Sequence logos for all C-X7-C-X5-C-X3-H and C-X8-C-X5-C-X3-H zinc finger motifs identified in T. brucei (Tb), T. cruzi (Tc) and L. major (Lm) that were recognized by at least either SMART (Sm00356), Pfam (FP00642) or Interpro (IPR000571). Conserved amino acids that were used for the evaluation of unknown CCCH motifs are shown in red (very conserved, used in first filtering) and green (conserved, used in second filtering). The sequence logos were created using WEBLOGO [48].