

**Supplementary Table S1:** Mitochondrial gene-sets

<b>Citrate cycle (TCA cycle):</b> ACLY, ACO1, ACO2, CS, DLAT, DLD, DLST, FH, IDH1, IDH2, IDH3A, IDH3B, IDH3G, MDH1, MDH2, OGDH, OGDHL, PC, PCK2, PDHA1, PDHA2, PDHB, PDHX, SDHA, SDHB, SDHC, SDHD, SUCLA2, SUCLG1, SUCLG2
<b>Fatty acid biosynthesis:</b> ACACA, ACACB, FASN, MCAT, OXSM
<b>Fatty acid elongation:</b> ACAA2, ECHS1, HADH, HADHA, HADHB, MECR
<b>Fatty acid metabolism:</b> ACAA1, ACAA2, ACADL, ACADM, ACADS, ACADSB, ACADVL, ACAT1, ACOX3, ACSL1, ACSL4, ACSL5, ALDH1B1, ALDH2, ALDH3A2, ALDH7A1, ALDH9A1, CPT1A, CPT1B, CPT1C, CPT2, ECHS1, ECI1, ECI2, EHHADH, GCDH, HADH, HADHA, HADHB
<b>Synthesis and degradation of ketone bodies:</b> ACAT1, BDH1, HMGCL, HMGCS2, OXCT1
<b>Ubiquinone and other terpenoid-quinone biosynthesis:</b> COQ2, COQ3, COQ5, COQ6, COQ7
<b>Oxidative phosphorylation:</b> ATP5A1, ATP5B, ATP5C1, ATP5D, ATP5F1, ATP5G1, ATP5G2, ATP5G3, ATP5I, ATP5J, ATP5J2, ATP5L, ATP5O, ATP6, COX1, COX10, COX11, COX15, COX17, COX2, COX3, COX4I1, COX4I2, COX5A, COX5B, COX6A1, COX6A2, COX6B1, COX6B2, COX6C, COX7A1, COX7A2, COX7A2L, COX7B, COX7C, COX8A, CYC1, CYTB, ND1, ND2, ND3, ND4, ND4L, ND5, ND6, NDUFA1, NDUFA10, NDUFA12, NDUFA13, NDUFA2, NDUFA3, NDUFA4, NDUFA5, NDUFA6, NDUFA7, NDUFA8, NDUFA9, NDUFA10, NDUFB1, NDUFB10, NDUFB11, NDUFB2, NDUFB3, NDUFB5, NDUFB6, NDUFB7, NDUFB8, NDUFB9, NDUFC1, NDUFC2, NDUFS1, NDUFS2, NDUFS3, NDUFS4, NDUFS5, NDUFS7, NDUFS8, NDUFV1, NDUFV2, NDUFV3, PPA2, SDHA, SDHB, SDHC, SDHD, TCIRG1, UQCR10, UQCR11, UQCRC1, UQCRC2, UQCRCFS1, UQCRRH, UQCRRQ
<b>Pyrimidine metabolism:</b> AK3, CTPS2, DHODH, DUT, NME1, NME1-NME2, NME2, NME3, NME4, NME6, NT5C, NT5C3, NT5M, NUDT2, PNPT1, TK2, TXNRD1, TXNRD2
<b>Alanine, aspartate and glutamate metabolism:</b> ABAT, ADSL, AGXT, AGXT2, ALDH4A1, ALDH5A1, CPS1, GLS, GLS2, GLUD1, GOT2, GPT2
<b>Glycine, serine and threonine metabolism:</b> AGXT, AGXT2, ALAS1, ALAS2, ALDH7A1, AMT, CHDH, DLD, DMGDH, GATM, GCAT, GLDC, GLYCTK, MAOA, MAOB, SARDH, SHMT1, SHMT2
<b>Valine, leucine and isoleucine degradation:</b> ABAT, ACAA1, ACAA2, ACAD8, ACADM, ACADS, ACADSB, ACAT1, ALDH1B1, ALDH2, ALDH3A2, ALDH6A1, ALDH7A1, ALDH9A1, AUH, BCAT1, BCAT2, BCKDHA, BCKDHB, DBT, DLD, ECHS1, EHHADH, HADH, HADHA, HADHB, HIBADH, HIBCH, HMGCL, HMGCS2, HSD17B10, IVD, MCCC1, MCCC2, MCEE, MUT, OXCT1, PCCA, PCCB
<b>Valine, leucine and isoleucine biosynthesis:</b> BCAT1, BCAT2, IARS2, LARS2, PDHA1, PDHA2, PDHB, VARS2
<b>Lysine degradation:</b> AADAT, AASS, ACAT1, ALDH1B1, ALDH2, ALDH3A2, ALDH7A1, ALDH9A1, BBOX1, DLST, ECHS1, EHHADH, GCDH, HADH, HADHA, OGDH, OGDHL
<b>Arginine and proline metabolism:</b> AGMAT, ALDH18A1, ALDH1B1, ALDH2, ALDH3A2, ALDH4A1, ALDH7A1, ALDH9A1, ARG2, CKMT2, CPS1, GATM, GLS, GLS2, GLUD1, GOT2, LAP3, MAOA, MAOB, NAGS, OAT, OTC, PRODH, PRODH2, PYCR1
<b>Tryptophan metabolism:</b> AADAT, ACAT1, ALDH1B1, ALDH2, ALDH3A2, ALDH7A1, ALDH9A1, CCBL2, ECHS1, EHHADH, GCDH, HADH, HADHA, KYNU, MAOA, MAOB, OGDH, OGDHL, WARS2
<b>beta-Alanine metabolism:</b> ABAT, ACADM, ALDH1B1, ALDH2, ALDH3A2, ALDH7A1, ALDH9A1, ECHS1, EHHADH, HADHA, HIBCH, MLYCD
<b>Pyruvate metabolism:</b> ACACA, ACACB, ACAT1, ACSS1, ACYP2, ALDH1B1, ALDH2, ALDH3A2, ALDH7A1, ALDH9A1, DLAT, DLD, HAGH, LDHA, LDHAL6B, LDHB, LDHD, MDH1, MDH2, ME1, ME2, ME3, PC, PCK2, PDHA1, PDHA2, PDHB, PDHX
<b>Glyoxylate and dicarboxylate metabolism:</b> ACAT1, ACO1, ACO2, CS, GLYCTK, HAO2, MCEE, MDH1, MDH2, MUT, PCCA, PCCB
<b>Propanoate metabolism:</b> ABAT, ACACA, ACACB, ACADM, ACAT1, ACSS1, ACSS3, ALDH1B1, ALDH2, ALDH3A2, ALDH6A1, ALDH7A1, ALDH9A1, ECHS1, EHHADH, HADHA, HIBCH, LDHA, LDHAL6B, LDHB, MCEE, MLYCD, MUT, PCCA, PCCB, SUCLA2, SUCLG1, SUCLG2
<b>Butanoate metabolism:</b> ABAT, ACADS, ACAT1, ACSM1, ACSM2A, ACSM3, ACSM4, ACSM5, ALDH5A1, BDH1, ECHS1, EHHADH, HADH, HADHA, HMGCL, HMGCS2, L2HGDH, OXCT1, PDHA1, PDHA2, PDHB
<b>One carbon pool by folate:</b> ALDH1L1, ALDH1L2, AMT, ATIC, MTFMT, MTHFD1, MTHFD1L, MTHFD2, MTHFD2L, SHMT1, SHMT2
<b>Porphyrin and chlorophyll metabolism:</b> ALAS1, ALAS2, COX10, COX15, CPOX, EARS2, FECH, FTH1, FTMT, HCCS, MMAB, PPOX

**Supplementary Table S2:** PARIS-derived gene P-values for the lipid metabolism mitochondrial gene-set for primary open-angle glaucoma (POAG, 3430 cases) and normal tension glaucoma (NTG, 717 cases) versus controls (3108) in the combined GLAUGEN – NEIGHBOR dataset.

Lipid metabolism mitochondrial gene	LD block count	Gene P-value		Other pathway?
		POAG	NTG	
<i>ACOX3</i>	21	1.00	1.00	
<i>ACACB</i>	16	0.55	0.24	Carbohydrate metabolism
<i>ALDH9A1</i>	15	0.54	0.23	Carbohydrate metabolism
<i>ACSL1</i>	15	0.11	<b>0.020</b>	
<i>MCAT</i>	13	1.00	1.00	
<i>ALDH7A1</i>	13	0.34	1.00	Carbohydrate metabolism
<i>BDH1</i>	12	< 0.0001	< 0.0001	Carbohydrate metabolism
<i>ALDH1B1</i>	12	<b>0.046</b>	0.20	Carbohydrate metabolism
<i>CPT1B</i>	10	0.19	1.00	
<i>HMGCS2</i>	9	1.00	1.00	Carbohydrate metabolism
<i>EHHADH</i>	8	<b>0.044</b>	<b>0.0009</b>	Carbohydrate metabolism
<i>ECI2</i>	8	0.24	1.00	
<i>ACADS</i>	7	1.00	1.00	Carbohydrate metabolism
<i>ACACA</i>	6	0.27	0.06	Carbohydrate metabolism
<i>CPT2</i>	5	0.09	1.00	
<i>ECHS1</i>	5	<b>0.004</b>	<b>0.002</b>	Carbohydrate metabolism
<i>CPT1A</i>	5	0.13	1.00	
<i>ACAA2</i>	5	<b>0.024</b>	<b>0.0008</b>	
<i>CPT1P</i>	5	<b>0.037</b>	1.00	

Nb: Only genes with at least 5 LD blocks are presented.

**Supplementary Table S3:** PARIS-derived gene P-values for the carbohydrate metabolism mitochondrial gene-set for primary open-angle glaucoma (POAG, 3430 cases) and normal tension glaucoma (NTG, 717 cases) versus controls (3108) in the combined GLAUGEN – NEIGHBOR dataset .

Carbohydrate metabolism mitochondrial gene	LD block count	Gene P-value		Other pathway?
		POAG	NTG	
<i>ME3</i>	43	<b>0.0003</b>	0.30	
<i>ABAT</i>	31	<b>0.047</b>	0.31	
<i>SUCLG2</i>	26	0.77	1.00	
<i>PCCA</i>	25	1.00	0.45	
<i>ACSS3</i>	17	1.00	0.09	
<i>ACACB</i>	16	0.55	0.23	Lipid metabolism
<i>MLYCD</i>	16	0.27	0.25	
<i>ALDH9A1</i>	15	0.55	0.24	Lipid metabolism
<i>ACYP2</i>	14	<b>0.001</b>	0.43	
<i>ACO1</i>	14	1.00	1.00	
<i>ACSM5</i>	13	1.00	1.00	
<i>ALDH7A1</i>	13	0.36	1.00	Lipid metabolism
<i>ALDH5A1</i>	13	0.13	<b>0.024</b>	
<i>IDH3A</i>	12	0.27	1.00	
<i>BDH1</i>	12	< 0.0001	< 0.0001	Lipid metabolism
<i>ALDH1B1</i>	12	<b>0.046</b>	0.20	Lipid metabolism
<i>ACSM2A</i>	11	0.12	1.00	
<i>ACSS1</i>	11	1.00	0.00	
<i>PCHC</i>	10	1.00	1.00	
<i>HMGCS2</i>	9	1.00	1.00	Lipid metabolism
<i>HUMNDME</i>	9	0.28	0.26	
<i>SDHC</i>	8	1.00	1.00	
<i>PCK2</i>	8	1.00	1.00	
<i>HIBCH</i>	8	1.00	<b>0.025</b>	
<i>IDH1</i>	8	0.14	0.12	
<i>EHHADH</i>	8	0.05	0.00	Lipid metabolism
<i>HAOX2</i>	7	1.00	1.00	
<i>ACADS</i>	7	1.00	1.00	Lipid metabolism
<i>GLOD2</i>	7	1.00	1.00	
<i>IDH3B</i>	7	1.00	1.00	
<i>SDHA</i>	7	<b>0.013</b>	<b>0.012</b>	
<i>LDHAL6B</i>	6	1.00	<b>0.001</b>	
<i>IDH2</i>	6	0.12	1.00	
<i>LDHD</i>	6	1.00	<b>0.004</b>	
<i>ACACA</i>	6	0.27	0.06	Lipid metabolism
<i>PDHA2</i>	6	1.00	1.00	
<i>OGDHL</i>	5	0.11	1.00	
<i>ECHS1</i>	5	<b>0.004</b>	<b>0.003</b>	Lipid metabolism
<i>PCB</i>	5	0.02	1.00	
<i>LDHB</i>	5	1.00	<b>0.003</b>	
<i>L2HGDH</i>	5	1.00	1.00	
<i>DLTS</i>	5	<b>0.037</b>	1.00	
<i>ACSM1</i>	5	1.00	1.00	
<i>HAGH</i>	5	1.00	1.00	

NB: Only genes with at least 5 LD blocks are presented.