

Simple Epidemic Models with Segmentation Can Be Better than Complex Ones (SUPPLEMENTARY DOCUMENT)

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Abstract

In this supplementary document, we provide additional experimental results that are not included in the main paper, which is titled *Simple Epidemic Models with Segmentation Can Be Better than Complex Ones*.

Keywords: COVID-19, Epidemic models, Segmentation, Minimum Description Length

Figure 1. In Figure 1, we provide the input and estimated event sequences when the NLLD model is used with our proposed segmentation scheme.

Figure 2, 3, and 4. We demonstrate the effectiveness by measuring how segmentation affects the model complexity and fitting error of the three considered epidemic models.

⁵ We provide the results when the NLLD, LLD, and SIR models are used in Figure 2, Figure 3, and Figure 4, respectively.

Figure 5, 6, and 7. In addition, we examine the effectiveness of our greedy segmentation scheme based on the MDL principle by comparing it with the incremental segmentation method, in terms of the model complexity and fitting error of the three

¹⁰ considered epidemic models. We provide the results when the NLLD, LLD, and SIR models are used in Figure 5, Figure 6, and Figure 7, respectively.

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Preprint submitted to Journal of L^AT_EX Templates

July 15, 2021

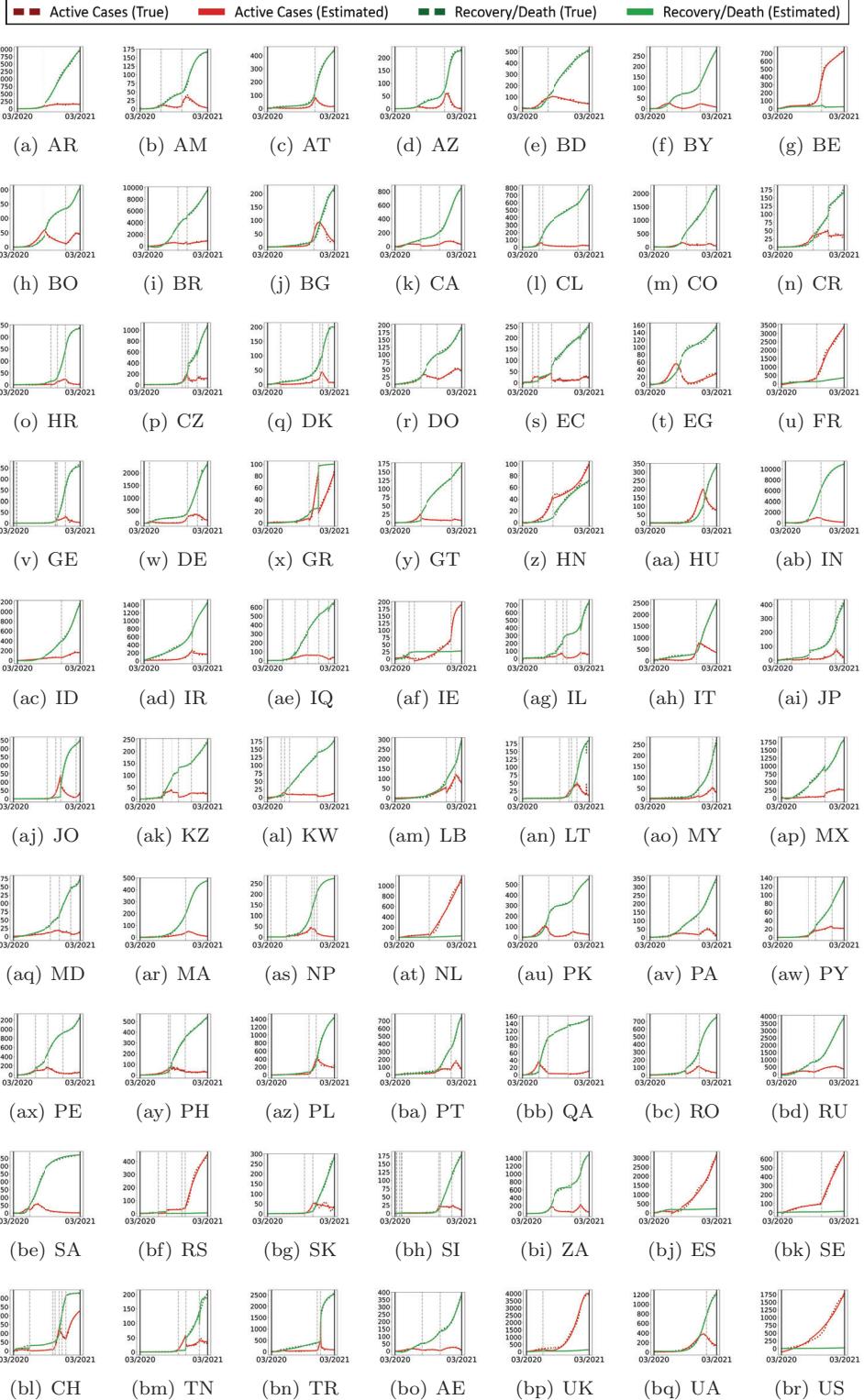


Figure 1: The true and estimated event sequences when the NLLDmodel is used with our proposed segmentation scheme. The scale of the values in the y-axis are thousands ($10^3 \times$).

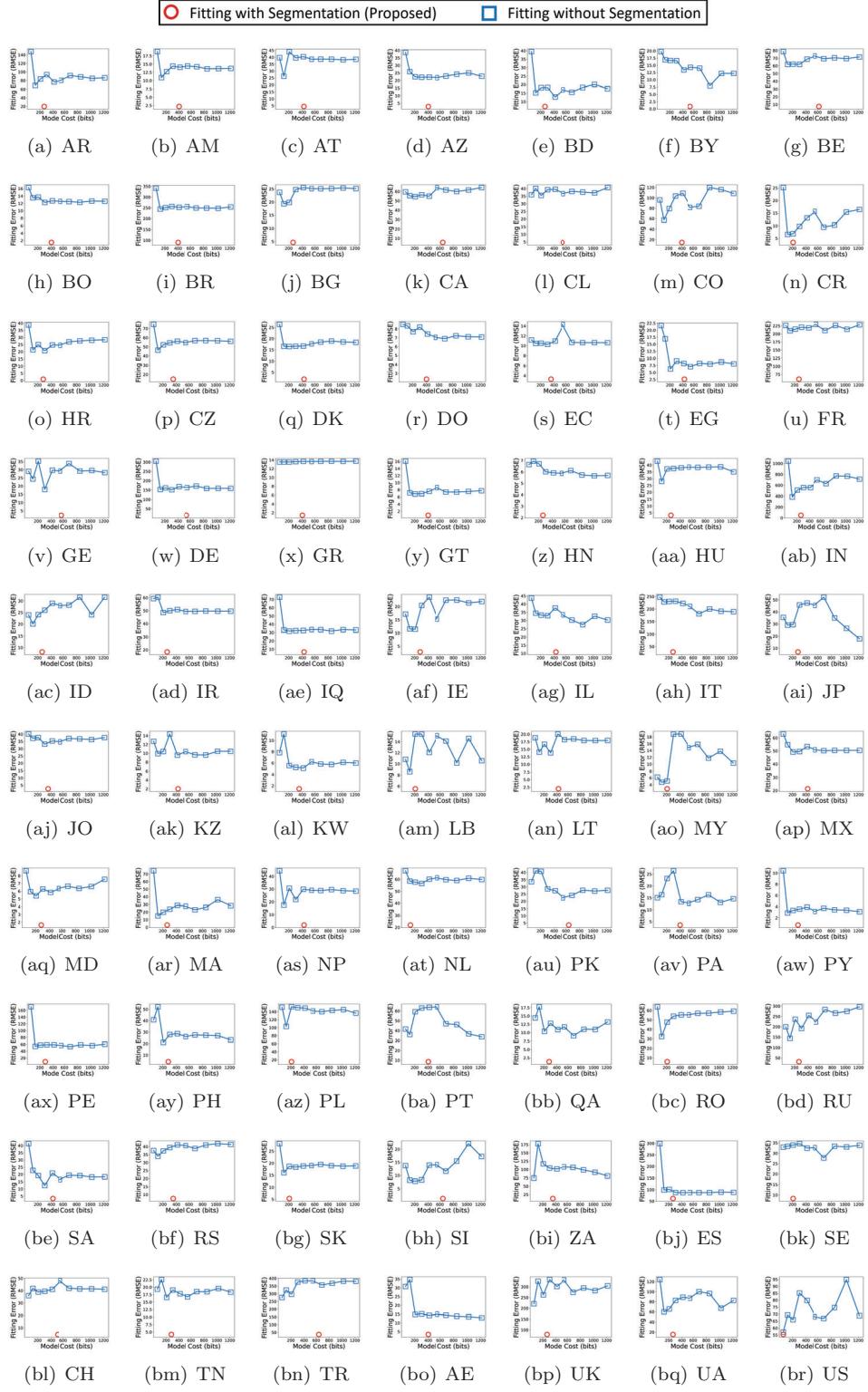


Figure 2: The true and estimated event sequences when the NLLDmodel is used with our proposed segmentation scheme. The scale of the values in the y-axis are thousands ($10^3 \times$).

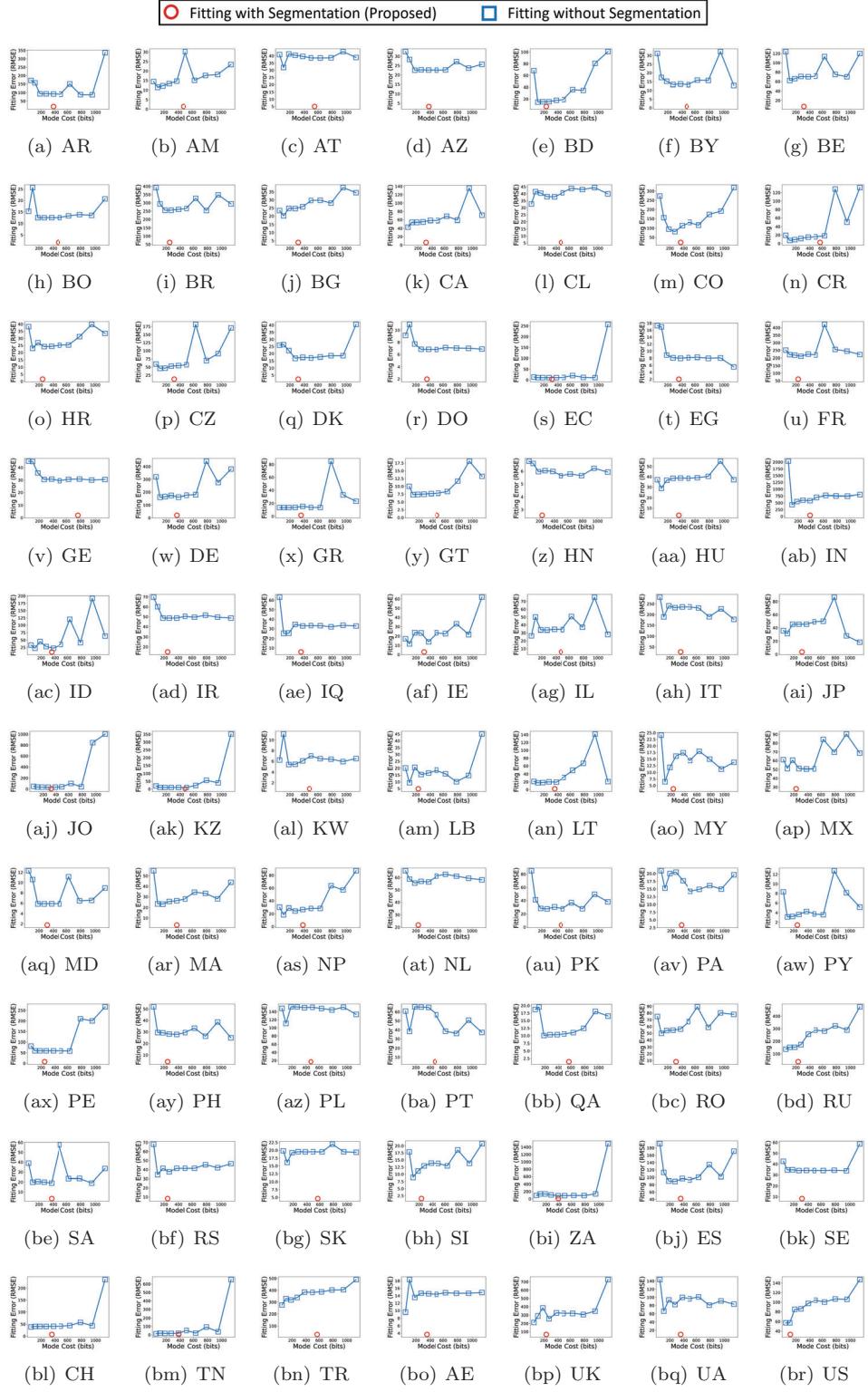


Figure 3: The true and estimated event sequences when the LLDmodel is used with our proposed segmentation scheme. The scale of the values in the y-axis are thousands ($10^3 \times$).

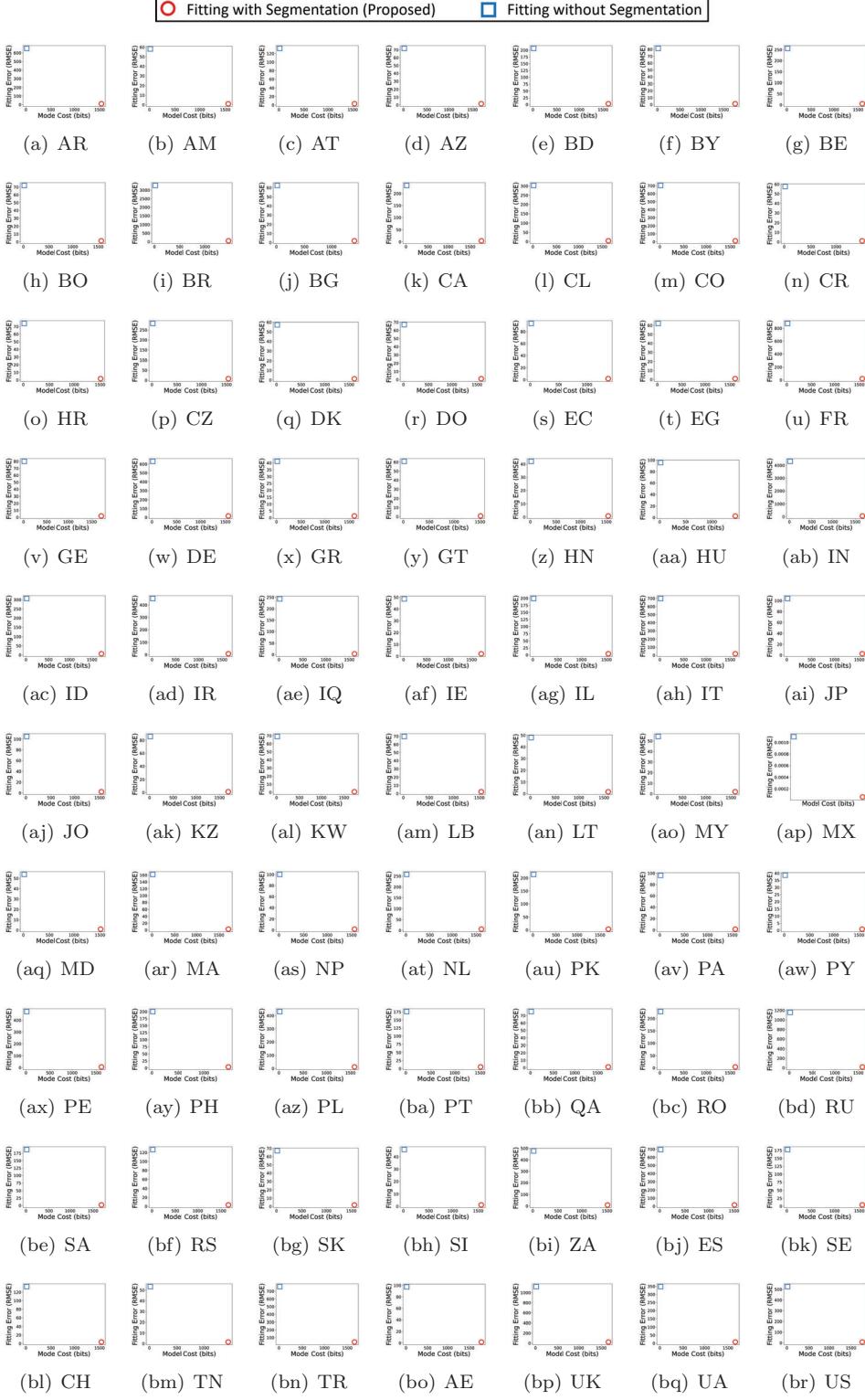


Figure 4: The true and estimated event sequences when the SIR model is used with our proposed segmentation scheme. The scale of the values in the y-axis are thousands ($10^3 \times$).

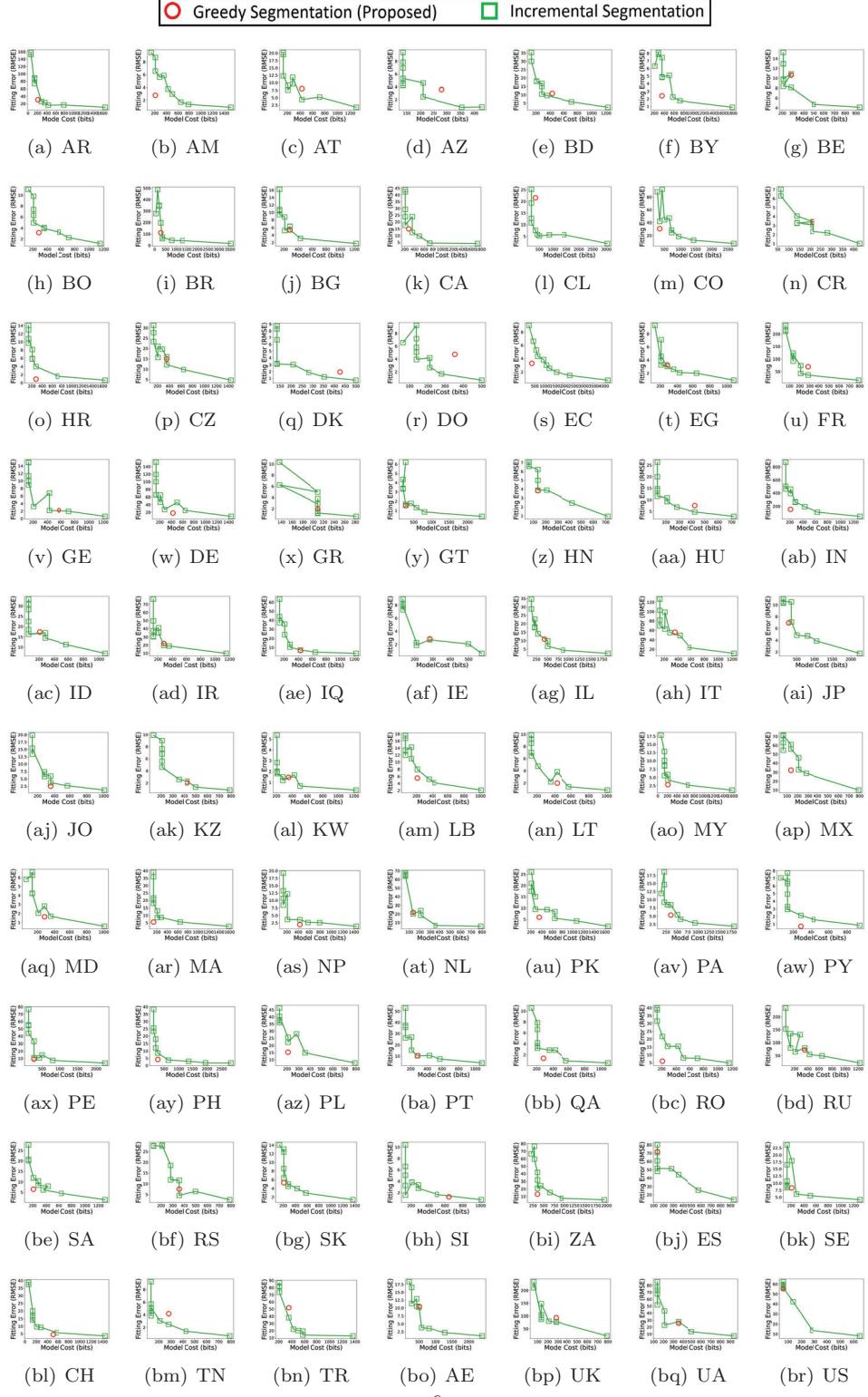


Figure 5: The true and estimated event sequences when the NLLDmodel is used with our proposed segmentation scheme. The scale of the values in the y-axis are thousands ($10^3 \times$).

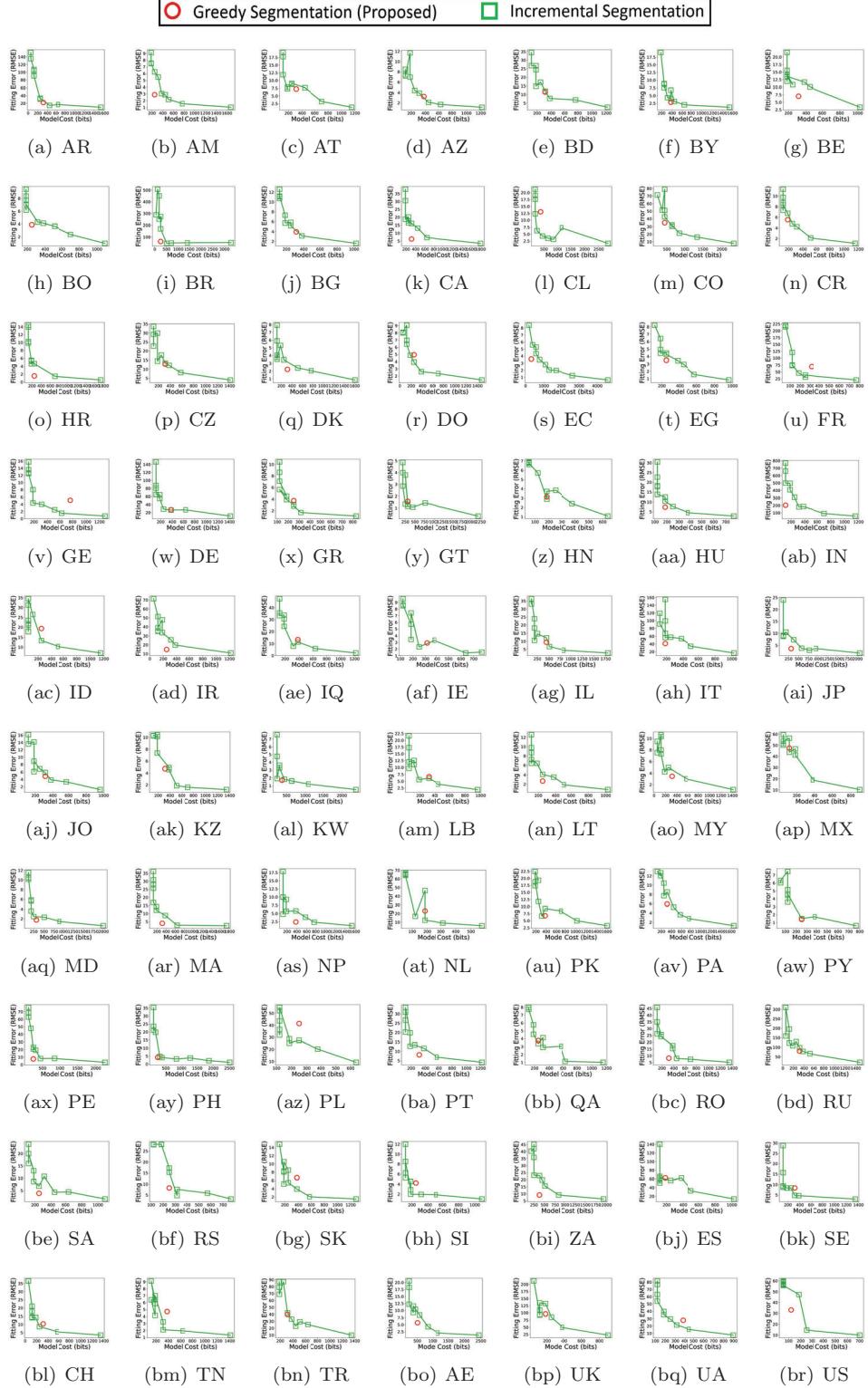


Figure 6: The true and estimated event sequences when the LLDmodel is used with our proposed segmentation scheme. The scale of the values in the y-axis are thousands ($10^3 \times$).

○ Greedy Segmentation (Proposed) □ Incremental Segmentation

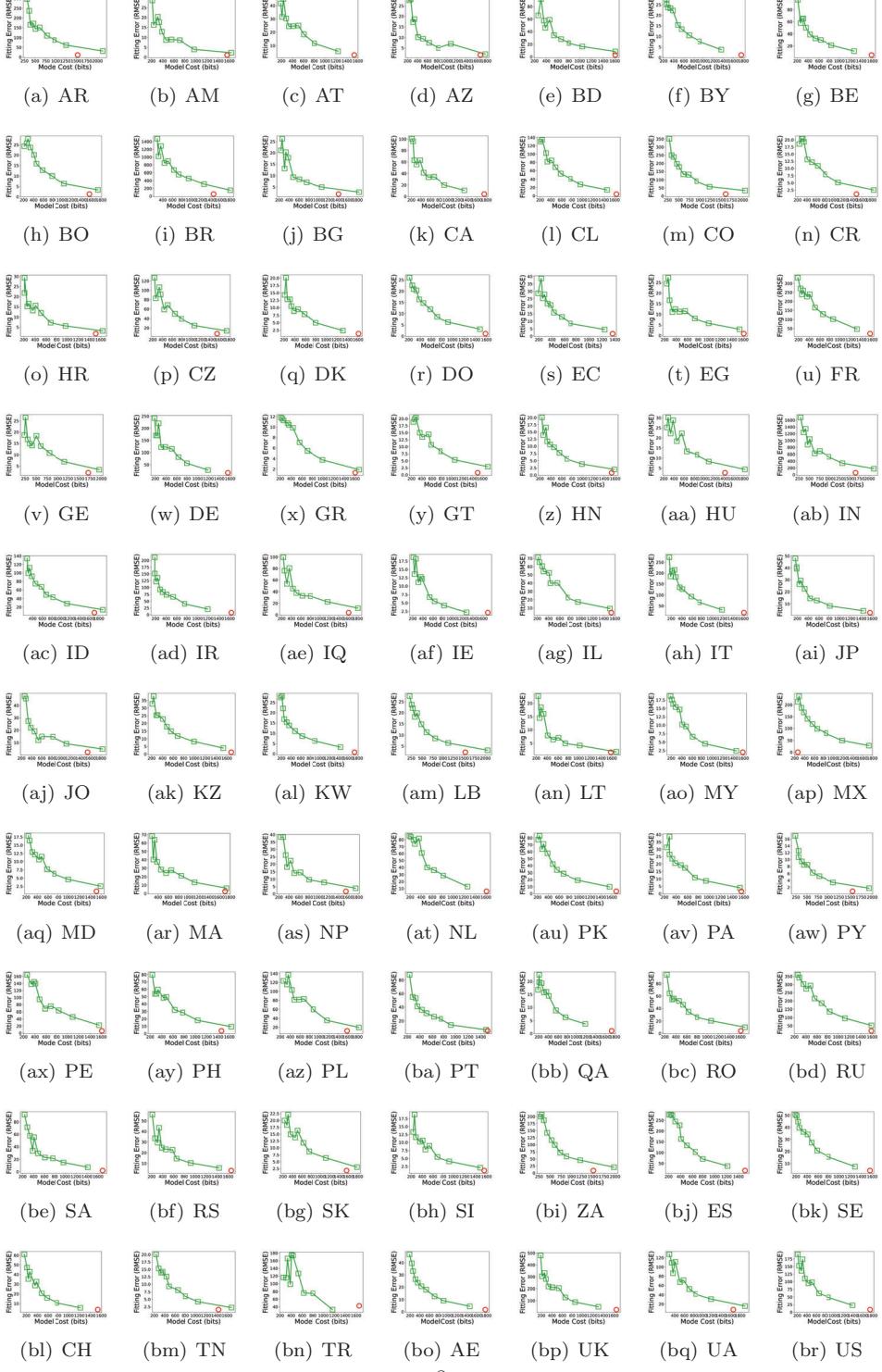


Figure 7: The true and estimated event sequences when the SIR model is used with our proposed segmentation scheme. The scale of the values in the y-axis are thousands ($10^3 \times$).