

Additional file 3. ORF (Open Reading Frame) coverage of reconstructed metabolic models

Organism	¹ Model	Coverage (%)	Ref
<i>Escherichia coli</i>	iAF1260	28.0	Feist <i>et al</i> (2007)
<i>Saccharomyces cerevisiae</i>	iND750	12.0	Duarte <i>et al</i> (2004)
	iMM904	14.4	Mo <i>et al</i> (2009)
<i>Aspergillus nidulans</i>	iHD666	7.0	David <i>et al</i> (2008)
<i>Aspergillus niger</i>		6.1	David <i>et al</i> (2003)
<i>Arabidopsis thaliana</i>	AraGEM	5.2	de Oliveira Dal'Molin <i>et al</i> (2010)
<i>Mus musculus</i>		3.0	Selvarasu <i>et al</i> (2010)
<i>Homo sapiens</i>	Recon 1	6.3	Duarte <i>et al</i> (2007)
<i>Acinetobacter baumannii</i>	AbyMBEL891	17.3	Kim <i>et al</i> (2010)
<i>Helicobacter pylori</i>	iIT341	26.7	Schilling <i>et al</i> (2008)

1. There are metabolic models which do not have its own name.

References

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