

COL0 CONTROL

	maximum	minimum
Mal_Store	0.808000	0.39600
Fum_Store	1.414000	0.99000
Starch_Store	5.858000	5.34600
Im_CO2	101.000000	101.00000
FumHA_c	1.414000	0.00000
Fd_DASH_NADPR_h	210.000000	210.00000
ATPase_h	90.000000	90.00000
Tr_TPT3	208.771000	0.00000
RBC_h	91.777200	81.52725
PGAM_c	-43.380000	-996.11700
Enol_c	-43.380000	-996.11700
PEPC2_c	-9.222800	-147.29050
MalDH1_c	249.292429	-73.04000
FumHA_c	1.414000	0.00000
PGAK_h	223.347000	196.65300
GAPDH2_h	95.782000	-109.34000
Tr_TPT2	208.771000	0.00000
GAPDH3_c	69.910000	0.00000
Tr_TPT1	949.771000	0.00000

COL0 COLD

	maximum	minimum
Mal_Store	2.626000	2.178000
Fum_Store	4.848000	4.356000
Starch_Store	10.504000	9.900000
Im_CO2	101.000000	101.000000
FumHA_c	4.848000	0.000000
Fd_DASH_NADPR_h	180.000000	140.000000
ATPase_h	100.000000	90.174000
Tr_TPT3	247.465333	0.000000
RBC_h	105.672384	87.235459
PGAM_c	-7.328485	-1000.000000
Enol_c	-7.328485	-1000.000000
PEPC2_c	-2.798539	-160.283333
MalDH1_c	424.964737	-75.566667
FumHA_c	4.848000	0.000000
PGAK_h	251.633333	176.373333
GAPDH2_h	148.079000	-178.964333
Tr_TPT2	247.465333	0.000000
GAPDH3_c	102.569000	0.000000
Tr_TPT1	994.426667	0.000000

FUM2 CONTROL

	maximum	minimum
Mal_Store	2.626	2.178
Fum_Store	0.000	0.000
Starch_Store	8.118	8.118
Im_CO2	100.156	100.156
FumHA_c	0.000	0.000
Fd_DASH_NADPR_h	210.000	210.000
ATPase_h	90.000	90.000
Tr_TPT3	180.902	0.000
RBC_h	87.294	87.294
PGAM_c	-48.274	-969.332
Enol_c	-48.274	-969.332
PEPC2_c	-12.862	-114.254
MalDH1_c	149.586	-48.274
FumHA_c	0.000	0.000
PGAK_h	210.000	210.000
GAPDH2_h	55.065	-55.271
Tr_TPT2	180.902	0.000
GAPDH3_c	52.938	0.000
Tr_TPT1	921.058	0.000

FUM2 COLD

	maximum	minimum
Mal_Store	5.252000	4.752000
Fum_Store	0.000000	0.000000
Starch_Store	12.665333	12.276000
Im_CO2	95.000000	95.000000
FumHA_c	0.000000	0.000000
Fd_DASH_NADPR_h	180.000000	170.431333
ATPase_h	100.000000	92.977333
Tr_TPT3	207.579333	0.000000
RBC_h	94.265429	89.080000
PGAM_c	-34.769333	-969.691333
Enol_c	-34.769333	-969.691333
PEPC2_c	-3.412857	-147.549667
MalDH1_c	241.629905	-49.746333
FumHA_c	0.000000	0.000000
PGAK_h	227.898667	208.761333
GAPDH2_h	48.183667	-90.810333
Tr_TPT2	207.579333	0.000000
GAPDH3_c	61.825000	0.000000
Tr_TPT1	934.922000	0.000000