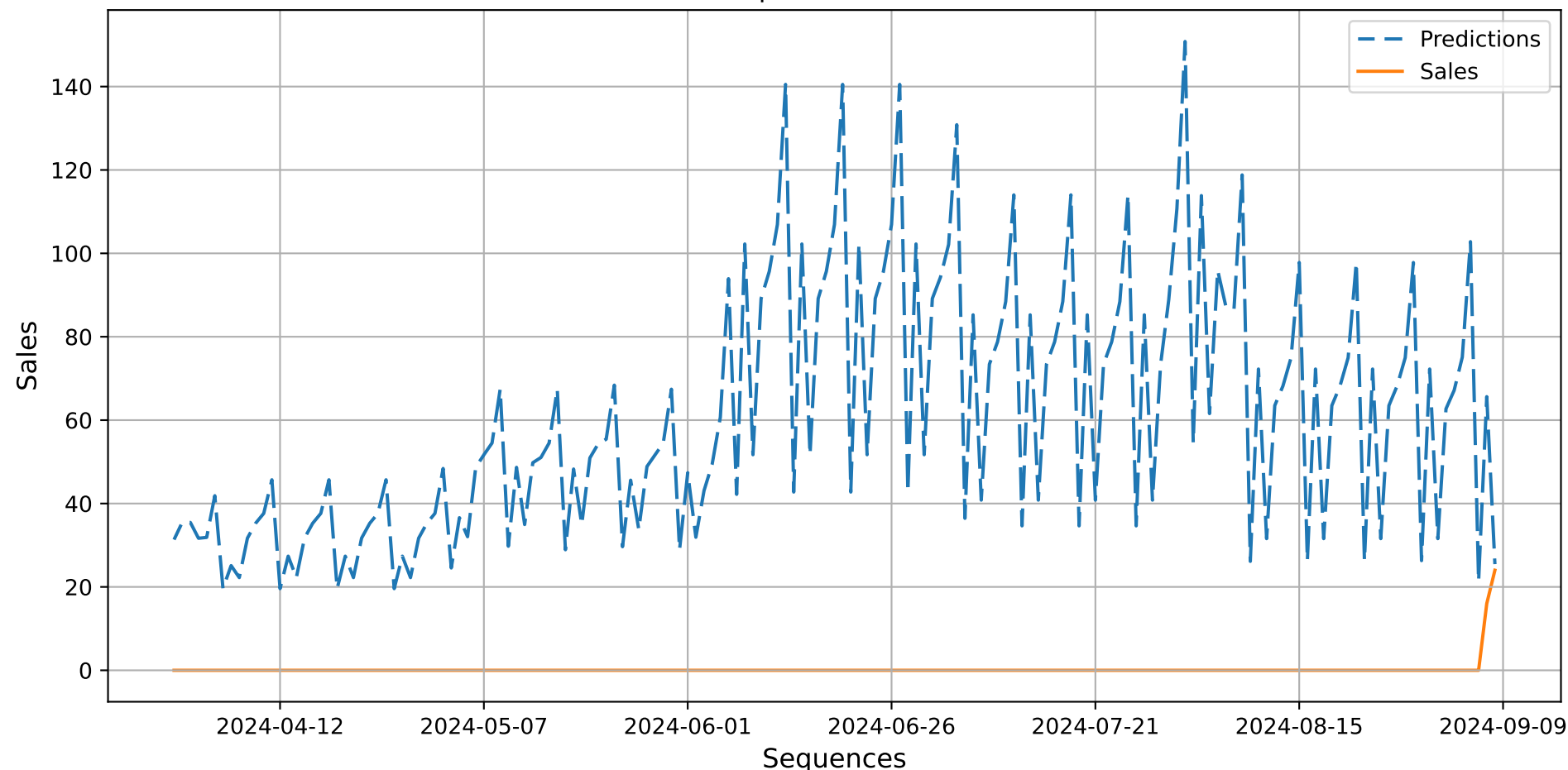# Test dataset predictions compared to sales for 4th worst performing article
704300501: LIME ÄNGLAMARK 3STK

Number of features: 351
Sequence length: 5
Number of epochs: 5
Learning rate: 0.0001
Loss function: mean squared error
Batch size: 32
Number of neurons: 351
Dropout rate: 0.05

Sequences

Mean absolute error: 60.31
Mean absolute percentage error: 270190977717928928.00
Root mean squared error: 67.60
Coefficient of determination: -905.0960