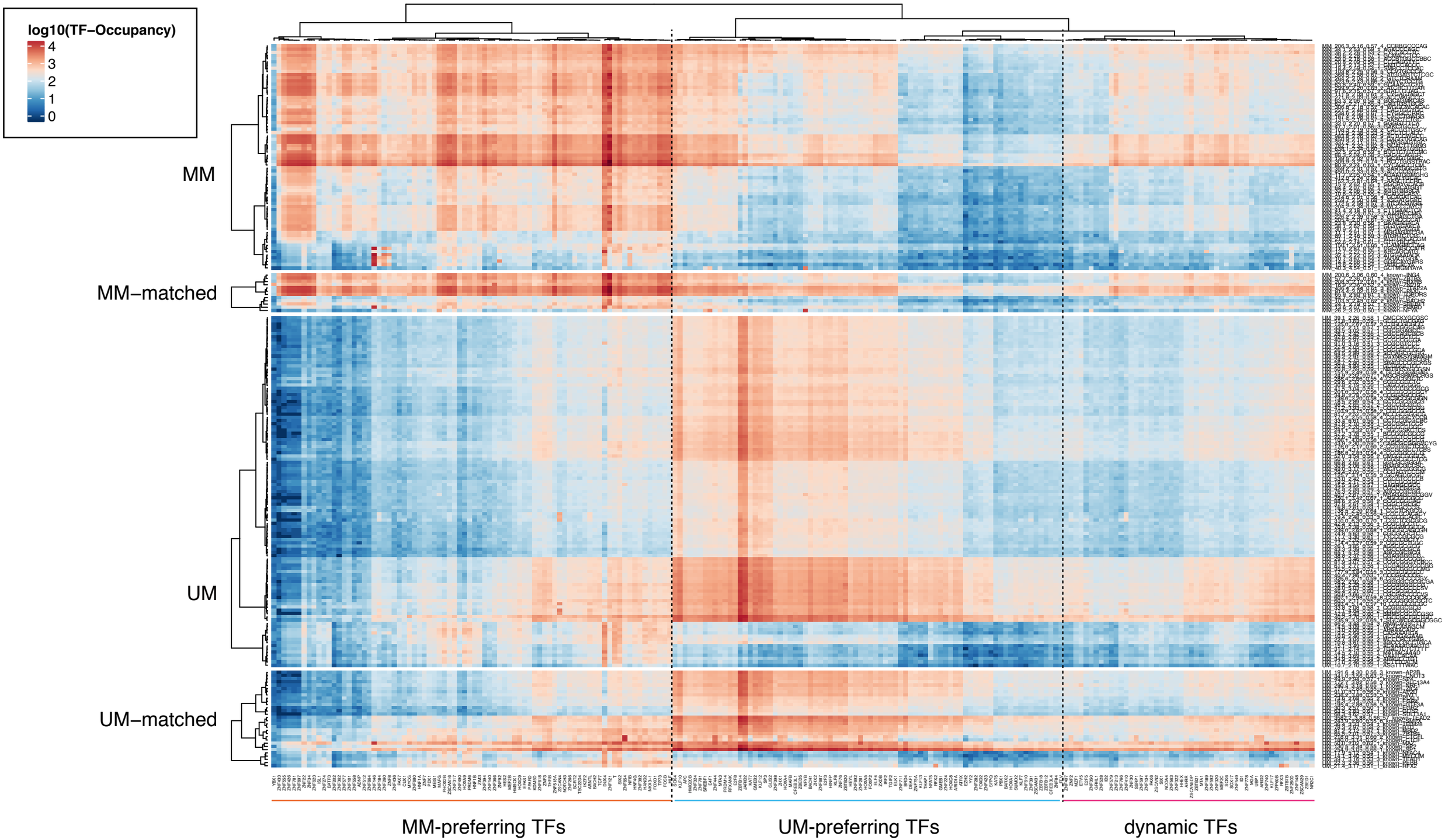Occupancy of TFs ChIP-seq peaks at identified motif sites