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Supplemental Data

Autosomal-Dominant Multiple Pterygium Syndrome Is Caused by Mutations in *MYH3*

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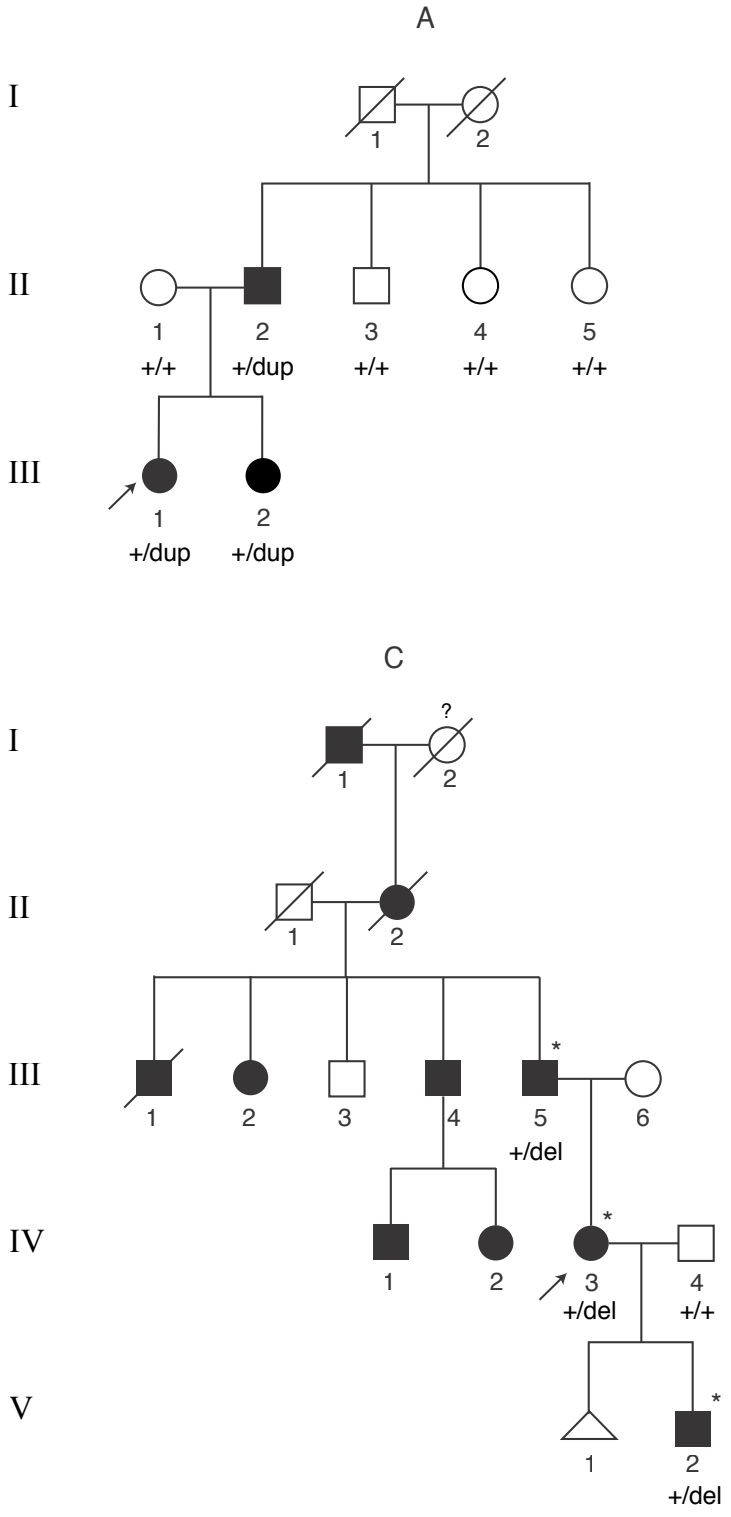
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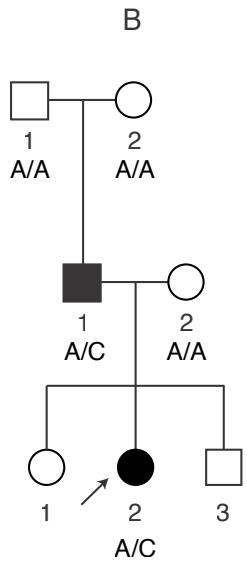
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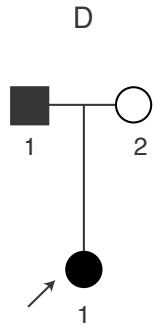
D
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Figure S1. Pedigrees of families of persons initially diagnosed with Distal Arthrogryposis type 8.

Case identifiers for each individual shown correspond to those in Table 1 and Figures 1 and 2. Each pedigree depicts a multiplex family (i.e., multiple persons affected with DA8) and mutation statuses confirmed by Sanger sequencing are provided below each individual. In family A, dup = c.3214_3216dup [p.(Asn1072dup)]. In family B, C = c.3224A>C [p.(Gln1075Pro)]. In family C, del = c.727_729del [p.(Ser243del)]. No mutation in *MYH3* was identified in family D.