

Supplementary file 1 A: Model equations

Reaction nr	reaction	Function	Present in model(s)
1 PTS_Glc Glucose proton symporter	GLC_Ext + PEP -> G6P + PYR; FBP Pi_Int	$\frac{\text{PTS_Glc} \cdot \frac{\text{Pi_Int}}{\text{Pi_Int} + \text{kaPi_Int_PTS_Glc}} \cdot \frac{\text{kiFBP_PTS_Glc}}{\text{FBP} + \text{kiFBP_PTS_Glc}} \cdot \text{kcat_PTS_Glc} \cdot \frac{\text{GLC_Ext}}{\text{kmGLC_PTS_Glc}} \cdot \frac{\text{PEP}}{\text{kmPEP_PTS_Glc}}}{\left(1 + \frac{\text{GLC_Ext}}{\text{kmGLC_PTS_Glc}}\right) \cdot \left(1 + \frac{\text{PEP}}{\text{kmPEP_PTS_Glc}}\right) + \left(1 + \frac{\text{G6P}}{\text{kmG6P_PTS_Glc}}\right) \cdot \left(1 + \frac{\text{PYR}}{\text{kmPYR_PTS_Glc}}\right) - 1}$	1,2,3,4,5,6
2 PGI Glucose-6-phosphate isomerase	G6P = F6P	$\frac{\text{PGI} \cdot \left(\text{kcat_PGI} \cdot \frac{\text{G6P}}{\text{kmG6P_PGI}} - \frac{\text{kcat_PGI}}{\text{Keq_PGI}} \cdot \frac{\text{F6P}}{\text{kmG6P_PGI}} \right)}{1 + \frac{\text{G6P}}{\text{kmG6P_PGI}} + \frac{\text{F6P}}{\text{kmF6P_PGI}}}$	1,2,3,4,5,6
3 PFK Phosphofructokinase	F6P + ATP -> FBP + ADP	$\frac{\text{PFK} \cdot \text{kcat_PFK} \cdot \frac{\text{ATP}}{\text{kmF6P_PFK}} \cdot \frac{\text{F6P}}{\text{kmATP_PFK}}}{\left(1 + \frac{\text{ATP}}{\text{kmF6P_PFK}}\right) \cdot \left(1 + \frac{\text{F6P}}{\text{kmATP_PFK}}\right) + \left(1 + \frac{\text{ADP}}{\text{kmFBP_PFK}}\right) \cdot \left(1 + \frac{\text{FBP}}{\text{kmADP_PFK}}\right) - 1}$	1,2,3,4,5,6
4 FBA Fructose-bisphosphate aldolase	GAP + Pi_Int + NAD = DGP + NADH	$\frac{\text{FBA} \cdot \left(\text{kcat_FBA} \cdot \frac{\text{FBP}}{\text{kmFBP_FBA}} - \frac{\text{kcat_FBA}}{\text{Keq_FBA}} \cdot \frac{\text{GAP}^2}{\text{kmFBP_FBA}} \right)}{1 + \frac{\text{FBP}}{\text{kmFBP_FBA}} + \frac{\text{GAP}}{\text{kmGAP_FBA}} + \left(\frac{\text{GAP}}{\text{kmGAP_FBA}} \right)^2}$	1,2,3,4,5,6
5 GAPDH Glyceraldehyde-3-phosphate dehydrogenase	DGP + ADP = PEP + ATP	$\frac{\text{GAPDH} \cdot \left(\text{kcat_GAPDH} \cdot \frac{\text{GAP}}{\text{kmGAP_GAPDH}} \cdot \frac{\text{NAD}}{\text{kmNAD_GAPDH}} \cdot \frac{\text{Pi_Int}}{\text{kmPi_Int_GAPDH}} - \frac{\text{kcat_GAPDH}}{\text{Keq_GAPDH}} \cdot \frac{\text{DGP}}{\text{kmGAP_GAPDH}} \cdot \frac{\text{NADH}}{\text{kmNAD_GAPDH}} \right)}{\left(1 + \frac{\text{GAP}}{\text{kmGAP_GAPDH}}\right) \cdot \left(1 + \frac{\text{Pi_Int}}{\text{kmPi_Int_GAPDH}}\right) \cdot \left(1 + \frac{\text{NAD}}{\text{kmNAD_GAPDH}}\right) + \left(1 + \frac{\text{DGP}}{\text{kmDGP_GAPDH}}\right) \cdot \left(1 + \frac{\text{NADH}}{\text{kmNADH_GAPDH}}\right) - 1}$	1,2,3,4,5,6

6 ENO Enolase	PEP + ADP -> PYR + ATP; FBP Pi_Int	$\text{ENO} \cdot \left(\frac{\text{DGP}}{\text{kmDGP_ENO}} \cdot \frac{\text{ADP}}{\text{kmADP_ENO}} \cdot \frac{\text{kcat_ENO}}{\text{Keq_ENO}} \cdot \frac{\text{PEP}}{\text{kmDGP_ENO}} \cdot \frac{\text{ATP}}{\text{kmADP_ENO}} \right)$ $\left(1 + \frac{\text{DGP}}{\text{kmDGP_ENO}} \right) \cdot \left(1 + \frac{\text{ADP}}{\text{kmADP_ENO}} \right) + \left(1 + \frac{\text{PEP}}{\text{kmPEP_ENO}} \right) \cdot \left(1 + \frac{\text{ATP}}{\text{kmATP_ENO}} \right) - 1$	1,2,3,4,5,6
7 PYK Pyruvate kinase	PYR + CoA + NAD = AcCoA + NADH; GAP	$\frac{\text{PYK} \cdot \text{FBP}}{\text{FBP} + \text{kaFBP_PYK}} \cdot \frac{\text{kiPi_Int_PYK}^{n\text{PYK}}}{\text{Pi_Int}^{n\text{PYK}} + \text{kiPi_Int_PYK}^{n\text{PYK}}} \cdot \text{kcat_PYK} \cdot \frac{\text{ADP}}{\text{kmADP_PYK}} \cdot \frac{\text{PEP}}{\text{kmPEP_PYK}}$ $\left(1 + \frac{\text{ADP}}{\text{kmADP_PYK}} \right) \cdot \left(1 + \frac{\text{PEP}}{\text{kmPEP_PYK}} \right) + \left(1 + \frac{\text{ATP}}{\text{kmATP_PYK}} \right) \cdot \left(1 + \frac{\text{PYR}}{\text{kmPYR_PYK}} \right) - 1$	1,2,3,4,5,6
8 LDH Lactate dehydrogenase	PYR + NADH -> LAC + NAD; FBP Pi_Int* O2	$\frac{\text{LDH} \cdot \text{FBP}}{\text{FBP} + \text{kaFBP_LDH}} \cdot \frac{\text{kiPi_Int_LDH}}{\text{Pi_Int} + \text{kiPi_Int_LDH}} \cdot \frac{\text{kiO2_LDH}}{\text{O2} + \text{kiO2_LDH}} \cdot \text{kcat_LDH} \cdot \frac{\text{NADH}}{\text{kmPYR_LDH}} \cdot \frac{\text{PYR}}{\text{kmNADH_LDH}}$ $\left(1 + \frac{\text{NADH}}{\text{kmPYR_LDH}} \right) \cdot \left(1 + \frac{\text{PYR}}{\text{kmNADH_LDH}} \right) + \left(1 + \frac{\text{LAC}}{\text{kmLAC_LDH}} \right) \cdot \left(1 + \frac{\text{NAD}}{\text{kmNAD_LDH}} \right) - 1$	1*,2*,3*,4*,5,6
9 PDH Pyruvate dehydrogenase	PYR + CoA + NAD = AcCoA + NADH; GAP	$\text{PDH} \cdot \frac{\text{kiGAP_PDH}}{\text{GAP} + \text{kiGAP_PDH}} \cdot \left(\text{kcat_PDH} \cdot \frac{\text{NAD}}{\text{kmPYR_PDH}} \cdot \frac{\text{PYR}}{\text{kmCoA_PDH}} \cdot \frac{\text{CoA}}{\text{kmNAD_PDH}} \cdot \frac{\text{kcat_PDH}}{\text{Keq_PDH}} \cdot \frac{\text{AcCoA}}{\text{kmPYR_PDH}} \cdot \frac{\text{NADH}}{\text{kmCoA_PDH}} \right)$ $\left(1 + \frac{\text{NAD}}{\text{kmPYR_PDH}} \right) \cdot \left(1 + \frac{\text{PYR}}{\text{kmCoA_PDH}} \right) \cdot \left(1 + \frac{\text{CoA}}{\text{kmNAD_PDH}} \right) + \left(1 + \frac{\text{AcCoA}}{\text{kmAcCoA_PDH}} \right) \cdot \left(1 + \frac{\text{NADH}}{\text{kmNADH_PDH}} \right) - 1$	1,2,3,4,5,6
10 PTA_ACK Phosphotransacetylase	AcCoA + ADP -> ACE + ATP + CoA	$\frac{\text{PTA_ACK} \cdot \text{kcat_PTA_ACK} \cdot \frac{\text{AcCoA}}{\text{kmAcCoA_PTA_ACK}} \cdot \frac{\text{ADP}}{\text{kmADP_PTA_ACK}}}{\left(1 + \frac{\text{AcCoA}}{\text{kmAcCoA_PTA_ACK}} \right) \cdot \left(1 + \frac{\text{ADP}}{\text{kmADP_PTA_ACK}} \right) + \left(1 + \frac{\text{ACE}}{\text{kmACE_PTA_ACK}} \right) \cdot \left(1 + \frac{\text{ATP}}{\text{kmATP_PTA_ACK}} \right) \cdot \left(1 + \frac{\text{CoA}}{\text{kmCoA_PTA_ACK}} \right) - 1}$	1,2,3,4,5,6
11 ATPase	ATP -> ADP + Pi_Int	$\frac{\text{ATPase} \cdot \text{kcat_ATPase} \cdot \left(\frac{\text{ATP}}{\text{kmATP_ATPase}} \right)^{n\text{ATPase}}}{\left(\frac{\text{ATP}}{\text{kmATP_ATPase}} \right)^{n\text{ATPase}} + 1}$	2,4,6
12 NOXE NADH oxidase	O2 + 2 * NADH -> 2 * NAD	$\frac{\text{NOXE} \cdot \text{kcat_NOXE} \cdot \left(\frac{\text{O2}}{\text{kmNADH_NOXE}} \right)^2 \cdot \frac{\text{NADH}}{\text{kmO2_NOXE}}}{\left(1 + \frac{\text{O2}}{\text{kmNADH_NOXE}} \right)^2 \cdot \left(1 + \frac{\text{NADH}}{\text{kmO2_NOXE}} \right) + 1 + \frac{\text{NAD}}{\text{kmNAD_NOXE}}} - 1$	3,4,6

*Models marked with an asterisk contain the function for Lactate Dehydrogenase without the oxygen inhibition term $\frac{kiO2_LHD}{O2+kiO2_LDH}$

Supplementary file 1 B comparison of model additions

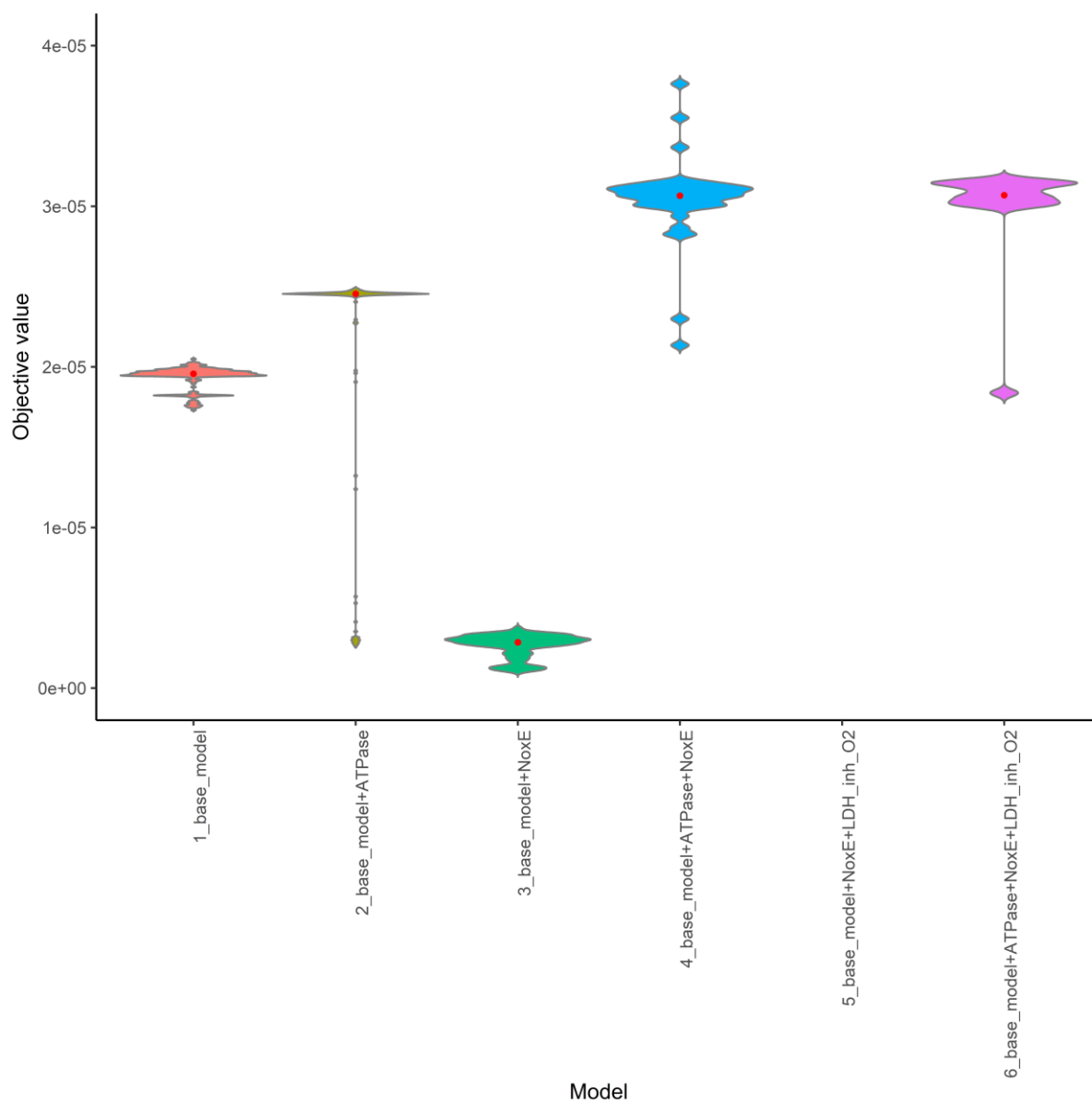


Figure 1 Zoomed in version of comparison of 100 parameter sets for the model with and without additions. Outliers on the top were removed to better show overall performance of the models.

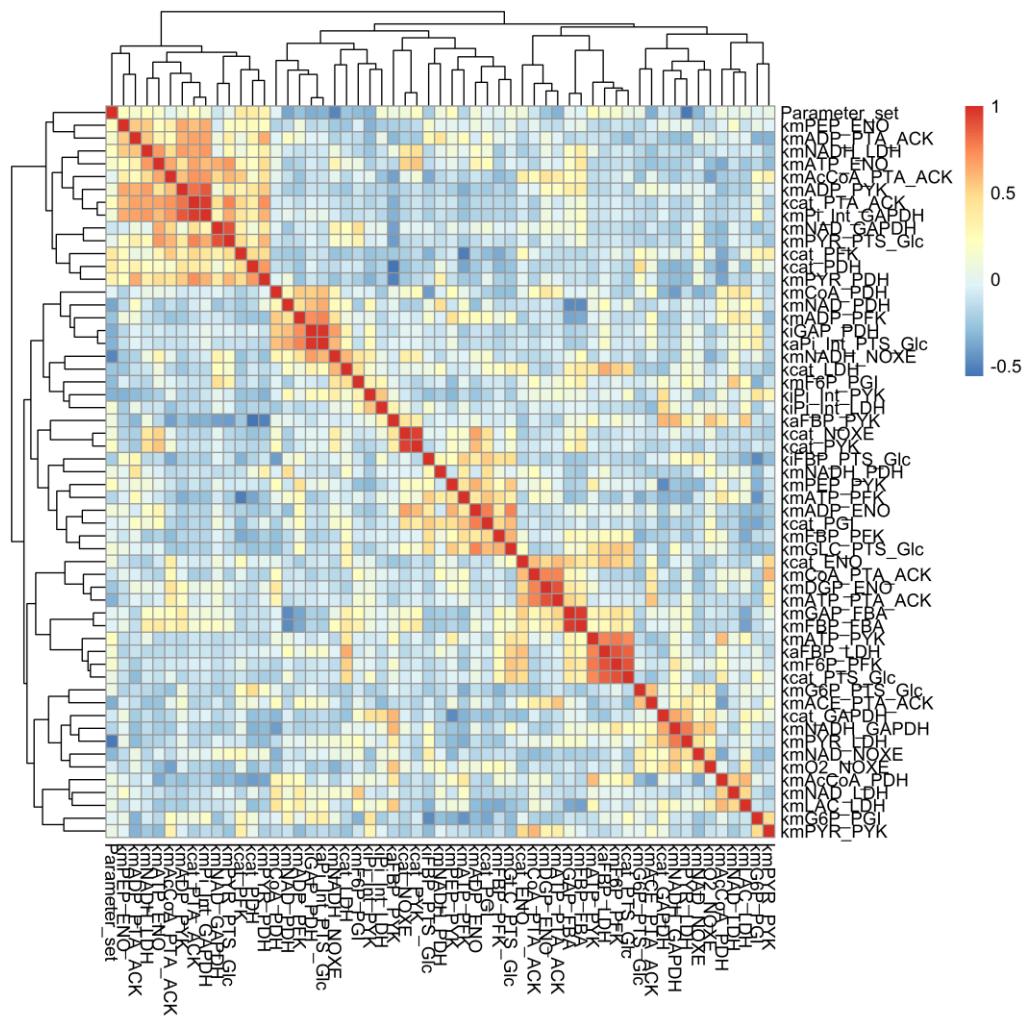
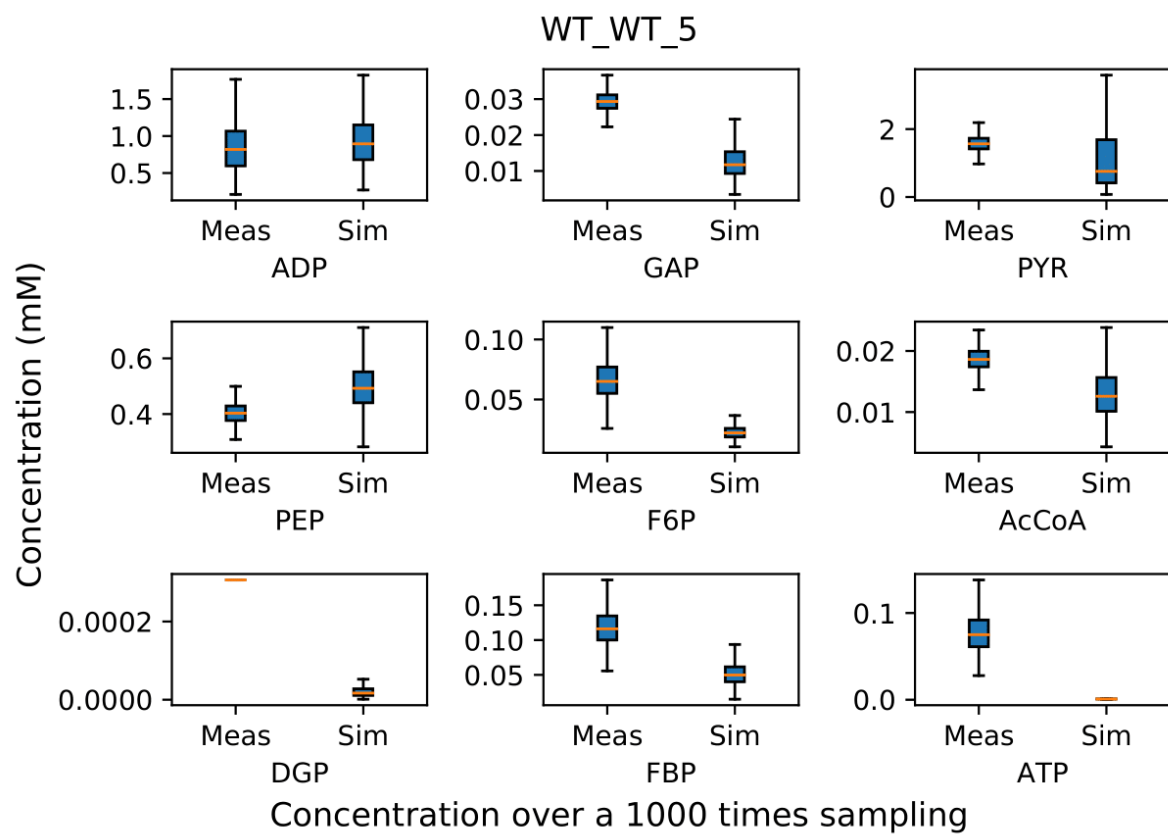
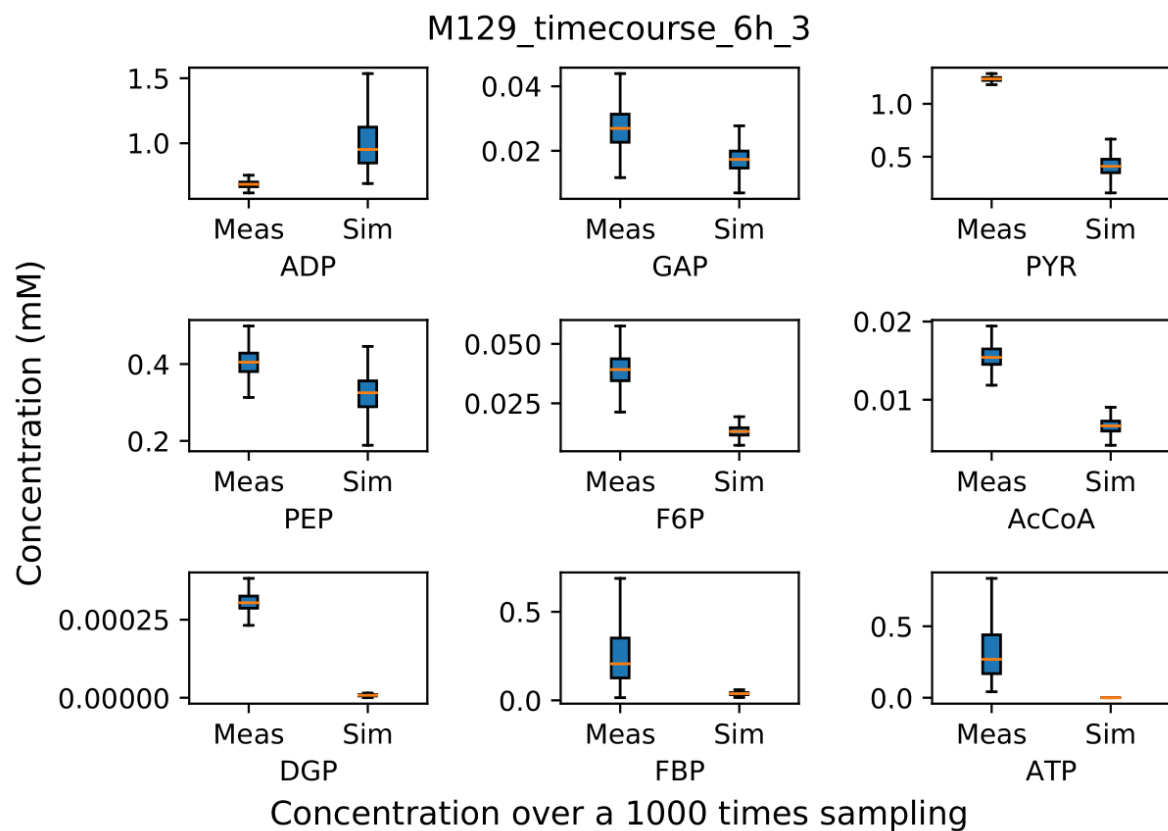


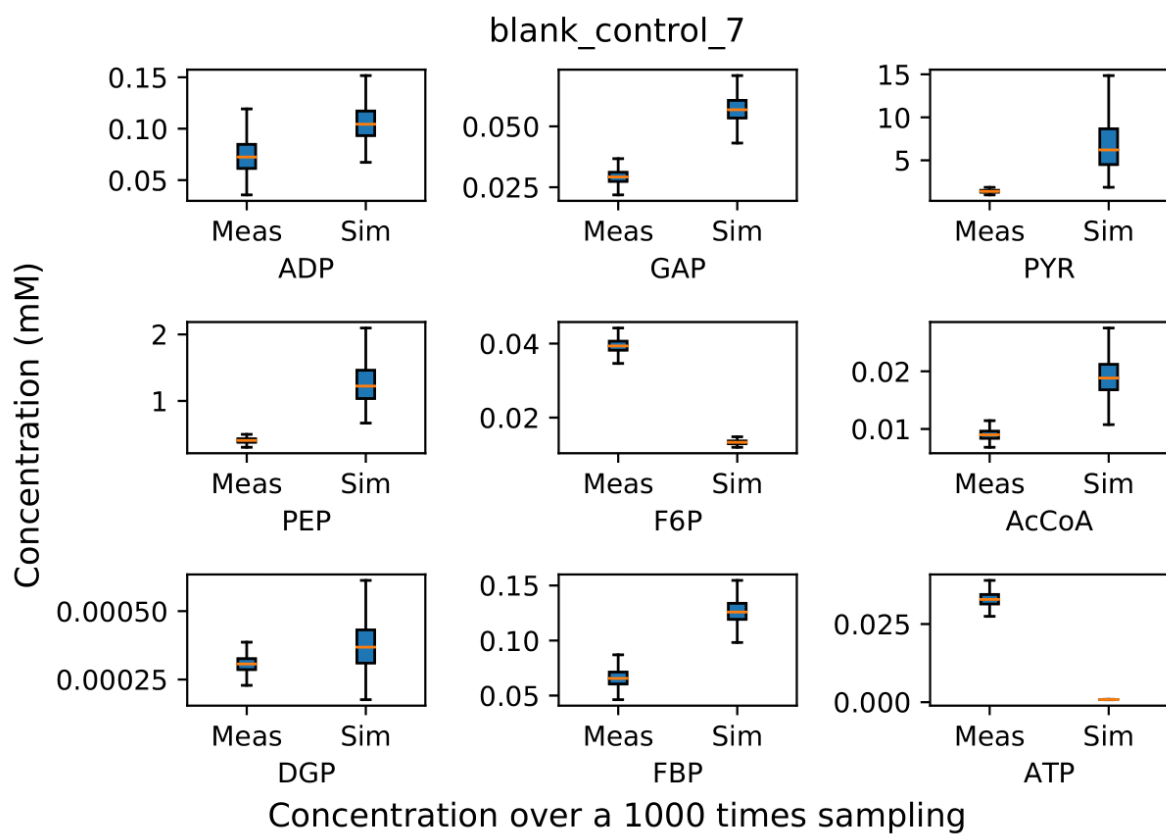
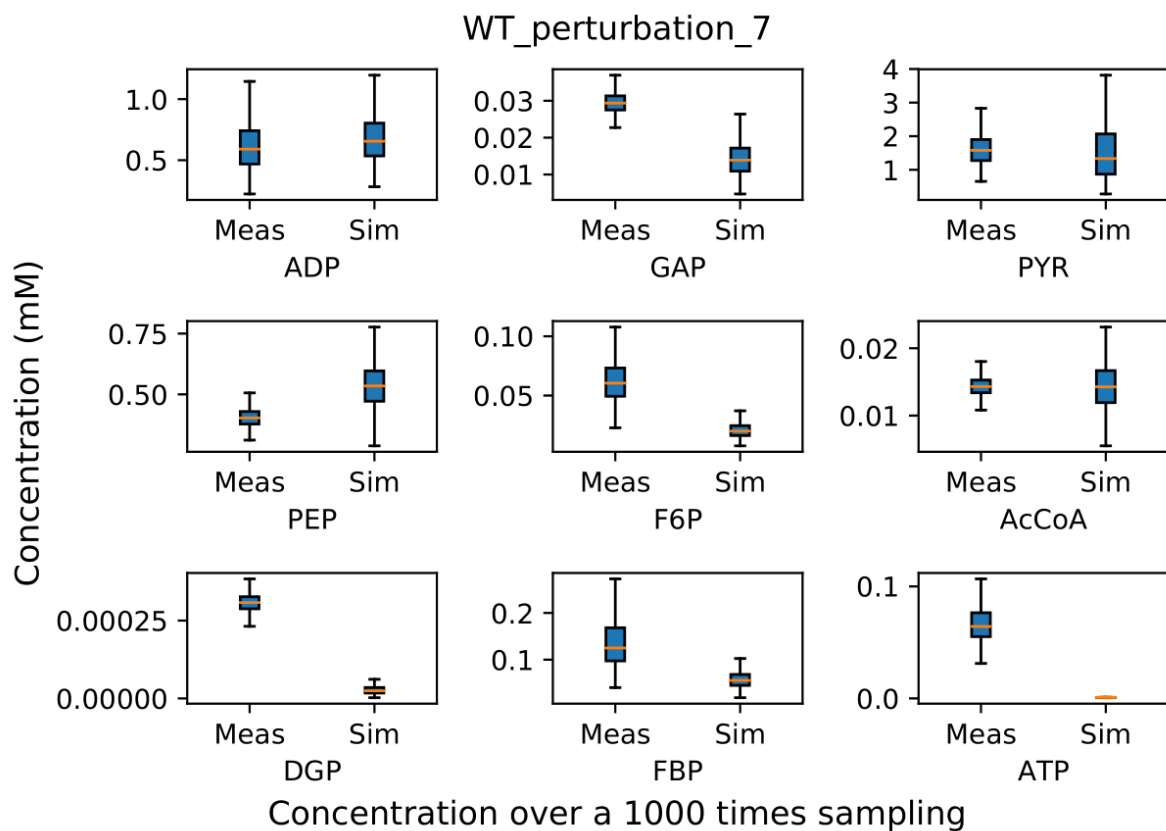
Figure 2 Correlation of the parameter sets for model 3 that includes NoxE for the best 10 performing parameter sets.

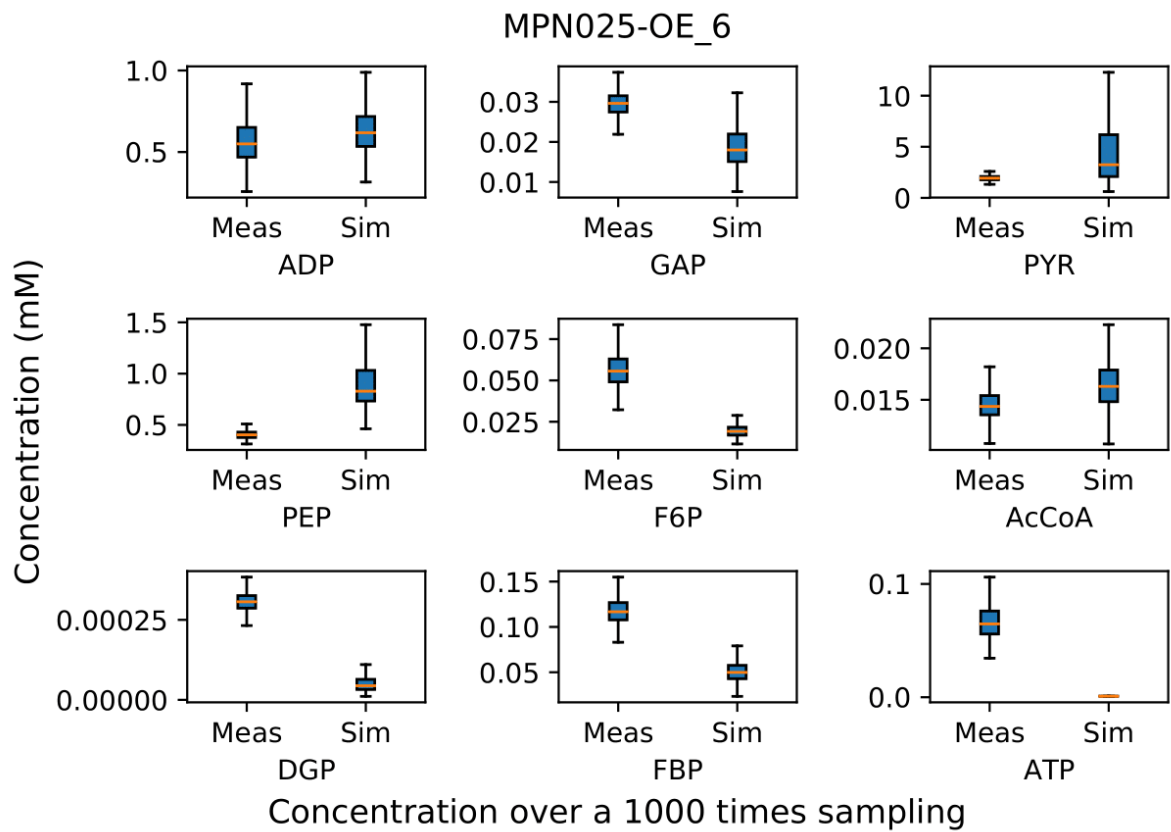
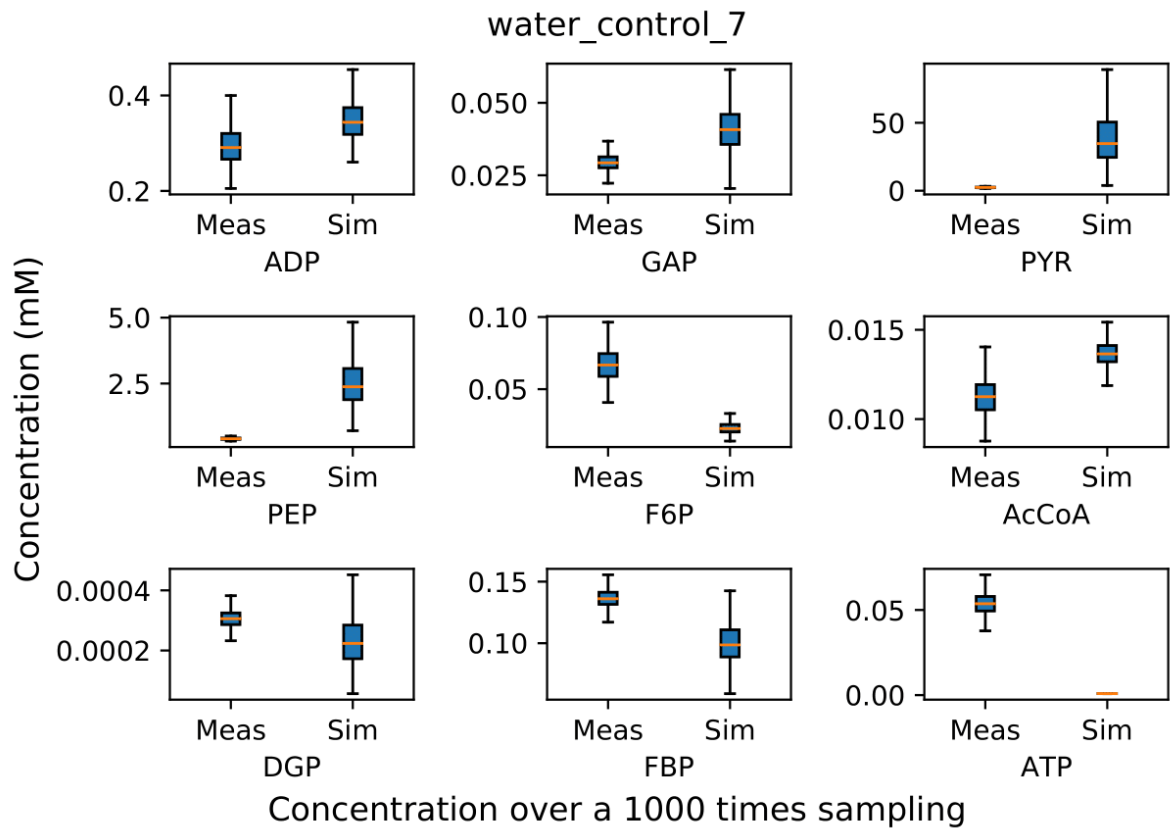
Parameter distributions for each of the 6 models over 100 parameter sets as well as heatmaps showing the correlation between parameters over all 100 fittings for each of the six model configurations are available upon request.

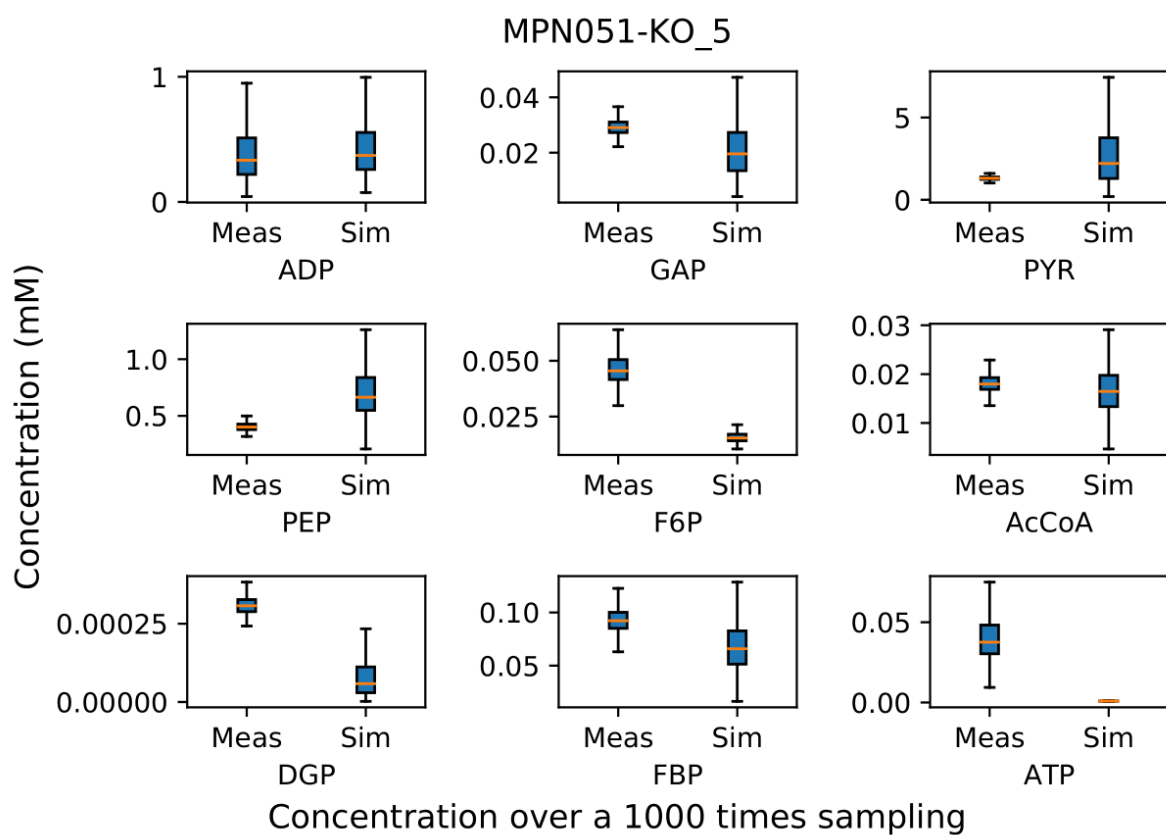
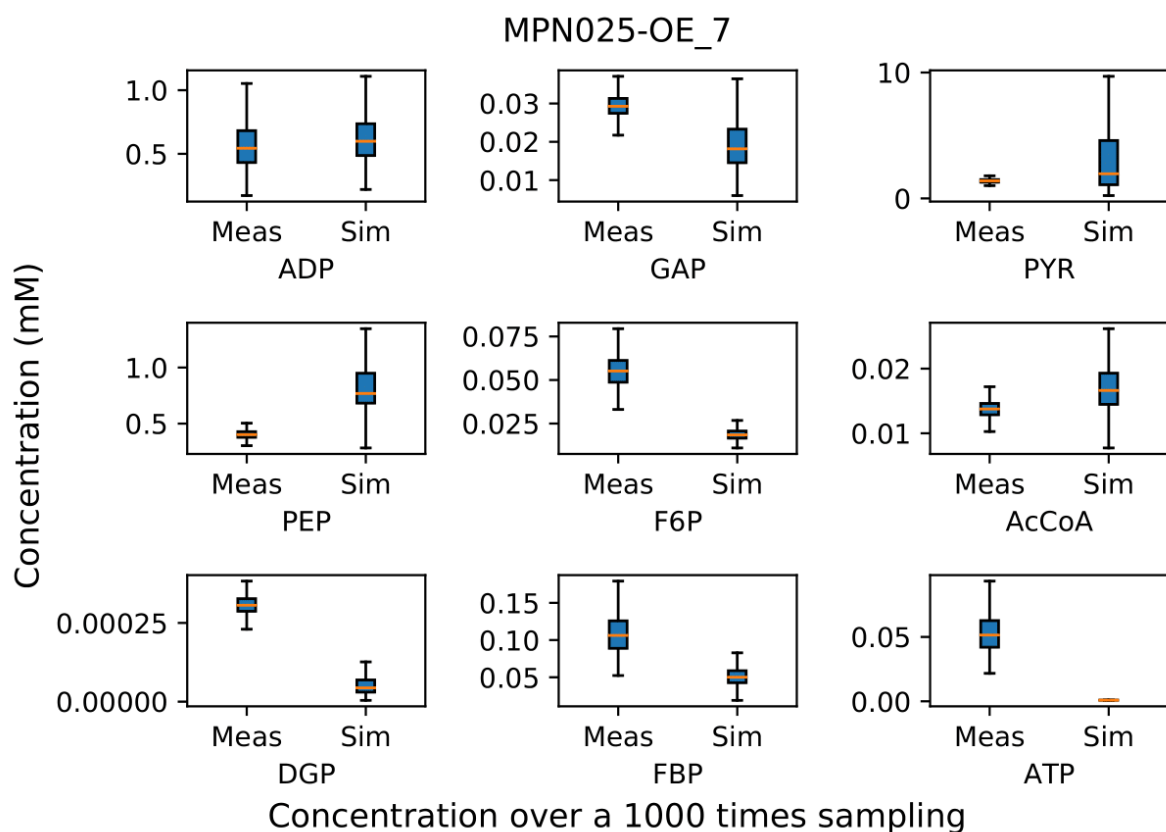
Supplementary file 1 C Simulating 40 independent samples.

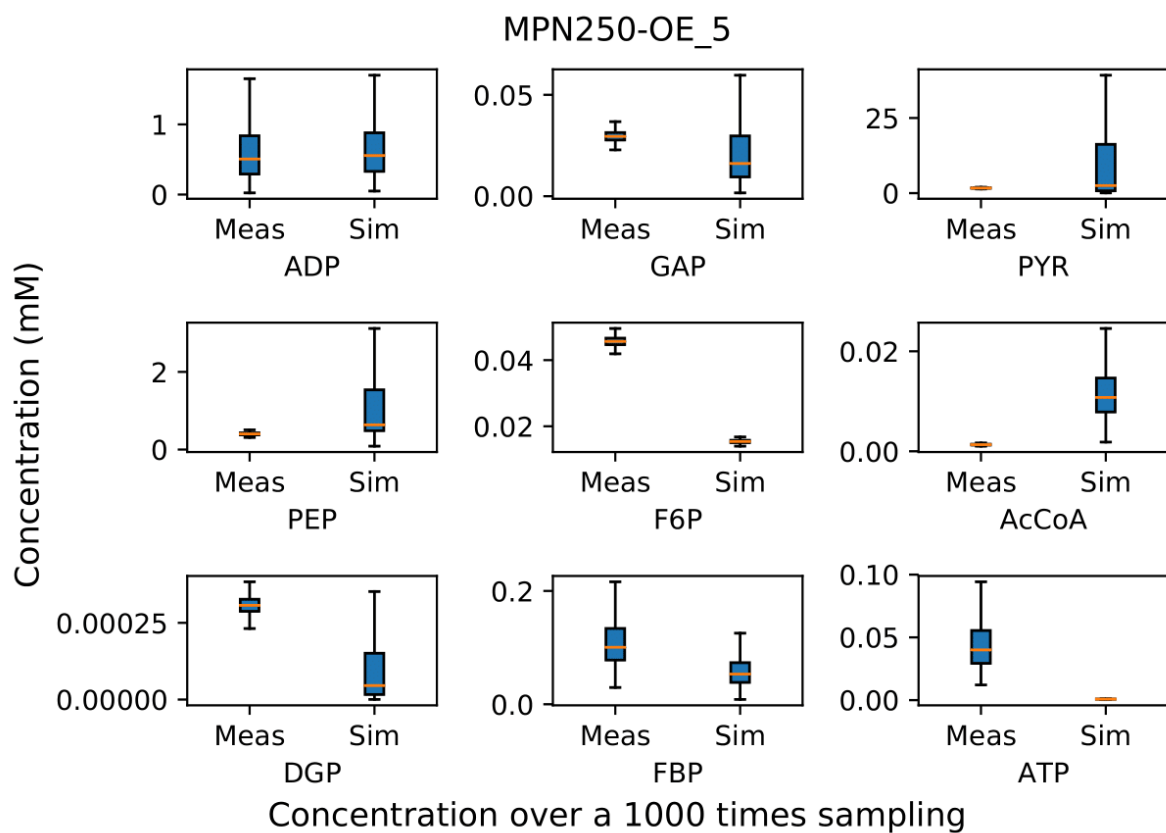
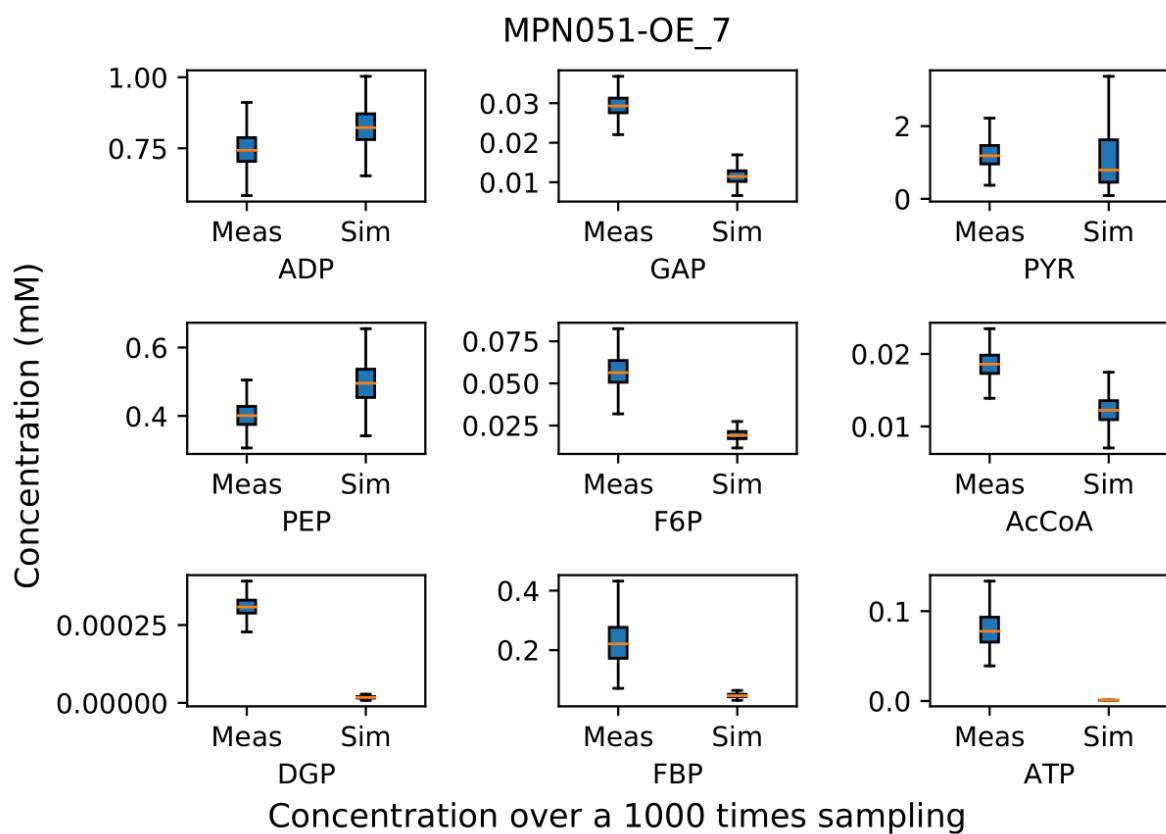
Metabolite concentrations measured and predicted using a 1000x sampling.

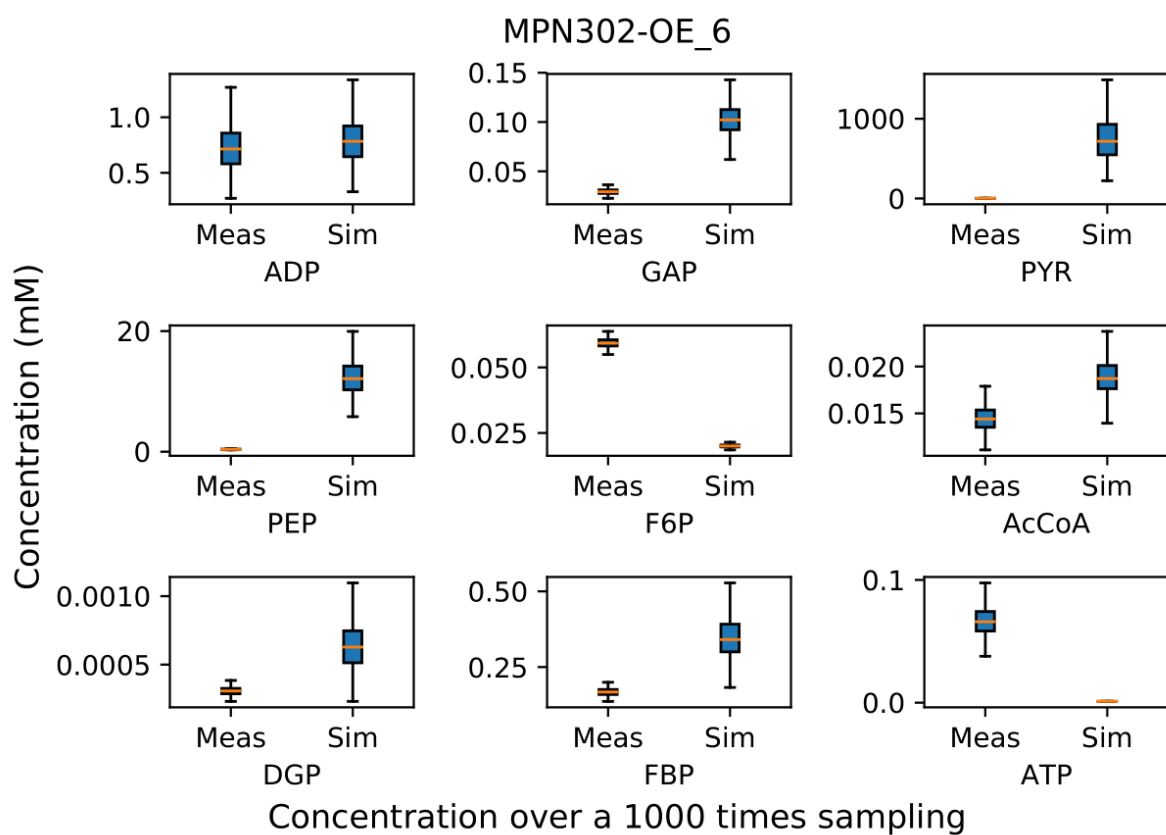
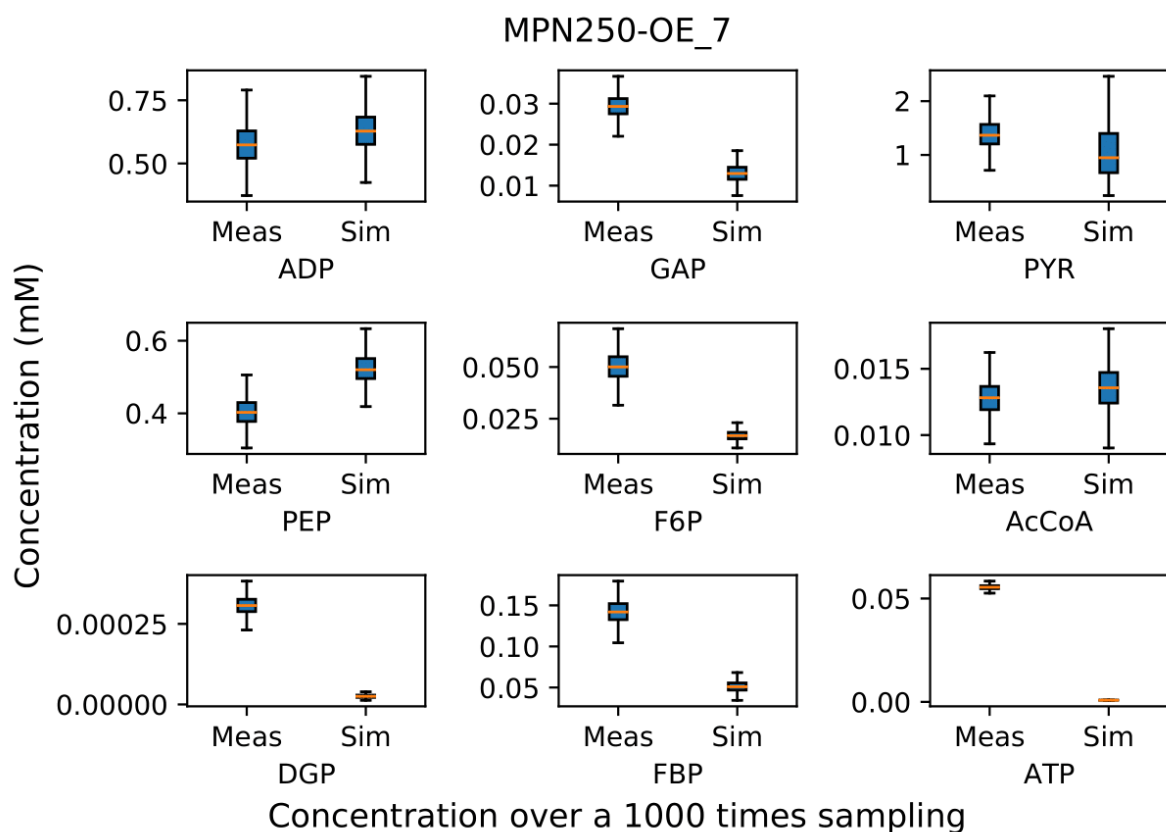


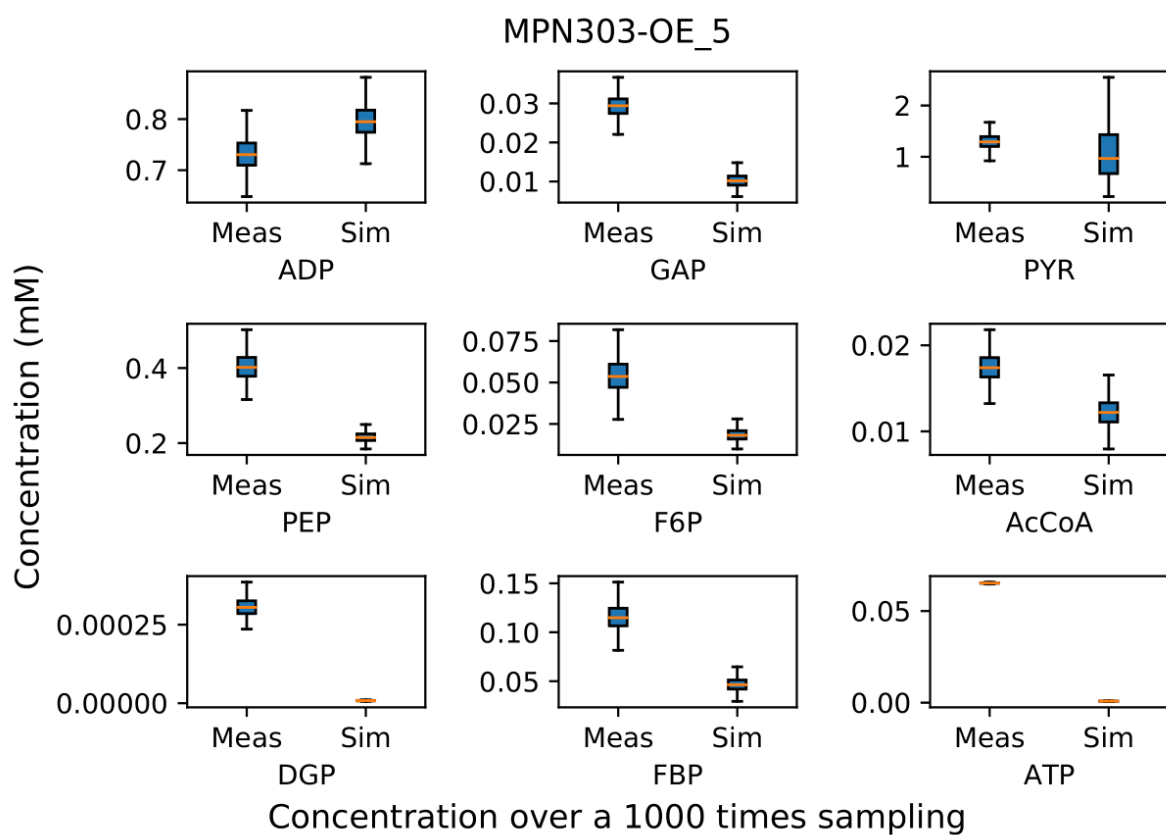
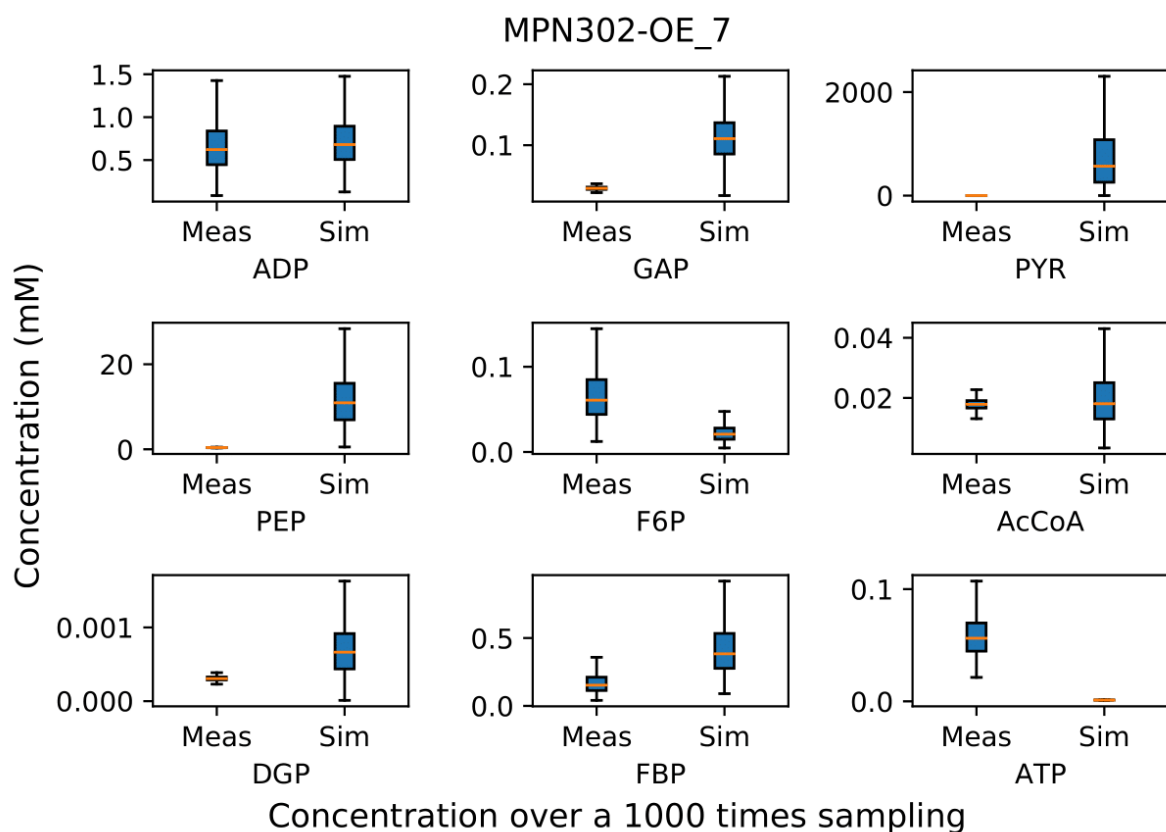


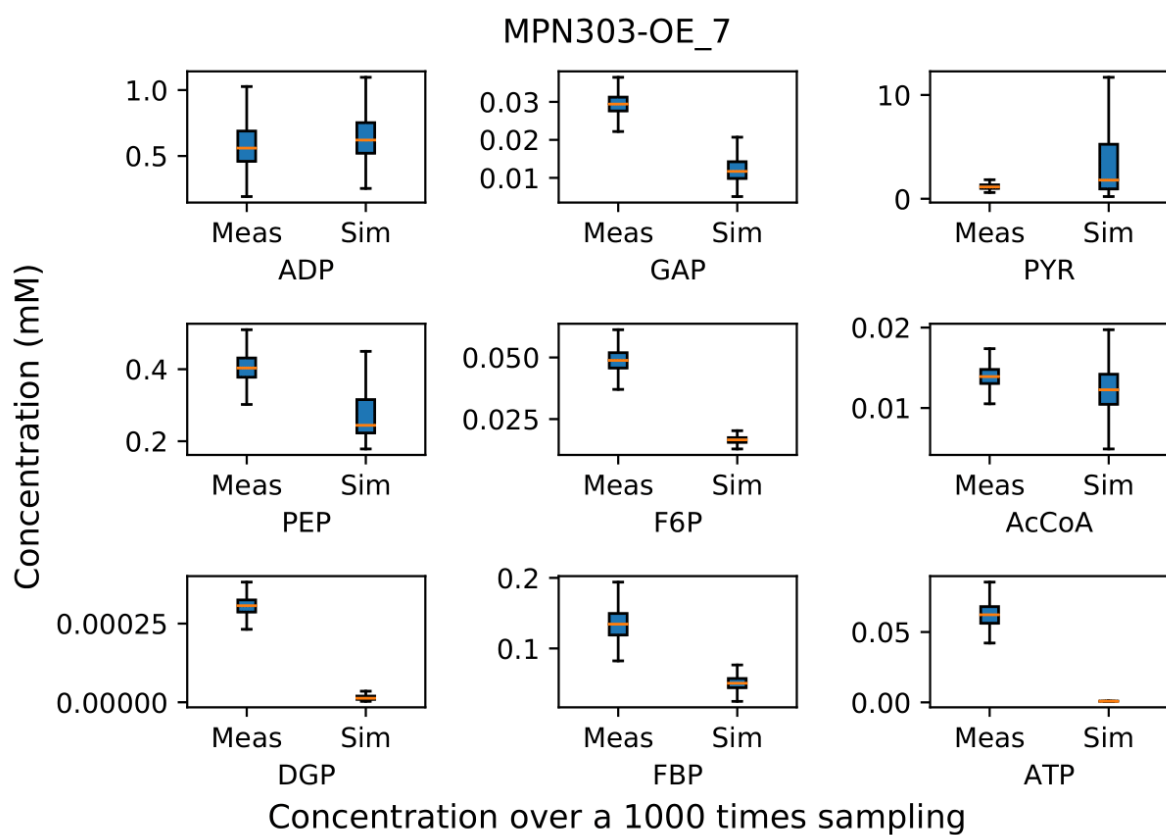
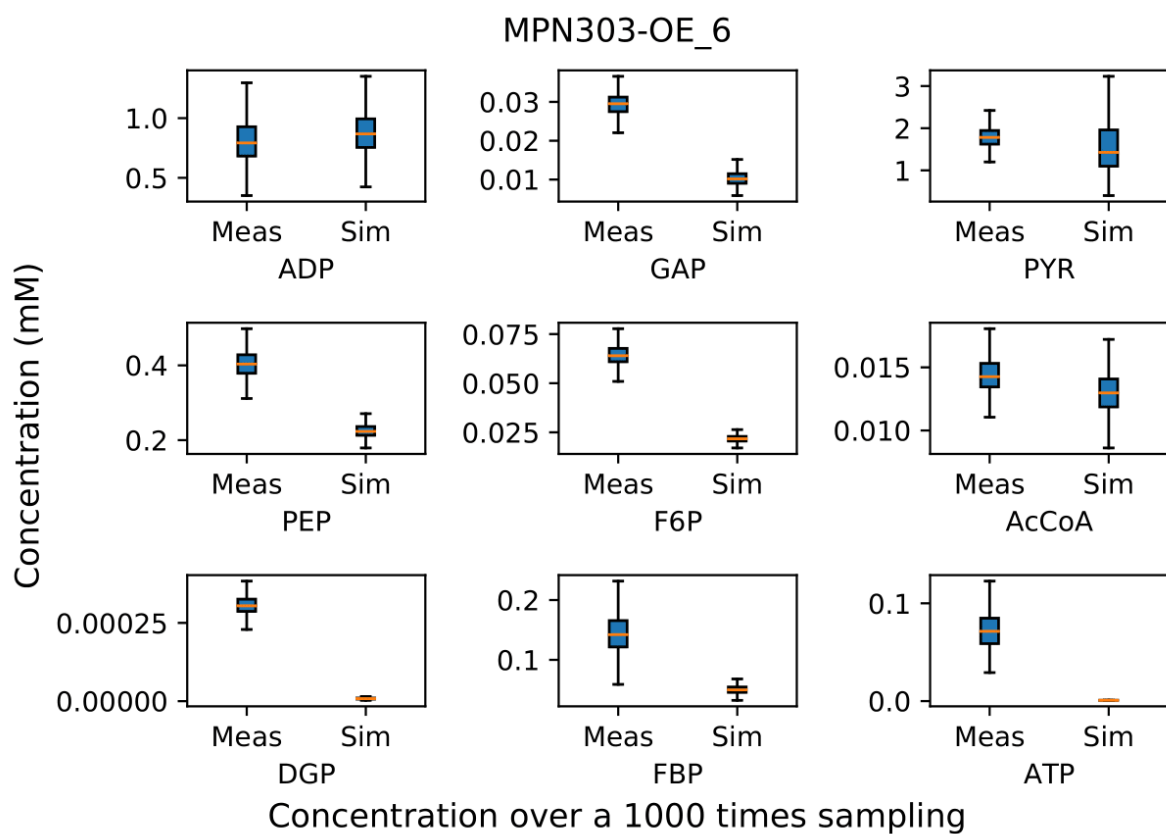


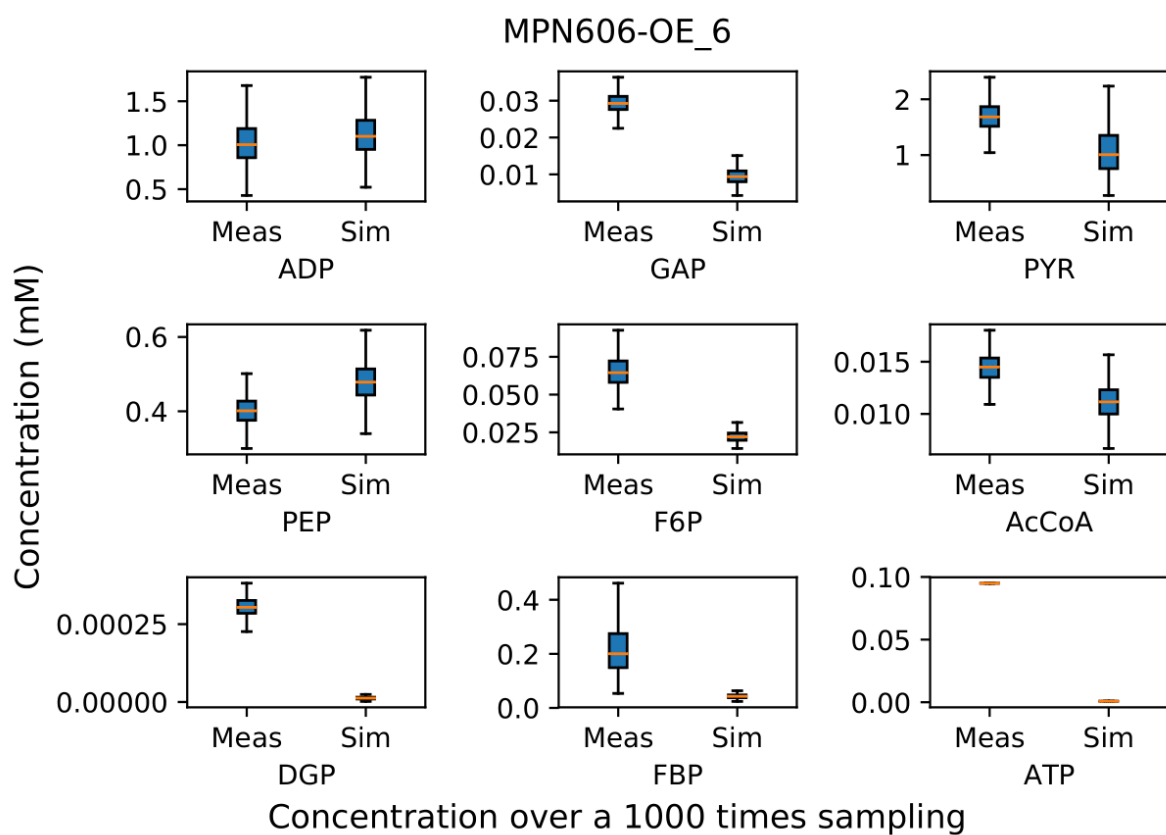
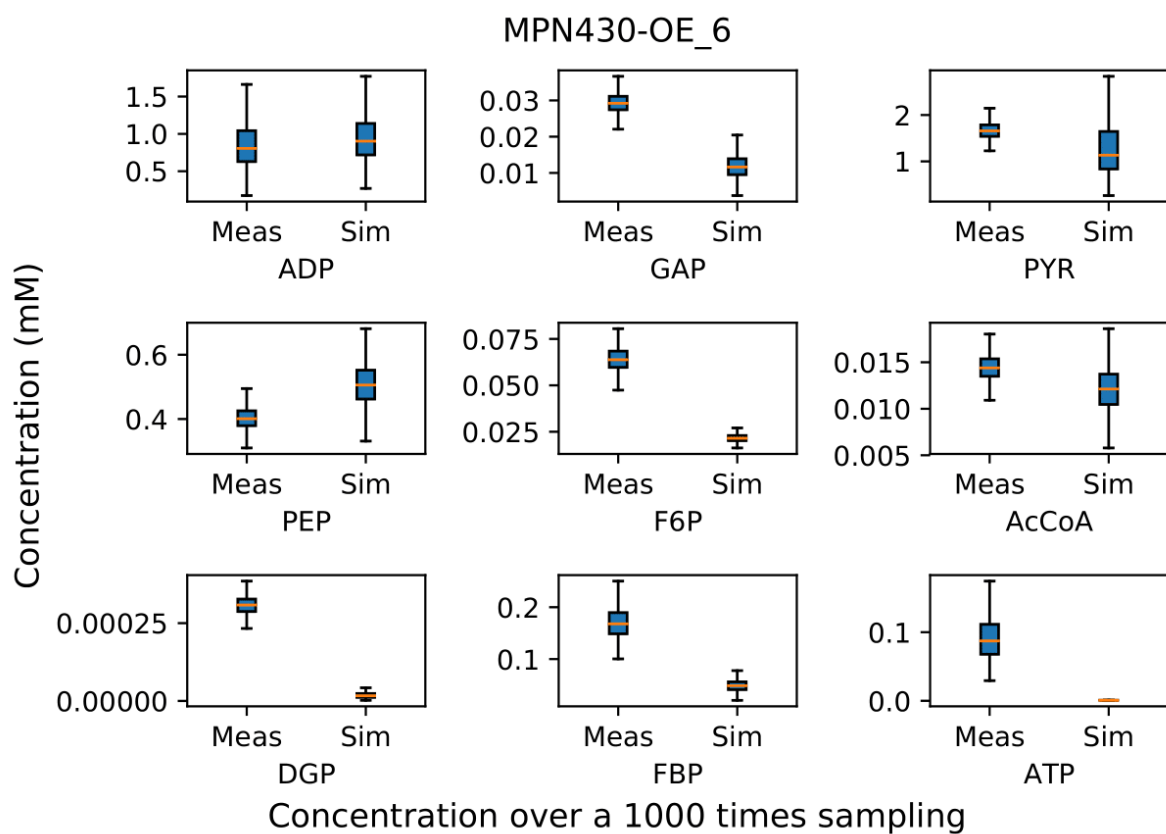


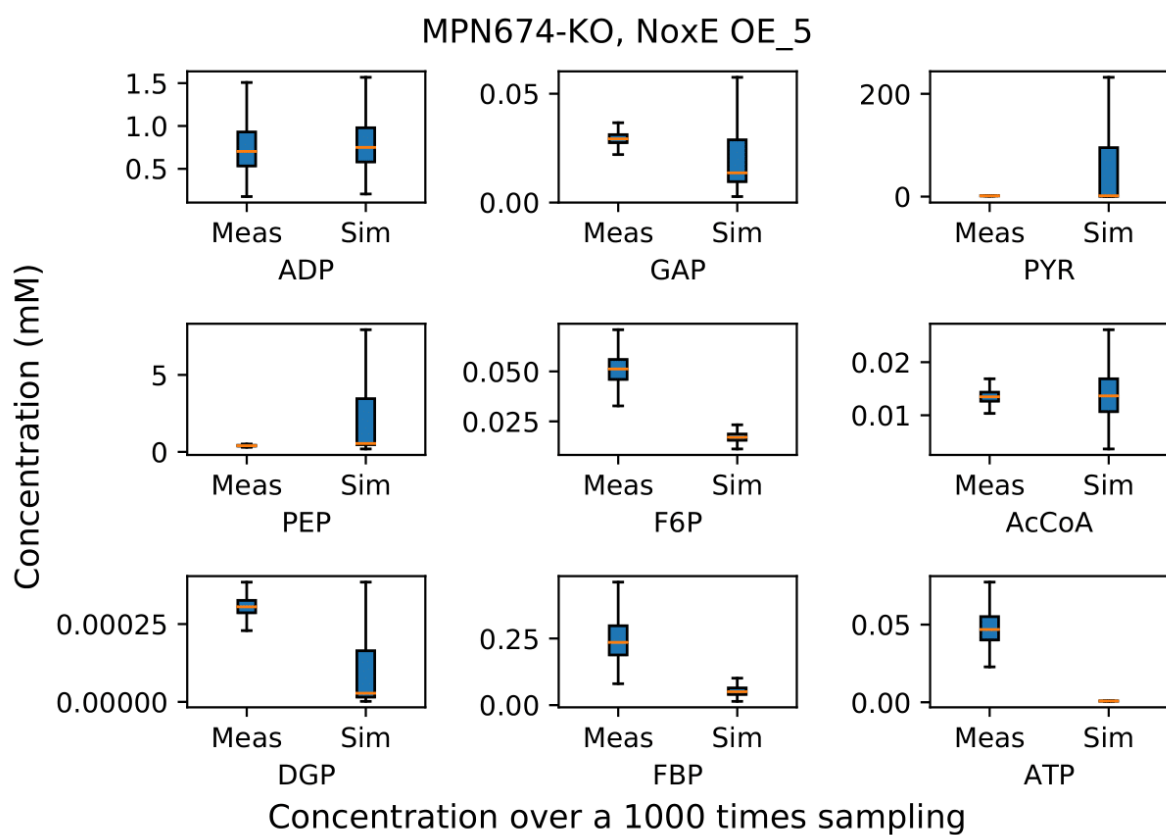
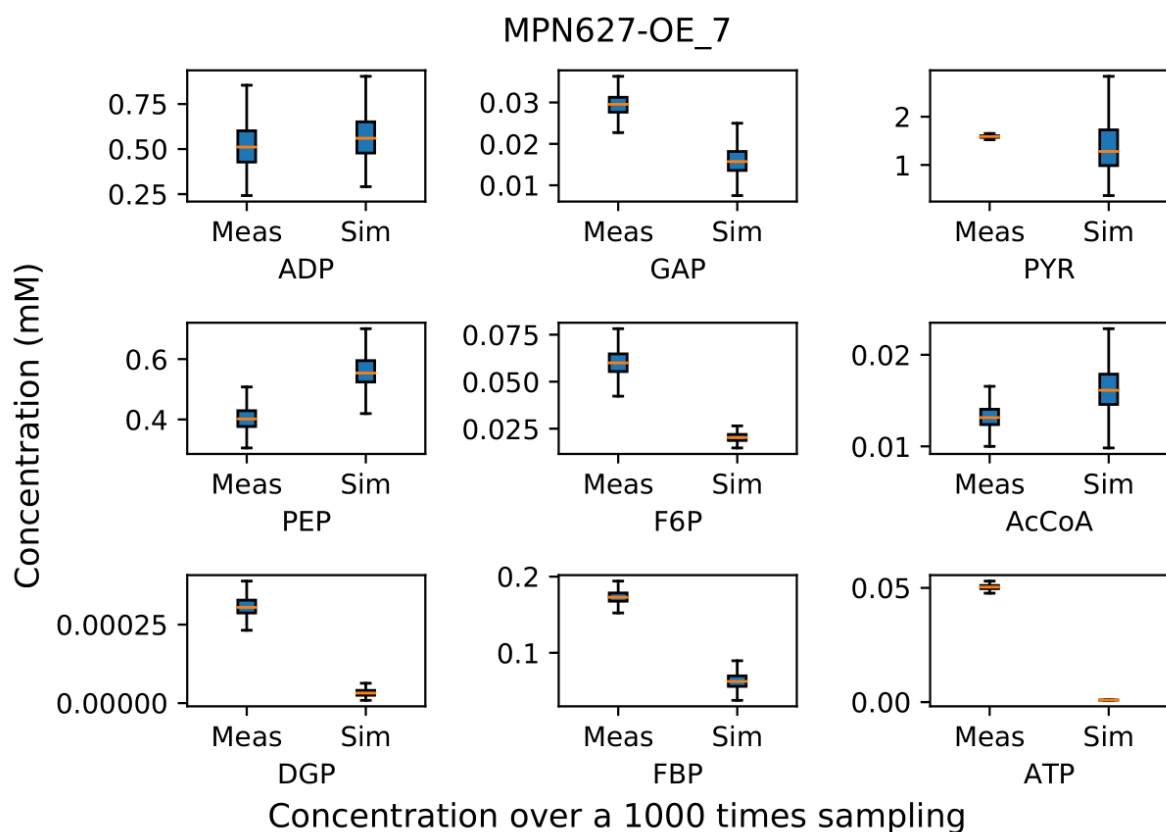




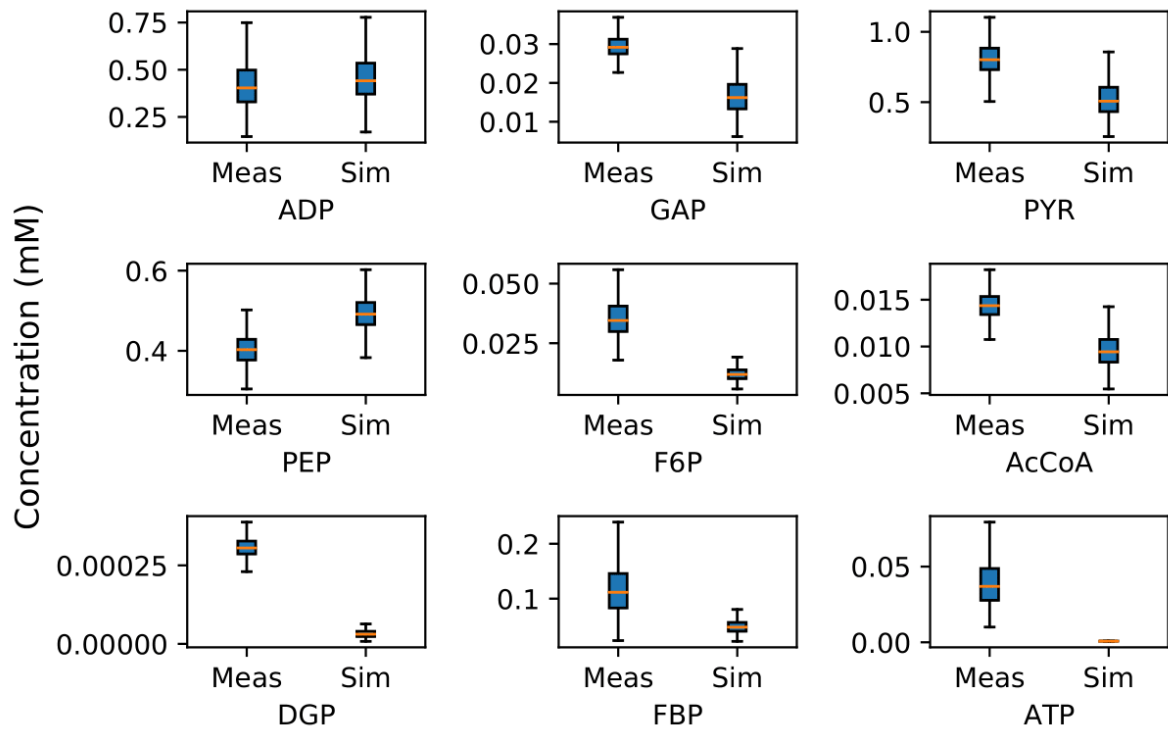




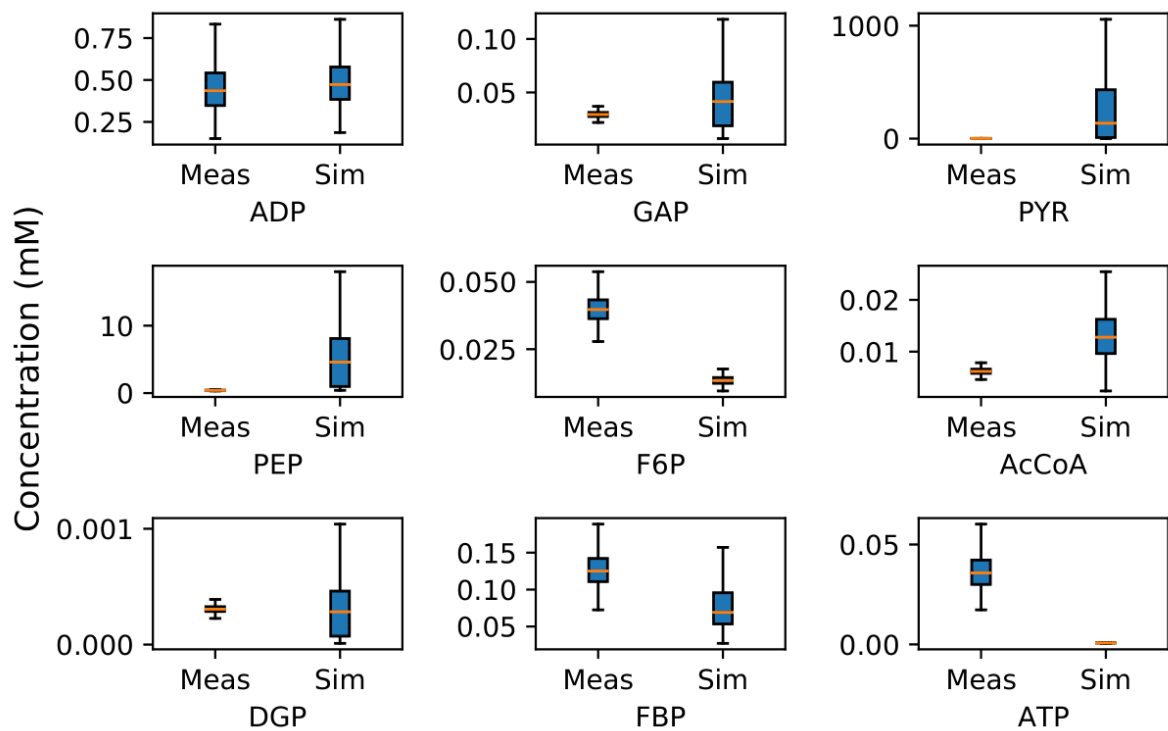


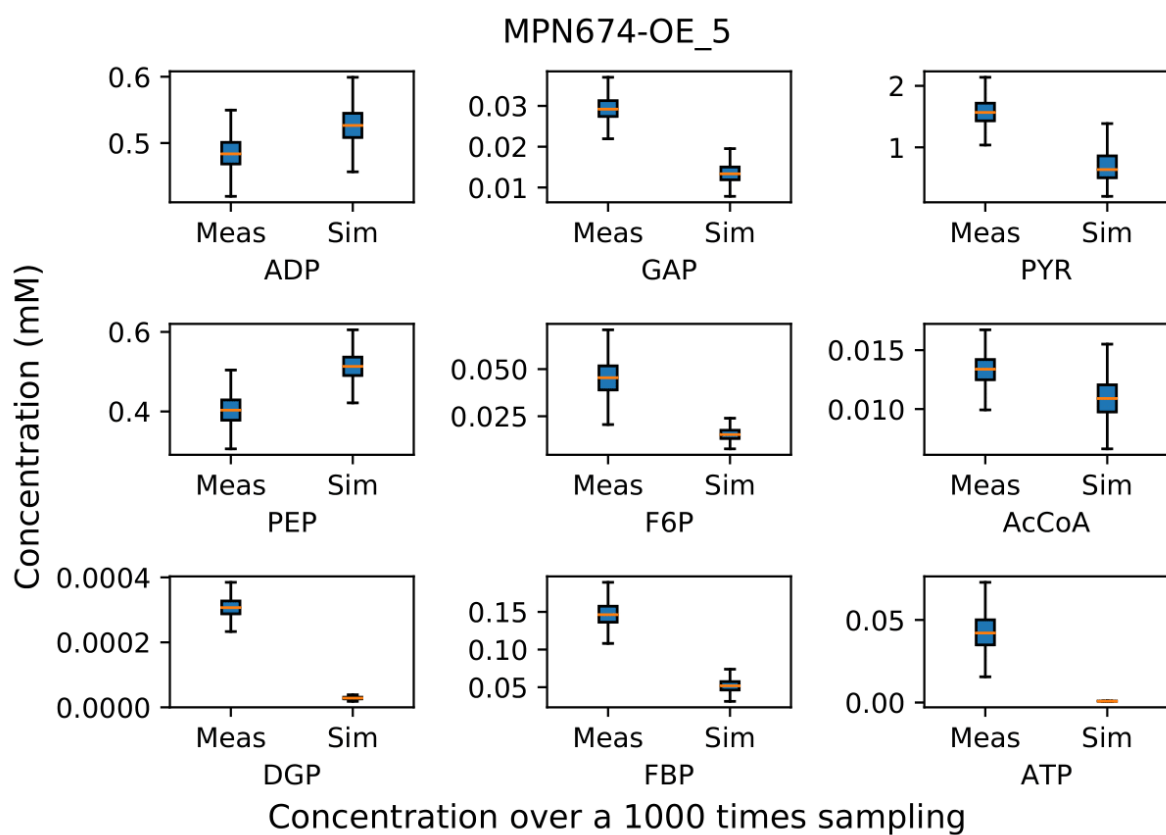
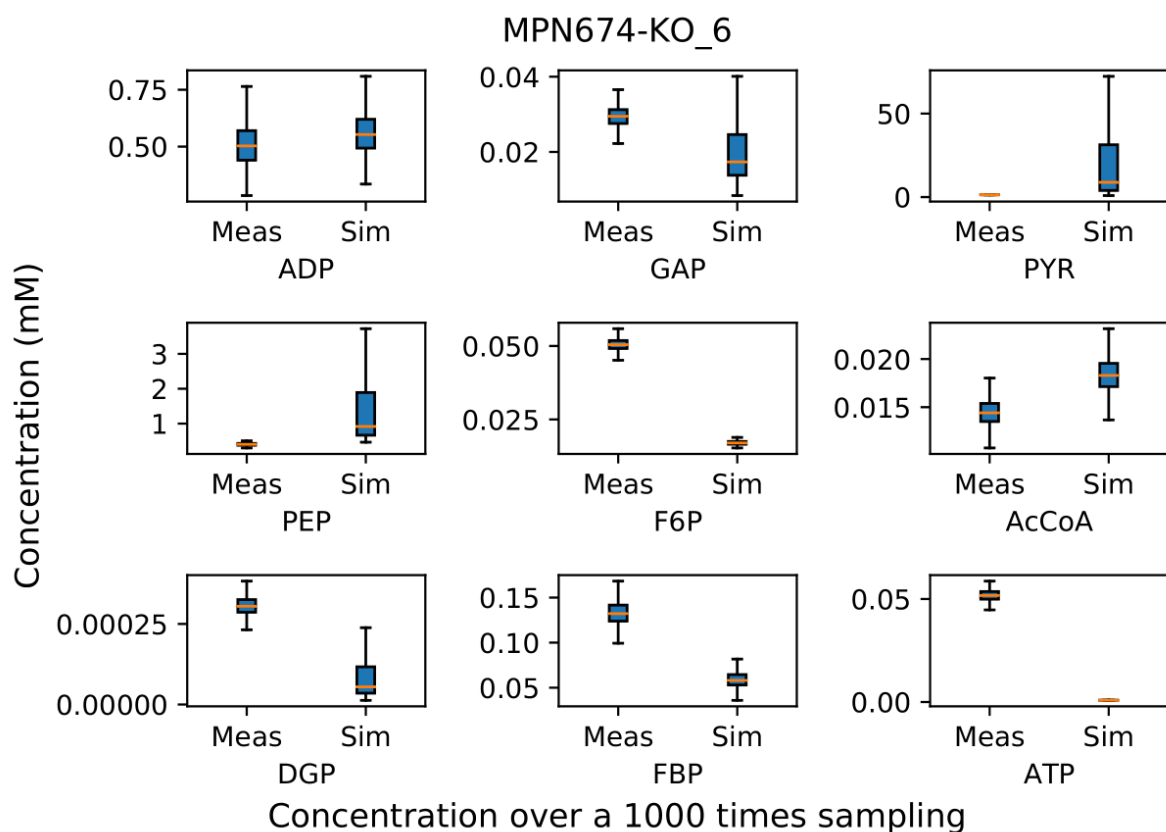


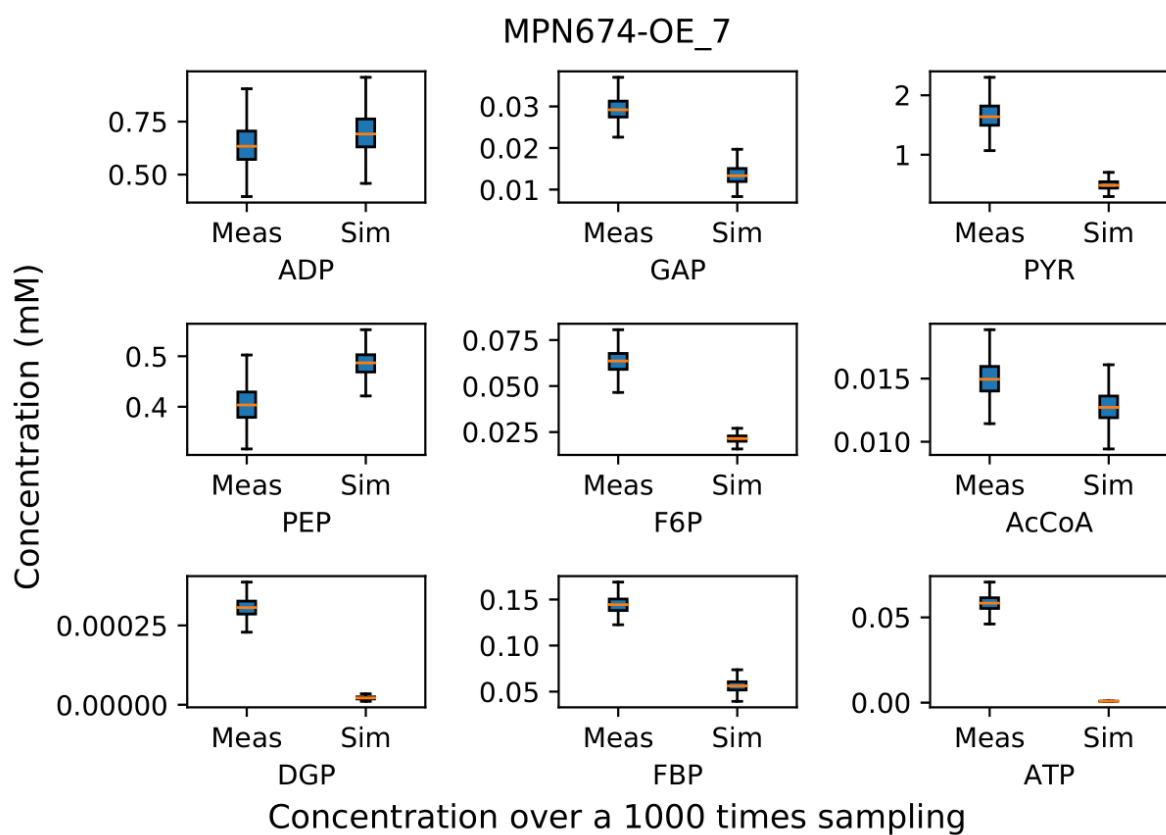
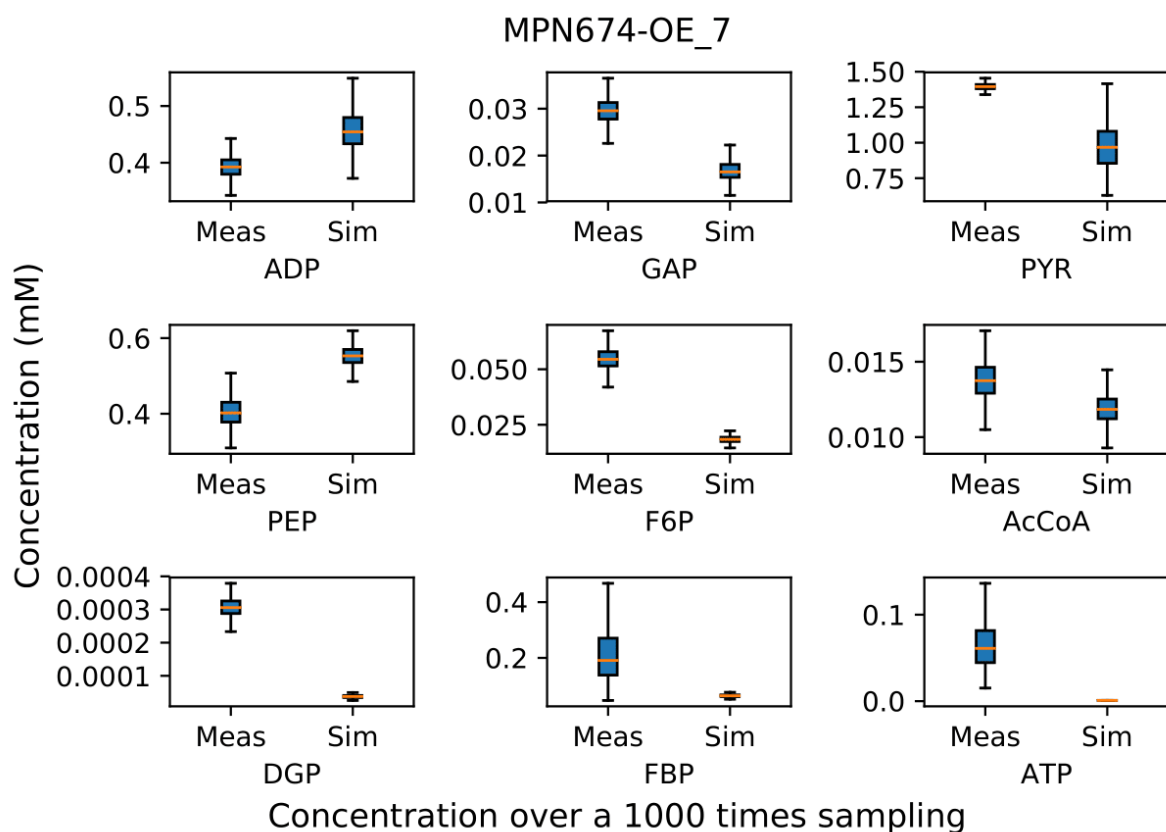
MPN674-KO, NoxE OE_6



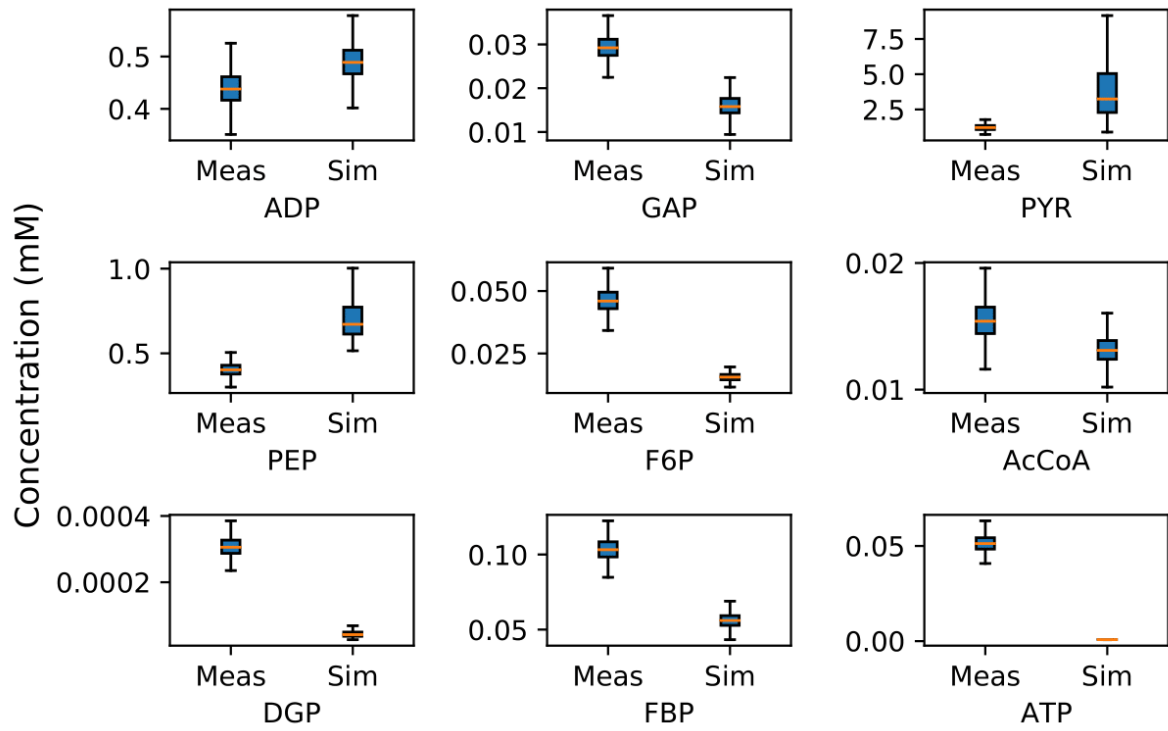
MPN674-KO_5





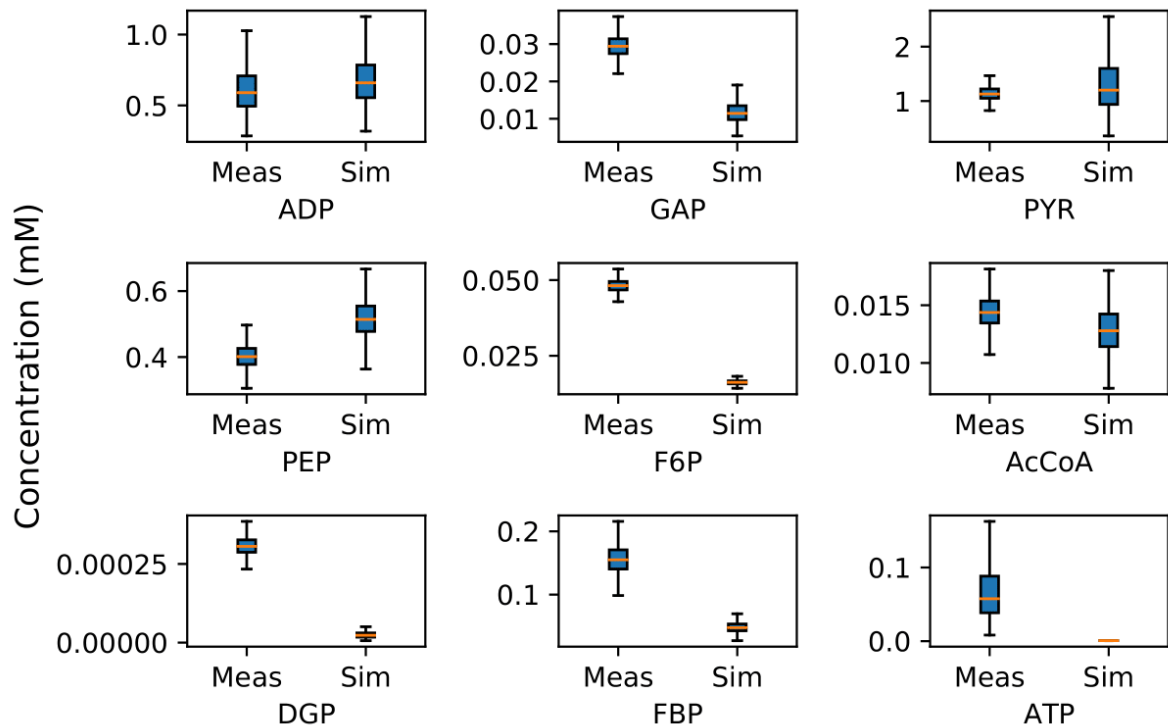


Tn051_Gly_perturbation_7

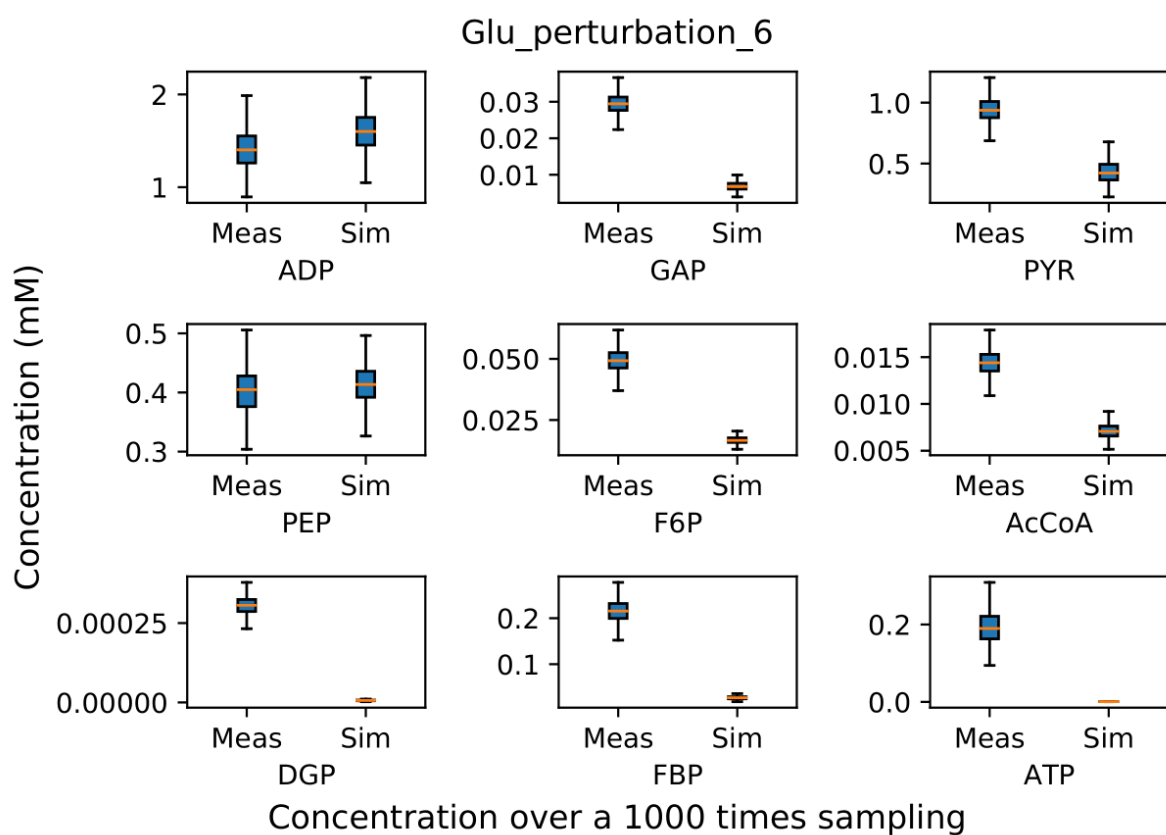
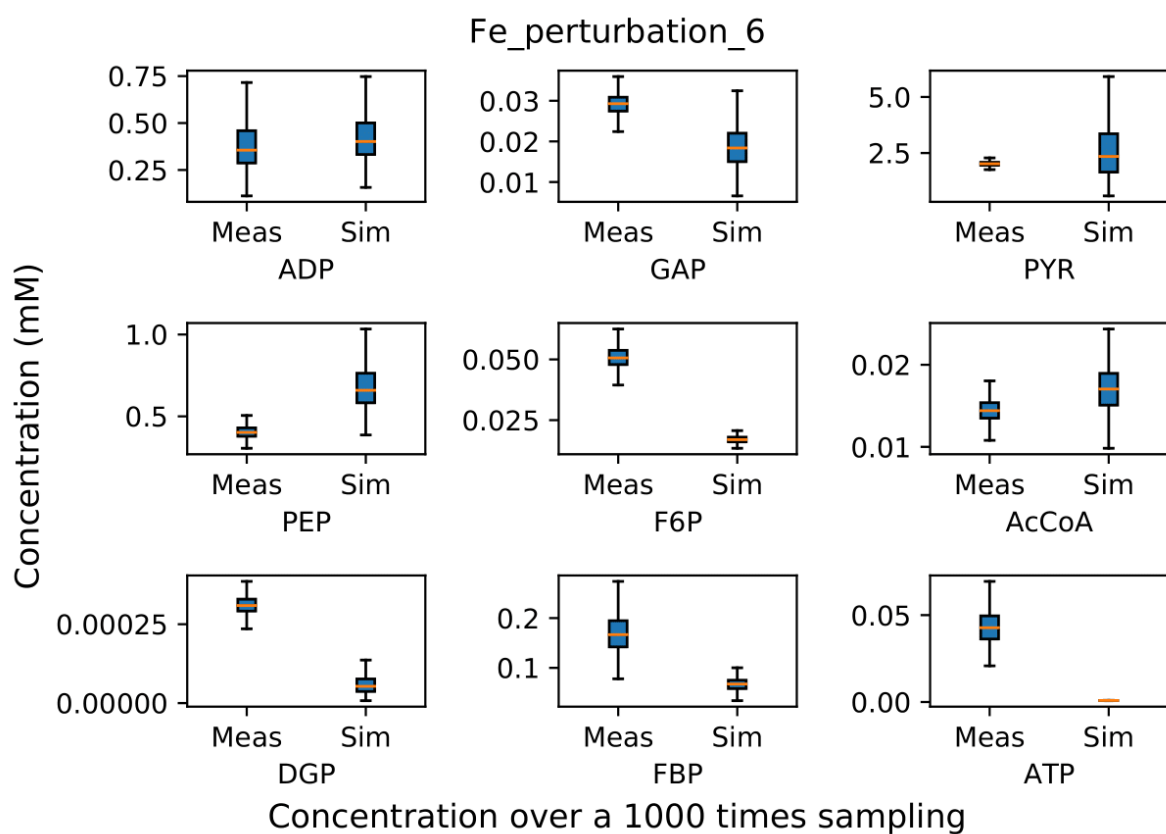


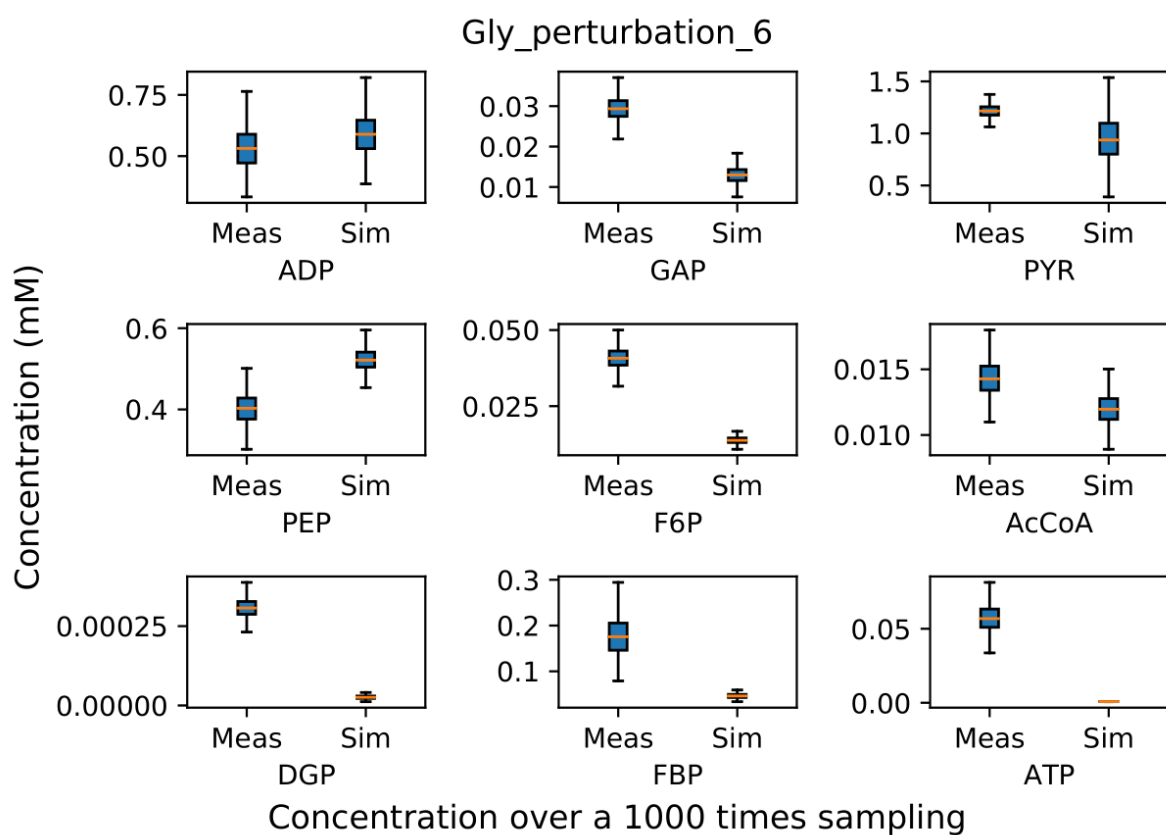
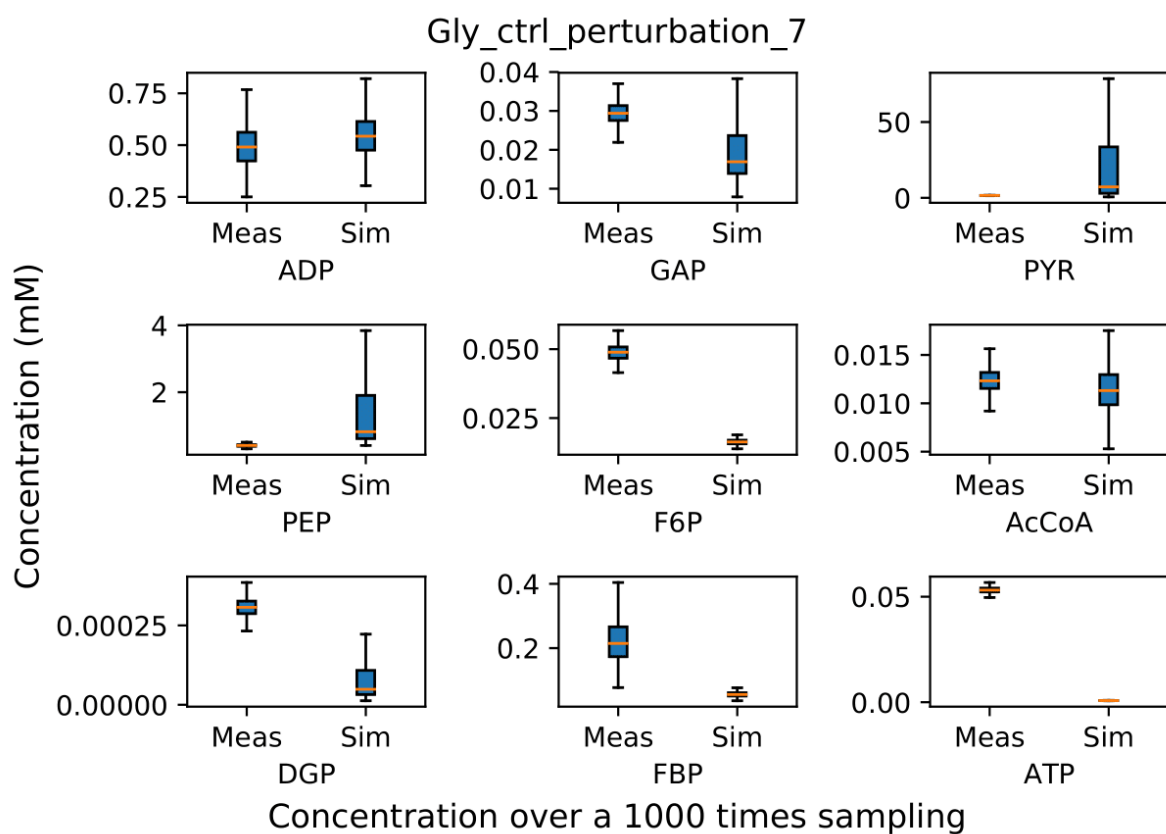
Concentration over a 1000 times sampling

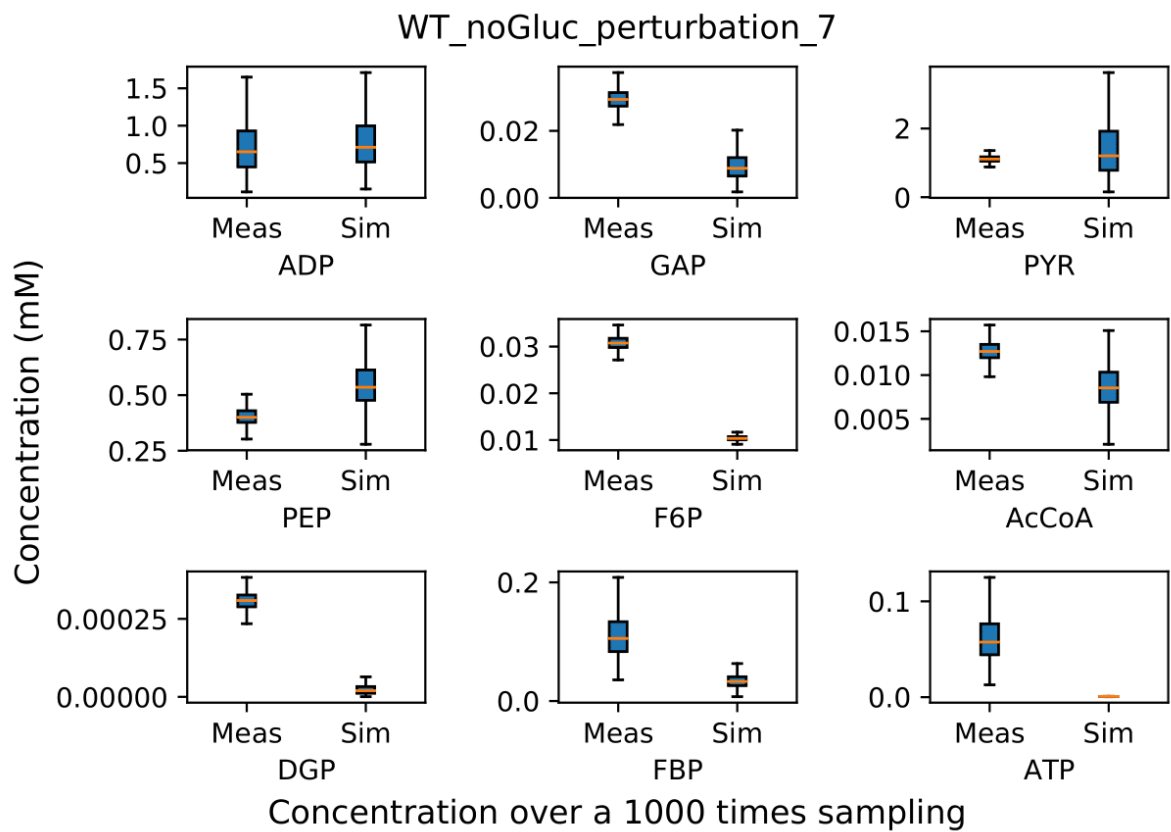
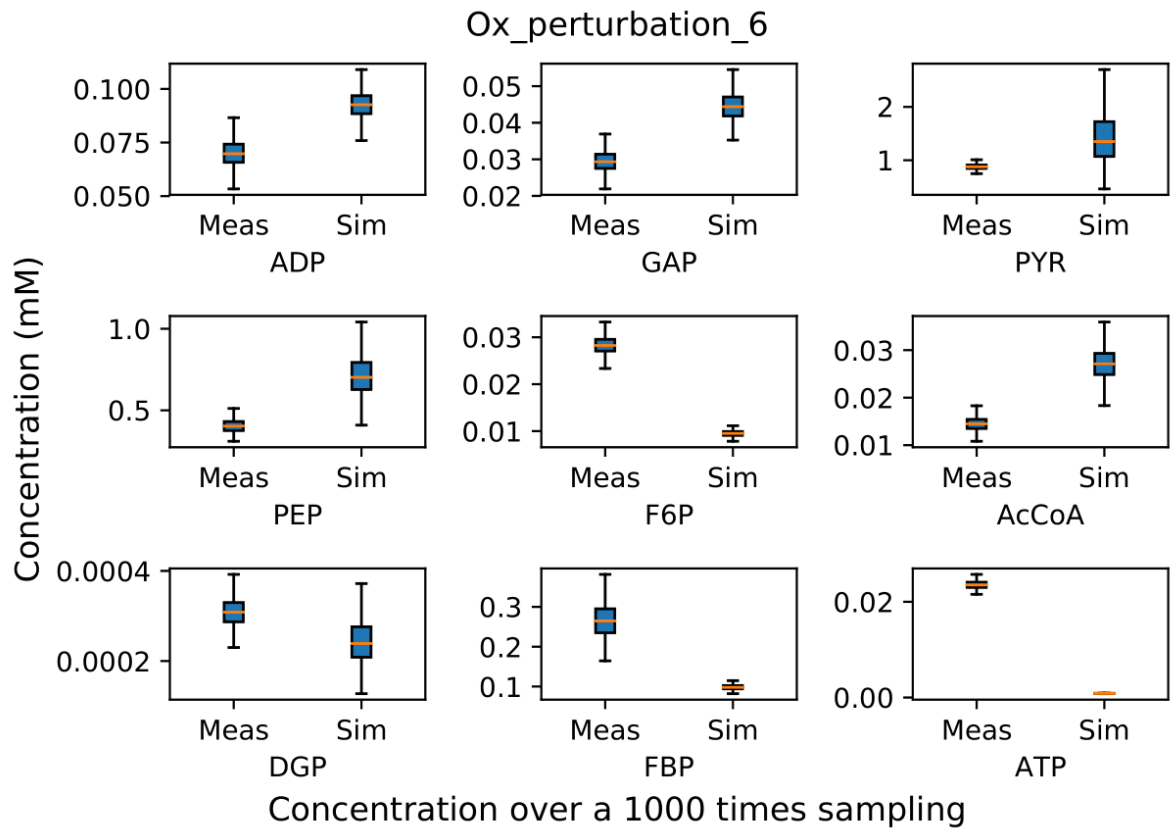
AA_perturbation_6

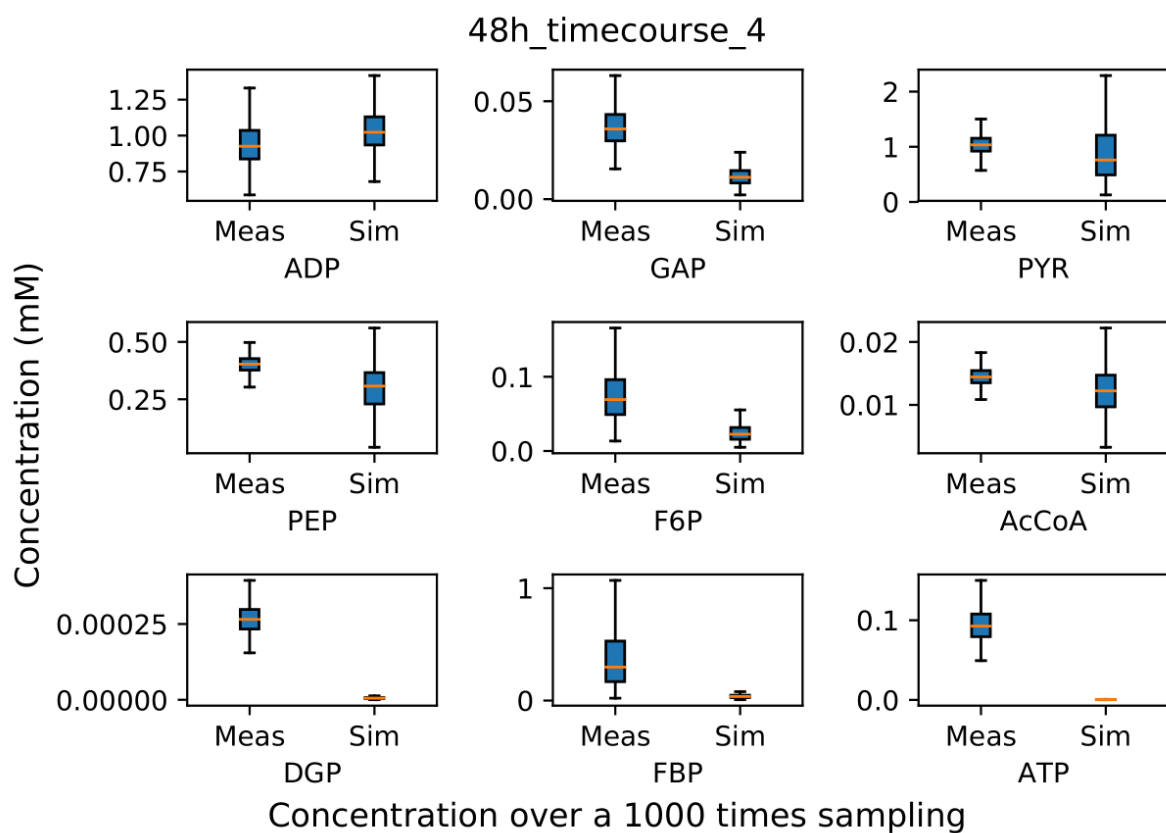
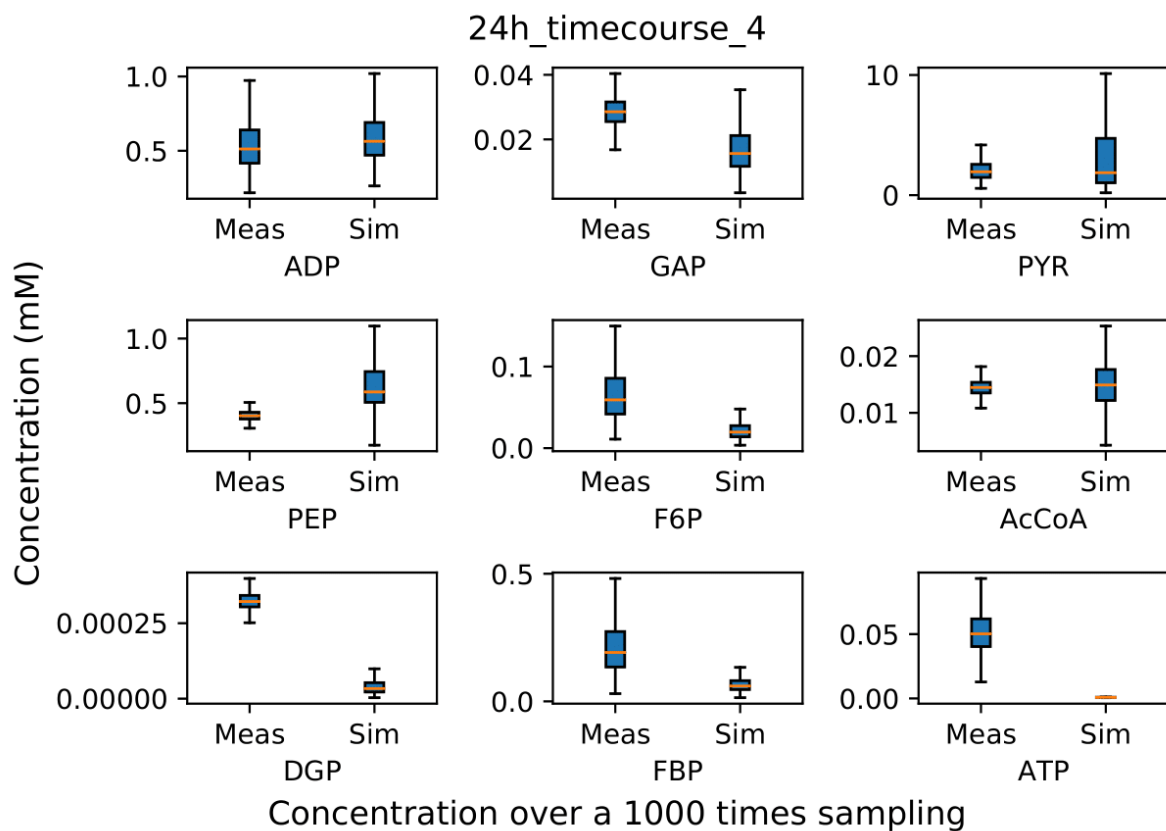


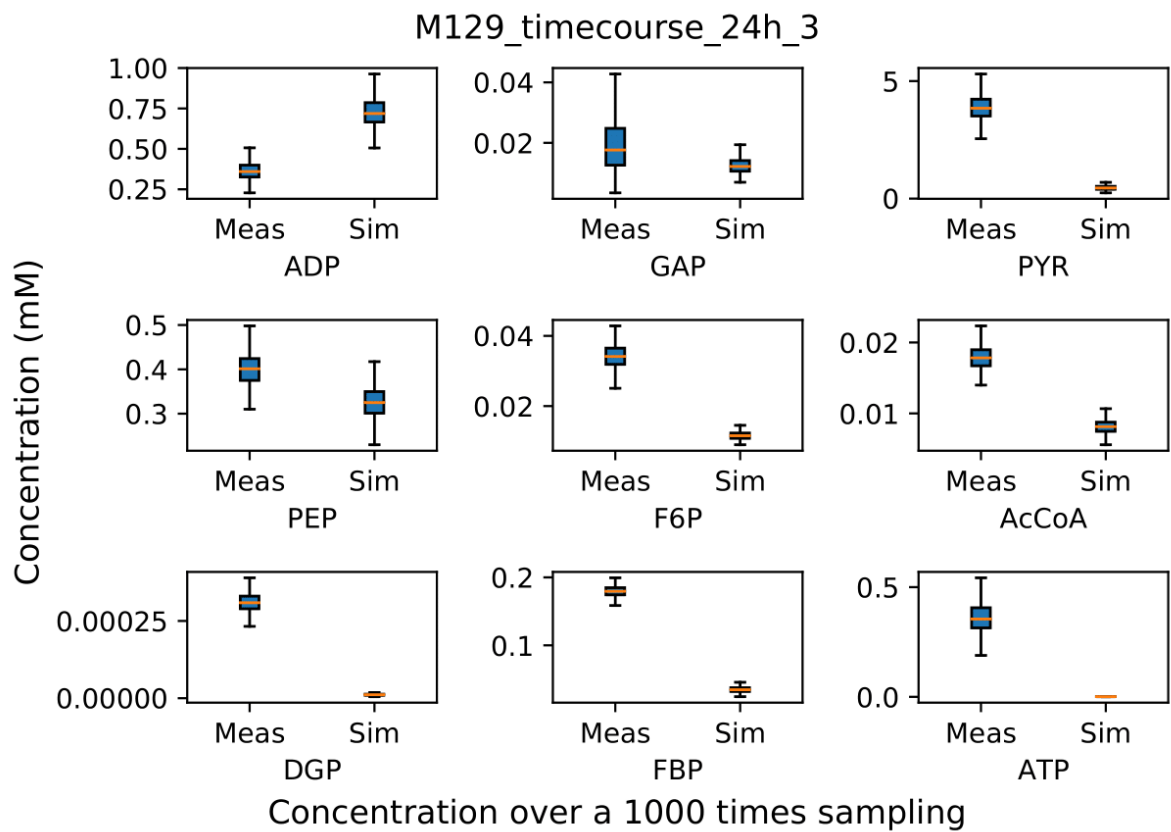
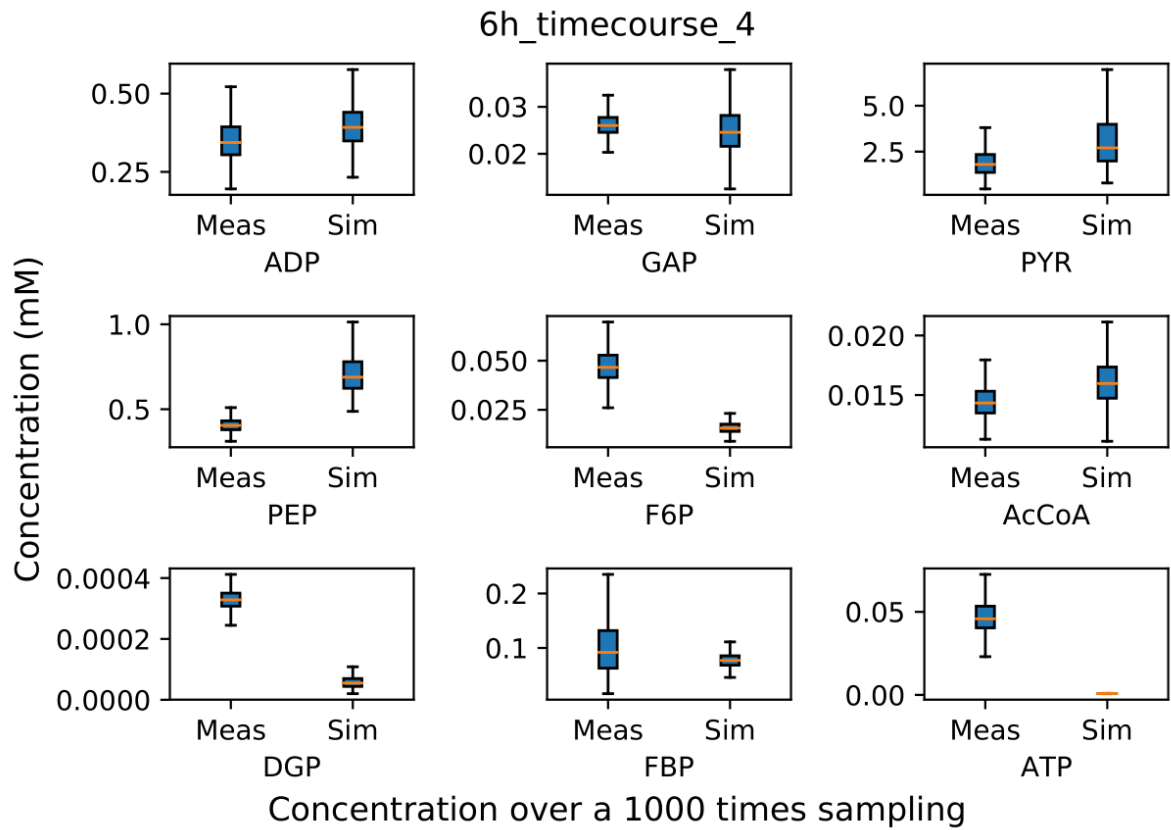
Concentration over a 1000 times sampling



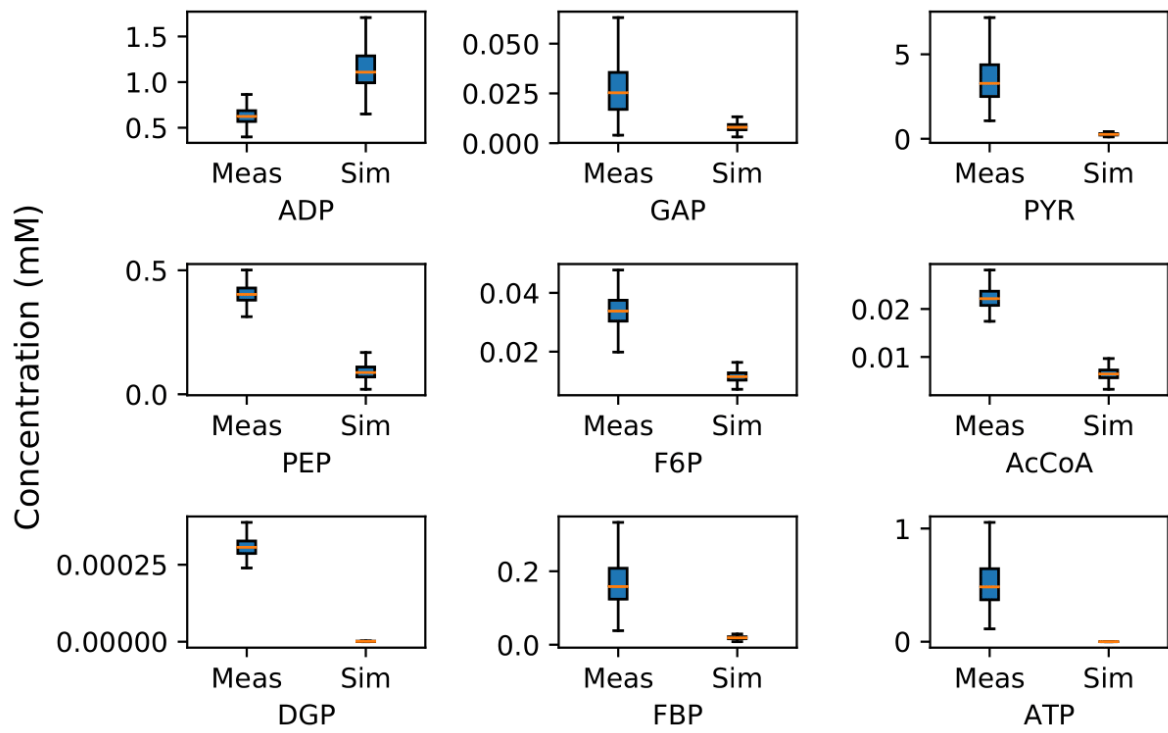






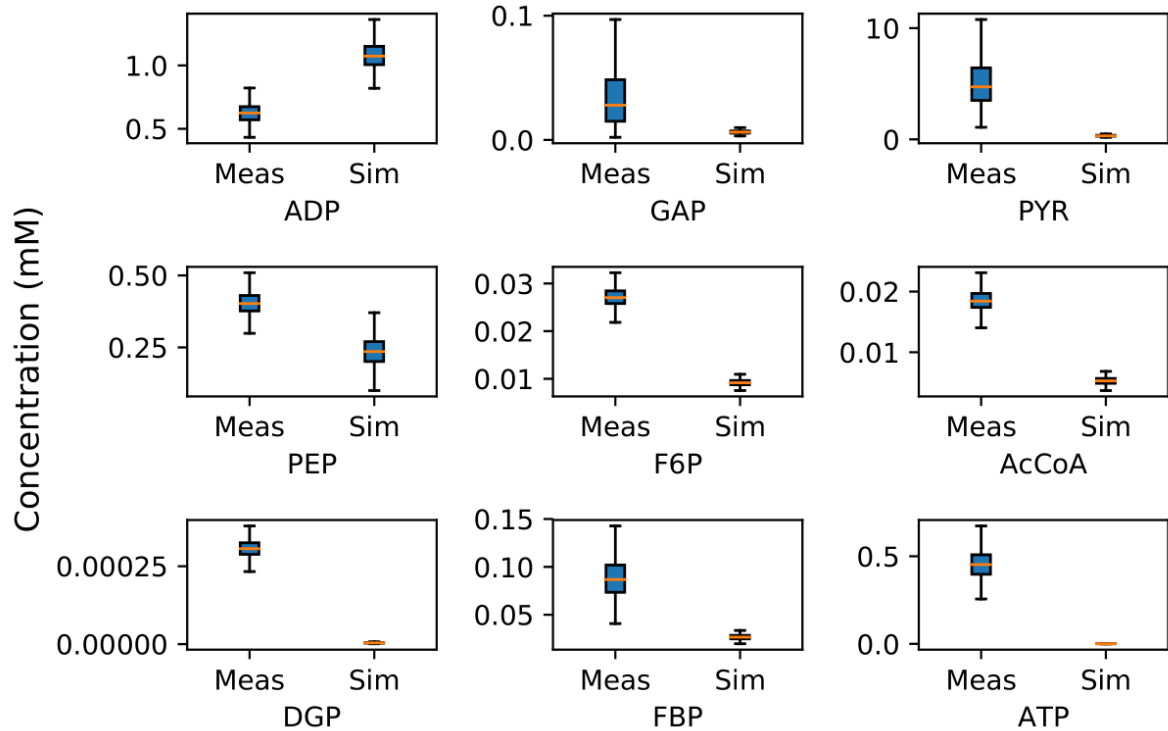


M129_timecourse_48h_3



Concentration over a 1000 times sampling

M129_timecourse_96h_3



Concentration over a 1000 times sampling

Supplementary file 1 D Metabolomics analysis.

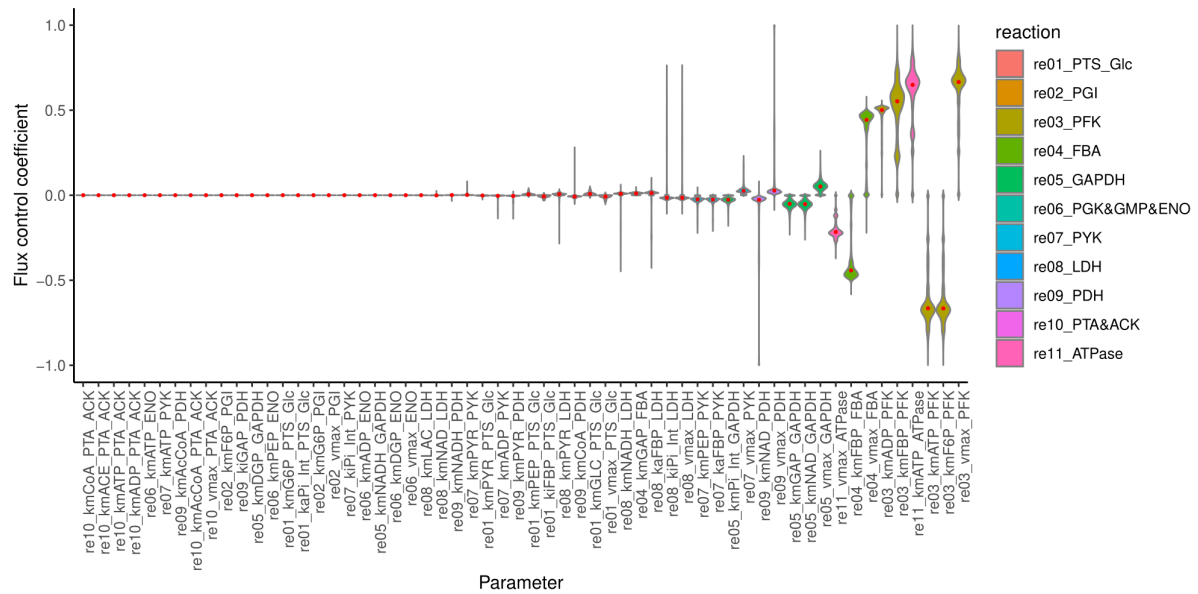


Figure 3 Local sensitivity analysis using 1000x sampling from the measurement distribution.

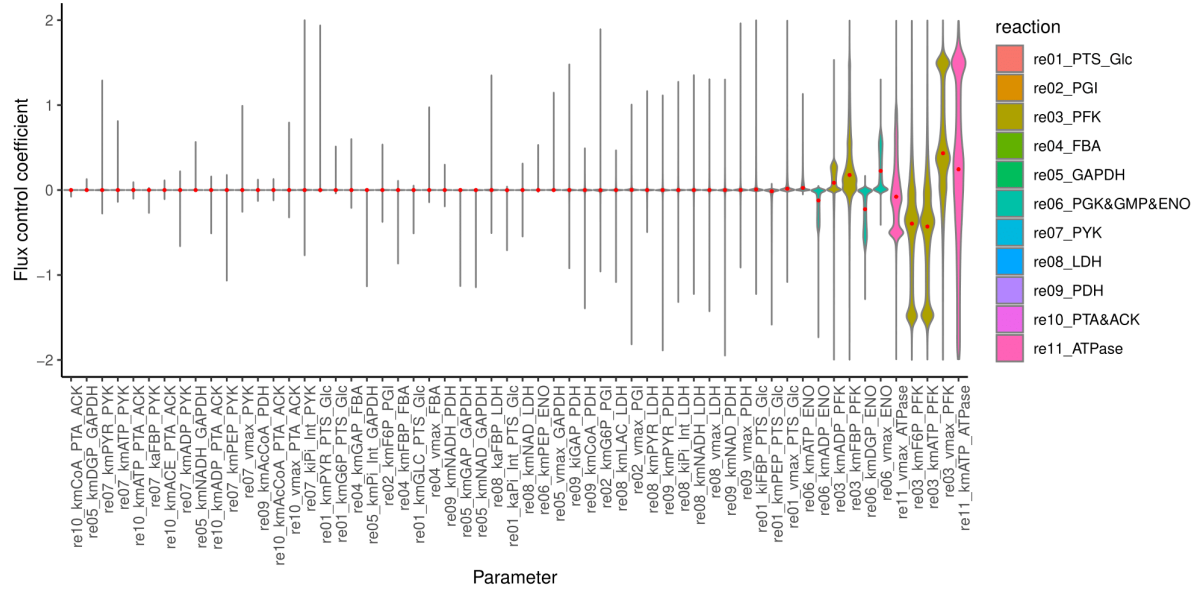


Figure 4 Global sensitivity analysis using a 100,000 Latin Hypercube sampling with the parameter search range.

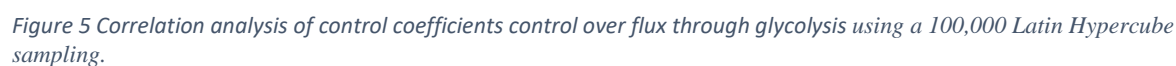


Figure 5 Correlation analysis of control coefficients control over flux through glycolysis using a 100,000 Latin Hypercube sampling.