



A Location-Mixture Autoregressive Model for online forecasting of lung tumor motion

Debdeep Pati*

Florida State University, Tallahassee, USA, debdeep@stat.fsu.edu

Daniel Cervone

Harvard University, Cambridge, MA, USA, dcervone@fas.harvard.edu

Natesh Pillai

Harvard University, Cambridge, MA, USA, pillai@fas.harvard.edu

Ross Berbeco

Harvard Medical School, Cambridge, MA, USA, rberbeco@lroc.harvard.edu

John Henry Lewis

Harvard Medical School, Cambridge, MA, USA, jhlewis@lroc.harvard.edu

Lung tumor tracking for radiotherapy requires real-time, multiple-step ahead forecasting of a quasi-periodic time series recording instantaneous tumor locations. We introduce a location-mixture autoregressive (LMAR) process that admits multimodal conditional distributions, fast approximate inference using the Expectation-Maximization algorithm and accurate multiple-step ahead predictive distributions. LMAR outperforms several commonly used methods in terms of out-of-sample prediction accuracy using clinical data from lung tumor patients. With its superior predictive performance and real-time computation, the LMAR model could be effectively implemented for use in current tumor tracking systems.

Keywords: radiotherapy; nonlinear; motifs; prediction