

Supplementary Figures and Tables for:

Proteome efficiency of metabolic pathways
in *Escherichia coli* increases along the nutrient flow

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This PDF includes Figure S1-S11 and Table S1-S6.

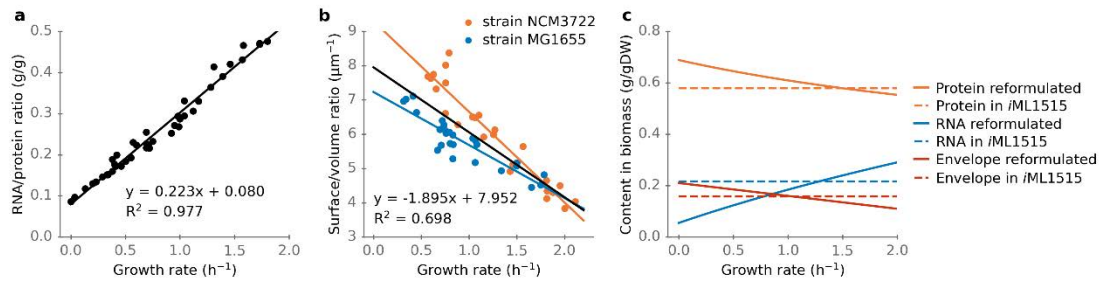


Fig. S1. Growth rate-dependent biomass composition. (a) Growth rate-dependent RNA/protein mass ratio; data from Ref.(1, 2). (b) Growth rate-dependent surface/volume ratio; data for wildtype strains in non-stressed growth conditions from Ref.(3). Surface/volume ratios are slightly different between the NCM3722 and MG1655 strains. Since the proteome data used in this study pertains to a third *E. coli* strain, BW25113, we used a regression model based on data from both strains (black line) to estimate the required production of cell envelope components. It should be noted that the re-estimated biomass composition here is not specific to any particular *E. coli* strain. (c) Growth rate-dependent biomass composition, reformulated considering the growth rate-dependent RNA/protein ratio in panel (a) and the surface/volume ratio in panel (b). The category “Envelope” includes murein, lipopolysaccharides, and lipids.

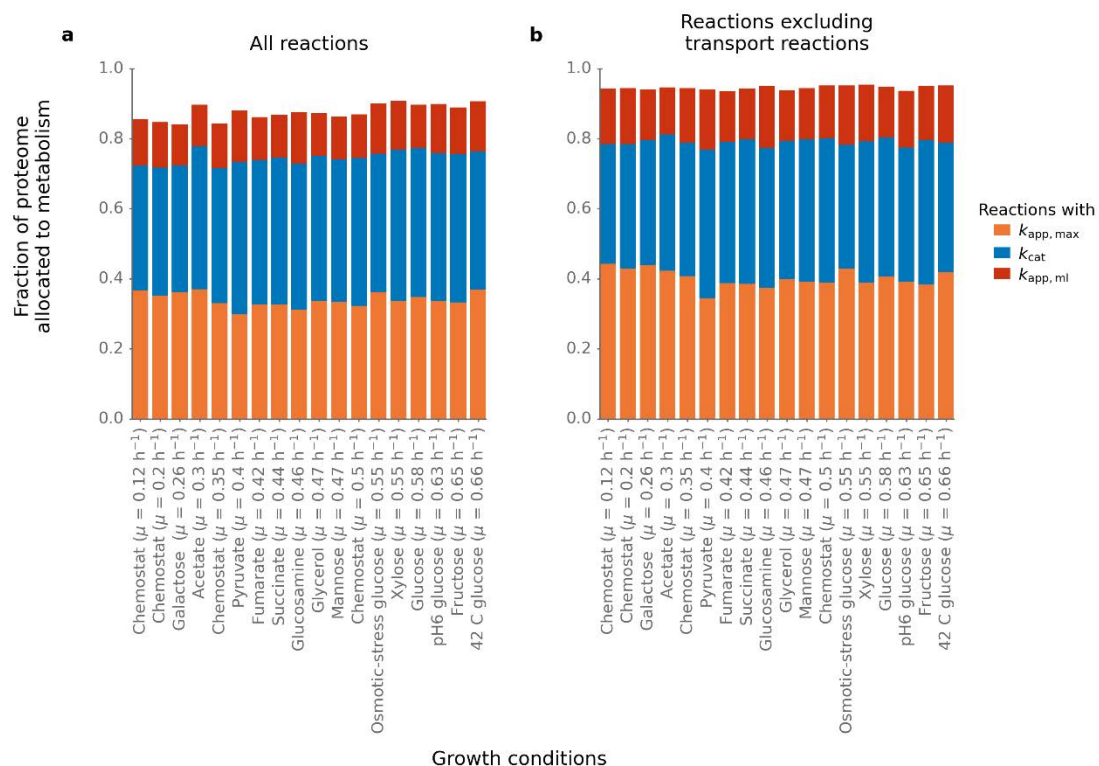


Fig. S2. Reactions with available enzyme turnover numbers cover more than 70% of the proteome fraction allocated to metabolism. Colors show the proteome fractions of enzymes with *in vivo* enzyme turnover number ($k_{app,max}$, in orange) or *in vitro* turnover number (k_{cat} , in blue). (a) All reactions in iML1515. (b) All non-transport reactions in iML1515.

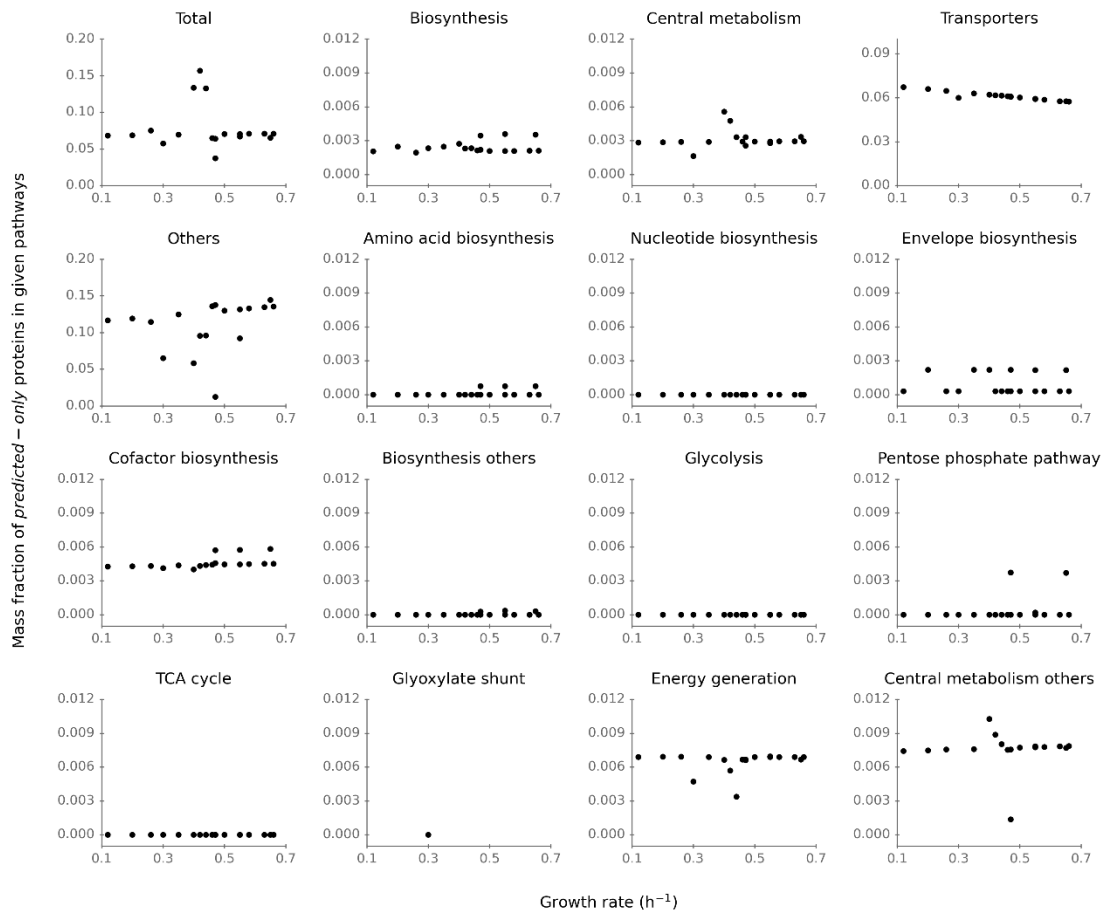


Fig. S3. The mass fraction of predicted-only proteins across pathways is generally low. Except for transporters and for proteins that cannot be assigned to particular pathways (others), the *predicted-only* proteins account for less than 1% of the predicted proteins in each pathway. We also note that the predictions of “others” and “transporters” are much lower than the measured data, and thus a relatively large fractions of predicted-only proteins in these two pathways does not change our conclusions.

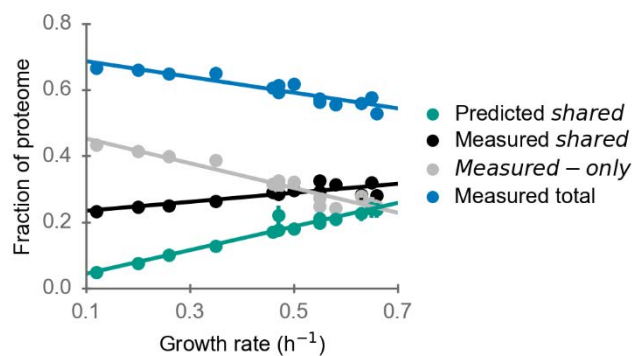


Fig. S4. Proteome efficiency of the whole metabolic network. With increasing growth rate, predicted and experimentally observed *shared* metabolic proteome fractions increase, while the *measured-only* metabolic proteome fraction decreases. Error bars extend from the 5th percentile to 95th percentile of 100 simulations, each with randomly perturbed turnover numbers.

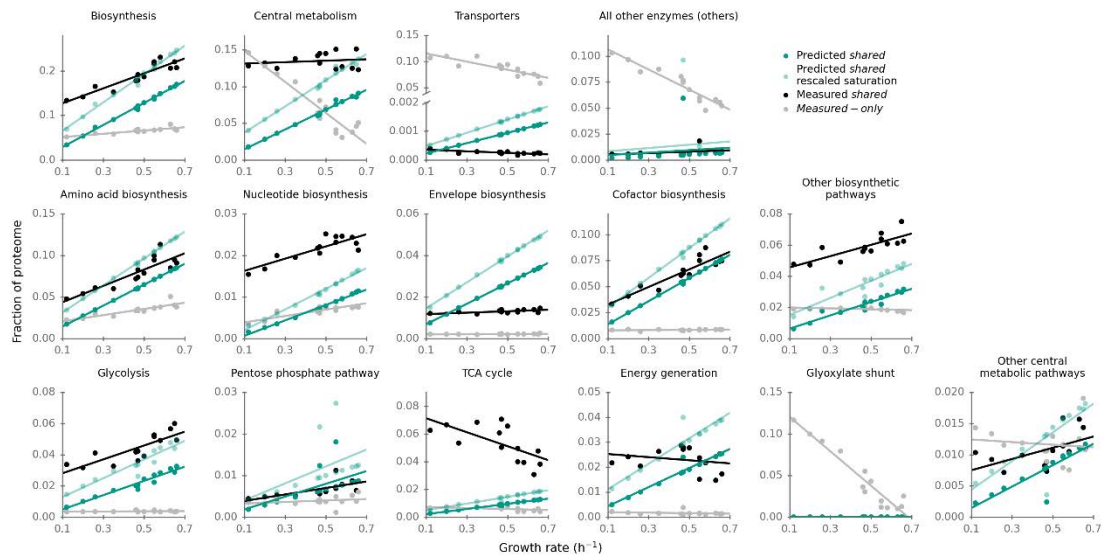


Fig. S5. Predictions of proteome allocated to pathways, including calculations based on growth rate-dependent expected enzyme saturation levels.

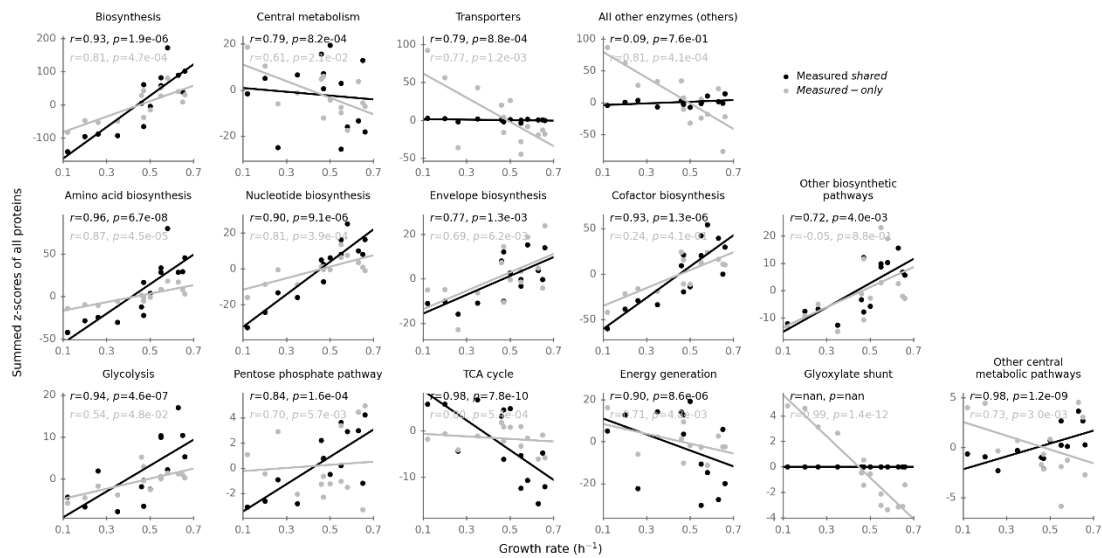


Fig. S6. Normalized proteome allocation trends of pathways based on z-scores. Pearson's r represents the correlation between the normalized proteome allocation and the total proteome fraction. The normalized proteome allocation of *shared* proteins is correlated with the total proteome fraction for all pathways; the only exception is "others", where most of the proteins are *measured-only* proteins.

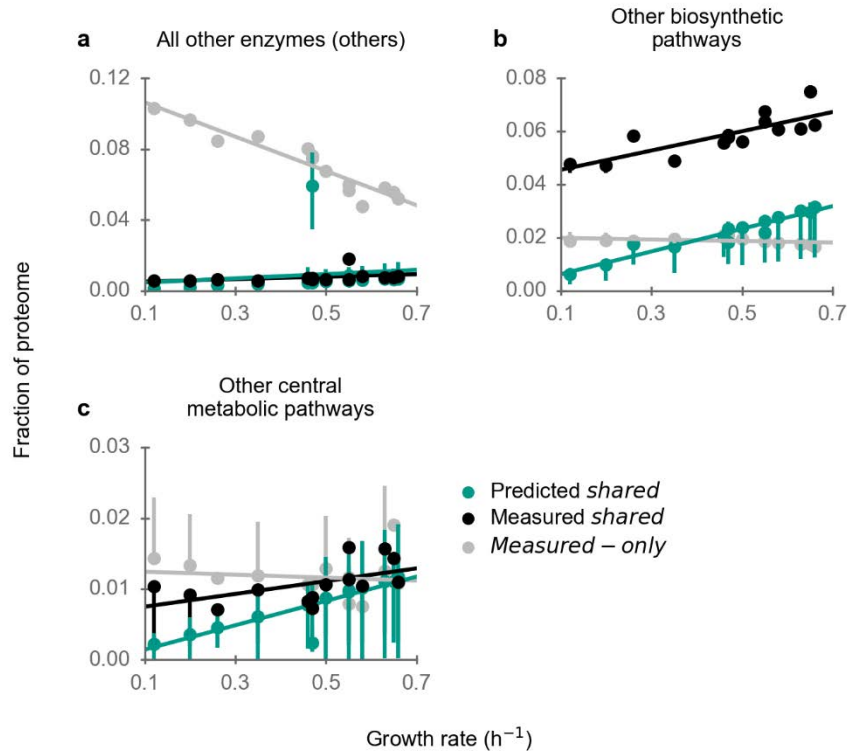


Fig. S7. The proteome efficiency of other pathways. (a) Combined proteome fraction of all enzymes in the *iML1515* model that are not part of the three pathway sets in Fig. 1b-1d (“others”). While the abundance of those proteins predicted to be active under optimality is similar to the expected abundance ($GMFE_{\text{pathway}} = 1.79$), a much higher proteome fraction is allocated to *measured-only* proteins. (b) Proteome investment into those biosynthetic proteins that are not part of the four pathway sets in Fig. 2. While predicted and observed concentrations of *shared* enzymes are strongly correlated ($r_{\text{pathway}}^2 = 0.60$; Table 1) and the abundance of individual enzymes can be well explained by the predictions ($r_{\text{individual}}^2 = 0.46$), the corresponding values differ on average by almost 3-fold ($GMFE_{\text{pathway}} = 2.91$). (c) Proteome investment into enzymes of central metabolism that are not part of the five pathway sets in Fig. 3. The prediction increases with growth rate, whereas the measured data is largely independent of growth rate. Error bars extend from the 5th percentile to 95th percentile of 100 simulations, each with randomly perturbed turnover numbers.

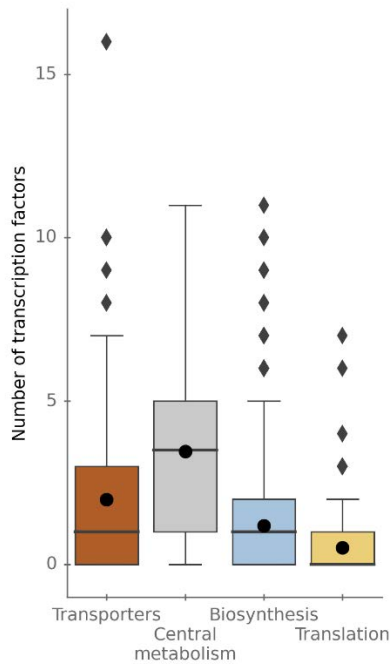


Fig. S8. Genes in more central cellular processes are regulated by fewer transcription factors than genes in peripheral processes. Genes encoding enzymes in biosynthesis pathways are regulated by fewer transcription factors than genes encoding transporters (one-sided Wilcoxon rank sum tests: $p < 10^{-10}$) and central metabolism ($p < 10^{-10}$). Genes in translation (here, all genes annotated with COG category “J”(4)) are also regulated by fewer transcription factors than genes in transporters ($p < 10^{-10}$) and central metabolism ($p < 10^{-10}$). Lines within boxes mark medians, dots indicate means. Boxes indicate 25% and 75% quantiles, whiskers extend from the boxes by 1.5x the interquartile range, diamonds are outliers.

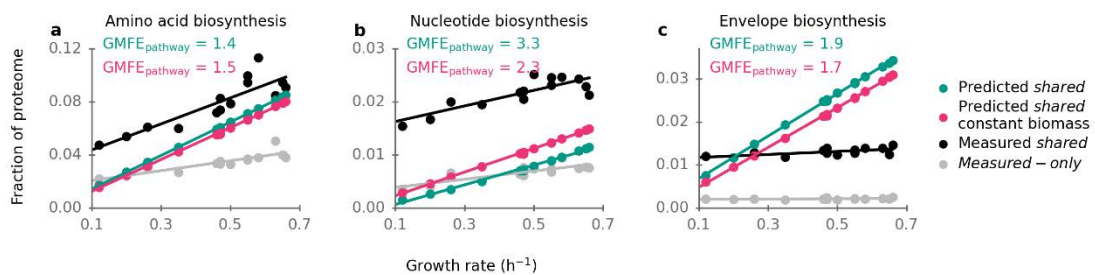


Fig. S9. Comparison of predictions assuming growth rate-dependent biomass composition versus constant biomass composition. (a) Amino acid biosynthesis pathway, (b) nucleotide biosynthesis pathway, and (c) cell envelope biosynthesis pathway. The composition of constant biomass was obtained from *iML1515* model.

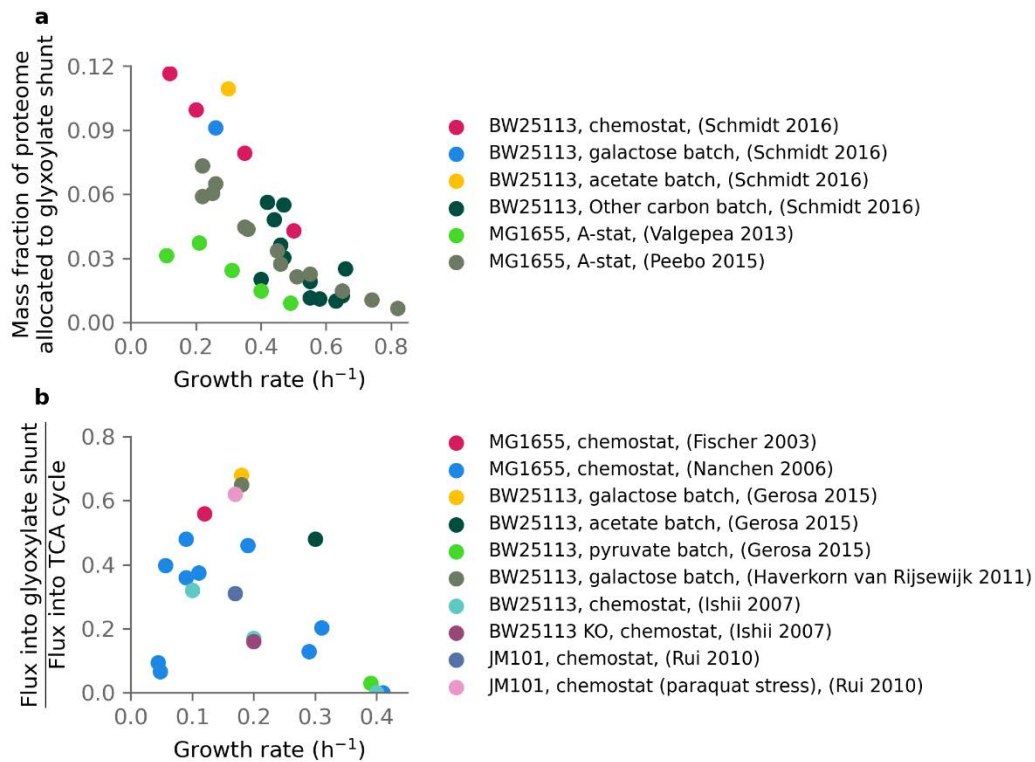


Fig. S10. Growth rate-dependent activity of the glyoxylate shunt. (a) Growth rate-dependent proteome allocation to the glyoxylate shunt, data from Refs. (5–7). **(b)** Growth rate-dependence of the flux through the glyoxylate shunt relative as a fraction of the flux into the TCA cycle. As the ^{13}C metabolic flux analysis usually lumps linear sequential reaction sequences into a single reaction, flux into the TCA cycle is quantified through ^{13}C measures of the flux of acetyl-CoA and oxaloacetate to isocitrate. The flux into the glyoxylate shunt is quantified through ^{13}C measures of the flux catalyzed by Isocitrate lyase (isocitrate \rightarrow succinate + glyoxylate). Data from Refs. (8–13). Both protein abundance and flux decrease with increasing growth rate. The glyoxylate shunt’s high protein abundance and large flux at low growth rates mark it as an important alternative to the TCA cycle in these conditions.

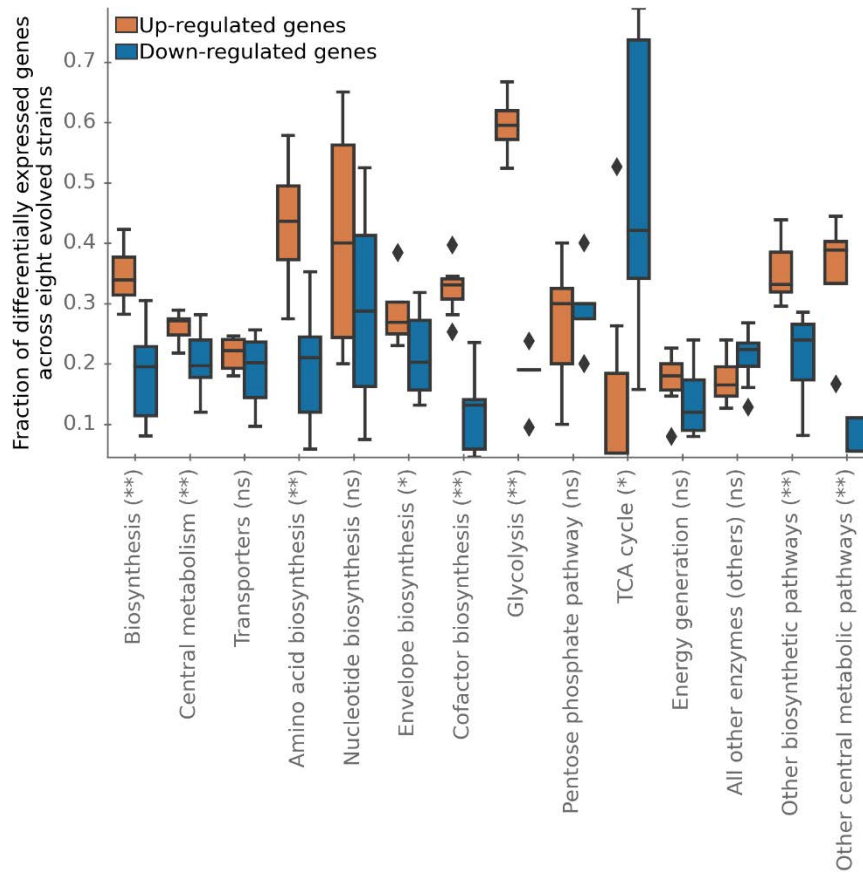


Fig. S11. Fraction of differentially expressed genes in evolved strains grown on glucose minimal medium across pathways. *P*-values of two-sided paired *t*-tests (*n* = 8) are denoted in brackets as follows: ** for *p* < 0.01, * for *p* < 0.1, and ns for *p* ≥ 0.1 (non-significant).

Supplementary References

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Table S2: Enzyme turnover numbers and their sources

Reaction id	value	direction	source ¹
DHORTS	14.30024	-1	$k_{app,max}$
OMPDC	58.99106	1	$k_{app,max}$
GSSD	9.681925	1	$k_{app,max}$
CS	31.49134	1	$k_{app,max}$
ICDHyr	10.0849	1	$k_{app,max}$
APRAUR	0.076305	1	$k_{app,max}$
DB4PS	0.091319	1	$k_{app,max}$
RBFK	0.086318	1	$k_{app,max}$
GLYK	1.422746	1	$k_{app,max}$
ASPTA	21.5778	-1	$k_{app,max}$
RBFSb	0.024042	1	$k_{app,max}$
SHCHD2	0.131828	1	$k_{app,max}$
CPPPGO	0.696853	1	$k_{app,max}$
ILETA	8.458744	-1	$k_{app,max}$
VALTA	19.05329	-1	$k_{app,max}$
ORPT	80.85754	-1	$k_{app,max}$
RBFSa	0.07372	1	$k_{app,max}$
PFK_3	108.1692	1	$k_{app,max}$
DHAD2	12.76649	1	$k_{app,max}$
PSCVT	8.659949	1	$k_{app,max}$
ANPRT	3.173382	1	$k_{app,max}$
CHORS	32.64963	1	$k_{app,max}$
IGPS	3.37822	1	$k_{app,max}$
DMATT	0.585096	1	$k_{app,max}$
GRTT	0.585096	1	$k_{app,max}$
UPP3S	0.912843	1	$k_{app,max}$
UPPDC1	0.068513	1	$k_{app,max}$
MOAT	10.66652	1	$k_{app,max}$
UDCPDPS	0.062705	1	$k_{app,max}$
USHD	16.5942	1	$k_{app,max}$
ENO	58.18488	1	$k_{app,max}$
MGSA	805.9263	1	$k_{app,max}$
IMPC	11.8847	-1	$k_{app,max}$
IMPD	1.399624	1	$k_{app,max}$
LPADSS	8.700117	1	$k_{app,max}$
GLUTRS	0.120344	1	$k_{app,max}$
TDSK	16.71266	1	$k_{app,max}$
PGI	129.9447	1	$k_{app,max}$
TMPPP	0.021426	1	$k_{app,max}$
PGK	114.571	-1	$k_{app,max}$
PGL	588.7476	1	$k_{app,max}$
DHQTi	40.31471	1	$k_{app,max}$
IPMD	9.643575	1	$k_{app,max}$
AIRC3	30.13264	-1	$k_{app,max}$
ADSL2r	10.24187	1	$k_{app,max}$
TMDS	4.639885	1	$k_{app,max}$
PPCDC	0.073955	1	$k_{app,max}$
DHAD1	38.41795	1	$k_{app,max}$
FMNAT	0.086318	1	$k_{app,max}$
OHPBAT	0.001678	1	$k_{app,max}$
KARA1	3.787365	-1	$k_{app,max}$
KARA2	1.258562	1	$k_{app,max}$
NADK	2.640962	1	$k_{app,max}$
SUCBZS	0.611389	1	$k_{app,max}$
DHBD	0.936895	1	$k_{app,max}$
ICHORT	0.113636	1	$k_{app,max}$
GLUTRR	12.14911	1	$k_{app,max}$
HEMEOS	0.445764	1	$k_{app,max}$
ADMDC	2.161437	1	$k_{app,max}$
LGTHL	64.4189	1	$k_{app,max}$
ARGSS	1.754436	1	$k_{app,max}$
AGPR	12.40352	-1	$k_{app,max}$
SPMS	0.993811	1	$k_{app,max}$
G1SAT	0.168925	1	$k_{app,max}$
KDOCT2	1.446841	1	$k_{app,max}$
EDTXS1	6.821223	1	$k_{app,max}$
DXPS	1.027865	1	$k_{app,max}$
CDPMEK	4.086811	1	$k_{app,max}$
EDTXS2	3.793857	1	$k_{app,max}$
MECDPS	3.386297	1	$k_{app,max}$
PGAMT	2.249193	-1	$k_{app,max}$
ICL	4.849202	1	$k_{app,max}$
UAGDP	7.491129	1	$k_{app,max}$

¹ Note $k_{app,max}$: in vivo enzyme turnover number estimated from experimental data by Heckmann *et al.* Ref. (31). $k_{app,ml}$: in vivo enzyme turnover number estimated from machine learning by Heckmann *et al.* Ref. (31). k_{cat} : in vitro enzyme turnover number from ccFBA by Desouki Ref. (27).transporter: transport reactions with no in vivo or in vitro parameters, set to 65 s⁻¹ according to Heckmann *et al.* Ref. (31).

UAGCVT	1.397778	1	K _{app,max}
GLUR	32.76217	-1	K _{app,max}
NNDPR	0.446503	1	K _{app,max}
DMQMT	0.021907	1	K _{app,max}
HPPK2	0.31375	1	K _{app,max}
PTPATi	1.286563	1	K _{app,max}
DXPRIi	3.079024	1	K _{app,max}
GTPCII2	0.068128	1	K _{app,max}
OHPHM	0.021907	1	K _{app,max}
GHMT2r	4.439337	1	K _{app,max}
DHORD2	38.53068	1	K _{app,max}
SDPDS	102.2286	1	K _{app,max}
ADCL	1.553082	1	K _{app,max}
TDPDRE	3.910464	1	K _{app,max}
TDPDRR	1.226032	1	K _{app,max}
UAPGR	5.838932	1	K _{app,max}
CTPS2	4.258782	1	K _{app,max}
PAPPT3	15.59233	1	K _{app,max}
UAGPT3	6.150296	1	K _{app,max}
GMHEPPA	23.31693	1	K _{app,max}
S7PI	1.63139	1	K _{app,max}
AGMHE	0.934696	1	K _{app,max}
GMHEPAT	1.806171	1	K _{app,max}
FADRx	0.088143	1	K _{app,max}
ACODA	10.97911	1	K _{app,max}
FCLT	1.301035	1	K _{app,max}
HBZOPT	0.567372	1	K _{app,max}
PNTK	0.227027	1	K _{app,max}
ADSS	1.84019	1	K _{app,max}
OCTDPS	1.287727	1	K _{app,max}
DPCOAK	1.333897	1	K _{app,max}
ASNS2	8.972534	1	K _{app,max}
KDOPS	0.763831	1	K _{app,max}
KDOPP	18.7249	1	K _{app,max}
DMPPS	0.148089	1	K _{app,max}
IPDPS	0.888064	1	K _{app,max}
DAPE	55.54165	1	K _{app,max}
DHDPRy	8.859222	1	K _{app,max}
HSK	151.8686	1	K _{app,max}
SDPTA	6.698489	-1	K _{app,max}
THRS	28.64601	1	K _{app,max}
HSST	8.539119	1	K _{app,max}
PPNDH	17.86531	1	K _{app,max}
DTMPK	7.94476	1	K _{app,max}
RHCCE	0.015614	1	K _{app,max}
G3PD2	42.97261	-1	K _{app,max}
GMPS2	2.395951	1	K _{app,max}
GALUi	1.309006	1	K _{app,max}
U23GAAT	1.39172	1	K _{app,max}
UAAGDS	4.160067	1	K _{app,max}
UAMAS	2.786797	1	K _{app,max}
UAMAGS	4.523614	1	K _{app,max}
P5CR	23.66027	1	K _{app,max}
DHPPDA2	0.076305	1	K _{app,max}
ASP1DC	0.041348	1	K _{app,max}
UGMDDS	8.68655	1	K _{app,max}
DBTS	0.003799	1	K _{app,max}
UHGADA	12.98895	1	K _{app,max}
ATPPRT	3.754474	1	K _{app,max}
HISTD	3.179596	1	K _{app,max}
DHDPS	7.126217	1	K _{app,max}
G1PACT	7.491129	1	K _{app,max}
GF6PTA	2.021826	1	K _{app,max}
IGPDH	3.333692	1	K _{app,max}
PRPPS	17.03327	1	K _{app,max}
GLUCYS	0.016512	1	K _{app,max}
PRATPP	9.319031	1	K _{app,max}
PRMICI	13.48006	1	K _{app,max}
GLUPRT	18.65284	1	K _{app,max}
SERAT	21.21567	1	K _{app,max}
NADS1	0.122167	1	K _{app,max}
BPNT	84.84513	1	K _{app,max}
ASAD	18.63376	-1	K _{app,max}
DAPDC	55.701	1	K _{app,max}
GARFT	57.1662	1	K _{app,max}
PPND	19.97671	1	K _{app,max}

METAT	0.054295	1	<i>k</i> _{app,max}
PRAMPC	9.319031	1	<i>k</i> _{app,max}
TMPK	0.206249	1	<i>k</i> _{app,max}
PPBNGS	0.158583	1	<i>k</i> _{app,max}
PPNCL2	0.073955	1	<i>k</i> _{app,max}
PRAGS _r	13.68985	1	<i>k</i> _{app,max}
PRFGS	10.23022	1	<i>k</i> _{app,max}
NNAT _r	5.322368	1	<i>k</i> _{app,max}
PSERT	37.41342	1	<i>k</i> _{app,max}
AIRC2	48.17105	1	<i>k</i> _{app,max}
GAPD	35.76496	1	<i>k</i> _{app,max}
HEX1	305.0672	1	<i>k</i> _{app,max}
GND	66.7612	1	<i>k</i> _{app,max}
MOAT2	10.66652	1	<i>k</i> _{app,max}
G6PDH2 _r	163.6787	1	<i>k</i> _{app,max}
ARGSL	8.567866	1	<i>k</i> _{app,max}
GLU5K	12.22192	1	<i>k</i> _{app,max}
IPPS	8.606867	1	<i>k</i> _{app,max}
MOHMT	0.029624	1	<i>k</i> _{app,max}
AHCYSNS	0.052337	1	<i>k</i> _{app,max}
SERAS _r	0.209118	1	<i>k</i> _{app,max}
CHRPL	0.543619	1	<i>k</i> _{app,max}
FBA3	29.27115	1	<i>k</i> _{app,max}
PMPK	0.03897	1	<i>k</i> _{app,max}
PSP_L	2693.142	1	<i>k</i> _{app,max}
ADK3	14.77674	1	<i>k</i> _{app,max}
ADSK	85.07205	1	<i>k</i> _{app,max}
SHSL1	7.32263	1	<i>k</i> _{app,max}
GMHEPK	1.806171	1	<i>k</i> _{app,max}
GTPCI	0.009085	1	<i>k</i> _{app,max}
CYTK1	14.9248	1	<i>k</i> _{app,max}
GK1	37.5363	1	<i>k</i> _{app,max}
DHFS	0.119841	1	<i>k</i> _{app,max}
ASPO6	1.248945	1	<i>k</i> _{app,max}
PRASCS _i	5.514622	1	<i>k</i> _{app,max}
DHPS2	0.371229	1	<i>k</i> _{app,max}
MTHFC	40.91791	1	<i>k</i> _{app,max}
MTHFD	40.91791	1	<i>k</i> _{app,max}
QULNS	0.724281	1	<i>k</i> _{app,max}
GLCTR1	17.76051	1	<i>k</i> _{app,max}
GLCTR3	10.85577	1	<i>k</i> _{app,max}
HEPT4	6.874304	1	<i>k</i> _{app,max}
CBLAT	0.204043	1	<i>k</i> _{app,max}
NH4 _{tp}	8586.02	1	<i>k</i> _{app,max}
GLYCLT4 _{pp}	0.661979	1	<i>k</i> _{app,max}
ACGK	6.482128	1	<i>k</i> _{app,max}
HMBS	0.194692	1	<i>k</i> _{app,max}
MEPCT	13.96947	1	<i>k</i> _{app,max}
DAAD	0.212326	1	<i>k</i> _{app,max}
GLCt2 _{pp}	266.8132	1	<i>k</i> _{app,max}
HSTPT	5.95604	1	<i>k</i> _{app,max}
THDPS	1.677663	1	<i>k</i> _{app,max}
GLYOX	2609.268	1	<i>k</i> _{app,max}
PRA _{li}	3.37822	1	<i>k</i> _{app,max}
MTHFR2	3.720112	1	<i>k</i> _{app,max}
MTAN	0.444461	1	<i>k</i> _{app,max}
AMAOT _r	0.000662	1	<i>k</i> _{app,max}
GTHS	0.016597	1	<i>k</i> _{app,max}
ACGS	25.92218	1	<i>k</i> _{app,max}
HEPT1	15.56121	1	<i>k</i> _{app,max}
HEPT2	4.951763	1	<i>k</i> _{app,max}
MOAT3C	15.80545	1	<i>k</i> _{app,max}
RHAT1	19.01035	1	<i>k</i> _{app,max}
GALT1	9.122473	1	<i>k</i> _{app,max}
GLCTR2	17.99057	1	<i>k</i> _{app,max}
HEPK1	9.747015	1	<i>k</i> _{app,max}
HEPT3	12.10135	1	<i>k</i> _{app,max}
HEPK2	12.15066	1	<i>k</i> _{app,max}
SHCHF	0.131828	1	<i>k</i> _{app,max}
SUCBZL	1.396733	1	<i>k</i> _{app,max}
PRAIS	14.58303	1	<i>k</i> _{app,max}
PANTS	0.028019	1	<i>k</i> _{app,max}
GCALDD	0.042906	1	<i>k</i> _{app,max}
HISTP	3.333692	1	<i>k</i> _{app,max}
KAS15	14.98418	1	<i>k</i> _{app,max}
T2DECAI	2.047433	1	<i>k</i> _{app,max}

3OAS161	1.876475	1	$k_{app,max}$
DHQS	38.1248	1	$k_{app,max}$
EAR161x	1.491904	1	$k_{app,max}$
DASYN160	111.4841	1	$k_{app,max}$
DASYN161	86.71167	1	$k_{app,max}$
DASYN181	44.69031	1	$k_{app,max}$
PGCD	37.46429	1	$k_{app,max}$
PG161abcpp	0.832095	1	$k_{app,max}$
3OAS141	1.876475	1	$k_{app,max}$
3OAR121	3.763232	1	$k_{app,max}$
ADSL1r	4.036562	1	$k_{app,max}$
AICART	11.8847	1	$k_{app,max}$
PG160abcpp	1.069765	1	$k_{app,max}$
DHBS	0.509428	1	$k_{app,max}$
PPGO3	0.184466	1	$k_{app,max}$
OMBZLM	0.013342	1	$k_{app,max}$
EAR141x	1.491904	1	$k_{app,max}$
EAR181x	0.507402	1	$k_{app,max}$
TPI	193.9066	1	$k_{app,max}$
PERD	0.016316	1	$k_{app,max}$
5DOAN	0.029921	1	$k_{app,max}$
3OAR120	4.480061	1	$k_{app,max}$
SO4t2pp	248.6991	1	$k_{app,max}$
DHNAOT4	0.356291	1	$k_{app,max}$
PSSA161	2.879332	1	$k_{app,max}$
PSD160	5.497192	1	$k_{app,max}$
PSD161	4.275621	1	$k_{app,max}$
COLIPAabcpp	0.808873	1	$k_{app,max}$
3OAR181	1.27989	1	$k_{app,max}$
EAR140x	1.350815	1	$k_{app,max}$
3OAS181	1.316407	1	$k_{app,max}$
AGPAT160	30.0267	1	$k_{app,max}$
PGSA160	12.65921	1	$k_{app,max}$
3OAR140	4.265518	1	$k_{app,max}$
3OAR160	3.192801	1	$k_{app,max}$
PGSA161	9.846652	1	$k_{app,max}$
3OAR141	3.763232	1	$k_{app,max}$
3OAR161	3.763232	1	$k_{app,max}$
EAR121x	1.491904	1	$k_{app,max}$
PE160abcpp	3.155509	1	$k_{app,max}$
PE161abcpp	2.454306	1	$k_{app,max}$
PG181abcpp	0.428799	1	$k_{app,max}$
PGSA181	5.074589	1	$k_{app,max}$
PSSA160	3.701974	1	$k_{app,max}$
PSSA181	1.484009	1	$k_{app,max}$
AGPAT181	12.03672	1	$k_{app,max}$
3OAR60	8.243372	1	$k_{app,max}$
3OAR80	8.243372	1	$k_{app,max}$
EAR100x	1.776085	1	$k_{app,max}$
EAR120x	1.776085	1	$k_{app,max}$
EAR160x	1.265761	1	$k_{app,max}$
3OAS121	1.876475	1	$k_{app,max}$
3OAR40	8.243372	1	$k_{app,max}$
3OAR100	8.243293	1	$k_{app,max}$
AGPAT161	23.3546	1	$k_{app,max}$
EAR40x	3.268021	1	$k_{app,max}$
PSD181	2.203657	1	$k_{app,max}$
EAR60x	3.268021	1	$k_{app,max}$
EAR80x	3.268021	1	$k_{app,max}$
PE181abcpp	1.264962	1	$k_{app,max}$
ZN2tpp	0.431349	1	$k_{app,max}$
CAT6pp	1.058014	-1	$k_{app,max}$
AMMQLT8	0.013342	1	$k_{app,max}$
DALat2pp	3.342165	1	$k_{app,max}$
FE2tpp	10.82699	1	$k_{app,max}$
MN2tpp	0.87601	1	$k_{app,max}$
MALCOAMT	0.002455	1	$k_{app,max}$
CU2tpp	0.897312	1	$k_{app,max}$
OGMEACPS	9.57E-05	1	$k_{app,max}$
OGMEACPR	5.26E-05	1	$k_{app,max}$
OPMEACPS	2.62E-05	1	$k_{app,max}$
MPTAT	0.077786	1	$k_{app,max}$
OPMEACPD	9.10E-05	1	$k_{app,max}$
EPMEACPR	2.09E-05	1	$k_{app,max}$
AOXsr2	0.001779	1	$k_{app,max}$
OGMEACPD	9.10E-05	1	$k_{app,max}$

EGMEACPR	2.09E-05	1	$k_{app,max}$
OPMEACPR	5.26E-05	1	$k_{app,max}$
PMEACPE	0.002756	1	$k_{app,max}$
ICYSDS	0.021592	1	$k_{app,max}$
I4FE4SR	0.01136	1	$k_{app,max}$
BMOCOS	0.010176	1	$k_{app,max}$
FESR	4.942637	1	$k_{app,max}$
BTS5	0.000739	1	$k_{app,max}$
LIPOS	0.000242	1	$k_{app,max}$
MPTSS	0.380772	1	$k_{app,max}$
LIPOCT	0.002578	1	$k_{app,max}$
VPAMTr	49.25294	-1	$k_{app,max}$
DHNPA2r	2.917904	1	$k_{app,max}$
SEPHCHCS	1.474107	1	$k_{app,max}$
DHNCOAS	0.052986	1	$k_{app,max}$
DHNCOAT	0.683201	1	$k_{app,max}$
TYRL	0.014337	1	$k_{app,max}$
MOCOS	0.011404	1	$k_{app,max}$
CYTDK2	3.633271	1	$k_{app,ml}$
NDPK5	5.004318	1	$k_{app,ml}$
NDPK5	3.760914	-1	$k_{app,ml}$
NDPK6	0.096211	1	$k_{app,ml}$
NDPK6	4.9978	-1	$k_{app,ml}$
NDPK8	3.948568	1	$k_{app,ml}$
NDPK8	2.493469	-1	$k_{app,ml}$
PYNP2r	6.931895	1	$k_{app,ml}$
PYNP2r	3.136084	-1	$k_{app,ml}$
ICDHyr	3.74751	-1	$k_{app,ml}$
ALATA_L2	3.880836	1	$k_{app,ml}$
DURIPP	3.250846	1	$k_{app,ml}$
ACALD	16.37094	1	$k_{app,ml}$
ACALD	8.194523	-1	$k_{app,ml}$
PTRCTA	3.899522	1	$k_{app,ml}$
CYSDS	2.611557	1	$k_{app,ml}$
MAN6PI	2.817264	-1	$k_{app,ml}$
TRPAS2	3.364753	-1	$k_{app,ml}$
ALATA_L	12.19887	-1	$k_{app,ml}$
PPM	2.86429	1	$k_{app,ml}$
ACP1_FMN	3.891823	1	$k_{app,ml}$
NDP3	5.563382	1	$k_{app,ml}$
CDPPH	5.226493	1	$k_{app,ml}$
NDP7	5.226493	1	$k_{app,ml}$
ALATA_D2	3.880836	1	$k_{app,ml}$
PMDPHT	0.188859	1	$k_{app,ml}$
GTHOr	2.287207	-1	$k_{app,ml}$
GLYCTO2	1.57191	1	$k_{app,ml}$
GLYCTO3	1.484281	1	$k_{app,ml}$
IPPMib	3.659187	1	$k_{app,ml}$
IPPMib	8.523155	-1	$k_{app,ml}$
GLYCTO4	1.484013	1	$k_{app,ml}$
ACHBS	7.395987	1	$k_{app,ml}$
ACLS	29.79637	1	$k_{app,ml}$
G3PD5	2.686021	1	$k_{app,ml}$
G1PP	0.056349	1	$k_{app,ml}$
TMDPP	2.991571	-1	$k_{app,ml}$
ECOAH6	4.236997	1	$k_{app,ml}$
ECOAH6	2.418356	-1	$k_{app,ml}$
ENO	53.81454	-1	$k_{app,ml}$
FBA	42.12004	-1	$k_{app,ml}$
HCO3E	6.099103	-1	$k_{app,ml}$
IMPC	4.277784	1	$k_{app,ml}$
ECOAH4	5.827645	1	$k_{app,ml}$
ECOAH4	3.17966	-1	$k_{app,ml}$
NTPP1	23.34322	1	$k_{app,ml}$
URIK1	3.576691	1	$k_{app,ml}$
URIK2	3.633271	1	$k_{app,ml}$
NTPP2	23.34322	1	$k_{app,ml}$
NTPP2	29.25939	1	$k_{app,ml}$
CYTDK1	3.576691	1	$k_{app,ml}$
AIRC3	7.516071	1	$k_{app,ml}$
TALA	39.83541	1	$k_{app,ml}$
ADSL2r	3.193393	-1	$k_{app,ml}$
TKT1	6.0799	-1	$k_{app,ml}$
RNDR1	0.923364	1	$k_{app,ml}$
RNDR4	0.00941	1	$k_{app,ml}$
LYSAM	3.126748	1	$k_{app,ml}$

LYSAM	3.126748	-1	K_app,ml
FTHFD	46.06887	1	K_app,ml
IPPM1a	3.336187	1	K_app,ml
IPPM1a	8.230083	-1	K_app,ml
G3PD7	2.589234	1	K_app,ml
EDD	22.88964	1	K_app,ml
NADH9	2.403306	1	K_app,ml
ADK4	4.506135	1	K_app,ml
ADK4	4.281521	-1	K_app,ml
INSK	3.979366	1	K_app,ml
OHPBAT	0.06708	-1	K_app,ml
NADDP	2.587898	1	K_app,ml
HEX4	3.469276	1	K_app,ml
KARA2	2.607869	-1	K_app,ml
ICHORS_copy2	7.732491	1	K_app,ml
ICHORS_copy2	6.314789	-1	K_app,ml
DHBD	2.377994	-1	K_app,ml
RBP4E	3.032085	-1	K_app,ml
TARTD	3.088357	1	K_app,ml
PDXPP	4.055824	1	K_app,ml
HACD6	3.339417	1	K_app,ml
HACD6	2.576447	-1	K_app,ml
ECOAH1	5.435808	1	K_app,ml
ECOAH2	5.843902	1	K_app,ml
ECOAH2	3.217962	-1	K_app,ml
AGPR	3.582858	1	K_app,ml
ECOAH3	5.841133	1	K_app,ml
ECOAH3	3.218444	-1	K_app,ml
ALTRH	2.85554	1	K_app,ml
GUI1	2.725522	1	K_app,ml
GUI1	2.725522	-1	K_app,ml
EDXS3	3.892001	1	K_app,ml
PGAMT	3.095659	1	K_app,ml
EDXS4	4.834386	1	K_app,ml
PMANM	2.995768	-1	K_app,ml
UAG2E	3.055023	-1	K_app,ml
GLYCL	8.970031	1	K_app,ml
CMPN	11.29015	1	K_app,ml
TRE6PH	9.31694	1	K_app,ml
ICHORS_copy3	7.712237	1	K_app,ml
HACD4	3.353535	1	K_app,ml
HACD4	2.35958	-1	K_app,ml
CINND0	2.766831	1	K_app,ml
DHPPD	2.733627	1	K_app,ml
HACD5	3.60068	1	K_app,ml
HACD5	2.500709	-1	K_app,ml
DHCIND	2.733627	1	K_app,ml
HACD7	3.281245	1	K_app,ml
HACD7	2.536963	-1	K_app,ml
CITL	6.392752	1	K_app,ml
GOFUCR	2.275287	1	K_app,ml
DNMPPA	2.750124	1	K_app,ml
SADT2	4.711474	1	K_app,ml
OPHBDC	1.522857	1	K_app,ml
AACPS1	0.473365	1	K_app,ml
GHMT2r	4.716639	-1	K_app,ml
LCARS	2.751624	1	K_app,ml
LCARS	18.49761	-1	K_app,ml
ALCD19	11.22635	1	K_app,ml
ALCD19	3.933228	-1	K_app,ml
RMPA	3.08298	-1	K_app,ml
SBTPD	2.728479	1	K_app,ml
SBTPD	2.501926	-1	K_app,ml
PFK_2	6.244474	1	K_app,ml
GLTPD	2.635642	1	K_app,ml
GLTPD	2.59837	-1	K_app,ml
ACACCT	2.846615	1	K_app,ml
BUTCT	2.752224	1	K_app,ml
TGBPA	3.29865	-1	K_app,ml
MANAO	2.559645	1	K_app,ml
MANAO	2.955656	-1	K_app,ml
TAGURr	2.578188	1	K_app,ml
TAGURr	2.951025	-1	K_app,ml
2DGULRx	2.17844	1	K_app,ml
DKGLCNR2y	2.448182	1	K_app,ml
2DGULRy	2.448182	1	K_app,ml

DHAPT	1.008448	1	K _{app,ml}
UMPK	4.601321	-1	K _{app,ml}
RZ5PP	5.385396	1	K _{app,ml}
DDPGALA	3.250094	-1	K _{app,ml}
DURIK1	6.997138	1	K _{app,ml}
G3PD6	2.589234	1	K _{app,ml}
NDPK7	4.9978	-1	K _{app,ml}
GLYAT	10.53765	1	K _{app,ml}
GLYAT	5.520045	-1	K _{app,ml}
ADNUC	3.447988	1	K _{app,ml}
F6PA	6.249798	1	K _{app,ml}
F6PA	11.29118	-1	K _{app,ml}
CRNBCT	3.236285	1	K _{app,ml}
CRNBCT	3.236285	-1	K _{app,ml}
CRNCBCT	3.236285	1	K _{app,ml}
CRNCBCT	3.236285	-1	K _{app,ml}
CRNCDH	4.078269	1	K _{app,ml}
CRNCDH	4.078269	-1	K _{app,ml}
IDOND	2.670143	1	K _{app,ml}
IDOND	17.8758	-1	K _{app,ml}
IDOND2	17.11058	1	K _{app,ml}
NADN	21.87028	1	K _{app,ml}
GLYCLTDx	2.401812	1	K _{app,ml}
NDPK2	19.10756	1	K _{app,ml}
NDPK2	6.490113	-1	K _{app,ml}
NDPK3	11.67335	1	K _{app,ml}
NDPK3	4.9978	-1	K _{app,ml}
HPYRRy	12.34104	1	K _{app,ml}
NDPK4	4.9978	-1	K _{app,ml}
HETZK	3.805523	1	K _{app,ml}
MCOATA	2.622024	-1	K _{app,ml}
AST	3.243138	1	K _{app,ml}
SGDS	6.736857	1	K _{app,ml}
CYNTHA	2.868002	1	K _{app,ml}
SDPTA	4.120056	1	K _{app,ml}
TMAOR2	2.26961	1	K _{app,ml}
DGK1	3.378429	-1	K _{app,ml}
DTMPK	3.421685	-1	K _{app,ml}
NDPK1	3.292598	-1	K _{app,ml}
DDPA	10.95666	1	K _{app,ml}
NMNDA	20.73942	1	K _{app,ml}
PGMT	22.12188	-1	K _{app,ml}
FRUK	63.07576	1	K _{app,ml}
TDPADGAT	4.807875	1	K _{app,ml}
MALS	32.55933	1	K _{app,ml}
X5PL3E	3.45401	1	K _{app,ml}
DOGULNR	3.051568	1	K _{app,ml}
BETALDHx	2.585938	1	K _{app,ml}
MLTG3	0.272841	1	K _{app,ml}
DKGLCNR1	2.508822	1	K _{app,ml}
NADTRHD	24.51586	1	K _{app,ml}
AACPS2	2.004406	1	K _{app,ml}
FMNRx2	2.27405	1	K _{app,ml}
ARAI	2.740122	-1	K _{app,ml}
FCI	2.646269	1	K _{app,ml}
FCI	2.646269	-1	K _{app,ml}
FCLK	2.90671	1	K _{app,ml}
FCLPA	3.027816	1	K _{app,ml}
FCLPA	2.841457	-1	K _{app,ml}
GSNK	4.019751	1	K _{app,ml}
AACPS3	1.470615	1	K _{app,ml}
AACPS4	1.572154	1	K _{app,ml}
UGLYCH	5.374962	1	K _{app,ml}
NTPP3	8.841867	1	K _{app,ml}
NTPP6	27.48631	1	K _{app,ml}
NTPP7	5.451964	1	K _{app,ml}
OP4ENH	4.119825	1	K _{app,ml}
HOPNTAL	3.265266	1	K _{app,ml}
NTPP8	8.705841	1	K _{app,ml}
3HPPPNH	2.676857	1	K _{app,ml}
PPCSCT	18.76042	1	K _{app,ml}
AKGDH	40.88314	1	K _{app,ml}
HPYRRx	0.352116	1	K _{app,ml}
GLYCDx	4.506491	1	K _{app,ml}
DCYTD	3.76805	1	K _{app,ml}
AMANK	3.752667	1	K _{app,ml}

UAGAAT	2.99576	-1	K _{app,ml}
GART	6.381109	1	K _{app,ml}
PRPPS	16.77529	-1	K _{app,ml}
H2SO	3.487541	1	K _{app,ml}
PRMICI	3.676523	-1	K _{app,ml}
GMPR	2.610772	1	K _{app,ml}
ASPK	3.322316	-1	K _{app,ml}
NTRIR2x	3.284036	1	K _{app,ml}
GARFT	13.45849	-1	K _{app,ml}
PHETA1	4.269688	1	K _{app,ml}
PHETA1	11.59656	-1	K _{app,ml}
ASCBPL	6.13234	1	K _{app,ml}
ADOCBLS	4.193462	1	K _{app,ml}
ADOCBIK	2.721728	1	K _{app,ml}
MCITL2	3.450673	1	K _{app,ml}
MCITL2	3.298899	-1	K _{app,ml}
DXYLK	2.892534	1	K _{app,ml}
MDH3	1.502434	1	K _{app,ml}
SOTA	3.569773	1	K _{app,ml}
DHCINDO	3.193946	1	K _{app,ml}
SGSAD	2.418219	1	K _{app,ml}
HKND DH	10.07529	1	K _{app,ml}
HKNTDH	9.829484	1	K _{app,ml}
PTA2	1.524221	1	K _{app,ml}
MDH2	15.33422	1	K _{app,ml}
ACOTA	6.936054	-1	K _{app,ml}
GPDDA5	13.05643	1	K _{app,ml}
PDX5PS	0.091514	1	K _{app,ml}
ENTCS	1.514259	1	K _{app,ml}
SULR	4.395033	1	K _{app,ml}
MANPGH	5.755779	1	K _{app,ml}
URDGLYCD	3.287935	1	K _{app,ml}
CBMKr	12.49795	1	K _{app,ml}
CBMKr	4.845883	-1	K _{app,ml}
PRAGSr	4.628824	-1	K _{app,ml}
AMANAPEr	4.245241	1	K _{app,ml}
AMANAPEr	4.245241	-1	K _{app,ml}
ACM6PH	2.703964	1	K _{app,ml}
NNATr	3.476746	-1	K _{app,ml}
MAN1PT2	4.021793	1	K _{app,ml}
AP5AH	11.17604	1	K _{app,ml}
GAPD	31.49729	-1	K _{app,ml}
MCITS	2.829793	1	K _{app,ml}
DM_amob_c	1.484089	1	K _{app,ml}
PPM2	2.709613	-1	K _{app,ml}
G6PDH2r	7.322026	-1	K _{app,ml}
ARGSL	5.957479	-1	K _{app,ml}
NADH5	2.403306	1	K _{app,ml}
MLTG2	5.614094	1	K _{app,ml}
MLTP3	2.619613	1	K _{app,ml}
MLTP3	1.569838	-1	K _{app,ml}
SPODM	1.887779	1	K _{app,ml}
TMK	3.529747	1	K _{app,ml}
NTPP4	14.51541	1	K _{app,ml}
GMPS	6.336127	1	K _{app,ml}
CAT	4.358878	1	K _{app,ml}
SERASr	2.660067	-1	K _{app,ml}
PPPND0	2.766831	1	K _{app,ml}
UACMAMO	2.462367	1	K _{app,ml}
FBA3	6.447201	-1	K _{app,ml}
LCADi	2.385278	1	K _{app,ml}
ACACT7r	9.086134	1	K _{app,ml}
ACACT7r	2.967914	-1	K _{app,ml}
ADK3	4.014544	-1	K _{app,ml}
ACACT2r	10.82815	1	K _{app,ml}
ACACT2r	3.631752	-1	K _{app,ml}
BSORy	2.362444	1	K _{app,ml}
DMSOR2	2.26961	1	K _{app,ml}
ACACT3r	8.939981	1	K _{app,ml}
ACACT3r	2.697898	-1	K _{app,ml}
ACACT4r	11.08831	1	K _{app,ml}
ACACT4r	3.64321	-1	K _{app,ml}
HSDy	10.56054	-1	K _{app,ml}
ACMAMUT	3.365882	1	K _{app,ml}
SSALy	12.18545	1	K _{app,ml}
DDGALK	3.806318	1	K _{app,ml}

AB6PGH	3.909703	1	K _{app,ml}
AHGDx	4.427314	1	K _{app,ml}
AHGDx	2.770437	-1	K _{app,ml}
3HCINNMH	2.676857	1	K _{app,ml}
BSORx	2.895214	1	K _{app,ml}
GP4GH	12.84317	1	K _{app,ml}
AADDGT	3.399368	1	K _{app,ml}
SPODMpp	3.15388	1	K _{app,ml}
PHYTSpp	12.8032	1	K _{app,ml}
PPTHpp	4.950185	1	K _{app,ml}
THD2pp	3.840832	1	K _{app,ml}
TMAOR2pp	2.447084	1	K _{app,ml}
FDMO	2.330453	1	K _{app,ml}
CHOLD	3.420031	1	K _{app,ml}
TKT2	3.049336	-1	K _{app,ml}
CYTK1	4.524805	-1	K _{app,ml}
DMSOR1pp	2.464278	1	K _{app,ml}
CYTK2	3.25772	-1	K _{app,ml}
ECOAH8	1.932477	1	K _{app,ml}
ECOAH8	2.418335	-1	K _{app,ml}
GLCDpp	4.155313	1	K _{app,ml}
HYD2pp	2.435902	1	K _{app,ml}
GK1	5.679219	-1	K _{app,ml}
HACD1	3.683205	1	K _{app,ml}
HACD1	2.561001	-1	K _{app,ml}
HACD2	3.581255	1	K _{app,ml}
HACD2	2.501425	-1	K _{app,ml}
ECOAH5	5.841235	1	K _{app,ml}
ECOAH5	3.218489	-1	K _{app,ml}
CYTDpp	2.540151	1	K _{app,ml}
CYTD2pp	3.52715	1	K _{app,ml}
ASPO3	1.043996	1	K _{app,ml}
GLYOX3	8.840878	1	K _{app,ml}
FDMO3	2.330453	1	K _{app,ml}
O16GLCT1	3.027211	1	K _{app,ml}
O16AP2pp	2.563836	1	K _{app,ml}
O16AP3pp	2.563836	1	K _{app,ml}
ACACT5r	8.939981	1	K _{app,ml}
ACACT5r	2.697898	-1	K _{app,ml}
O16A4Lpp	3.819881	1	K _{app,ml}
MALTATr	3.703914	1	K _{app,ml}
MALTATr	3.703914	-1	K _{app,ml}
DHBSH	7.098512	1	K _{app,ml}
MALDDH	2.52877	1	K _{app,ml}
ENTERES	17.49672	1	K _{app,ml}
TRSARr	4.313351	1	K _{app,ml}
TRSARr	34.25961	-1	K _{app,ml}
MTHFC	10.24086	-1	K _{app,ml}
MTHFD	8.114772	-1	K _{app,ml}
LYXI	2.791214	1	K _{app,ml}
LADGMDH	7.882974	1	K _{app,ml}
ALKP	4.550678	1	K _{app,ml}
ALAGLUE	2.794805	1	K _{app,ml}
ALAGLUE	2.794805	-1	K _{app,ml}
FE3HOXR2	3.487541	1	K _{app,ml}
ACACT6r	9.712107	1	K _{app,ml}
ACACT6r	3.645638	-1	K _{app,ml}
GGGABAH	4.619979	1	K _{app,ml}
FRULYSK	3.417939	1	K _{app,ml}
FDMO6	2.330453	1	K _{app,ml}
LIPAbcpp	2.50442	1	K _{app,ml}
LIPACabcpp	2.50442	1	K _{app,ml}
ECAP1pp	2.797934	1	K _{app,ml}
GGGABADr	2.886939	1	K _{app,ml}
GGGABADr	2.285701	-1	K _{app,ml}
ECAP2pp	2.546605	1	K _{app,ml}
ECA4OALpp	3.819881	1	K _{app,ml}
FE3Ri	2.640478	1	K _{app,ml}
FE3HOXR3	3.487541	1	K _{app,ml}
FEENTERR1	3.487541	1	K _{app,ml}
FEENTERR3	3.487541	1	K _{app,ml}
LA4NTpp	2.785555	1	K _{app,ml}
AGM4PA	3.051655	1	K _{app,ml}
RNDR4b	0.078608	1	K _{app,ml}
UDPGALPpp	5.821825	1	K _{app,ml}
UACGAMPpp	5.821825	1	K _{app,ml}

UACGALPpp	5.821825	1	k _{app,ml}
UGLCURPpp	5.821825	1	k _{app,ml}
GUR1PPpp	7.098512	1	k _{app,ml}
ACGAL1Ppp	21.87028	1	k _{app,ml}
AM4PCP	3.733613	1	k _{app,ml}
AM3PA	3.051655	1	k _{app,ml}
UM4PCP	3.733613	1	k _{app,ml}
CBLAT	3.411688	-1	k _{app,ml}
NTRIR4pp	2.517789	1	k _{app,ml}
NTRIR3pp	2.517789	1	k _{app,ml}
AGM3PA	3.051655	1	k _{app,ml}
ACANTHAT	3.407813	1	k _{app,ml}
FECRMR1	3.487541	1	k _{app,ml}
FECRMR2	3.487541	1	k _{app,ml}
TDP	4.813343	1	k _{app,ml}
3PEPTabcpp	2.526631	1	k _{app,ml}
4PEPTabcpp	2.525389	1	k _{app,ml}
AGM3Pt2pp	3.755548	1	k _{app,ml}
AGM4Pt2pp	3.755548	1	k _{app,ml}
AGMt2pp	3.755548	1	k _{app,ml}
ARBTNR1	3.487541	1	k _{app,ml}
ARBTNR2	3.487541	1	k _{app,ml}
METSXR1	2.295099	1	k _{app,ml}
METSXR2	5.168006	1	k _{app,ml}
DAPAL	3.73781	1	k _{app,ml}
AMPTASECG	11.68583	1	k _{app,ml}
ARBTNR3	3.487541	1	k _{app,ml}
FEOXAMR1	3.847276	1	k _{app,ml}
FEOXAMR2	3.847276	1	k _{app,ml}
FEOXAMR3	3.847276	1	k _{app,ml}
PTHRpp	4.226436	1	k _{app,ml}
FEENTERR2	3.487541	1	k _{app,ml}
FE3HOXR1	3.487541	1	k _{app,ml}
FECRMR3	3.487541	1	k _{app,ml}
FDMO2	2.330453	1	k _{app,ml}
LALGP	3.67872	1	k _{app,ml}
AGM4PApp	3.143364	1	k _{app,ml}
ARABDI	2.646269	1	k _{app,ml}
ARABDI	2.646269	-1	k _{app,ml}
R5PPpp	4.226436	1	k _{app,ml}
R1PK	12.56464	1	k _{app,ml}
GRXR	6.532341	1	k _{app,ml}
RNDR1b	0.292833	1	k _{app,ml}
R15BPK	12.72596	1	k _{app,ml}
RNDR2b	0.890565	1	k _{app,ml}
FACOA181	8.936187	1	k _{app,ml}
FACOA160	11.48934	1	k _{app,ml}
FACOA140	3.671227	1	k _{app,ml}
CPGMR1	3.487541	1	k _{app,ml}
CPGMR2	3.487541	1	k _{app,ml}
CPGMR3	3.487541	1	k _{app,ml}
AAMYL	6.124081	1	k _{app,ml}
AAMYLpp	4.282791	1	k _{app,ml}
GGPTRCS	2.435125	1	k _{app,ml}
23PDE2pp	3.540425	1	k _{app,ml}
23PDE4pp	3.540425	1	k _{app,ml}
MN6PP	4.896934	1	k _{app,ml}
3KGK	2.823988	1	k _{app,ml}
F6PP	4.692017	1	k _{app,ml}
FRULYSDG	10.78971	1	k _{app,ml}
FRULYSDG	3.768395	-1	k _{app,ml}
GDPMNH	3.105539	1	k _{app,ml}
GDPDPK	2.457914	1	k _{app,ml}
DNTPPA	1.573331	1	k _{app,ml}
NTPP11	4.533803	1	k _{app,ml}
ALLPI	4.119161	1	k _{app,ml}
ALLPI	4.119161	-1	k _{app,ml}
ALLK	3.550841	1	k _{app,ml}
ALLLULPE	4.154883	1	k _{app,ml}
ALLLULPE	3.402488	-1	k _{app,ml}
NADPHQR2	3.564066	1	k _{app,ml}
NADPHQR3	3.564066	1	k _{app,ml}
NADPHQR4	3.564066	1	k _{app,ml}
GLBRAN2	2.464644	1	k _{app,ml}
ASR	2.47837	1	k _{app,ml}
XTSNH	3.447988	1	k _{app,ml}

ECAtp	9.168891	1	k _{app,ml}
O16AT	5.35852	1	k _{app,ml}
O16GALFT	3.460466	1	k _{app,ml}
O16GLCT2	3.487541	1	k _{app,ml}
O16AUNDtp	8.982407	1	k _{app,ml}
O16AP1pp	2.867069	1	k _{app,ml}
AGM4PCP	3.733613	1	k _{app,ml}
4PCP	3.733613	1	k _{app,ml}
4PCPpp	3.280727	1	k _{app,ml}
FALDH2	2.672188	-1	k _{app,ml}
CBIAT	3.411688	-1	k _{app,ml}
PPK2	2.665989	1	k _{app,ml}
DHPTPE	5.442813	1	k _{app,ml}
DHPTPE	5.442813	-1	k _{app,ml}
PPDOy	2.810607	1	k _{app,ml}
PPA2	1.569796	1	k _{app,ml}
GNK	26.65476	1	k _{app,ml}
3NTD7pp	3.487722	1	k _{app,ml}
3NTD4pp	3.80098	1	k _{app,ml}
RMK	2.881141	1	k _{app,ml}
FHL	2.524572	1	k _{app,ml}
HYPGE	21.87028	1	k _{app,ml}
NTPP5	15.04929	1	k _{app,ml}
ACONMT	3.880165	1	k _{app,ml}
NADPPPS	7.680184	1	k _{app,ml}
THRD_L	27.91808	1	k _{app,ml}
NAMNPP	2.633579	1	k _{app,ml}
SELNPS	5.118439	1	k _{app,ml}
MTRPOX	2.482688	1	k _{app,ml}
AGM4PCPpp	3.280727	1	k _{app,ml}
CRNCAR	8.097315	1	k _{app,ml}
CRNCAR	8.097315	-1	k _{app,ml}
RNDR2	1.810068	1	k _{app,ml}
ANHMK	3.539172	1	k _{app,ml}
DHFR	2.448391	-1	k _{app,ml}
NMNN	11.29015	1	k _{app,ml}
XAND	2.26533	1	k _{app,ml}
HACD3	3.592268	1	k _{app,ml}
HACD3	2.493227	-1	k _{app,ml}
NNDMBRT	4.321261	1	k _{app,ml}
THRA2	66.09963	1	k _{app,ml}
PPK	2.665989	1	k _{app,ml}
SADH	14.35947	1	k _{app,ml}
E4PD	0.038994	-1	k _{app,ml}
IG3PS	5.01309	1	k _{app,ml}
PDH	71.8999	1	k _{app,ml}
DADK	4.360949	1	k _{app,ml}
DADK	2.375471	-1	k _{app,ml}
HYD3pp	2.435902	1	k _{app,ml}
AACPS5	1.294524	1	k _{app,ml}
MLTG4	0.272841	1	k _{app,ml}
UDPGDC	5.371394	1	k _{app,ml}
AMAOTr	3.170838	-1	k _{app,ml}
MMCD	4.067714	1	k _{app,ml}
UDPKAAT	3.291144	1	k _{app,ml}
UDPKAAT	3.65951	-1	k _{app,ml}
ULA4NFT	3.740215	1	k _{app,ml}
DKGLCNR2x	2.17844	1	k _{app,ml}
UPLA4FNT	4.922288	1	k _{app,ml}
UPLA4FNF	11.21595	1	k _{app,ml}
ULA4Ntpi	7.353592	1	k _{app,ml}
MCITD	3.417475	1	k _{app,ml}
GLCATr	3.703914	1	k _{app,ml}
GLCATr	3.703914	-1	k _{app,ml}
ALR2	2.438322	1	k _{app,ml}
ALAALAR	2.462285	-1	k _{app,ml}
HXAND	2.26533	1	k _{app,ml}
G2PPpp	4.226436	1	k _{app,ml}
ALDD3y	2.909703	1	k _{app,ml}
R5PP	3.409544	1	k _{app,ml}
CYSSADS	2.872551	1	k _{app,ml}
CPPPGO2	2.407467	1	k _{app,ml}
AGM3PH	2.831628	1	k _{app,ml}
AGM3PApp	3.143364	1	k _{app,ml}
LALDO3	2.75933	1	k _{app,ml}
NTP10	12.81817	1	k _{app,ml}

GDMANE	2.644533	1	K_app,ml
LPLIPAL2ATG1	3.251202	1	K_app,ml
PLIPA1E140pp	2.514084	1	K_app,ml
LPLIPAL2E140	3.055662	1	K_app,ml
PLIPA1E160pp	3.462475	1	K_app,ml
GALKr	10.22032	-1	K_app,ml
PLIPA1E161pp	2.514084	1	K_app,ml
AACPS8	2.004406	1	K_app,ml
AACPS9	0.00694	1	K_app,ml
LPLIPAL2E180	3.055662	1	K_app,ml
LPLIPAL2G161	3.055662	1	K_app,ml
LPLIPAL2G181	3.055662	1	K_app,ml
HXCT	2.752224	1	K_app,ml
AACPS6	2.004406	1	K_app,ml
2AGPEAT160	2.044477	1	K_app,ml
AACPS7	0.900429	1	K_app,ml
DURADx	2.372947	1	K_app,ml
DURADx	2.477274	-1	K_app,ml
ECOAH7	4.864677	1	K_app,ml
ECOAH7	2.713135	-1	K_app,ml
GALCTND	3.347427	1	K_app,ml
QMO2	1.483889	1	K_app,ml
QMO3	1.896808	1	K_app,ml
LPLIPAL2E141	3.055662	1	K_app,ml
LPLIPAL2E161	3.055662	1	K_app,ml
LDH_D	5.601872	1	K_app,ml
LPLIPAL2E181	3.055662	1	K_app,ml
YUMPS	3.840615	1	K_app,ml
YUMPS	3.606845	-1	K_app,ml
LPLIPAL2G180	3.055662	1	K_app,ml
LPLIPAL2G141	3.055662	1	K_app,ml
IPDDI	0.577649	-1	K_app,ml
T2DECAI	2.653291	-1	K_app,ml
2AGPGAT180	2.044477	1	K_app,ml
2AGPGAT181	2.044477	1	K_app,ml
2AGPEAT140	2.044477	1	K_app,ml
2AGPEAT120	2.044477	1	K_app,ml
CDAPPA161	5.716151	1	K_app,ml
2AGPEAT161	2.044477	1	K_app,ml
2AGPEAT141	2.044477	1	K_app,ml
2AGPGAT160	2.044477	1	K_app,ml
PGSA180	5.363519	1	K_app,ml
PSSA140	2.765518	1	K_app,ml
2AGPGAT141	2.044477	1	K_app,ml
FACOA180	7.34444	1	K_app,ml
LPLIPAL2ATG1	3.251202	1	K_app,ml
LPLIPAL2ATG1	3.251202	1	K_app,ml
LPLIPAL2ATG1	3.251202	1	K_app,ml
LPLIPAL2ATG1	3.251202	1	K_app,ml
LPLIPAL2ATG1	3.251202	1	K_app,ml
LPLIPAL2ATG1	3.251202	1	K_app,ml
FACOA100	7.34444	1	K_app,ml
FACOA180	0.001362	1	K_app,ml
FACOA160	7.34444	1	K_app,ml
CLPNS180pp	3.757662	1	K_app,ml
CLPNS180pp	2.569282	-1	K_app,ml
PLIPA1E120pp	2.514084	1	K_app,ml
CLPNS181pp	0.194469	1	K_app,ml
CLPNS181pp	2.569282	-1	K_app,ml
PLIPA1G120pp	2.514084	1	K_app,ml
FACOA141	7.34444	1	K_app,ml
CLPNS120pp	3.757662	1	K_app,ml
CLPNS120pp	2.569282	-1	K_app,ml
ALPATE160pp	2.404651	1	K_app,ml
FACOA161	10.52334	1	K_app,ml
ALPATG160pp	2.404651	1	K_app,ml
2AGPGAT120	2.044477	1	K_app,ml
LPLIPAL2ATE1	3.251202	1	K_app,ml
ALAALAD	3.76856	1	K_app,ml
2AGPEAT180	2.044477	1	K_app,ml
LPLIPAL2ATE1	3.251202	1	K_app,ml
LPLIPAL2ATE1	3.251202	1	K_app,ml
LPLIPAL2ATE1	3.251202	1	K_app,ml
LPLIPAL2ATE1	3.251202	1	K_app,ml
2AGPEAT181	2.044477	1	K_app,ml
2AGPGAT140	2.044477	1	K_app,ml

PLIPA1E181pp	2.514084	1	K_app,ml
PLIPA1G180pp	2.514084	1	K_app,ml
LPLIPAL2ATE1	3.251202	1	K_app,ml
LPLIPAL2ATE1	3.251202	1	K_app,ml
PLIPA1G181pp	2.514084	1	K_app,ml
2AGPGAT16l	2.044477	1	K_app,ml
AMPTASEPG	11.68583	1	K_app,ml
ADSL1r	3.15405	-1	K_app,ml
PGPP140pp	4.251784	1	K_app,ml
PGPP141pp	4.251784	1	K_app,ml
PGPP160pp	4.251784	1	K_app,ml
PGPP180pp	4.251784	1	K_app,ml
CLPNH141pp	7.098512	1	K_app,ml
CLPNH160pp	7.098512	1	K_app,ml
CLPNH161pp	7.098512	1	K_app,ml
AICART	3.406534	-1	K_app,ml
CLPNH181pp	7.098512	1	K_app,ml
PLIPA1E180pp	3.462475	1	K_app,ml
PLIPA1G140pp	2.514084	1	K_app,ml
PLIPA1G161pp	2.514084	1	K_app,ml
CLPNS160pp	0.389207	1	K_app,ml
CLPNS160pp	2.569282	-1	K_app,ml
CLPNS161pp	0.326188	1	K_app,ml
CLPNS161pp	2.569282	-1	K_app,ml
PAPA120pp	7.234931	1	K_app,ml
PAPA140pp	7.234931	1	K_app,ml
GGPTRCO	4.320293	1	K_app,ml
PAPA141pp	7.234931	1	K_app,ml
PAPA160pp	7.234931	1	K_app,ml
PAPA161pp	7.234931	1	K_app,ml
PAPA180pp	7.234931	1	K_app,ml
HPYRI	4.277075	1	K_app,ml
HPYRI	31.1465	-1	K_app,ml
HYD1pp	2.418675	1	K_app,ml
AACTOOR	2.496258	1	K_app,ml
ECAP3pp	2.546605	1	K_app,ml
TPRDCOAS	7.748391	1	K_app,ml
ACGAM1PPpp	21.87028	1	K_app,ml
THIORDXi	5.927244	1	K_app,ml
PDX5P02	1.510259	1	K_app,ml
MDDCP1pp	1.353983	1	K_app,ml
GUI2	2.725522	1	K_app,ml
GUI2	2.725522	-1	K_app,ml
PLIPA1A181pp	2.514084	1	K_app,ml
PGM	65.56609	1	K_app,ml
LPLIPAL2A141	2.809181	1	K_app,ml
MLTG5	0.272841	1	K_app,ml
G3PAT141	2.571933	1	K_app,ml
2DOXG6PP	4.038472	1	K_app,ml
OXAMTC	3.487541	1	K_app,ml
UDCPTppi	7.098512	1	K_app,ml
AGMH	2.831628	1	K_app,ml
G3PAT181	2.180277	1	K_app,ml
MPTG	1.881249	1	K_app,ml
MPTG2	0.156749	1	K_app,ml
MCTP1AApp	1.791982	1	K_app,ml
MCTP2AApp	0.216502	1	K_app,ml
CDAPPA160	5.716151	1	K_app,ml
CDAPPA141	5.716151	1	K_app,ml
MCTP1Bpp	0.320418	1	K_app,ml
PERD	0.078893	-1	K_app,ml
PGSA141	5.363519	1	K_app,ml
DM_sdrib_c	1.525218	1	K_app,ml
3OAR120	3.967411	-1	K_app,ml
3HAD80	2.407494	1	K_app,ml
3HAD120	2.407494	1	K_app,ml
DSBGGT	3.246589	1	K_app,ml
EAR180x	2.290886	1	K_app,ml
AGPAT120	6.658908	1	K_app,ml
AMPMS2	1.542061	1	K_app,ml
DM_oxam_c	7.098512	1	K_app,ml
ALR4x	2.499005	1	K_app,ml
PGPP120	4.251784	1	K_app,ml
DAGK120	5.427744	1	K_app,ml
PSSA180	2.765518	1	K_app,ml
PSSA141	2.765518	1	K_app,ml

PSD180	3.061142	1	K app,ml
LPLIPAL1G141	3.324473	1	K app,ml
LPLIPAL1G180	3.171706	1	K app,ml
LPLIPAL1G181	3.324473	1	K app,ml
ACONTb	14.12519	-1	K app,ml
PLIPA2A140pp	3.462475	1	K app,ml
PLIPA2A141pp	3.462475	1	K app,ml
PLIPA2A180pp	3.462475	1	K app,ml
PLIPA2E141pp	2.514084	1	K app,ml
PLIPA2E181pp	2.514084	1	K app,ml
PLIPA2G140pp	2.514084	1	K app,ml
3HAD161	2.407494	1	K app,ml
PLIPA2G160pp	2.514084	1	K app,ml
CU1Opp	2.852558	1	K app,ml
PLIPA2G181pp	2.514084	1	K app,ml
FADRx2	13.29396	1	K app,ml
LPLIPAL1E141	3.324473	1	K app,ml
LPLIPAL1E160	3.324473	1	K app,ml
LPLIPAL1E180	3.324473	1	K app,ml
3HAD121	2.407494	1	K app,ml
LPLIPAL1E181	3.324473	1	K app,ml
LPLIPAL1G160	3.324473	1	K app,ml
LPLIPAL1G161	3.324473	1	K app,ml
GTHRDHpp	2.907945	1	K app,ml
MDDCP2pp	0.082167	1	K app,ml
LPLIPAL1G120	3.324473	1	K app,ml
PLIPA2E140pp	2.514084	1	K app,ml
PLIPA2E160pp	2.514084	1	K app,ml
PLIPA2A161pp	2.514084	1	K app,ml
PLIPA2A181pp	2.514084	1	K app,ml
MDDCP3pp	0.235402	1	K app,ml
G3PAT140	2.571933	1	K app,ml
G3PAT160	2.853299	1	K app,ml
G3PAT161	2.619723	1	K app,ml
AGPAT140	6.658908	1	K app,ml
MLDCP2Bpp	3.160439	1	K app,ml
3OAR140	3.967411	-1	K app,ml
CDAPPA140	5.716151	1	K app,ml
MLDCP1App	0.103874	1	K app,ml
3OAR160	3.967411	-1	K app,ml
3HAD181	2.407494	1	K app,ml
PGPP160	2.609757	1	K app,ml
3HAD141	2.407494	1	K app,ml
PGPP180	4.251784	1	K app,ml
PGPP161	2.603426	1	K app,ml
PGPP181	1.266994	1	K app,ml
PSD140	3.061142	1	K app,ml
MICITDr	2.729069	1	K app,ml
MICITDr	2.729069	-1	K app,ml
CDAPPA180	5.716151	1	K app,ml
AGM4PH	2.831628	1	K app,ml
NTP12	12.44237	1	K app,ml
CDAPPA181	5.716151	1	K app,ml
PGSA140	5.363519	1	K app,ml
LPLIPAL1A141	3.324473	1	K app,ml
LPLIPAL1A180	3.324473	1	K app,ml
LPLIPAL1E140	3.324473	1	K app,ml
LPLIPAL1E161	3.324473	1	K app,ml
LPLIPAL1G140	3.324473	1	K app,ml
PSD141	3.061142	1	K app,ml
MLDCP2App	3.160439	1	K app,ml
AM4PA	3.051655	1	K app,ml
PGPP120pp	4.251784	1	K app,ml
PGPP181pp	4.251784	1	K app,ml
CLPNH120pp	7.098512	1	K app,ml
PAPA181pp	7.234931	1	K app,ml
CLPNH140pp	7.098512	1	K app,ml
PGPP140	4.251784	1	K app,ml
PGPP141	4.251784	1	K app,ml
MDDCP5pp	3.133974	1	K app,ml
MLTGY3pp	2.725129	1	K app,ml
MDDCP4pp	0.532646	1	K app,ml
CLPNS140pp	3.757662	1	K app,ml
CLPNS140pp	2.569282	-1	K app,ml
CLPNS141pp	3.757662	1	K app,ml
CLPNS141pp	2.569282	-1	K app,ml

G3PAT180	2.571933	1	K _{app,ml}
PLIPA1G160pp	2.514084	1	K _{app,ml}
PLIPA1G141pp	2.514084	1	K _{app,ml}
LPLIPAL2E160	3.055662	1	K _{app,ml}
LPLIPAL2G140	3.055662	1	K _{app,ml}
LPLIPAL2G160	3.055662	1	K _{app,ml}
PAPA160	7.234931	1	K _{app,ml}
PAPA180	7.234931	1	K _{app,ml}
PAPA181	7.234931	1	K _{app,ml}
LPLIPAL2G120	3.055662	1	K _{app,ml}
PETNT161pp	2.758737	1	K _{app,ml}
DAGK141	5.427744	1	K _{app,ml}
DAGK161	5.427744	1	K _{app,ml}
PGPP161pp	4.251784	1	K _{app,ml}
MDDEP3pp	0.230219	1	K _{app,ml}
MDDEP4pp	3.244021	1	K _{app,ml}
PAPA140	7.234931	1	K _{app,ml}
MLTGY1pp	2.725129	1	K _{app,ml}
3HAD40	2.407494	1	K _{app,ml}
PLIPA1E141pp	2.514084	1	K _{app,ml}
PAPA141	7.234931	1	K _{app,ml}
PAPA161	7.234931	1	K _{app,ml}
DSBAO1	3.832589	1	K _{app,ml}
DSBAO2	3.832589	1	K _{app,ml}
DSBDR	2.587281	1	K _{app,ml}
PLIPA2A120pp	3.462475	1	K _{app,ml}
PLIPA2A160pp	3.462475	1	K _{app,ml}
CLPNH180pp	7.098512	1	K _{app,ml}
3OAR60	3.967411	-1	K _{app,ml}
3OAR80	3.967411	-1	K _{app,ml}
MLDCP3App	0.174891	1	K _{app,ml}
MLDCP1Bpp	3.160439	1	K _{app,ml}
LPLIPAL2E120	3.055662	1	K _{app,ml}
ATHRDHr	2.400645	1	K _{app,ml}
ATHRDHr	16.03101	-1	K _{app,ml}
LSERDhr	2.452173	-1	K _{app,ml}
3OAR40	2.660884	-1	K _{app,ml}
3OAR100	3.967411	-1	K _{app,ml}
DSERDhr	3.00543	1	K _{app,ml}
DSERDhr	4.674328	-1	K _{app,ml}
3HAD60	2.745113	1	K _{app,ml}
3HAD100	2.407494	1	K _{app,ml}
3HAD140	2.745113	1	K _{app,ml}
3HAD160	2.745113	1	K _{app,ml}
PSD120	3.061142	1	K _{app,ml}
AGPAT180	6.658908	1	K _{app,ml}
AGPAT141	6.658908	1	K _{app,ml}
PAPA120	7.234931	1	K _{app,ml}
MDDEP1pp	0.192692	1	K _{app,ml}
MDDEP2pp	3.244021	1	K _{app,ml}
MLDEP1pp	1.717065	1	K _{app,ml}
DAGK180	5.427744	1	K _{app,ml}
MLDEP2pp	7.648413	1	K _{app,ml}
MLTGY2pp	2.725129	1	K _{app,ml}
PLIPA2E120pp	2.514084	1	K _{app,ml}
PLIPA2G141pp	2.514084	1	K _{app,ml}
MLTGY4pp	2.725129	1	K _{app,ml}
PLIPA2G161pp	2.514084	1	K _{app,ml}
LPLIPAL1A120	3.324473	1	K _{app,ml}
LPLIPAL1A140	3.324473	1	K _{app,ml}
LPLIPAL1A181	3.324473	1	K _{app,ml}
LPLIPAL1E120	3.324473	1	K _{app,ml}
3OAR180	2.660884	-1	K _{app,ml}
G3PAT120	2.571933	1	K _{app,ml}
PGSA120	5.363519	1	K _{app,ml}
3HAD180	2.407494	1	K _{app,ml}
CDAPPA120	5.716151	1	K _{app,ml}
OMMBLHX3	1.507884	1	K _{app,ml}
PSSA120	2.765518	1	K _{app,ml}
UM4PL	2.563139	1	K _{app,ml}
2MAHMP	3.462365	1	K _{app,ml}
NADHPO	4.510757	1	K _{app,ml}
ACOAD7f	2.822655	1	K _{app,ml}
ACOAD7f	5.366348	-1	K _{app,ml}
ACOAD8f	2.822655	1	K _{app,ml}
ACOAD8f	2.822655	-1	K _{app,ml}

PLIPA1A120pp	3.462475	1	K_app,ml
ACOAD5f	3.666903	1	K_app,ml
ACOAD5f	8.701169	-1	K_app,ml
PLIPA1A140pp	3.462475	1	K_app,ml
PLIPA1A141pp	3.462475	1	K_app,ml
PLIPA1A160pp	3.462475	1	K_app,ml
PLIPA1A161pp	2.514084	1	K_app,ml
PLIPA1A180pp	3.462475	1	K_app,ml
LPLIPAL2A120	2.809181	1	K_app,ml
ACOAD3f	3.639979	1	K_app,ml
ACOAD3f	8.652383	-1	K_app,ml
ACOAD2f	2.822655	1	K_app,ml
ACOAD2f	7.085496	-1	K_app,ml
ACOAD4f	2.822655	1	K_app,ml
ACOAD4f	7.085496	-1	K_app,ml
ACOAD6f	7.639588	-1	K_app,ml
KDUI	11.50452	1	K_app,ml
CTBTCAL2	2.501161	1	K_app,ml
OBTFL	2.620728	1	K_app,ml
FA161ACPHi	55.65879	1	K_app,ml
BGLA1	4.74562	1	K_app,ml
FA141ACPHi	55.65879	1	K_app,ml
FA160ACPHi	55.65879	1	K_app,ml
GDPMNP	16.35714	1	K_app,ml
OPHHX3	1.507884	1	K_app,ml
FA120ACPHi	55.65879	1	K_app,ml
FA80ACPHi	55.65879	1	K_app,ml
FA140ACPHi	55.65879	1	K_app,ml
FA100ACPHi	55.65879	1	K_app,ml
CRNDAL2	2.501161	1	K_app,ml
CRNCAL2	2.501161	1	K_app,ml
PLIPA2E161pp	2.514084	1	K_app,ml
PLIPA2E180pp	2.514084	1	K_app,ml
FRULYSE	3.410595	1	K_app,ml
FRULYSE	3.410595	-1	K_app,ml
PLIPA2G120pp	2.514084	1	K_app,ml
PLIPA2G180pp	2.127453	1	K_app,ml
LPLIPAL1A161	3.324473	1	K_app,ml
LPLIPAL1A160	3.324473	1	K_app,ml
LPLIPAL2A140	2.809181	1	K_app,ml
LPLIPAL2A160	2.809181	1	K_app,ml
LPLIPAL2A161	2.809181	1	K_app,ml
LPLIPAL2A180	2.809181	1	K_app,ml
LPLIPAL2A181	2.809181	1	K_app,ml
DM_aacald_c	7.098512	1	K_app,ml
UM3PL	2.563139	1	K_app,ml
GALCTLO	2.881793	1	K_app,ml
POR5	3.980006	1	K_app,ml
POR5	6.197212	-1	K_app,ml
HACD8	11.80702	1	K_app,ml
HACD8	8.339206	-1	K_app,ml
CTECOAI6	6.982129	-1	K_app,ml
SELGTHR	3.487541	1	K_app,ml
SELGTHR3	2.723451	1	K_app,ml
THRA	4.288814	1	K_app,ml
DGUNC	3.786562	1	K_app,ml
XYLK2	2.797534	1	K_app,ml
GLDBRAN2	2.334182	1	K_app,ml
S2FE2SR	2.15723	1	K_app,ml
CDGUNPD	3.530083	1	K_app,ml
ACACT8r	2.697898	1	K_app,ml
ACACT8r	3.686624	-1	K_app,ml
CTECOAI7	6.982129	1	K_app,ml
CTECOAI7	11.53259	-1	K_app,ml
3AMACHYD	33.13717	1	K_app,ml
FDH4pp	25.79707	1	K_app,ml
FDH5pp	0.00855	1	K_app,ml
ACPPAT140	2.410531	1	K_app,ml
APG3PAT140	4.890185	1	K_app,ml
LDGUNPD	3.321879	1	K_app,ml
APG3PAT141	4.890185	1	K_app,ml
MOADSUX	1.495763	1	K_app,ml
APG3PAT181	3.416439	1	K_app,ml
APH140	3.12568	1	K_app,ml
APH160	3.12568	1	K_app,ml
SCYSDS	3.198283	1	K_app,ml

S2FE2SS	2.059549	1	K_app,ml
S4FE4SR	2.467245	1	K_app,ml
CCGS	2.838964	1	K_app,ml
REPHACCOAI	4.078866	1	K_app,ml
REPHACCOAI	4.078866	-1	K_app,ml
ACPPAT141	2.410531	1	K_app,ml
ACPPAT161	2.087464	1	K_app,ml
ACPPAT180	2.410531	1	K_app,ml
APG3PAT160	4.541992	1	K_app,ml
APG3PAT161	4.15706	1	K_app,ml
APG3PAT180	4.890185	1	K_app,ml
MECDPDH5	1.869982	1	K_app,ml
CDGR	2.505639	1	K_app,ml
RNTR1c2	2.218289	1	K_app,ml
RNTR2c2	2.260405	1	K_app,ml
S2FE2ST	2.68241	1	K_app,ml
BWCOS	2.326042	1	K_app,ml
RNTR3c2	1.99679	1	K_app,ml
RNTR4c2	1.498421	1	K_app,ml
BWCOGDS1	3.699292	1	K_app,ml
BWCOGDS2	3.699292	1	K_app,ml
CDGS	3.322615	1	K_app,ml
DHNPTE	4.361322	1	K_app,ml
DHNPTE	4.361322	-1	K_app,ml
DHMPTR	2.523332	1	K_app,ml
I2FE2SS	1.504195	1	K_app,ml
I2FE2ST	1.493889	1	K_app,ml
OXCOAHDH	2.613249	1	K_app,ml
DHACOAH	4.178044	1	K_app,ml
DHACOAH	4.178044	-1	K_app,ml
I2FE2SS2	1.502792	1	K_app,ml
MSAR	2.427211	1	K_app,ml
I4FE4ST	1.517756	1	K_app,ml
S4FE4ST	2.68241	1	K_app,ml
FOMETRI	3.875567	1	K_app,ml
S2FE2SS2	2.059549	1	K_app,ml
SELGTHR2	2.723451	1	K_app,ml
GDPTPDP	3.524073	1	K_app,ml
COLIPAKpp	3.829001	1	K_app,ml
ACPPAT120	2.410531	1	K_app,ml
CPMPS	0.099392	1	K_app,ml
LIPATPT	3.449978	1	K_app,ml
LIPAMPL	2.586752	1	K_app,ml
ALDD19xr	4.68611	-1	K_app,ml
MOGDS	1.489004	1	K_app,ml
CPH4S	3.212318	1	K_app,ml
WCOS	2.326042	1	K_app,ml
SHCHCS3	1.552127	1	K_app,ml
VPAMTr	4.407651	1	K_app,ml
PYK4	1.48396	1	K_app,ml
NADPHXD	2.314239	1	K_app,ml
NADPHXE	3.527344	1	K_app,ml
NADPHXE	3.527344	-1	K_app,ml
DM_4crsol_c	1.524038	1	K_app,ml
3OXCOAT	3.170324	1	K_app,ml
URACPAH	5.20523	1	K_app,ml
FE3DHBZS3R	3.665524	1	K_app,ml
DXYLTD	3.444396	1	K_app,ml
DXYLTD	3.444396	-1	K_app,ml
2DDARAA	3.116683	1	K_app,ml
2DDARAA	2.689897	-1	K_app,ml
2DGULRGx	2.17844	1	K_app,ml
2DGULRGy	2.448182	1	K_app,ml
AP4AS	2.655165	1	K_app,ml
GNP	1.572874	1	K_app,ml
ALLTAMH2	6.517188	1	K_app,ml
UGCIAMH	10.07596	1	K_app,ml
POAACR	2.992232	1	K_app,ml
METGLCUR	3.104906	1	K_app,ml
ACOAD1fr	9.260675	1	K_app,ml
FEENTERES	7.485505	1	K_app,ml
FEENTERR4	3.665524	1	K_app,ml
ACPPAT160	2.23661	1	K_app,ml
ALR2x	2.731465	1	K_app,ml
OMPHHXy	1.508531	1	K_app,ml
13PPDH2	2.791839	1	K_app,ml

13PPDH2	3.283804	-1	K_app,ml
APCS	2.507151	1	K_app,ml
ACPPAT181	1.990575	1	K_app,ml
SQVOSI	6.062962	1	K_app,ml
6D6SPA	9.53266	1	K_app,ml
ThDPAT	3.651278	1	K_app,ml
GHBDHx	2.477119	1	K_app,ml
GHBDHx	2.757048	-1	K_app,ml
APPLDhr	2.499005	1	K_app,ml
APPLDhr	2.827125	-1	K_app,ml
APH141	3.12568	1	K_app,ml
AI2K	1.660644	1	K_app,ml
G2PP	45.26527	1	K_app,ml
APH161	3.12568	1	K_app,ml
DHNPA2r	5.202772	-1	K_app,ml
SHGO	4.325558	1	K_app,ml
ACOXT	5.120754	1	K_app,ml
ACOXT	5.120754	-1	K_app,ml
2HESR	3.911573	1	K_app,ml
CHOLID	3.274975	1	K_app,ml
NADHXD	2.317967	1	K_app,ml
4ABZGLUH	4.851053	1	K_app,ml
4ABZGLUH	2.580011	-1	K_app,ml
APH180	3.12568	1	K_app,ml
NADHXE	3.527344	1	K_app,ml
NADHXE	3.527344	-1	K_app,ml
PHDA	4.501681	1	K_app,ml
TEO2M	3.591815	1	K_app,ml
PACOAT	5.182569	1	K_app,ml
PYK2	1.483969	1	K_app,ml
PYK3	1.484309	1	K_app,ml
APH181	3.12568	1	K_app,ml
DHPTDNR	2.959104	1	K_app,ml
DHPTDNRN	2.301584	1	K_app,ml
PCNO	2.794064	1	K_app,ml
PCNO	2.719302	-1	K_app,ml
PYK6	1.484091	1	K_app,ml
HPACOAT	5.182569	1	K_app,ml
DC6PDA	8.290936	1	K_app,ml
5DGLCNR	33.42809	1	K_app,ml
PAI2I	1.941962	1	K_app,ml
CURR	2.671109	1	K_app,ml
LKDRA	3.122071	1	K_app,ml
LKDRA	2.912668	-1	K_app,ml
METNA	3.817243	1	K_app,ml
METNA	3.817243	-1	K_app,ml
SDKGR	32.2425	1	K_app,ml
SDKGR	2.874367	-1	K_app,ml
XYHDL	2.822777	1	K_app,ml
E4PP	3.4766	1	K_app,ml
PUACGAMS	2.664868	1	K_app,ml
PUACGAMS	3.220147	-1	K_app,ml
RUSPP	3.226787	1	K_app,ml
6D6SFK	3.439494	1	K_app,ml
3SLAR	2.702708	1	K_app,ml
MOX	1.488573	1	K_app,ml
ARHGDx	4.789896	1	K_app,ml
ARHGDx	4.750349	-1	K_app,ml
CPL	11.76008	1	K_app,ml
CYSTA	2.456687	1	K_app,ml
FE3DCITR5	4.112798	1	K_app,ml
4HTHRA	8.34468	1	K_app,ml
4HTHRA	1.505022	-1	K_app,ml
I2FE2SR	1.485261	1	K_app,ml
GAPP	12.69453	1	K_app,ml
INOSTO	2.437373	1	K_app,ml
INOSTO	2.866705	-1	K_app,ml
THZPSN3	0.037488	1	K_app,ml
HPYRP	3.857914	1	K_app,ml
PSURIK	3.356983	1	K_app,ml
PAI2T	1.575669	1	K_app,ml
APG3PAT120	4.890185	1	K_app,ml
QUINDHyI	4.070744	1	K_app,ml
DHCURR	2.671109	1	K_app,ml
RHMND	4.664837	1	K_app,ml
RHMND	4.664837	-1	K_app,ml

3HBZCT	3.283962	1	k _{app,ml}
CXSAMS	4.072984	1	k _{app,ml}
PNSPA	4.831707	1	k _{app,ml}
4HTHRK	1.512763	1	k _{app,ml}
AMPNTAT	3.693533	1	k _{app,ml}
APH120	3.12568	1	k _{app,ml}
SQGH	2.964042	1	k _{app,ml}
QUINDH	3.095731	1	k _{app,ml}
PETNT181pp	2.758737	1	k _{app,ml}
ARMEPNS	2.604305	1	k _{app,ml}
LCARSyi	16.67119	1	k _{app,ml}
MOCD5	1.491555	1	k _{app,ml}
RPNTPH	6.806072	1	k _{app,ml}
PRCPD	8.589033	1	k _{app,ml}
UDPGPT	2.703733	1	k _{app,ml}
UDPGPT	2.703733	-1	k _{app,ml}
OMMBLXy	2.28037	1	k _{app,ml}
OPHHXy	2.585025	1	k _{app,ml}
PROD3	2.562292	1	k _{app,ml}
F1PP	1.483336	1	k _{app,ml}
ACP1p	2.77876	1	k _{app,ml}
3HPADHi	2.464206	1	k _{app,ml}
2HPTCOAT	6.153794	1	k _{app,ml}
MC6PH	6.1778	1	k _{app,ml}
FORCT	3.793628	1	k _{app,ml}
FORCT	3.793628	-1	k _{app,ml}
PACCOAE	2.314845	1	k _{app,ml}
OXDHCOAT	2.665219	1	k _{app,ml}
HADPCOADH3	2.508011	1	k _{app,ml}
HADPCOADH3	2.915597	-1	k _{app,ml}
DM_mththf_c	1.508031	1	k _{app,ml}
PYROX	2.79138	1	k _{app,ml}
BMOGDS1	3.699292	1	k _{app,ml}
MPTS	1.490435	1	k _{app,ml}
BMOGDS2	3.699292	1	k _{app,ml}
OCTNLL	1.483907	1	k _{app,ml}
XPPT	131	1	k _{cat}
HXPRT	7.89946	1	k _{cat}
SHK3Dr	31.35933	1	k _{cat}
SHK3Dr	11445.67	-1	k _{cat}
DHORTS	40.62444	1	k _{cat}
DURIPP	145	-1	k _{cat}
ACS	20.26667	1	k _{cat}
MAN6PI	64.4952	1	k _{cat}
PPA	108.0008	1	k _{cat}
TRPAS2	42	1	k _{cat}
PPCK	65	1	k _{cat}
ME1	134.4	1	k _{cat}
ALAR	27.6	1	k _{cat}
ALAR	6	-1	k _{cat}
ALATA_L	17.1	1	k _{cat}
XYLK	3427.5	1	k _{cat}
RBK	10.5	1	k _{cat}
PPM	173	-1	k _{cat}
ASPTA	82.47143	1	k _{cat}
FBP	19.1	1	k _{cat}
GLGC	370	1	k _{cat}
PYK	135.6429	1	k _{cat}
ASPISO	172	1	k _{cat}
ASPISO	130	-1	k _{cat}
GTHOr	441.95	1	k _{cat}
ILETA	48	1	k _{cat}
DHORD5	49.64	1	k _{cat}
VALTA	19	1	k _{cat}
ORPT	1.95	1	k _{cat}
TRPS2	4.45194	1	k _{cat}
PSCVT	1.94667	-1	k _{cat}
PFL	12.8	1	k _{cat}
ANS	64.03322	1	k _{cat}
FRD2	79.6	1	k _{cat}
FRD3	79.6	1	k _{cat}
POX	167	1	k _{cat}
CHORM	147457.6	1	k _{cat}
PTAr	1341.326	1	k _{cat}
PTAr	1419.466	-1	k _{cat}
ACKr	632.6667	1	k _{cat}

ACKr	942.75	-1	k _{cat}
LEUTAi	290	1	k _{cat}
TMDPP	984	1	k _{cat}
FBA	10.85	1	k _{cat}
HCO3E	17000	1	k _{cat}
PP5	1160.137	1	k _{cat}
PGI	1340.56	-1	k _{cat}
PGK	529.8	1	k _{cat}
EDA	72.15063	1	k _{cat}
NTPTP1	4000	1	k _{cat}
RPE	1180.013	1	k _{cat}
RPE	1300	-1	k _{cat}
TALA	5.67	-1	k _{cat}
PUNP1	38.785	1	k _{cat}
PUNP1	110	-1	k _{cat}
PUNP2	38.785	1	k _{cat}
PUNP2	110	-1	k _{cat}
PUNP3	15	1	k _{cat}
PUNP3	100.1667	-1	k _{cat}
TKT1	21.92635	1	k _{cat}
PUNP5	50.945	1	k _{cat}
PUNP5	111.6667	-1	k _{cat}
PUNP6	50.945	1	k _{cat}
PUNP6	111.6667	-1	k _{cat}
PUNP7	50.61333	1	k _{cat}
PUNP7	110	-1	k _{cat}
G6PDA	1800	1	k _{cat}
ALCD2x	24.37023	1	k _{cat}
ALCD2x	11.08	-1	k _{cat}
RNDR3	8.115	1	k _{cat}
ALDD2y	0.04	1	k _{cat}
RMI	68	1	k _{cat}
RMI	1717	-1	k _{cat}
TMDK1	114.104	1	k _{cat}
NADH10	0.077	1	k _{cat}
MDH	1035	1	k _{cat}
MDH	525	-1	k _{cat}
ABTA	3.34559	1	k _{cat}
FUM	290.0556	1	k _{cat}
FUM	40.10714	-1	k _{cat}
PYDXNK	0.3	1	k _{cat}
PYAM5PO	14.86	1	k _{cat}
SSALx	31.7	1	k _{cat}
ASPT	180	1	k _{cat}
SERD_L	490	1	k _{cat}
THRD	135	1	k _{cat}
NMNAT	0.00089	1	k _{cat}
KARA1	0.00417	1	k _{cat}
HMPK1	0.00012	1	k _{cat}
ACCOAC	65.25	1	k _{cat}
GLXCL	18.9	1	k _{cat}
GLCS1	284	1	k _{cat}
GLCP	50.5	1	k _{cat}
RBK_L1	40.42	1	k _{cat}
RBP4E	9.27384	1	k _{cat}
NTD2	368.25	1	k _{cat}
GLCRAL	27.3	1	k _{cat}
NTD4	368.25	1	k _{cat}
NTD5	294.632	1	k _{cat}
KAS14	3	1	k _{cat}
AGMT	3	1	k _{cat}
NTD6	294.66	1	k _{cat}
PYDXPP	10.25	1	k _{cat}
ARGDC	5.02	1	k _{cat}
ECOAH1	1280	-1	k _{cat}
G1SAT	0.59	-1	k _{cat}
UGLT	340.75	1	k _{cat}
UGLT	283	-1	k _{cat}
OAADC	2502.333	1	k _{cat}
MNNH	5.88	1	k _{cat}
DDGLK	26.68667	1	k _{cat}
XYLI2	16.91052	1	k _{cat}
XYLI2	11.085	-1	k _{cat}
PMANM	4.15	1	k _{cat}
GLUR	43.33	1	k _{cat}
UAG2E	3.625	1	k _{cat}

HCYSMT	0.0375	1	k _{cat}
MACPD	3	1	k _{cat}
UDPG4E	18	1	k _{cat}
UDPG4E	750	-1	k _{cat}
LACZ	247.5	1	k _{cat}
XYLI1	68.93588	1	k _{cat}
XYLI1	7.7	-1	k _{cat}
PPC	540	1	k _{cat}
ALDD4	26.95	1	k _{cat}
G3PT	160	1	k _{cat}
PYDXK	2.15	1	k _{cat}
HEX7	31.385	1	k _{cat}
URIC	24.7	1	k _{cat}
RMPA	38.3	1	k _{cat}
TGBPA	2.1	1	k _{cat}
DMSOR1	79.9	1	k _{cat}
PGLYCP	94	1	k _{cat}
G1PTT	5.925	1	k _{cat}
TDPGDH	3.4	1	k _{cat}
ADCS	0.53	1	k _{cat}
UDPGALM	1.5	1	k _{cat}
UMPK	107.399	1	k _{cat}
DDPGALA	0.94	1	k _{cat}
ALLTN	310.85	1	k _{cat}
UPPRT	3.85333	1	k _{cat}
NTD9	368.25	1	k _{cat}
NTD1	294.646	1	k _{cat}
NTD7	422.6667	1	k _{cat}
NDPK7	106	1	k _{cat}
TRPS3	2.09	1	k _{cat}
NTD10	368.25	1	k _{cat}
ADPT	9.33	1	k _{cat}
GTPDPK	2.4825	1	k _{cat}
ADK1	585.4	1	k _{cat}
ADK1	783	-1	k _{cat}
NDPK4	80.15	1	k _{cat}
ADA	18.5	1	k _{cat}
DADA	6.8	1	k _{cat}
ADD	5.5	1	k _{cat}
AGDC	95.875	1	k _{cat}
ASNN	800.0064	1	k _{cat}
ASPCT	417	1	k _{cat}
MCOATA	1580	1	k _{cat}
DAPE	75.5	-1	k _{cat}
GSPMDA	2.1	1	k _{cat}
GSPMDS	4.2	1	k _{cat}
GUAD	649	1	k _{cat}
TMAOR1	114	1	k _{cat}
DGK1	70.5	1	k _{cat}
NDPK1	38	1	k _{cat}
TRDR	16.955	1	k _{cat}
NNAM	0.93	1	k _{cat}
P5CD	19.15	1	k _{cat}
PGMT	92.5	1	k _{cat}
SPMDAT1	2.958	1	k _{cat}
KG6PDC	51	1	k _{cat}
SPMDAT2	2.958	1	k _{cat}
TRE6PP	14.3	1	k _{cat}
BETALDH _y	276	1	k _{cat}
ADPRDP	117	1	k _{cat}
ACPS1	0.7	1	k _{cat}
ARAI	39.36357	1	k _{cat}
M1PD	8.5	1	k _{cat}
M1PD	125	-1	k _{cat}
G3PD2	291	1	k _{cat}
GALS3	118.25	1	k _{cat}
AMPN	61	1	k _{cat}
ACCOAL	79.57143	1	k _{cat}
CYANST	245	1	k _{cat}
GMAND	5	1	k _{cat}
CYTD	4.8	1	k _{cat}
DCTPD	1.24	1	k _{cat}
UAGAAT	9.16667	1	k _{cat}
CSND	125.4667	1	k _{cat}
GLUN	1270	1	k _{cat}
SERAT	115.217	-1	k _{cat}

CYSS	24	1	k _{cat}
PAPSR	3.5	1	k _{cat}
ASPK	29.26818	1	k _{cat}
ASAD	199.932	1	k _{cat}
OCBT	2300.433	1	k _{cat}
OCBT	90	-1	k _{cat}
GPDDA1	3.2	1	k _{cat}
GPDDA2	1.1	1	k _{cat}
ACBIPGT	0.058	1	k _{cat}
TRPS1	3.85906	1	k _{cat}
HPPPND0	18.11	1	k _{cat}
GLYCK2	350	1	k _{cat}
ACOTA	1.55	1	k _{cat}
CYTDH	11.6	1	k _{cat}
ETHAAL	30	1	k _{cat}
AP4AH	250	1	k _{cat}
ADNCYC	1.67	1	k _{cat}
MI1PP	3.79667	1	k _{cat}
PPAKr	657.4	1	k _{cat}
PPAKr	618.5	-1	k _{cat}
TAUDO	2.34	1	k _{cat}
NTP3	422.9569	1	k _{cat}
PPM2	173	1	k _{cat}
ME2	66.6	1	k _{cat}
CBPS	3.15755	1	k _{cat}
ACOATA	0.30284	1	k _{cat}
ACOATA	0.30284	-1	k _{cat}
MLTG1	0.24	1	k _{cat}
AMALT3	94.275	1	k _{cat}
ACNML	7.7	1	k _{cat}
FTHFLi	3.6	1	k _{cat}
UDPGD	1.95425	1	k _{cat}
GLCRD	9.66583	1	k _{cat}
TDPAGTA	5.66	1	k _{cat}
GLUDC	4.74727	1	k _{cat}
HSDy	8.367	1	k _{cat}
CYSTL	1.5	1	k _{cat}
AMALT1	112.825	1	k _{cat}
MLTP2	4.1	1	k _{cat}
MLTP2	52	-1	k _{cat}
PACCOAL	30.85	1	k _{cat}
ACGAMT	1.2	1	k _{cat}
H2Otex	112	-1	k _{cat}
FACOAL160t2	0.029	1	k _{cat}
FACOAL180t2	0.029	1	k _{cat}
NO3R1bpp	392.5	1	k _{cat}
NO3R2pp	2.5	1	k _{cat}
NTP3pp	422.9569	1	k _{cat}
NO3R2bpp	392.5	1	k _{cat}
TMAOR1pp	258	1	k _{cat}
TRE6PS	5.2	1	k _{cat}
DRPA	294.5	1	k _{cat}
TKT2	20.43874	1	k _{cat}
UDCPDP	9	1	k _{cat}
TREH	199	1	k _{cat}
ATPS4rpp	270	1	k _{cat}
ATPS4rpp	217	-1	k _{cat}
CYTK2	41.66667	1	k _{cat}
O2tex	491	1	k _{cat}
NTD8pp	246.8733	1	k _{cat}
CSnt2pp	165	1	k _{cat}
TREHpp	199	1	k _{cat}
ACGAMK	0.03167	1	k _{cat}
METS	0.12	1	k _{cat}
ASPO5	5.55	1	k _{cat}
ASPO4	2.6	1	k _{cat}
FDMO4	0.86	1	k _{cat}
L_LACD3	178.7143	1	k _{cat}
L_LACD2	178.7143	1	k _{cat}
CYANSTpp	245	1	k _{cat}
MCPST	12.5	1	k _{cat}
UDCPDPpp	9	1	k _{cat}
NODOy	670	1	k _{cat}
GALM2pp	18400	1	k _{cat}
NODOx	670	1	k _{cat}
NHFRBO	14.9	1	k _{cat}

42A12BOOXpp	33.1032	1	k _{cat}
TYROXDApp	28.89711	1	k _{cat}
RNDR3b	2.5	1	k _{cat}
FEROpp	0.7125	1	k _{cat}
LACZpp	33.8	1	k _{cat}
GPDDA1pp	3.2	1	k _{cat}
GPDDA5pp	1.8	1	k _{cat}
FLVR	47.87692	1	k _{cat}
PSP_Lpp	26.8	1	k _{cat}
TYRpp	62.4	1	k _{cat}
PAPSR2	3.5	1	k _{cat}
FACOAL80t2p	0.029	1	k _{cat}
PPGPPDP	3	1	k _{cat}
GUACYC	28.7	1	k _{cat}
23PDE9pp	1.97	1	k _{cat}
SARCOX	37.43333	1	k _{cat}
FACOAL60t2p	0.029	1	k _{cat}
ASNpp	800.0064	1	k _{cat}
GLUNpp	1270	1	k _{cat}
NTD12	294.648	1	k _{cat}
NTD10pp	368.25	1	k _{cat}
NTD1pp	294.646	1	k _{cat}
NTD2pp	368.25	1	k _{cat}
NTD3pp	368.25	1	k _{cat}
NTD4pp	368.25	1	k _{cat}
NTD5pp	294.632	1	k _{cat}
NTD6pp	294.66	1	k _{cat}
NTD7pp	422.6667	1	k _{cat}
NTD9pp	368.25	1	k _{cat}
NH4tex	236	1	k _{cat}
PEAMNOpp	14.92857	1	k _{cat}
GAL1PPpp	0.0348	1	k _{cat}
NTPP9	0.16	1	k _{cat}
NTPP10	0.16	1	k _{cat}
ARGDCpp	5.02	1	k _{cat}
URIH	9.3	1	k _{cat}
INSH	0.043	1	k _{cat}
FACOAL100t2	0.029	1	k _{cat}
GPDDA3pp	2.8	1	k _{cat}
GPDDA4pp	3.3	1	k _{cat}
GPDDA2pp	1.1	1	k _{cat}
FACOAL120t2	0.029	1	k _{cat}
FACOAL140t2	0.029	1	k _{cat}
HCYSMT2	0.38	1	k _{cat}
FALDH2	51.7	1	k _{cat}
CO2tex	380	1	k _{cat}
CBIAT	0.0775	1	k _{cat}
NTD12pp	294.648	1	k _{cat}
3NTD2pp	0.23	1	k _{cat}
3NTD9pp	8.04	1	k _{cat}
GLUDy	2136	1	k _{cat}
GLUDy	19466.67	-1	k _{cat}
23PDE7pp	7.63	1	k _{cat}
GLYCLTDy	7.48333	1	k _{cat}
LYSDC	405.5	1	k _{cat}
NTD3	368.25	1	k _{cat}
GTPDPDP	0.023	1	k _{cat}
DHFR	3.1513	1	k _{cat}
TYRTA	31.075	1	k _{cat}
TYRTA	64.16667	-1	k _{cat}
SHKK	35	1	k _{cat}
GLUSy	24.55	1	k _{cat}
GUAPRT	12.4865	1	k _{cat}
ABUTD	7.7	1	k _{cat}
E4PD	20	1	k _{cat}
DMSOR2pp	79.9	1	k _{cat}
DUTPDP	22.6625	1	k _{cat}
NO3R1pp	392.5	1	k _{cat}
NTP5	317.8481	1	k _{cat}
DPR	25	1	k _{cat}
NTD8	246.8733	1	k _{cat}
GALCTD	22	1	k _{cat}
PUNP4	50.61333	1	k _{cat}
PUNP4	100.1667	-1	k _{cat}
G6PP	700	1	k _{cat}
GLNS	6	1	k _{cat}

ACACT1r	15.8	1	k _{cat}
ACACT1r	159.5909	-1	k _{cat}
NTD11pp	368.25	1	k _{cat}
PDE1	63.4	1	k _{cat}
AMALT2	123.625	1	k _{cat}
MLTP1	5.4	1	k _{cat}
MLTP1	52	-1	k _{cat}
ALAALAR	15.13	1	k _{cat}
ADNK1	8.09733	1	k _{cat}
UDPGPpp	71	1	k _{cat}
PPPGO	3.35443	1	k _{cat}
LDH_D2	875	1	k _{cat}
SUCOAS	25.66667	1	k _{cat}
SUCOAS	21	-1	k _{cat}
CYSDDS	72	1	k _{cat}
NTP1	430.5998	1	k _{cat}
GPDDA4	3.3	1	k _{cat}
GALKr	15.66727	1	k _{cat}
G1PPpp	117	1	k _{cat}
LDH_D	320.5	-1	k _{cat}
AMALT4	84.05	1	k _{cat}
NTD11	368.25	1	k _{cat}
IPDDI	19.41364	1	k _{cat}
ORNDC	17.22986	1	k _{cat}
UPP3MT	38	1	k _{cat}
FACOAL181t2	0.029	1	k _{cat}
EAR141y	130.2	1	k _{cat}
CFAS160G	0.39765	1	k _{cat}
FACOAL141t2	0.029	1	k _{cat}
FACOAL161t2	0.029	1	k _{cat}
SERD_D	13.1	1	k _{cat}
FLVRx	28.85	1	k _{cat}
FMNRx	6	1	k _{cat}
FORtex	272	-1	k _{cat}
PFK	83.6016	1	k _{cat}
PDXSPOi	5.185	1	k _{cat}
PGM	255.2021	-1	k _{cat}
CYTBO3_4pp	814.4	1	k _{cat}
GLYCK	350	1	k _{cat}
GPDDA3	2.8	1	k _{cat}
TPI	9000	-1	k _{cat}
EAR181y	130.2	1	k _{cat}
DASYN140	8.92	1	k _{cat}
DASYN180	8.92	1	k _{cat}
DASYN141	8.92	1	k _{cat}
RPI	2100	1	k _{cat}
RPI	50	-1	k _{cat}
SFGTHi	17.505	1	k _{cat}
PYDAMK	0.55	1	k _{cat}
DHORDfum	7.8	1	k _{cat}
FACOAE120	50	1	k _{cat}
DSBCGT	25.2	1	k _{cat}
EAR180y	130.2	1	k _{cat}
ACONTb	44.8	1	k _{cat}
GLCP2	50.5	1	k _{cat}
SUCDi	167	1	k _{cat}
LALDO2x	266	1	k _{cat}
ASNS1	1.89833	1	k _{cat}
EAR80y	130.2	1	k _{cat}
EAR120y	130.2	1	k _{cat}
EAR121y	130.2	1	k _{cat}
EAR40y	130.2	1	k _{cat}
EAR161y	130.2	1	k _{cat}
3OAS140	3	1	k _{cat}
EAR160y	130.2	1	k _{cat}
PDE4	0.00096	1	k _{cat}
EAR100y	130.2	1	k _{cat}
ACONTa	5.3	1	k _{cat}
ACONTa	44.8	-1	k _{cat}
CFAS160E	0.39765	1	k _{cat}
CFAS180E	0.39765	1	k _{cat}
CFAS180G	0.39765	1	k _{cat}
DAGK140	2.8	1	k _{cat}
DAGK160	1.7	1	k _{cat}
3OAS60	3	1	k _{cat}
3OAS160	3	1	k _{cat}

EAR140y	130.2	1	k _{cat}
3OAS80	3	1	k _{cat}
3OAS100	3	1	k _{cat}
3OAS120	3	1	k _{cat}
LSERDHR	4.2	1	k _{cat}
DASYN120	8.92	1	k _{cat}
EAR60y	130.2	1	k _{cat}
DAGK181	2.8	1	k _{cat}
NTP11	0.22	1	k _{cat}
3OAR180	71	1	k _{cat}
3OAS180	3	1	k _{cat}
ACOAD6f	6	1	k _{cat}
NADH16pp	1517	1	k _{cat}
NADH18pp	1517	1	k _{cat}
NADH17pp	1517	1	k _{cat}
DTARTD	7.3	1	k _{cat}
OXCDC	88	1	k _{cat}
CTECOAI8	47	1	k _{cat}
CTECOAI8	12	-1	k _{cat}
CTECOAI6	63	1	k _{cat}
SELR	387	1	k _{cat}
FLDR2	273.5	1	k _{cat}
THFAT	18.4	1	k _{cat}
ALDD19xr	96.8	1	k _{cat}
MMM	48	1	k _{cat}
GTHPi	221.7	1	k _{cat}
ACOAD1f	9.6	1	k _{cat}
26DAHtex	65	1	transporter
26DAHtex	65	-1	transporter
HCINNMtex	65	1	transporter
HCINNMtex	65	-1	transporter
HPPPntex	65	1	transporter
HPPPntex	65	-1	transporter
ACTex	65	1	transporter
ACTex	65	-1	transporter
ACALDtex	65	1	transporter
ACALDtex	65	-1	transporter
ACMANAtex	65	1	transporter
ACMANAtex	65	-1	transporter
ACMUJMtex	65	1	transporter
SUCct2_2pp	65	1	transporter
SUCFUMtpp	65	1	transporter
SUCFUMtpp	65	-1	transporter
SUCptspp	65	1	transporter
SULabcpp	65	1	transporter
TSULabcpp	65	1	transporter
TARTRt7pp	65	1	transporter
TARTRt7pp	65	-1	transporter
TAURabcpp	65	1	transporter
THMDt2pp_cc	65	1	transporter
NMNTt7pp	65	1	transporter
GLUtex	65	1	transporter
GLUtex	65	-1	transporter
Kt2pp	65	1	transporter
GLYtex	65	1	transporter
GLYtex	65	-1	transporter
GLYALDtex	65	1	transporter
GLYALDtex	65	-1	transporter
AKGt2rpp	65	1	transporter
AKGt2rpp	65	-1	transporter
XYLt2pp	65	1	transporter
GLYBtex	65	1	transporter
GLYBtex	65	-1	transporter
GLYCtex	65	1	transporter
GLYCtex	65	-1	transporter
GLYC3Ptex	65	1	transporter
GLYC3Ptex	65	-1	transporter
GLYCLTtex	65	1	transporter
GLYCLTtex	65	-1	transporter
GTPtex	65	1	transporter
GTPtex	65	-1	transporter
GUAAtex	65	1	transporter
GUAAtex	65	-1	transporter
Htex	65	1	transporter
Htex	65	-1	transporter
H2tex	65	1	transporter
H2tex	65	-1	transporter
H2Otex	65	1	transporter
O2tpp	65	1	transporter
O2tpp	65	-1	transporter
PYRt2rpp	65	1	transporter
PYRt2rpp	65	-1	transporter
THMabcpp	65	1	transporter
THMDt2pp_cc	65	1	transporter
THMDt2pp_cc	65	-1	transporter
THRabcpp	65	1	transporter

TREptspp	65	1	transporter
XANtpp	65	1	transporter
XANtpp	65	-1	transporter
DAPtex	65	1	transporter
DAPtex	65	-1	transporter
TRPt2rpp	65	1	transporter
TRPt2rpp	65	-1	transporter
TYRt2rpp	65	1	transporter
TYRt2rpp	65	-1	transporter
URIt2pp_copy	65	1	transporter
URIt2pp_copy	65	1	transporter
URIt2pp_copy	65	-1	transporter
VAlabcpp	65	1	transporter
12PPDStex	65	1	transporter
12PPDStex	65	-1	transporter
DDGLCNtex	65	1	transporter
DDGLCNtex	65	-1	transporter
XTSnt2rpp	65	1	transporter
XTSnt2rpp	65	-1	transporter
XYLabcpp	65	1	transporter
XYLut2pp	65	1	transporter
THRt4pp	65	1	transporter
GALURtex	65	1	transporter
GALURtex	65	-1	transporter
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FALDtex	65	-1	transporter
CU2tex	65	1	transporter
CU2tex	65	-1	transporter
FECRMUtex	65	1	transporter
CLtex	65	1	transporter

CLtex	65	-1	transporter
CYSDtex	65	1	transporter
CYSDtex	65	-1	transporter
CYSDabcpp	65	1	transporter
GMPtex	65	1	transporter
GMPtex	65	-1	transporter
IMPtex	65	1	transporter
IMPtex	65	-1	transporter
G3Ptex	65	1	transporter
G3Ptex	65	-1	transporter
CD2abcpp	65	1	transporter
G3PGtex	65	1	transporter
G3PGtex	65	-1	transporter
G3PEtex	65	1	transporter
G3PEtex	65	-1	transporter
G3PCtex	65	1	transporter
G3PCtex	65	-1	transporter
UMPtex	65	1	transporter
UMPtex	65	-1	transporter
G1Ptex	65	1	transporter
G1Ptex	65	-1	transporter
CUt3	65	1	transporter
ZN2t3pp	65	1	transporter
SKMtex	65	1	transporter
SKMtex	65	-1	transporter
MNt2pp	65	1	transporter
GLCabcpp	65	1	transporter
MNtex	65	1	transporter
MNtex	65	-1	transporter
GAL1Ptex	65	1	transporter
GAL1Ptex	65	-1	transporter
O2Stex	65	1	transporter
O2Stex	65	-1	transporter
Kt3pp	65	1	transporter
HOMt2pp	65	1	transporter
HOMtex	65	1	transporter
HOMtex	65	-1	transporter
SO2tex	65	1	transporter
SO2tex	65	-1	transporter
SO2tpp	65	1	transporter
SO2tpp	65	-1	transporter
CYStpp	65	1	transporter
DAMPtex	65	1	transporter
DAMPtex	65	-1	transporter
DCMPtex	65	1	transporter
DCMPtex	65	-1	transporter
DGMPtex	65	1	transporter
DGMPtex	65	-1	transporter
DTMPtex	65	1	transporter
DTMPtex	65	-1	transporter
ACSERTpp	65	1	transporter
ACSERTex	65	1	transporter
ACSERTex	65	-1	transporter
2AGPE120tipp	65	1	transporter
G3PGabcpp	65	1	transporter
G3Plabcpp	65	1	transporter
G3PSabcpp	65	1	transporter
2AGPA120tipp	65	1	transporter
2AGPG180tipp	65	1	transporter
ALAALAtex	65	1	transporter
ALAALAtex	65	-1	transporter
2AGPG141tipp	65	1	transporter
2AGPG161tipp	65	1	transporter
2AGPG181tipp	65	1	transporter
PGP120abcpp	65	1	transporter
MALtex	65	1	transporter
MALtex	65	-1	transporter
NO3tex	65	1	transporter
NO3tex	65	-1	transporter
PGP161abcpp	65	1	transporter
PGP181abcpp	65	1	transporter
PA120abcpp	65	1	transporter
URAt2pp	65	1	transporter
TTDCEAtexi	65	1	transporter
HDCEAtexi	65	1	transporter
PA160abcpp	65	1	transporter
ALAALAabcpp	65	1	transporter
HEXt2rpp	65	1	transporter
HEXt2rpp	65	-1	transporter
G3PEabcpp	65	1	transporter
G3PCabcpp	65	1	transporter
PA181abcpp	65	1	transporter
12DGR141tipp	65	1	transporter
PROGLYabcpp	65	1	transporter
GTHRDabcpp	65	1	transporter
2AGPG140tipp	65	1	transporter
2AGPG160tipp	65	1	transporter
D_LACT2pp	65	1	transporter
D_LACT2pp	65	-1	transporter
RIBtex	65	1	transporter
RIBtex	65	-1	transporter

THMtex	65	1	transporter
THMtex	65	-1	transporter
XYLUtex	65	1	transporter
XYLUtex	65	-1	transporter
ALLtex	65	1	transporter
ALLtex	65	-1	transporter
FRUpts2pp	65	1	transporter
PGP140abcpp	65	1	transporter
34dhpactex	65	1	transporter
34dhpactex	65	-1	transporter
G6Pt6_2pp	65	1	transporter
MALTHXtexi	65	1	transporter
12PPDStpp	65	1	transporter
12PPDStpp	65	-1	transporter
PROtex	65	1	transporter
PROtex	65	-1	transporter
FE3HOXUtex	65	1	transporter
FE3DHBZStone	65	1	transporter
H2tpp	65	1	transporter
H2tpp	65	-1	transporter
23DAPPAtex	65	1	transporter
23DAPPAtex	65	-1	transporter
FEOXAMtonex	65	1	transporter
GAMptspp	65	1	transporter
GSNT2pp	65	1	transporter
GSNtex	65	1	transporter
GSNtex	65	-1	transporter
SBTtex	65	1	transporter
SBTtex	65	-1	transporter
FORtex	65	1	transporter
PHET2rpp	65	1	transporter
PHET2rpp	65	-1	transporter
HG2t3pp	65	1	transporter
MELIBt3ipp	65	1	transporter
GLCtex_copy2	65	1	transporter
GLCtex_copy2	65	-1	transporter
NAt3pp	65	1	transporter
MINOHPtexi	65	1	transporter
12PPDRtpp	65	1	transporter
12PPDRtpp	65	-1	transporter
K2L4Aabcpp	65	1	transporter
AKGtex	65	1	transporter
AKGtex	65	-1	transporter
ALLNTtex	65	1	transporter
ALLNTtex	65	-1	transporter
ASNtex	65	1	transporter
ASNtex	65	-1	transporter
F6Pt6_2pp	65	1	transporter
PPALtpp	65	1	transporter
PPALtpp	65	-1	transporter
GAM6Pt6_2pp	65	1	transporter
FE3HOXabcpp	65	1	transporter
PPALtex	65	1	transporter
PPALtex	65	-1	transporter
CA2tex	65	1	transporter
CA2tex	65	-1	transporter
CUtex	65	1	transporter
CUtex	65	-1	transporter
12PPDRtex	65	1	transporter
12PPDRtex	65	-1	transporter
PROGLYtex	65	1	transporter
PROGLYtex	65	-1	transporter
F6Ptex	65	1	transporter
F6Ptex	65	-1	transporter
GAMAN6Ptex	65	1	transporter
GAMAN6Ptex	65	-1	transporter
PE141abcpp	65	1	transporter
PG140abcpp	65	1	transporter
PG141abcpp	65	1	transporter
12DGR140tipp	65	1	transporter
PG180abcpp	65	1	transporter
12DGR160tipp	65	1	transporter
PGP141abcpp	65	1	transporter
PGP160abcpp	65	1	transporter
12DGR180tipp	65	1	transporter
PGP180abcpp	65	1	transporter
PA140abcpp	65	1	transporter
12DGR120tipp	65	1	transporter
PG120abcpp	65	1	transporter
PE140abcpp	65	1	transporter
2AGPE140tipp	65	1	transporter
2AGPE160tipp	65	1	transporter
2AGPE180tipp	65	1	transporter
2AGPE141tipp	65	1	transporter
2AGPE161tipp	65	1	transporter
2AGPA161tipp	65	1	transporter
2AGPA181tipp	65	1	transporter
2AGPG120tipp	65	1	transporter
PA141abcpp	65	1	transporter
PA161abcpp	65	1	transporter
PA180abcpp	65	1	transporter

PE120abcpp	65	1	transporter
PE180abcpp	65	1	transporter
CGLYtex	65	1	transporter
CGLYtex	65	-1	transporter
GTHOxtex	65	1	transporter
GTHOxtex	65	-1	transporter
TDSR1	65	1	transporter
CGLYabcpp	65	1	transporter
2AGPE181tipp	65	1	transporter
2AGPA140tipp	65	1	transporter
2AGPA160tipp	65	1	transporter
2AGPA180tipp	65	1	transporter
2AGPA141tipp	65	1	transporter
12DGR161tipp	65	1	transporter
12DGR181tipp	65	1	transporter
TDSR2	65	1	transporter
GTHRDtex	65	1	transporter
GTHRDtex	65	-1	transporter
GTHRDabc2pp	65	1	transporter
NI2tpp	65	1	transporter
COBAL2t3pp	65	1	transporter
23DAPPAt2pp	65	1	transporter
CA16pp	65	1	transporter
GLYb2pp	65	1	transporter
5DGLCNt2rpp	65	1	transporter
5DGLCNt2rpp	65	-1	transporter
GALCNT2pp	65	1	transporter
GALCNTltex	65	1	transporter
GALCNTltex	65	-1	transporter
DSERT2pp	65	1	transporter
ALAt4pp	65	1	transporter
LYSt3pp	65	1	transporter
CA2t3pp	65	1	transporter
THRT2pp	65	1	transporter
ARGt3pp	65	1	transporter
MALDtex	65	1	transporter
MALDtex	65	-1	transporter
OROTtex	65	1	transporter
OROTtex	65	-1	transporter
PPAt4pp	65	1	transporter
OROT12_2pp	65	1	transporter
MALDt2_2pp	65	1	transporter
5DGLCNtex	65	1	transporter
5DGLCNtex	65	-1	transporter
CHL2pp	65	1	transporter
GLYt4pp	65	1	transporter
FRULYSt2pp	65	1	transporter
TUNGStex	65	1	transporter
TUNGStex	65	-1	transporter
PSCLYSt2pp	65	1	transporter
LYSt2pp	65	1	transporter
COBAL2tpp	65	1	transporter
GLYt2pp	65	1	transporter
ALAt2pp	65	1	transporter
BALAt2pp	65	1	transporter
MG2tpp	65	1	transporter
SMTRtex	65	1	transporter
SMTRtex	65	-1	transporter
NI2t3pp	65	1	transporter
PPAtex	65	1	transporter
PPAtex	65	-1	transporter
GALCNTlt2pp	65	1	transporter
THYMt3pp	65	1	transporter
CLT3_2pp	65	1	transporter
PSCLYStex	65	1	transporter
PSCLYStex	65	-1	transporter
THYMtex	65	1	transporter
THYMtex	65	-1	transporter
FE2t3pp	65	1	transporter
MN2t3pp	65	1	transporter
MG2t3_2pp	65	1	transporter
MG2t3_2pp	65	-1	transporter
CD2tpp	65	1	transporter
ETHAt2pp	65	1	transporter
FE3abcpp	65	1	transporter
LALADGLUtex	65	1	transporter
LALADGLUtex	65	-1	transporter
LALALGLUtp	65	1	transporter
3HPPtex	65	1	transporter
3HPPtex	65	-1	transporter
CMtpp	65	1	transporter
FUSAtpp	65	1	transporter
RFAMPtex	65	1	transporter
RFAMPtex	65	-1	transporter
MINCYCtex	65	1	transporter
MINCYCtex	65	-1	transporter
MINCYCtpp	65	1	transporter
MEOHtrpp	65	1	transporter
MEOHtrpp	65	-1	transporter
DOXRBCNtex	65	1	transporter
DOXRBCNtex	65	-1	transporter

DOXRBCNtpp	65	1	transporter
SELTex	65	1	transporter
SELTex	65	-1	transporter
ETOHtrpp	65	1	transporter
ETOHtrpp	65	-1	transporter
MEOHtex	65	1	transporter
MEOHtex	65	-1	transporter
5MTRtpp	65	1	transporter
TTRCYCtex	65	1	transporter
TTRCYCtex	65	-1	transporter
NOVBCNtpp	65	1	transporter
CMtex	65	1	transporter
CMtex	65	-1	transporter
TTRCYCtpp	65	1	transporter
LALALGLUtex	65	1	transporter
LALALGLUtex	65	-1	transporter
LALADGLUtp	65	1	transporter
SLNttx	65	1	transporter
SLNttx	65	-1	transporter
3HPPtpp	65	1	transporter
RFAMPtpp	65	1	transporter
ALAtpp	65	1	transporter
CH4tex	65	1	transporter
CH4tex	65	-1	transporter
TARTRDtex	65	1	transporter
TARTRDtex	65	-1	transporter
BTNt2ipp	65	1	transporter
BTNttx	65	1	transporter
BTNttx	65	-1	transporter
DHBSZ3FEabc	65	1	transporter
DXYLONtpp	65	1	transporter
DXYLONtpp	65	-1	transporter
2DGLCptspp	65	1	transporter
ARBTptspp	65	1	transporter
ARBTtex	65	1	transporter
DMSOtpp	65	1	transporter
DMSOtpp	65	-1	transporter
PYDXtex	65	1	transporter
PYDXtex	65	-1	transporter
PYDXtpp	65	1	transporter
SUCASPtpp	65	1	transporter
SUCASPtpp	65	-1	transporter
23DOGULNt4f	65	1	transporter
SUCMALtpp	65	1	transporter
SUCMALtpp	65	-1	transporter
PYDXNtex	65	1	transporter
PYDXNtex	65	-1	transporter
PYDXNtpp	65	1	transporter
DXYLONtex	65	1	transporter
DXYLONtex	65	-1	transporter
PYDAMtex	65	1	transporter
PYDAMtex	65	-1	transporter
PYDAMtpp	65	1	transporter
PSURitex	65	1	transporter
PSURitex	65	-1	transporter
CHLt3pp	65	1	transporter
TART2_3pp	65	1	transporter
SUCTARTtpp	65	1	transporter
SUCTARTtpp	65	-1	transporter
CHTBSptspp	65	1	transporter
GLYBt3pp	65	1	transporter
Al2tpp	65	1	transporter
Ftpp	65	1	transporter
CHTBStex	65	1	transporter
CHTBStex	65	-1	transporter
TRPtipp	65	1	transporter
TYRtipp	65	1	transporter
DHPStr	65	1	transporter
SQtr	65	1	transporter
DHPStex	65	1	transporter
DHPStex	65	-1	transporter
SPMDt3pp	65	1	transporter
FMNtex	65	1	transporter
FMNtex	65	-1	transporter
PHEtipp	65	1	transporter
MALT3pp	65	1	transporter
CITt3pp	65	1	transporter
QUIN2tpp	65	1	transporter
QUIN2tpp	65	-1	transporter
GLYC2Pabcpp	65	1	transporter
CRNDtex	65	1	transporter
CRNDtex	65	-1	transporter
CLBtex	65	1	transporter
CLBtex	65	-1	transporter
CHOLATEtpp	65	1	transporter
4ABZGLUtr	65	1	transporter
4ABZGLUtr	65	-1	transporter
SELabcpp	65	1	transporter
CYSItpp	65	1	transporter
FADt	65	1	transporter
PUACGAMtr	65	1	transporter

Pit2rpp	65	1	transporter
FADtex	65	1	transporter
FADtex	65	-1	transporter
AI2abcpp	65	1	transporter
4HBZt3pp	65	1	transporter
LEUt2rpp	65	1	transporter
LCTStpp	65	1	transporter
ILEt3pp	65	1	transporter
METt3pp	65	1	transporter
GSNt3pp	65	1	transporter
CLtipp	65	1	transporter
2DGLCtex	65	1	transporter
2DGLCtex	65	-1	transporter
Pitpp	65	1	transporter
LEUt3pp	65	1	transporter
INSt3pp	65	1	transporter
INDOLEt2rpp	65	1	transporter
SQtex	65	1	transporter
SQtex	65	-1	transporter
ILEt2rpp	65	1	transporter
GLYtpp	65	1	transporter
GLYtpp	65	-1	transporter
URATEtex	65	1	transporter
URATEtex	65	-1	transporter
CBL1tonex	65	1	transporter
CBItonex	65	1	transporter
PSURitpp	65	1	transporter
CH4t2pp	65	1	transporter
METGLCURt2p	65	1	transporter
METGLCURtex	65	1	transporter
METGLCURtex	65	-1	transporter
FMNt	65	1	transporter
AI2tex	65	1	transporter
AI2tex	65	-1	transporter
CUt2pp	65	1	transporter
Ftex	65	1	transporter
Ftex	65	-1	transporter
CYSitex	65	1	transporter
CYSitex	65	-1	transporter
SERtpp	65	1	transporter
THRt2rpp	65	1	transporter
SERt2rpp	65	1	transporter
URATEtpp	65	1	transporter
CS1tex	65	1	transporter
CS1tex	65	-1	transporter
ADOCBLtonex	65	1	transporter
URAtpp	65	1	transporter
ASPtpp	65	1	transporter
ASPtpp	65	-1	transporter
FUMt1pp	65	1	transporter
FUMt1pp	65	-1	transporter
SUCct1pp	65	1	transporter
SUCct1pp	65	-1	transporter
FUSAtex	65	1	transporter
FUSAtex	65	-1	transporter
MEPNtex	65	1	transporter
MEPNtex	65	-1	transporter
MEPNabcpp	65	1	transporter
NOVBCNtex	65	1	transporter
NOVBCNtex	65	-1	transporter
4ABZGLUtex	65	1	transporter
4ABZGLUtex	65	-1	transporter
QUIN2tex	65	1	transporter
QUIN2tex	65	-1	transporter
SLNtabcpp	65	1	transporter
VALt2rpp	65	1	transporter
VALt3pp	65	1	transporter
MN2tipp	65	1	transporter

Table S4. Proteome efficiency of pathways on all carbon (glycolytic and gluconeogenic) sources.

Pathway	Pathway expression (for <i>shared</i> proteins) ($n = 18$) ^a			<i>measured-only</i> fraction ^b (median across 18 conditions) ($f_{\text{measured-only}}$)	individual <i>shared</i> proteins; median across 18 conditions ^c		
	r_{pathway}^2 (1)	p (1)	GMFE _{pathway} (2)		$r_{\text{individual}}^2$ (4)	p (4)	n^d
Measures (1)-(4)							
Transporters	(-) 0.49 ^e	1.2×10^{-03}	3.306	0.997	0.1	0.688	4
Central metabolism	0.50	9.6×10^{-04}	2.394	0.306	0.15	3.2×10^{-03}	56
Glycolysis	0.75	4.0×10^{-06}	3.437	0.081	0.3	0.084	11
Pentose phosphate pathway	0.19	0.074	1.922	0.384	0.3	0.338	5
TCA cycle	0.35	0.01	5.795	0.096	0.38	0.033	12
Glyoxylate shunt	1	-	nan	1	1	1	2
Energy generation	(-) 0.004 ^e	0.801	1.571	0.052	0.13	0.058	28
Central metabolism others	0.68	2.5×10^{-05}	1.506	0.549	0.99	0.058	3
Biosynthesis	0.78	1.2×10^{-06}	1.769	0.265	0.44	1.7×10^{-30}	229
Amino acid	0.76	2.9×10^{-06}	1.386	0.31	0.45	8.6×10^{-11}	72
Nucleotide	0.58	2.6×10^{-04}	3.422	0.231	0.15	0.045	28
Cell envelope	0.38	6.3×10^{-03}	1.809	0.138	0.38	2.3×10^{-05}	40
Cofactor	0.84	1.2×10^{-07}	1.232	0.114	0.62	5.5×10^{-16}	70
Biosynthesis others	0.12	0.155	3.877	0.255	0.44	8.5×10^{-05}	29
Translation	0.87	1.2×10^{-03}	1.35	0	0.99	0.063	3
Others (other metabolic enzymes)	0.005	0.781	1.619	0.917	0.14	0.045	30
Total metabolism	0.50	9.6×10^{-04}	1.807	0.52	0.35	3.4×10^{-30}	308

^a Values reflect the local optimality of complete pathways across conditions. $n = 18$ indicates the number of all carbon (glycolytic and gluconeogenic) sources analyzed.

^b Mass fraction of *measured-only* (un-predicted but observed) proteins relative to all proteins in the pathway.

^c These columns reflect the local optimality compared across individual proteins within each pathway at a given growth condition; values are medians across the $n = 18$ growth conditions.

^d Number of proteins in each pathway or pathway set.

^e Negative correlation coefficient r_{pathway} .

Table S5: Data source for pathway annotation

Pathway	Source or notes
Biosynthesis	Ecocyc pathway ontology: "Biosynthesis" in the first hierarchy
Amino acid	Ecocyc pathway ontology: "Amino Acid Biosynthesis" in the secondary hierarchy
Nucleotide	Ecocyc pathway ontology: "Nucleoside and Nucleotide Biosynthesis" in the secondary hierarchy excluding the "Purine Nucleotide Salvage" and "Pyrimidine Nucleotide Salvage" in the fourth hierarchy
Cell envelope	Ecocyc pathway ontology: "Fatty Acid and Lipid Biosynthesis" and "Cell Structure Biosynthesis" in the secondary hierarchy
Cofactor	Ecocyc pathway ontology: "Cofactor, Carrier, and Vitamin Biosynthesis" in the secondary hierarchy
Biosynthesis_others	Genes in ecocyc pathway ontology: "Biosynthesis" in the first hierarchy but are not in the mentioned 4 pathways
Precursors and energy generation	Ecocyc pathway ontology: "Generation of Precursor Metabolites and Energy" in the first hierarchy
Glycolysis	Ecocyc pathway ontology: "Glycolysis" in the secondary hierarchy
Pentose phosphate pathway	Ecocyc pathway ontology: "Pentose Phosphate Pathways" in the secondary hierarchy
TCA cycle	Ecocyc pathway ontology: "TCA cycle" in the secondary hierarchy
Glyoxylate shunt	Gene aceA, aceB, and glcB (added manually)
Energy	Ecocyc pathway ontology: "Electron Transfer Chains" and "ATP biosynthesis" in the secondary hierarchy
Precursors_energy_others	Genes in ecocyc pathway ontology: "Generation of Precursor Metabolites and Energy" in the first hierarchy but are not in the mentioned 5 pathways
Transport	iML1515 annotation (genes that transport metabolites across both inner and outer membrane)
Others	Genes in iML1515 model but are not in all the mentioned pathways

