

Short-Term Memory Conjunctive Binding in Alzheimer's Disease: A Systematic Review and Meta-Analysis

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ABSTRACT

Objective: Short-term memory (STM) binding tests assess the ability to temporarily hold conjunctions between surface features, such as objects and their colors (i.e., feature binding condition), relative to the ability to hold the individual features (i.e., single feature condition). Impairments in performance of these tests have been considered cognitive markers of Alzheimer's disease (AD). The objective of the present study was to conduct a meta-analysis of results from STM binding tests used in the assessment of samples mapped along the AD clinical continuum. **Methods:** We searched PubMed, Scopus and Web of Science for articles that assessed patients with AD (from preclinical to dementia) using the STM binding tests and compared their results with those of controls. From each relevant article, we extracted the number of participants, the mean and standard deviations from single feature and of feature binding conditions. Results across studies were combined using standardized mean differences (effect sizes) to produce overall estimates of effect. **Results:** The feature binding condition of the STM binding showed large effects in all stages of AD. However, small sample sizes across studies, the presence of moderate to high heterogeneity and cross-sectional, case-controls designs decreased our confidence in the current evidence. **Conclusions:** To be considered as a cognitive marker for AD, properly powered longitudinal designs and studies that clearly relate conjunctive memory tests with biomarkers (amyloid and tau) are still needed.

Key points

Question: What is the magnitude of difference in short-term memory conjunctive binding tests between controls and patients in the Alzheimer's disease continuum? **Findings:** We found large differences in all Alzheimer's disease stages compared with controls, from the effect size of -1.10 (preclinical stage) to -2.40 (dementia stage). **Importance:** Short-term memory conjunctive binding test can contribute to the early detection of Alzheimer's disease. **Next Steps:** Longitudinal studies are needed relating the conjunctive tests to biomarkers (amyloid and tau).

Keywords: short-term memory binding; memory binding; conjunctive memory; working memory; Alzheimer's Disease; familial Alzheimer's Disease; mild cognitive impairment; preclinical Alzheimer's disease; subjective cognitive decline; systematic review; meta-analysis.

Over the last two decades, the role of feature binding in memory has attracted considerable interest and is now better understood (Luck & Vogel, 1997; Treisman, 2006; Vogel, Woodman, & Luck, 2001; Wheeler & Treisman, 2002). Binding, or the building of a mental representation of combinations of features such as color and shape, names and shapes, or word pairs, supports memory widely, across systems i.e., short-term and working memory (Allen, Baddeley, & Hitch, 2006; Baddeley, 2007a) and long term memory (LTM) (Buschke, 2014; Moses, Cole, & Ryan, 2005), and domains (i.e., verbal (Baddeley, 2001, 2007b; Baddeley, Hitch, & Allen, 2009) and visual (Baddeley, Allen, & Hitch, 2011; Hitch, Allen, & Baddeley, 2020; Hollingworth & Rasmussen, 2010; Logie, Brockmole, & Vandembroucke, 2009; Shimi & Logie, 2019)). Different models have been proposed to explain how binding supports the representation, formation, and use of memory. The two most prevalent are the slot model (Luck & Vogel, 2013; Rouder, Morey, Morey, & Cowan, 2011) and the resource model (Bays, Wu, & Husain, 2011; Heinen et al., 2016; Liang et al., 2016), although a comparison of these models is outside the scope of the current paper.

In this systematic review and meta-analysis, we focus on the formation and temporary retention in short-term or working memory of arbitrary combinations of features referred to as temporary conjunctive binding. This contrasts with research that has focused on the learning of associations between features (e.g., Moses & Ryan, 2006; Barnett et al., 2015; Bier et al., 2008; Blackwell, et al., 2004; Liang et al., 2016; for reviews see Zimmer, Mecklinger, & Lindenberger, 2006; Schneegans & Bays, 2019). However, whereas there is a substantial and growing volume of research on short-term binding in healthy adults (for a recent review see Hakim, Awh, & Vogel, 2021), there are relatively fewer studies of temporary feature binding in patient groups (see e.g. Parra et al., 2015; van Geldorp, Parra & Kessels, 2014). In particular, we focus on how the temporary binding of visual features is impacted by pathological aging, in particular Alzheimer's disease (AD). AD is a neurodegenerative disease that progressively impairs cognition and functionality (Albert et al., 2011; McKhann et al., 2011), and it is the main cause of dementia in older adults (Ferri et al., 2005; Prince et al., 2014; Reitz & Mayeux, 2014; Cao et al., 2020).

Relational and conjunctive binding

Individual features can be bound in memory by means of two mechanisms: relational and conjunctive. Relational binding refers to the ability to associate stimuli in memory, whereby the individual elements forming such associations retain their original identity (Mayes et al., 2007). Conjunctive memory binding, on the other hand, refers to the ability to integrate stimuli or their features into unified representations (see Moses & Ryan, 2006 for evidence from long-term memory and Wheeler & Treisman, 2002 for evidence from working memory). To recognize a blue car in the car park, we do not need to recall the association between 'blue' and 'car' as separate features; rather we remember the car as a unique object (a blue car) whose

identity differs from that of other neighboring objects. While altering a constituent part in a conjunctive memory representation leads to the formation of a new identity (a red car is different from a blue car), changing a part in a relational representation modifies the nature of the association but not the identity of its parts (see for example Mayes et al., 2007; Moses & Ryan, 2006). The dissociation between relational and conjunctive bindings has been shown in a series of single-case studies of patients with specific impairments on relational but not on conjunctive binding and vice versa (Vargha-Khadem et al., 1997; Baddeley et al., 2010; Parra et al., 2009a, 2011a, Parra et al., 2015a; Jonin et al., 2019), and this dissociation is relevant because it translates into distinct aging effects and associations with brain areas. A further crucial dissociation is between long-term learning of conjunctive bindings (e.g. learning that your car is blue) and temporary combinations of features that may change on a moment to moment basis, such as the color of cars around you on a busy motorway, or whether a participant in an experiment or cognitive test is presented with a blue square and a red circle to remember on one trial, but a blue circle and a red square on a subsequent trial.

Relational binding is affected by normal aging, both in LTM (Naveh-Benjamin et al., 2007; Naveh-Benjamin et al., 2004ab; for a review, see Old & Naveh-Benjamin, 2008) and short-term memory (STM) (Chen & Naveh-Benjamin, 2012; Cowan et al., 2006; Fandakova et al., 2014; Mitchell et al., 2000). Short-term, or temporary conjunctive binding, on the other hand, has consistently shown to be insensitive to age (Bastin, 2018; Brockmole & Logie, 2013; Brockmole et al., 2008; Brown et al., 2017; van Geldorp et al., 2015; Hoefeijzers et al., 2017; Isella et al., 2015; Killin et al., 2018; Kirmsse et al., 2018; Yassuda et al., 2020). It is also important to note that temporary conjunctive binding is not affected by literacy (Yassuda et al., 2020).

Relational and conjunctive memory bindings are also subsumed by different neuronal activations. Relational binding requires the work of the hippocampus (Gold et al., 2006;

Hannula et al., 2006; Kan et al., 2007; Monti et al., 2015; Nichols et al., 2006; Olsen et al., 2012; Olson et al., 2006; Yonelinas, 2013), whereas conjunctive STM binding does not (Baddeley et al., 2010; Parra et al., 2014; Piekema et al., 2010; Staresina & Davachi, 2010; Valdés Hernández et al., 2020; Xu, 2007). In the latter case, there are short intervals between study and test phase, such as one second, and a small number of items in the study display for subsequent recognition or recall (e.g., Jeneson et al., 2012).

Piekema et al. (2010) found that the medial temporal lobe (MTL) was not activated when people perform intrinsic intra-item bindings (color-object), but the inter-item associations yielded MTL activation. Visual short-term memory for conjunctive bindings seems instead to be associated with posterior areas of the brain, especially regions within the parietal and occipital lobes (Parra et al., 2014; Todd & Marois, 2005; Shafritz et al., 2002; Song & Jiang, 2006; Staresina & Davachi, 2010; Xu, 2007).

Conjunctive short-term memory binding paradigms

Different paradigms assessing conjunctive STM binding have been used in clinical settings with patients. The main difference between them relies on the retrieval method: recognition or free recall. In recognition tasks using the change detection paradigm, participants assess a test screen and decide if the stimuli are the same or different from those presented in the previous screen (study phase). In free recall, participants are asked to say aloud the names of the stimuli they have just seen in the study screen. In addition to these differences, the tasks vary in terms of presentation time, in the study screen and in the number of items presented per trial.

Several studies compared single feature condition versus feature binding condition. The single feature condition refers to a task in which the stimuli are presented as individual features (color-only, shape-only). The participant should memorize and retrieve each individual feature. For

instance, in the shape-only task using the change detection paradigm, participants should memorize shapes (study phase) and then recognize if the shapes presented in the test phase are the same or different. The feature binding condition, on the other hand, requires participants to memorize and retrieve features integrated within object representations, such as the specific color-shape combinations for colored shapes or colored objects (i.e., bindings).

Conjunctive binding as a cognitive marker for AD

The identification of a sensitive and specific cognitive marker of AD will ultimately aid its differential diagnosis and assist the early detection of the disease, as well as its follow-up (Logie, Parra, & Della Sala, 2015). Improving early diagnosis and care of patients with dementia is a current primary target for the National Institutes of Health in the US and for the National Health Service within the UK, and will no doubt continue to be a national and international priority. It has been suggested that the STM binding test could significantly contribute to the early detection of AD (Logie, Parra, & Della Sala, 2015), showing not only high sensitivity but also high specificity for AD (Costa et al., 2017; Martínez, Trujillo, Arévalo, Ibáñez, & Cardona, 2019; Rentz et al., 2013). Therefore, it is important to review how conjunctive binding tests can detect impairments in each stage of the AD continuum (Sperling et al., 2011; 2013).

Previous reviews addressed this issue only partly. Rentz et al. (2013) carried out a selective review about tests promising to detect preclinical AD, showing that poor performance in tests like the Memory Capacity, Face-Name Association, Spatial Pattern Separation and Discrimination and Transfer was associated with the presence of biomarkers for AD. In addition, a range of studies has found that the dual-tasking and the STM binding tests could discriminate preclinical AD patients from controls (e.g., Della Sala, Foley, Parra & Logie,

2011; Kaschel, Logie, Kazén, & Della Sala, 2009; Logie, Cocchini, Della Sala, & Baddeley, 2004). Fuller et al. (2019) reviewed the literature on cognitive and biological markers of familial AD, showing that structural and functional brain abnormalities could be found in preclinical AD patients, as well as cerebral spinal fluid biomarkers. In addition, they highlighted cognitive impairments in preclinical AD patients, among which were deficits on the STM binding test. Martínez et al. (2019) conducted a broader review on theoretical cognitive models of conjunctive binding, also reviewing differences in test paradigms, brain areas associated with them, and the clinical use of the conjunctive STM binding in dementia. Pavisic et al. (2020) presented arguments to use the visual binding tasks, including relational and conjunctive types, in clinical settings. However, none of these reviews specifically addressed the issue of the clinical use of conjunctive STM binding in the AD continuum, from preclinical to dementia stages. Moreover, none of the previous reviews was conducted using transparent and reproducible methods or performed a meta-analysis of the results of multiple studies. As early detection of AD was the main goal driving the development of the conjunctive memory binding tasks, it is essential to review the evidence available to establish where the current knowledge sits and inform future research.

We aimed to analyze studies that have reported on one of the three stages of AD described in the literature: i) preclinical/subjective cognitive decline, during which patients do not show impairments in common cognitive measures (Dubois et al., 2016; Jessen et al., 2014; Koppa, et al., 2015a; Reisberg et al., 2010; Sperling et al., 2011) ii) prodromal: mild cognitive impairment (MCI - Mitchell and Shiri-Feshki, 2009; Petersen, 2004), which defines patients with high risk to convert to AD and iii) the clinical stage of full blown dementia (Jack et al., 2018; McKhann et al., 2011).

To perform a meta-analysis of studies using the STM binding in the context of AD is not an ill-posed question which confounded dementia with specific diseases (Della Sala & Morris,

2020), like AD, since the STM binding deficits have been suggested to be specific to AD. Other reasons for a review on a specific cognitive marker of AD are that 1) AD is the most common cause of dementia (Prince et al., 2014); 2) biomarkers and neuroimaging techniques are expensive and not sufficiently available in most developing countries, or even in remote areas of developed countries. Therefore, searching for cognitive markers of AD, especially in early stages, is an important goal.

Objectives

The objectives of the present study were to systematically review the current evidence in the literature and combine the results of studies assessing the clinical use of the STM conjunctive binding tests in the context of the AD clinical continuum in a meta-analysis.

Methods

This systematic review and meta-analysis was conducted according to the recommendations of the Cochrane Handbook for Reviews of Interventions (Higgins et al., 2019) and reported in adherence with the principles of the Preferred Reporting Items for Systematic Reviews and Meta-Analyses (PRISMA) statement (Moher et al., 2009).

Criteria for considering studies for this review

Types of studies

Studies that used a conjunctive STM binding test with patients at any stage of AD were considered in this review. We considered AD on a continuum according to the level of

cognitive impairment from preclinical/subjective cognitive decline, to MCI up to full blown dementia. It is worth mentioning that most of these studies preceded the recent framework proposed for the biological definition of AD (Jack et al., 2018). We did not filter by the type of study design, but only cross-sectional case-control studies were found.

Participants

Adults at one of the following stages of AD preclinical/subjective cognitive decline, MCI or AD dementia. Furthermore, in this review we are using the term AD in a broader sense, acknowledging that the reviewed studies used different diagnostic criteria, such as the McKhan et al.'s (1984), McKhan et al.'s (2011) and the biological definition of Jack et al. (2018). In addition, studies involving the familial variant of AD recruited patients with genetic mutations E280A-PSEN1 that leads to early-onset autosomal dominant AD (Lopera et al., 1997). Different criteria for MCI were also considered, such as Petersen et al. (2004), Winblad et al. (2004), Albert et al. (2011) and for the familial MCI the Acosta-Baena et al. (2011).

Types of interventions

We aimed to analyze the performance of patients within the AD-continuum and related healthy controls on conjunctive STM binding tests. Different paradigms were included: change detection (Parra et al., 2010a, 2010b, 2011b, 2015b, 2017a, 2019; Koppa et al., 2015b; Della Sala et al., 2016; Pietto et al., 2016; Fernández et al., 2018; Cecchini et al., 2020; Kozlova et al., 2020; Norton et al., 2020; Valdés Hernández et al., 2020; Cecchini et al., 2021; Fernández & Parra, 2021; Martínez-Flores et al., 2021), free recall (Parra et al., 2009b; Della Sala et al., 2012; Cecchini et al., 2017a; Cecchini et al., 2020) and cued recall modalities (Guazzo et al.,

2020), different set sizes (i.e., from 2 to 4 items per screen), different stimuli (e.g., unnameable shapes or objects) and different amount of trials (from 6 to 32 trials).

Type of outcome measures

The primary outcome measures were the effect sizes for the scores of the STM binding tasks in controls and patients in AD continuum.

Search methods for identification of studies

Electronic searches

Comprehensive searches were designed using appropriate subject headings and free text terms. We searched PubMed, Scopus and Web of Science on 08 June 2020 with a combination of terms for memory binding (“short-term memory binding”, “memory binding”, “conjunctive memory”, “working memory binding”) and appropriate terms for dementia (“Alzheimer’s disease”, “mild cognitive impairment” and “subjective cognitive decline”). We performed another search on 24 November 2021 to check for new papers. No restrictions on date or language of publication were applied to the searches. All references were exported to StArt (Fabbri et al., 2016) for recording and deduplication. Further details of electronic searches are given in online Supplemental Material (S1).

Searching other resources

The reference lists of all selected studies were screened for additional studies and experts in the field contacted for further reports. Lists of included and excluded studies are fully presented in the online Supplemental Material (S2 and S3, respectively).

Data collection and analysis

Selection of studies

The titles and abstracts identified by the search strategies for eligibility were assessed and any disagreements were resolved by discussion. Articles were selected according to the following inclusion criteria: 1) studies assessing the conjunctive memory binding; 2) studies with patients at any stage of the AD continuum (pre-clinical/subjective cognitive decline; MCI; AD dementia). Articles were not deemed suitable for inclusion if they 1) dealt with other neuroscience or cognitive topics; 2) dealt with relational, associative, or other types of binding; 3) were reviews, opinion articles, single-case reports, or conference proceedings; 4) did not include an AD group. All potentially relevant articles were retrieved in full.

Data extraction and management

The search results were extracted to the StArt program (Fabbri et al., 2016), which was used to manage and select the articles. We extracted information on the testing procedure and methods (items per trial, quantity of trials, the encoding time, the type of single feature condition and the measure used in the study, sample size), the characteristics of participants (mean age and education level) and the target condition (diagnostic criteria used to categorize patients according to the stages of AD). The single feature condition refers to a task in which the stimuli are presented as isolated features (color-only, shape-only or objects and colors). To assess the

participants' performance on the single feature condition and feature binding condition of the STM binding tasks, we extracted mean scores for each tested group along with standard deviations and the number of participants. Parra et al. (2017a) study used the same MCI-FAD sample from Pietto et al. (2016), therefore the former was excluded from the meta-analysis.

Assessment of risk of bias and quality of evidence

The risk of bias of included studies was assessed using the ROBINS-I criteria (Higgins et al., 2019; Sterne et al., 2016). The confidence in the certainty of identified evidence was assessed using the Grading of Recommendations, Assessment, Development and Evaluation (GRADE) criteria (Guyatt et al., 2011; Higgins et al., 2019). The level of certainty was lowered if one of the following aspects were present: risk of bias, inconsistency, indirectness, imprecision, or publication bias. All studies included in this review were case-controls studies, which were considered as having low certainty of the effect estimation from the start due to their study design. We considered the 95% confidence intervals: if the lower and upper bound had different meanings (e.g., large effect for the upper bound and no effect for the lower bound), lowering the certainty was considered based on imprecision. The level of certainty was increased if the studies showed a large effect size or when it was considered plausible that the confound variables had undermined the potential effect size. Both the imprecision and increment in certainty judgements were based primarily on the feature binding conditions, as they were the focus of the included studies. The results of the risk of bias assessment and grading of evidence for each of the included studies are presented in online Supplemental Material (S4).

Measures of treatment effect

To assess the performance of controls and AD patients on the STM binding tasks we used mean differences and standard deviations of single feature condition and feature binding condition tasks. We included both tasks because the single feature condition was used as a control task against which the feature binding condition was compared to assess the presence of binding deficits. Age and level of education were considered possible confounded variables and notated in the Characteristics of Included Studies Table (see Table 1).

Unit of analysis issues

Each comparison between controls and AD patients was considered relevant. When a study had more than one comparison, such as comparing controls with preclinical AD and controls with AD, both comparisons were considered in the meta-analysis. The same occurred when a study used more than one experiment, comparing controls and patients in two different experimental settings. When a study had different test characteristics, such as comparing controls and patients with two or three items per screen, each comparison was considered as one unit of analysis.

Dealing with missing data

Mean and standard deviation values were not reported in ten papers (Della Sala et al., 2012; Fernández et al., 2018; Fernández & Parra, 2021; Martínez-Florez et al., 2021; Norton et al., 2020; Parra et al., 2009b; Parra et al., 2010a; Parra et al., 2017a; Parra et al., 2015b; Pietto et al., 2016). The authors of these papers, who were contacted for further information, kindly provided the relevant sets of data. We did not get the data from Fernández and Parra (2021),

but they used the same sample from Fernández et al. (2018), adding five more participants in each group, therefore, only the Fernández et al.'s (2018) data were included in the meta-analyses.

Assessment of heterogeneity

Heterogeneity between studies was assessed visually by inspection of the forest plots and statistically using the I^2 (on a scale from 0% to 100%) and the Chi-squared statistics (Higgins et al., 2019). The I^2 represents the variability in the effects that is due to heterogeneity (Borenstein et al., 2017). If $I^2 = 0$, all variability in the effect sizes is explained within studies instead of between studies (heterogeneity) (Huedo-Medina et al., 2006). An I^2 less than 40% was considered to indicate low inconsistency; 50% to 75% indicated moderate inconsistency, and greater than 75% considerable inconsistency (Higgins & Thompson, 2002; Higgins et al., 2019).

Assessment of reporting biases

We assessed the presence of publication bias, which refers to the fact that studies with negative results have less chance to be published, using funnel plots when at least ten studies assessing the same outcomes were identified. We acknowledge, however, that poor methodological study design (e.g., case-control design) can be an important source of funnel plot asymmetry.

Data synthesis

As studies used different number of items for the STM binding tasks, we calculated standardized mean differences. If appropriate, the results of included studies were combined in

random-effects meta-analyses to produce overall estimates of effect. The decision to combine data in meta-analyses was dependent upon the availability of outcome data and the heterogeneity observed between studies. The random-effects model assumes that the effect sizes represent a random sample from a distribution of these effect sizes, considering that there may be different effect sizes underlying different studies (Borenstein et al., 2010). We decided to use random effects models because between-study heterogeneity was expected due to the variability in experimental methods and sample characteristics. We combined studies using the inverse of the variance weighted approach and presented summary estimates alongside 95% confidence intervals.

To assess the effect size, the Hedges' g formula was used due to differences in sample sizes between the groups and because this formula is better for small sample sizes when compared to the Cohen's d (Cooper et al., 2009). To interpret the effect sizes, we used the following criteria (Cohen, 1988): 0.2 = small, 0.5 = medium and 0.8 = large.

Separate meta-analyses were performed according to the different stages of AD (i.e., preclinical AD /subjective cognitive decline versus controls, MCI versus controls, sporadic and familial AD versus controls).

The analyses were performed using the meta package v4.16-1 (Schwarzer, 2007) in R v.4.0.2 and the Review Manager (2020) software version 5.4.1.

'Summary of findings' table

The main results of this systematic review are shown in the 'Summary of findings' Table (Table 1). The table presents the magnitude of effects of the single feature condition and feature binding condition according to the different stages of AD, the total number of studies and participants, and information on the quality of evidence.

Subgroup analysis

The main aspects of the included studies that could increase heterogeneity and therefore reduce the strength of the conclusions were investigated using subgroup analyses. The following subgroup analyses were performed: 1) single feature condition vs. feature binding condition; 2) change detection vs. free recall tasks; 3) familial AD vs. sporadic AD; 4) Shape-color vs. color-color binding tasks; 5) titration vs. no-titration difficulty of the tasks between controls and patients. All these analyses were performed within each diagnostic group (e.g., preclinical/subjective cognitive decline, MCI and AD dementia). Analyses 2 to 5 were done for single feature conditions and feature binding conditions separately. A p-value ≤ 0.10 was considered significant (Richardson et al., 2019).

Results

Description of studies

Literature search results

In total, 320 reports were identified by the search strategies. Of the 320 identified reports, 301 were subsequently excluded for the following reasons: 103 were duplicates; 145 investigated neuroscience or cognitive topics not relevant to the purpose of this review; 27 focused on relational, associative, or other types of binding; 17 were reviews or opinion papers; 7 did not include an AD clinical group; 2 were conference proceedings, single-cases or we did not have access to the data. At last, one paper was included from perusing the lists of references. In total, 20 articles were considered in this review. The included and excluded studies are described in

online Supplemental Material (S2 and S3). Figure 1 presents the PRISMA flow diagram of study selection.

[INSERT FIGURE 1 HERE]

Included studies

Table 1 presents the summary of findings for the main comparisons and Table 2 presents the characteristics of the included studies. The studies varied in terms of the modality of the STM binding assessment, using either free recall, change detection or cued recall. They also used different set sizes in the task, ranging from 2 to 4 items per trial, and some studies titrated the cognitive load by using different set sizes for the control and experimental groups. All studies reviewed used small sample sizes, except for Martínez-Flores et al. (2021), which included 109 controls and 45 MCIs. The other studies varied from 6 to 37 participants within each group, with a mean sample size of 22.24 and a median of 23 participants per group.

Stimuli presentation time varied from 500 milliseconds to 12 seconds across studies. Concerning the change detection tasks, the number of trials varied from 32 to 100 across studies and most of them used 2000ms duration of the study phase, with only a few using 500ms. Across studies, the performance of participants in the change detection task was measured either using the number of correct responses, the proportion of correct responses, using A' (a measure of sensitivity, see Xu, 2002, pg. 1264 for the formula to calculate it), hit minus false alarms (corrected recognition) or Beta (β) (Stanislaw & Todorov, 1999). For the free recall

task, all studies used 6 trials and the same study time per feature (1.5 seconds). For the cued recall task, a 12-second delay between encoding and recall phases was used.

[INSERT TABLE 1 HERE]

[INSERT TABLE 2 HERE]

Meta-analysis results

AD / Familial AD versus Controls, single feature condition

Figure 2 shows the results of the single feature condition of the STM binding test for the comparison between sporadic/familial AD (343 patients in total) and controls (332 controls in total). The overall pooled standardized mean difference was -1.18 (95% CIs -1.61, -0.76). Substantial heterogeneity was observed between studies ($I^2 = 84\%$, $p < 0.01$). The analysis of the funnel plot (S5.1, online Supplemental Material) shows that there is no clear evidence of publication bias.

[INSERT FIGURE 2 HERE]

AD / Familial AD versus Controls, feature binding condition

Figure 3 shows the results of the feature binding condition of the STM binding test for the comparison between patients with sporadic/familial AD (390 patient in total) and controls (374 controls in total). Substantial heterogeneity was observed between studies ($I^2 = 78\%$, $p < 0.01$).

The overall pooled standardized mean difference was -2.41 (95% CIs -2.82, -1.99) ranging from -4.17 to -1.34 across studies. The analysis of the funnel plot (S5.2, online Supplemental Material) shows that there is no clear evidence of publication bias.

[INSERT FIGURE 3 HERE]

MCI / MCI-FAD versus controls, single feature condition

Figure 4 shows the results of the single feature condition of the STM binding test for the comparison between patients with MCI and MCI-FAD (249 in total) and controls (379 in total). Moderate heterogeneity was observed between studies ($I^2 = 53%$, $p < 0.01$). The overall pooled standardized mean difference was -1.08 (95% CIs -1.35, -0.81); effect sizes ranged from -1.97 to -0.37 across studies. The analysis of the funnel plot (S5.3, online Supplemental Material) shows that there is no clear evidence of publication bias.

[INSERT FIGURE 4 HERE]

MCI / MCI-FAD versus controls, feature binding condition

Figure 5 shows the results of the feature binding condition of the STM binding test for the comparison between patients with MCI and MCI-FAD (363 in total) and controls (565 in total). The overall pooled standardized mean difference was -1.07 (95% CI -1.32, -0.82); effect sizes ranged from -2.52 to -0.61 across studies. Moderate heterogeneity ($I^2 = 63%$, $p < 0.01$) was

observed across studies. It is worth noting that two studies used different set sizes and compared either two or three stimuli per screen (Parra et al., 2019; Valdés Hernández et al., 2020). In each study, the number of stimuli presented per screen did not affect the estimate of effect with similar standardized mean differences between the experimental and control groups. The analysis of the funnel plot (S5.4, online Supplemental Material) shows evidence of asymmetry with two studies outside the plot.

[INSERT FIGURE 5 HERE]

Preclinical AD and subjective cognitive decline versus controls, single feature condition

Figure 6 shows the results of the shape-only condition of the change detection modality of the STM binding test between patients with subjective cognitive decline (SCD)/preclinical AD (111 in total) and controls (130 in total). The overall standardized mean difference mean was -0.33 (95% CI -0.59, -0.07), with effect sizes ranging from -0.51 to -0.21 across studies. No statistical heterogeneity was observed across studies ($I^2 = 0\%$, $p = 0.93$).

[INSERT FIGURE 6 HERE]

Preclinical AD and subjective cognitive decline versus controls, feature binding tasks

Figures 7 present the results of the shape-color binding condition of the change detection modality of the STM binding task for patients with subjective cognitive decline and preclinical AD (111 in total) and for controls (130 in total). The overall standardized mean difference was

-1.10 (95% CI -1.48, -0.73) with effect sizes ranging from -1.52 to -0.56 across studies. Low inconsistency was observed across studies ($I^2 = 46\%$, $p = 0.11$).

[INSERT FIGURE 7 HERE]

Subgroup analyses

All subgroups' analyses with the forest plots are presented in online Supplemental Material (S6).

Subgroup 1: single feature condition vs. feature binding condition

The test for subgroup differences suggests that there is a statistically significant subgroup effect comparing the single feature condition and the feature binding condition in AD and preclinical patients (both $p < 0.001$). The feature binding condition showed larger effects than the single feature condition. Although heterogeneity was evident in both comparisons [possibly explained by the variation between the tasks (free recall, change detection, difficulty titration or no-titration, etc.)], it is worth noting that almost all studies had negative effects (AD and preclinical showing worse performance than controls). The comparison of the effects of single feature condition and feature binding condition in the MCI group did not show statistically significant subgroup differences ($p = 0.96$).

Subgroup 2: Change detection vs. free recall

The subgroup analyses comparing the effects between change detection and free recall tasks in AD patients showed no effect in both single feature condition and feature binding condition tasks (single feature condition $p = 0.52$; feature binding condition $p = 0.66$). As only one study

used the free recall task (Cecchini et al., 2020) to assess patients in the MCI stage group and none used the free recall task in the preclinical stage of AD, it proved unfeasible to conduct subgroup analyses for patients in these groups.

Subgroup 3: Familial vs. sporadic AD

We were not able to conduct meaningful subgroup analyses comparing the effects between sporadic and familial AD because of the small number of participants in the familial AD group (63 familial AD vs. 280 sporadic AD in single feature condition and 63 familial AD vs. 327 sporadic AD in the feature binding condition). In addition, only one study assessed familial AD at the preclinical/subjective cognitive decline stage (Koppara et al., 2015b) or MCI stage (Pietto et al., 2016).

Subgroup 4: Shape-color vs. color-color binding

Shape-colour and colour-colour binding tasks did not show statistically significant effect differences in the AD group ($p = 0.67$). Only limited data on color-color task compared with the shape-color task were available in the AD and preclinical groups, and no study used the color-color task for MCI patients.

Subgroup 5: Difficulty titration vs. no-titration

The titration method involves presenting fewer to-be-remembered items within visual arrays (i.e., smaller set sizes) for patients than for controls. This procedure was used so the ability to hold temporary conjunctions could be compared properly between the groups controlling for working memory load as informed by single feature performance (i.e., equated across groups).

In AD group, the comparison between the titration vs. no-titration methods showed statistically significant results in the single feature condition ($p = 0.01$). As expected, the studies that did not titrated the task difficulty showed larger effects. However, the effects were similar in the feature binding condition ($p = 0.21$), meaning that even in an easier task, AD patients showed a binding deficit. None of the studies with MCI patients titrated the task difficulty. For the preclinical group, only three studies used the titration method (Parra et al., 2010a; 2011b; 2015b) and two used no-titration (Koppara et al., 2015b; Norton et al., 2020), hampering the possibility of conducting any meaningful analysis.

Discussion

In this systematic review, we assessed evidence from 20 published studies and performed a meta-analysis on 19 studies on the performance of patients at different stages of the AD continuum (from preclinical to dementia stage) using the STM binding task. The reviewed studies were published between 2009 and 2021. In total, 864 patients at different stages of AD and 1,069 controls were assessed with the feature binding condition and 703 patients at different stages of AD and 841 controls with the single feature condition.

It is worth noting that some studies included more than one experiment, had more than one group (e.g., two groups of patients), did more than one session, or compared the groups using tasks with different set sizes. A total of 38 comparisons were performed with the feature binding condition (shape-color binding or object-color binding) and 32 with the single feature condition tasks.

Different testing paradigms were used, with variation in presentation time, recall strategy, stimuli type or quantity displayed on the screen. Essentially, three types of retrieval strategy

were used by the authors of the studies included in this review: 1) recognition during change detection; 2) free recall; 3) cued recall. For the change detection two different versions were used across studies: a computerized version using the E-prime program and an analogue flash-card version. Examples of the various tasks used in the different studies entering the analyses can be found in online Supplemental Material (S7).

The mean effect size of the single feature condition was small in the preclinical stage and increased significantly in the MCI and AD dementia groups. The feature binding condition, on the other hand, had a large mean effect size even when comparing controls and preclinical AD, and the effect size increased in comparison with AD at dementia stage. MCI patients showed a similar effect size when compared with preclinical AD in the feature binding condition. This could be related to the fact that most preclinical AD patients in the study had a genetic mutation that leads 100% to AD dementia (Lopera, 1997), but the MCI groups were more heterogeneous, as expected. Notwithstanding the higher risk to develop AD dementia (Mitchell & Shiri-Feshki, 2009), some MCI patients could convert to other dementia types, stay stable or return to normality (Galluzzi et al., 2013; Ganguli et al., 2011; Grande et al., 2016; Overton et al., 2019; Roberts & Knopman, 2013). This occurs because MCI is essentially a cognitive status, not a disease itself, and it is diagnosed using essentially cognitive and functional measures and self-reported cognitive complaints. Furthermore, none of the studies with people with MCI titrated the task difficulty between control and MCI group (i.e., the groups did the same task) and most used a task with 3 items per trial. This can have overloaded the patients' working memory, hampering their performance not only in the feature binding condition task (Parra et al., 2019). As the reviewed studies did not include biomarkers to ascertain AD pathology, the heterogeneity in the MCI group should be expected. In the next sections we will provide an in-depth examination of the evidence according to the three different stages of AD.

Alzheimer's disease dementia

The meta-analyses indicated a large mean effect comparing controls and AD dementia patients. The effect size was larger in the feature binding conditions (-2.40) compared with the single feature ones (-1.18), indicating that the AD groups showed much more difficulty to hold bound information (i.e., shape-color, color-color or object-color bindings) than to hold individual features, and that binding deficits could not be explained by a general working memory deficit (i.e., they are specific deficits). In addition, the studies with both sporadic and familial AD dementia showed similar results. That is important because it suggests that STM binding deficits are independent of the disease variant (Parra et al., 2011), and hence evidence drawn from familial AD could help interpret behaviors observed during STM binding tests in sporadic AD. However, it should be noted that future studies will be needed to ascertain the relationship between genotype and phenotypes in AD (Holmes, 2002).

Mild cognitive impairment stage

The single feature condition in the studies showed effect sizes from -0.37 (set size two) to -1.46 (set size three), that is, small to large effects. The MCI group showed a performance similar to that of controls in the shape-only condition when two items per screen were used, but a significant deficit when three items were used. It was argued (Kozlova et al., 2020; Parra et al., 2019) that the deficit to bind shape-color would be more apparent using a smaller set size, in which patients would show similar performance in the shape-only condition, but deficits in the shape-color binding task. The effect sizes on feature binding conditions were large in all studies, while the mean effect sizes of the single feature condition and feature binding condition were similar (-0.95 and -1.06, respectively). That is, the MCI groups showed significant difficulties to hold bound and single feature information temporarily in the memory,

suggesting that the memory load used in such studies did not allow separating the general underlying working memory deficits present in these samples from the specific binding impairments previously found. Strategies to address this methodological caveat have been discussed in Parra et al. (2019).

These results could be further explained by the small sample sizes in the studies, but also by the fact that most of the MCI patients in the studies did not have biomarker data and did not follow the new biological criteria for AD (Jack et al., 2018). It is likely that the MCI groups were too heterogeneous. The conclusions from the MCI analyses highlight the importance to work with AD biomarkers in new studies with STM binding tests and MCI groups. Without long-term follow up assessments, it is not possible to ascertain that MCI patients assessed in such studies were in the AD biological continuum, making it difficult to determine the predictive role of STM binding deficits in the prodromal stages of the disease.

Pre-clinical and subjective cognitive decline stages

The meta-analyses indicated a large mean effect size comparing controls and pre-clinical patients in the feature binding condition (-1.10), and a small mean effect size in the single feature condition (-0.33). The studies showed a pattern of specific binding deficits in the preclinical stages of AD, in which the patients show a deficit to hold temporarily conjunctions, but not single items. All studies with asymptomatic carriers of the presenilin mutation and patients with subjective cognitive decline used a set size with three items per screen.

If the STM binding test can assist the detection of AD in the preclinical stage, it would be expected that this task should be related to the biomarkers that define the AD pathology. Three studies examined the comparison of STM binding test data with biomarkers (Cecchini et al., 2021; Norton et al., 2020; Parra et al., 2017b). Parra et al. (2017b) presented a poster in which

controls with amyloid PET (n=39) were assessed. The sample was split in two groups: strong binders and weak binders using the performance on the change detection shape-color binding task. The authors showed that the weak binders group had more amyloid burden in the parietal-occipital-temporal regions and fusiform gyrus when compared with the strong binders (Parra et al., 2017b). Norton et al. (2020) studied a sample of controls, asymptomatic carriers of the presenilin-1 E280A mutation and familial MCI; the STM binding test significantly correlated with amyloid deposition in the brain ($r = -0.50$, $p = 0.03$), but not with tau deposition in the inferior temporal lobe ($r = -0.30$, $p = 0.21$) or the entorhinal cortex ($r = -0.26$, $p = 0.27$). Cecchini et al. (2021) showed that the STM binding was the only cognitive task that discriminated groups with and without amyloid deposition when compared with RAVLT and Short Cognitive Performance Test (SKT) (episodic memory tests). Therefore, the STM binding test seems to be related to amyloid deposition in the brain, but the relation with tau or the entorhinal activity still need to be better understood.

Heterogeneity analyses

The heterogeneity was high in the comparisons between controls and AD patients in both conditions (single feature and feature binding). Four studies showed a particularly large effect sizes in the feature binding condition (Della Sala et al., 2016; Fernández et al., 2018; Guazzo et al., 2020; Kozlova et al., 2020). If these studies were dropped from analyses, the heterogeneity would be significantly reduced (to $I^2 = 48\%$). The study by Guazzo et al. (2020) showed the largest effect size, and this could be related to differences in the paradigm: it was the only study that tested memory with cued recall. In addition, the large effect size of the other three discrepant studies could be related to the set size used, as the controls and AD patients were presented with a task with the same set size (only two items per screen). This was not the

case for the majority of studies with the change detection task in which the AD group was presented with a smaller set size than the controls (Parra et al., 2010a, 2010b; Parra et al., 2011b; Parra et al., 2015b), with the exception of Cecchini et al. (2020). However, in this last study, 16 trials were carried out, while the other studies with the change detection used 32 trials, which possibly increased the observed effect.

The strategy to titrate the working memory load across groups could also explain the high heterogeneity in the single feature condition tasks as well. Parra et al. (2011b), for instance, showed a positive effect, with controls showing worse performance than the AD patients (although these differences were non-significant). This occurred due to the use of a titration method in which controls were presented with a more challenging task than patients, thereby equating performance on the single feature conditions across groups. Studies that did not use titration, such as Cecchini et al. (2020), showed negative and large effect sizes. In addition, in a task with the same and easiest set size (two items per screen), the control group showed performance close to ceiling, with low standard deviations, which increased the effect sizes compared to the other groups. Sample characteristics also could be related to the heterogeneity found. AD patients were not taking cholinesterase inhibitors in the study of Fernández et al. (2018), which could have improved their cognitive performance (Rockwood, 2004; Wilkinson et al., 2004). It seems, also, that this group was heterogeneous, and probably some patients were in more advanced stages of the disease. This was observed in the cognitive measures, especially in the Mini-Mental State Exam (MMSE) (mean = 22.6, SD = 4.1) and Addenbrooke's Cognitive Examination - Revised (ACE-R) (mean = 65.3, SD = 16.9).

The high variability in the AD groups comparisons could be expected. Besides differences in age (from around 45 to 76 years old in mean age), education (from around 6.4 to 14.7 years of formal schooling), and the differences in methods cited above, it is impossible to ascertain that all patients were in the same stage of the disease. In addition, the disease itself is heterogeneous,

with patients showing different clinical and cognitive presentations (Binetti et al., 1993; Lam et al., 2013; Martorelli et al., 2019).

Caveats and suggestions for future studies

It is important to highlight some limitations of the memory binding studies included in this systematic review and to discuss which further evidence is necessary to accrue before STM binding tests could be reliably used in broader clinical settings.

Some studies with asymptomatic carriers of the presenilin mutation and with subjective cognitive decline patients (Koppara et al., 2015b; Parra et al., 2015b; Parra et al., 2011b; Parra et al., 2010a) argued that the STM binding can detect very early signs of the disease, therefore the task should be a good predictor of the dementia evolution. However, longitudinal studies should be carried out to verify whether the STM binding test is a good predictor of dementia in patients with suspected AD. Only one such study could be gleaned from the literature (Martínez-Flores et al., 2021). However, in this study the authors did not test the STM binding as a predictor of MCI to AD dementia conversion but used machine learning techniques to extract classifiers distinguishing between controls and MCI at baseline and tested their accuracy at the follow up assessment (i.e., by comparing MCI vs controls).

Another important issue is the fact that all studies reviewed used small sample sizes (median of 23 participants per group). Therefore, studies with larger samples would be recommended. In addition, most of the papers shared common authors; for instance, Parra authored 17 out of 19 articles, while Della Sala authored 13. The binding tasks should be used by different research groups, with no affiliation or relation with the original authors of the binding paradigm to assess independent replicability. Also, to facilitate the use of the STM binding test by other research groups, the tasks should be available on a free and open-access website. To address this issue, all STM binding test versions used by the authors of this review are now available from

<https://www.strath.ac.uk/research/subjects/psychology/cognition/appliedcognitionlab/visualshort-termmemorybindingtestvstmbt/>.

Studies testing the psychometric properties of the tasks should also be conducted, performing test-retest and inter-rater reliability, verify the validity and correlate the STM binding with other neuropsychological tests to better understand precisely what cognitive functions are being assessed.

Some aspects of the tasks could be improved. Around 8% of men and 0.4% of women show color-blindness (Birch, 2012), especially to distinguish red and green. To avoid difficulties driven from color blindness, a color-blind palette could be used to tailor a new version of the change detection task. In addition, it is important to note that in the change detection task participants could pay attention to $x-1$ stimuli, the x representing the number of items on the screen. For instance, as the colors swap between the shapes in the feature binding condition with two stimuli, to detect a change from previous screen participants could focus on only one stimulus. As this could not be clear in the instructions for the participants, some of them could be aware of that, and some not, causing an extra source of variation. That could be one of the causes for the large standard deviations in some studies, and maybe an instruction making this explicit could avoid this problem.

Studies varied also in terms of task titration. Many studies used a different set size to compare control and patient groups (Parra et al., 2015b; Parra et al., 2009b; Della Sala et al., 2012; Parra et al., 2011b; Valdés Hernández et al., 2020; Parra et al., 2010a, 2010b), the former doing a more difficult task than the latter. However, some variation within each group would be expected, since not every person has the same working memory capacity. It would probably be better to titrate the task individually, from one to four or five stimuli, as four is the average of stimuli limit storage in the visual STM (Cowan, 2001, 2010). In addition, some studies

suggested that the best set size for identifying MCI or AD using the change detection task is two stimuli per trial (Kozlova et al., 2020; Parra et al., 2019), while to detect preclinical AD three items per trial proved more sensitive (Koppara et al., 2015b; Parra et al., 2010a; Parra et al., 2011b). Therefore, in a clinical setting, both set sizes should be used, but this would significantly increase the time spent to do the task.

The same question about the number of stimuli applies to the free recall tasks. In addition, the relation between the single feature condition and feature binding condition tasks scores should be better investigated in the free recall paradigm. For instance, if a patient can recall three items out of four in a trial, she/he will achieve 75% in performance in the single feature condition, but only 50% in feature binding condition. This occurs because in the feature binding condition only the correct binding between objects and colors is scored, but in the single feature condition the participant score in each color or object recalled individually. Therefore, a binding cost would be expected, that is, to have worse performance in the feature binding condition task when compared with the single feature condition, and this could be amplified in the clinical groups, as presented by Cecchini et al. (2020). To address this question, the number of stimuli could vary from one or two stimuli to four/five, as explained before.

Another important topic of discussion is the specificity of the binding deficits. If binding deficits are specific to AD, as suggested by many studies here reviewed, the best measure to detect AD should be the binding cost, which represents the difference (or proportion) between single feature condition and the feature binding condition tasks. However, the best variable to detect AD is the feature binding condition alone, and even when the performance in the single feature condition was the same as controls, the variable used to discriminate the groups was still the feature binding condition alone, not the binding cost (Cecchini et al., 2020; Della Sala et al., 2012; Parra et al., 2010a, 2010b; Parra et al., 2011b). One possible explanation could be that this occurred due to a general working memory deficit in the AD groups in addition to a

binding deficit, decreasing the performance in both conditions (single feature and feature binding), interfering with the binding cost variable. However, if this is the case, it would be still necessary to create norms for non-pathological performance for the feature binding condition. Yassuda et al. (2020), for instance, showed that the performance between healthy participants divided in age groups was not different in the feature binding condition of the free recall task with two items per screen, but had differences using three items. Probably with more items the age impact in the performance would become more prominent due to general working memory deficits related to aging (Brockmole & Logie, 2013; for reviews see Logie & Morris, 2015).

To claim that a test is specific to one type of dementia (e.g., AD), it is not enough to compare it only with controls. A few studies compared AD with other types of dementia, and most of them used the free recall task (Cecchini et al., 2017a; Della Sala et al., 2012; Guazzo et al., 2020). Only one study used the change detection task (Kozlova et al., 2020). However, the differential diagnosis between AD and Parkinson's disease used in this latter study is not the most challenging in the clinical setting and these pathologies typically do not present with similar cognitive deficits (Bronnick et al., 2007). As far as we are aware, no article was published to this date comparing, for instance, AD and other dementia types (besides Parkinson's) using the change detection task, but this information was found in posters, conferences abstracts and in a dissertation. The posters from Yassuda et al. (2018), Cecchini et al. (2017b) and the Cecchini's dissertation (Cecchini & Yassuda, 2017) informed that the change detection and free recall modalities of the STM binding test showed a different pattern when controls, AD, and behavioral variant frontotemporal dementia (bvFTD) groups were compared. Only AD patients showed impairments in the free recall task [(Controls = bvFTD) > AD], while AD and bvFTD showed impairment in the change detection task [(Controls > (AD = bvFTD))]. In another poster, Cecchini et al. (2017c) showed that the free recall version

of the STM binding test could differentiate AD from amnesic bvFTD. That is important because AD and amnesic bvFTD showed the same pattern of episodic memory deficits on the RAVLT test, thus the free recall binding task could be used to differentiate AD from bvFTD even when the latter show episodic memory deficits, which is not unusual (Hornberger et al., 2010; van den Berg et al., 2020). Therefore, free recall and change detection tasks could assess binding through different brain networks.

With that in mind, it is necessary to better understand each paradigm of the STM binding. For instance, the color-color binding paradigms does not seem to assess the same ability as the shape-color binding (Wheeler & Treisman, 2002). Color-color binding probably is a form of relational binding because the combination does not define an object, whereas shape-color binding is conjunctive (Parra et al., 2011a). However, the color-color showed similar results as shape-color binding, with similar effect sizes in the meta-analyses. In the paradigm used by (Guazzo et al., 2020), it was also not clear how the cross-modal binding (i.e., binding visual and auditory stimuli) would be different from the relational binding. Without better understanding of each task, it is not possible to know, for instance, if AD and bvFTD patients are performing similarly but for different reasons. As AD patients have hippocampal atrophy and difficulties in binding conjunctive information, they can show poor performance on relational (hippocampal) and conjunctive (not hippocampal) binding. The same can occur with asymptomatic carriers of the presenilin mutation, which showed impairments in the color-color (Parra et al., 2011b) and shape-color binding (Koppara et al., 2015b; Parra et al., 2010a). The bvFTD patients, on the other hand, could show difficulties in the change detection task due to a disruption in networks related to the frontal lobe (e.g., salience network) (Filippi et al., 2013; Seeley et al., 2008; Yu et al., 2016), that probably is associated with this task (Parra et al., 2017a; Pietto et al., 2016). These discrepancies suggest that the free recall and change detection tasks may rely on different brain areas, and it is not clear if they assess the same construct or

how they assess conjunctive binding. As far as we are aware, all imaging studies to this date have used the change detection task.

One of the studies included in this review showed discrepant results (Norton et al., 2020). In this study with the STM binding test, the authors found a significant effect of condition (shape-only x shape-color binding) and a significant effect of group (controls x asymptomatic carriers x MCI-FAD), but no significant interaction between them ($p = 0.29$). That is, asymptomatic carriers and MCI-FAD patients had similar deficits on both conditions when compared with controls. In addition, the shape-only condition showed higher correlation with amyloid deposition in the brain and tau deposition on entorhinal and parietal inferior areas when compared to the feature binding condition. Lastly, the accuracy to detect amyloid and tau positivity was higher for the shape-only condition than for the shape-color one. These behavioral results contradict the initial hypotheses and results from previous findings (Parra et al., 2010a; Parra et al., 2015b). However, from a biomarker point of view, Norton et al. (2020) reported that performance on the shape-color condition correlated with Pittsburgh compound-B (PIB) (at a similar level as the shape condition) and that association was only significant when restricting the analysis to the asymptomatic carriers. They argue that a reason why such a correlation decreased as patients progressed to the tau stages is that performance on the task was reaching floor (in addition to a small sample size). This evidence calls for more studies to document not only “what” should be assessed along the continuum of AD but also “when”.

Conclusions

The articles reviewed showed evidence that patients in the AD continuum present with deficits in holding feature bindings in STM. The difficulties can be found even at preclinical stages of the disease. However, the studies used a cross-sectional design with small sample sizes, in

addition to a moderate to high heterogeneity, which could have contributed to some inconsistencies between the results. Future studies should investigate the STM binding test as predictor of dementia in asymptomatic participants in longitudinal designs and investigate how the task relates to AD biomarkers (amyloid and tau) in larger samples.

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Tables

Table 1. Summary of findings for the main comparisons. The clinical use of short-term memory binding

Patient or population: patients with Alzheimer's disease (AD)

Setting: clinical setting in primary or secondary healthcare practices

Comparison: controls and AD patients assessed using the short-term memory binding test

Outcomes		Effect size (95% CI)	Number of participants (studies)	Quality of evidence (GRADE)
Preclinical AD and subjective cognitive decline	Single feature condition	-0.33 (-0.59; -0.07)	241 (5)	⊕⊕⊕○ Moderate
	Feature binding condition	-1.10 (-1.48; -0.73)	241 (5)	
Mild cognitive impairment	Single feature condition	-1.08 (-1.35; -0.81)	628 (8)	⊕⊕⊕○ Moderate
	Feature binding condition	-1.07 (-1.32; -0.82)	928 (9)	
AD in dementia stage	Single feature condition	-1.18 (-1.61; -0.76)	675 (11)	⊕⊕⊕○ Moderate
	Feature binding condition	-2.41 (-2.82; -1.99)	764 (13)	

CI: confidence interval; Effect Size was measured using the Hedges' g formula.

GRADE Working Group grades of evidence

High quality: Further research is very unlikely to change our confidence in the estimate of effect.

Moderate quality: Further research is likely to have an important impact on our confidence in the estimate of effect.

Low quality: Further research is very likely to have an important impact on our confidence in the estimate of effect.

Very low quality: We are very uncertain about the estimate.

Table 2. Characteristics of included studies.

	STM binding modality	Sample	N	Age Mean(SD)	Education Mean(SD)	Items per trial	Trials	Study / encoding time	Single feature condition	M	
	Change detection	HC	30	40.9(9.3)	9.5(3.2)	3	32	2000	Shape-only	Perce	
		AC	30	35.6(6.6)	9.3(4.4)	3					and A
	Change detection (color-color)	HC_2	29	39.55(8.82)	9.21(2.90)	3	32	2000	Colors-only	Perce	
		AC	25	37.24(5.21)	9.12(3.68)	3					corre
al.,	Change Detection	HC	23	68.00(8.31)	14.39(3.07)	3	32	2000	Shape-only	Hit m	
		SCD	19	66.79(7.58)	16.53(3.03)	3					alarm
	Change detection	HC	21	39.30(83)	10.30(27)	3	32	2000	Shape-only	Perce	
		AC	18	35.10(5.5)	10.20(3.9)	3					corre
,	Change detection	HC	27	37.0(6.5)	9.9(4.1)	3	32	2000	Shape-only	A'	
		AC	19	37.5(6.5)	11.4(3.8)	3					
al.,	Change detection	HC	23	68.00(8.31)	14.39(3.07)	3	32	2000	Shape-only	Hit m	
		MCI	23	72.82(4.37)	13.27(2.96)	3					alarm
2016	Change detection	HC	14	67.21(10.14)	16.50(1.99)	3	100	500	Shape-only	Hit m	
		HC_2	10	44.30(5.60)	11.30(13.90)	3					alarm
		MCI	13	73.08(9.01)	14.08(4.44)	3					
	MCI-FAD	10	44.40(3.20)	7.30(4.10)	3						
Change detection	HC	10	44.30(5.60)	11.30(13.90)	3	100	500	Shape-only	Hit m		
	MCI-FAD	10	44.40(3.20)	7.30(4.10)	3					alarm	
2019		HC	25	74.73(4.74)	10.84(5.02)	2	32	2000	Shape-only		

et al.,	Change detection	HC_2	29	72.34(3.76)	11.00(5.11)	3	32	2000	Shape-only	Perce
		MCI	27	75.07(5.30)	10.86(5.80)	2				
		MCI_2	23	75.43(5.77)	9.43(2.90)	3				
et al.,	Change detection	HC	25	76.24(5.37)	15.08(3.58)	3	32	2000	Shape-only	Perce
		MCI	21	74.00(5.49)	13.57(3.88)	2				
al.,	Change Detection	HC	23	67.83(6.06)	12.83(4.06)	2	32	2000	Shapes and colors	Perce
		MCI	24	70.33(6.89)	9.54(5.82)	2				
al.,	Free Recall	HC	21	67.83(6.06)	12.83(4.06)	3	6	9000	Objects and Colors	Perce
		MCI	24	70.33(6.89)	9.54(5.82)	3				
,	Change detection	HC	27	37.0(6.5)	9.9(4.1)	3	32	2000	Shape-only	A'
		MCI-FAD	6	44.8(1.4)	9.3(3.4)	3				
ores	Change Detection	HC	109	66.53(7.15)	12.55(4.07)	2	32	2000	Shape-only	Perce
		HC	109	66.53(7.15)	12.55(4.07)	3				
		HC_2	38	66.10(7.12)	13.81(4.05)	2				
		HC_2	38	66.10(7.12)	13.81(4.05)	3				
		MCI	45	68.33(7.09)	11.28(3.67)	2				
		MCI	45	68.33(7.09)	11.28(3.67)	3				
		MCI_2	18	67.83(6.33)	12.38(3.58)	2				
al.,	Change Detection	HC	18	72.39(5.84)	12.33(4.97)	2	16	2000	None (only bound condition)	Numb
		MCI	30	73.67(4.69)	9.57(4.76)	2				
	Free recall	HC	23	69.78(6.47)	7.08(2.81)	3	6	9000	Objects and Colors	Perce
		HC_2	20	69.35(6.02)	7.25(2.97)	4				
		AD	23	73.26(6.09)	6.39(3.34)	2				
		AD	21	73.33(6.71)	6.81(3.66)	2				
	Change detection	HC	30	40.9(9.3)	9.5(3.2)	3	32	2000	Shape-only	Perce
FAD	22	45.2(4.8)	8.5(4.2)	2						

	Change detection	HC	14	70.71(4.30)	15.57(3.32)	3	32	2000	Shape-only	A' and
		AD	14	76.29(5.78)	12.71(3.77)	2				
	Change detection (color-color)	HC	14	70.71(4.30)	15.57(3.32)	3	32	2000	Colors-only	Perce
		HC_2	29	39.55(8.82)	9.21(2.90)	3				
		AD	14	76.29(5.78)	12.71(3.77)	2				
t al.,	Free recall	HC	20	69.35(6.21)	7.25(2.97)	4	6	12000	Objects and Colors	Perce
		AD	15	72.93(5.79)	7.13(3.74)	2				
	Change detection	HC	21	39.30(83)	10.30(27)	3	32	2000	Shape-only	Perce
		FAD	19	47.50(6.4)	7.30(3.7)	2				
t al.,	Change detection	HC	33	73.87(8.51)	13.30(3.32)	2	32	2000	Shape-only	Perce
		AD	33	75.24(7.72)	13.27(3.17)	2				
al.,	Free Recall	HC	32	67.84(6.82)	12.25(3.69)	3	6	9000	Objects and Colors	Perce
		AD	35	71.40(7.96)	10.09(5.41)	3				
t al.,	Change detection (color-color)	HC	13	68(4.2)	18.2*	2	32	2000	Colors-only	Perce
		AD	13	67(2.6)	13.4*	2				
al.,	Change detection	HC	31	69.10(8.42)	14.42(2.87)	2	32	2000	Shape-only	Perce
		AD	24	72.58(8.16)	14.67(2.87)	2				
l.,	Cued recall	HC	24	74.54(4.12)	10.20(3.47)	2	12	3000	Shape-only	Perce
		HC_2	24	74.75(3.92)	9.56(2.90)	3				
	Change Detection	AD	24	76.29(5.18)	9.08(1.18)	2	32	2000	Shapes and colors	Perce
		HC	23	67.83(6.06)	12.83(4.06)	2				
al.,	Free Recall	AD	37	71.14(7.58)	10.05(5.23)	2	6	9000	Objects and colors	Perce
		HC	21	67.83(6.06)	12.83(4.06)	3				
al.,	Change Detection	AD	37	71.14(7.58)	10.05(5.23)	2	16	2000	None (only bound condition)	Numb
		HC	18	72.39(5.84)	12.33(4.97)	2				
		AD	23	73.78(5.99)	8.57(4.15)	2				

Change Detection	HC	18	69.00(3.60)	17.1*	2	32	2000	Colors-only	Perce corre
	AD	18	68.00(2.20)	13.6*	2				

Legend. STM = short-term memory; HC = healthy controls; AD = Alzheimer's disease; MCI = mild cognitive impairment; MCI = mild cognitive impairment; AC = asymptomatic carrier of the presenilin-1 E280A mutation; FAD = familial AD (mutation of the presenilin-1 E280A in the dementia stage); MCI-FAD = MCI in familial AD; SCD = subjective cognitive decline. The samples presented in this table are only related to the AD continuum, from normal controls to AD dementia; other types of dementia on these studies were omitted. * Standard deviation for Education was not presented.

SUPPLEMENTAL MATERIAL

Short-Term Memory Conjunctive Binding in Alzheimer's Disease: A Systematic Review and Meta-Analysis

S1. Details of the terms used to search the electronic databases.

The terms combinations used to search the databases were:

“short-term memory binding” AND “Alzheimer's disease”

“short-term memory binding” AND “mild cognitive impairment”

“short-term memory binding” AND “subjective cognitive decline”

“memory binding” AND “Alzheimer's disease”

“memory binding” AND “mild cognitive impairment”

“memory binding” AND “subjective cognitive decline”

conjunctive memory AND “Alzheimer's disease”

conjunctive memory AND “mild cognitive impairment”

conjunctive memory AND “subjective cognitive decline”

“working memory binding” AND “Alzheimer's disease”

“working memory binding” AND “mild cognitive impairment”

“working memory binding” AND “subjective cognitive decline”

S2. List of studies included in meta-analyses

Cecchini, M.A., Yassuda, M.S., Bahia, V.S., de Souza, L.C., Guimarães, H.C., Caramelli, P., et al. (2017a). Recalling feature bindings differentiates Alzheimer's disease from frontotemporal dementia. *Journal of Neurology*, 264(10), 2162–2169.

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S3.2 Relational, associative or other types of binding

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S3.5 Poster, meeting abstract, single-case or no access to data

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S4. Quality of evidence and risk of bias assessment of the included studies following GRADE guidelines

Study design	Risk of bias	Inconsistency	Indirectness	Imprecision	Publication bias	Observations	Effect size (95%CI)	Quality
Parra, Abrahams, Fabi, Logie, Luzzi & Della Sala (2009). Two groups of controls and AD patients were compared.								
-1. Observational, case-control.	-1. Impossibility of blinding procedures or lack of allocation concealment	No serious inconsistencies.	No indirectness observed	No serious imprecision.	No publication bias identified.	Increased from low to moderate quality based on large effect sizes.	Single feature condition HC x AD (1) = -1.61 (-2.28; -0.94) HC x AD (2) = -0.51 (-1.13; 0.12) Feature binding condition HC x AD (1) = -2.89 (-3.74; -2.04) HC x AD (2) = -2.22 (-3.02; -1.43)	Moderate
Parra, Abrahams, Logie, & Della Sala (2010). Controls and AD patients were compared.								
-1. Observational, case-control.	-1. Impossibility of blinding procedures or lack of allocation concealment	No serious inconsistencies.	No indirectness observed	No serious imprecision.	No publication bias identified.	Increased from low to moderate quality based on large effect sizes.	Single feature condition HC x AD (ss2) = -1.17 (-1.98; -0.35) HC x AD (ss3) = -2.66 (-3.72; -1.61) Feature binding condition HC x AD (ss2) = -2.47 (-3.48; -1.45) HC x AD (ss3) = -2.50 (-3.52; -1.48)	Moderate
Parra, Abrahams, Logie, Mendez, Lopera & Della Sala (2010). Controls, asymptomatic carriers and FAD patients were compared.								
-1. Observational, case-control.	-1. Impossibility of blinding procedures or lack of allocation concealment	No serious inconsistencies.	No indirectness observed	No serious imprecision.	No publication bias identified.	Increased from low to moderate quality based on large effect sizes.	Single feature condition HC x AC = -0.24 (-0.75; 0.27) HC x FAD = -0.13 (-0.68; 0.43) Feature binding condition HC x AC = -1.52 (-2.10; -0.94) HC x FAD = -1.65 (-2.29; -1.01)	Moderate
Parra, Della Sala, Abrahams, Logie, Méndez & Lopera (2011). Controls, sporadic and familial patients were compared.								
-1. Observational, case-control.	-1. Impossibility of blinding procedures or lack of allocation concealment	No serious inconsistencies.	No indirectness observed	No serious imprecision.	No publication bias identified.	Increased from low to moderate quality based on large effect sizes.	Single feature condition HC x SAD = 0.41 (-0.34; 1.16) HC x FAD = 0.04 (-0.58; 0.65) Feature binding condition HC x SAD = -1.35 (-2.19; -0.52) HC x FAD = -1.48 (-2.19; -0.78)	Moderate

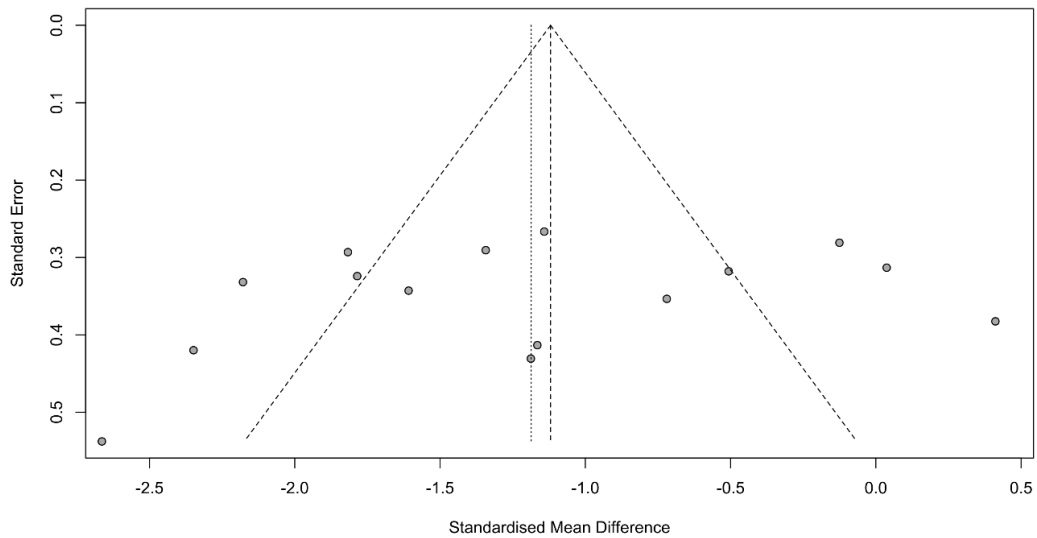
Della Sala, Parra, Fabi, Luzzi & Abrahams (2012). Controls and AD were compared.								
-1. Observational, case-control.	-1. Impossibility of blinding procedures or lack of allocation concealment	No serious inconsistencies.	No indirectness observed	No serious imprecision.	No publication bias identified.	Increased from low to moderate quality based on large effect sizes.	Single feature condition HC x AD = -0.72 (-1.41; -0.03) Feature binding condition HC x AD = -2.89 (-3.87; -1.90)	Moderate
Koppara, Frommann, Polcher, Parra, Maier, Jessen, Klockgether & Wagner (2015). Controls, Subjective cognitive decline and MCI were compared.								
-1. Observational, case-control.	-1. Impossibility of blinding procedures or lack of allocation concealment	No serious inconsistencies.	No indirectness observed	No serious imprecision.	No publication bias identified.	Increased from low to moderate quality based on large effect sizes.	Single feature condition HC x SCD = -0.21 (-0.82; 0.40) HC x MCI = -1.16 (-1.79; -0.54) Feature binding condition HC x SCD = -0.85 (-1.49; -0.21) HC x MCI = -1.33 (-1.97; -0.68)	Moderate
Parra, Saarimaki, Bastin, Londono, Pettit, Lopera, Della Sala & Abrahams (2015). Controls, Asymptomatic carriers and FAD patients were compared.								
-1. Observational, case-control.	-1. Impossibility of blinding procedures or lack of allocation concealment	No serious inconsistencies.	No indirectness observed	No serious imprecision.	No publication bias identified.	Increased from low to moderate quality based on large effect sizes.	Single feature condition HC x AC = -0.51 (-1.15; 0.13) HC x FAD = -2.35 (-3.17; -1.52) Feature binding condition HC x AC = -1.06 (-1.73; -0.38) HC x FAD = -2.09 (-2.88; -1.31)	Moderate
Della Sala, S., Kozlova, I., Stamate, A., & Parra, M. A. (2016). Controls and AD patients were compared.								
-1. Observational, case-control.	-1. Impossibility of blinding procedures or lack of allocation concealment	No serious inconsistencies.	No indirectness observed	No serious imprecision.	No publication bias identified.	Increased from low to moderate quality based on large effect sizes.	Single feature condition HC x AD = -1.14 (-1.66; -0.62) Feature binding condition HC x AD = -3.89 (-4.73; -3.05)	Moderate
Pietto, Parra, Trujillo, Flores, García, Bustin, Richly, Manes, Lopera, Ibáñez & Baez (2016). Controls, MCI and MCI-FAD were compared.								
-1. Observational, case-control.	-1. Impossibility of blinding procedures or lack of allocation concealment	No serious inconsistencies.	No indirectness observed	-1. 95%CI upper and lower bounds with different meaning.	No publication bias identified.		Single feature condition HC x MCI = -0.86 (-1.66; -0.07) HC x MCI-FAD = -0.98 (-1.91; -0.04) Feature binding condition HC x MCI = -0.90 (-1.69; -0.10) HC x MCI-FAD = -0.89 (-1.81; 0.04)	Low

Cecchini, Yassuda, Bahia, et al. (2017). Controls and AD patients were compared.								
-1. Observational, case-control.	-1. Impossibility of blinding procedures or lack of allocation concealment	No serious inconsistencies.	No indirectness observed	No serious imprecision.	No publication bias identified.	Increased from low to moderate quality based on large effect sizes.	Single feature condition HC x AD = -1.82 (-2.39; -1.24) Feature binding condition HC x AD = -1.49 (-2.03; -0.94)	Moderate
Fernandez, Orozco, Agamennoni, Schumacher, Sanudo, Biondi & Parra (2018). Controls and AD patients were compared.								
-1. Observational, case-control.	-1. Impossibility of blinding procedures or lack of allocation concealment	No serious inconsistencies.	No indirectness observed	No serious imprecision.	No publication bias identified.	Increased from low to moderate quality based on large effect sizes.	Single feature condition HC x AD = -1.19 (-2.03; -0.34) Feature binding condition HC x AD = -4.01 (-5.42; -2.60)	Moderate
Parra, Calia, García, Olazarán-Rodríguez, Hernandez-Tamames, Alvarez-Linera, Della Sala, Fernandez & Guinea (2019). Two groups of controls and MCI patients were compared.								
-1. Observational, case-control.	-1. Impossibility of blinding procedures or lack of allocation concealment	No serious inconsistencies.	No indirectness observed	No serious imprecision.	No publication bias identified.	Increased from low to moderate quality based on large effect sizes.	Single feature condition HC x MCI (ss2) = -0.37 (-0.92; 0.18) HC x MCI (ss3) = -1.46 (-2.08; -0.84) Feature binding condition HC x MCI (ss2) = -1.25 (-1.85; -0.65) HC x MCI (ss3) = -1.21 (-1.80; -0.61)	Moderate
Cecchini et al. (2020). Controls, MCI and AD patients were compared.								
-1. Observational, case-control.	-1. Impossibility of blinding procedures or lack of allocation concealment	No serious inconsistencies.	No indirectness observed	No serious imprecision.	No publication bias identified.	Increased from low to moderate quality based on large effect sizes.	Single feature condition HC x MCI (CD) = -1.19 (-1.81; -0.57) HC x MCI (FR) = -1.95 (-2.64; -1.25) HC x AD (CD) = -1.34 (-1.91; -0.77) HC x AD (FR) = -2.18 (-2.83; -1.53) Feature binding condition HC x MCI (CD) = -1.58 (-2.24; -0.92) HC x MCI (FR) = -2.52 (-3.32; -1.72) HC x AD (CD) = -1.83 (-2.45; -1.22) HC x AD (FR) = -2.48 (-3.17; -1.80)	Moderate
Guazzo, Allen, Baddeley & Della Sala (2020). Controls and AD patients were compared.								
-1. Observational, case-control.	-1. Impossibility of blinding procedures or lack of allocation concealment	No serious inconsistencies.	No indirectness observed	No serious imprecision.	No publication bias identified.	Increased from low to moderate quality based on large effect sizes.	Feature binding condition HC x AD = -4.17 (-5.21; -3.14)	Moderate

Kozlova, Parra, Titova, Gantman, & Sala (2020). Controls, amnesic MCI and AD patients were compared.								
-1. Observational, case-control.	-1. Impossibility of blinding procedures or lack of allocation concealment	No serious inconsistencies.	No indirectness observed	No serious imprecision.	No publication bias identified.	Increased from low to moderate quality based on large effect sizes.	Single feature condition HC x aMCI = -1.26 (-1.94; -0.59) HC x AD = -1.79 (-2.42; -1.15) Feature binding condition HC x aMCI = -2.36 (-3.16; -1.56) HC x AD = -3.37 (-4.22; -2.53)	Moderate
Norton et al. (2020). Controls, asymptomatic carriers and MCI patients were compared.								
-1. Observational, case-control.	-1. Impossibility of blinding procedures or lack of allocation concealment	No serious inconsistencies.	No indirectness observed	-1. 95%CI upper and lower bounds with different meaning.	No publication bias identified.		Single feature condition HC x AC = -0.49 (-1.08; 0.11) HC x MCI = -1.97 (-2.99; -0.95) Feature binding condition HC x AC = -0.56 (-1.16; 0.04) HC x MCI = -0.75 (-1.66; 0.16)	Low
Valdéz-Hernández, Clark, Wang, Guazzo, Calia, Pattan, Starr, Della Sala & Parra (2020). Controls and MCI patients were compared.								
-1. Observational, case-control.	-1. Impossibility of blinding procedures or lack of allocation concealment	No serious inconsistencies.	No indirectness observed	-1. 95%CI upper and lower bounds with different meaning.	No publication bias identified.		Single feature condition HC x MCI = -0.66 (-1.25; -0.06) Feature binding condition HC x MCI (ss2) = -0.82 (-1.43; -0.22) HC x MCI (ss3) = -0.79 (-1.39; -0.18)	Low
Cecchini et al. (2021). Controls, MCI and AD patients were compared.								
-1. Observational, case-control.	-1. Impossibility of blinding procedures or lack of allocation concealment	No serious inconsistencies.	No indirectness observed	No serious imprecision.	No publication bias identified.	Increased from low to moderate quality based on large effect sizes.	Feature binding condition HC x MCI = -0.63 (-1.23; -0.03) HC x AD = -1.34 (-2.03; -0.65)	Moderate
Martínez-Florez et al. (2021). Controls and MCI patients were compared.								
-1. Observational, case-control.	-1. Impossibility of blinding procedures or lack of allocation concealment	No serious inconsistencies.	No indirectness observed	-1. 95%CI upper and lower bounds with different meaning.	No publication bias identified.		Single feature condition HC x MCI (t1 ss3) = -0.67 (-1.03; -0.32) HC x MCI (t2 ss3) = -1.07 (-1.67; -0.48) Feature binding condition HC x MCI (t1 ss2) = -0.87 (-1.23; -0.51) HC x MCI (t1 ss3) = -0.64 (-1.00; -0.29) HC x MCI (t2 ss2) = -0.64 (-1.22; -0.07) HC x MCI (t2 ss3) = -0.61 (-1.18; -0.03)	Low

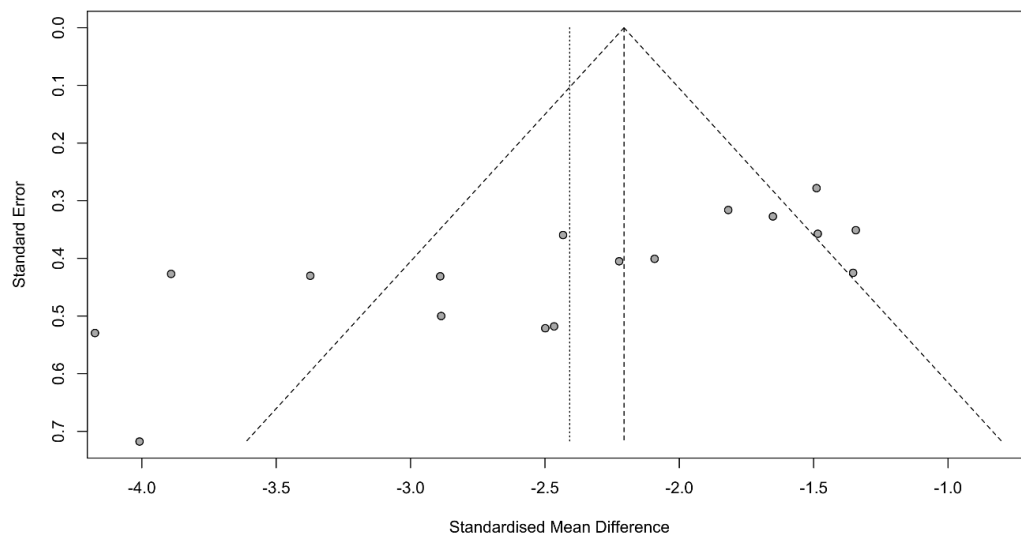
S5. Funnel Plots

S5.1 Performance of people with AD / Familial AD versus Controls, single feature condition.



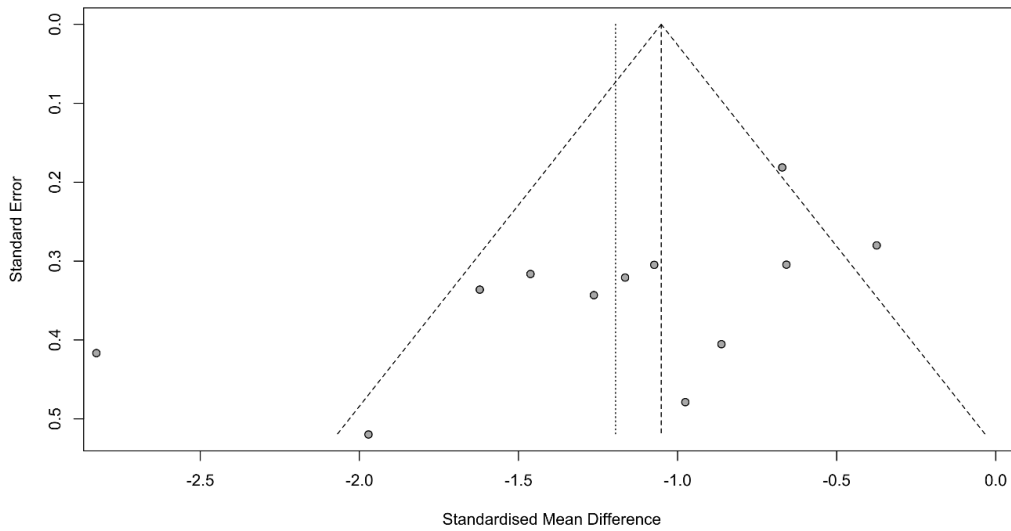
Note. Funnel plot of studies assessing the difference of the single feature condition of the short-term memory binding test between patients with sporadic/familial Alzheimer's disease and controls.

S5.2 Performance of people with AD / Familial AD versus Controls, feature binding condition.



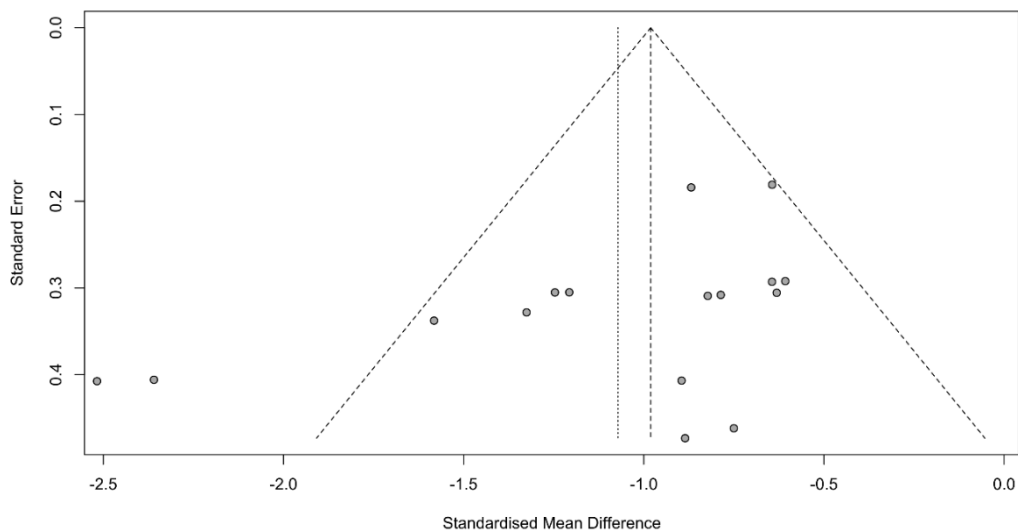
Note. Funnel plot of studies assessing the difference of the feature binding condition of the short-term memory binding test between patients with sporadic/familial Alzheimer's disease and controls.

S5.3 Performance of people with MCI / MCI-FAD versus controls, single feature condition.



Note. Funnel plot of studies assessing the difference of the single feature condition of the short-term memory binding test between people with MCI and MCI-FAD and controls.

S5.4 Performance of people with MCI / MCI-FAD versus controls, feature binding condition.

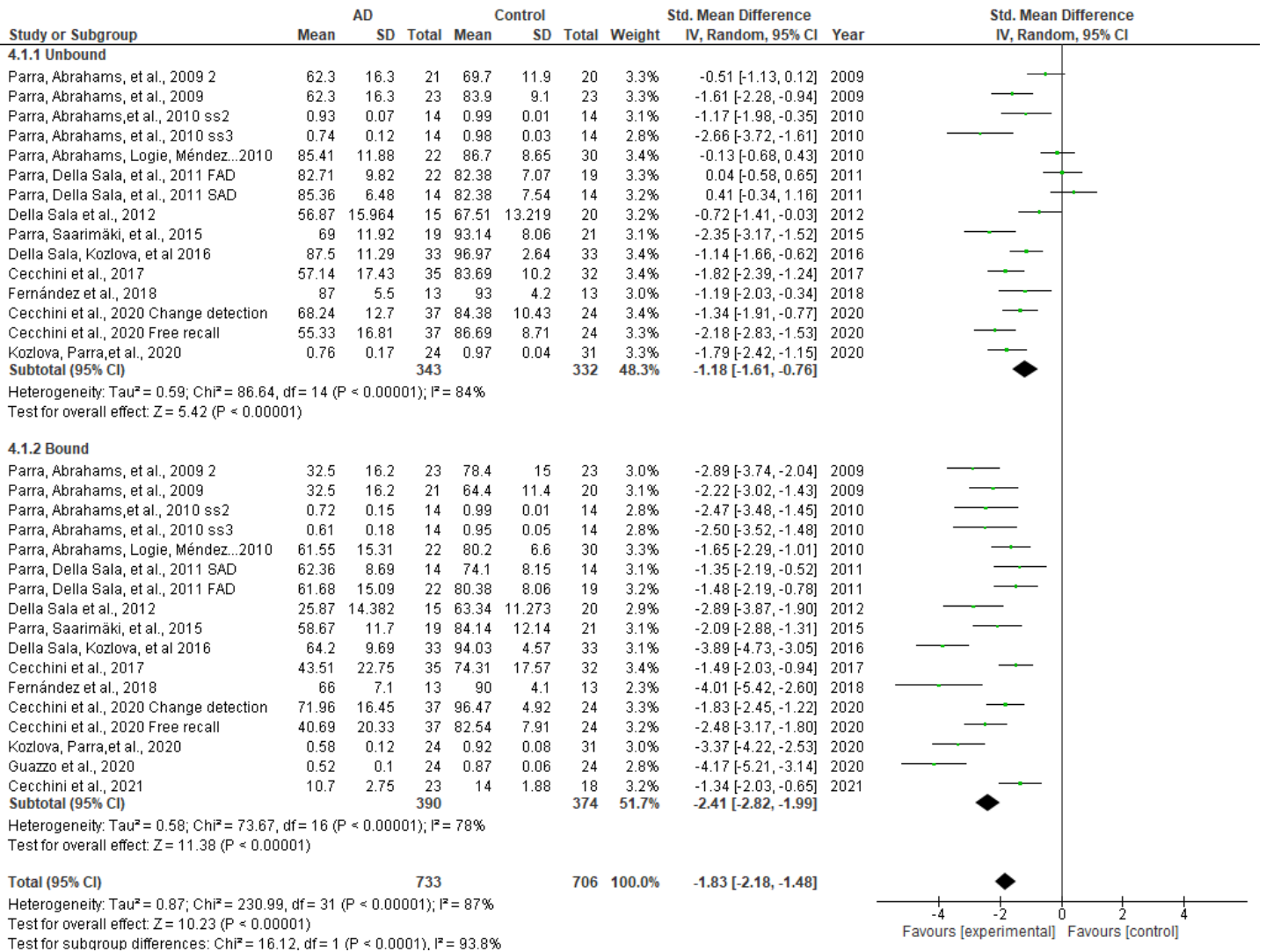


Note. Funnel plot of studies assessing the difference of the feature binding condition of the short-term memory binding test between patients with MCI and MCI-FAD and controls.

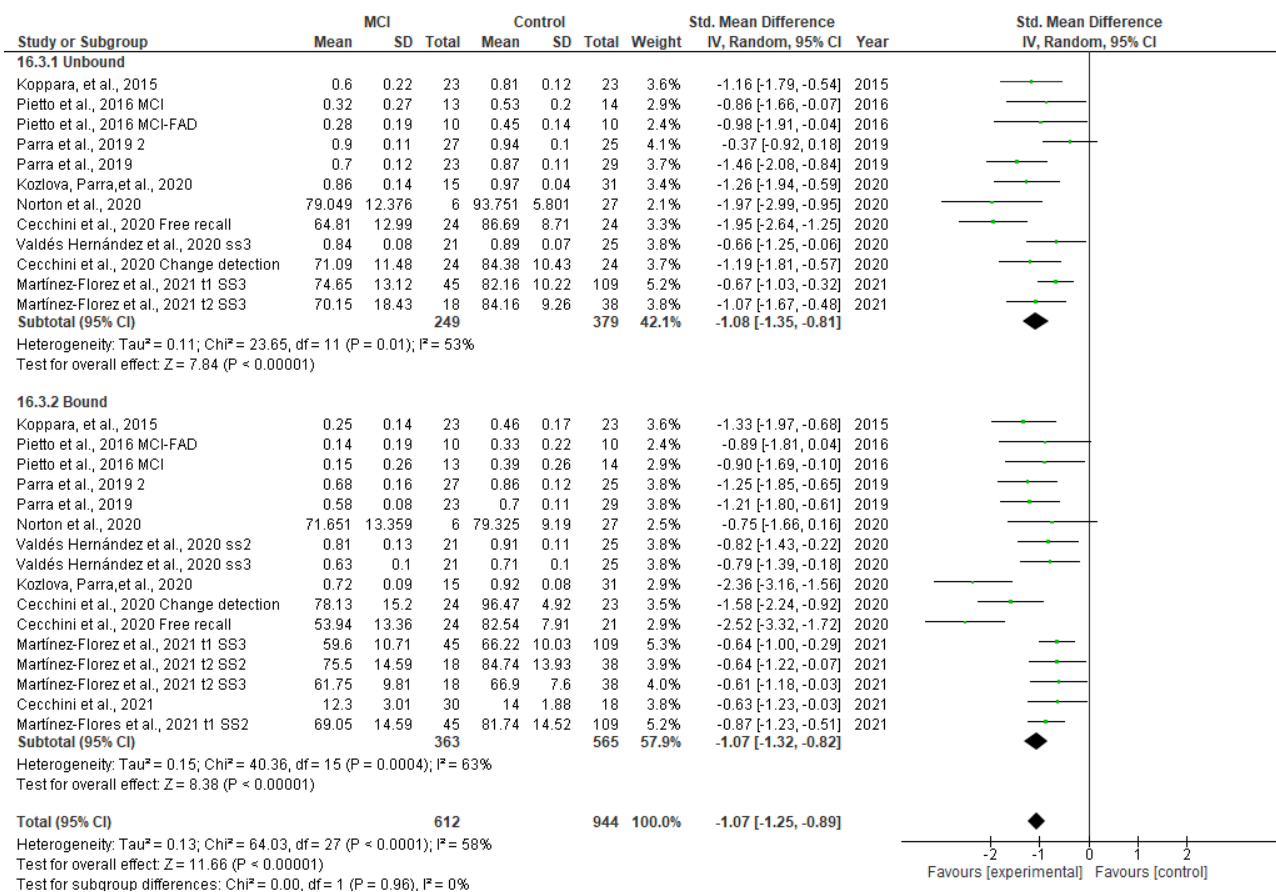
S6. Subgroup analyses

S6.1 Single feature vs. feature binding conditions

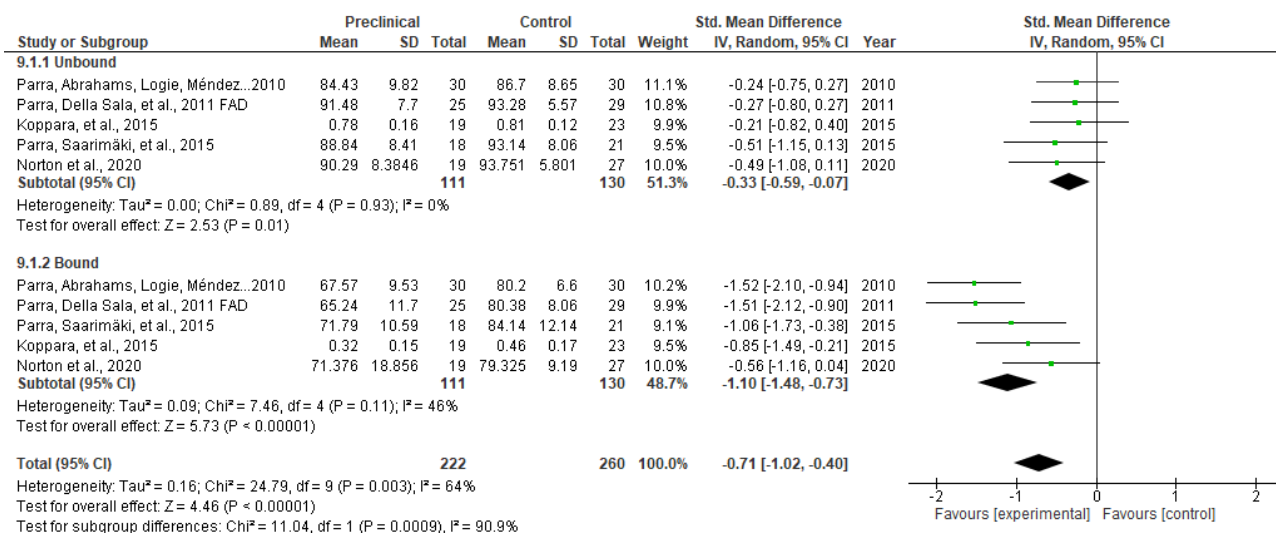
AD



MCI

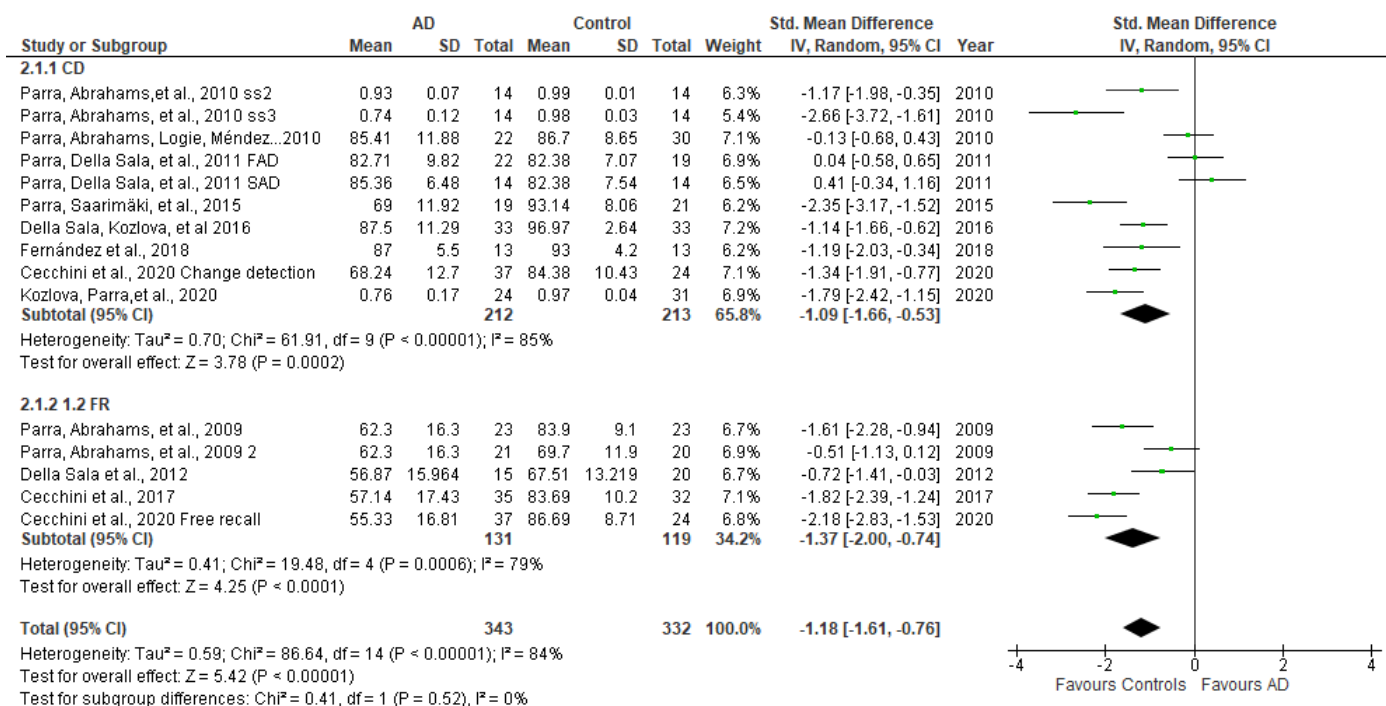


Preclinical

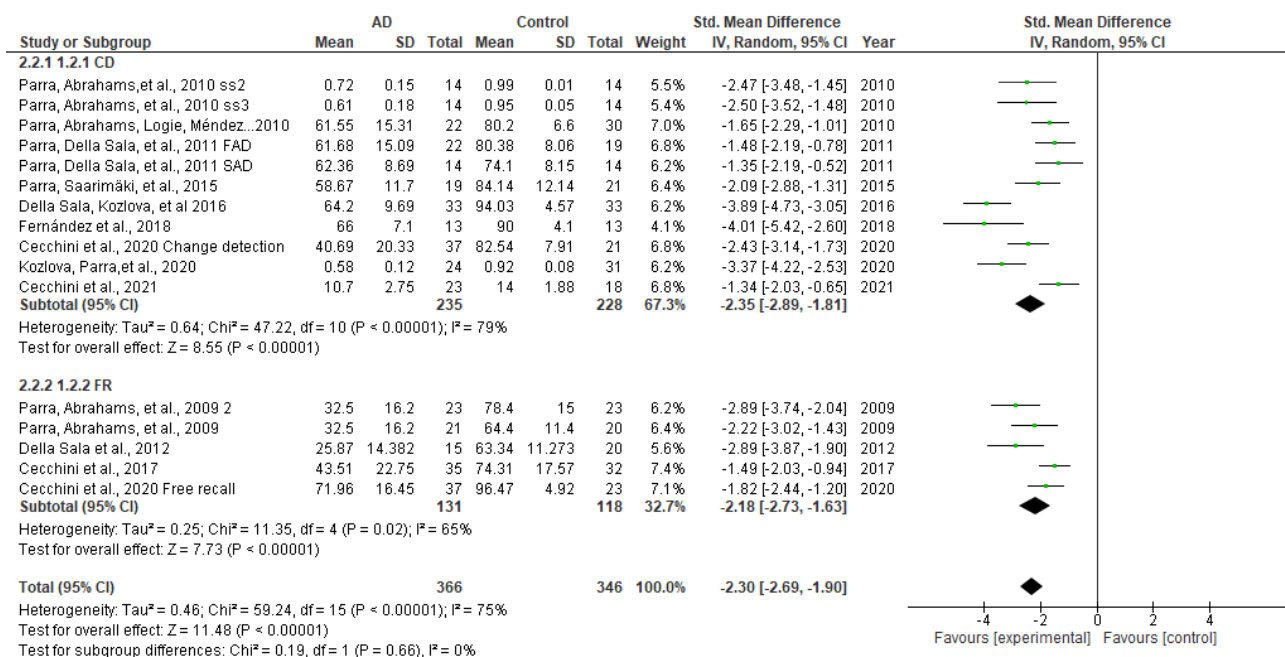


S6.2 Change detection vs. Free recall tasks

AD – Single feature



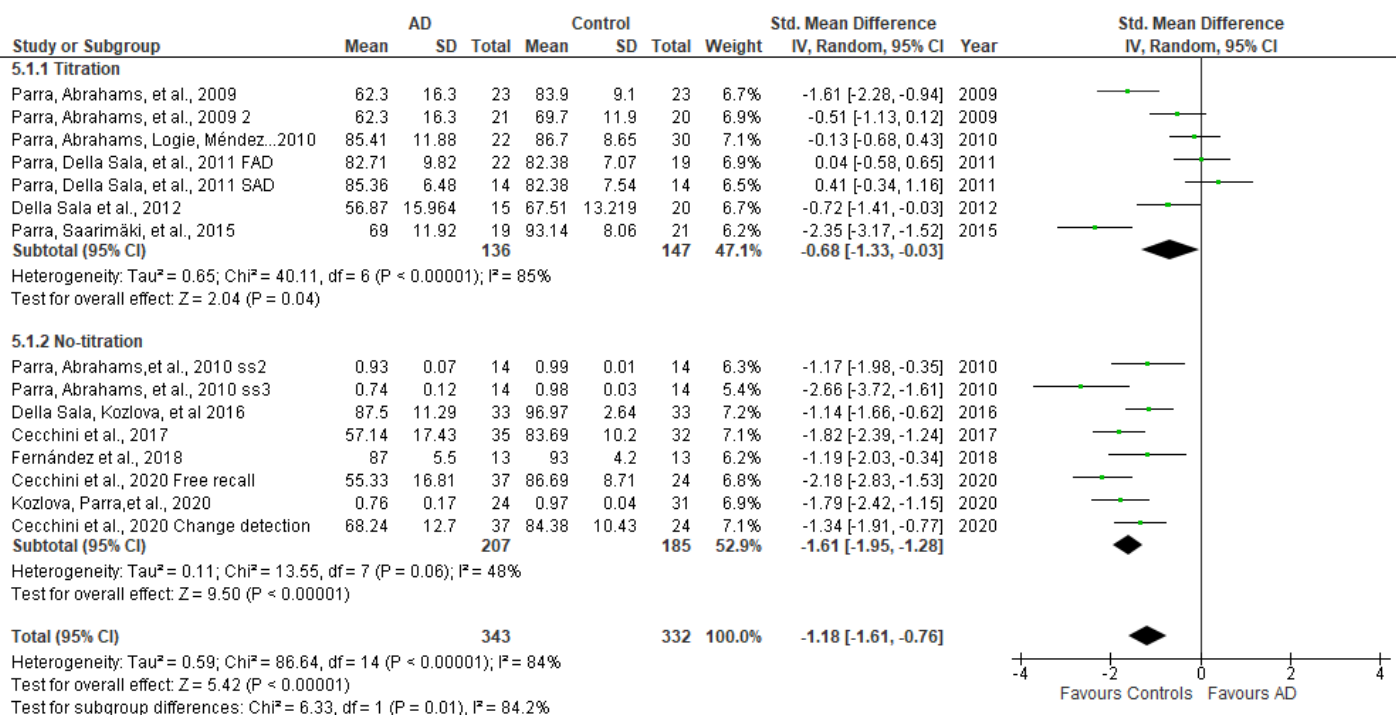
AD – feature binding



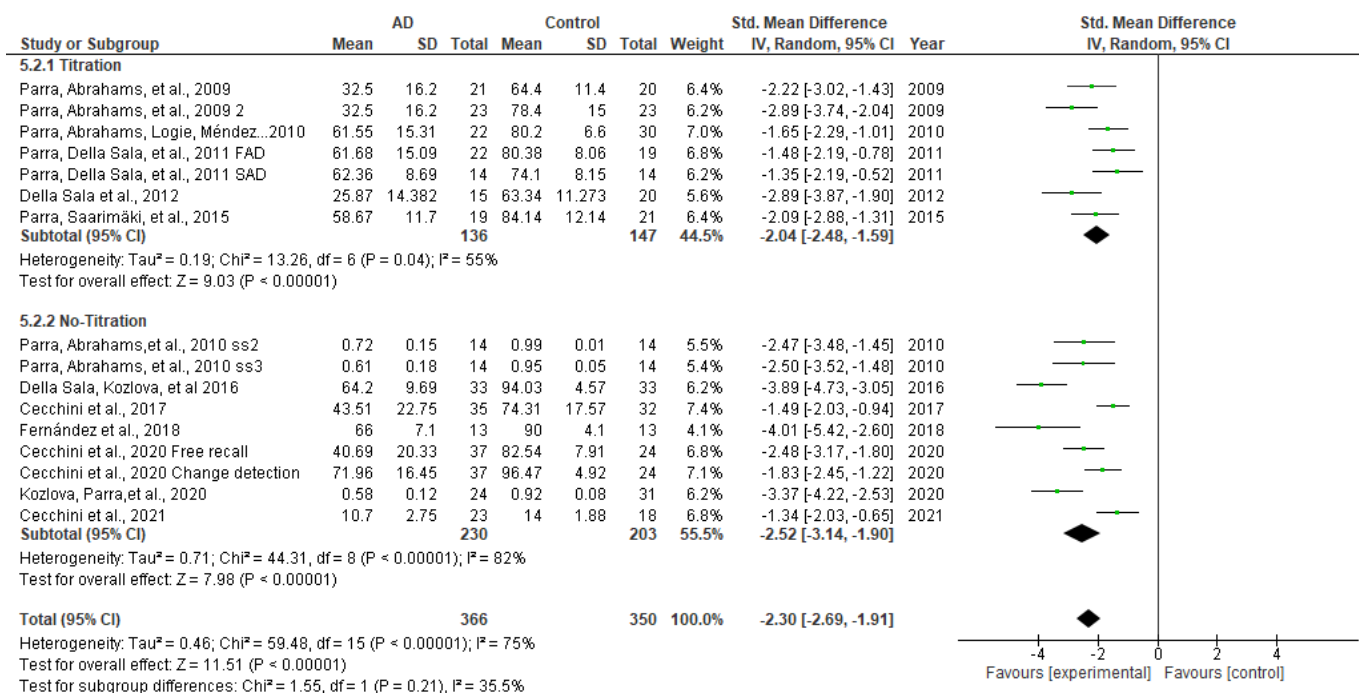
S6.3 Difficulty titration vs. no-titration

AD

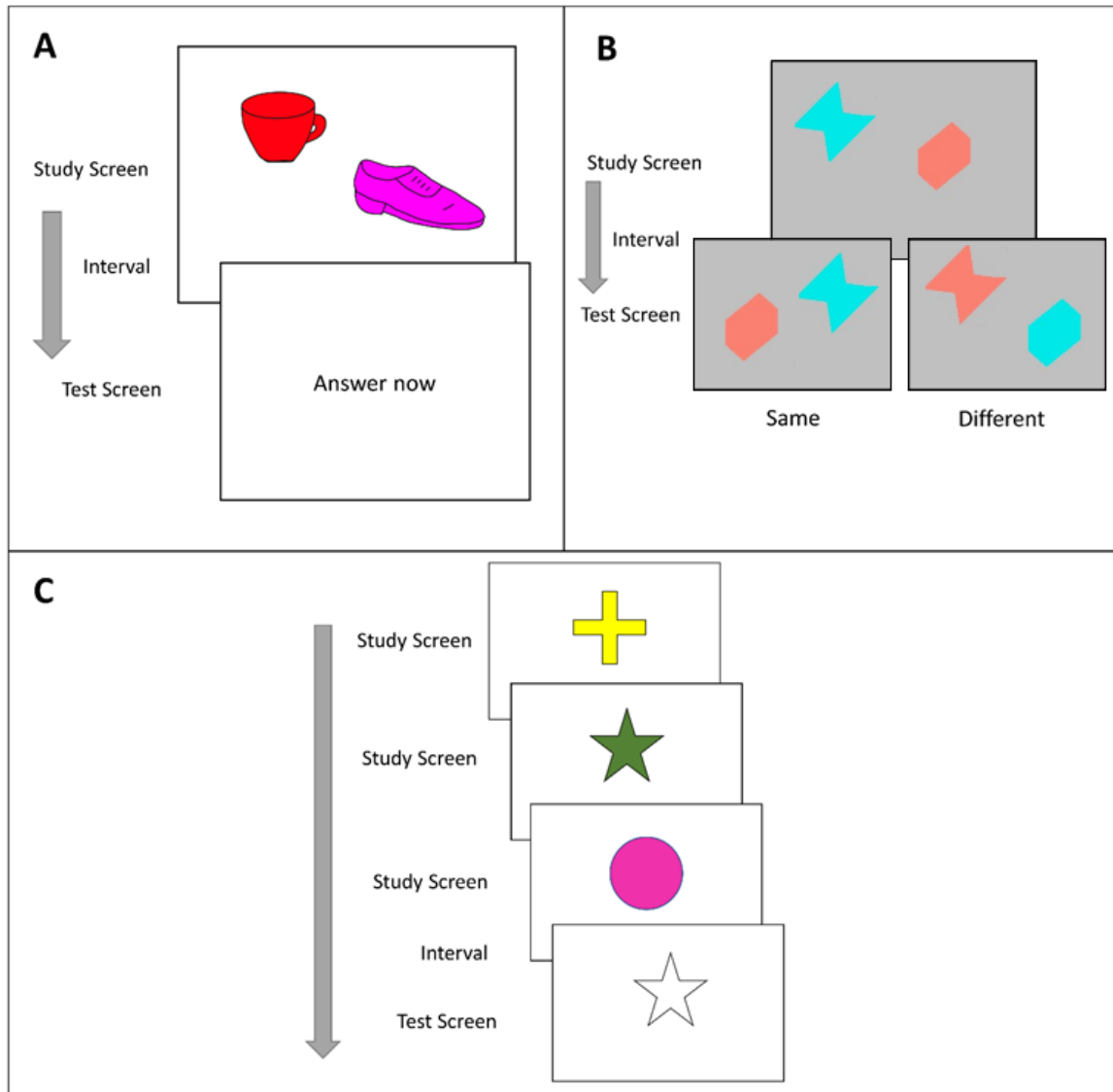
Single feature



Feature binding



S7. Short-Term Memory Conjunctive Binding Tasks Examples



Note. Examples of paradigms used in short-term memory conjunctive binding tasks. **A:** Free recall task, as used in *Della Sala et al., 2012; Parra et al., 2009b; Cecchini et al., 2017a; 2020*. Participants were presented with nameable stimuli (objects and colors) and after a delay they should say aloud the combination of objects and colors previously seen. **B:** Change detection task, as used in *Parra et al., 2010a; 2010b; 2011b; 2015b; 2017a; 2019; Koppara et al., 2015b; Della Sala et al., 2016; Pietto et al., 2016; Fernández et al., 2018; Kozlova et al., 2020; Norton et al., 2020; Valdés Hernández et al., 2020; Cecchini et al., 2020; Martínez-Flores et al., 2021; Cecchini et al., 2021; Fernández & Parra, 2021*. In test phase, participants should recognize if the combinations between shapes and colors are the same or different as shown in the study phase. **C:** Cued recall paradigm, in a continuous presentation method, as used in *Guazzo et al., 2020*. A sequence of shapes was presented, and their colors were presented verbally through headphones. On test phase, participants should recall the missing feature (e.g., in this example, the color green).

Variations in the tasks were found, including: the exposure time at study or test phase, the study-test interval, the activity performed during this interval (e.g., articulatory suppression), the number of stimuli per trial and the type of stimuli on change detection tasks (unnamable shapes-colors binding or color-color binding).