

GENERAL INFORMATION		FORMULATION					FEATURES					CASE STUDIES			AVAILABILITY		
METHOD	PUBLICATION URL	YEAR	FRAMEWORK	DESCRIPTION	SCALABILITY	TYPE	GPRs	DELETIONS	HETEROLOGOUS INSERTION	OVERLAPPING EXPRESSION	REGULATION	EXTRA FEATURES	ORGANISMS	MODELS	APPLICATIONS	SOFTWARE PACKAGES	DEPENDS
OptKnock	http://www.ncbi.nlm.nih.gov/pubmed/14595777	2003	BILEVEL	MIBLP - converted to single level MILP using duality theory	LOW	Exact	NO	YES	NO	NO	NO		<i>E. coli</i>	central carbon metabolism model (Edwards and Palsson, 2000)	Succinate, Lactate and 1,3-propanediol	Cobra toolbox, OptFlux	Matlab (Cobra)
OptStrain	http://www.ncbi.nlm.nih.gov/pubmed/15520298	2004	BILEVEL	MIBLP - converted to single level MILP using duality theory	LOW	Exact	NO	YES	YES	NO	NO		<i>E. coli</i> <i>C. acetobutylicum</i>	iJR904 and (Papoutsakis 1984; Desai et al, 1999)	Hydrogen and vanillin		
OptGene	http://www.ncbi.nlm.nih.gov/pubmed/16375763	2005	BILEVEL	Bi-level Heuristic. GA generates genotypes whose phenotypes are predicted by LP (FBA)	HIGH	Heuristic	NO	YES	NO	NO	NO		<i>S. cerevisiae</i>	IFF708	Vanillin, Glycerol and Succinate	OptFlux, Cobra toolbox	Matlab (Cobra)
OptReg	http://www.ncbi.nlm.nih.gov/pubmed/16199194	2006	BILEVEL	MIBLP - converted to single level MILP using duality theory	LOW	Exact	NO	YES	NO	YES	NO		<i>E. coli</i>	iJR904	Ethanol		
MMF	http://www.sciencedirect.com/science/article/pii/S10967176060067X	2006	EMA	EFMs	LOW	Exact	NO	NO	NO	NO	NO		<i>E. coli</i>	core of intermediary metabolism (Carlson and Srienc, 2004)	Biomass		
EAs/SAs	http://www.ncbi.nlm.nih.gov/pubmed/19038030	2008	BILEVEL	Bi-level Heuristic. EA/SA generate genotypes whose phenotypes are predicted by LP (FBA)	HIGH	Heuristic	YES	YES	NO	NO	NO		<i>E. coli</i> <i>S. cerevisiae</i>	iJR904 IND750	Succinate and Lactate	OptFlux	
GDLS	http://www.ncbi.nlm.nih.gov/pubmed/19690565	2009	BILEVEL	MIBLP - converted to single level MILP using duality theory	MEDIUM	Heuristic	YES	YES	NO	NO	NO		<i>E. coli</i>	iAF1260	Acetate and Succinate	Cobra toolbox; author's homepage: http://crab.rutgers.edu/~dslu/n/gdls/	Matlab
CIED	http://www.ncbi.nlm.nih.gov/pubmed/19633125	2009	BILEVEL	Bi-level Heuristic. Evolutionary cipher	HIGH	Heuristic	YES	YES	NO	NO	NO		<i>E. coli</i>	iJR904	Flavones		
FluxDesign	http://www.ncbi.nlm.nih.gov/pubmed/20035624	2009	EMA	EFMs	LOW	Exact	NO	YES	NO	YES	NO		<i>C. glutamicum</i> <i>A. niger</i>	(simplified models)	Lysine and Enzymes		
RobustKnock	http://www.ncbi.nlm.nih.gov/pubmed/20031969	2010	Multi-LEVEL	Max-min MIBLP - converted to single level MILP using duality theory	LOW	Exact	NO	YES	NO	NO	NO		<i>E. coli</i>	iJR904	Hydrogen, Acetate, Formate and Fumarate	Author's homepage: http://www.cs.technion.ac.il/~tomersh/methods.html	
OptORF	http://www.biomedcentral.com/1752-0509/4/53/	2010	BILEVEL	MIBLP - converted to single level MILP using duality theory	LOW	Exact	YES	YES	NO	YES	YES		<i>E. coli</i>	IMC1010 (v2)	Ethanol, isobutanol and 2-phenylethanol		
OptForce	http://www.ncbi.nlm.nih.gov/pubmed/20419153	2010		LP	LOW	Exact	NO	NO	NO	YES	NO		<i>E. coli</i>	iAF1260	Succinate	Available in the author's homepage upon request: http://maranas.che.psu.edu/submit/optforce.htm	
CASOP	http://www.ncbi.nlm.nih.gov/pubmed/20303369 http://subs.emis.de/LNI/Proceedings/Proceedings173/P_173.pdf?orjin=publication_detail#page=22	2010	EMA	EFMs	LOW	Heuristic	NO	YES	NO	YES	NO		<i>E. coli</i>	central carbon metabolism model	Succinate	CellNetAnalyzer	Matlab
CASOP-GS	http://www.ncbi.nlm.nih.gov/pubmed/21147248	2010	EMA	EFMs	HIGH	Heuristic	NO	YES	NO	YES	NO		<i>E. coli</i>	iAF1260	Succinate and histidine	Available on request (including source)	
cMCS	http://www.ncbi.nlm.nih.gov/pubmed/21147248	2011	EMA	EFMs	LOW	Exact	NO	YES	NO	NO	NO		<i>E. coli</i>	(Trinh et al., 2008)	Ethanol	CellNetAnalyzer	Matlab
BIMOMA	http://www.ncbi.nlm.nih.gov/pubmed/21949695	2011	BILEVEL	MIQCP	MEDIUM	Exact	NO	YES	NO	NO	NO		<i>E. coli</i>	iJR904	Glutamate and pyruvate		
SimOptStrain	http://www.ncbi.nlm.nih.gov/pubmed/21949695	2011	BILEVEL	MILP	LOW	Exact	YES	YES	YES	NO	NO		<i>E. coli</i>	iJR904	Succinate and glycerol		
EMILIO	http://www.ncbi.nlm.nih.gov/pubmed/21414417	2011		Iterative LP (ILP)	HIGH	Exact	NO	YES	NO	YES	NO		<i>E. coli</i>	iAF1260	Succinate, glutamate and L-serine		
EAs/SAs Regulation	http://www.ncbi.nlm.nih.gov/pubmed/21144882	2011	BILEVEL	(EA/SA) FBA/SR-FBA	HIGH	Heuristic	YES	YES	NO	NO	YES		<i>E. coli</i>	iAF1260	Ethanol	OptFlux plugin	
GDMO	http://www.ncbi.nlm.nih.gov/pubmed/23044547	2012	BILEVEL	NSGA-II based search, FBA evaluation	HIGH	Heuristic	YES	YES	NO	NO	NO		<i>E. coli</i>	iAF1260	Succinate and acetate		
GDBB	http://www.ncbi.nlm.nih.gov/pubmed/22543499	2012	BILEVEL	MIBLP converted to MILP based implementation. TBB, prevents the branch and bound algorithm from reaching optimal solution and stops at near-optimality.	HIGH	Heuristic	YES	YES	NO	YES	NO		<i>E. coli</i>	iAF1260	Succinate and acetate	Author's homepage: http://crab.rutgers.edu/~dslu/n/gddb	Matlab
EAs/SAs Over-under expression	http://online.liebertpub.com/dp/abs/10.1089/cmb.2011.0265	2012	BILEVEL	(EA/SA) FBA/MOMA/ROOM	HIGH	Heuristic	YES	YES	NO	YES	NO		<i>E. coli</i>	iJR904	Succinate and Lactate	OptFlux	

rcMCS	http://www.biomedcentral.com/1752-0509/6/103/abstract	2012	EMA	BLP(Binary Linear Programming)	LOW	Exact	YES	YES	NO	NO	YES	<i>E. coli</i>	(Trinh et al., 2008)	Ethanol	
MOMAknock	http://www.ncbi.nlm.nih.gov/pubmed/23368729	2013	BILEVEL	MIBQP - converted to single level MILP using duality theory	LOW	Exact	YES	YES	NO	NO	NO	<i>E. coli</i>	IAF1260	Succinate	
Redirector	http://www.ncbi.nlm.nih.gov/pubmed/23341769	2013	BILEVEL	MIBLP - converted to single level MILP using duality theory	MEDIUM	Heuristic	YES	NO	NO	YES	NO	<i>E. coli</i>	IAF1260	myristoyl-CoA	Available from author's site: https://github.com/bionomicron/Redirector.git
OptSwap	http://online.liebertpub.com/doi/abs/10.1089/ind.2013.0005	2013	MULTI-LEVEL (similar to RobustKnock)	Max-min MIBLP - converted to single level MILP using duality theory	LOW	Exact	NO	YES	NO	NO	NO	<i>E. coli</i>	IO1366	production of a wide spectrum of products	Swapping of the cofactor specificity for oxyreductase enzymes
ReacKnock	http://www.ncbi.nlm.nih.gov/pubmed/24348984/	2013	BILEVEL	MIBLP - converted to single level MILP using Karush-Kuhn-Tucker for continuous inner problems.	LOW	Exact	NO	YES	NO	NO	NO	<i>E. coli</i>	IAF1260	Succinate, ethanol and threonine	
SMET	http://online.liebertpub.com/doi/10.1002/biot.201200233/abstract	2013	EMA	cMCS	LOW	Exact	NO	YES	NO	YES	NO	<i>E. coli</i>	(Rizk and Liao, 2009)	DAHP	
FastPros	http://www.ncbi.nlm.nih.gov/pubmed/24257186	2013	Multi-Step	Exhaustive computation of dual mutants, LP for calculation of u-targets followed by iterative single mutant addition for best. Screening via OptKnock (MILP)	HIGH	Heuristic	NO	YES	NO	NO	NO	<i>E. coli</i>	IAF1260	production of every cytosolic metabolite	http://www.shimizu-ist.osaka-u.ac.jp/shimizu_lab/FastPros/ MatLab, COBRA
CosMos	http://online.liebertpub.com/doi/10.1002/biot.201200316/full	2013	BILEVEL	MILP	LOW	Exact	YES	YES	NO	YES	NO	<i>E. coli</i> <i>S. cerevisiae</i>	IO1366 IMM904	Chorismate, ethanol, succinate, L-amino acids	
MCS Enumerator	http://www.ploscompbiol.org/article/info%3Adoi%2F10.1371%2Fjournal.pcbi.1003378	2014	EMA	MILP	HIGH	Exact	NO	YES	NO	NO	NO	<i>E. coli</i>	IAF1260	Fumarate and serine	CellNetAnalyzer Matlab
k-OptForce	http://www.ploscompbiol.org/article/info%3Adoi%2F10.1371%2Fjournal.pcbi.1003487 Curran1	2014	BILEVEL	MILP	LOW	Exact	NO	YES	NO	YES	NO	<i>E. coli</i> <i>S. cerevisiae</i>	IAF1260 + (Chassagnole et al., 2002) IAZ900 + (van Eunen et al., 2012)	L-serine and TAL	Hybrid stoichiometric and kinetic