

Online Supplementary Material

A founder mutation in Myosin-Binding-Protein C with an early onset and a high penetrance in males

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Table S1. Markers selected for haplotype analysis.

| Marker ID | Alleles | Position | MAF Global | MAF Iberian |
|-----------------------|----------------|-----------------|-------------------|--------------------|
| <i>rs7120013</i> | G/A | 46,155,938 | 0.375 | 0.411 |
| <i>MYBPC3 c.2149-</i> | G/A | 47,360,235 | 0 | 0 |
| <i>rs4882135</i> | A/T | 48,334,369 | 0.354 | 0.449 |
| <i>rs671299</i> | C/G | 56,307,532 | 0.349 | 0.309 |
| <i>rs7116652</i> | C/A | 58,236,065 | 0.404 | 0.421 |
| <i>rs10792299</i> | T/G | 60,721,699 | 0.163 | 0.280 |
| <i>rs11605489</i> | G/A | 62,472,357 | 0.455 | 0.365 |

1. MAF: minor allele frequency, HSA11: human chromosome 11

FIGURE LEGEND:

Figure S1. Splicing Analysis