

1 **Title:** Selective processing of auditory evoked responses with iterative-  
2 randomized stimulation and averaging: A strategy for evaluating the time-  
3 invariant assumption.

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21 **Abstract:**

22 The recording of auditory evoked potentials (AEPs) at fast rates allows the study  
23 of neural adaptation, improves accuracy in estimating hearing threshold and may  
24 help diagnosing certain pathologies. Stimulation sequences used to record AEPs  
25 at fast rates require to be designed with a certain jitter, i.e., not periodical. Some  
26 authors believe that stimuli from wide-jittered sequences may evoke auditory  
27 responses of different morphology, and therefore, the time-invariant assumption  
28 would not be accomplished. This paper describes a methodology that can be  
29 used to analyze the time-invariant assumption in jittered stimulation sequences.  
30 The proposed method [Split-IRSA] is based on an extended version of the  
31 iterative randomized stimulation and averaging (IRSA) technique, including  
32 selective processing of sweeps according to a predefined criterion. The  
33 fundamentals, the mathematical basis and relevant implementation guidelines of  
34 this technique are presented in this paper. The results of this study show that  
35 Split-IRSA presents an adequate performance and that both fast and slow  
36 mechanisms of adaptation influence the evoked-response morphology, thus both  
37 mechanisms should be considered when time-invariance is assumed. The  
38 significance of these findings is discussed.

39 **Keywords:** randomized stimulation and averaging (RSA), jitter, deconvolution,  
40 evoked potentials, time-invariant, ABR, MLR, SOA.

41

42 **Highlights:**

43       ▪ Split-IRSA disentangles overlapping evoked potentials of different  
44       morphology.

45       ▪ Split-IRSA allows analysis of time-invariant assumption in jittered stimuli.

46       ▪ Both fast and slow mechanisms of adaptation influence the time-invariant  
47       assumption.

48

49 **Text body:**

50

## 1. INTRODUCTION

51 The conventional auditory evoked potential (AEP) recording method consists in  
52 the periodical presentation of stimuli and the average of their associated auditory  
53 neural responses (sweeps) in order to increase the signal-to-noise ratio (SNR)  
54 (Thornton, 2007). The conventional method presents the limitation that the period  
55 of stimulation (i.e., the inverse of the stimulation rate) must be greater than the  
56 averaging window, avoiding sweeps to be overlapped (Wong and Bickford,  
57 1980); otherwise it would not be mathematically possible to recover the transient  
58 evoked response (Kjaer, 1980). This rate limitation implies that auditory  
59 brainstem responses (ABR) and middle latency responses (MLR) cannot be  
60 recorded with the conventional technique at rates faster than 100 Hz and 10 Hz,  
61 respectively, considering standard averaging windows of 10 ms in ABR and  
62 100 ms in MLR signals. However, the recording of these signals at higher rates  
63 present several advantages, such as the study of neural adaptation (Burkard et  
64 al., 1990; Lasky, 1997), the diagnosis of certain pathologies (Jiang et al., 2000;  
65 Yagi and Kaga, 1979) and better performance in hearing threshold estimation  
66 (Leung et al., 1998).

67 The maximum length sequence (MLS) technique was developed by Eysholdt and  
68 Schreiner (1982) to overcome the rate limitation imposed by the conventional  
69 technique. This technique was extensively used not only to record AEPs at fast  
70 stimulation rates, when the responses are overlapped (Burkard and Palmer,  
71 1997; Eggermont, 1993; Lasky et al., 1995), but also to analyze the linear and  
72 non-linear interaction components of otoacoustic emissions (de Boer et al., 2007;

73 Hine et al., 1997; Hine et al. 2001; Lineton et al., 2006). Stimulus-onset  
74 asynchrony (SOA) , i.e. the distribution of time intervals between adjacent stimuli,  
75 are multiples of a minimum pulse interval in MLS sequences, which leads to  
76 stimulation sequences of a large jitter (Burkard et al., 1990; Özdamar et al, 2007).  
77 The jitter of a stimulation sequence determines dispersion of the SOA distribution.

78 Several techniques have emerged to deconvolve overlapped AEPs using narrow-  
79 jittered stimulation sequences. Some of the most relevant techniques are quasi-  
80 periodic sequence deconvolution (QSD) (Jewett et al., 2004), continuous loop  
81 averaging deconvolution (CLAD) (Delgado and Özdamar, 2004; Özdamar and  
82 Bohórquez, 2006), and least-squares deconvolution (LSD) (Bardy et al., 2014a).  
83 These techniques have been successfully used in several research applications  
84 (Bardy et al., 2014b; Bohórquez and Özdamar, 2008; Özdamar et al., 2007). The  
85 major limitation of these techniques is that obtaining efficient, narrow-jittered  
86 stimulation sequences may require an extensive search, since they must  
87 accomplish frequency-domain restrictions to avoid noise amplification in the  
88 deconvolution process (Jewett et al., 2004; Özdamar and Bohórquez, 2006).

89 A recently published paper describes iterative randomized stimulation and  
90 averaging (IRSA), which allows AEPs to be recorded at fast rates using narrow-  
91 jittered sequences (Valderrama et al., 2014c). This is achieved by the estimate  
92 and removal of the interference associated with overlapping responses through  
93 iterations in the time-domain, providing better estimates of the response in  
94 succeeding iterations. This technique assumes that the AEP morphology is time-  
95 invariant, i.e., all stimuli evoke the same neural response, which may constrain  
96 the flexibility of this technique in certain applications.

97 Despite the great effort in developing different methodologies to record AEPs at  
98 fast rates using narrow-jittered sequences, it is still controversial whether or not  
99 stimulation sequences of a wide jitter are a problem. Some authors believe that  
100 stimuli in high-jittered sequences may evoke auditory responses of different  
101 morphology as a consequence of the effects of neural adaptation, contradicting  
102 therefore the time-invariant assumption (Jewett et al., 2004, Özdamar and  
103 Bohórquez, 2006; Valderrama et al., 2014b). However, to the best of our  
104 knowledge, we have not found any technique that allows evaluation of the time-  
105 invariant assumption.

106 This paper describes an extended version of IRSA [Split-IRSA] which allows  
107 selective averaging and processing when AEPs of different morphology are  
108 recorded. In this study, the performance of this technique is assessed with both  
109 artificially synthesized and real experiments. The Split-IRSA technique is applied  
110 to evaluate the time-invariant assumption on ABR and MLR signals recorded with  
111 16 ms-jittered stimulation sequences. The results of this study show that (a) the  
112 Split-IRSA technique presents an adequate performance, (b) the time-invariant  
113 assumption in auditory responses recorded on jittered stimulation sequences can  
114 be evaluated following a methodology based on Split-IRSA, and (c) the  
115 morphology of individual sweeps in ABR and MLR signals is influenced by both  
116 fast and slow mechanisms of adaptation. The potential of this method and the  
117 significance of the findings obtained in this study are discussed.

## 2. METHODS

118

119 This section presents the basis and the mathematical formulation of the Split-  
120 IRSA technique, the protocols followed in the recording of real  
121 electroencephalograms (EEGs), and the objectives, hypotheses and procedures  
122 of the experiments.

### 123 2.1. Split-IRSA

124 The fundamentals for the Split-IRSA algorithm are very similar to those of IRSA,  
125 described in detail in Valderrama et al. (2014c). AEPs are estimated in Split-IRSA  
126 through an iterative process in the time domain. Each iteration includes  
127 estimation of the interference associated with overlapping responses, subtraction  
128 of this interference from the recorded EEG, and re-estimation of the AEPs. Better  
129 AEPs estimates can be obtained recursively since improved AEPs estimates lead  
130 to a better interference estimate, which leads to more accurate AEPs estimates.  
131 The precision of the AEPs estimates increases with the number of iterations. In  
132 contrast to IRSA, this updated formulation [Split-IRSA] allows selective  
133 processing of sweeps, and therefore, AEPs of different morphology can be  
134 separately estimated.

135 Stimulation sequences are generated in Split-IRSA as the combination of  
136 independent sub-sequences, each of them based on randomized stimulation, in  
137 which the SOA of the stimuli vary randomly according to a predefined probability  
138 distribution (Valderrama et al., 2012). The Split-IRSA technique retrieves the  
139 time-invariant component of the AEPs belonging to each sub-sequence, i.e., it is  
140 assumed that all stimuli from the same sub-sequence evoke the same AEP.

141 The mathematical formulation for the Split-IRSA technique is outlined below. Let  
 142  $[\mathbf{s}_1(n), \mathbf{s}_2(n), \dots, \mathbf{s}_T(n)]$  ( $n = 1, \dots, N$ ) be  $T$  sub-sequences, each of them  
 143 composed of  $[K_1, K_2, \dots, K_T]$  stimuli that evoke, respectively,  $T$  AEPs of different  
 144 morphology, represented by  $[\mathbf{x}_1(j), \mathbf{x}_2(j), \dots, \mathbf{x}_T(j)]$  ( $j = 1, \dots, J$ ), where  $N$  and  $J$   
 145 represent, respectively, the length in samples of the EEG and the averaging  
 146 window. The recorded EEG  $\mathbf{y}(n)$ , can be modeled as the summation of the  
 147 convolutions (\*) of each sub-sequence with their corresponding AEP plus noise:  
 148  $\mathbf{y}(n) = \mathbf{s}_1(n) * \mathbf{x}_1 + \mathbf{s}_2(n) * \mathbf{x}_2 + \dots + \mathbf{s}_T(n) * \mathbf{x}_T + noise.$  (1)

149 The AEPs corresponding to each sub-sequence ( $\tau = 1, \dots, T$ ) in the iteration  $i$ ,  
 150  $\hat{\mathbf{x}}_{\tau,i}(j = 1, \dots, J)$ , are estimated in Split-IRSA according to

$$151 \hat{\mathbf{x}}_{\tau,i}(j) = \frac{1}{K_\tau} \cdot \sum_{k=1}^{K_\tau} \mathbf{y}_{\tau,k}(j + \mathbf{m}_\tau(k)), \quad (2)$$

152 where  $\mathbf{y}_{\tau,k}$  represents the EEG in which the auditory responses adjacent to the  
 153 stimulus  $k$  (from the sub-sequence  $\tau$ ) are suppressed; and  $\mathbf{m}_\tau$  is a trigger vector  
 154 that includes the samples of the EEG in which the stimuli of the sub-sequence  $\tau$   
 155 occur ( $k = 1, \dots, K_\tau$ ). The  $\mathbf{y}_{\tau,k}$  signals can be obtained for each stimulus  $k$  at each  
 156 sub-sequence  $\tau$  by suppressing from the recorded EEG the AEPs estimated on  
 157 the preceding iteration ( $i - 1$ ) corresponding to all sub-sequences ( $t = 1, \dots, T$ )  
 158 and by adding the AEP corresponding to the stimulus  $k$  of the sub-sequence  $\tau$ :

$$159 \mathbf{y}_{\tau,k}(n) = \mathbf{y}(n) - \sum_{t=1}^T [\mathbf{s}_t(n) * \hat{\mathbf{x}}_{t,i-1}] + \mathbf{s}_{\tau,k}(n) * \hat{\mathbf{x}}_{\tau,i-1}, \quad (3)$$

160 where  $\mathbf{s}_{\tau,k}$  represents the stimulation sequence for the stimulus  $k$  of the sub-  
 161 sequence  $\tau$ . Considering  $\mathbf{z}_i(n)$  as the EEG on the iteration  $i$  with all AEPs  
 162 estimated on the preceding iteration suppressed:  $\mathbf{z}_i(n) = \mathbf{y}(n) - \sum_{t=1}^T [\mathbf{s}_t(n) * \hat{\mathbf{x}}_{t,i-1}]$ , then equation (3) can be rewritten as



164  $\mathbf{y}_{\tau,k}(n) = \mathbf{z}_i(n) + \mathbf{s}_{\tau,k}(n) * \hat{\mathbf{x}}_{\tau,i-1}.$  (4)

165 Hence, the sections of  $\mathbf{y}_{\tau,k}$  corresponding to the averaging window can be  
 166 obtained as

167  $\mathbf{y}_{\tau,k}(j + \mathbf{m}_{\tau}(k)) = \mathbf{z}_i(j + \mathbf{m}_{\tau}(k)) + \mathbf{s}_{\tau,k}(j + \mathbf{m}_{\tau}(k)) * \hat{\mathbf{x}}_{\tau,i-1}.$  (5)

168 The  $\mathbf{s}_{\tau,k}(n)$  signal can be expressed as  $\delta(n - \mathbf{m}_{\tau}(k))$ , where  $\delta(n)$  represents the  
 169 Dirac delta function, with the value 1 for  $n = 0$ , and 0 otherwise. Since  $\delta(n) * \mathbf{f} =$   
 170  $\mathbf{f}$ , for whatever function  $\mathbf{f}$ , equation (5) can be expressed as

171  $\mathbf{y}_{\tau,k}(j + \mathbf{m}_{\tau}(k)) = \mathbf{z}_i(j + \mathbf{m}_{\tau}(k)) + \delta(n - \mathbf{m}_{\tau}(k) + \mathbf{m}_{\tau}(k)) * \hat{\mathbf{x}}_{\tau,i-1} =$   
 172  $\mathbf{z}_i(j + \mathbf{m}_{\tau}(k)) + \hat{\mathbf{x}}_{\tau,i-1}.$  (6)

173 Therefore, from equation (2), the AEP estimate on the iteration  $i$  can be obtained  
 174 as

175  $\hat{\mathbf{x}}_{\tau,i}(j) = \frac{1}{K_{\tau}} \cdot \sum_{k=1}^{K_{\tau}} [\mathbf{z}_i(j + \mathbf{m}_{\tau}(k)) + \hat{\mathbf{x}}_{\tau,i-1}] = \hat{\mathbf{x}}_{\tau,i-1} + \frac{1}{K_{\tau}} \cdot \sum_{k=1}^{K_{\tau}} \mathbf{z}_i(j + \mathbf{m}_{\tau}(k)).$  (7)

176 Similar to IRSA, we have found in simulations and real data that Split-IRSA might  
 177 present problems of instability, where succeeding iterations lead to worse AEP  
 178 estimates. Instability might be especially relevant in low-jittered stimulation  
 179 sequences in which the averaged SOA is significantly lower than the averaging  
 180 window, e.g., with a high-degree of overlap. Problems of instability can be solved  
 181 using a correction factor ( $\alpha$ ) that weights the correction  $\frac{1}{K_{\tau}} \cdot \sum_{k=1}^{K_{\tau}} \mathbf{z}_i(j + \mathbf{m}_{\tau}(k))$   
 182 made on the preceding AEP estimate. Low  $\alpha$  values ensure convergence, but  
 183 require a greater number of iterations to converge. The greatest  $\alpha$  that avoids  
 184 instability is optimal. Thus, the inclusion of this correction factor onto equation (7)  
 185 leads to:

186  $\hat{\mathbf{x}}_{\tau,i}(j) = \hat{\mathbf{x}}_{\tau,i-1} + \alpha \cdot \frac{1}{K_{\tau}} \cdot \sum_{k=1}^{K_{\tau}} \mathbf{z}_i(j + \mathbf{m}_{\tau}(k)).$  (8)

187 The number of iterations can be defined either as a fixed value  $I$  ( $\hat{x}_\tau = \hat{x}_{\tau,I} \forall \tau$ ) or  
188 automatically considering whether the differences between AEP estimates in  
189 succeeding iterations are negligible ( $\hat{x}_\tau = \hat{x}_{\tau,i} \Leftrightarrow \hat{x}_{\tau,i} \approx \hat{x}_{\tau,i-1} \forall \tau$ ).

190 Figure 1 illustrates an example of the performance of the Split-IRSA technique  
191 under a simulation framework. In this example, a stimulation sequence  $s(n)$  was  
192 generated containing 4000 stimuli in which the SOA varied randomly between 20  
193 and 30 ms [short SOA sub-sequence:  $s_1(n)$ ] and between 60 to 70 ms [long SOA  
194 sub-sequence:  $s_2(n)$ ]. Figure 1.A shows the histogram of the SOA of this  
195 stimulation sequence, where the sub-sequences  $s_1(n)$  and  $s_2(n)$  can be  
196 identified. Figure 1.B shows the configuration settings of this simulation  
197 experiment. Figure 1.B.1 shows a frame of the first 20.000 samples of  $s(n)$ , using  
198 a sampling frequency of 25 kHz. In this segment, long- and short-SOA stimuli can  
199 be visually identified. Figures 1.B.2 and 1.B.3 show, respectively, the triggers  
200 corresponding to each sub-sequence. In these sub-sequences, the first three  
201 elements of the trigger vectors [ $m_1$  and  $m_2$ ] are labeled as a reference. An  
202 artificially synthesized EEG was generated as the summation of the convolutions  
203 of the sub-sequences  $s_1(n)$  and  $s_2(n)$  with two high-quality real MLR signals of  
204 different morphology:  $x_1$  and  $x_2$ . The  $x_1$  and  $x_2$  signals are shown next to the first  
205 stimulus in each sub-sequence. These signals were recorded from two normal  
206 hearing subjects (males, 28 and 26 yr, respectively) using 4800 stimuli presented  
207 at 70 dB HL at an average rate of 40 Hz and processed by the IRSA technique.  
208 The artificially synthesized EEG [ $y(n)$ ], along with the triggers corresponding to  
209 both sub-sequences, are shown in figure 1.B.4. In this experiment, passband-  
210 filtered noise (Butterworth, 4<sup>th</sup> order, [30-200] Hz) was added to  $y(n)$  at a SNR of

211 -5 dB (figure 1.B.5). Figure 1.C presents the normalized energy of the averaged  
212 residual, evaluated as  $\frac{1}{N} \sum_{n=1}^N z_i(n)^2$ , at different number of iterations for different  
213  $\alpha$  values. This figure shows that the  $\alpha$  parameter can be used to control  
214 convergence and avoid instability. In this example,  $\alpha$  values 1.3 and 1.0 cause  
215 instability, where the averaged residual increases in succeeding iterations. In  
216 contrast, the averaged residual for  $\alpha$  values 0.8 and 0.1 decreases with the  
217 number of iterations, which means that better estimates of the responses are  
218 obtained recursively. This figure shows that although both  $\alpha$  values 0.8 and 0.1  
219 tend to converge, the convergence for  $\alpha$  value 0.1 requires a larger number of  
220 iterations, i.e., it is less efficient. This simulation shows that  $\alpha$  equal to 0.8 and 5  
221 iterations are appropriate values to obtain accurate estimates of the signals  $x_1$   
222 and  $x_2$ . Figures 1.D.1 and 1.D.2 show, respectively, the AEP estimates for  $x_1$  and  
223  $x_2$  at the second, fifth and tenth iteration for  $\alpha$ -value of 1.3. These figures show  
224 an example of instability, where worse estimates of the responses are obtained  
225 in succeeding iterations, i.e., the root-mean-square (RMS) error between the  
226 template and the MLR estimate increases in succeeding iterations. Figures 1.E.1  
227 and 1.E.2 show, respectively, the first three estimates of the signals  $x_1$  and  $x_2$  for  
228 an  $\alpha$  value 0.8. In this example, when the  $\alpha$  value is selected appropriately, better  
229 estimates are obtained recursively, i.e., the RMS error decreases with increasing  
230 iterations [convergence scenario].

231 A software routine programmed in MATLAB (The Mathworks, Inc., Natick, MA)  
232 that implements the Split-IRSA technique is available as supporting information  
233 in this paper (Appendix A).

## 234 **2.2. EEG recording and processing**

235 The EEG recording process consisted in the presentation of stimuli to a subject  
236 and the recording of their associated neural response through surface disposable  
237 electrodes (Ambu Neuroline 720, Ambu A/S, Denmark) placed on the skin at  
238 different positions on the head. The positive electrode was placed at the high-  
239 forehead, the negative electrode at the ipsilateral mastoid and the reference  
240 electrode at the low-forehead. The interelectrode impedance was below 5 k $\Omega$  in  
241 all recordings. Stimuli consisted of 100  $\mu$ s-duration, monophasic clicks delivered  
242 in rarefaction polarity at 70 dB HL (corresponding to 103.54 dB peak-to-peak  
243 equivalent sound pressure level) through the Etymotic ER-3A insert earphones  
244 (Etymotic Research, Inc., Elk Grove Village, IL). Calibration was carried out  
245 according to the ISO-389 standard, using an Artificial Ear type 4153 2-cc acoustic  
246 coupler (Brüel & Kjær Sound & Vibration Measurements A/S, Nærum, Denmark).  
247 The recording sessions took place in the MRC Institute of Hearing Research  
248 (Royal South Hants Hospital, Southampton, United Kingdom), in a sound-  
249 shielded screening booth prepared to attenuate electrical and electromagnetic  
250 interference. Subjects were comfortably seated in order to minimize  
251 electromyogenic noise. The signal recorded by the electrodes was 86 dB  
252 amplified (gain x20.000) and bandpass filtered by a 24 dB/Octave slope filter with  
253 a bandwidth of [0.5-3500] Hz. The amplified EEG was sampled at 25 kHz and  
254 quantized with a resolution of 16 bits. Digitized EEGs were digitally filtered by a  
255 4<sup>th</sup> order Butterworth filter ([200-2000] Hz for ABR and [30-1500] for MLR). Group  
256 delays introduced by the insert earphones (0.81 ms) (Eiberling et al., 2012) and  
257 by both analog and digital filters were digitally compensated. Data processing

258 was carried out by custom-designed scripts implemented in MATLAB. The  
259 features of the AEP recording system used in this study are presented in  
260 Valderrama et al. (2014a).

261 Analysis of AEPs consisted in the measurement of their most relevant  
262 components in terms of latencies and amplitudes. Latencies were measured as  
263 the time difference in milliseconds from stimulus onset to the occurrence of the  
264 components. Amplitudes were estimated in ABR as the difference in microvolts  
265 between the top of the peak and the following trough, whereas in MLR,  
266 amplitudes were measured as the difference between the positive and negative  
267 peaks of the wave complex (Burkard and Don, 2007).

268 The recording protocols followed in the experiments of this work were in  
269 accordance with the Code of Ethics of the World Medical Association (Declaration  
270 of Helsinki) for experiments involving humans, and were approved by the  
271 Research Ethics Committee established by the Health Research Authority  
272 (Reference No. RHM ENT0082).

## 273 **2.3. Description of the experiments**

### 274 2.3.1. Rationale

275 Three experiments were carried out with the double purpose of evaluating the  
276 performance of the Split-IRSA technique and the validity of the time-invariant  
277 assumption in the recording of ABR and MLR signals with 16 ms-jittered  
278 randomized stimulation sequences.

### 279 2.3.2. Subjects

280 All subjects tested on the experiments of this study were volunteers, reported no  
281 history of auditory dysfunction and presented normal hearing sensitivity at octave  
282 frequencies ([250-8000] Hz). These subjects were paid and gave written consent  
283 to participate.

### 284 2.3.3. Experiment 1

285 The first experiment compares ABR and MLR real signals recorded on one  
286 subject (male, 30 yr) at different rates in two scenarios.

287 In scenario 1, ABR signals were recorded at 16 different rates using 1 ms-jittered  
288 sequences:  $SOA_{15-16}$  (65 Hz),  $SOA_{14-15}$  (69 Hz),  $SOA_{13-14}$  (74 Hz),  $SOA_{12-13}$  (80  
289 Hz),  $SOA_{11-12}$  (87 Hz),  $SOA_{10-11}$  (95 Hz),  $SOA_{9-10}$  (105 Hz),  $SOA_{8-9}$  (118 Hz),  
290  $SOA_{7-8}$  (133 Hz),  $SOA_{6-7}$  (154 Hz),  $SOA_{5-6}$  (182 Hz),  $SOA_{4-5}$  (222 Hz),  $SOA_{3-4}$   
291 (286 Hz),  $SOA_{2-3}$  (400 Hz),  $SOA_{1-2}$  (667 Hz),  $SOA_{0-1}$  (2000 Hz); and MLR signals  
292 were recorded at 4 different rates using 4 ms-jittered sequences:  $SOA_{12-16}$  (71  
293 Hz),  $SOA_{8-12}$  (100 Hz),  $SOA_{4-8}$  (167 Hz) and  $SOA_{0-4}$  (500 Hz). A large number of  
294 stimuli were used in each stimulation sequence in order to obtain signals of  
295 sufficient quality. In ABR signals, sequences  $SOA_{15-16}$  to  $SOA_{9-10}$  included 12,500  
296 stimuli, while sequences  $SOA_{8-9}$  to  $SOA_{0-1}$  contained 20,000 stimuli. The larger  
297 number of stimuli in higher-rate sequences was used to accommodate the loss  
298 of SNR due to the reduction of amplitude of the components as a consequence  
299 of adaptation (Hine et al., 2001). In MLR signals, all sequences contained 50.000  
300 stimuli. ABR and MLR signals on this scenario were processed by the IRSA  
301 technique (Valderrama et al., 2014c). The number of iterations for ABR and MLR

302 signals were, respectively 50 and 500. The value of  $\alpha$  was 0.8 at all rates for ABR  
303 signals, except for the sequences  $SOA_{5-6}$ ,  $SOA_{4-5}$ ,  $SOA_{3-4}$  and  $SOA_{2-3}$ , where  $\alpha$   
304 was 0.5. In MLR signals, the  $\alpha$ -value for  $SOA_{12-16}$  and  $SOA_{8-12}$  was 0.3; for  $SOA_{4-}$   
305  $8$ ,  $\alpha$  was 0.5; and for  $SOA_{0-4}$ ,  $\alpha$  was 0.8. We tested in simulations that these  
306 parameters were appropriate to obtain accurate ABR and MLR estimates.

307 In scenario 2, ABR and MLR signals were estimated on the same subject and at  
308 the same stimulation rates as for scenario 1 from a single EEG corresponding to  
309 a stimulation sequence  $SOA_{0-16}$  (jitter of 16 ms) of 200,000 stimuli. In ABR, each  
310 stimulus was categorized in 1 ms-jittered sub-sequences according to their  
311 preceding stimulus:  $s_1$  ( $SOA_{0-1}$ : preceding SOA belongs to the interval [0-1]),  
312  $s_2$  ( $SOA_{1-2}$ ),  $s_3$  ( $SOA_{2-3}$ ), ...,  $s_{16}$  ( $SOA_{15-16}$ ). Equally, the processing of MLR  
313 signals included the categorization of the stimuli according to the intervals:  $s_1$   
314 ( $SOA_{0-4}$ : preceding SOA belongs to the interval [0-4]),  $s_2$  ( $SOA_{4-8}$ ),  $s_3$  ( $SOA_{8-12}$ )  
315 and  $s_4$  ( $SOA_{12-16}$ ). Since randomized stimulation sequences used in this  
316 experiment were distributed according to uniform distributions, the number of  
317 stimuli that belonged to each sub-sequence was approximately 12,500 in ABR  
318 signals (200,000/16), and 50,000 stimuli in MLR signals (200,000/4). ABR and  
319 MLR signals were processed with Split-IRSA, as described in section 2.1 of this  
320 paper. The number of iterations ( $I$ ) and the  $\alpha$ -value were, respectively,  $I = 50$   
321 and  $\alpha = 0.8$  in ABR; and  $I = 500$  and  $\alpha = 0.8$  in MLR. Experiments in simulations  
322 validated the value of these parameters.

323 The morphology of the ABR and MLR signals obtained in both described  
324 scenarios was compared in terms of amplitudes and latencies. The morphology  
325 of the auditory responses obtained at different rates on the two scenarios is

326 expected to be influenced by both fast and slow mechanisms of adaptation. On  
327 the one hand, the morphology of ABR and MLR signals obtained on scenario 1  
328 is expected to be in accordance with several previous studies in which ABR and  
329 MLR signals are recorded at fast rates (Lasky, 1997; Özdamar et al., 2007; Yagi  
330 and Kaga, 1979). On the other hand, there is not sufficient literature to  
331 hypothesize the ABR and MLR waveforms on scenario 2. If fast mechanisms of  
332 adaptation (with a time-constant of a few milliseconds) prevail over slow  
333 mechanisms (with a time-constant of several tens of milliseconds), the  
334 morphology of the AEPs in scenario 2 will be similar to those in scenario 1, since  
335 the morphology of the responses would be strongly influenced by the preceding  
336 SOA. In contrast, if slow mechanisms of adaptation prevail over fast mechanisms,  
337 then the AEPs corresponding to different sub-sequences would be similar, since  
338 the morphology of the response to each stimulus would not be very much  
339 influenced by its preceding SOA, but by the averaged SOA of several  
340 milliseconds in advanced.

#### 341 2.3.4. Experiment 2

342 The objective of experiment 2 is to analyze the performance of the Split-IRSA  
343 technique in order to validate the experimental results obtained in experiment 1.  
344 This analysis was carried out through a simulation, in which the acquisition  
345 settings of experiment 1 were reproduced. This study was performed for ABR  
346 and MLR signals, both with and without added noise.

347 First, a  $SOA_{0-16}$  randomized stimulation sequence of 200.000 stimuli was  
348 generated. Each stimulus from this sequence was categorized into sub-



349 sequences as described in scenario 2 in experiment 1, i.e., in the study with ABR  
350 signals there were 16 sub-sequences of 1 ms jitter:  $s_1$  (SOA<sub>0-1</sub>),  $s_2$  (SOA<sub>1-2</sub>), ...,  
351  $s_{16}$  (SOA<sub>15-16</sub>); and in the study with MLR signals, there were 4 sub-sequences  
352 of 4 ms jitter:  $s_1$  (SOA<sub>0-4</sub>),  $s_2$  (SOA<sub>4-8</sub>), ...,  $s_4$  (SOA<sub>12-16</sub>). Second, two artificially  
353 synthesized EEGs (one for each scenario) were built as the convolution of the  
354 stimuli belonging to each sub-sequence with the corresponding ABR/MLR signals  
355 obtained in experiment 1 on scenarios 1 and 2. These artificially synthesized  
356 EEGs represent the overlapping evoked potentials without any type of noise or  
357 artifacts. Finally, the ABR/MLR signals corresponding to each sub-sequence  
358 were estimated from these synthesized EEGs using the Split-IRSA technique at  
359 the iterations  $I = [0, 10, 20, 50]$  in ABR, and  $I = [0, 10, 20, 50, 100, 200, 500]$  in  
360 MLR. The  $\alpha$ -value used in these simulations was the same as in experiment 1,  
361 i.e.,  $\alpha = 0.8$  in both ABR and MLR signals. The error between the original  
362 ABR/MLR signals (templates) and the estimated signals was calculated in terms  
363 of RMS value.

364 The same study was repeated including filtered noise (4<sup>th</sup> order Butterworth, [200-  
365 2000] Hz for ABR and [30-1500] for MLR) added to the synthesized EEGs at a  
366 RMS value similar to the recorded real EEG. This RMS value was estimated on  
367 the recorded EEG after digital filtering (4<sup>th</sup> order Butterworth, [200-2000] Hz for  
368 ABR and [30-1500] for MLR). The estimated RMS values were 1.7  $\mu$ V for ABR  
369 and 3.5  $\mu$ V for MLR. In ABR signals, the SNRs on the noisy EEGs were -29.2 dB  
370 in scenario 1 and -30.2 dB in scenario 2. In MLR, the SNR-values were -17.8 dB  
371 in scenario 1 and -23.4 dB in scenario 2.

### 372 2.3.5. Experiment 3

373 In this experiment, we analyzed the morphology of ABR and MLR signals evoked  
374 by stimuli that belong to different rate-subsets from stimulation sequences of  
375 16 ms-jitter in order to evaluate the time-invariant assumption.

376 8 subjects (5 males,  $27 \pm 4$  yr) participated in this study. Each subject was  
377 presented a randomized stimulation sequence  $SOA_{0-16}$  of 60.000 stimuli. A single  
378 EEG was recorded from each subject. These EEGs were digitally filtered (4<sup>th</sup>  
379 order Butterworth) using a bandwidth [200-2000] Hz for the ABR analysis and  
380 [30-1500] Hz for MLR. Sub-sequences were defined as described in scenario 2  
381 on experiment 1 of this paper:  $s_1$  ( $SOA_{0-1}$ ),  $s_2$  ( $SOA_{1-2}$ ), ...,  $s_{16}$  ( $SOA_{15-16}$ ) in ABR;  
382 and  $s_1$  ( $SOA_{0-4}$ ),  $s_2$  ( $SOA_{4-8}$ ), ...,  $s_4$  ( $SOA_{12-16}$ ) in MLR. ABR and MLR signals  
383 were estimated from each rate-subset using the Split-IRSA technique, as  
384 described in section 2.1 of this paper, using  $\alpha = 0.8$ ,  $I = 50$  in ABR and  $I = 500$   
385 in MLR. In addition, we used as reference the ABR/MLR signal obtained from the  
386 complete stimulation sequence, assuming that all stimuli from the sequence  
387 evoked the same response. These signals were obtained using the IRSA  
388 technique ( $\alpha = 0.8$ ,  $I = 50$  in ABR and  $I = 500$  in MLR) (Valderrama et al.,  
389 2014c).

390 The latencies and amplitudes of waves III and V were measured on ABR signals.  
391 In MLR, we measured the latencies for the Na, Pa, Nb and Pb components and  
392 the amplitudes for the Na-Pa, Pa-Nb and Nb-Pb wave-complexes. The influence  
393 of the average rate in each sub-sequence on the morphology of ABR/MLR signals  
394 was evaluated through linear correlation hypothesis tests, considering the slope  
395 equal to zero as the null hypothesis of the tests.

396 The inter-subject variability of the fast adaptation was analyzed in each subject  
397 for each parameter as the difference of latencies and ratio of amplitudes between  
398 the averaged values corresponding to the intervals [1-8] ms and [8-16] ms, i.e.  
399  $L_{[1-8]}-L_{[8-16]}$  and  $A_{[1-8]}/A_{[8-16]}$ , both in ABR and MLR signals. These parameters  
400 evaluate the changes on the waveform morphology depending solely on the  
401 previous SOA, thus directly associated with the fast adaptation. The Pb  
402 component was excluded from this analysis because of insufficient clear  
403 measures of this component, especially at high rates.

### 404 **3. RESULTS**

#### 405 **3.1. Experiment 1**

406 Figure 2 shows a comparison of the morphology of ABR and MLR signals  
407 obtained from one subject at different rates in two different recording-scenarios.  
408 The ABR signals used in this study, along with an analysis of the latency and  
409 amplitude of the wave V component, are presented in figures 2.A.1, 2.A.2 and  
410 2.A.3 respectively. Comparison of the morphology of ABR signals in both  
411 scenarios show remarkable differences. In scenario 1, as rate increases, the  
412 latency of the ABR components increases and the amplitude decreases, which  
413 is consistent with several previous studies (Jiang et al., 2009; Stone et al., 2009).  
414 However in scenario 2, the latency of wave V seems to be unaltered by rate, and  
415 the slope of the linear regression curve of the wave V amplitude obtained at each  
416 SOA range is lower than in scenario 1, which suggests that as rate increases, the  
417 amplitude of wave V decreases more slowly. ABR signals of both scenarios

418 obtained at rates faster than 400 Hz ( $SOA_{2-3}$ ) showed a high-level of adaptation  
419 and no wave V component could be identified.

420 Figure 2.B.1 shows the MLR signals obtained in this study. The Na, Pa, Nb and  
421 Pb components are labeled on the  $SOA_{8-12}$  MLR signal on this figure. All  
422 components could be identified at all rates, except Nb and Pb at 500 Hz ( $SOA_{0-4}$ )  
423 in both scenarios. The values of latency and amplitude of the MLR components  
424 obtained in scenario 1 are consistent with those reported on previous studies, in  
425 which MLR signals were recorded at fast rates (Özdamar et al., 2007). Figure  
426 2.B.2 shows the latencies and a linear regression analysis for the Na, Pa, Nb and  
427 Pb components at different rates. This analysis shows that, while Na latency is  
428 similar in both scenarios, the latency drift in the rest of the components is more  
429 accentuated in scenario 1 than in scenario 2. Analysis of amplitudes for the wave  
430 complexes Na-Pa, Pa-Nb and Nb-Pb is presented on figures 2.B.3, 2.B.4 and  
431 2.B.5 respectively. These figures show that, although amplitudes decrease as  
432 rate increases in both scenarios, amplitudes in scenario 1 present a greater value  
433 and the slope of the linear regression analysis is steeper in scenario 1 than in  
434 scenario 2. Data shown in this experiment is obtained from a single subject. A  
435 more robust study of amplitudes and latencies is presented in experiment 3 of  
436 this paper.

### 437 **3.2. Experiment 2**

438 Figure 3 shows the MLR signals used as reference (templates) and the MLR  
439 estimates by the Split-IRSA technique at a different number of iterations in a  
440 simulation study. Figures 3.A.1 and 3.A.2 show, respectively, the results of this  
441 study when no noise is added to the synthesized EEG in scenarios 1 and 2. These

442 figures show that the accuracy of the MLR estimates increases with the number  
443 of iterations. The MLR estimates obtained with 500 iterations in both scenarios  
444 approximate accurately the original templates (errors lower than  $0.0002 \mu V_{RMS}$  in  
445 all cases). Figures 3.B.1 and 3.B.2 show the results of a similar study in which  
446 noise was added to the synthesized EEG at a similar RMS value as in a real  
447 situation. As in the no-noise case, the accuracy of the MLR estimates increases  
448 with the number of iterations. Although the MLR estimates obtained with 500  
449 iterations in panel B present greater error-values than in the case of EEGs without  
450 added noise (panel A), these MLR estimates approximate the morphology of the  
451 original templates with sufficient accuracy to estimate correctly the amplitudes  
452 and latencies of the main components of these signals.

453 A similar study was carried out with ABR signals. The results of this study are  
454 consistent with those obtained in the study with MLR signals. These results  
455 indicate the ABR estimated by Split-IRSA after 50 iterations in both scenarios fit  
456 perfectly the templates (error estimates  $<0.00001 \mu V_{RMS}$ ) when no noise is added  
457 to the synthesized EEG. The ABR estimates in both scenarios when noise is  
458 added to the EEG present a higher level of noise, but the morphology of these  
459 estimates approximates the original templates. The figures that present the  
460 morphology of these ABR estimates are available as supplementary material in  
461 Appendix B. This appendix also includes tables with the RMS errors between the  
462 templates and the ABR/MLR estimates obtained in each scenario at each  
463 iteration analyzed in this study.

464 The results of this experiment point out that (a) the Split-IRSA technique is able  
465 to estimate accurately templates of different morphology in different jittering

466 conditions, and (b) the parameters  $\alpha$ -value and number of iterations selected on  
467 experiment 1 in this paper ( $I = 50$  in ABR,  $I = 500$  in MLR,  $\alpha = 0.8$ ) are  
468 appropriate.

### 469 **3.3. Experiment 3**

470 Figure 4 shows the grand-average ABR and MLR waveforms from a set of 8  
471 normal hearing subjects. Subject 2 was not included in the grand-average ABR  
472 waveforms since no clear components could be identified. Thick lines in the upper  
473 section on each panel represent the ABR and MLR signals obtained directly from  
474 the SOA<sub>0-16</sub> stimulation sequences, considering that all stimuli evoked the same  
475 response (time-invariant assumption). The main components of ABR and MLR  
476 are labeled on these signals. The rest of the lines represent the ABR/MLR  
477 responses corresponding to different rate-subsets obtained by the Split-IRSA  
478 technique, e.g., the ABR waveform corresponding to SOA<sub>15-16</sub> is obtained from  
479 the auditory responses corresponding to stimuli whose preceding SOA belonged  
480 to the interval [15-16] ms. This figure allows an overall study of the morphology  
481 of these signals across subjects. This figure shows that the morphology of ABR  
482 signals at different rate-subsets is very similar to the signal obtained from the  
483 complete stimulation sequence (upper-panel line), except for the ABRs obtained  
484 at very fast rates, i.e., SOA<sub>2-3</sub> and higher rates, where the latencies of the main  
485 components increase and their amplitude decrease significantly. On MLR  
486 signals, their morphology vary across different rate-subsets, especially at higher  
487 rates. The individual ABR and MLR signals obtained in each subject are available  
488 as supplementary material (appendix C).

489 Figure 5 and table 1 show the results of the linear regression analysis of the  
490 latencies (L) and amplitudes (A) of the main components of ABR (panel A) and  
491 MLR (panel B) signals versus the SOA intervals. The linear regression analyses  
492 in panel A show, on one hand, absence of statistically significant evidence for  
493 latencies and amplitudes being influenced by rate in the [4-16] ms SOA interval,  
494 and on the other, statistically significant evidence of variations on the amplitudes  
495 in the [0-8] ms SOA interval. These results point out that the time-invariant  
496 assumption is accomplished in ABR along the [4-16] ms SOA interval, but not at  
497 the fastest rates. The linear regression analyses in panel B show statistically  
498 significant evidence of variations of the morphology of MLR signals at different  
499 SOA intervals, thus indicating that the time-invariant assumption is not  
500 accomplished.

501 The inter-subject variability of the fast adaptation is analyzed in figure 6. This  
502 figure shows a significant variability across subjects. For instance: (a) subjects  
503 S1, S7 and S8 show a larger fast adaptation on the latency of ABR wave III than  
504 subjects S4, S5 and S6; (b) subject S4 shows a particular low fast adaptation on  
505 the amplitude of ABR waves III and V; (c) S4 is also the only subject in which the  
506 latency of the ABR wave V and the MLR Na components decreased at high rates;  
507 and, (d) subjects S1 and S2 show a lower fast adaptation than the rest of the  
508 subjects on the latency of the MLR Pa and Nb components. In addition, this study  
509 shows a large variability across different parameters within the same subject. For  
510 example, subject S1 is the subject showing the largest fast adaptation on the Na  
511 latency, but it is also the subject presenting the lowest fast adaptation on the  
512 latency of the Pa and Nb components.

## 4. DISCUSSION

513

514 This paper presents a full description of the iterative-randomized stimulation and  
515 averaging Split (Split-IRSA) technique. The fundamentals of this technique are  
516 similar to IRSA, described in Valderrama et al. (2014c), with the difference that  
517 Split-IRSA includes selective processing of responses, i.e., each response can  
518 be individually processed and categorized according to a predefined criteria.  
519 Split-IRSA allows, therefore, overlapping auditory evoked responses of different  
520 morphology to be obtained by an iterative procedure in the time domain. The  
521 main advantages of the Split-IRSA technique are: (a) stimulation sequences are  
522 based on randomized stimulation, which allows the amount of jitter to be under  
523 control; (b) this technique includes a mechanism to control convergence ( $\alpha$ -  
524 value); (c) Split-IRSA is easy to implement (programming code attached on  
525 appendix A of this paper); and (d) it allows selective processing of auditory  
526 responses.

527 The performance of the Split-IRSA technique was validated in this paper through  
528 experiments with both simulation and real data. The results of these experiments  
529 point out that this technique presents an adequate performance when the  $\alpha$ -value  
530 and the number of iterations are correctly defined. The simulation study  
531 presented in experiment 2 shows that the AEP estimates obtained with Split-IRSA  
532 on the first iteration (blue signals on figure 3 and in appendix B on this paper)  
533 were not accurate, i.e., they present a morphology different from the template  
534 signal. This is consistent with results presented in Valderrama et al. (2014c),  
535 where we found that interference associated with overlapping responses  
536 introduces an artifact in the AEP estimate which cannot be reduced by averaging



537 when the amount of jitter of the stimulation sequence is lower than the dominant  
538 period of the recorded AEPs (i.e., 2 ms in ABR and 25 ms in MLR). Thus, a single  
539 iteration was not sufficient to obtain accurate AEP estimates. The results of  
540 experiment 2 show that more accurate ABR/MLR estimates can be obtained  
541 recursively. The results of experiments 1 and 3 in this paper point out that the  
542 Split-IRSA technique has allowed real ABR and MLR signals of different  
543 morphologies to be recorded simultaneously at very rapid rates using narrow-  
544 jittered stimulation sub-sequences.

545 The flexible nature of Split-IRSA is appropriate for research purposes. In this  
546 paper, we have used this technique to analyze the variations in the morphology  
547 of ABR and MLR signals across different rate-subsets in 16 ms-jittered  
548 stimulation sequences in order to evaluate the time-invariant assumption all along  
549 the stimulation sequence. This topic may be of interest as time-invariance is  
550 assumed in all techniques that process evoked potentials (Bardy et al., 2014a;  
551 Jewett et al., 2004, Özdamar and Bohórquez, 2006), and secondly, it is still not  
552 clear whether or not the amount of jitter of a stimulation sequence is a critical  
553 parameter to be considered when assuming that each stimulus evokes the same  
554 ABR/MLR response (Jewett et al., 2004, Özdamar and Bohórquez, 2006). As far  
555 as we are concerned, the methodology presented in this paper is the first attempt  
556 to analyze the time-invariant assumption in real ABR and MLR signals obtained  
557 in a specific jittered stimulation sequence.

558 Analysis of ABR and MLR waveforms obtained in scenarios 1 and 2 in  
559 experiment 1 provide evidence that both fast and slow mechanisms of adaptation  
560 interact when presenting jittered stimuli. These fast and slow mechanisms of

561 adaptation have been observed in a number of animal studies (Chimento and  
562 Schreiner, 1991; Eggermont, 1985; Javel, 1996; Yates et al., 1985; Westerman  
563 and Smith, 1984) and in ABR signals recorded with long- and short-SOA  
564 distributions (Valderrama et al., 2014b). If ABR/MLR waveforms in scenarios 1  
565 and 2 were similar, it would be suggested that fast mechanisms of adaptation  
566 prevail over slow mechanisms, since the morphology of the response would be  
567 mostly influenced by the SOA of the preceding stimulus. In contrast, if ABR and  
568 MLR waveforms in scenario 2 were similar among themselves (and different to  
569 those obtained in scenario 1), that would indicate that slow mechanisms of  
570 adaptation prevail over fast mechanisms, since the morphology of the ABR/MLR  
571 signal would be determined by an averaged stimulation rate corresponding to  
572 several preceding stimuli. The results obtained in experiment 1 show that, in ABR  
573 signals on scenario 2, the latency of wave V remained constant across most of  
574 the sub-rates and that the amplitude decreased at a lower rate than in scenario  
575 1. These results highlight the significant role of slow mechanisms of adaptation.  
576 The morphology of MLR signals in scenario 2 present significant variations  
577 among themselves, as a consequence of the fast mechanisms of adaptation,  
578 however in comparison with the MLRs on scenario 1, latencies seem less  
579 dependent on rate, amplitudes are smaller, and decrease with rate more slowly.  
580 These results point out the effects of both fast and slow mechanisms of  
581 adaptation.

582 The results obtained in experiment 3 are consistent with those obtained in  
583 experiment 1. These results show that the MLR waveforms obtained at different  
584 rate-subsets present significant variations as a consequence of the

585    aforementioned fast and slow mechanisms of adaptation. This variability  
586    indicates that the time-invariant assumption is not accomplished all along the  
587    stimulation sequence. A direct consequence of this deviation from the time-  
588    invariant behavior is a degradation of the quality of the recordings, since the  
589    components are not phase-locked when the sweeps are averaged. The variability  
590    of the latencies observed in this study suggests that a possible strategy to  
591    improve the quality of the recordings could be the adjustment of the time-axis in  
592    each individual sweep in order to average phase-locked auditory responses.

593    In contrast to MLR, this study did not show differences in the morphology of ABR  
594    signals obtained at rate-subsets down to SOA<sub>4-5</sub> (equivalent rate of 222 Hz),  
595    which shows the influence of the slow mechanisms of adaptation and that the  
596    time-invariant assumption is accomplished in this SOA range ([4-16] ms). The  
597    amplitudes of the ABR signals obtained at faster sub-rates present a significant  
598    decrease, indicating the prevalence of fast mechanisms of adaptation. The  
599    influence of the fast adaptation is particularly relevant at very fast rates, as in the  
600    SOA<sub>1-2</sub> sub-sequence the ABR components could be detected in only a few  
601    subjects, and no subject showed any clear component at the SOA<sub>0-1</sub> sub-  
602    sequence. The strong influence of the fast mechanisms of adaptation at these  
603    very fast rates could be associated with the refractory period of the neurons of  
604    the auditory pathway (Alvarez et al., 2011).

605    The results obtained in this study contradict the classical approach that claims  
606    that wide-jittered stimulation sequences can be a problem when assuming time-  
607    invariance of the response, since large SOA variations would evoke responses  
608    of different morphology. This classical approach only considers the fast

609 mechanisms of adaptation. In contrast, this study highlights that both fast and  
610 slow mechanisms of adaptation influence the morphology of the evoked  
611 responses in jittered sequences, and therefore, both mechanisms should be  
612 considered when evaluating the time-invariant assumption in jittered stimulation  
613 sequences.

614 The mechanisms of adaptation have been attributed different functionalities in the  
615 auditory system. For example, the adaptive processes at different levels of the  
616 auditory pathway have been proven to enhance novelty detection (Ulanovsky et  
617 al., 2009), and to improve the neural coding accuracy by accommodating the rate-  
618 level function of the neurons to the characteristics of the input sound (Dean et al.,  
619 2005; Wen et al., 2009). The evaluation of the time-constants of the fast and slow  
620 mechanisms of adaptation observed in this study could have a potential clinical  
621 application in the future.

622 Future research could also investigate the manner in which the SOA jitter  
623 distribution influences the fast and slow adaptation mechanisms. The  
624 understanding of this relationship could help design stimulation sequences with  
625 prevalence of the slow mechanisms of adaptation, thus accomplishing the time-  
626 invariance assumption.

627

## 5. CONCLUSIONS

628 This paper describes in detail the Split-iterative randomized stimulation and  
629 averaging (Split-IRSA) technique. This technique allows overlapping AEPs of  
630 different morphology to be disentangled through an iterative procedure in the

631 time-domain. The results obtained with real and synthesized data indicate that  
632 the performance of this technique is robust when the parameter that controls  
633 convergence ( $\alpha$ -value) and the number of iterations are adequately selected. A  
634 new strategy was designed to evaluate the time-invariant assumption on the AEP  
635 morphology in jittered sequences. The results point out that both fast and slow  
636 mechanisms of adaptation influence the AEP morphology, and therefore, both  
637 mechanisms should be taken into account when time-invariance is assumed.

### 638 **Declaration of interest**

639 The authors report no conflict of interest.

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### 648 **Supplementary data**

649 Supplementary data associated with this article can be found, in the online  
650 version, at [URL].

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## 777 **Figure Legends**

- 778 • Figure 1. Performance and parameters involved on the Split-IRSA technique.

779 (A) Histogram of the inter-stimulus interval (SOA) of an example stimulation  
780 sequence  $s(n)$ . The sub-sequences  $s_1(n)$  and  $s_2(n)$  are marked on the figure.

781 (B) Parameter settings of this experiment. (C) Normalized energy ( $\mu V^2$ ) of the  
782 averaged residual,  $\frac{1}{K_\tau} \cdot \sum_{k=1}^{K_\tau} z_i(j + \mathbf{m}_\tau(k))$ , at different iterations and  $\alpha$ -values.

783 This figure shows that instability problems (normalized energy increases with  
784 the number of iterations) can be avoided by selecting an appropriate value of  
785  $\alpha$ . (D.1 and D.2) Evoked potential estimates at different iterations under  
786 instability: worse estimates are obtained in succeeding iterations. (E.1 and  
787 E.2) Evoked potential estimates at different iterations in a convergence  
788 scenario: better estimates are obtained in succeeding iterations, e.g., error  
789 between the original template and the estimates decrease as iterations  
790 increase.

- 791 • Figure 2. Comparison of the morphology of ABR and MLR signals recorded  
792 from one subject (scenario 1) by narrow-jittered stimulation sequences and  
793 processed by IRSA and (scenario 2) by a single 16 ms-jittered stimulation  
794 sequence and processed by the Split-IRSA technique in different subsets of  
795 stimuli. (A.1) ABR signals obtained at different average SOA (Av SOA) in each  
796 scenario. (A.2 and A.3) Latency (ms) and amplitude ( $\mu V$ ) of wave V and linear  
797 regression analysis evaluated at different rates in scenarios 1 and 2. (B.1)  
798 MLR signals obtained in each scenario and rate. (B.2) Latencies (ms) and  
799 linear regression analysis measured on the components Na, Pa, Nb and Pb  
800 at different rates in each scenario. (B.3, B.4 and B.5) Amplitudes ( $\mu V$ ) and

801 linear regression analysis of the waves complexes Na-Pa, Pa-Nb and Nb-Pb  
802 at different rates in both scenarios.

803 • Figure 3. MLR signals estimated by the Split-IRSA technique at a different  
804 number of iterations in a simulation study that reproduces the acquisition  
805 settings of experiment 1 when no noise is added to the synthesized EEG  
806 (panel A) and when noise is added at a similar RMS value as in a real situation  
807 (panel B). Errors between the MLR estimates obtained at 500 iterations and  
808 the original templates are shown in  $\mu V_{RMS}$ .

809 • Figure 4. Grand-average ABR (panel A) and MLR (panel B) waveforms from  
810 a set of 8 normal hearing subjects. Thick lines represent the ABR/MLR signals  
811 obtained from the complete sequence  $SOA_{0-16}$ , and standard lines show the  
812 responses obtained at each rate-subset by the Split-IRSA technique.

813 • Figure 5. Latencies (L) and amplitudes (A) of the main components of ABR  
814 (panel A) and MLR (panel B) signals obtained at the average SOA ( $Av$  SOA)  
815 of different rate-subsets. In panel A, the black and grey lines represent a linear  
816 regression analysis between the SOA intervals [4-16] and [0-8] ms,  
817 respectively. In panel B, the black line shows the linear regression analysis  
818 for the [0-16] ms SOA interval. The statistical analysis of these hypothesis  
819 tests are shown in table 1.

820 • Figure 6. Inter-subject variability of the fast adaptation. The fast adaptation  
821 was measured as the difference of latencies (in ms) and ratio of amplitudes  
822 between the averaged values corresponding to the intervals [1-8] ms and [8-  
823 16] ms, i.e.  $L_{[1-8]} - L_{[8-16]}$  and  $A_{[1-8]}/A_{[8-16]}$ , both in ABR (panel A) and MLR (panel

824 B) signals. Black boxes represent the estimates measured on the Grand-  
825 Average (GA) ABR/MLR waveforms, while the white boxes are the estimates  
826 for each individual subject.

## 827 **Table Legends**

- 828 • Table 1. Statistic parameters of the linear regression hypothesis tests  
829 presented on figure 5. Legend: N, number of observations; r, correlation  
830 coefficient;  $R^2$ , coefficient of determination;  $p$ -value, probability of rejecting the  
831 null hypothesis; a, angle slope; b, y-intercept; SE, standard error. \* represents  
832  $p$ -value < 0.05; # represents  $p$ -value  $\approx$  0.05.