

Compound RYR1 heterozygosity with
MH-susceptibility, central cores and multiminicore disease in one family.

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❖ **Proband** : female, born 1977

First investigations at age 3 because of delayed motor milestones : 'myopathy with cores'

2004 Pt contacts centre for NMD, University of Antwerp because of child wish : new investigations

- **Clinical findings :** +/- invalidating weakness shoulder/hip girdles

- **Muscle biopsy 2004**

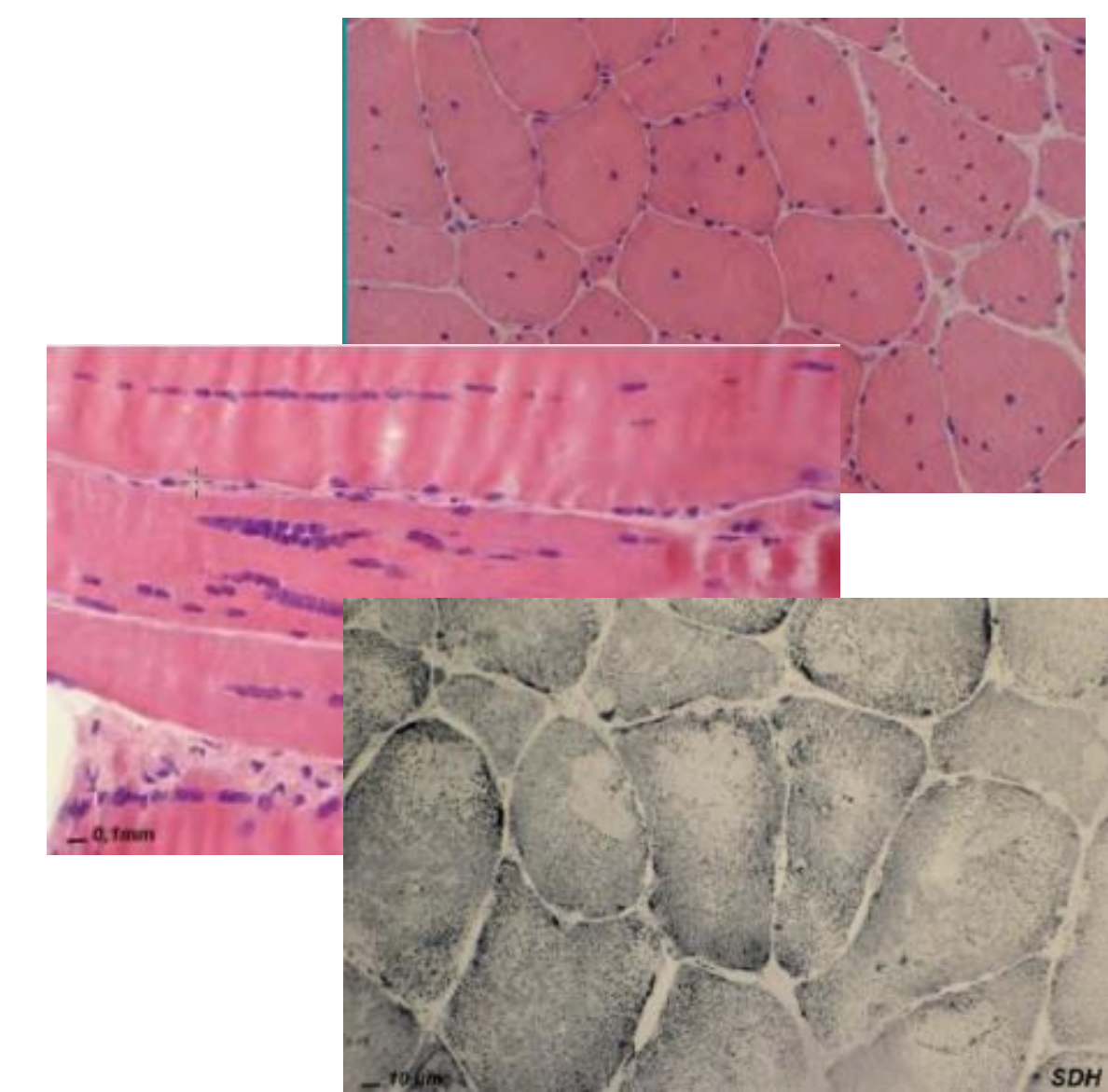
- Fatty infiltration, fibrosis, necrotic fibres, centralization of several nuclei in numerous fibres
- Almost type 1 fibre 'uniformity'
- EM : multiple core-like myofibrillar alterations corresponding to 'multi-minicore type lesions, focal
- loss of striations, rods in atrophic fibres
- Conclusion : **"Multi-minicore disease + rods"**

- **IVCT 2004:** 16 mN at 2 mM caff, 35 mN at 2% Hal, 20 mN at 75 microM 4CmC : **MHS**

- **106 exon RYR1 sequencing 2005** CHU Grenoble **proband** : 20 polymorphisms, 4 'variants of unknown significance including p.Val4849Ile , no 'causative mutations'.

- **Western blot analysis Basel 2013 : 50% less RYR1 protein** compared to control

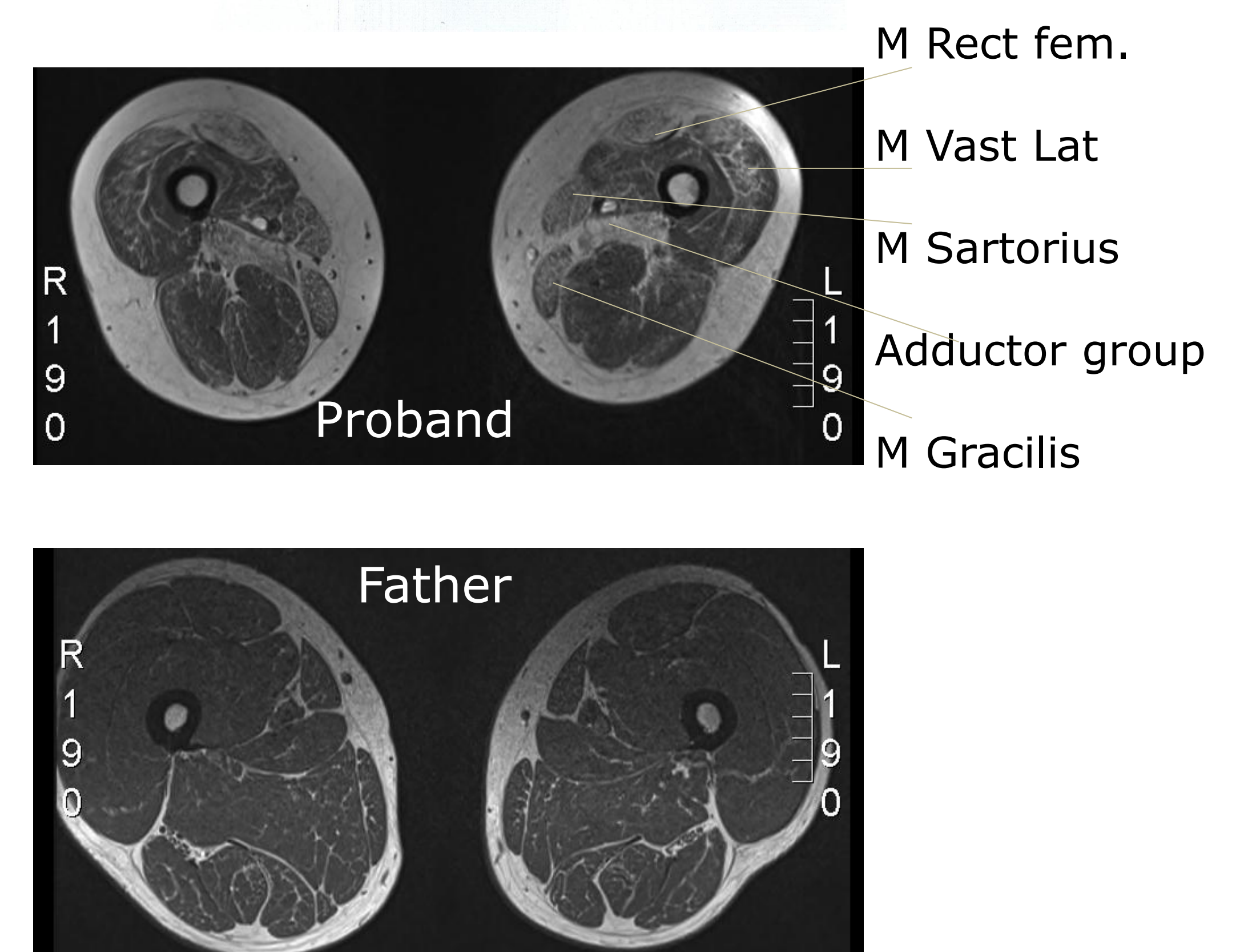
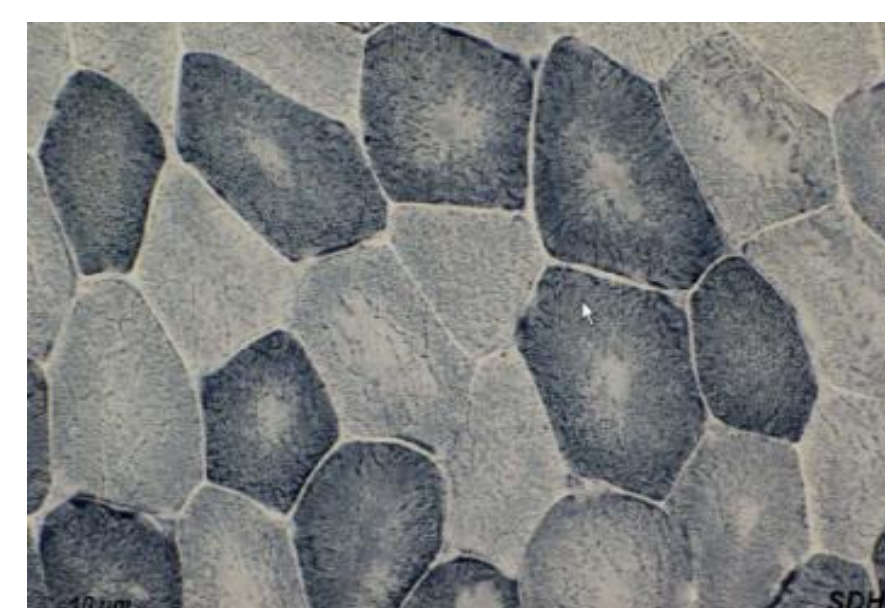
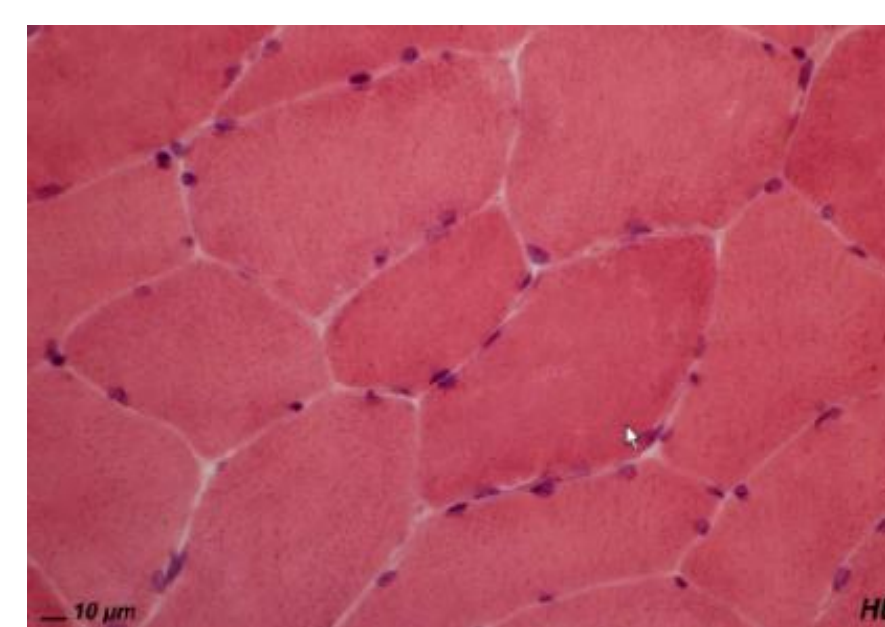
- **Muscle MRI 2013** : symm T1-weighted signal abnormalities in select muscle



fragme nt	AA	Variant +/-	Poly morphisms +/-
01	1-138	-	Leu18Leu
02	123-138	-	-
03	315-500	-	Ala359ala
04	470-709	-	Ser565ser
05	860-844	-	Pro56pro
06	824-1049	-	His579his, Thr198Thr
07	992-1020	-	Ile1152ile
08	1154-1347	-	-
09	1333-1352	-	-
10	1517-1713	Ile157Ival	-
11	1950-1918	-	-
12	187-1011	-	Gly200Gly, Lys203Lys
13	2054-2536	-	Gly200Gly
14	2213-2487	-	-
15	2415-2592	-	Arg262ala
16	2561-2707	-	Thr285Thr, Ile270Ile, Asp273Asp, Asp273Glu, Glu277Glu
17	2720-2912	-	-
18	2699-3101	-	Ser2630ser
19	3052-3559	-	Pro30pro
20	3240-3442	Arg336His	Glu332Glu
21	3442-3609	-	-
22	3593-3784	-	Glu3583Gln
23	3768-3948	Thr393Gly	-
24	3938-4154	-	-
25	4112-4234	-	-
26	4232-4471	-	-
27	4424-4641	-	-
28	4468-4931	-	-
29	4629-4808	-	-
30	4803-4973	Val484Ile	-
31	5033-5041	-	-

❖ **Father of proband**

- **No** clinical symptoms/complaints
- Histology : Normal fibre type distribution (1 & 2)
- No centralization of nuclei. **Central cores** on histology
- IVCT : 16 mN at 2 mM caff, 35 mN at 2% Hal, 20 mN at 75 microM 4CmC : **MHS**
- MRI : normal
- **Western blot analysis** : RYR1 **equal** to control
- **106 exon RYR1 sequencing**
 - variant p.Val4849Ile , polymorphism p.Glu3583Gln.

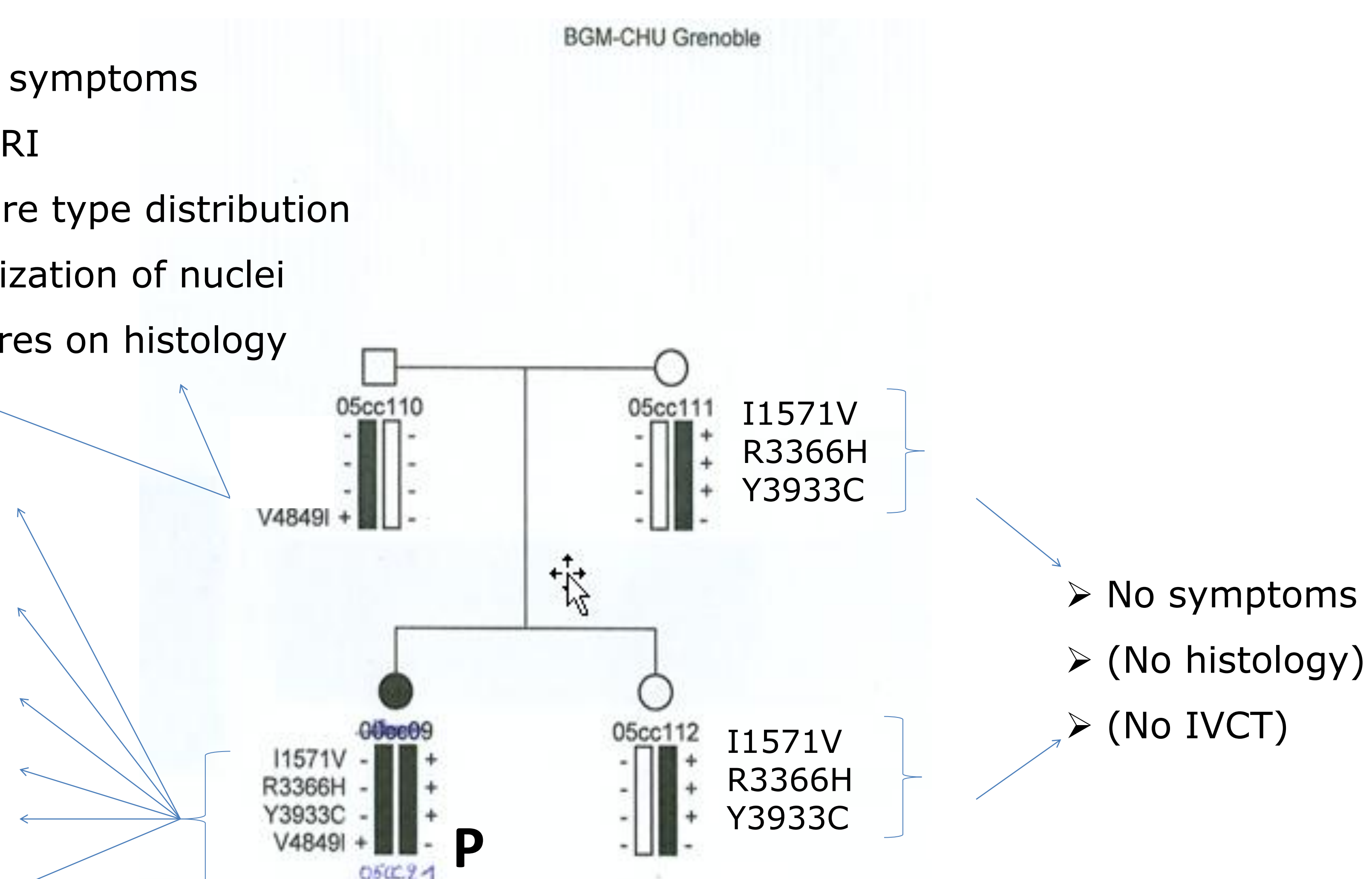


❖ **Mother and sister of proband** : no consent for investigations except DNA testing

- variants p.Ile1571Val, p.Arg3366His, p.Tyr3933Cys polymorphism p.Gly2060Cys.

Present assessment

- Final assessment**
- No clinical symptoms
 - Normal MRI
 - Normal fibre type distribution
 - No centralization of nuclei
 - Central cores on histology
 - MHS
- Invalidating girdle weakness
- Selective muscle involvement on MRI
- Histological findings of myopathy
- Multi-minicore disease + rods
- 50% less RYR1 expression
- MHS
-
- V4849I +
- I1571V
R3366H
Y3933C
V4849I



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