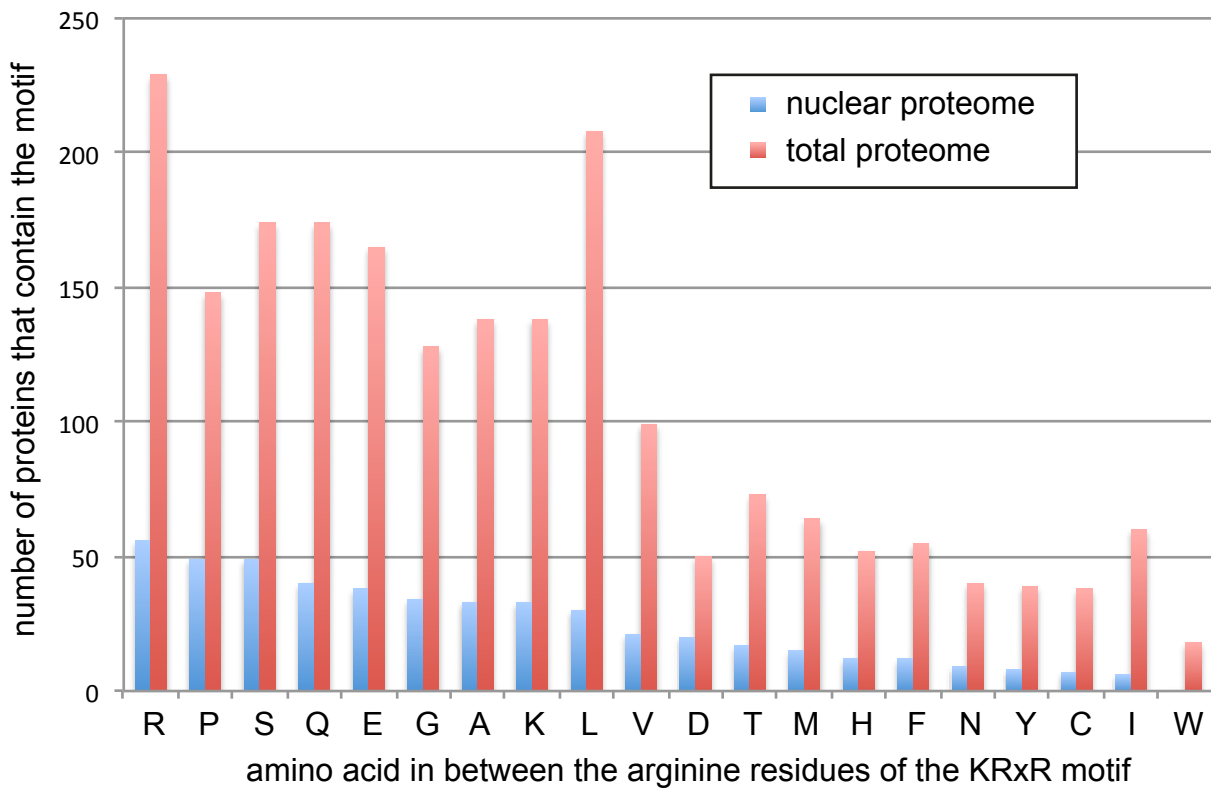


Figure S1



S1 Fig: The nuclear proteome and the total proteome of *T. brucei* Treu927 was searched for the KRxR motif with x being filled successively by all amino acids (x-axis). Note that the total number of proteins with the individual motifs is higher than the number of proteins that contain the KRxR motif, because some proteins contain several mixed motifs.