

Clinical and genetic profiles of 168 Russian patients with hypophosphatemic rickets.

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Introduction: The aim of this study was to clinically characterize and perform genetic analysis of 168 cases with HR.

Patients:

- n= 168 patients with the diagnosis of HR
- age, from 1 months to 56 years
- female n=111, male n=57
- 52 familial and 116 sporadic cases from 137 families

Methods:

1. Custom Ion AmpliSeq™ «Rickets panel» gene panel:

- 22 genes (*ALPL*, *ATP6V0A4*, *ATP6V1B1*, *CASR*, *CLCN5*, *CLCNKB*, *CYP24A1*, *CYP27B1*, *CYP2R1*, *DMP1*, *ENPP1*, *FGF23*, *GALNT3*, *KL*, *LRP5*, *PHEX*, *PTHR1*, *SLC2A2*, *SLC34A1*, *SLC34A3*, *SLC9A3R1*, *VDR*);

- ~83 kb, 409 amplicons, coverage 98,5%.

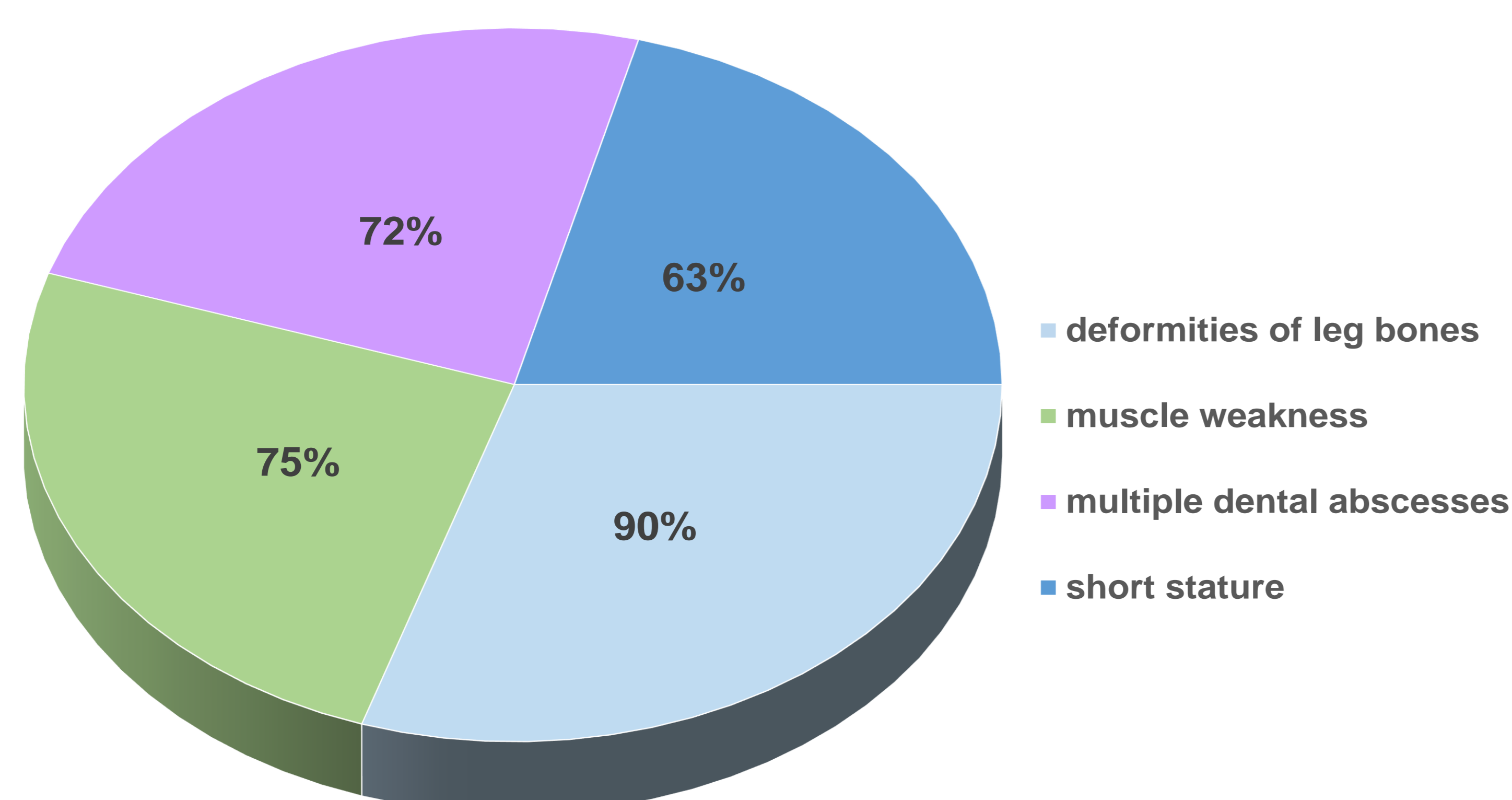
2. PGM semiconductor sequencer (Ion Torrent, Life Technologies).

3. Bioinformatics: Torrent Suite (Ion Torrent, Life Technologies), ANNOVAR version 2013 Feb21 (annovar.openbioinformatics.org)

Results:

- The mean age at diagnosis was 7.5 years (aged from 2 month to 17 years).

Clinical symptoms of HR



- The mean height SDS, $-2.3 \pm 1,8$
- The mean RSS= 4.5 points (range: 1.5-10)



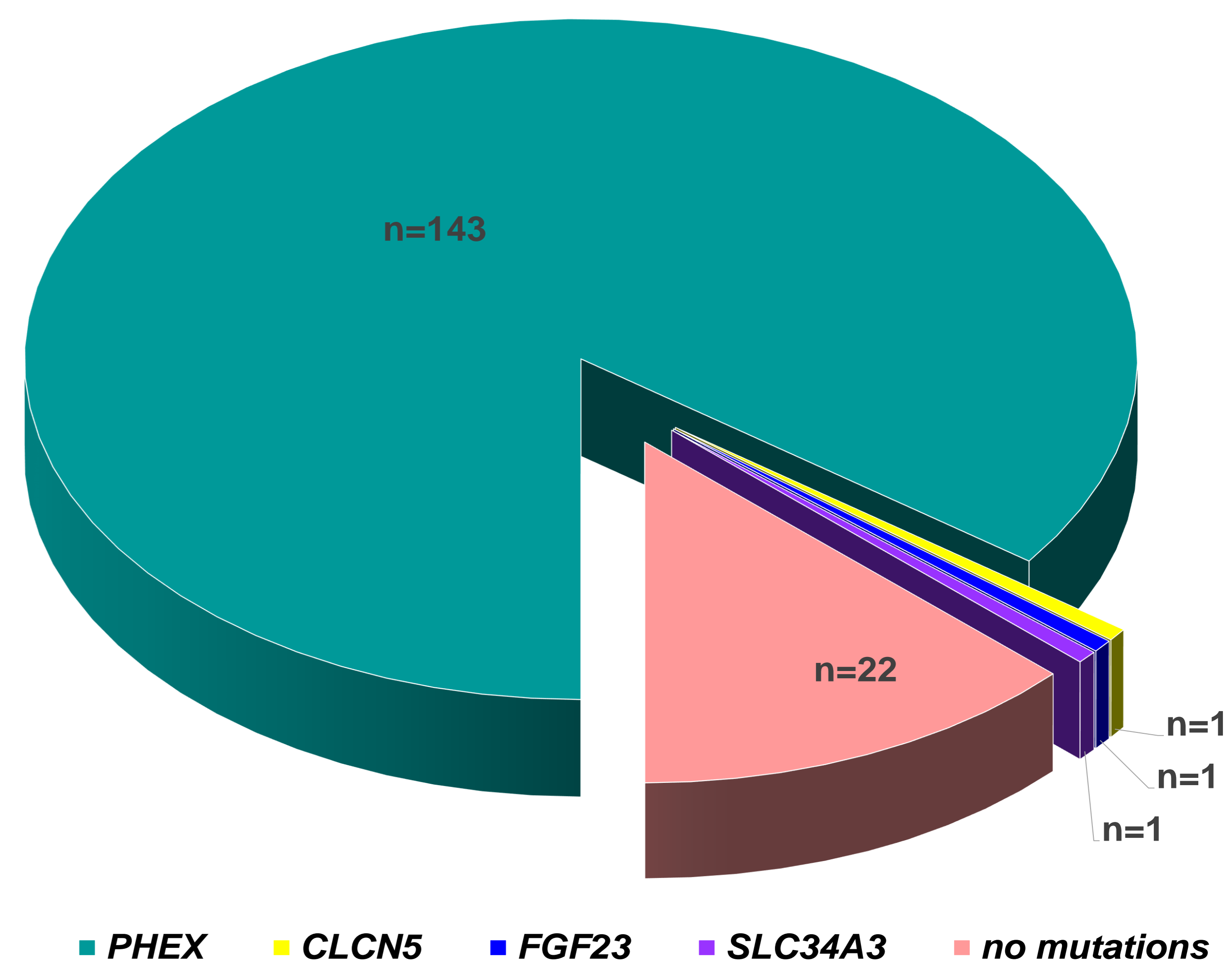
Figure 1. Radiographs of wrists and knees of a 3-year-old male with severe rickets (RSS = 10 points).

Financial support



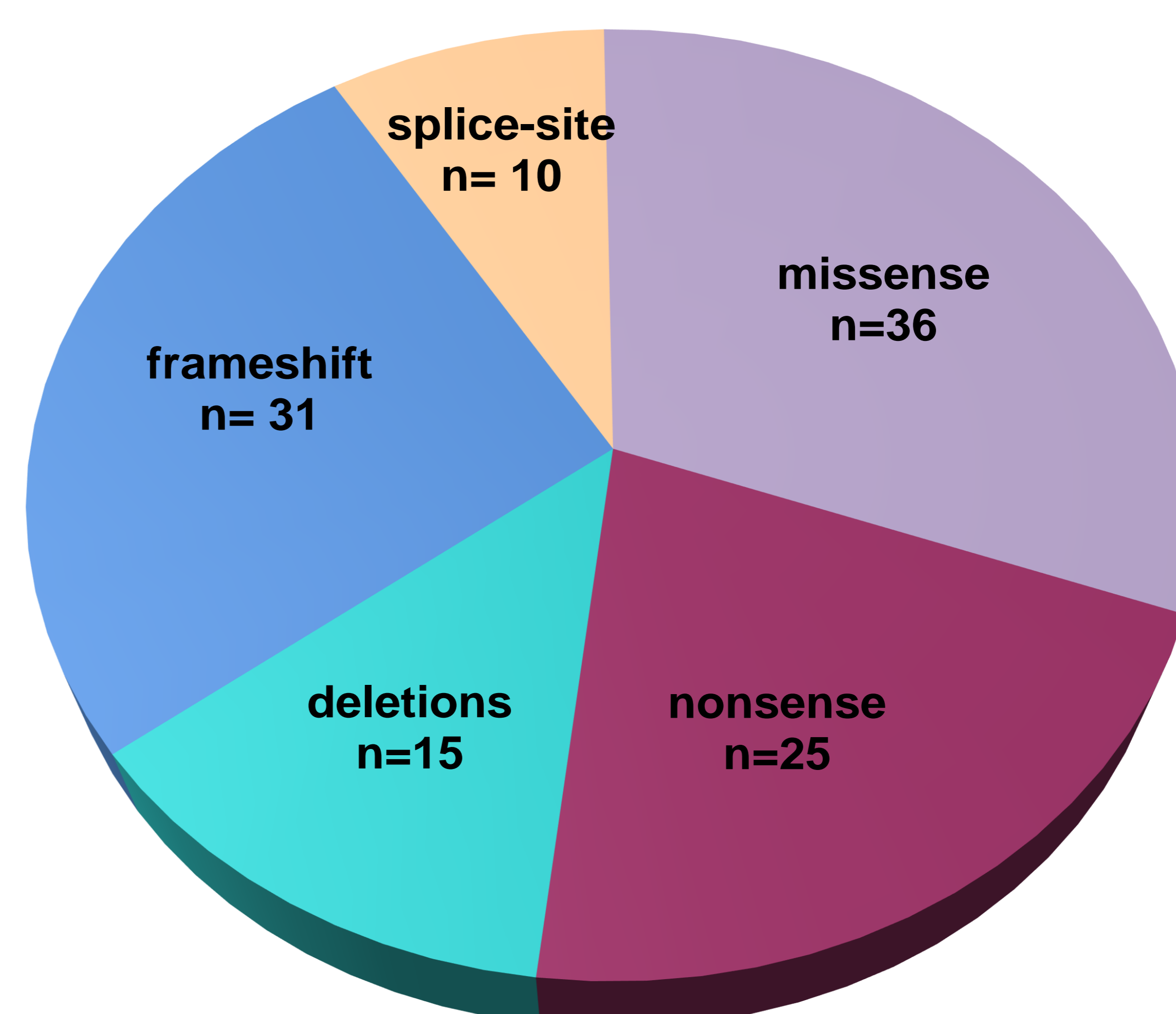
- Mutations were identified in 92.3% of familial and 84.4% of sporadic cases.

Spectrum of mutations



- In 143 probands mutations were detected in *PHEX*, 70 of which were novel.

Spectrum of *PHEX* mutations (n=117)



Conclusion:

- This study showed the predominance of *PHEX* mutations among the patients with HR in Russia.
- We identified 117 *PHEX* mutations, including 70 unreported ones.
- Earlier diagnostic and treatment can result in less severe complications and improve the quality of life of patients with HR.

