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1. SUPPLEMENTARY MATERIAL

1.1. Figure S1

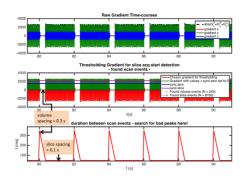


Figure 51: Determining scan timing from gradient logging for Phillips data (sync.method = 'gradient_log'). Besides cardiac and respiratory data, the Phillips SCANPHYSLOG-file stores time courses of the x,y,e-gradients at a coarse temporal resolution (2 ms) (top). Their regularity, however, is sufficient to infer on slice and volume repetitions, and does not change over subjects. Thus, for each study, one only has to determine once which gradient shows the highest regularity (e.g. sync.grad_direction = 'y', green curve), set lower thresholds for irrelevant and slice peaks (middle, sync.zero = 1500, blue line; sync.silce = 2200, mogenta line), as well as volume peaks, if they differ from slice peaks (not shown, sync.vol = []). If volume peaks are not discernible, the non-equidistant slice spacing between volumes can be used to identify volume onsets (bottom, red curve, sync.vol_spacing = 0.3 seconds).

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