CORRIGENDUM

BTKbase, mutation database for X-linked agammaglobulinemia


The authors of the above paper wish to note that the scale of the topmost panel in Figure 1 of the above paper as published is incorrect. The corrected Figure, including the legend, is published below.

Figure 1. Distribution of the mutations in the BTK protein (mutation spectrum). (a) Domain organization. The number of different types of mutations in the BTK; (b) mutations not causing frameshift, i.e., missense and inframe insertions and deletions (the four inframe mutations are indicated below the X-axis), (c) all the frameshift mutations (nonsense, insertions and deletions), (d) nonsense mutations, (e) frameshift deletions and (f) frameshift insertions. In (c) and (f) four mutations at residue 70 are presented. These mutations in unrelated families occur in a stretch of seven A nucleotides, and exact localization of a singel A insertion in unknown.