

| Disease/Condition name | Method | No of detectable genes/markers | Genes | Price (EUR) |
|---|-------------------|--------------------------------|--|-------------|
| Citrin Deficiency | NGS | 1 | SLC25A13 | 803 |
| Citrullinemia, type 1 | NGS | 21 | ASS1 | 803 |
| Fatty Acid Oxidation Disorder | NGS | 21 | ACAD9, ACADM, ACADS, ACADVL, CPT1A, CPT2, ETFA, ETFB, ETFDH, GLUD1, HADH, HADHA, HADHB, HMGCL, HMGCS2, HSD17B10, LPIN1, PPARG, SLC22A5, SLC25A20, TAZ | 1070 |
| | MLPA | 1 | SLC22A5 | 310 |
| Glutaric Aciduria, type 1 | Sanger Sequencing | 1 | GCDH | 535 |
| Glutaric Aciduria, type 2 | Sanger Sequencing | 1 | EFTA | 535 |
| Glycogen Storage Disease | NGS | 25 | AGL, ALDOA, ENO3, FBP1, G6PC, GAA, GBE1, GYG1, GYS1, GYS2, LAMP2, LDHA, PFKM, PGAM2, PGK1, PGM1, PHKA1, PHKA2, PHKB, PHKG2, PRKAG2, PYGL, PYGM, SLC2A2, SLC37A4 | 1070 |
| Hemochromatosis | NGS | 5 | HAMP, HFE, HJV, SLC40A1, TFR2 | 1070 |
| Lysosomal Storage Disease | NGS | 50 | AGA, ARSA, ARSB, ASAHI, CLN3, CLN5, CLN6, CLN8, CTNS, CTSA, CTSC, CTSD, CTSK, DNAJC5, FUCA1, GAA, GALC, GALNS, GBA, GLA, GLB1, GM2A, GNPTAB, GNPTG, GNS, GUSB, HEXA, HEXB, HGSNAT, HYAL1, IDS, IDUA, LAMP2, LIPA, MAN2B1, MANBA, MCOLN1, MFSD8, NAGA, NAGLU, NEU1, NPC1, NPC2, PPT1, PSAP, SGSH, SLC17A5, SMPD1, SUMF1, TPP1 | 1338 |
| | MLPA | 1 | GLA | 310 |
| | MLPA | 1 | IDS | 310 |
| | MLPA | 2 | NPC1, NPC2 | 310 |
| | MLPA | 1 | HEXA | 310 |
| Metabolic Myopathy and Rhabdomyolysis | NGS | 44 | ABHD5, ACAD9, ACADM, ACADVL, AGL, ALDOA, AMPD1, CAV3, CPT2, ENO3, ETFA, ETFB, ETFDH, GAA, GBE1, GYG1, GYS1, HADHA, HADHB, ISCU, LDHA, LPIN1, OPA1, OPA3, PFKM, PGAM2, PGK1, PGM1, PHKA1, PNPLA2, PRKAG2, POLG, POLG2, PYGM, RRM2B, RYR1, SCN4A, SLC22A5, SLC25A20, SUCLA2, TAZ, TK2, TWNK, TYMP | 1070 |
| Methylmalonic Aciduria and Homocystinuria | NGS | 23 | ABCD4, ACSF3, AMN, CBS, CD320, CUBN, GIF, IVD, LMBRD1, MCEE, MLYCD, MMAA, MMAB, MMACHC, MMADHC, MTHFR, MTR, MTRR, MUT, SUCLA2, SUCLG1, TCN1, TCN2 | 1070 |
| | MLPA | 1 | MLYCD | 310 |

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|---|-------------------|--------------------------------|---|-------------|
| Porphyria | NGS | 9 | ALAD, ALAS2, CPOX, FECH, HFE, HMBS, PPOX, UROD, UROS | 1070 |
| | MLPA | 7 | ALAD, CPOX, FECH, HMBS, PPOX, UROD, UROS | 710 |
| Short-Chain Acyl-CoA Dehydrogenase (SCAD) Deficiency | Sanger Sequencing | 1 | ACADS | 535 |
| Smith Lemli Opitz Syndrome | Sanger Sequencing | 1 | DHCR7 | 525 |
| Urea Cycle Disorder | NGS | 10 | ARG1, ASL, ASS1, CPS1, NAGS, OAT, OTC, SLC7A7, SLC25A13, SLC25A15 | 1070 |
| | MLPA | 1 | OTC | 310 |
| Very Long Chain Acyl-CoA Dehydrogenase (VLCAD) Deficiency | NGS | 1 | ACADVL | 803 |
| Whole Exome Sequencing (WES) | NGS | | Solo sample | 1567 |
| | | | Trio samples | 3100 |
| Whole Genome Sequencing (WGS) | NGS | | Solo sample | 4500 |
| | | | Trio samples | 8800 |

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 and WGS 15 weeks

Online ordering, specimen requirements and shipping instructions are available at www.asperbio.com.