BCAT2 aKG-	-glutamate metabolism -glutamate metabolism -glutamate metabolism
VDAC3 VDAC1 VDAC2	VDAC VDAC VDAC VDAC
TIMM10 TIMM9 TIMM8B	Tim9-Tim10 complex Tim9-Tim10 complex Tim8-Tim13 complex
TIMM13 TIMM8A TIMM50	Tim8-Tim13 complex Tim8-Tim13 complex Tim23
TIMM23 TIMM21 TIMM17B	Tim23 Tim23 Tim23
TIMM17A TIMM10B TIMM22 TOMM40	Tim23 Tim22 Tim22 TOM complex
TOMM20 TOMM22 TOMM40L	TOM complex TOM complex TOM complex TOM complex
TOMM5 TOMM70A TOMM6	TOM complex TOM complex TOM complex
TOMM7 TMEM256 TMEM126A	TOM complex TMEM256 TMEM126A
TMEM11 TMED8 TMED1 TMED3	TMEM11 TMEDs TMEDs TMEDs
TMED3 TMED7 TMED5 TMED5 TMED9	TMEDS TMEDS TMEDS TMEDS TMEDS
TMED4 TMED2 TMED10	TMEDs TMEDs TMEDs
IDH3A IDH3G IDH3B CS	TCA TCA TCA TCA
FH MDH2 OGDH	TCA TCA TCA TCA
SUCLG2 DLD ACO2	TCA TCA TCA
SUCLG1 DLST SIRT5 SIRT3	TCA TCA SirT5 SirT3
SLC25A S LC25 family, aKG-	
SLC25A32 SLC25A1 SLC25A6	SLC25 family SLC25 family SLC25 family
SLC25A33 SLC25A29 SLC25A5	SLC25 family SLC25 family SLC25 family
SLC25A4 SLC25A30 SLC25A35	SLC25 family SLC25 family SLC25 family
SLC25A15 SLC25A10 SLC25A42	SLC25 family SLC25 family SLC25 family
SLC25A46 SLC25A40 SLC25A3 SLC25A24	SLC25 family SLC25 family SLC25 family SLC25 family
SLC25A24 SLC25A12 SLC25A26 SLC25A44	SLC25 family SLC25 family SLC25 family SLC25 family
SLC25A19 SLC25A22 SLC25A3	SLC25 family SLC25 family SLC25 family
SAMM50 MTX1 MTX2 RHOT1	SAM SAM SAM RHOT1
PRDX5 PRDX3 PHB	PRDX5 PRDX3 PHB PHB2 SLP2
STOML2 PHB2 PEX11B	PHB_PHB2_SLP2 PHB_PHB2_SLP2 PEX11B
HSPA9 TIMM44 GRPEL1 GRPEL2	PAM PAM PAM PAM
OXA1L OPA3 NME3	OXA1 OPA3 NME3
MTCH2 APOOL IMMT	MTCH2 MICOS complex MICOS complex
APOO CHCHD6 CHCHD3 QIL1	MICOS complex MICOS complex MICOS complex MICOS complex
MINOS1 CHCHD4 GFER	MICOS complex MICOS complex MIA Mean_RF_score MIA 1.0
HADH ACADVL ACADS	FAO FAO FAO 0.6
ACAD9 ECHS1 ACADSB ACADM	FAO FAO FAO FAO
ACAA2 ACAT1 HADHA	FAO 0.0 FAO FAO FAO
HADHB ERLIN2 ETFA	FAO Erlin-2 ETF system
ETFB ETFDH CYCS COX6A1	ETF system ETF system Cytochrome C Complex IV
MT-CO1 MT-CO3 COX6B1	Complex IV Complex IV Complex IV
COX7A2 COX7B COX7C COX8A	Complex IV Complex IV Complex IV Complex IV
MT-CO2 COX5B COX5A	Complex IV Complex IV Complex IV Complex IV
COX4I1 COX6C MT-CYB	Complex IV Complex IV Complex III
UQCR10 UQCR11 UQCRH CYC1	Complex III Complex III Complex III Complex III
UQCRC1 UQCRC2 UQCRQ	Complex III Complex III Complex III
UQCRFS1 UQCRB SDHC SDHB	Complex III Complex III Complex II
SDHA SDHD MT-ND2	Complex II Complex II Complex II Complex I
NDUFV3 MT-ND4 MT-ND5	Complex I Complex I Complex I
MT-ND1 NDUFA7 NDUFS6 NDUFC1	Complex I Complex I Complex I Complex I
NDUFA10 NDUFB2 NDUFS2	Complex I Complex I Complex I
NDUFA6 NDUFS4 NDUFA8 NDUFA12	Complex I Complex I Complex I
NDUFB9 NDUFA3 NDUFB3	Complex I Complex I Complex I Complex I
NDUFA1 NDUFV1 NDUFB4	Complex I Complex I Complex I
NDUFA11 NDUFS1 NDUFV2 NDUFB5	Complex I Complex I Complex I
NDUFA2 NDUFA9 NDUFS5	Complex I Complex I Complex I Complex I
NDUFB11 MT-ND3 NDUFB7	Complex I Complex I Complex I
NDUFA5 NDUFB8 NDUFS7	Complex I Complex I Complex I
NDUFB10 NDUFA13 NDUFS8 NDUFB1	Complex I Complex I Complex I Complex I
NDUFS3 NDUFC2;NDUFC2-KCTD14 NDUFB6	Complex I Complex I Complex I
CPT1A CPT2 TMLHE ALDH9A1	Carnitine shuttle Carnitine shuttle Carnitine biosynthesis
ALDH9A1 ATPIF1 ATP5J ATP5E	Carnitine biosynthesis ATPIF1 ATP synthase ATP synthase
MT-ATP6 ATP5S MP68	ATP synthase ATP synthase ATP synthase
MT–ATP8 ATP5D ATP5J2	ATP synthase ATP synthase ATP synthase
USMG5 ATP5C1 ATP5J2 ATP5I	ATP synthase ATP synthase ATP synthase ATP synthase
ATP5L ATP5H ATP5A1	ATP synthase ATP synthase ATP synthase
ATP50 ATP5B ATP5F1 ARL8B	ATP synthase ATP synthase ATP synthase ARL8B
ARL8B AGK	ARL8B AGK