



Figure S2. Patient pool and *dystrophin* mutational patterns among the individuals with in-frame large deletions. (A) Sources of the clinical data on individuals with in-frame large deletions in *dystrophin*. (B) The proportion of deletions present to all theoretically possible in-frame large deletions ($N = 1408$) in each data source. (C) Frequency of in-frame large deletions starting or ending at exons 3 to 13 and exons 44 to 55 regions, compared to all other exons. $***p < 0.0001$, n.s. indicates a non-significant difference.