



Figure S3. Frequency of deletions starting and/or ending at a given exon was significantly high in a proximal (exons 3 to 13, p -value = 0.000015, f = 20.64776) and a distal (exons 44 to 55, p -value = 0.000069, f = 17.2277) region. 88.12% of individuals harbored deletions starting and/or ending within these two regions of the dystrophin. The average frequency of in-frame deletions starting and/or ending at a given region and all other exons was compared using f -statistic.