

## DIAGNOSIS OF SPINAL MUSCULAR ATROPHY THROUGH QPCR METHOD



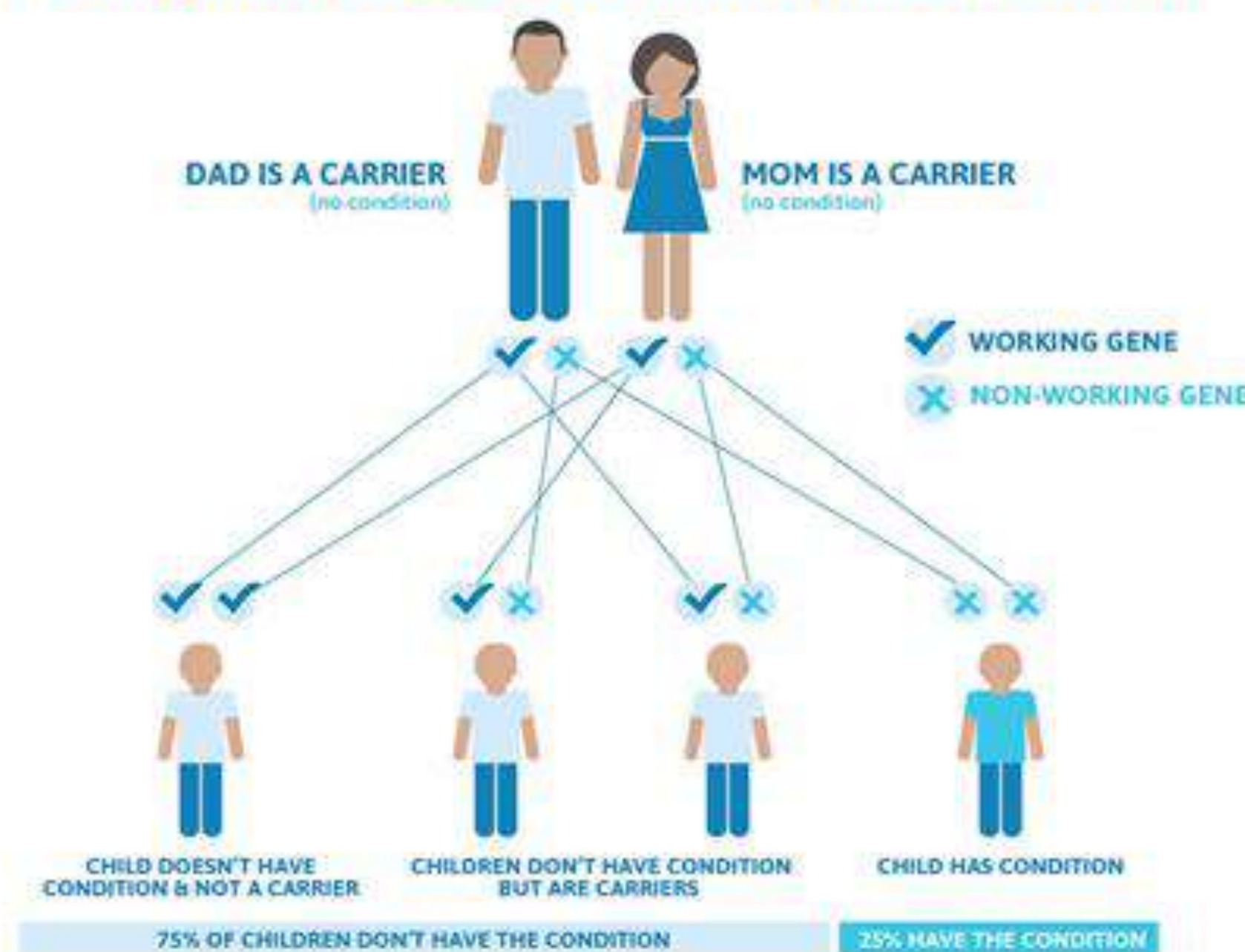
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### Introduction

Spinal muscular atrophy (SMA) is a progressive neuromuscular disease inherited in an autosomal recessive way. The prevalence of SMA in the RM constitutes  $8.43 \pm 0,15:100000$  population. 95% of SMA is caused by deletion of exon 7 of SMN1. In carrier couples there is a 25% chance of offspring with SMA.

### Autosomal Recessive Inheritance Pattern



### Purpose

Diagnosis of SMA through qPCR method (caused by deletion of exon 7 SMN1) in Human Molecular Genetics Lab. This will reduce the time of diagnosis and offer the possibility to identify the carriers of deletion.

### Material and methods.

60 DNAs representing 15 couples from control group and 10 families (mother, father and child affected or suspected) were diagnosed for determining the status of exon 7 SMN1 by qPCR method, melting curve (2 replicates for SMN1 exon 7, 1 replicate for the ALB exon 12). The DNA concentration was measured by spectrophotometry. EvaGreen was used as a DNA-binding dye.

**Keywords:** molecular genetics, diagnosis, method, screening, SMA

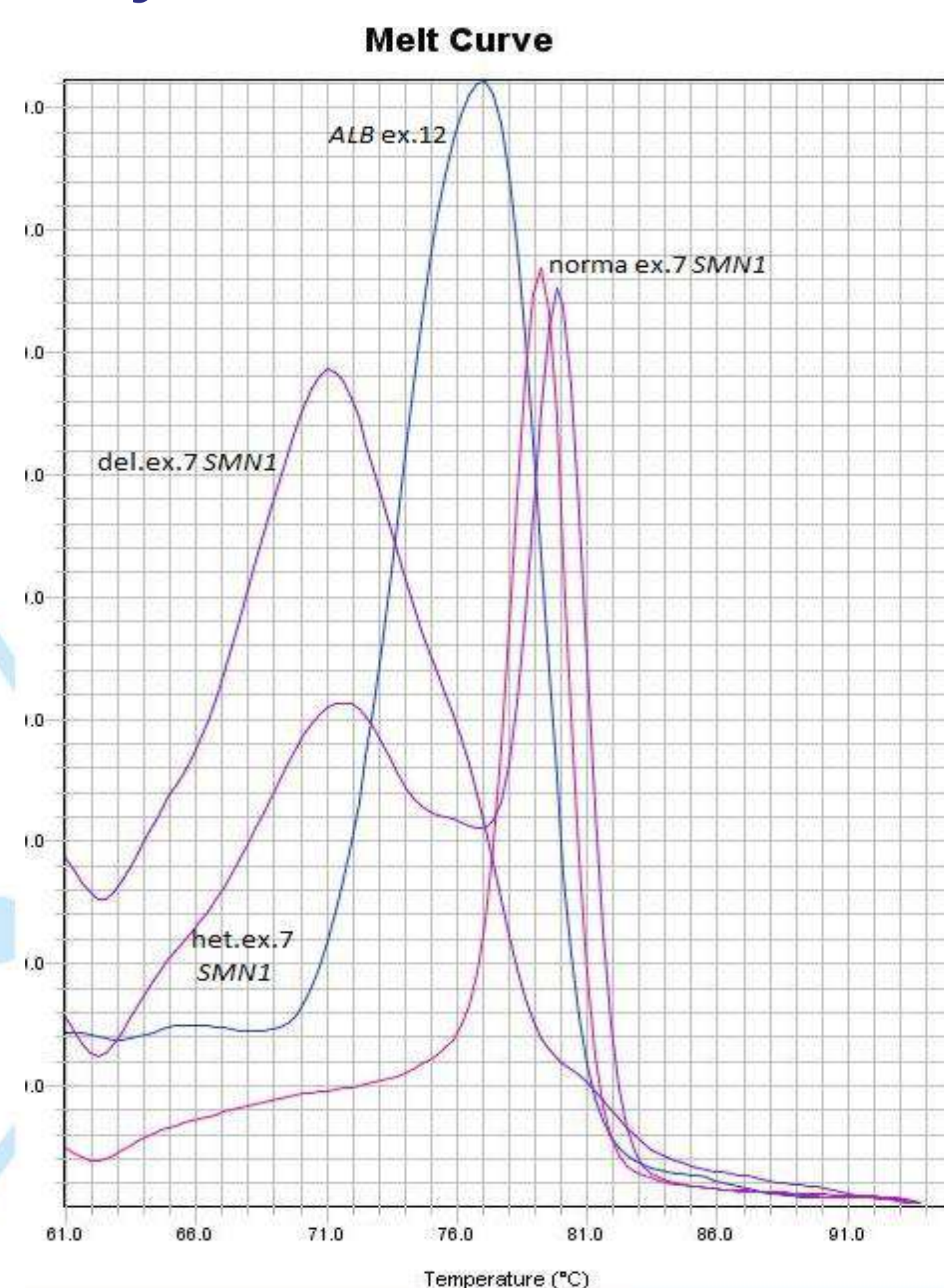


Fig. 1: Melt curve profile for each status of SMN1 7 exon

### Results

Diagnosis of SMA is available through different methods. The molecular genetic diagnosis by PCR-RFLP is expensive and time consuming than qPCR method. For all DNA samples, amplification occurred for both exon 12 ALB and exon 7 SMN1. According to the melting curves, in families with history of SMA 9 DNAs with heterozygous status were identified and 7 DNA with exon deletion 12 have the status normal for exon 7 gene SMN1 and for 2 DNAs the reaction did not take place. For 22 persons from control group the exon7 SMN1 was determined to be present and for 8 persons was determined heterozygous status (5 women and 3 men). Among those who are heterozygous, 2 people form the same couple.

### Conclusions

Considering the presence of treatment the diagnosis as soon as possible is needed and QPCR is an effective method for this: prenatal for families in which the history of SMA is already present, for newborns (newborn screening) and even in the family planning process (carrier screening).

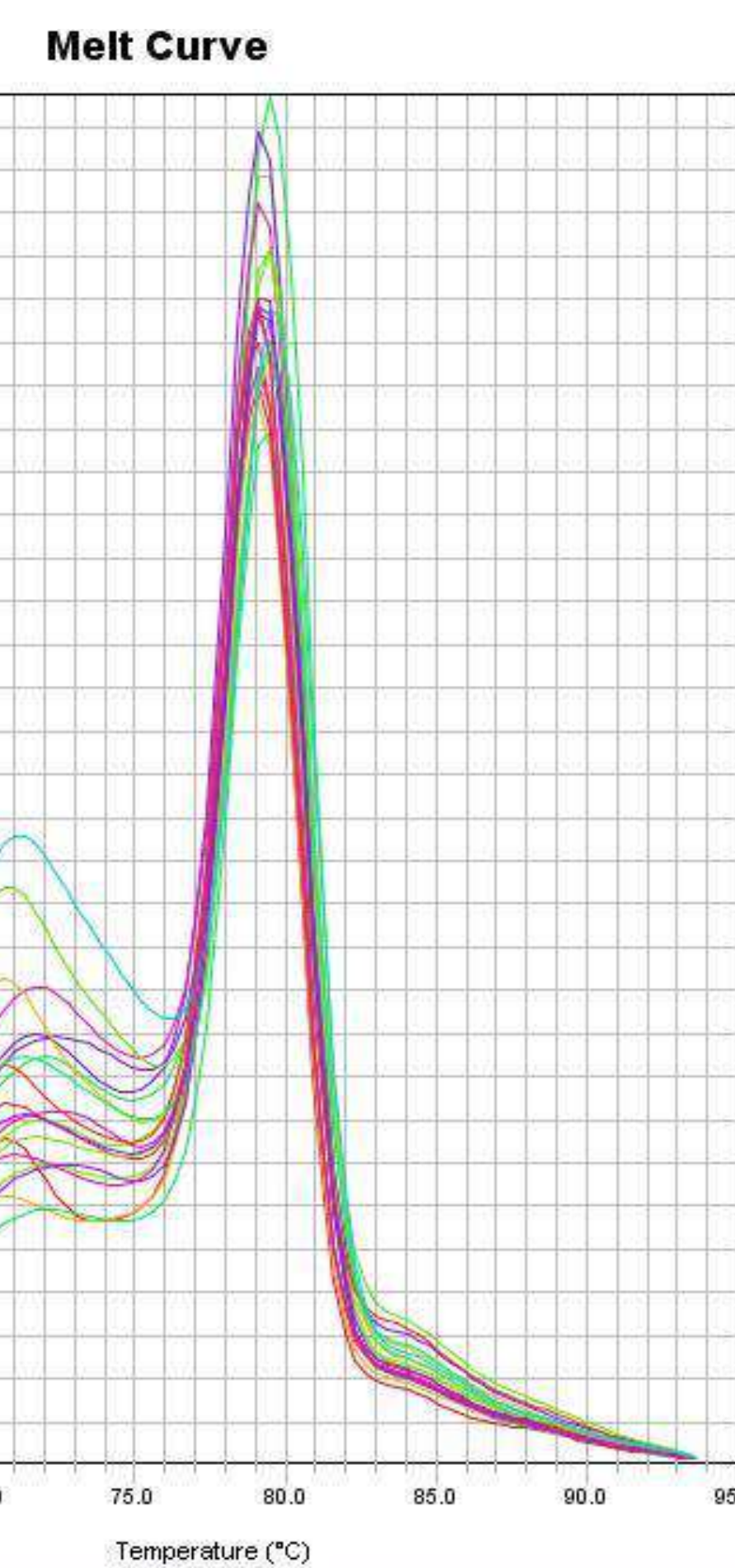


Fig. 2: Melt curve profile for normal exon 7 SMN1 status

Exon 7 status of SMN1 gene in all samples

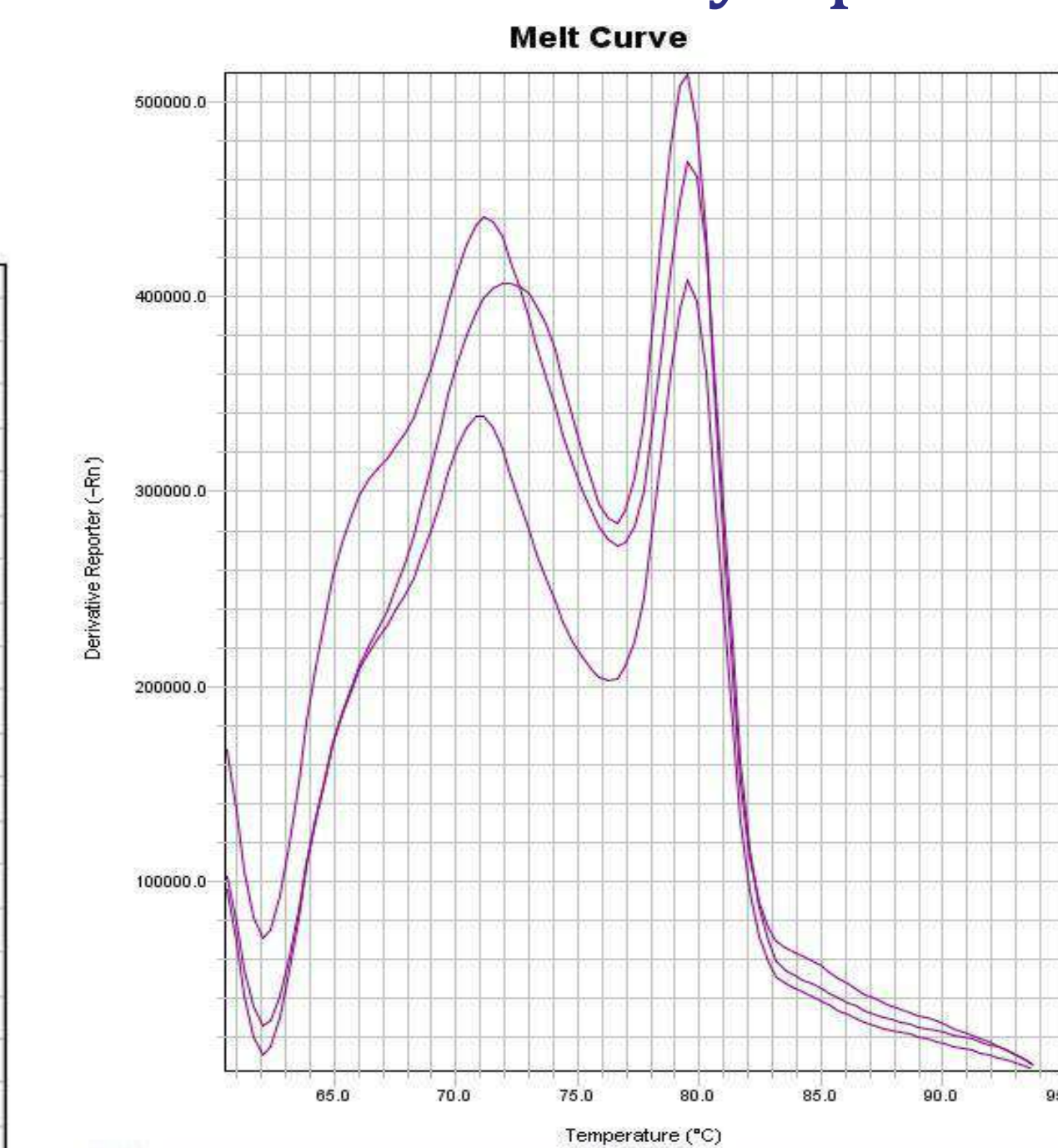
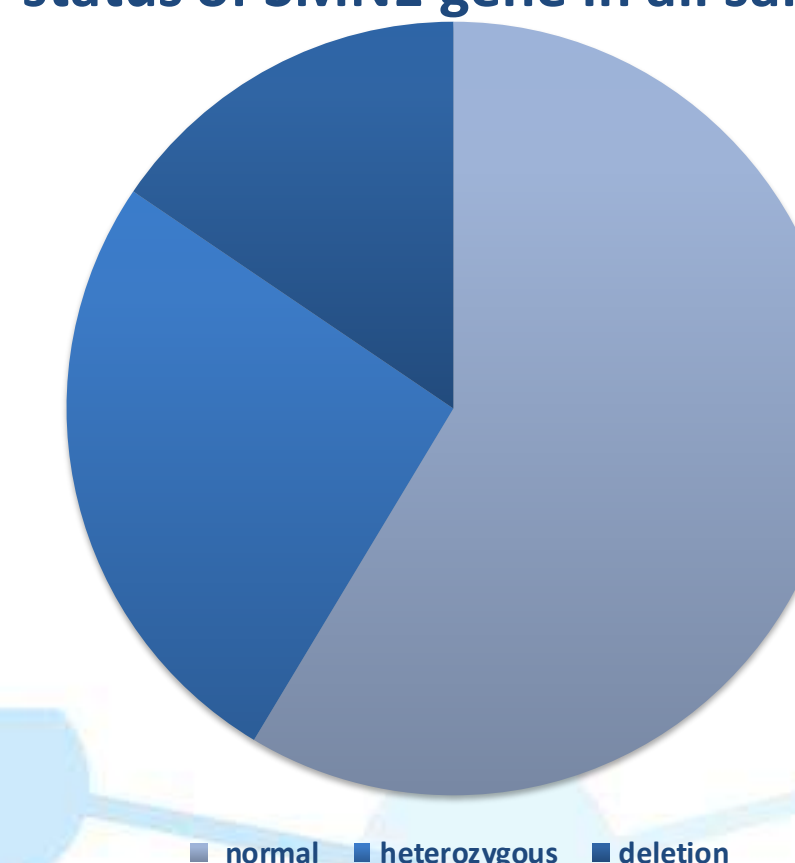


Fig. 3: Melt curve profile for SMN1 7 exon carrier deletion from couples of control group

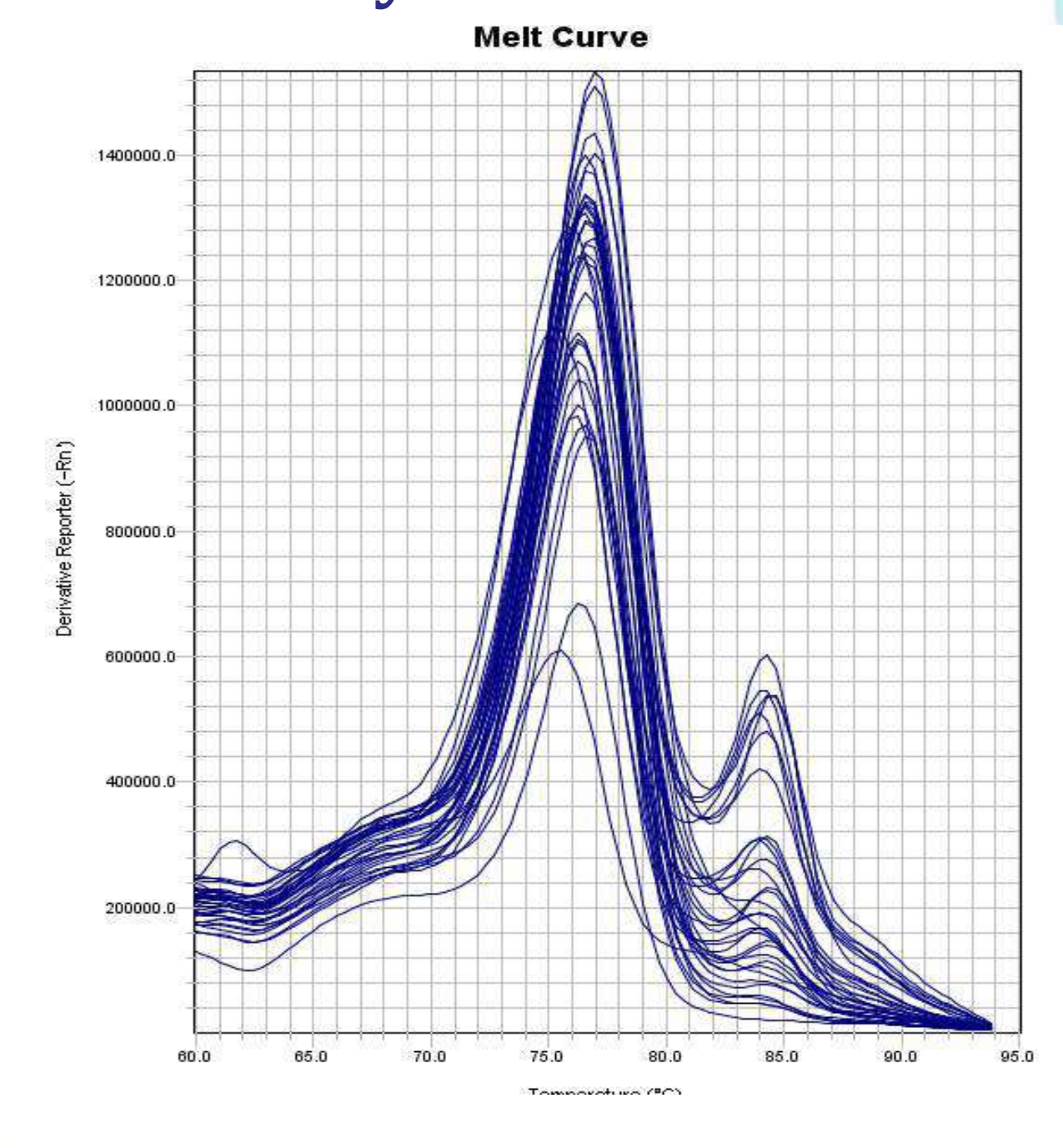


Fig. 4: Melt curve profile for ALB, 12 exon of all samples

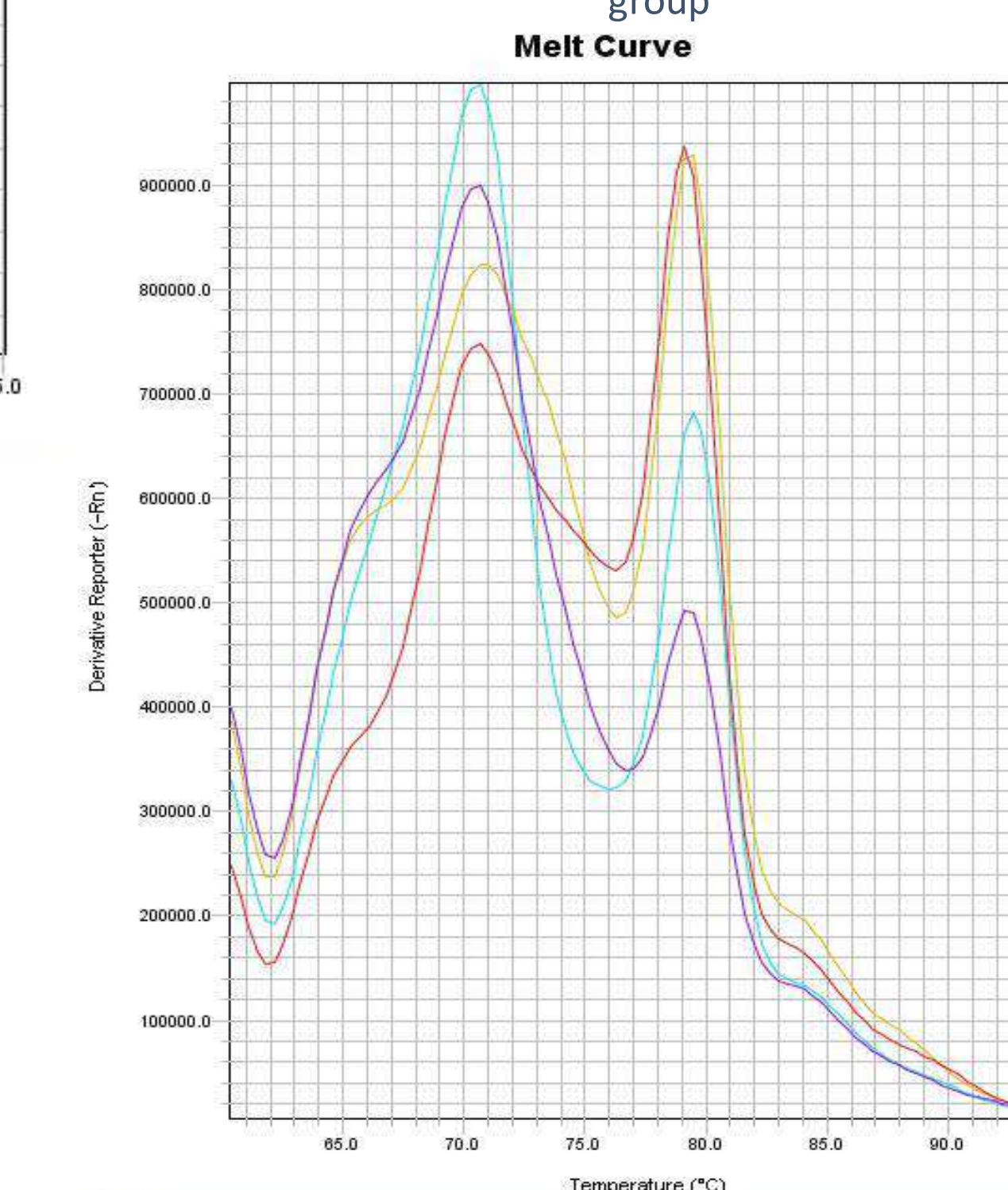


Fig. 5: Melt curve profile for SMN1 7 exon carrier deletion from families with historic of SMA

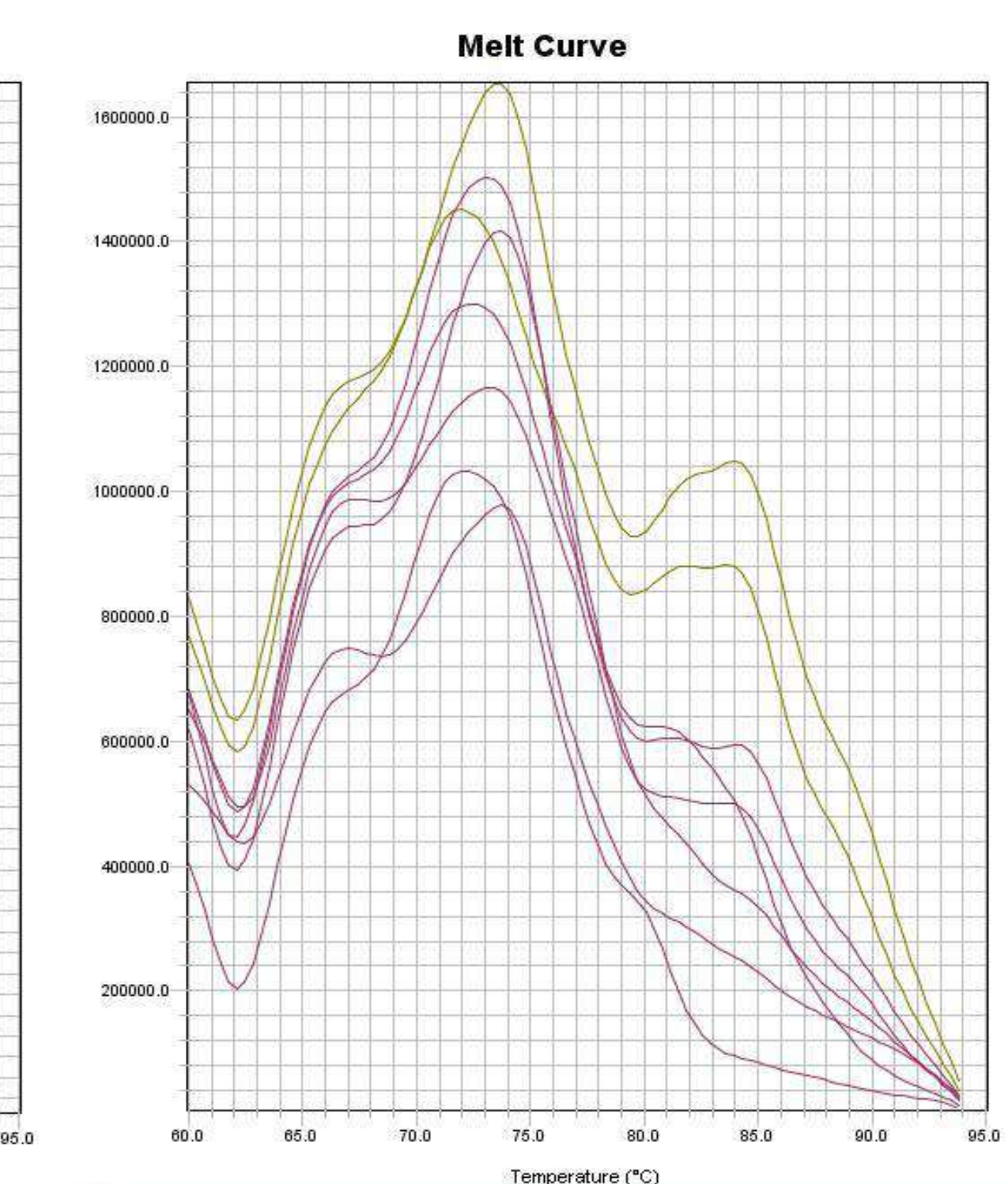


Fig. 6: Melt curve profile for SMN1 7 exon deletion of probands from families with historic of SMA