NeuroImage xxx (2009) xxx-xxx



Contents lists available at ScienceDirect

### NeuroImage



journal homepage: www.elsevier.com/locate/ynimg

# Separating event-related BOLD components within trials: The partial-trial design revisited

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#### ARTICLE INFO

 9
 Article history:

 10
 Received 7 January 2009

 11
 Revised 25 April 2009

 12
 Accepted 28 April 2009

 13
 Available online xxxx

 16

5

6 7

#### ABSTRACT

Many event-related fMRI designs involve multiple successive events occurring within a trial, spaced closely 18 in time (e.g., in cued set-shifting paradigms). Yet, it is notoriously difficult to separate the activation 19 components to these sequentially ordered events, given the long evolution time of the BOLD response. One 20 approach to deal with this problem is to omit the second of two successive events (S1 and S2) in a certain 21 proportion of 'partial S1-only' trials. The present article describes a novel method that extends the basic 22 partial-trial design in several ways. As a central new feature it introduces two different delay intervals 23 between S1 onset and S2 presentation, or, in case of S1-only trials, S2 omission. The analysis is based on three 24 BOLD response regressors, one synchronized with S1 onset for short S1-S2 delay trials, another one 25 synchronized with S1 onset for long S1-S2 delay trials, and a third synchronized with S2 onset. The two 26 estimated S1-related activation time courses are then assessed by 'temporal profiling' based on the 27 parameterization of onset latencies, peak latencies, and the area under the curves. Based on this information 28 it is possible (1) to distinguish transient activity elicited with S1 onset from delay-related activity and (2) to 29 identify the activation profile associated with possible 'nogo-type' activity caused by S2 omission. Despite 30 these two new important possibilities, some caution is still advised when interpreting data from the 31 proposed partial-trial design. Yet, in contrast to previous methods, it is possible to identify ambiguous data 32 patterns and, by following an explicit decision scheme, to avoid erroneous conclusions. 33

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#### 25

#### 38 Introduction

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Rapid event-related functional imaging has become a widely used 39 technique that enables the adoption of most paradigms from expe-40 rimental psychology without the need for severely compromising 41 modifications to task designs (e.g., extremely long trial and inter-trial 42durations). Specifically, such imaging designs are suited to extract 43 event-related BOLD response estimates associated with different 44 event, or trial types that occur in a randomly intermixed fashion 45 (Burock, Buckner, Woldorff, Rosen, and Dale, 1998; Dale and Buckner, 46 47 1997; Glover, 1999; Josephs and Henson, 1999). A good example for such trial-type mixing as opposed to trial-type blocking is the set-48 shifting paradigm in its various forms (Corbetta and Shulman, 2002; 49 Monsell, 2003; Pessoa, Kastner, and Ungerleider, 2003; Wager, 50Jonides, and Reading, 2004). In such experiments the key comparison 5152of interest is between trials in which a cognitive set is repeated from 53the previous trial (repeat condition) and trials in which the current set

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1053-8119/\$ – see front matter  $\textcircled{\sc 0}$  2009 Published by Elsevier Inc. doi:10.1016/j.neuroimage.2009.04.075

is changed (switch condition). As repeat and switch trials necessarily 54 occur in an intermixed sequence, an event-related analysis is man- 55 datory and, importantly, it is also perfectly feasible. 56

Yet, the logic of rapid event-related designs breaks down if the 57 contrasted event types cannot be randomly intermixed, that is, when 58 different within-trial events occur in a fixed order. For instance, in 59 task switching experiments researchers have been interested in 60 preparatory processes that are engaged when the upcoming task is 61 indicated by an advance task cue (S1) followed by an imperative 62 target stimulus (S2) which has to be selectively processed within the 63 previously cued task context (Brass and von Cramon, 2002; Braver, 64 Reynolds, and Donaldson, 2003; Bunge, Kahn, Wallis, Miller, and 65 Wagner, 2003; Ruge et al., 2005). Thus, within a given trial, S1 and S2 66 do necessarily occur in a fixed order. Similar situations are given in 67 other paradigms like, for instance, movement preparation (e.g., Toni 68 et al., 2002) or working memory (e.g., Curtis and D'Esposito, 2003). 69 In contrast to standard designs with randomized event order, designs 70 with fixed event order require additional measures to be able to 71 obtain separate BOLD response estimates related to the S1, the delay 72 period, and the S2. The fundamental problem one faces under such 73 conditions is rooted in the extremely long evolution time of the 74 canonical event-related BOLD response which is estimated to be 20 s 75 or more in duration, thus implicating a massive overlap of successive 76

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Fig. 1. Graphical representation of the extended partial-trial design, including the generation of synthetic BOLD data and their analysis using FIR basis sets (for details, see the main text).

BOLD components. Two general approaches have been used to dealwith this problem.

One approach relies on the realization of long and variable 79 intervals between S1 and S2 (Curtis and D'Esposito, 2003), ranging, 80 for instance, between 9 and 18 s (Rowe, Toni, Josephs, Frackowiak, and 81 Passingham, 2000) or between 4 and 12 s (Sakai and Passingham, 82 2003). To obtain separate estimates for transient S1-related activity, 83 delay-related activity, and S2-related activity, model regressors for 84 85 each of the three BOLD components must be included in a General Linear Model (GLM), each convolved with an assumed, or empirically 86 87 derived (Postle, Zarahn, and D'Esposito, 2000) hemodynamic response function. 88

The other method, suited for much shorter S1-S2 intervals, is 89 based on the use of so-called partial trials (Ollinger, Corbetta, and 90 Shulman, 2001; Ollinger, Shulman, and Corbetta, 2001; Serences, 91 2004; Shulman et al., 1999).<sup>1</sup> The basic setup requires the implemen-92tation of a single S1-S2 interval (e.g. 2.5 s), but the S2 is omitted in a 93 certain proportion of trials (i.e., 20-33%). This way it is possible to 94obtain independent estimates for S1-related activation and, by 95partialling out the S1-related component, also for S2-related activa-96

tion. In contrast to experimental designs that rely on long inter-event 97 durations, the use of short S1–S2 intervals in the partial-trial design 98 seems to be advantageous for several reasons. First, the total 99 experiment time decreases considerably, or, in other words, the 100 number of trials one can realize within a reasonable amount of time is 101 greater, and so is statistical power. Second, the partial-trial design 102 avoids another problem associated with long S1–S2 intervals, which is 103 that long S1–S2 intervals are likely to induce neural and mental 104 processes during the delay period that might not be directly task- 105 related (except for certain situations, such as working memory tasks 106 with high load levels).

However, there are also two major disadvantages of the standard 108 partial-trial method. First, S2 omission can potentially cause 'nogo- 109 type' artifacts, that is, neural activity which is specifically elicited by 110 the omission of the S2. Importantly, there is no way to distinguish such 111 nogo-related BOLD activation from genuine S1-related activation. 112 Also, nogo-related BOLD activation cannot be partialled out with res- 113 pect to S2-related activation, since the nogo-related component is, by 114 definition, absent in full S1–S2 trials. Consequently, the estimation of 115 S2-related activation becomes distorted (cf., Fig. 3C in this paper). 116 Second, it is not possible to distinguish S1-related activation that is 117 transiently elicited with S1 onset from delay-related activation that is 118 maintained throughout the S1-S2 interval. To be able to distinguish 119 these two important cases some sort of reference BOLD response 120 would be crucially needed because of the well-known fact that there 121 exists no generic BOLD response shape (i.e., a canonical hemodynamic 122 response function or HRF) independent of brain regions and subjects 123 (Bellgowan, Saad, and Bandettini, 2003; Formisano and Goebel, 2003; 124 Huettel and McCarthy, 2001; Saad, Ropella, Cox, and DeYoe, 2001). For 125 instance, if brain region A showed a greater BOLD response width than 126

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<sup>&</sup>lt;sup>1</sup> Alternatively, some researchers have employed an approach using S1–S2 intervals that are randomly varied within a relatively narrow range of, for instance, 2.5–5s (Luks, Simpson, Feiwell, and Miller, 2002) or even 0–1.5s (Gruber, Karch, Schlueter, Falkai, and Goschke, 2006). This approach crucially relies on the assumption that S1-related and S2-related activation is independent of the length of the S1–S2 interval (Serences, 2004). This assumption is violated in case of delay-related BOLD activation that is sustained between S1 and S2 as such activation persists for a longer duration with a corresponding increase in the S1–S2 interval. Unfortunately, it is impossible to tell from the data whether or not delay-related activation was present, and thus, whether or not the obtained BOLD estimates are valid.

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brain region B, it would not be legitimate to conclude that region A isactivated in a more sustained way.

In the present paper, a novel partial-trial method is proposed that 129 130aims to overcome the two major limitations mentioned above. The key innovation is to enable better estimation of a reference BOLD response 131 from which to detect effects related to both S2 omission and transient 132vs. sustained S1 activation. Specifically, the approach extends the 133partial-trial design which is extended by introducing at least two 134different S1-S2 delay intervals, in combination with the use of 135separate S1-related model regressors for the different S1-S2 interval 136levels. After model estimation, we implemented a 'temporal profiling' 137 method similar to the approach previously used in the context of a 138full-trial design (Ruge, Brass, Lohmann, and von Cramon, 2003). 139Specifically, the two S1-related regressor estimates are cross-refer-140 enced based on parameterizations of the respective onset latency, 141 peak latency, and area under the curve. Thereby, it becomes possible 142 to distinguish BOLD activation patterns reflecting neural activity (1) 143

elicited transiently following S1 onset, (2) maintained throughout the 144 delay period, or (3) triggered by S2 omission specifically in S1-only 145 trials. The basic rationale of this extended partial-trial method is first 146 developed by using synthetic data. Subsequently, the method is 147 applied to a real data set to demonstrate that the fine-grained analysis 148 of temporal activation profiles is feasible under realistic conditions. 149

#### Methods (synthetic data)

The main goal of the simulations described here, was to develop the 151 basic rationale of the extended partial-trial design. The simulations 152 were implemented with the R software package (R-Development- 153 Core-Team, 2005). All simulations were performed within the same 154 basic GLM-based deconvolution scheme without an assumed BOLD 155 response shape for the model regressors (Ollinger et al., 2001). This 156 model is equivalent to the finite-impulse-response (FIR) basis set 157 implemented in the SPM software. Synthetic BOLD time courses were 158





**Fig. 2.** Graphical representation of the different possible patterns of neural activity in partial-trial designs for full trials (A–C) and for partial trials (D–F). (A) Transient S1-related activity plus optional sustained delay-related activity; (B) S2-related activity or S1S2-interval termination-related activity; (C) transient S1-related and S2-related activity (or S1S2-interval termination-related activity plus optional sustained delay-related activity; (D) Transient S1-related activity plus optional sustained delay-related activity; (P) Transient S1-related activity plus optional sustained delay-related activity; (F) Transient S1-related and S2-omission-related activity (or S1S2-interval termination-related activity) plus optional sustained delay-related activity; (F) Transient S1-related and S2-omission-related activity (or S1S2-interval termination-related activity) plus optional sustained delay-related activity.

Please cite this article as: Ruge, H., et al., Separating event-related BOLD components within trials: The partial-trial design revisited, NeuroImage (2009), doi:10.1016/j.neuroimage.2009.04.075

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created at a sampling rate of one time step every 1.0 s by convolving a
canonical BOLD response model with the sum of input functions for
neural activity associated with different events (for an example, see
Fig. 1 left-hand side).

The canonical BOLD response was modeled according to the gamma function given by equation 1 using the following parameters: a1 = 11.0, a2 = 12.0, b1 = 0.35, b2 = 0.9, c = 0.1, d1 = a1\*b1, and d2 = a2\*b2.

BOLD(t) = 
$$(t/d1)^{a1}e^{-(t-d1/b1)} - c(t/d2)^{a2}e^{-(t-d2/b2)}$$
 (1)

The neural input functions for the different event types were created as described below and were meant to cover all the different neural activity patterns that are theoretically possible in partial-trial designs (for an overview, see Fig. 2).

The neural input function for transient S1-onset-related activity 173had the duration of 1 time step. For delay-related activity, two 174different neural input functions were optionally implemented to cover 175the two most plausible scenarios. One such profile was designed to 176mimic constant neural activity throughout the entire S1-S2 delay as 177 would be expected, for instance, in case of persistent visual 178 stimulation by a flickering stimulus contrast. To account for the 179180 refractoriness of the BOLD signal, we created graded amplitude profiles given by [1, 0.66] covering 2 time steps for short S1–S2 delay 181 trials and [1, 0.66, 0.66, 0.66, 0.66] covering 5 time steps for long S1-S2 182 delay trials. The alternative delay-related activity profile was designed 183 to mimic preparatory processes as would be expected, for instance, in 184 185the selective attention paradigm. Specifically, we created U-shaped amplitude profiles given by [1.2, 0.6] for short S1-S2 delay trials and 186 [1.2, 0.6, 0.3, 0.5, 0.8] for long S1–S2 delay trials. For both delay-related 187 activity profiles, the exact values were chosen rather arbitrarily. 188 189 Importantly, though, other versions not reported here, yielded highly 190comparable results. As will become clear in the light of the actual results, the two activity profiles that we do report also produced 191 qualitatively highly similar results. The neural input function for S2-192related activity was set to a fixed length of 1 time step. To simulate the 193194 impact of nogo-type activity in response to S2 omission, a neural input function was added assuming neural activity with the duration of 1 195

time step and time-locked to the end of the S1–S2 interval exclusively 196 for partial S1-only trials. Additionally, we considered neural activity 197 associated with the termination of the delay interval, which was 198 modeled by introducing neural input with the duration of 1 time step 199 and time-locked to the end of the S1-S2 interval for both full S1-S2 200 trials and S1-only trials. This form of neural activity has typically been 201 ignored in empirical and modeling studies, but there are important 202 reasons to take this type of activity into account (Shulman et al., 203 2002). Importantly, although delay termination occurs at the same 204 point in time as both, potential nogo-type processes and S2-related 205 processes, it can still be uniquely characterized. First, in contrast to 206 nogo-type processes which occur exclusively in S1-only trials due to 207 S2 omission, delay termination does also occur in full S1-S2 trials. 208 Second, in contrast to S2-related processes, which occur exclusively in 209 full S1-S2 trials, delay termination does additionally occur in S1-only 210 trials. 211

In a second step of each simulation, a GLM was estimated to predict 212 the synthetic BOLD time course via two regressors time-locked to S1 213 onset (labeled "R1\_1" for the short interval and "R1\_2" for the long 214 interval; see Fig. 1, right-hand side) and one regressor time-locked to 215 S2 onset (labeled "R2"), plus as constant term for the overall mean 216 activation level. Regressors associated with S1 and S2 each covered an 217 interval of 20 s always starting from the respective stimulus onsets of 218 S1 and S2. No particular BOLD response shape was assumed for the 219 regressors. Thus, for each regressor estimate 20 free parameters were 220 to be determined through the GLM estimation process (see Fig. 1, 221 bottom panel). A first set of simulations aimed at demonstrating some 222 of the more general properties of the method. These simulations were 223 run without noise added to the synthetic BOLD time courses to obtain 224 a clear graphical representation of the results (adding noise did not 225 systematically alter the results). Subsequent simulations were run 226 with noise added to demonstrate some relevant statistical properties 227 of the results. The noise was composed of Gaussian noise (amplitude 228 0.7) plus sine waves at 1 Hz (amplitude 0.3) and 0.2 Hz (amplitude 229 0.3). See Figs. 4 and 5 for the contributions of signal (i.e., the 230 amplitudes of the neural activity components) relative to noise. 231

The synthetic BOLD time courses were based on a sequence of 144  $_{232}$  trials Two thirds of all trials (96) were full S1–S2 trials and one third of  $_{233}$ 



Fig. 3. Results of four simulations run for the extended partial-trial design assuming different single S1-related neural activity components for each simulation. The S2-related neural activity component was always present and identical across all 4 simulations. Note, that for the simulation of nogo-type activation due to S2 omission (panel C) the expected S1-related BOLD activation equals zero as genuine S1-related neural activity was absent.

Please cite this article as: Ruge, H., et al., Separating event-related BOLD components within trials: The partial-trial design revisited, NeuroImage (2009), doi:10.1016/j.neuroimage.2009.04.075

all trials (48) were partial S1-only trials (S2 omitted). The S1–S2 delay 234 235 interval was randomly assigned to be either 2 s or 5 s. Following the S2 (or S2 omission), another constant 'blank' interval of 2.5 s was added 236 237to keep the simulated trials as close as possible to the empirical study design described later in this paper. Thus, trials could have a total 238duration of either 4.5 s or 7.5 s depending on the delay interval. The 239total of 144 trials was equally split into short and long S1-S1 delay 240trials. The ITI duration was determined by the variable duration of 48 241 242'no-event' trials (i.e., blank intervals) randomly interspersed and 243 matched in length to the experimental trials (4.5 or 7.5 s, depending 244on the S1–S2 delay), to again follow the structure of the empirical study design. Since no-event trials were allowed to occur in direct 245sequence, the distribution of ITIs comprised intervals that could be 246247twice or, rarely, even three times the duration of one trial, thereby approximating an exponential ITI distribution. The ITI was constructed 248 in this particular way to enable two alternative estimation procedures 249 for the removal of inter-trial BOLD overlap. First, it is possible to 250 implement an 'implicit baseline' estimation procedure without 251explicit modeling of baseline activity during the ITI period (Ollinger 252et al., 2001). Second, it is possible to explicitly estimate activity during 253the no-event trials and later subtract this 'baseline' activation from 254activity during the experimental trials, separately for the two S1-S2 255256delay conditions (Burock et al., 1998). Since both estimation procedures produced qualitatively similar results, we only report in detail 257the results from the more commonly used implicit baseline estimation 258procedure (the same holds for the empirical data). For the simula-259tions, we also tested different ITI distributions. Since other recom-260261mended ITI distributions (Hagberg, Zito, Patria, and Sanes, 2001) yielded qualitatively similar results, we do not further elaborate on 262this matter in the present paper. 263

#### 264 Results (synthetic data)

#### 265 Simulation set I – the basics

The goal of the first set of simulations (no noise added) was to demonstrate the basic characteristics of the fitting procedure for the 4 basic activity components in isolation (results depicted in Fig. 3).

Figs. 3A and B depict the results for transient activity time-locked 269to S1 onset and delay-related activity, respectively. First, and not 270surprisingly, GLM estimation is perfectly suited to obtain distortion-271272free time course estimates for all 3 model regressors. Second, and more importantly, the BOLD estimates R1\_1 and R1\_2 can be cross-273referenced and compared, thereby enabling us to determine the 274degree to which S1-related regressor estimates reflect delay-related 275activation. In comparison to transient S1-related activation, delay-276277related activation leaves BOLD onset latencies unaffected, but R1\_2 reaches its maximum later, resulting in a greater area under the 278curve as compared to R1\_1. As further substantiated in the context of 279the second set of simulations, the difference between the area under 280the curves for R1\_2 vs. R1\_1 ('area-difference index'), but not the 281 282peak latency shift, can be used as an unambiguous measure of delay-283related activation.

Figs. 3C and D depict the results for nogo-type activity and delay-284termination-related activity time-locked to the end of the S1-S2 delay 285interval. These two types of activity result in identical R1\_1 and R1\_2 286 287estimates. Importantly, this pattern can be easily distinguished from genuine S1-related activity time-locked to the onset of S1 presenta-288 tion. The crucial difference lies in the relationship between BOLD 289 onset latencies for R1\_1 and R1\_2. In contrast to genuine S1-related 290activity, which is characterized by equal BOLD onset latencies for R1\_1 291and R1\_2 (Figs. 3A and B), both, nogo-type activity and termination-292related activity result in a temporal shift of R1\_2 relative to R1\_1. This 293shift is indicated by delays of onset latencies and peak latencies for 294R1\_2 relative to R1\_1 (equal to the time difference between the two 295296 respective S1-S2 intervals). Yet, despite identical estimates for R1\_1

and R1\_2, nogo-type activity and termination-related activity have 297 strikingly different impacts on R2. Specifically, nogo-type activity does 298 not only load on regressors R1\_1 and R1\_2, but also loads negatively 299 on R2, thereby leading to a severe distortion. Thus, the R2 estimate 300 should not be interpreted in this context. Unfortunately, since we do 301 not have a priori expectations regarding the exact strength of S2- 302 related activation, it is not possible to determine the presence or 303 absence of S2 distortion (unless R2 becomes clearly negative with 304 respect to fixation, which strongly suggests distortion due to S2 305 omission). If this were the case, R2 distortion could be used as an 306 indicator to distinguish between nogo-type activity and termination- 307 related activity.

#### Simulation set II – the significance of the area-difference index 309

The second set of simulations aimed at demonstrating the central 310 relevance of the area-difference index (ADI; i.e., the difference of the 311 area under the curve for R1\_2 minus R1\_1 expressed as the percentage 312



**Fig. 4.** Simulation results for different neural activity components occurring in isolation (A–D). The panels on the left depict the neural activity patterns for short (gray) and long delay trials (black). The panels on the right depict the corresponding BOLD estimates R1\_1 (short delay; gray) and R1\_2 (long delay; black). Included also are the area-difference indices ( $\Delta$  area)  $\pm$ 95% confidence interval, representing the difference between the area under the curves for R1\_2 minus R1\_1.

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**Fig. 5.** Simulation results for three types of combinations (A–C) of different neural activity components (see sub-headings for the specific components). For each of the three combination types, one activity component was held constant while the relative contribution of another, variable, component was increased in three steps (rows 1 to 3). The panels on the left depict the neural activity patterns for short (gray) and long delay trials (black). The panels on the right depict the corresponding BOLD estimates R1\_1 (short delay; gray) and R1\_2 (long delay; black). Included also are the area-difference indices ( $\Delta$  area)  $\pm$  95% confidence interval, representing the difference between the area under the curves for R1\_2 minus R1\_1. Note, that to achieve a better graphical fit, the *y*-axes of the neural activity graphs were adaptively scaled with changing contributions of the respective variable activity component; the absolute strength of the respective constant activity component was factually constant (and not variable as suggested by the graphical representation).

increase relative to the total area under both curves). Specifically, we 313 show that the ADI is both, necessary and sufficient for determining the 314 presence of delay-related activity. This demonstration also required 315 the inclusion of the ADI estimation error (expressed in terms of the 316 95% confidence interval). The confidence interval was computed 317 318 based on 40 independently generated synthetic BOLD time courses with noise added. Since the R2 estimate is not relevant in this context, 319 we only report the results for R1\_1 and R1\_2. Also, we did not 320 distinguish between nogo-type activity and termination-related 321 activity, as R1\_1 and R1\_2 estimates are identical for these cases. 322

323 We started by computing the ADI for single activity components (Figs. 4A–D). Clearly, the ADI is significant for different types of delay-324 related activation (Figs. 4C, D), but not for transient activity time-325locked to the onset of S1 or time-locked to the end of the S1-S2 326 interval (Figs. 4A, B). Yet, at first sight, the ADI does not seem to be the 327 only parameter that discriminates delay-related activation from the 328 other two cases. Specifically, only delay-related activation seems to be 329 specifically characterized by equal onset latencies for R1\_1 and R1\_2 330 plus delayed peak latency for R1\_2 relative to R1\_1. Importantly, 331 however, this conclusion is invalid, as becomes clear when combina-332 tions of different activity components are taken into consideration. 333 Fig. 5 depicts the results of this analysis. 334

Fig. 5A depicts results that demonstrate that the ADI is necessary to determine the presence of delay-related activation. In the underlying simulation the neural activity component for transient activation time-locked to the onset of S1 was held constant while the amplitude 338 of the termination-related component was increased in three steps 339 (i.e., the delay-related component was consistently zero). Crucially, 340 increasing the relative contribution of the termination-related 341 component also increases the peak latency of R1\_2 relative to R1\_1 342 whereas onset latencies stay constant. This is exactly the same pattern 343 as for delay-related activation (see Figs. 4C–D). By contrast, the ADI is 344 still perfectly suited to discriminate delay-related activation (ADI 345 significantly greater than 0) from all three cases in which delay- 347 related activity was absent (ADI not significantly different from 0).<sup>2</sup> 347

Figs. 5B, C depict results that demonstrate that the ADI is also 348 sufficient to determine the presence of delay-related activation 349 irrespective of other overlapping activity components, be it additional 350 transient activity time-locked to S1 onset (Fig. 5B) or additional 351 termination-related activity (Fig. 5C). Fig. 5B shows the impact of 352 increasing the relative neural activity component for transient S1- 353 onset-locked activity in three steps. Fig. 5C shows the impact of 354 increasing the relative neural activity component for termination- 355

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<sup>&</sup>lt;sup>2</sup> One might argue that the striking twin-peak structure created by adding a termination-related activity component (Fig. 5A) is clearly different from the single-peak structure created by delay-related activation (Fig. 4C). Yet, this caveat is invalidly based on a description of superficial properties of the curves. For instance, reducing the long S1–S2 interval from 5 to 3s would easily merge the two peaks into one single peak.

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related activity in three steps while the delay-related activity 356 357 component stays constant. Clearly, greater relative contributions of both, the transient S1-onset-related component and the termination-358 359related component imply smaller ADI values. Yet, the simulation also shows that the estimation error decreases proportionally. Thus, the 360 ADI seems well suited to identify a delay-related activity component 361 irrespective of the strength of other overlapping transient activity 362 components. 363

364To summarize, the simulation results depicted in Figs. 4 and 5 365 demonstrate that the extended partial-trial design is well suited to extract important details about the underlying neural activity com-366 ponents based on evaluating both, the relative onset delay of R1\_2 367 relative to R1\_1 and the ADI index. By contrast, other, more superficial 368 properties of the R1\_1 and R1\_2 estimates, like single-peak vs. twin-369 peak structure or the extent of the relative peak delay, are not suited 370 for valid inferences about the underlying neural activity components. 371 For instance, the BOLD estimates in Figs. 5A (row 3) and C (row 3) look 372 quite similar, yet the underlying neural activity components are 373 decisively different as becomes clear when taking into account the ADI 374 index. 375

#### 376 Methods (empirical data)

#### 377 Material and procedure

For empirical validation of the extended partial-trial method, we 378 devised three different versions of a visual selective attention para-379 380 digm, in which S1 and S2 were two visual stimulus events (see Fig. 6). Additionally, an auditory stimulus event (a sound that could easily be 381 discriminated from scanner noise; duration 300 ms) was included to 382 383 mark either the start or the termination of the S1-S2 interval. The delay interval separating S1 and S2 was variable (2.0 or 5.0 s). The S1 384385was a randomly chosen attentional cue indicating the currently relevant color (blue vs. green '+') and the S2 was a target–distractor 386 pair (blue and green 'O' located unpredictably to the left and right of 387 the screen center). Participants had to indicate the location of the 388 relevant 'O' (as defined by the preceding color cue) by pressing a 389 spatially compatible response key with their right index or middle 390 finger The cue was displayed for 500 ms, followed by a black fixation 391 cross for the remainder of the cue–target interval (not shown in Fig. 392 6). The target–distractor pair remained on the screen until response 393 execution or until the response deadline was reached after 1.25 s. The 394 end of a trial was reached after the response deadline had elapsed 395 (irrespective of the actual time of response) and after the presentation 396 of the fixation cross for another 1.25 s. Trials were separated by a 397 variable inter-trial interval (ITI) during which the black fixation cross 398 was displayed (for details, see below).

For each subject the total number of trials was 192, split into 128 400 'full S1–S2' trials and 64 'partial S1-only' trials. The total of 192 trials 401 was equally split into short and long S1–S1 delay trials. Equivalent to 402 the simulations, the ITI duration was determined by the variable 403 duration of 64 'no-event' trials randomly interspersed and matched in 404 length to the experimental trials, approximating an exponential 405 distribution of ITIs (for details, see Simulation methods). The expe- 406 riment took approximately 25 min and was run without interruption 407 within a single scanning session. 408

Fig. 6 depicts the three different experimental versions that were 409 run for three different groups of participants. First, the three experi-410 ment versions differed with respect to the time point of auditory 411 stimulation. The sound was played either with S1 onset (experiment 412 version C) or with S1S2-interval termination (experiment versions A 413 and B). Thus, we expected to detect auditory cortex activation that was 414 either transient and time-locked to the onset of the S1 (C) or transient 415 and time-locked to the termination of the S1–S2 interval (A and B). By 416 contrast, the type of the visual cue (S1) was the same across all 417 experiment versions. Thus, we expected transient activation time-418



**Fig. 6.** Experimental design used for generating the empirical data. There were 3 different experiment versions (panels A–C). These versions differed with respect to the time point of sound presentation and with respect the type of target (S2) omission in case of partial S1-only trials. CTI = Cue (S1)–Target (S2) interval. The cue was a blue or green fixation cross presented at time point zero (here, gray standing for green). The target was a pair of blue and green circles (black = blue and gray = green in reality). The black fixation cross (black was also black in reality) was displayed throughout the entire trial.

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locked to the onset of the S1 plus possible delay-related activation
across all experiment versions, particularly within brain areas that
support the preparatory control of attention.

422 Second, the three experiment versions differed with respect to the type of S2 omission in partial trials. For one group of participants 423 (experiment version B), the target/distractor pair was completely 424 omitted in partial trials (Fig. 6B). In this group, the termination of the 425S1-S2 interval was marked by the sound event. Thus it was still 426 427 possible to identify the end of the delay period. For another group of participants (experiment versions A and C) target and distractor were 428 429 indistinguishably presented in the same color in partial trials (Figs. 6A 430 and C). For both types of S2 omission, participants were instructed not to respond (since a correct response was not defined). Both types of S2 431432 omission were intended to exclude target selection and response selection/generation processes. Hence, in all three experiment ver-433 sions, the S2 event (i.e. entailing those processes that occur in full 434trials only) was not simply defined by the presentation of a second 435visual stimulus, but rather by the implementation of target selection 436 and response selection/generation processes, which were only pos-437 sible when distinguishable target and distractor stimuli were dis-438 played. Yet, specifically in experiment versions A and C, S1S2-interval 439termination (i.e., entailing those processes that occur both, in full trials 440 441 and partial trials at the end of the S1–S2 interval) was defined by the onset of the two "O" stimuli irrespective of whether they were 442 presented in the same color (partial trials) or in different colors (full 443 trials). The presentation of two indistinguishable "O" stimuli in partial 444 trials was intended to create a specific pattern of neural activity in the 445446 ventral visual stream driven by both, the visual processing of the S1 (attentional cue) and the visual processing of the two "O" stimuli 447 irrespective of their respective colors in both, partial and full trials. 448 Thus, we expected a combination of S1-related activation time-locked 449 450to the onset of the attentional cue plus activation elicited by the two "O" stimuli that marked the termination of the S1-S2 interval. 451

#### 452 Subjects

13 right-handed human participants with no evidence of neurological compromise took part in this study (age range: 20–28; 9 females,
4 males). All participants gave informed consent according to the
guidelines set by the Dresden University of Technology Ethics Committee. The participants were paid €8 for each hour of participation.

#### 458 Imaging procedure

Whole-brain images were acquired on a Siemens 3 T whole-body 459Trio System (Erlangen, Germany) with a 16 channel circularly 460 461 polarized head coil. Headphones (NordicNeuroLab) and earplugs dampened scanner noise. Both structural and functional images were 462 acquired for each participant. High-resolution structural images 463  $(1.0 \times 1.0 \times 1.0 \text{ mm})$  were acquired using an MP-RAGE T1-weighted 464 sequence  $(TR = 1900 \text{ ms}, TE = 2.26 \text{ ms}, TI = 900 \text{ ms}, flip = 9^\circ).$ 465466 Functional images were acquired using a gradient echo planar 467 sequence (TR = 1620 ms, TE = 30 ms, flip =  $80^\circ$ , interleaved slice acquisition, slice gap = 0), sensitive to blood oxygen level-dependent 468 (BOLD) magnetic susceptibility. Each volume contained 26, 5.0 mm 469thick slices (in-plane resolution  $4.0 \times 4.0$  mm). 470

The experiment was controlled by Eprime 1.2 software (Psychology Software Tools) run on a Windows-XP PC. Stimuli were projected
to participants via Visuastim digital goggles (Resonance Technology,
Inc.; Northridge, USA). A fiber-optic, light-sensitive key press was used
to record participants' behavioral responses.

#### 476 Data analysis

The empirical data set was analyzed with SPM 5 for pre-processing and for the initial FIR model estimation step. For the subsequent finegrained temporal profiling of time course estimates, the R software 479 package was used. Preprocessing included slice-time correction, rigid 480 body movement correction (3 translation, 3 rotation parameters), 481 normalization of the functional images by directly registering the 482 mean functional image to the standard MNI EPI template image 483 provided by SPM 5 (the resulting interpolated spatial resolution was 484  $4 \times 4 \times 4$  mm), and smoothing of the functional images (Gaussian 485 Kernel, FWHM=4 mm).

The estimation procedure was analogous to that used for the 487 synthetic data. Specifically, we included two S1-related model 488 regressors (R1\_1 and R1\_2; time-locked to the onset of S1) and one 489 S2-related model regressor (R2; time-locked to the onset of S2). We 490 also examined the effects of a slightly different GLM estimation 491 approach that included two separate regressors for S2-related activity 492 at the two delay intervals. This analysis was meant to account for 493 possible distortions due to differential non-linear summation effects 494 of S1-related and S2-related BOLD activation associated with the two 495 different delay intervals. Since the two GLM versions yielded 496 qualitatively similar results, we only report the results from the 497 GLM that included a single regressor for S2-related activity. Model 498 regressors were based on FIR basis sets including 21 time steps 499 covering an interval of 34 s. In an initial whole-brain analysis, voxels 500 were identified for each subject that exhibited significant S1-related 501 activation (based on an F-test for systematic variance across the 42 502 data points estimated for R1\_1 and R1\_2 with p(F) < 0.001). 503

Finally, we attempted to directly identify voxels that were 504 specifically associated with activity elicited by S2 omission. To this 505 end, we computed an additional GLM based on a canonical basis set 506 including the assumed hemodynamic response function provided by 507 SPM5 (no derivatives). We included two model regressors, one for 508 partial trials and another one for full trials. Both regressors were 509 synchronized with the termination of the S1–S2 interval to be able to 510 capture the following two activity components. In particular, the 511 regressor estimate for partial trials was assumed to capture nogo-type 512 activation and/or termination-related activation, whereas the regres- 513 sor estimate for full trials was assumed to capture termination-related 514 activation and/or S2-related activation. The rationale was that voxels 515 exhibiting stronger activation for partial trials compared to full trials 516 would be involved in processes specific of S2 omission.<sup>3</sup> Notably, to 517 anticipate the results, this estimation procedure did not reveal voxels 518 that exhibited the activation pattern predicted for nogo-type activity. 519

Based on the initial whole-brain activation map, several repre- 520 sentative regions of interest (ROIs) were defined on the group level 521 according to the peak voxel in each anatomically defined region 522 (auditory cortex, inferior occipito-temporal cortex; posterior IPS; 523 PMC; Pre-SMA). For each ROI, the peak voxel was selected for each 524 subject within a radius of 12 mm (i.e., 3 voxels) centered around the 525 peak voxel identified on the group level. For each of these ROIs (at 526 their subject-specific peak voxels) the BOLD estimates R1\_1, R1\_2, and 527 R2 generated by SPM were further examined in a subsequent analysis 528 using the R software package. In particular, BOLD estimates R1\_1 and 529 R1\_2 were analyzed to characterize the temporal profile of S1-related 530 activation. As described in detail in the Simulation section of this 531 paper, four different temporal profiles can be distinguished theore- 532 tically. First, transient S1-related activation is characterized by equal 533 onset latencies and equal peak latencies for R1\_1 compared to R1\_2. 534 Moreover, in this case the area-difference index should not be 535 significantly different from zero. Second, delay-related activation is 536 characterized by a significant area-difference index (ADI). Third, a 537

<sup>&</sup>lt;sup>3</sup> Such a conclusion, however, must be handled with caution as the differences between the two regressor estimates might occur for other reasons. One such reason might be that in full trials the S2 induces a BOLD activation decrease relative to S2 omission in partial trials. Another reason might be overlapping activation elicited earlier during the trial, causing a misfit of the canonical hemodynamic response function that might be more pronounced for full trials than for partial trials due to a disadvantageous S2-related BOLD increase in full trials.

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BOLD activation profile reflecting neural activity associated with 538 539either S1S2-interval termination or S2 omission is characterized by parallel shifts of onset and peak latencies for R1\_2 relative to R1\_1 540541equal to the temporal difference between the two S1-S2 intervals (i.e. 3 s). In this case, the ADI should not be significantly different from 542zero. Fourth, a combination of transient activation associated with 543both, S1 onset and S1S2-interval termination or S2 omission is cha-544racterized by equal onset latencies, but delayed peak latency for R1\_2 545546compared to R1\_1. At the same time, the ADI must not be significantly 547different from zero.

548The ROIs were selected based on a priori anatomical hypotheses. A 549first ROI, primary auditory cortex, was obviously selected to track 550BOLD activation elicited by the auditory stimulus. One group of 551subjects participating in experiment version C (N=5) received brief auditory stimulation at the onset of the visual S1. Thus, we expected 552transient S1-related BOLD activation in auditory cortex time-locked to 553 554 the onset of S1. Another group of subjects participating in experiment versions A and B (N=8) received brief auditory stimulation at the end 555of the S1-S2 interval in both, full and partial trials. Thus, we expected 556 transient delay-termination-related BOLD activation in auditory 557cortex. A second ROI was located in an exemplary region within the 558 ventral visual stream (the most strongly activated in voxel in posterior 559560temporal cortex). This ROI was selected to track BOLD activation 561elicited time-locked to both, S1 onset (the attentional cue) and delayinterval termination (marked by the presentation of the two "O" 562stimuli irrespective of color in experiment versions A and C with 563 N=8). A third set of ROIs was selected to track either transient S1-564565related activation or delay-related activation associated with an assumed top-down attentional signal according to the currently 566 presented color cue. Such activation was expected for all three expe-567riment versions (N = 13). Based on the selective attention literature 568 569(Corbetta and Shulman, 2002; Pessoa et al., 2003; Wager et al., 2004; 570Yantis and Serences, 2003), we examined activation time courses from 571the pre-supplementary motor area (pre-SMA), the dorsal pre-motor cortex (dPMC), and the posterior intra-parietal sulcus region (pIPS). 572

To determine onsets and peaks in real data time courses, a non-573standard analysis strategy needed to be applied. In contrast to the 574 575situation with noise-free synthetic time courses, the presence of high noise levels in real data poses a considerable challenge. Thus, in a first 576 data processing step, the original time course estimates obtained for 577 each subject were re-sampled using the jackknife procedure (Efron, 578 5791981; Maertens and Pollmann, 2005; Ruge et al., 2003). Jackknife resampling generates new time courses by averaging the original data 580 across subjects, but leaving out each subject once. As a consequence, 581 the jackknifed time courses are much smoother than single-subject 582583time courses, but at the same time they fully preserve information 584regarding cross-subject variability. Therefore, parameters like onsets and peaks can be determined much more reliably and jackknife 585statistics can be used for the assessment of statistical significance. A 586 second analysis step was performed specifically for onset detection, 587 which is particularly delicate even for relatively smooth jackknifed 588 589time courses. To determine BOLD onsets, a three-parameter ramp 590function given by Eq. (2) was fitted to the jackknifed time courses.

 $\label{eq:linear} \begin{array}{l} \mbox{If (time<ONSET)then amplitude} = \mbox{INTERCEPT}_{baseline} \\ \mbox{else amplitude} = (\mbox{INTERCEPT}_{baseline} + \mbox{(time} - \mbox{ONSET})) \\ * \mbox{SLOPE}_{rising_{f} lank} \end{array} \tag{2}$ 

59**2** 

The three free parameters were INTERCEPT<sub>baseline</sub>, ONSET, and SLOPE<sub>rising\_flank</sub>. Using the non-linear fitting tool implemented in the R software package, the ramp function given by Eq. (2) was fitted to the jackknifed time course estimates for R1\_1 and R1\_2 within a time range starting 2 time steps before S1 presentation (time point zero) and ending at the respective peak latencies (see Fig. 7 for an example).



**Fig. 7.** Exemplary visualization of the method used for detection of BOLD response onsets. The onset was determined by fitting a 3-parameter (INTERCEPT = baseline activity, ONSET, and SLOPE) linear ramp function to the time course estimates.

In case of two peaks, the first one was chosen as reference point (e.g., 599 Figs. 8B–D). 600

#### Results (empirical data)

Table 1 reports the MNI coordinates for each ROI for each subject.602Fig. 8 depicts for each ROI the estimated time courses for R1\_1, R1\_2,603and R2. Table 2 reports onset latencies, peak latencies and area-604difference indices for each ROI.605

For auditory cortex, the results are clear-cut and conform to the 606 hypotheses. When the sound was presented in synchrony with S1 607 onset, we observed transient S1-related activation (Fig. 8A) as indexed 608 by equal onset latencies and equal peak latencies for R1\_1 and R1\_2. In 609 contrast, when the sound was presented in synchrony with the 610 termination of the S1–S2 interval (Fig. 8B), we observed the expected 611 3 s shift of both, onsets and peaks for R1\_2 as compared to R1\_1. In 612 both cases the area-difference index was, as expected, not significantly 613 different from zero. 614

For the other 4 ROIs, two different types of activation patterns were 615 found. Pre-SMA and dorsal PMC (Figs. 8C and E) showed all signs of 616 delay-related activation, indexed by similar onset latencies for R1\_1 617 and R1\_2, shifted peak latency for R1\_2 relative to R1\_1, and, most 618 importantly, the area-difference index was significantly different from 619 zero. In contrast, the area-difference index was not significantly 620 different from zero for pTEMP and pIPS (Figs. 8D and F), whereas 621 onset latencies and peak latencies exhibited the same pattern as for 622 Pre-SMA and dPMC. This observation suggests that pTEMP and pIPS 623 did not exhibit significant delay-related activation. Instead, the overall 624 temporal profile suggests that these areas exhibit a combination of 625 transient S1-related activation synchronized with S1 onset together 626 with either nogo-type activation or delay-termination-related 627 activation. 628

#### Discussion

The aim of this paper was to evaluate the power and the 630 limitations of an extended version of the standard partial-trial 631 method for separating BOLD components associated with narrowly 632 spaced within-trial events. Our analysis indicates clear advantages of 633 the proposed methodological extension over the standard version of 634 the partial-trial design introduced earlier (Ollinger et al., 2001; 635 Ollinger et al., 2001; Serences, 2004; Shulman et al., 1999). At the 636 same time we also show that some limitations still remain and that 637 the obtained results need to be interpreted with caution.

The first of two major improvements is that it becomes possible to 639 distinguish transient S1-related and delay-related activation. The 640 second improvement is that we can now identify the pattern of BOLD 641 activation that indicates nogo-type activation due to S2 omission in 642 partial S1-only trials. This is an important advantage over the standard 643 partial-trial method which does not provide any means to determine 644 whether time course estimates of S1-related BOLD activation might 645

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Fig. 8. Empirical results for 6 regions of interest. Four different activation patterns could be distinguished. A: transient S1-related activation time-locked to S1 onset; B: transient S1-related activation time-locked to the end of the S1–S2 interval; C and E: delay-related activation during the S1–S2 interval. D and F: combination of transient S1-related activation time-locked to S1 onset plus transient activation time-locked to the end of the S1–S2 interval; C and E: delay-related activation during the S1–S2 interval. D and F: combination of transient S1-related activation time-locked to S1 onset plus transient activation time-locked to the end of the S1–S2 interval.

reflect a methodological artifact due to S2 omission rather than a true
functional component associated with S1 processing occurring in
both, S1-only and full S1–S2 trials.

Furthermore, the method was successfully applied to an empirical 649 650 data set, demonstrating its feasibility under conditions that require the discrimination of temporal BOLD response profiles under realistic, 651 hence noisy conditions. A particularly delicate matter in this respect is 652 653 the extraction of BOLD onsets and the comparison of onset latencies across conditions. Yet, reliable results could be obtained by using 654 655 jackknife re-sampling of time course estimates in combination with fitting a simple linear ramp function to the jackknifed time courses. 656 For a group size of 13 subjects, the 95% confidence interval for onset 657 latencies and differences between onset latencies was below 0.9 s for 658 the examined ROIs. This suggests a sufficiently good sensitivity to 659 660 detect even smaller onset differences that are expected for much smaller differences between short and long S1–S2 intervals (>0.9 s) 661 than realized in the current study (3 s). Obviously, for larger group 662sizes, sensitivity should improve even more. Besides parameterization 663 of onset latencies, the area-difference index, rather than the peak 664 latency difference, turned out to be another highly relevant parameter 665 that is specifically important to determine the amount of delay-666 related activation. Based on this index we could identify significant 667 delay-related activation components for areas like Pre-SMA and 668 dPMC, but not for others like pIPS and posterior temporal cortex 669 which exhibited only transient S1-related activation. 670

A comprehensive power analysis of the extended partial-trial method by itself and in comparison to the standard partial-trial method (i.e., 2 regressors for S1-related activity in the extended design vs. 1 regressor in the standard version) or methods based on a wider range of S1–S2 delay intervals is beyond the scope of this paper. 675 Yet, even without formal analyses, a few relevant factors can be 676 expected to influence statistical power by affecting the number of 677 trials per total experiment time. In the present study it took 678 approximately 25 min to acquire the fMRI data for 192 experimental 679 trials, given delay intervals of 2 s and 5 s and a mean ITI of 2 s 680 (resulting from the randomly interspersed no-event trials). As 681 mentioned above, shortening the S1-S2 delay intervals should be 682 feasible and, if the spared time was invested in increasing the total 683 number of trials, statistical power would likely benefit. Finally it 684 should be noted that the improved estimation power for reconstruct- 685 ing neural activity components underlying S1-related BOLD activation 686 comes at the cost of reduced detection power for the two S1-related 687 model regressors at the two different S1-S2 delay intervals as 688 compared to the standard partial-trial method based on only a single 689 regressor in association with a single delay interval. 690

Despite its merits, the extended partial-trial method still faces 691 remaining limitations. Specifically, the same temporal activation 692 profile indicative of nogo-type activity can also arise due to func- 693 tionally meaningful processes associated with the termination of the 694 S1–S2 interval (cf., Shulman et al., 2002). If such an activation pattern 695 is observed, one might choose to refrain from drawing any con- 696 clusions about the functional role of the affected brain region. A 697 possible solution, though not systematically investigated in the pre- 698 sent paper, might be to use distinct model regressors for partial trials 699 and for full trials synchronized to the time point of S2 omission and 700 S2 presentation, respectively. Voxels exhibiting relatively stronger 701 activation for the partial-trial regressor might be associated with 702 nogo-type activation rather than termination-related activation (see 703

MNI brain coordinates (x,y,z) for different regions of interest for each subject in the three experiment versions (Exp A, Exp B, and Exp C).

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Subject no.	Auditory cortex Sound at S1-onset		Auditory cortexS2 interval termination Sound at S1-S2 interval termination		Post. temp. cortex S2 interval termination Visual stimulus at S1-onset and S1-S2 interval termination		Pre-SMA		Left pIPS			Left dPMC					
	Exp A	Exp B Exp C	Exp A	Exp B	Exp C	Exp A	Exp B	Exp C	Exp A	Exp B	Exp C	Exp A	Exp B	Exp C	Exp A	Exp B	Exp C
01	-		-60 - 28			-40-88			0 12 56			-24-52 52			-36-8		
02	-		-56 - 24			-28 -92			-84			-32 - 48			-48 - 8		
03	-		-48 - 16			o -44-84			4 4 56			-28 - 52			-28 - 4		
04		-	δ	-52-288		4	-			-40		60	-36 - 48		52	-40 - 4	
05		-		-48 -24 4			-						-28 - 48			52 36 0 52	
06		-		-56 - 20 0			-			-40 56			-28 - 48 52			-36-8 48	
07		-		-60 - 16 - 4			-			4060			- 32 - 52 52			-40 - 4 44	
08		-		-52-240			-			4 - 4 60							
		-32-5640			-32-12 52												
09		-56-160			-			-44 -72 4			4064			-28-52 44			-28-844
10		-48 -28 0			-			-32 - 88 -4			4 4 64			-32 - 44			-28-860
11		-48 -20 8			-			-36 - 84			-4464			-28-52			-28-452
12		-60 - 20 0			-			-36 -88 8			- 12 12 52			-24 - 64			-24056
13		-52 - 16 -4			-			-32-844			-8060			-36-60			-28 - 12
Mean		-53 -20 1		-54 - 233			-36-8	35 3		-235	9		-30 - 52 51	1		-33-651	1

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### 12

Table 2

t2.1

Mean values  $\pm$  95% confidence interval.

	Onset latency			Peak latency	Area-difference index		
	R1_1	R1_2	Difference	R1_1	R1_2	Difference	R1_2 - R1_1
Auditory cortex (sound at S1-onset)	$1.41\pm0.65$	$1.17\pm0.36$	$0.24\pm0.49$	$4.11 \pm 1.03$	$3.95\pm0.62$	$-0.16 \pm 0.69$	$20.10\pm74.20$
Auditory cortex (sound at	$4.14\pm0.90$	$7.41 \pm 0.82$	$3.27\pm0.46$	$7.33 \pm 0.36$	$10.57\pm0.21$	$3.24 \pm 0.24$	$-4.37 \pm 17.78$
S1S2-interval termination)							
Left post. temp. cort.	$1.61\pm0.11$	$1.35\pm0.68$	$-0.26 \pm 0.72$	$6.96 \pm 0.15$	$10.73\pm0.14$	$3.77\pm0.17$	$11.13\pm19.18$
Left pIPS	$1.22\pm0.57$	$1.23\pm0.37$	$-0.01\pm0.56$	$6.74 \pm 0.28$	$10.56\pm0.20$	$3.82\pm0.33$	$11.72\pm19.25$
Pre-SMA	$1.23\pm0.56$	$1.34 \pm 0.46$	$-0.11 \pm 0.63$	$6.59 \pm 0.31$	$10.32\pm0.20$	$3.72\pm0.31$	$26.00 \pm 10.03$
Left dPMC	$1.01\pm0.64$	$0.71\pm0.80$	$-0.30 \pm 0.87$	$6.39 \pm 0.48$	$10.18\pm0.29$	$\textbf{3.80} \pm \textbf{0.32}$	$\textbf{27.19} \pm \textbf{12.61}$

R1 1: BOLD response estimate for S1-related activation at 2 s cue-target interval.

t2 12 R1\_2: BOLD response estimate for S1-related activation at 5 s cue-target interval.

Empirical methods, for further details). Alternatively, one might 704 choose to clarify ambiguous activation patterns through experi-705 mental means (cf., Goghari and MacDonald, 2008). Specifically, it 706 seems worthwhile to consider experimental manipulations that 707 selectively affect either nogo-type or termination-related processes. 708 For instance, manipulating the proportion of S1-only trials might 709 be a good way to influence the strength of nogo-type neural res-710 ponses, which should be stronger for less frequent S1-only trials. 711 712 Following a similar rationale, one might be willing to accept the 713argument that a design with a high proportion of S1-only trials (e.g. 33%) would make nogo-type responses unlikely to occur at all. 714 In fact, in the present empirical study, for which the partial-trial 715 proportion was 33%, we could identify only a single brain area 716 717 (auditory cortex) that exhibited the temporal activation profile indicative of purely nogo-related activation.<sup>4</sup> Yet, this activation 718 719 pattern could clearly be attributed to the auditory stimulation that 720 marked the termination of the S1-S2 interval. The situation was less clear-cut for areas like pIPS and posterior temporal cortex 721 722 which exhibited a combination of transient S1-related activation and either nogo-related or delay-termination-related activation. 723 While the presence of the transient S1-related activity component 724 can be inferred without doubt, it cannot be decided if the second 725 overlapping activity component was due to S2 omission or delay-726 interval termination. 727

To conclude, we believe that the remaining interpretative 728ambiguities should not be turned into an argument against the 729 proposed extended partial-trial design, given that it provides 730 731 important improvements with respect to existing approaches. Importantly though, these limitations should always be kept in 732 mind to avoid erroneous interpretation. Put into a more general 733 perspective, the extended partial-trial approach is arguably the 734 better choice in comparison to the full-trial method with variable 735 736 but narrowly distributed event spacing. As pointed out earlier (Serences, 2004), such an approach yields distorted time course 737 estimates when delay-related activation is present - without being 738 able to tell from the observed data whether or not that might be the 739 case. The other alternative would be to use a design with a wider 740 741 distribution of events spacing and explicit modeling of a delayrelated BOLD component. While such a design might be a good 742 choice for certain questions (e.g., manipulation within working 743memory), it does not seem to be well suited for other experimental 744paradigms in which long event spacing might introduce intervening 745 processes that can potentially mask neural activity associated with 746 the processes of genuine interest. 747

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Acknowledgments		751

We wish to thank Kerstin Raum for her assistance in recruiting the 752 participants and collecting the data. The research was supported by 753 RO1 GRANT MH066078. 754

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Please cite this article as: Ruge, H., et al., Separating event-related BOLD components within trials: The partial-trial design revisited, NeuroImage (2009), doi:10.1016/j.neuroimage.2009.04.075

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<sup>&</sup>lt;sup>4</sup> It should be noted that we implemented this study with an unusually high proportion of 33% partial trials as compared to previous studies based on the standard partial trial design that included smaller proportions ranging from 20 to 25%. As designs with smaller proportion of partial trials make S2 omission more surprising, it might well be that a nogo-type response is elicited under such conditions.

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