT_PET**

In symptomatic MAPT mutation carriers, [18F]-fluorodeoxyglucose (FDG)-PET has showed asymmetric temporal lobe hypometabolism (46). Frontotemporal hypometabolism on FDG-PET was reported in two asymptomatic MAPT mutation carriers in the same family who had only mild speech change at the time of examination. These participants were in a transitional stage from normal neurological functioning to overt FTLD and one became symptomatic later, suggesting that FDG-PET changes can be detected when asymptomatic MAPT mutation carriers get close to phenoconversion. However, specificity of FDG-PET to differentiate between various neurodegenerative diseases at the early stages remains unknown (96).

Given the heterogeneity of tau pathology observed in MAPT mutations, tau PET (e.g., 18F-AV-1451 PET) was explored as a biomarker for various forms of tau pathology within a similar patient population. Subtypes of MAPT mutations were associated with different AV-1451 uptake patterns due to different types of underlying tauopathies. The mutations inside exon 10 (i.e., S305N, P301L, and N279K) with 4R tau pathology had a low level of AV-1451 binding in Tau PET in asymptomatic MAPT mutation carriers. Mutations outside exon 10 (i.e., V337M and R406W) with mixed 3R/4R tau pathology that are more likely to produce AD-like tau pathology presented had a high magnitude of binding of AV-1451 in symptomatic MAPT mutation carriers (37). In symptomatic MAPT V337M mutation carriers, the tau accumulation assessed by [18F] AV1451 tracer was also associated with regional brain atrophy by structural MRI (97). Tau PET studies in asymptomatic MAPT mutation carriers are very limited. One asymptomatic MAPT N279K mutation carrier presented low level of AV-1451 uptake, one of two asymptomatic MAPT R406W mutation carriers had little to no signal and the other one had high level of AV-1451 uptake (37). General conclusions about the tau PET in asymptomatic MAPT mutation carriers remain inconclusive due to small number of cases reported so far.

Tracers associated with dopaminergic function were applied in MAPT mutation carriers, especially in the N279K mutation type, which may present with parkinsonism as the first symptom. Dopaminergic dysfunction was shown early in asymptomatic MAPT N279K mutation carriers via 2b-carbomethoxy-3b-(4-trimethylstannylphenyl) tropane (11C-CFT)-PET (34, 98). Gliial activity was elevated in the frontal cortex, the posterior cingulate cortex, and the occipital cortex of asymptomatic MAPT mutation carriers on [11C] DAA1106 PET, and acetylcholinesterase activity was reduced in the temporoparietal cortex using [11C] N-methylpiperidin-4-yl acetate PET in three asymptomatic MAPT mutation carriers (34). These case studies with small sample sizes will need to be replicated in larger cohorts.

**GRN_PET**

In asymptomatic GRN mutation carriers, hypometabolism was detected in right medial, ventral frontal cortex, andinsula on FDG-PET (59). A longitudinal investigation reported that hypometabolism in the left middle temporal gyrus at baseline is associated with greater decrease in metabolism in the frontotemporal lobes and thalamus in asymptomatic GRN mutation carriers compared to non-carriers (26).

**C9orf72_PET**

A radioligand for the microtubule-associated protein tau (18F-Flortaucipir) was found to show increased binding in semantic variant primary progressive aphasia (svPPA) (99), which is associated with underlying TDP-43 pathology. In contrast, a recent study reported that none or limited 18F-Flortaucipir retention was found in symptomatic C9orf72 mutation carriers, suggesting that 18F-Flortaucipir binding in svPPA patients is not a general TDP-43 related phenomenon (100). To date, a PET tracer for the in vivo detection of TDP-43 pathology does not exist.

**CONCLUSION**

This review underscores the importance of imaging biomarkers for accurate prediction of symptom onset and tracking of disease progression during the presymptomatic stage of familial FTLD. The application of advanced neuroimaging techniques in monogenic familial FTLD represents a unique model to detect the natural history of specific proteinopathies and clinical phenotypes. They may also provide a direct comparison across different gene groups in the future. Furthermore, multiple MRI and PET modalities may provide information on the different aspects and stages of the neurodegenerative disease process in a single individual. Many of the studies discussed in this review focus on a single imaging biomarker. Thus, it is not possible to compare biomarkers on their accuracy in prediction of onset of clinical symptoms and tracking disease progression in familial FTLD. The potential application of multi-model imaging is to provide evidence of very early asymptomatic alterations by collectively including cortical atrophy, white matter integrity loss, functional alteration, as well as brain metabolic changes. There are several ongoing prospective multisite studies (e.g., GENFI, ARTFL, and LEFFTDS) involving the familial FTLD kindred. These longitudinal investigations in large cohorts on the genotype-specific imaging profiles for MAPT, GRN, and C9orf72 at different time points during the disease course would provide insights into their potential use as prognostic biomarkers for clinical trials of disease-modifying therapies.

**AUTHOR CONTRIBUTIONS**

QC data collection, analysis and interpretation of the data, drafting the manuscript. KK design or conceptualization of the study, data collection, analysis and interpretation of the data, drafting the manuscript.
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Conflict of Interest: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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