

| Disease/Condition name                | No of detectable genes/markers | Genes  | Method              | Price (EUR)  |
|---------------------------------------|--------------------------------|--|---------------------|--------------|
| Alpha-Thalassemia                     | 2/7                            | HBA1, HBA2   | PCR                 | 91           |
|                                       | 2                              | HBA1, HBA2   | MLPA                | 310          |
| Beta-Thalassemia, Sickle Cell Disease | 1                              | HBB  | Sanger Sequencing   | 257          |
|                                       |                                |  | MLPA                | 310          |
| Coagulation Disorders                 | 17                             | F2, F5, F8, F9, F10, F11, F12, F13A1, FGA, FGB, FGG, GGCX, LMAN1, MCFD2, SERPINC1, VKORC1, VWF                         | NGS                 | 1030         |
|                                       |                                |  | NGS with CNV        | 1400         |
| Fanconi Anemia                        | 17                             | BRCA2, BRIP1, ERCC4, FANCA, FANCB, FANCC, FANCD2, FANCE, FANCF, FANCG, FANCI, FANCL, FANCM, PALB2, RAD51C, SLX4, XRCC2 | NGS<br>NGS with CNV | 1030<br>1400 |
|                                       | 4                              | FANCA, FANCB, FANCD2, PALB2  | MLPA                | 1070         |
| Hereditary Sideroblastic Anemia       | 9                              | ABCB7, ALAS2, GLRX5, HSPA9, PUS1, SLC19A2, SLC25A38, TRNT1, YARS2  | NGS<br>NGS with CNV | 1030<br>1400 |
| Thrombocytopenia                      | 14                             | ADAMTS13, ANKRD26, CYCS, GATA1, GP1BA, GP1BB, GP9, ITGA2B, ITGB3, MASTL, MPL, MYH9, RUNX1, WAS                         | NGS                 | 1030         |
|                                       |                                |  | NGS with CNV        | 1400         |
| Whole Exome Sequencing (WES)          |                                | Solo sample  | NGS                 | 1300         |
|                                       |                                |  | NGS with CNV        | 1700         |
|                                       |                                | Trio samples   | NGS                 | 2600         |
|                                       |                                |  | NGS with CNV        | 3400         |
| Whole Genome Sequencing (WGS)         |                                | Solo sample  | NGS                 | 3635         |
|                                       |                                | Trio samples   |                     | 7449         |

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### Service includes

- DNA extraction
- Genotyping
- Bioinformatic analysis of detected variants
- Detected variants available in Excel format (on request)
- Validation of phenotype/diagnosis associated known or possibly pathogenic variants
- Clinical and biological interpretation
- Results delivery by registered mail (on request)
- TAT 2-9 weeks, excl. WES 10 weeks and WGS 15 weeks

Online ordering, specimen requirements and shipping instructions are available at [www.asperbio.com](http://www.asperbio.com).