Supplementary Figure 2. Representative p67 N-linked glycosylation sites. Potentially glycosylated Asparagines (N) are in red and are flanked by consensus amino acids in blue. The value crossing the default threshold of 0.5, represents a predicted glycosylated site (as long as it occurs in the required sequon Asn-Xaa-Ser/Thr without Proline at Xaa). The 'potential' score is the averaged output of nine networks. For further information, the jury agreement column indicates how many of the nine networks support the prediction.