Supplementary material to article by L. Guerra et al. “A Postzygotic ATP2A2 Novel Mutation Identified by Next-generation Sequencing in Mosaic Darier Disease

Fig. S1. (a) Illumina data showing 14 of 44 reads (31.8%) bearing the c.2256_2258delCTA mutation. Sanger sequencing validates the mutation only in the DNA purified from affected epidermis. (b) Nucleotide peaks corresponding to the mutant allele are indicated by asterisks.