

```

model=modelOri;
model = changeRxnBounds(model,'EX_glc(e)',0,'1');
model = changeRxnBounds(model,'EX_succ(e)',-20,'1');
checkObjective(model);
FBAsolution = optimizeCbModel(model,'max');
FBAsolution.f
model = changeRxnBounds(model,'EX_o2(e)',0,'1');
FBAsolution = optimizeCbModel(model,'max');
FBAsolution.f

```

Warning: checkObjective will be depreciated because the function name is a misnomer, please use printObjective instead

summaryT =

23x5 table

Coefficient	Metabolite	metID	Reaction	RxnID
-1.496	3pg[c]	3	Biomass_Ecoli_core_N(w/GAM)-Nmet2	13
-3.7478	accoa[c]	10	Biomass_Ecoli_core_N(w/GAM)-Nmet2	13
59.81	adp[c]	13	Biomass_Ecoli_core_N(w/GAM)-Nmet2	13
4.1182	akg[c]	14	Biomass_Ecoli_core_N(w/GAM)-Nmet2	13
-59.81	atp[c]	17	Biomass_Ecoli_core_N(w/GAM)-Nmet2	13
3.7478	coa[c]	21	Biomass_Ecoli_core_N(w/GAM)-Nmet2	13
-0.361	e4p[c]	23	Biomass_Ecoli_core_N(w/GAM)-Nmet2	13
-0.0709	f6p[c]	26	Biomass_Ecoli_core_N(w/GAM)-Nmet2	13
-0.129	g3p[c]	33	Biomass_Ecoli_core_N(w/GAM)-Nmet2	13
-0.205	g6p[c]	34	Biomass_Ecoli_core_N(w/GAM)-Nmet2	13
-0.2557	gln-L[c]	36	Biomass_Ecoli_core_N(w/GAM)-Nmet2	13
-4.9414	glu-L[c]	38	Biomass_Ecoli_core_N(w/GAM)-Nmet2	13
-59.81	h2o[c]	41	Biomass_Ecoli_core_N(w/GAM)-Nmet2	13
59.81	h[c]	43	Biomass_Ecoli_core_N(w/GAM)-Nmet2	13
-3.547	nad[c]	50	Biomass_Ecoli_core_N(w/GAM)-Nmet2	13
3.547	nadh[c]	51	Biomass_Ecoli_core_N(w/GAM)-Nmet2	13
13.028	nadp[c]	52	Biomass_Ecoli_core_N(w/GAM)-Nmet2	13
-13.028	nadph[c]	53	Biomass_Ecoli_core_N(w/GAM)-Nmet2	13
-1.7867	oaa[c]	58	Biomass_Ecoli_core_N(w/GAM)-Nmet2	13
-0.5191	pep[c]	59	Biomass_Ecoli_core_N(w/GAM)-Nmet2	13
59.81	pi[c]	60	Biomass_Ecoli_core_N(w/GAM)-Nmet2	13
-2.8328	pyr[c]	62	Biomass_Ecoli_core_N(w/GAM)-Nmet2	13
-0.8977	r5p[c]	66	Biomass_Ecoli_core_N(w/GAM)-Nmet2	13

ans =

0.8401

ans =

NaN

