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The Benson Complex Figure Test detects deficits in visuoconstruction and visual memory in symptomatic familial frontotemporal dementia: A GENFI study

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ABSTRACT

Objective: Sensitive cognitive markers are still needed for frontotemporal dementia (FTD). The Benson Complex Figure Test (BCFT) is an interesting candidate test, as it assesses visuospatial, visual memory, and executive abilities, allowing the detection of multiple mechanisms of cognitive impairment. To investigate differences in BCFT Copy, Recall and Recognition in presymptomatic and symptomatic FTD mutation carriers, and to explore its cognitive and neuroimaging correlates.

Method: We included cross-sectional data from 332 presymptomatic and 136 symptomatic mutation carriers (GRN, MAPT or C9orf72 mutations), and 290 controls in the GENFI consortium. We examined gene-specific differences between mutation carriers (stratified by CDR® NACC-FTLD score) and controls using Quade's / Pearson X^2 tests. We investigated associations with neuropsychological test scores and grey matter volume using partial correlations and multiple regression models respectively.

Results: No significant differences were found between groups at CDR® NACC-FTLD 0–0.5. Symptomatic *GRN* and *C9orf72* mutation carriers had lower Copy scores at CDR® NACC-FTLD \geq 2. All three groups had lower Recall scores at CDR® NACC-FTLD \geq 2, with *MAPT* mutation carriers starting at CDR® NACC-FTLD \geq 1. All three groups had lower Recognition scores at CDR® NACC FTLD \geq 2. Performance correlated with tests for visuoconstruction, memory, and executive function. Copy scores correlated with frontal-subcortical grey matter atrophy, while Recall scores correlated with temporal lobe atrophy.

Conclusions: In the symptomatic stage, the BCFT identifies differential mechanisms of cognitive impairment depending on the genetic mutation, corroborated by gene-specific cognitive and neuroimaging correlates. Our findings suggest that impaired performance on the BCFT occurs relatively late in the genetic FTD disease process. Therefore its potential as cognitive biomarker for upcoming clinical trials in presymptomatic to early-stage FTD is most likely limited.

1. Introduction

Frontotemporal dementia (FTD) is one of the most prevalent forms of early-onset dementia. Its clinical profile is typically characterized by disturbances in behaviour (behavioural variant; bvFTD) and language (primary progressive aphasia; PPA), with cognitive deficits in executive function and social cognition commonly seen. In contrast, episodic memory and visuospatial abilities are relatively spared [1-2]. FTD has an autosomal dominant inheritance pattern in around a third of cases, with mutations in progranulin (GRN), microtubule-associated protein tau (MAPT), and chromosome 9 open reading frame 72 (C9orf72) the most common causes of familial FTD [3]. As the mutations cause brain atrophy in distinct as well as overlapping anatomical brain regions, the associated phenotypes are often rather heterogeneous [4]. The clinical presentation associated with GRN mutations includes bvFTD, nonfluent variant PPA, atypical parkinsonism, and corticobasal syndrome (CBS) [5,6]. The cognitive profile commonly shows executive dysfunction, speech and language disorders, amnestic deficits and apraxia, consistent with frontal, temporal and parietal lobe involvement [7,8]. Patients with MAPT mutations commonly present with bvFTD or atypical parkinsonism (CBS or progressive supranuclear palsy, PSP) [9], with early and prominent naming and memory recall deficits as a result of symmetrical anteromedial temporal lobe atrophy [4,8]. Lastly, the C9orf72 repeat expansion is associated with a clinical phenotype of bvFTD, amyotrophic lateral sclerosis (ALS), and FTD-ALS [10,11]. The pattern of cognitive impairment is often widespread, including deficits in language, attention, mental processing speed, executive function and immediate memory recall [8], due to atrophy of the frontal and temporal, as well as posterior cortical and subcortical (e.g., cerebellum and thalamus) areas [12,13].

In recent years, research in the familial FTD field has increasingly

focussed on the presymptomatic stage, as the critical time-window for treatment most likely lies prior to overt symptom onset, when the pathological damage is still low. With promising therapeutic avenues leading to disease-modifying therapy trials, the identification of robust clinical biomarkers is of utmost importance [12]. Interestingly, previous neuropsychological studies show that subtle cognitive decline is present in the presymptomatic stage of FTD (up to 10 years prior to overt disease onset), with gene-specific cognitive profiles for *GRN*, *MAPT* and *C9orf72* [14–17]. This suggests that presymptomatic neuropsychological assessment may provide sensitive cognitive markers indicative of disease, onset and progression.

One particular neuropsychological instrument, the Benson Complex Figure test (BCFT), is an interesting candidate for familial FTD. Being part of the National Alzheimer's Coordinating Centre (NACC) FTD-module neuropsychological battery [18], performance on the BCFT relies on multiple cognitive functions, including visuospatial abilities, visual memory, and executive functions such as organization and working memory. Most studies into the BCFT have looked into differences between patients with bvFTD, patients with AD, and healthy controls, demonstrating a trend for those with bvFTD to score lower on figure copying than controls [19–20]. Moreover, poor figure copy correlated with specific cognitive mechanisms (i.e. spatial planning and working memory) and neuroanatomical atrophy substrates (i.e. dorsolateral prefrontal cortex) in bvFTD [19]. Until now, research into the BCFT in presymptomatic FTD has been lacking.

The aim of the present study was therefore to: 1) investigate cross-sectional differences in the BCFT (copy, recall and recognition) between presymptomatic FTD mutation carriers, symptomatic FTD mutation carriers and cognitively unimpaired controls; 2) explore associations between the BCFT and other neuropsychological tests, and 3) examine associations between the BCFT and grey matter (GM) volume. Additionally, we investigated normative data and relationships with age, sex and education from the cognitively unimpaired control group.

¹ See Appendix 1 for the full list of GENFI consortium members.

2. Method

2.1. Participants

We included baseline data of 758 participants from genetically confirmed FTD families with either a GRN or MAPT pathogenic variant, or C9orf72 repeat expansion, recruited within the GENFI 2 fifth data freeze between March 2015 and May 2019. We determined clinical status according to established diagnostic criteria [1-2,21] and a standardized clinical assessment, including medical and family history taking, extensive neuropsychological assessment covering the major cognitive domains (see Neuropsychological assessment below), and MR imaging of the brain [14]. DNA genotyping was performed locally at each research site. >30 repeats in C9orf72 was considered to be pathogenic [22]. The total sample consisted of 332 presymptomatic mutation carriers (*GRN* = 143; *MAPT* = 59; *C9orf72* = 130), 136 symptomatic mutation carriers (GRN = 41; MAPT = 23; C9orf72 = 72), and 290 noncarriers that were used as reference group (GRN = 122; MAPT = 57; C9orf72 = 111). The clinical diagnoses in symptomatic mutation carriers were as follows: bvFTD (n = 91), PPA (n = 21), ALS or FTD-ALS (n = 91), PPA (n = 21), ALS or FTD-ALS (n = 91), PPA (n = 21), ALS or FTD-ALS (n = 91), PPA (n = 21), ALS or FTD-ALS (n = 91), PPA (n = 21), ALS or FTD-ALS (n = 91), PPA (n = 21), ALS or FTD-ALS (n = 91), PPA (n = 21), ALS or FTD-ALS (n = 91), PPA (n = 21), ALS or FTD-ALS (n = 91), PPA (n = 21), ALS or FTD-ALS (n = 91), PPA (n = 21), ALS or FTD-ALS (n = 91), PPA (n = 21), ALS or FTD-ALS (n = 91), PPA (n = 21), ALS or FTD-ALS (n = 91), PPA (n = 21), ALS or FTD-ALS (n = 91), ALS or FTD-ALS (n = 91), PPA (n = 21), ALS or FTD-ALS (n = 91), ALS (= 15), PSP (n = 2), dementia not otherwise specified (n = 2), and other (n = 5). We administered the global CDR® NACC-FTLD global score [23] as a measure of disease severity. Knowledgeable informants answered questions about behavioural and cognitive symptoms as well as the participant's activities of daily living in a structured interview which included two questionnaires (Cambridge Behavioural Inventory -Revised (CBI-R) [24] and Frontotemporal Dementia Rating Scale (FRS) [25]. Unless presymptomatic mutation carriers had undergone predictive testing at their own request, the clinical investigators were blinded to their genetic status. We obtained written informed consent from all participants at study enrolment. Ethical committees at each research site approved the study. This study was conducted in accordance with the declaration of Helsinki.

2.2. Benson complex figure test

The BCFT [19] is part of the standardized GENFI neuropsychological battery and consists of 3 conditions: *Copy* (in which the figure has to be copied from an example – see Fig. 1), *Recall* (in which the figure has to be drawn from memory after a 10–15 min interval), and *Recognition* (in which the target figure has to be recognised amongst three distractor figures). Scoring follows the NACC FTD-criteria [18]. Total scores for both Copy and Recall range from 0 to 16; each of the eight elements can receive a maximum score of two when both accuracy and placement are correct. A bonus point – adding up to a maximum score of 17 – is given when the figure is well-drawn (i.e., each element must be accurately drawn, all elements must be properly placed, all elements must be drawn in proper proportions, all connections between elements must be clean, and no extraneous lines may be present). Recognition is either scored as correct (score 1) or incorrect (score 0).

2.3. Neuropsychological assessment

Global cognitive functioning was screened by means of the Mini-Mental State Examination (MMSE) [26], whilst other cognitive tests performed within the larger GENFI neuropsychological battery measured executive function (letter fluency [27]), Trail Making Test (TMT) [28], D-KEFS Color-Word Interference Test [29]), memory (Free and Cued Selective Reminding Test (FCSRT) [30]), and visuoconstructive abilities (WASI Block Design [31]).

2.4. MRI acquisition and (pre)processing

Volumetric T1-weighted MR images were acquired on a 3 T scanner in 698 participants (Philips Achieva n=191, Siemens Prisma n=191, Siemens Trio n=178, Siemens Skyra n=136, GE Discovery MR750 n=2). All images were subjected to strict visual quality control, after which

Appendix 3. Neuroimaging correlates of the BCFT. Abbreviations: GRN, progranulin; MAPT, microtubule-associated protein tau; C9orf72, chromosome 9 open reading frame 72; BCFT, Benson Complex Figure Test; L, left; R, right. *only clusters >50 voxels were reported; **uncorrected *p* < 0.001.

| Gene | BCFT | Cluster | T | $P_{FWE\text{-corrected}}$ | MNI coordinates | | | Region |
|---------|---------|---------|------|----------------------------|-----------------|-----|-----|---------------------------|
| | | | | | х | у | z | |
| GRN | Сору | 4 | 5.15 | 0.031 | -20 | -22 | 10 | Thalamus L |
| | Recall* | 7354 | 6.82 | < 0.001 | 2 | 32 | 27 | Anterior cingulate R |
| | | 2964 | 7.00 | < 0.001 | -33 | -28 | -10 | Hippocampus L |
| | | 1726 | 6.35 | < 0.001 | -15 | -58 | 39 | Precuneus L |
| | | 1605 | 6.45 | < 0.001 | -40 | 15 | 2 | Frontal operculum L |
| | | 1542 | 6.57 | < 0.001 | 44 | 2 | 2 | Anterior insula R |
| | | 943 | 6.13 | < 0.001 | 20 | -22 | -20 | Parahippocampal gyrus R |
| | | 637 | 5.72 | < 0.001 | -32 | -63 | -38 | Cerebellum L |
| | | 590 | 5.77 | < 0.001 | -22 | 3 | -16 | Basal forebrain L |
| | | 238 | 6.33 | < 0.001 | 2 | 10 | -12 | Subcallosal area R |
| | | 216 | 5.96 | < 0.001 | 33 | 14 | -24 | Temporal pole R |
| | | 206 | 5.45 | < 0.001 | 22 | -70 | -36 | Cerebellum R |
| | | 191 | 5.54 | < 0.001 | -26 | 40 | -12 | Anterior orbital gyrus L |
| | | 128 | 5.95 | < 0.001 | -50 | 39 | -8 | Inferior frontal gyrus L |
| | | 128 | 5.55 | < 0.001 | 51 | -57 | -12 | Inferior temporal gyrus R |
| | | 117 | 5.59 | 0.001 | -2 | -88 | 14 | Cuneus L |
| | | 113 | 5.58 | 0.001 | -39 | -16 | 45 | Precentral gyrus L |
| | | 82 | 5.52 | 0.001 | -44 | 26 | 10 | Inferior frontal gyrus L |
| | | 62 | 5.43 | 0.003 | -34 | 18 | -16 | Posterior orbital gyrus L |
| | | 50 | 5.50 | 0.004 | -8 | -96 | 18 | Occipital pole L |
| | | 50 | 5.26 | 0.004 | 24 | 58 | 4 | Superior frontal gyrus R |
| MAPT | Copy** | 57 | 3.74 | < 0.001 | 46 | -70 | -22 | Cerebellum R |
| | | 44 | 3.69 | < 0.001 | 52 | -45 | -40 | Cerebellum R |
| | Recall* | 185 | 6.41 | < 0.001 | -21 | 4 | -38 | Temporal pole L |
| | | 89 | 6.08 | < 0.001 | -18 | -36 | 3 | Hippocampus L |
| | | 85 | 5.80 | < 0.001 | -33 | -32 | -2 | Hippocampus L |
| | | 67 | 6.29 | 0.001 | 24 | 30 | -9 | Hippocampus R |
| | | 57 | 5.75 | 0.001 | 33 | -18 | -12 | Hippocampus R |
| C9orf72 | Copy** | 22 | 3.26 | 0.001 | -26 | 12 | 30 | Middle frontal gyrus L |
| | Recall* | 593 | 6.34 | < 0.001 | -28 | -20 | -14 | Hippocampus L |
| | | 100 | 5.49 | < 0.001 | 27 | -22 | -12 | Hippocampus R |
| | | 58 | 5.56 | 0.002 | 18 | -12 | -12 | Hippocampus R |

15 participants were excluded from further analysis due to inadequate image quality. The DICOM images were subsequently corrected for gradient nonlinearity distortions and converted to NifTI format. These images were then analysed using the standard Voxel-Based Morphometry (VBM) pipeline in Statistical Parametric Mapping 12 (SPM12; Functional Imaging Laboratory, University College London, London, UK; www.fil.ion.ucl.ac.uk/spm) implemented in Matlab R2018a (Mathworks, USA). In the first pre-processing step, the T1-weighted images were normalized to a template space and segmented into GM, white matter (WM) and cerebrospinal fluid (CSF), after which they were rigidly aligned. We calculated total intracranial volume (TIV) by adding GM, WM and CSF. Secondly, the segmentations were spatially normalized to a DARTEL template by applying the flow fields of all the individual scans. Images were smoothed using a 6 mm full width at half maximum (FWHM) isotropic Gaussian kernel. At every preprocessing step, images were visually inspected.

2.5. Statistical analysis

We performed statistical analyses using SPSS Statistics 25.0 (IBM Corp., Armonk, NY, USA) and GraphPad Prism 5 (La Jolla, California, USA). Alpha was set at 0.05 across all comparisons, unless otherwise specified, and two-tailed analyses were performed. We compared continuous demographic data between groups by means of one-way ANOVA with post hoc Bonferroni comparisons for normally distributed data, or Kruskal-Wallis tests with post hoc Mann-Whitney *U* tests in case of non-normally distributed data. Between-group differences in sex distribution were analysed using Pearson X² tests. In our reference (healthy control) group, we calculated cumulative frequencies, percentile scores, and performance across age, sex and education for BCFT Copy, Recall and Recognition. We used Spearman rank correlations to explore the relationships between the BCFT Copy and Recall, and age and education. The square root of eta squared $(\sqrt{\eta^2})$ was used to investigate the relationship between age and education, and BCFT Recognition. We explored the differences in BCFT Copy and Recall and sex by means of Mann-Whitney U tests, and sex differences in BCFT Recognition by means of a Pearson X² test. As BCFT Copy and Recall scores were non-normally distributed, we examined gene-specific (GRN, MAPT, C9orf72) differences between presymptomatic mutation carriers (CDR® NACC-FTLD global score 0 and 0.5), symptomatic mutation carriers (CDR® NACC-FTLD global score ≥ 1) and controls by means of Quade's rank analysis of covariance - adjusting for the effect of age, sex, years of education, and family clustering. We performed Pearson X2 tests to compare BCFT Recognition scores between groups. We investigated associations between BCFT Copy and Recall with neuropsychological test scores per mutation by means of partial correlations, controlling for the effect of age, sex, years of education, and family

clustering. We explored the relationship between each BCFT test score and GM volume by means of multiple regression models in SPM12 (University College London, London, UK). Age, sex, scanner and TIV were entered as covariates. We set the statistical threshold at p < 0.05, adjusted for multiple comparisons with familywise error (FWE) correction. The uncorrected statistical threshold was set at p < 0.001 (minimum cluster size > 10 voxels).

3. Results

3.1. Demographic and clinical data

Demographic and clinical data are shown in Table 1. Controls were significantly younger than symptomatic mutation carriers (GRN U = 1655.5, MAPT U = 1374.5; C9orf72 U = 3190.5; all p < 0.001), while presymptomatic MAPT mutation carriers were younger than controls (U = 6043, p < 0.001). All presymptomatic mutation carriers were younger than symptomatic mutation carriers (p < 0.001). There were fewer females in the symptomatic C9orf72 group than in the control (X(1) =9.69, p < 0.001) or presymptomatic groups (GRN X(1) = 13.21, MAPT X (1) = 7.18; $\textit{C9orf72} \ \text{X}(1) = 9.40$; all p < 0.007). Symptomatic GRN and symptomatic *C9orf72* were lower educated than controls [F(6,751) =5.74, p < 0.001). MMSE scores were lower in symptomatic mutation carriers than in controls (GRN U = 949, MAPT U = 654; C9orf72 U = 1909; all p < 0.001) and all presymptomatic groups (all p < 0.001). No differences were found amongst the symptomatic or presymptomatic groups (all p > 0.05). CDR® NACC-FTLD scores were higher in symptomatic mutation carriers than in presymptomatic mutation carriers and controls (all p < 0.001), and presymptomatic mutation carriers also had higher CDR® NACC-FTLD scores than controls (GRN U = 15,660, MAPT U = 6380; *C9orf72* U = 13,050; all p < 0.001). Behavioural symptoms were higher in symptomatic mutation carriers than in presymptomatic mutation carriers and controls (all p < 0.001), but also higher in presymptomatic C9orf72 mutation carriers compared to controls (CBI-R U = 12,625.5, p = 0.053; FRS U = 10,677.5, p < 0.001) and presymptomatic *GRN* mutation carriers (CBI-R U = 6121.5, p = 0.008; FRS U = 4967.5, p = 0.004).

3.2. Normative data non-carriers (reference group)

Appendix 2 shows the reference groups' cumulative frequencies (Appendix 2.1), percentile scores (Appendix 2.2), and performance across age, sex and education (Appendix 2.3) for the BCFT Copy, Recall and Recognition. Scores for Copy ranged between 9 and 17; scores for Recall ranged between 6 and 17. 94.5% of controls were able to identify the correct figure in the Recognition trial. Performance below 14 for the Copy trial and below 8 for the Recall trial would be considered outside

Table 1

Demographic and clinical data of the mutation carriers and controls. Values indicate: count (percentage) or mean (standard deviation). Abbreviations: *GRN*, progranulin; *MAPT*, microtubule-associated protein tau; *C9orf72*, chromosome 9 open reading frame 72; MMSE, Mini-Mental State Examination; CDR, clinical dementia rating; NACC, National Alzheimer's Coordinating Center; FTLD, frontotemporal lobar degeneration; CBI-R, Cambridge Behavioural Inventory – Revised; FRS, Frontotemporal Dementia Rating Scale; BCFT, Benson Complex Figure Test.

| Mutation | GRN | | MAPT | | C9orf72 | | Controls | |
|-------------------------------|----------------|-------------|----------------|-------------|----------------|-------------|-------------|--|
| Clinical status | Presymptomatic | Symptomatic | Presymptomatic | Symptomatic | Presymptomatic | Symptomatic | | |
| n | 143 | 41 | 59 | 23 | 130 | 72 | 290 | |
| Age, y [range] | 46.4 (12.4) | 64.0 (8.7) | 39.6 (10.6) | 59.0 (7.1) | 44.7 (13.4) | 62.5 (7.7) | 46.2 (12.9) | |
| Sex, female | 91 (63.6) | 19 (46.3) | 36 (61.0) | 10 (43.5) | 78 (60.0) | 27 (37.5) | 168 (57.9) | |
| Education, y | 14.8 (3.5) | 11.9 (3.5) | 14.4 (3.0) | 13.8 (3.9) | 14.4 (3.0) | 13.0 (3.6) | 14.5 (3.4) | |
| MMSE | 29.4 (1.1) | 22.1 (6.2) | 29.5 (0.9) | 23.8 (6.3) | 29.2 (1.2) | 24.2 (5.0) | 29.4 (1.1) | |
| CDR© NACC-FTLD | 0.2(0.3) | 1.7 (0.9) | 0.2(0.3) | 1.8 (0.9) | 0.2(0.3) | 2.0 (0.9) | 0 (0) | |
| CBI-R | 4.5 (6.9) | 52.3 (28.8) | 7.1 (10.2) | 59.0 (36.6) | 7.4 (9.2) | 65.3 (31.3) | 5.4 (7.9) | |
| FRS, percentage | 96.7 (8.3) | 51.7 (27.9) | 94.4 (9.4) | 44.5 (26.9) | 93.8 (9.7) | 36.8 (29.4) | 96.2 (7.9) | |
| BCFT - copy | 16.5 (1.0) | 14.2 (3.7) | 16.1 (1.1) | 14.3 (3.5) | 16.0 (1.5) | 13.6 (3.5) | 16.3 (1.3) | |
| BCFT - recall | 12.9 (2.7) | 7.7 (4.5) | 12.9 (2.8) | 5.9 (5.3) | 12.8 (2.8) | 8.1 (4.4) | 13.2 (2.8) | |
| BCFT – recognition, % correct | 94.4 | 65.8 | 96.6 | 65.2 | 93.1 | 66.7 | 94.5 | |

the normal range (i.e. \leq 5th percentile). Age ($r_s(288) = -0.09$, p = 0.125) and education ($r_s(288) = 0.11$, p = 0.068) were not significantly associated with BCFT Copy. However, there was a significant correlation between both age ($r_s(288) = -0.45$, p < 0.001) and education ($r_s(288) = 0.13$, p = 0.031) and BCFT Recall. There was a strong positive correlation between BCFT Recognition and age ($\sqrt{\eta^2} = 0.88$); the correlation with education was weak ($\sqrt{\eta^2} = 0.26$). Women had higher BCFT Copy scores than men (mean rank women: 153.9 vs. men: 133.87; U = 8829.5, p = 0.014), whereas there were no sex differences in Recall (U = 9233.5, p = 0.147). Also BCFT Recognition scores did not differ between males and females (X(1) = 1.40, p = 0.237).

3.3. Group differences of the BCFT

Figure 1 shows the group differences in the BCFT Copy, Recall and Recognition between *GRN*, *MAPT* and *C90rf72* mutation carriers according to CDR® NACC-FTLD global score.

For the BCFT Copy, no significant differences were found between groups at CDR® NACC-FTLD global score =0 [F(3,529) $=1.170,\,p=0.321]$ or 0.5 [F(3,370) $=0.751,\,p=0.522].$ However, there were significant differences between groups at CDR® NACC-FTLD global score ≥ 1 [F(3,426) $=10.128,\,p<0.001],$ with both GRN and C9orf72 mutation carriers having lower Copy scores than controls (p=0.001 and p<0.001, respectively). No differences were seen in the MAPT mutation group. Performing a sub-analysis in the CDR® NACC-FTLD global score ≥ 1 group (stratifying into scores of 1, 2 and 3) demonstrated significant differences from a score of 2 onwards in both GRN and C9orf72 (but not MAPT) mutation carriers: at CDR® NACC-FTLD global scores of both 2 and 3 GRN and C9orf72 mutation carriers had lower Copy scores than controls (all p<0.001).

For the BCFT Recall, there were similarly no significant differences between groups at CDR® NACC-FTLD global score = 0 [F(3,529) = 2.390, p = 0.068] or 0.5 [F(3,370) = 1.279, p = 0.281]. However, significant differences were seen between groups at CDR® NACC-FTLD global score ≥ 1 [F(3,426) = 20.469, p < 0.001]: all mutation carrier groups (GRN, MAPT and C9orf72) had significantly lower Recall scores than controls (all p < 0.001). Performing additional sub-analyses in the CDR® plus NACC FTLD score ≥ 1 group (stratified into scores of 1, 2 and 3) demonstrated significant differences in the CDR® NACC-FTLD global score = 1 group in the MAPT mutation carriers only (lower Recall scores than controls: p = 0.024). At CDR® NACC-FTLD global scores of 2 and 3, all mutation carrier groups had lower Recall scores than controls (p-values for scores 2 and 3 respectively: GRN, p = 0.007, p < 0.001; MAPT, p = 0.065, p < 0.001; C9orf72, p = 0.024, p < 0.001). No significant differences at any time point were found between mutation carrier

groups (GRN vs. MAPT, p = 0.872; MAPT vs. C9orf72, p = 0.608; C9orf72 vs. GRN, p = 1.000).

For the BCFT Recognition, there were no significant differences between groups at CDR® NACC-FTLD global score =0 [X(3) =2.982, p=0.394] or 0.5 [X(3) =4.381, p=0.223]. Significant differences between groups were seen at CDR® NACC-FTLD global score ≥ 1 [X(3) =52.924, p<0.001], with all mutation carrier groups having significantly lower Recognition scores than controls (all p<0.001), although no significant differences were found between mutation carrier groups (*GRN* vs. *MAPT*, p=0.830; *MAPT* vs. *C9orf72*, p=0.794; *C9orf72* vs. *GRN*, p=0.974). Additional sub-analyses in the CDR® NACC-FTLD global score ≥ 1 group (stratified into scores of 1, 2 and 3) demonstrated no significant differences at CDR® NACC-FTLD global score =1, but significant differences were seen between all mutation carrier groups and controls at a score of 2 (*GRN* vs. control, p<0.001; *MAPT* vs. control, p=0.038; *C9orf72* vs. controls, p<0.001) and a score of 3 (all comparisons p<0.001).

3.4. Cognitive correlates of the BCFT

Partial correlation coefficients between the BCFT Copy and Recall test score and other relevant neuropsychological tests within the GENFI battery are shown in Table 2. Irrespective of the underlying mutation, both BCFT Copy and Recall test scores correlated significantly with TMT part B and WASI Block Design (p < 0.05). FCSRT immediate and delayed recall also correlated significantly with both BCFT Copy and Recall in every genetic group (p < 0.01), apart from Copy in *C9orf72* mutation carriers. In this mutation, but not in *GRN* and *MAPT*, significant correlations were found between BCFT Copy and Recall test scores and D-KEFS Color-Word Interference Test card III and the letter fluency test (p < 0.05).

3.5. Neuroimaging correlates of the BCFT

The relationships between BCFT Copy and Recall and GM volume are displayed in Fig. 2 and Appendix 3. VBM analyses demonstrated different structures to be involved in BCFT Copy depending on the mutation involved: in *GRN* mutation carriers worse performance correlated with GM atrophy of the left thalamus (p < 0.05 FWE corrected), in *MAPT* mutation carriers with atrophy of the right cerebellum, and in *C9orf72* repeat expansion carriers with atrophy of the left middle frontal gyrus (both p < 0.001 uncorrected). In all mutation carriers, worse BCFT Recall score correlated with atrophy of the temporal lobe, especially the hippocampus (p < 0.05 FWE corrected). In *MAPT* mutation carriers there was additional involvement of the left temporal pole,

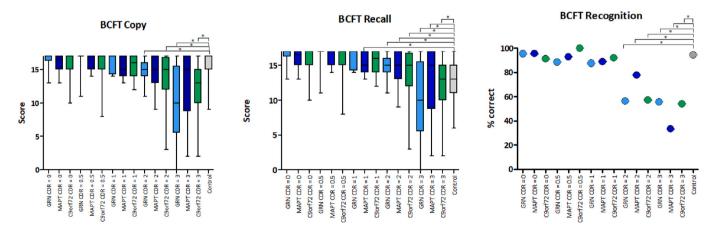


Fig. 1. BCFT Copy, Recall and Recognition data stratified by CDR plus NACC FTLD global score (0, 0.5, 1, 2 and 3) in GRN, MAPT and C9orf72 mutation carriers. Boxplots (for BCFT Copy and Recall) visualize mean (with whiskers representing min-max) scores per clinical group. * p < 0.05. Abbreviations: BCFT, Benson Complex Figure Test; GRN, progranulin; MAPT, microtubule-associated protein tau; C9orf72, Chromosome 9 open reading frame 72; CDR, Clinical Dementia Rating Scale.

Table 2

Partial correlation coefficients (corrected for age, sex, years of education, and family clustering) in GRN, MAPT and C9orf72 mutation carriers between Benson Complex Figure Copy and Recall and other neuropsychological test scores. Significant correlations are displayed in bold; * p < 0.05, ** p < 0.01, *** p < 0.001. Abbreviations: BCFT, Benson Complex Figure Test; GRN, progranulin, MAPT, microtubule-associated protein tau; C9orf72, Chromosome 9 open reading frame 72; TMT, Trailmaking Test; FCSRT, Free and Cued Selective Reminding Test, D-KEFS, Delis-Kaplan Executive Function System; WASI, Wechsler Abbreviated Scale of Intelligence.

| Mutation | GRN | GRN | | MAPT | | C9orf72 | |
|--|---------|----------|----------|---------|----------|----------|--|
| BCFT | Сору | Recall | Сору | Recall | Сору | Recall | |
| TMT part B | -0.18* | -0.49*** | -0.45*** | -0.35** | -0.40*** | -0.31*** | |
| FCSRT immediate recall | 0.41*** | 0.54*** | 0.43*** | 0.72*** | 0.06 | 0.26** | |
| FCSRT delayed recall | 0.40*** | 0.54*** | 0.52*** | 0.76*** | 0.04 | 0.33*** | |
| D-KEFS Color-Word Interference Test card III | -0.06 | -0.21* | -0.13 | -0.21 | -0.25** | -0.28*** | |
| Letter fluency | 0.15 | 0.13 | -0.04 | 0.14 | 0.19* | 0.22** | |
| WASI Block Design | 0.28** | 0.37*** | 0.33** | 0.32** | 0.31*** | 0.31*** | |

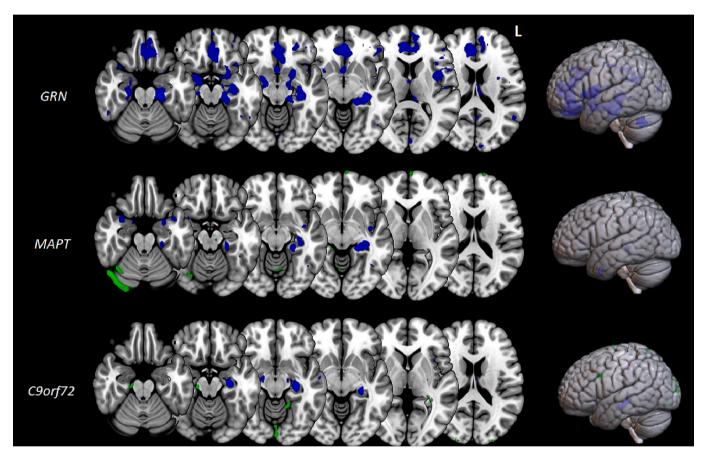


Fig. 2. Neuroimaging correlates of the BCFT Copy and Recall. VBM analyses demonstrated lower scores in BCFT Copy (in green) and BCFT Recall (in blue) to be correlated with lower grey matter volume in *GRN* mutation carriers (top), *MAPT* mutation carriers (middle) and *C9orf72* repeat expansion carriers (bottom). We set the statistical threshold at p < 0.05 (FWE-corrected) for *GRN* copy and all recall conditions, and p < 0.001 (uncorrected) for *MAPT* and *C9orf72* copy. Abbreviations: L, left; *GRN*, progranulin; *MAPT*, microtubule-associated protein tau; *C9orf72*, chromosome 9 open reading frame 72. (For interpretation of the references to color in this figure legend, the reader is referred to the web version of this article.)

whilst in *GRN* mutation carriers, there was also involvement outside of the temporal lobe, including the anterior cingulate, anterior insula, frontal and parietal lobes in particular (both p < 0.05 FWE corrected).

4. Discussion

In this study of a large cohort of participants from genetic FTD families, we have shown lower scores compared to healthy controls in the BCFT Copy, Recall and Recognition abilities of symptomatic mutation carriers, with different profiles depending on the genetic mutation involved. GRN and C9orf72 – but not MAPT – mutation carriers had lower BCFT Copy performance at a CDR® NACC-FTLD global scores of 2

and 3 whereas all mutation carriers had lower BCFT Recall and Recognition scores than controls at those stages, with the addition of earlier impairment of Recall in *MAPT* mutation carriers (from CDR® NACC-FTLD global score of 1). Cognitive correlates of the BCFT Copy and Recall included tests for visuoconstruction, verbal memory, and executive function. Furthermore, lower BCFT Copy score was associated with atrophy of fronto-subcortical areas, while lower BCFT Recall score correlated with predominantly (medial) temporal lobe atrophy.

Our results demonstrate visuoconstructive deficits in FTD mutation carriers only from the moderate dementia stage onwards, reflected in lower BCFT Copy performance at CDR® NACC-FTLD global score of 2 and 3. This is in contrast with a previous study, that showed progressive

decline in BCFT Copy after the CDR = 0.5 stage in patients with bvFTD [32]. A potential explanation for this discrepancy is the use of the original CDR [33], which does not include the behaviour and language domains, and therefore is likely less sensitive for early changes in FTD, i. e. patients with original CDR = 0.5 potentially score higher on the CDR® NACC-FTLD [23], which was used in our study. In our patient sample, BCFT Copy performance was not affected in asymptomatic and prodromal mutation carriers (i.e. CDR® NACC-FTLD global scores 0 and 0.5), which is in line with an earlier study that did not find visuoconstructive decline in presymptomatic mutation carriers [15]. Interestingly, our findings also suggest gene-specific patterns in BCFT Copy performance, in that both GRN and C9orf72 mutation carriers, but not MAPT mutation carriers, had lower scores than controls in the moderate to severe dementia stages. Deficits in visuoconstructive functioning have been described in both symptomatic GRN and C9orf72-related FTD previously [34-36]. Results in the presymptomatic stage have been mixed, with some studies showing early decline [22,37], but not others [38-39]. A recent study into cognitive composites for familial FTD suggested BCFT Copy as part of the neuropsychological battery best discriminating C9orf72 mutation carriers from controls, whereas BCFT Recall was amongst the tests best differentiating MAPT mutation carriers from controls [40]. The latter, as well as our findings, confirms the presence of early memory decline in particularly MAPT mutations, as has also been found in previous studies [8,41]. In contrast to studies demonstrating verbal memory deficits in presymptomatic MAPT [15-16,42], we only found significant differences in BCFT Recall (i.e. visual memory) from CDR® NACC-FTLD global score of 1. A potential explanation for this discrepancy could be the difference between performances on verbal versus visual memory tests. Because of the early semantic memory involvement in MAPT-related FTD [43], language-led tests could be more sensitive to change in the presymptomatic stage than visuoconstructive-mediated tests.

The cognitive and neuroimaging correlates of the BCFT Copy and Recall showed both cross-mutation as well as mutation-specific patterns. Irrespective of the underlying mutation, BCFT scores correlated with tests for visuoconstruction, verbal memory, and executive function, with stronger executive function involvement in C9orf72. These findings suggest two important aspects about the BCFT, namely that it - as previous research suggested [19-20] - assess multiple cognitive functions, allowing the exploration of differential mechanisms of cognitive impairment in familial FTD, and also specifically taps into frontallymediated skills in C9orf72. This is an interesting finding, as BCFT Copy performance indeed correlated with atrophy of the left middle frontal gyrus in this mutation. Although early atrophy of the thalamus and cerebellum is commonly regarded as the neuroimaging signature of C9orf72 [14,44], the associations we found with the thalamus (in GRN) and cerebellar (in MAPT) atrophy confirm that subcortical involvement is also present in the other two FTD genetic groups [45], and leads to lower visuoconstructive scores. In all mutation carriers, worse BCFT Recall score correlated with atrophy of the (medial) temporal lobe. This is not a surprising finding, given the pivotal role of the hippocampus in memory recall, and indeed previous studies into the Rey Complex Figure Test, similar to the BCFT, have related recall performance to medial temporal lobe structures including the hippocampus [46]. In MAPT mutation carriers there was specific involvement of the temporal pole. This finding coincides with the lower BCFT Recall performance relatively early in the disease process of this mutation, confirming MAPT-FTD as a predominantly temporal-predominant disease [14].

Key strengths of our study are the large sample sizes of presymptomatic and symptomatic *GRN*, *MAPT* and *C9orf72* mutation carriers and non-carriers from the same families. Not only is the non-carrier group an ideal control group as they have the same genetic and social background as the mutation carriers, we were also able to generate new normative data and relationships with age, sex and education for the BCFT. Despite large numbers, some groups (especially *MAPT* mutation carriers) remain relatively small when dividing the sample according to

CDR® NACC-FTLD global scores, so that replication in other familial FTD cohorts (e.g., ALLFTD, DINAD) is warranted. We were unable to detect any changes in the CDR® NACC-FTLD global score = 0.5 group, which might have been the result of the heterogeneous nature of this category, likely including mutation carriers without overt dementia symptoms as well as people with primary psychiatric disorders and early-stage PPA, in which it is difficult to detect clinical features [23]. Directions for future research include modifications to traditional scoring methods (i.e., accuracy and placement), such as incorporating process (e.g., direction and order of drawing) and/or digital scoring methods to increase test sensitivity in early disease stages [47] and to allow the measurement of the different cognitive processes that the BCFT relies on (i.e., visuospatial abilities, visual memory, and executive functions such as organization and working memory) but currently cannot be separated.

5. Conclusion

Our study showed lower BCFT Copy, Recall and Recognition performance in symptomatic FTD mutation carriers in comparison to non-carriers. We demonstrated copy deficits in symptomatic *GRN* and *C90rf72* mutation carriers, whereas recall was affected in the early-symptomatic period in *MAPT* mutation carriers, suggesting differential mechanisms of cognitive impairment depending on the genetic mutation involved, which was corroborated by specific cognitive and neuro-imaging correlates. Performance on this brief and easy-to-apply test may aid in differential diagnosis in genetic FTD, but its potential as candidate cognitive biomarker for upcoming clinical trials is most likely limited as impaired performance on the BCFT occurs relatively late in the genetic FTD disease process.

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Declaration of Competing Interest

RSV has served in Advisory boards Meetings for Wave Life Sciences, Ionis and Novo Nordisk and received personal fees for participating in educational activities from Janssen, Roche Diagnostics, and Neuraxpharma and funding to her institution for research projects from Biogen and Sage Pharmaceuticals. The other authors declare that they have no competing interests.

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Appendix B. Cumulative frequencies, percentile scores, and performance across age, sex and education for the Benson Complex Figure Test (BCFT) in the reference (non-carrier) group (n = 290)

Appendix 2.1 – Cumulative frequencies for the BCFT Copy, Recall and Recognition in the reference group.

| BCFT Copy | | | BCFT Recall | | | |
|-----------|----------|--------------------------|-------------|----|--------------------------|--|
| Score | n | Cumulative frequency (%) | Score | n | Cumulative frequency (%) | |
| 9 | 1 | 0.3 | 6 | 4 | 1.4 | |
| 11 | 1 | 0.7 | 7 | 5 | 3.1 | |
| 12 | 1 | 1.0 | 8 | 14 | 7.9 | |
| 13 | 8 | 3.8 | 9 | 10 | 11.4 | |
| 14 | 18 | 10.0 | 10 | 20 | 18.3 | |
| 15 | 49 | 26.9 | 11 | 22 | 25.9 | |
| 16 | 13 | 31.4 | 12 | 30 | 36.2 | |
| 17 | 199 | 100 | 13 | 41 | 50.3 | |
| | BCFT Rec | ognition | 14 | 41 | 64.5 | |
| Score | n | Cumulative frequency (%) | 15 | 46 | 80.3 | |
| 0 | 16 | 5.5 | 16 | 10 | 83.8 | |
| 1 | 274 | 100 | 17 | 47 | 100 | |

Appendix 2.2 – Percentile scores of the BCFT Copy and Recall in the reference group.

| Percentile | BCFT Copy | BCFT Recall | |
|------------|-----------|-------------|--|
| 5th | 14 | 8 | |
| 10th | | 9 | |
| 20th | 15 | 11 | |
| 30th | 16 | 12 | |
| 40th | 17 | 13 | |
| 50th | | | |
| 60th | | 14 | |
| 70th | | 15 | |
| 80th | | | |
| 90th | | 17 | |

Appendix 2.3 - BCFT performance across age, sex, and education in the reference group. Abbreviation: SD, standard deviation.

| | BCFT Copy | | BCFT Recall | | BCFT Recognition | |
|-------------------|-----------|---------------------|-------------|--------------------|------------------|--|
| | n | Mean (SD); [range] | n | Mean (SD); [range] | n (% correct) | |
| Age group (years) | | | | | | |
| 18.1-29.9 | 33 | 16.4 (1.2); [13–17] | 33 | 15.0 (2.4); [6–17] | 32 (97.0) | |
| 30.0-39.9 | 68 | 16.1 (1.5); [9–17] | 68 | 13.6 (2.4); [8–17] | 65 (95.6) | |
| 40.0-49.9 | 83 | 16.5 (1.1); [12–17] | 83 | 13.2 (2.7); [6–17] | 78 (94.0) | |
| 50.0-59.9 | 54 | 16.5 (0.9); [14–17] | 54 | 13.2 (2.7); [7–17] | 51 (94.4) | |
| 60.0-69.9 | 42 | 15.9 (1.3); [13–17] | 42 | 11.7 (2.7); [7–17] | 40 (95.2) | |
| 70.0–85.0 | 10 | 15.6 (2.0); [11–17] | 10 | 10.2 (3.2); [7–17] | 8 (80.0) | |
| Education (years) | | | | | | |
| 0–9 | 24 | 16.1 (1.2); [14–17] | 24 | 12.1 (2.8); [8–17] | 23 (95.8) | |
| 10-12 | 60 | 16.1 (1.2); [13–17] | 60 | 13.2 (2.6); [7–17] | 54 (90.0) | |
| 13-16 | 135 | 16.2 (1.5); [9–17] | 135 | 13.1 (2.9); [6–17] | 129 (95.6) | |
| ≥17 | 71 | 16.5 (0.9); [14–17] | 71 | 13.6 (2.6); [6–17] | 68 (95.8) | |
| Sex | | | | | | |
| Female | 168 | 16.4 (1.1); [12–17] | 168 | 13.0 (2.7); [6–17] | 161 (95.8) | |
| Male | 122 | 16.0 (1.5); [9–17] | 122 | 13.4 (2.9); [6–17] | 113 (92.6) | |

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