Suppl. Table S3: Cost estimate by the authors. Calculation of costs is based on the premise that the laboratory already has all of the necessary instruments and analysis software.

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| **Method** | **Purpose** | **Cost estimate per sample** |
| NGS/WES based gene panel | 1st line routine, SNV and CNV analysis | ~ $300 depending on target size and sequencing device |
| MLPA *CFH* gene cluster | 2nd line CNV detection  Nota bene: Exon coverage is incomplete with the commercially available MLPA kit (P236, MRC Holland, Amsterdam, Netherlands) | ~ $60 including costs for sex matched MLPA control sample.  Gene rearrangements without copy number changes will not be identified by MLPA |
| Molecular combing  of *CFH* gene cluster | 2nd line SV detection in unsolved cases or those cases where more precise information of haplotypes is needed and NGS/MLPA analyses is insufficient to reconstruct both alleles | ~ $50 including HMW DNA extraction, non-recurring investment in fluorescent probe design needs to be done in addition |
| Target enrichment (Samplix) and long-read sequencing  (ONT or PacBio) | validation of conspicuous SV findings | ~ $220 enrichment and ~ $220 sequencing costs (depends on the pooling strategies, a long read WGS would be ~ $1700) |
| PCR and Sanger sequencing | any validation or  segregation within families of SNV or SV (breakpoints) | less than $15 |