

**Supporting Table S1: Sources of biochemistry integrated into PlantSEED**

Database/Model	Species	Owner/Reference	Date / Version	Number of Compounds	Number of Reactions
KEGG	N/A	KEGG (1, 2)	12/03/18	17,793	14,039
MetaCyc	N/A	PathwayTools (3)	22.5	19,172	27,414
EcoCyc	<i>Escherichia coli</i>	PathwayTools (3)	16.1	3,355	2,411
PlantCyc	N/A	PMN (4, 5)	7.0	3,919	4,436
AraCyc	<i>Arabidopsis thaliana</i>	PMN (5)	10.0	4,229	4,008
PoplarCyc	<i>Populus trichocarpa</i>	PMN (4)	4.0	2,318	2,301
SoyCyc	<i>Glycine max</i>	PMN	2.0	2,796	2,657
ChlamyCyc	<i>Chlamydomonas reinhardtii</i>	PMN	03/03/10	1,718	1,875
BrachyCyc	<i>Brachypodium distachyon</i>	Gramene (6, 7)	2.0	2,348	3,104
MaizeCyc	<i>Zea mays</i>	Gramene (8)	2.0	2,219	2,358
RiceCyc	<i>Oryza sativa</i>	Gramene (9)	2.0.1	1,859	1,955
SorghumCyc	<i>Sorghum bicolor</i>	Gramene (6, 7)	1.0.1	1,773	1,873
iAF1260	<i>Escherichia coli</i>	Feist <i>et al.</i> (2007) (10)		1,041	2,064
iAF692	<i>Methanosarcina barkeri</i>	Feist <i>et al.</i> (2006) (11)		562	613
iIN800	<i>Saccharomyces cerevisiae</i>	Nookaew <i>et al.</i> (2008) (12)		683	1,053
iJR904	<i>Escherichia coli</i>	Reed <i>et al.</i> (2003) (13)		629	921
iMA945	<i>Salmonella spp.</i>	AbuOun <i>et al.</i> (2009) (14)		1,032	1,960
iMM904	<i>Saccharomyces cerevisiae</i>	Mo <i>et al.</i> (2009) (15)		712	1,401
iRR1083	<i>Salmonella typhimurium</i> LT2	Raghunathan <i>et al.</i> (2009) (16)		759	1,086
iSB619	<i>Staphylococcus aureus</i> N315	Becker <i>et al.</i> (2005) (17)		614	639
iSO783	<i>Shewanella oneidensis</i> MR-1	Pinchuk <i>et al.</i> (2010) (18)		634	774
iAbaylyiv4	<i>Acinetobacter baylyi</i> ADP1	Durot <i>et al.</i> (2008) (19)		699	867
*	<i>Bacillus subtilis</i>	Goelzer <i>et al.</i> (2008) (20)		475	504
iGT196	<i>Buchnera aphidicola</i>	Thomas <i>et al.</i> (2009) (21)		740	210
iIT341	<i>Helicobacter pylori</i>	Thiele <i>et al.</i> (2005) (22)		411	473
iJN746	<i>Pseudomonas putida</i> KT2440	Nogales <i>et al.</i> (2008) (23)		706	915
iMO1056	<i>Pseudomonas aeruginosa</i> PAO1	Oberhardt <i>et al.</i> (2008) (24)		750	864
iND750	<i>Saccharomyces cerevisiae</i>	Duarte <i>et al.</i> (2004) (25)		650	1,038

iNJ661	<i>Mycobacterium tuberculosis</i> H37Rv	Jamshidi & Palsson (2007) (26)	761	951
iPS189	<i>Mycoplasma genitalium</i>	Suthers <i>et al.</i> (2009) (27)	277	262
iRS1563	<i>Zea mays</i>	Saha <i>et al.</i> (2011) (28)	1,812	1,949
iRS1597	<i>Arabidopsis thaliana</i>	Saha <i>et al.</i> (2011) (28)	1,759	1,837
iYO844	<i>Bacillus subtilis</i>	Oh <i>et al.</i> (2007) (29)	776	1,016
*	<i>Chlamydomonas reinhardtii</i>	Boyle & Morgan (2009) (30)	266	485
*	<i>Chlamydomonas reinhardtii</i>	Manichaikul <i>et al.</i> (2009) (31)	124	238
*	<i>Chlamydomonas reinhardtii</i>	Chang <i>et al.</i> (2011) (32)	1,164	2,084
C4GEM	<i>Zea mays</i>	D'al Molin <i>et al.</i> (2010) (33)	1,207	1,227
*	<i>Arabidopsis thaliana</i>	Mintz-Oron <i>et al.</i> (2012) (34)	1,181	3,382
*	<i>Arabidopsis thaliana</i>	Poolman <i>et al.</i> (2009) (35)	1,224	1,354
AraGEM	<i>Arabidopsis thaliana</i>	D'al Molin <i>et al.</i> (2010) (36)	1,546	1,590
AlgaGEM	<i>Chlamydomonas reinhardtii</i>	D'al Molin <i>et al.</i> (2011) (37)	1,662	1,713

The numbers of compounds and reactions are listed for each source *after* integration, and may not reflect the numbers seen in the literature. In addition, the number of reactions includes compartmentalized reactions, which may be duplicates. \*No unique identifier for these models was described in the literature.

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