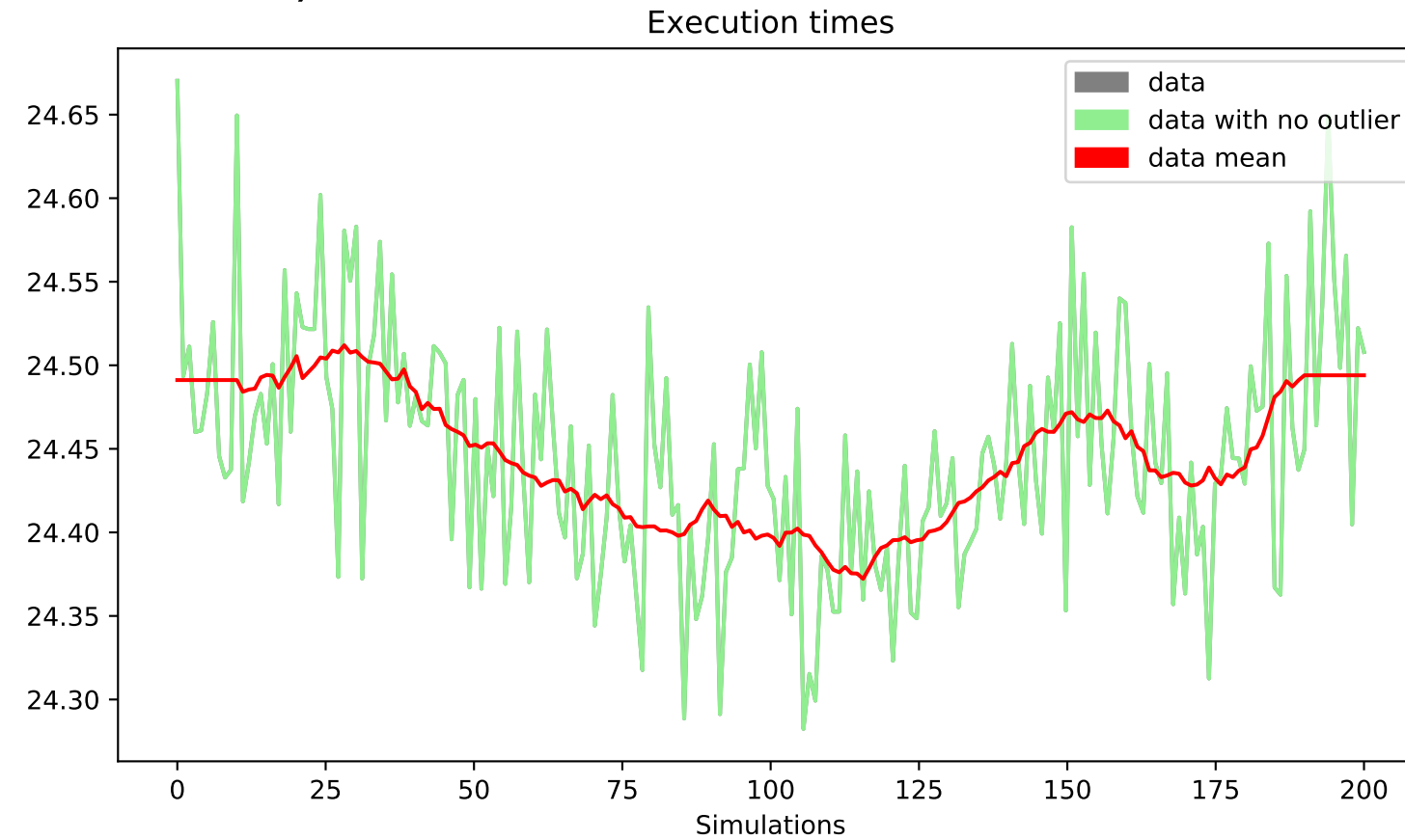
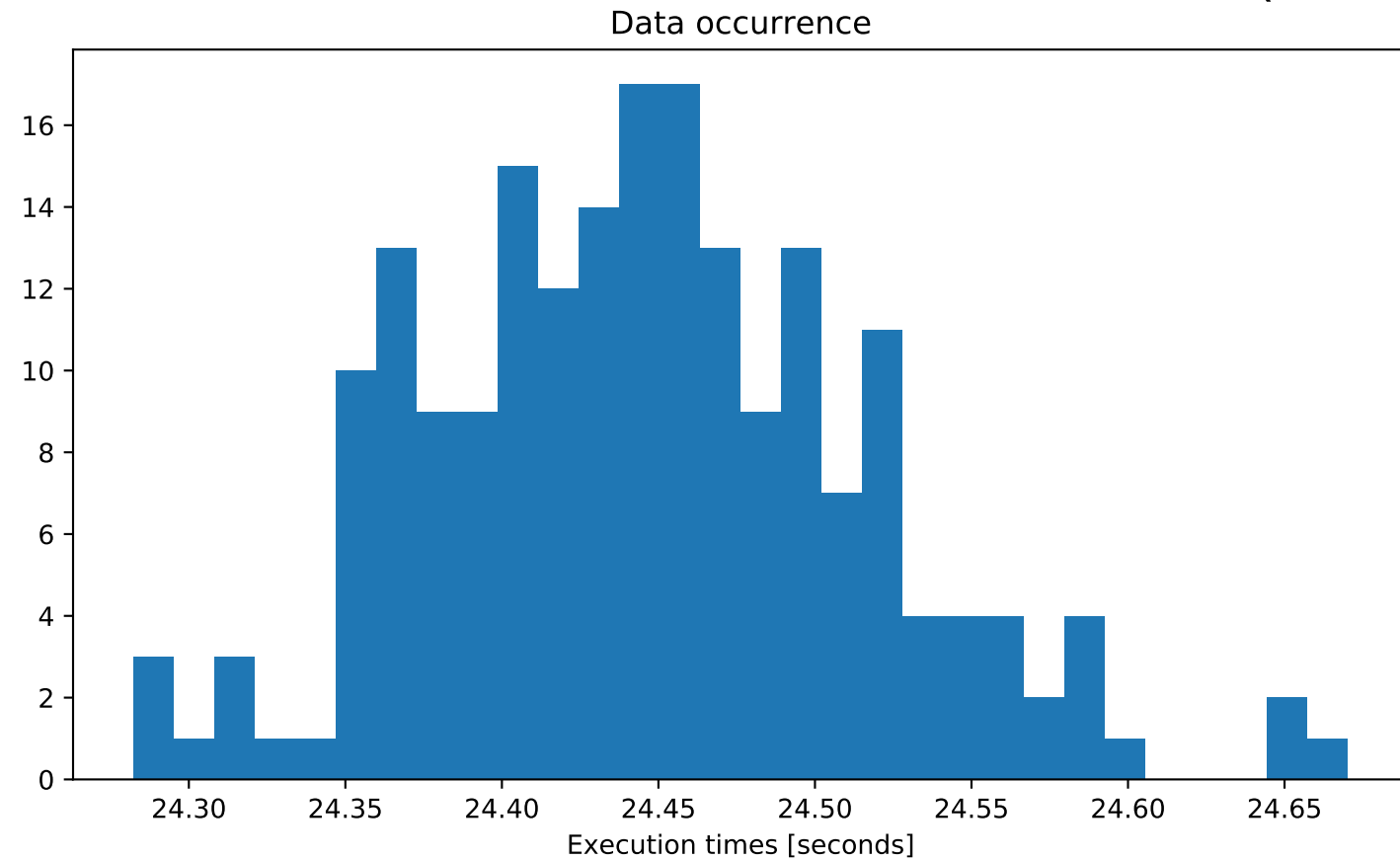
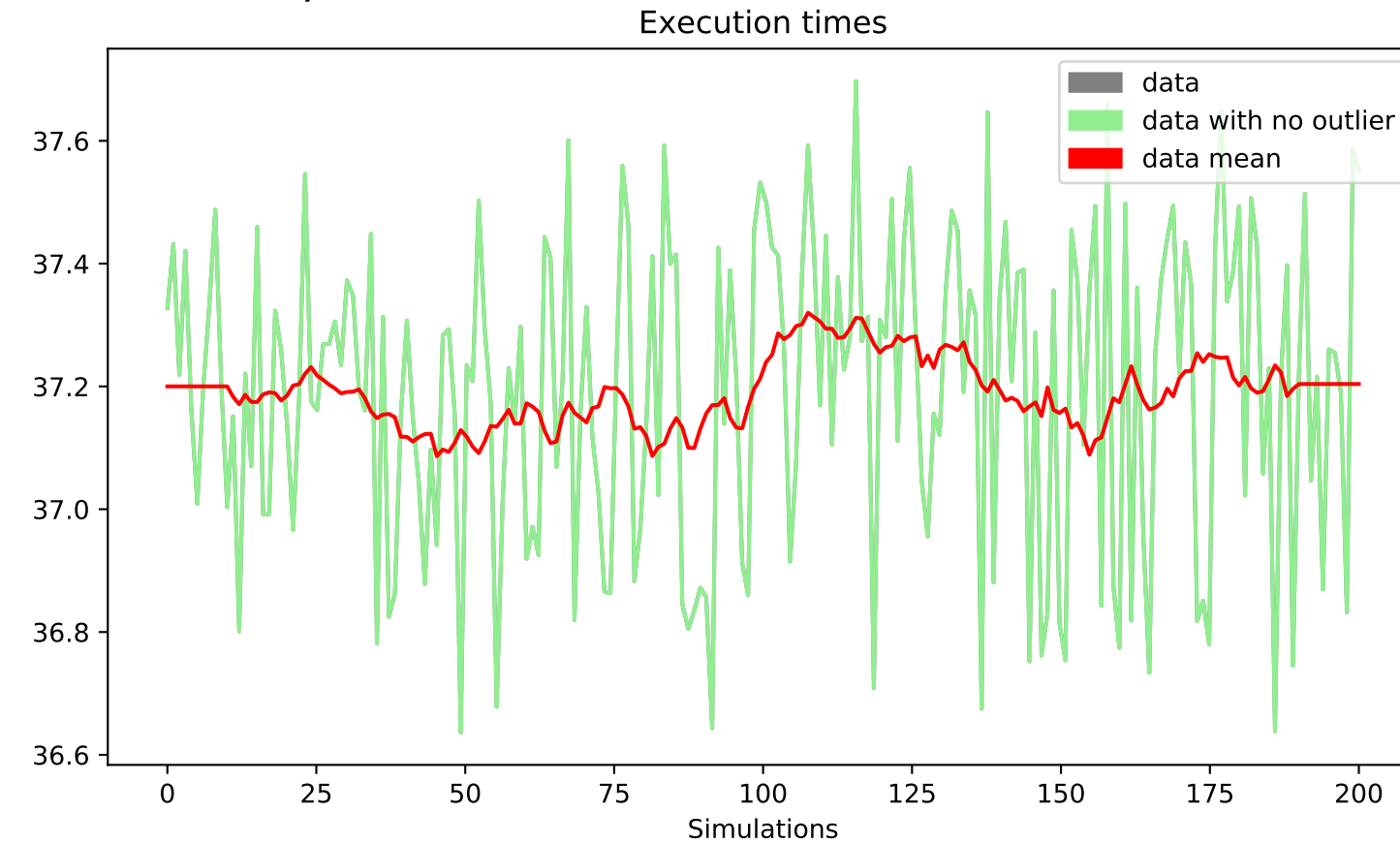
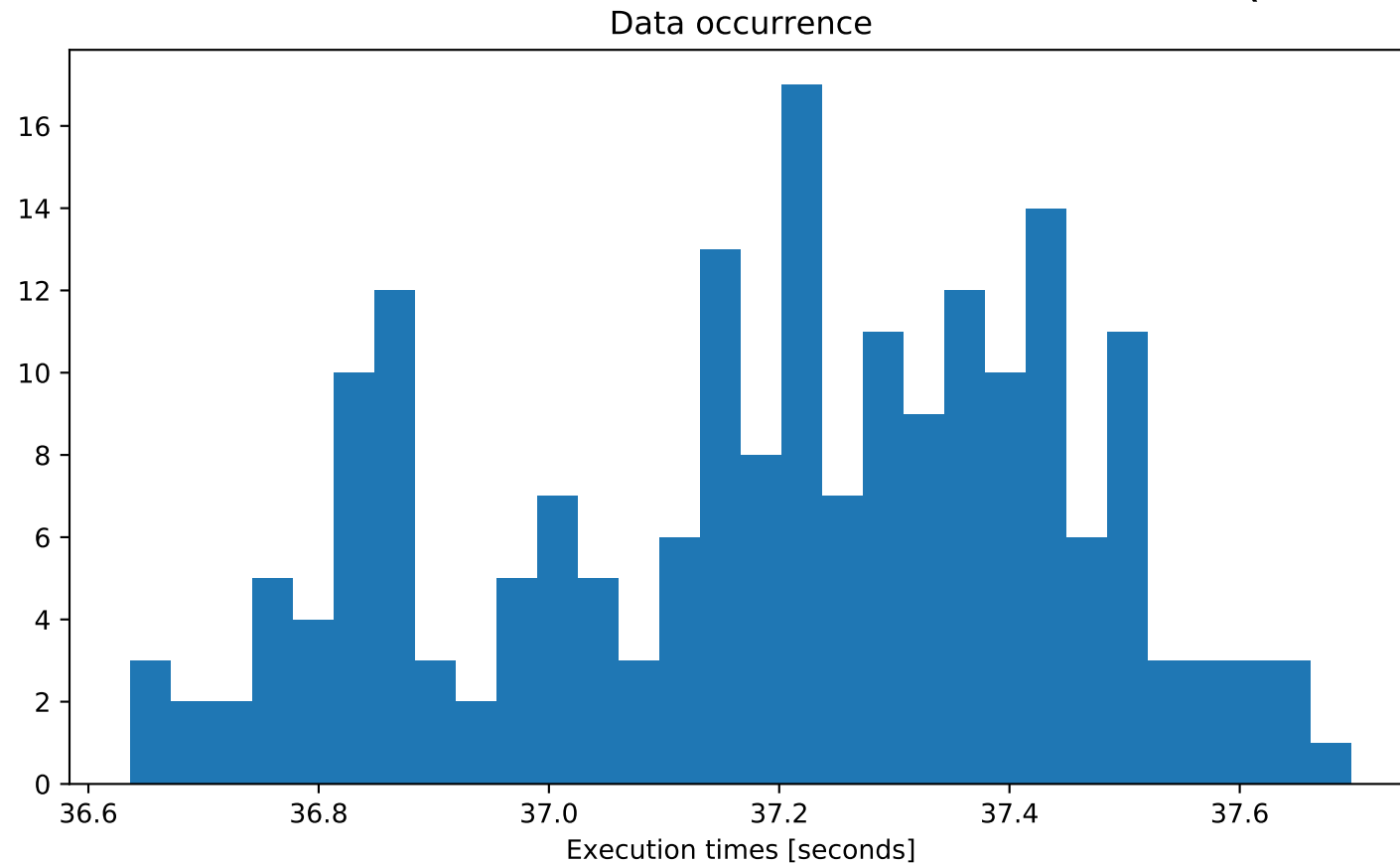


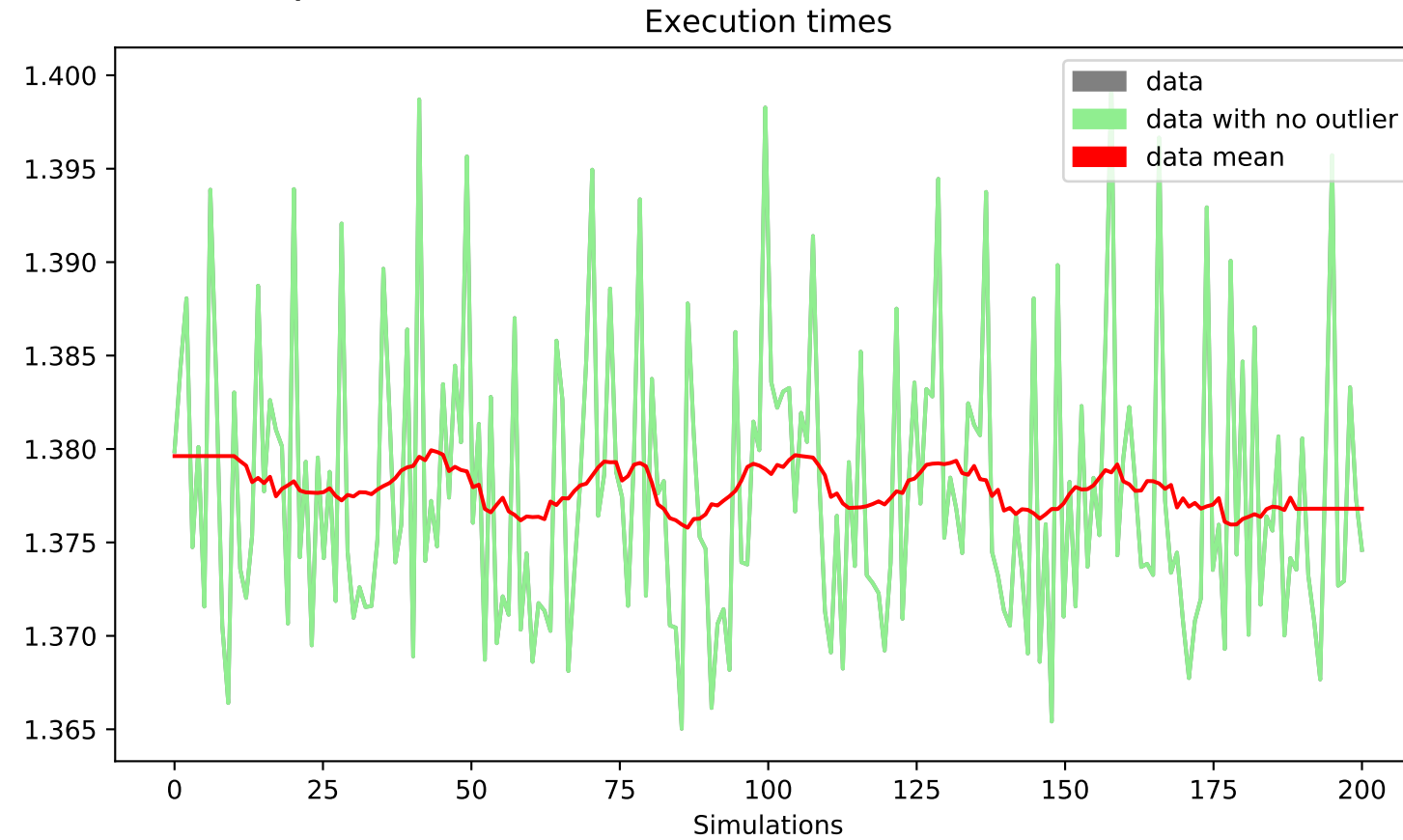
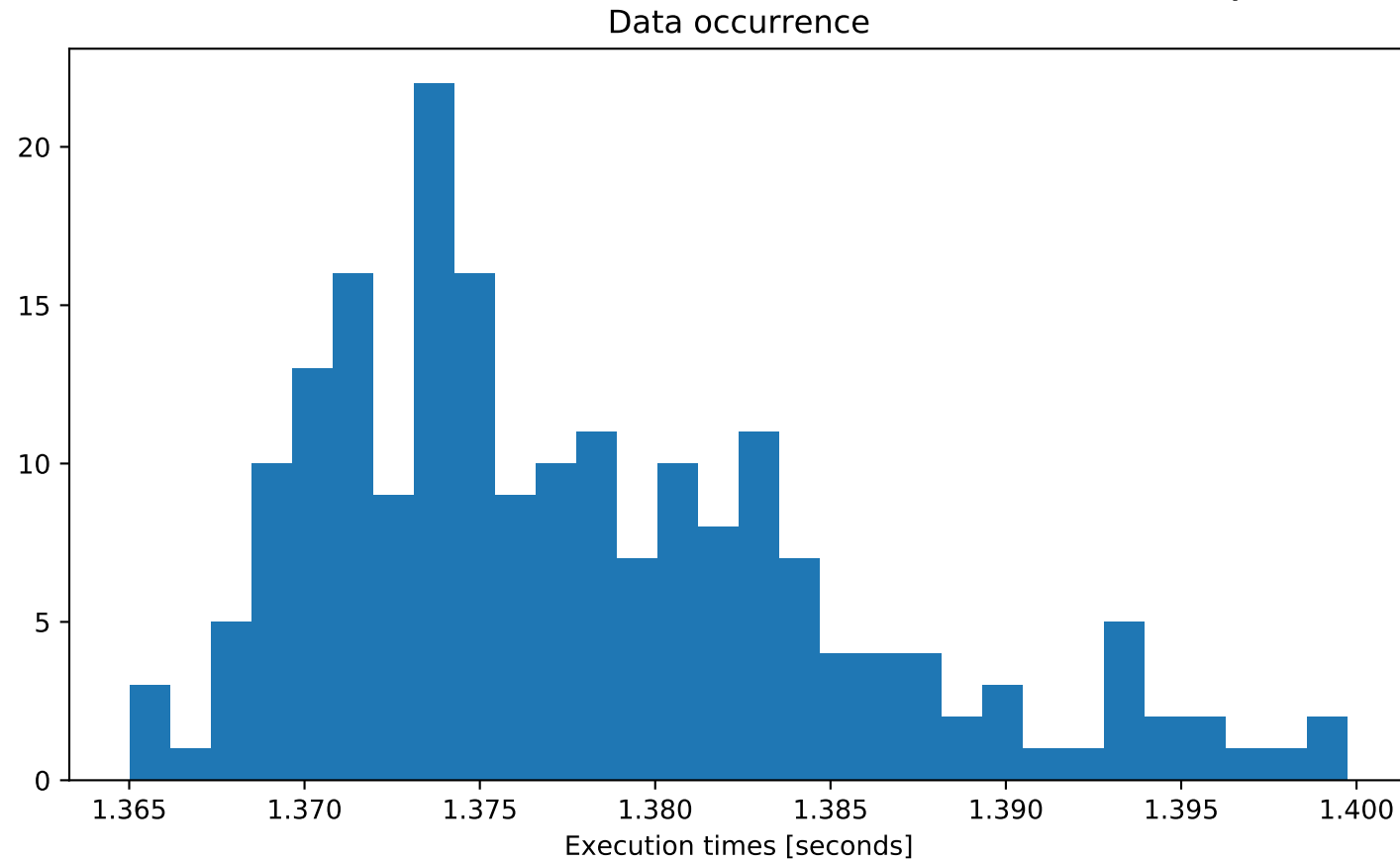
# 2MM (IID with p-value 0.309)



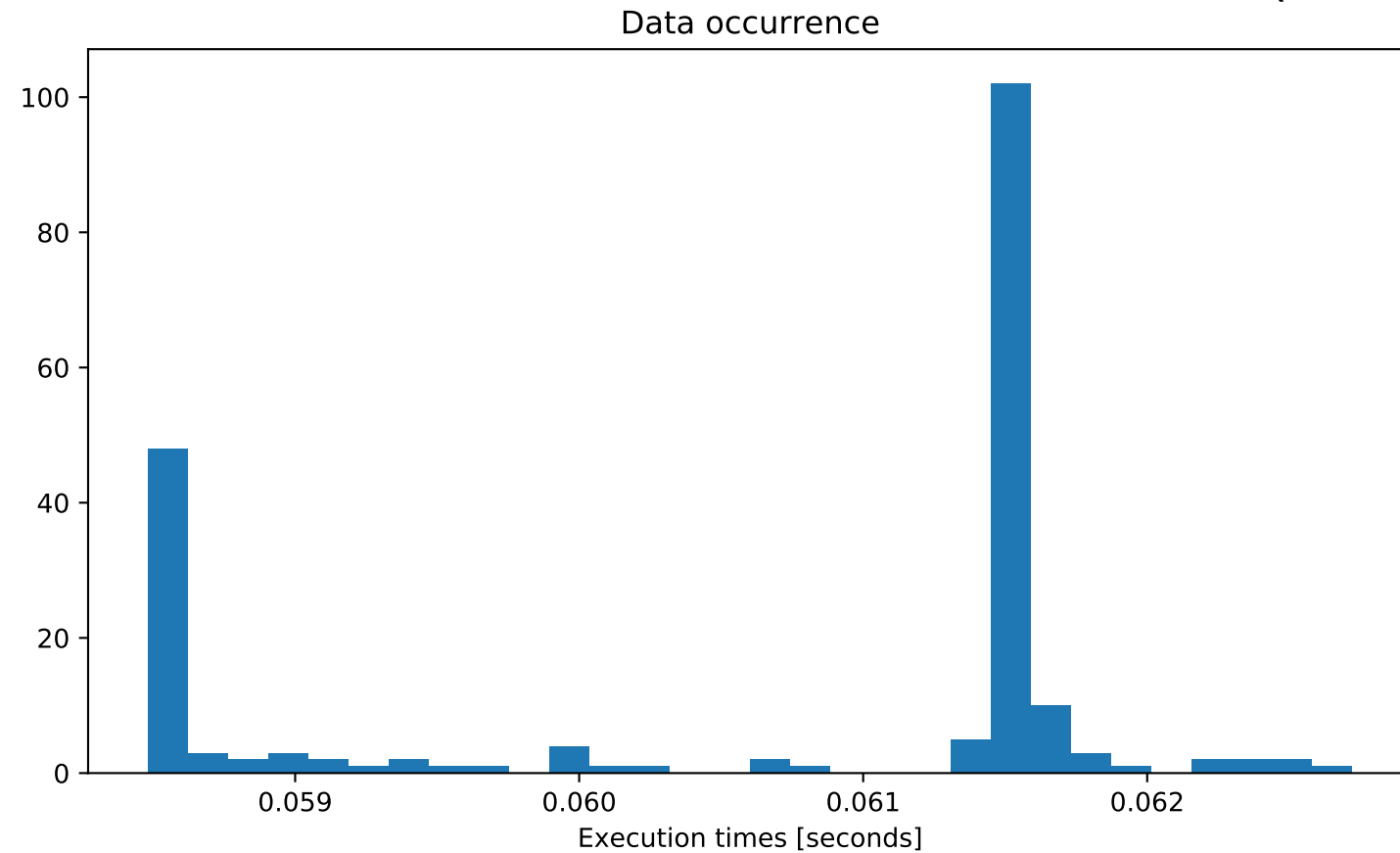
# 3MM (IID with p-value 0.5782)



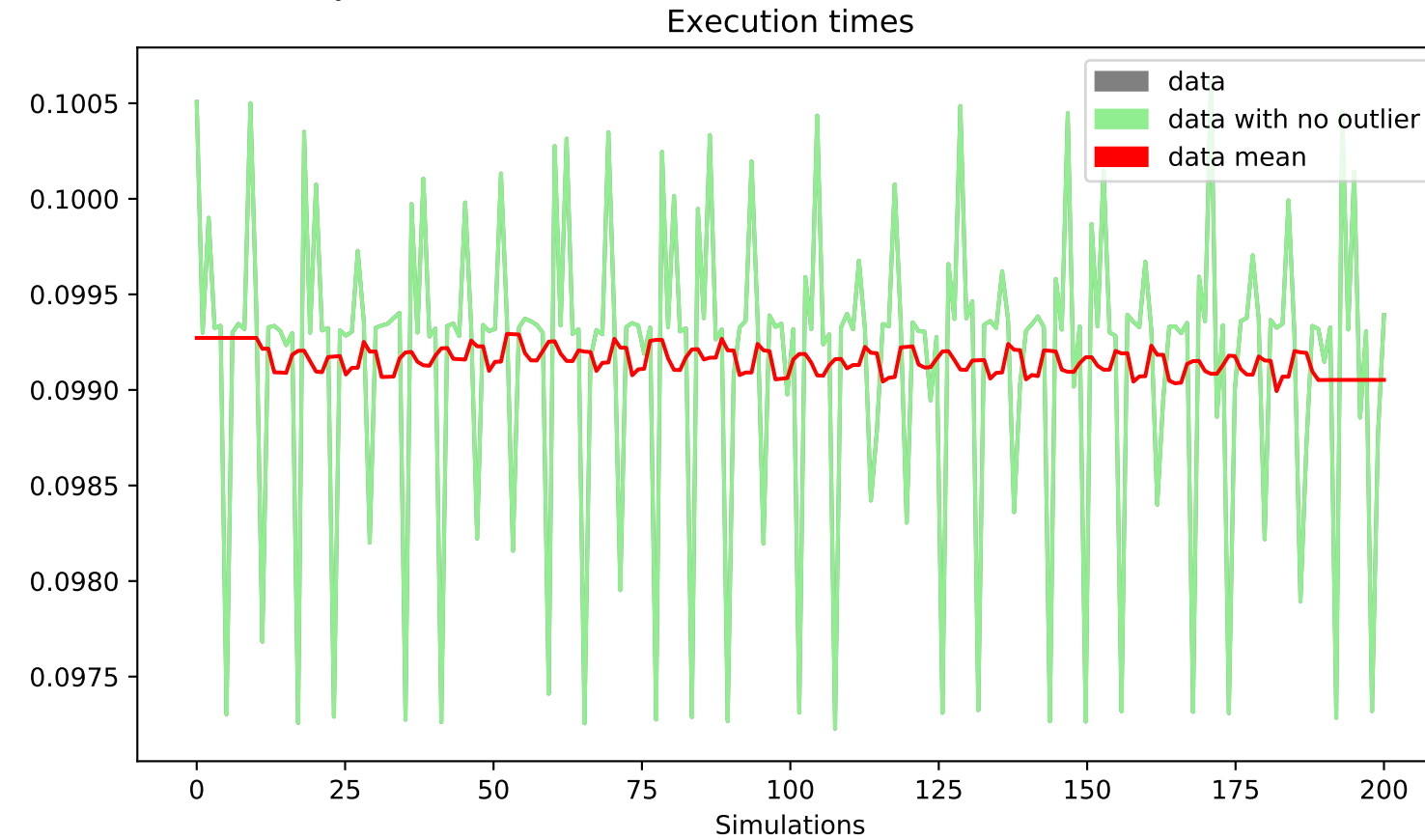
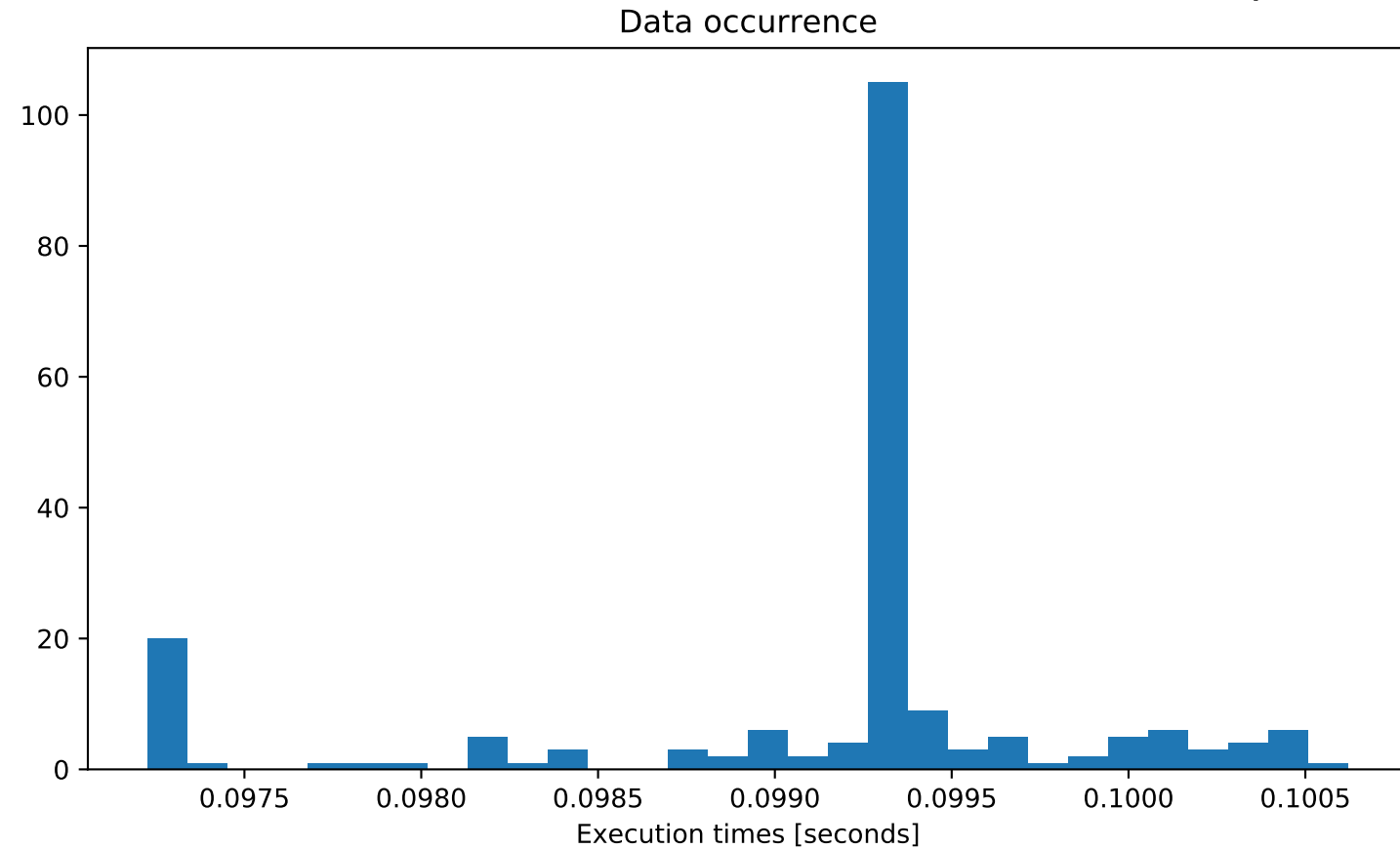
# ADI (IID with p-value 0.5226)



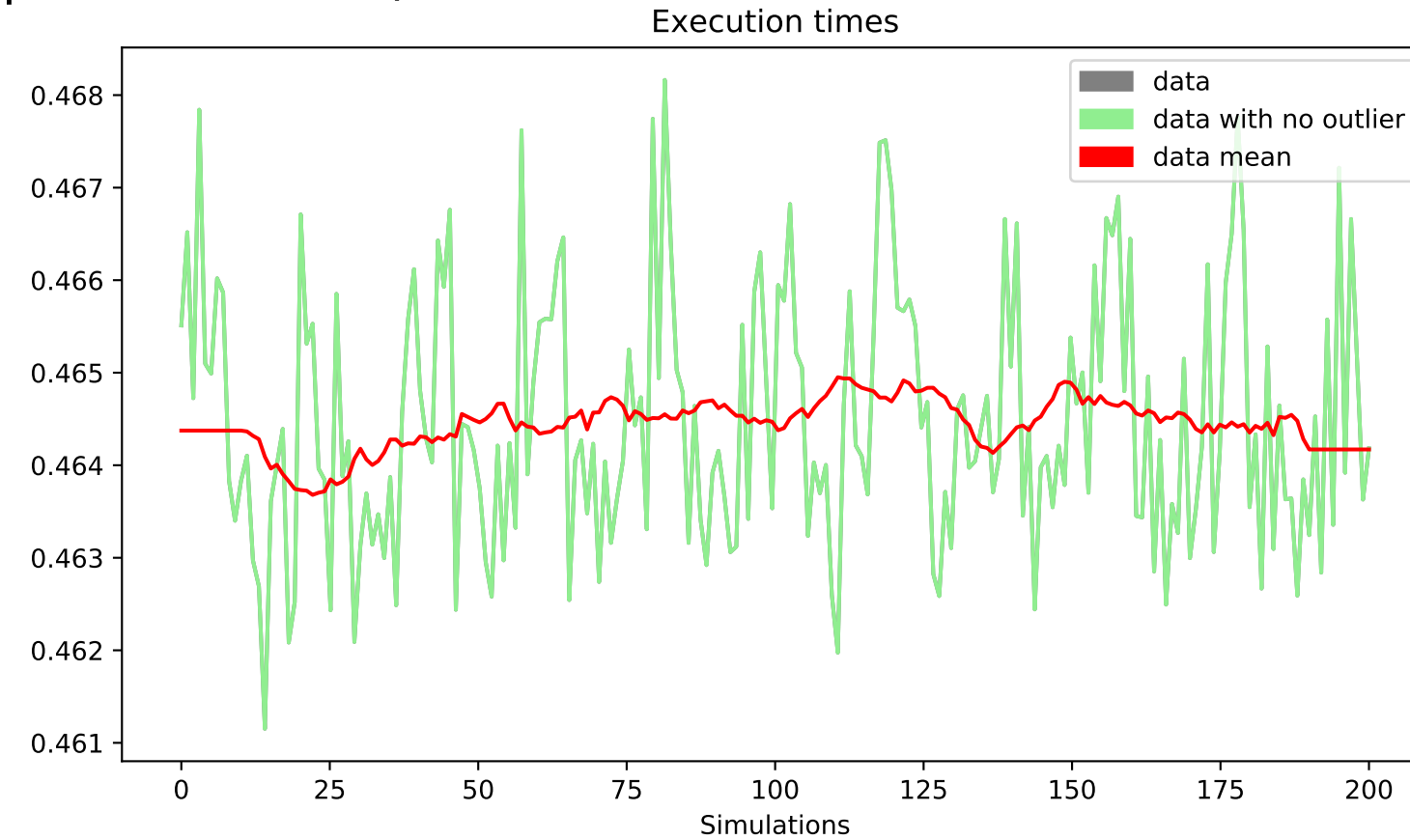
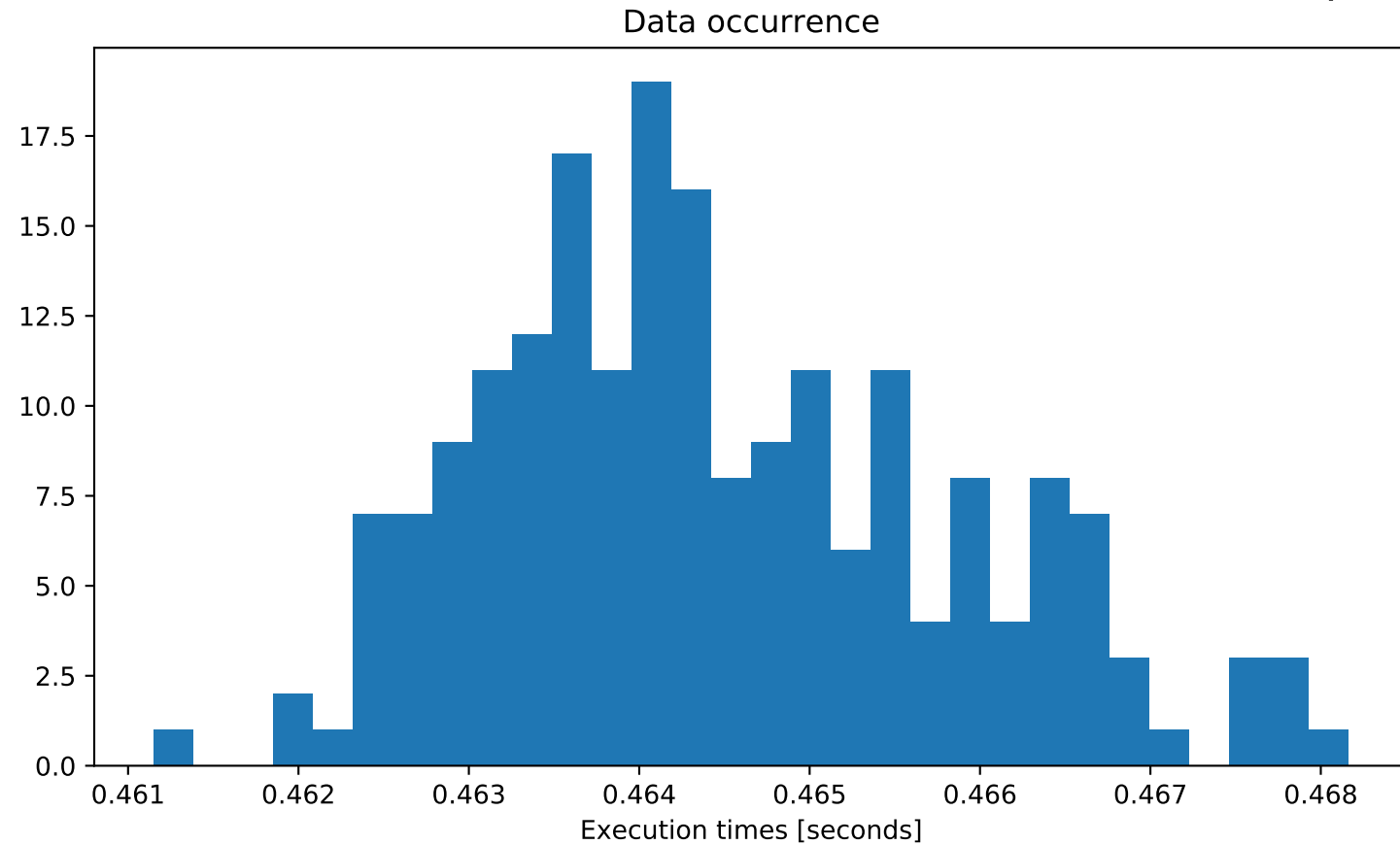
# ATAX (IID with p-value 0.539)



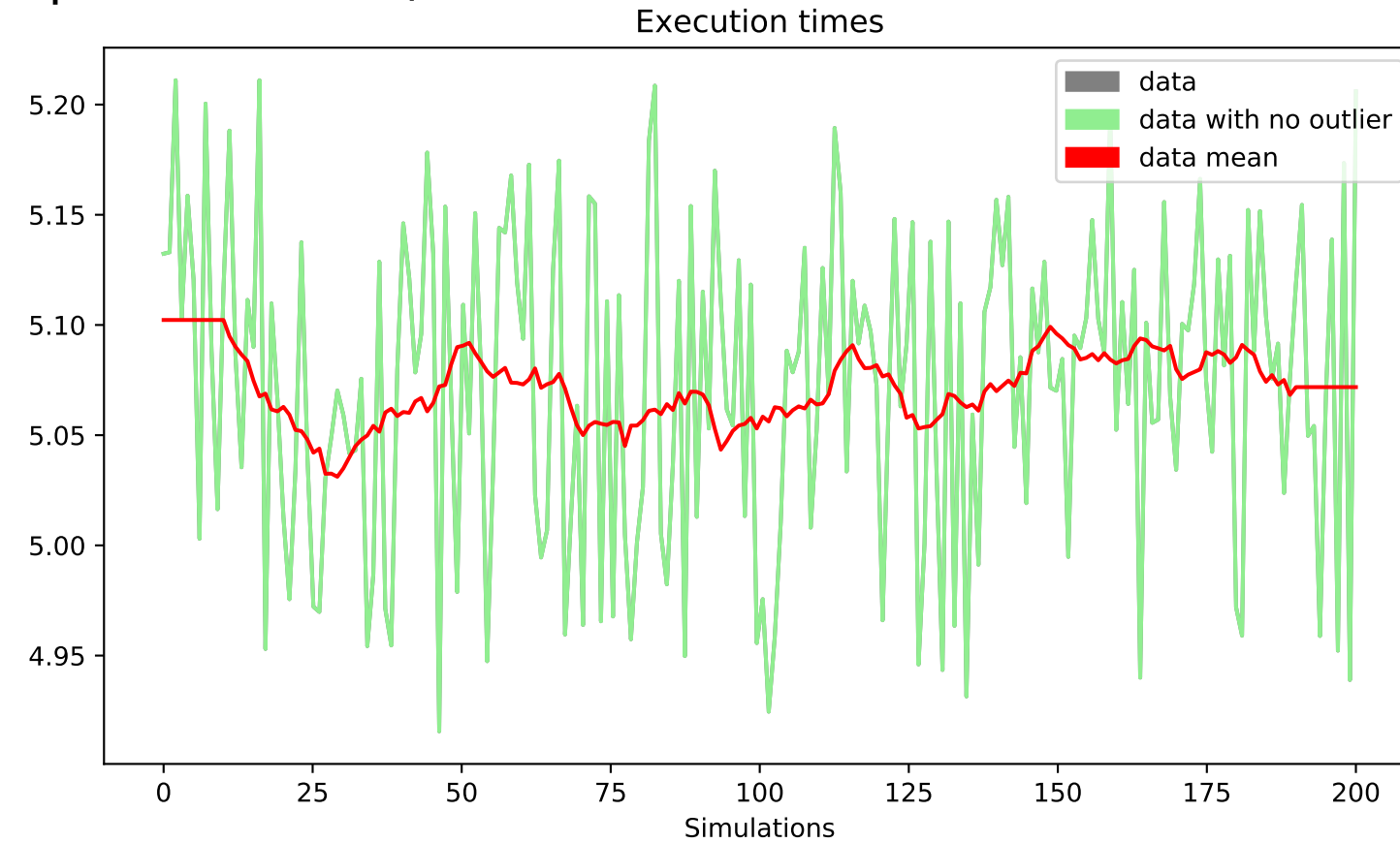
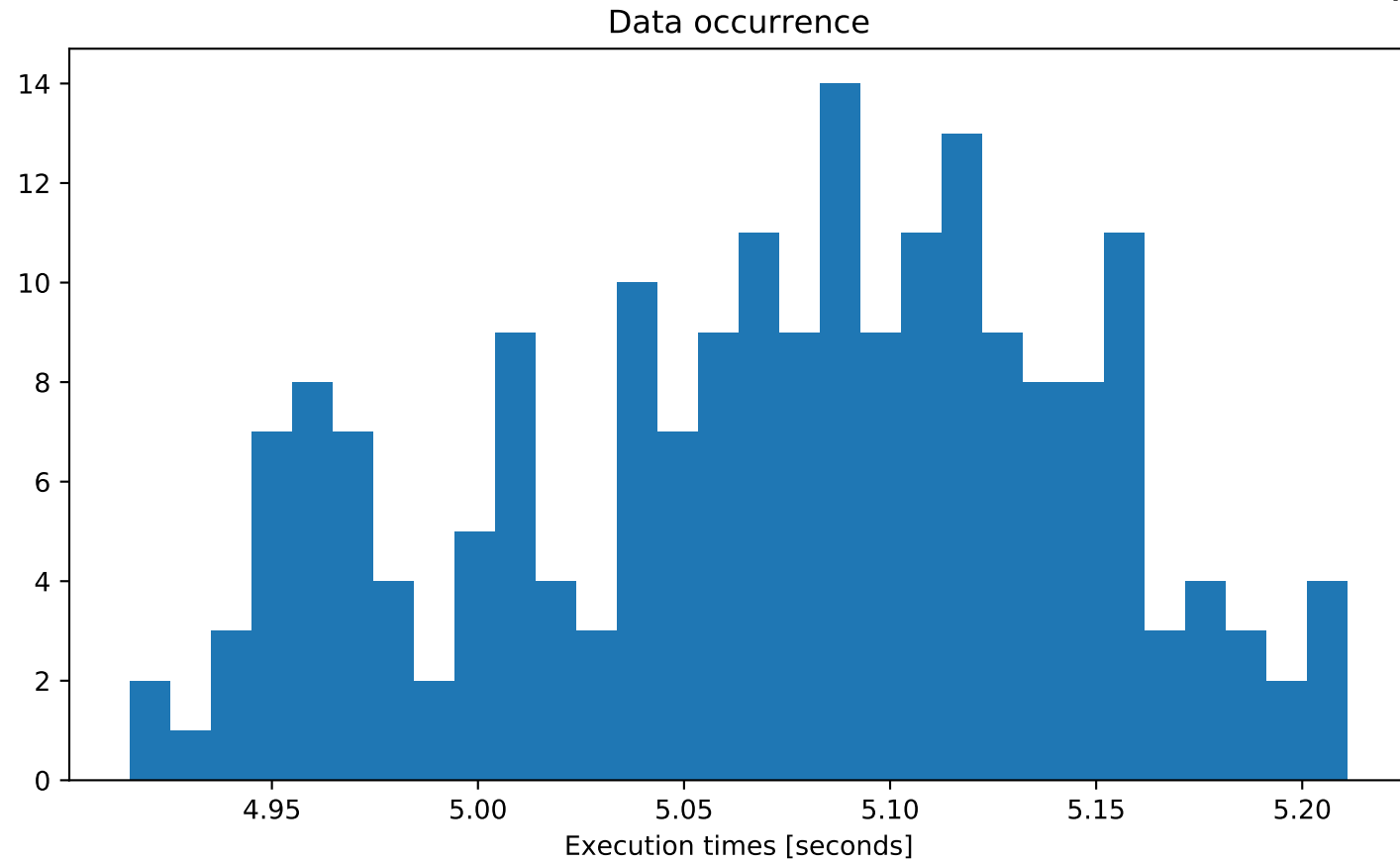
# BICG (IID with p-value 0.8058)



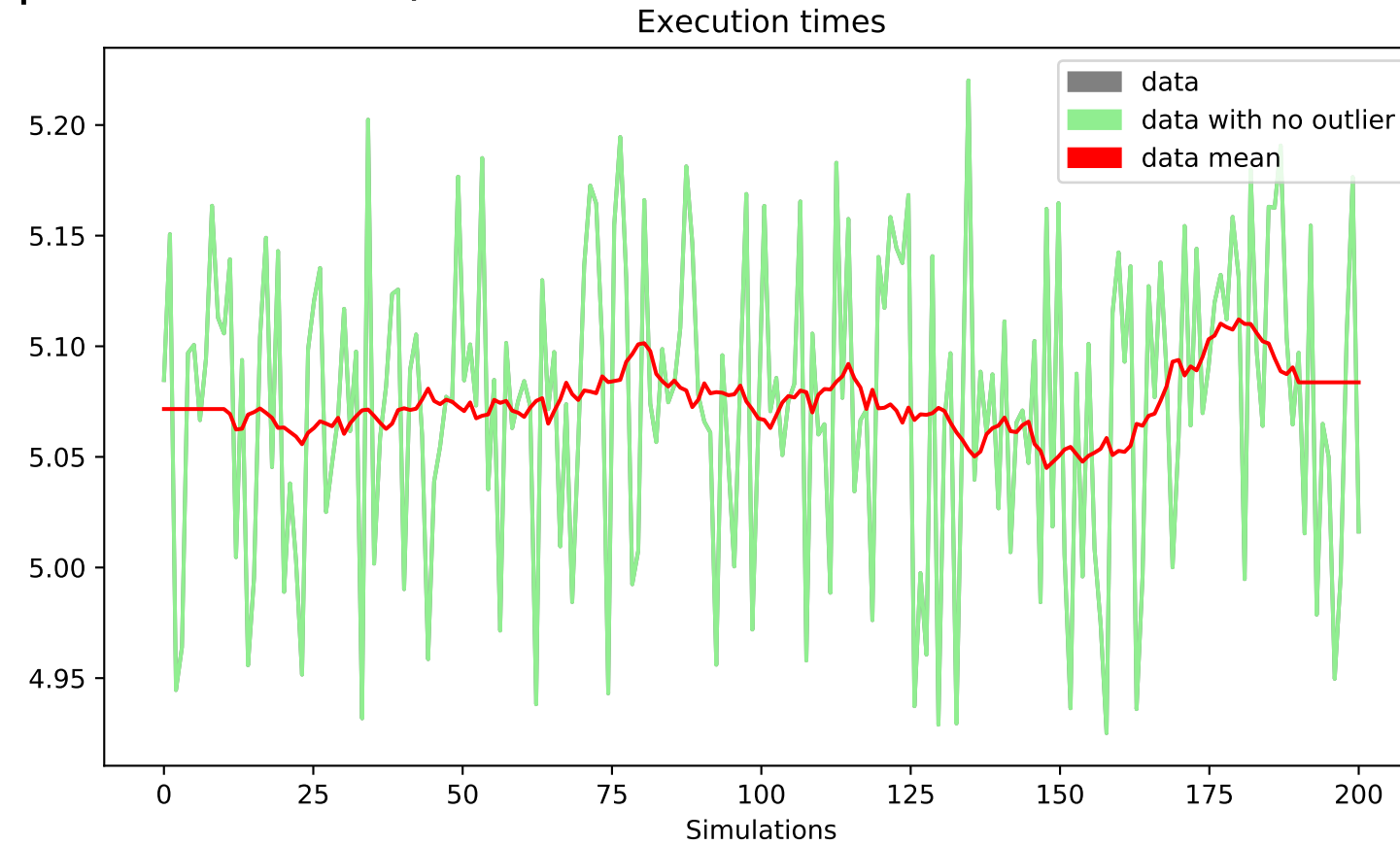
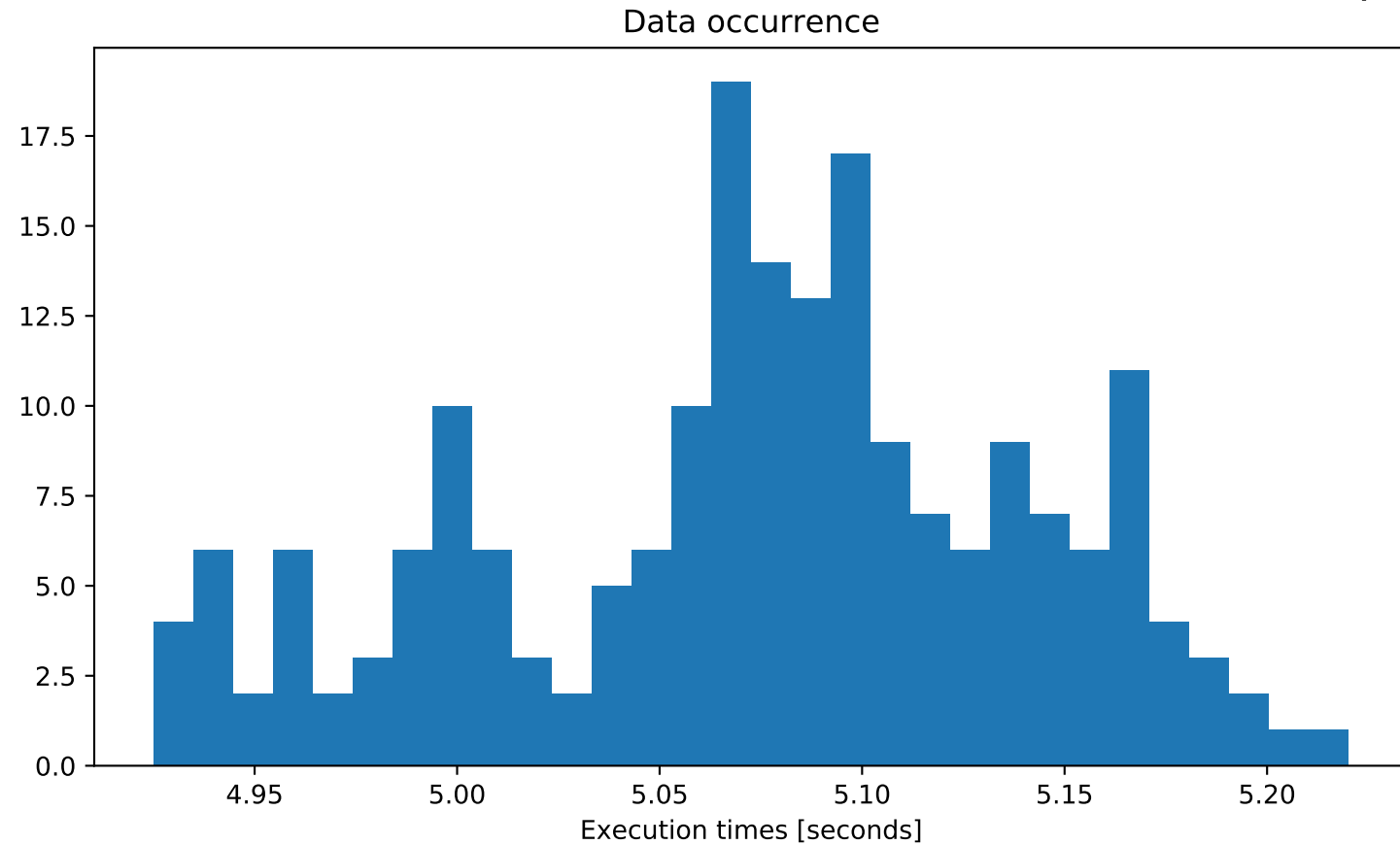
# CHOLSKY (IID with p-value 0.4038)



# CORRELATION (IID with p-value 0.753)

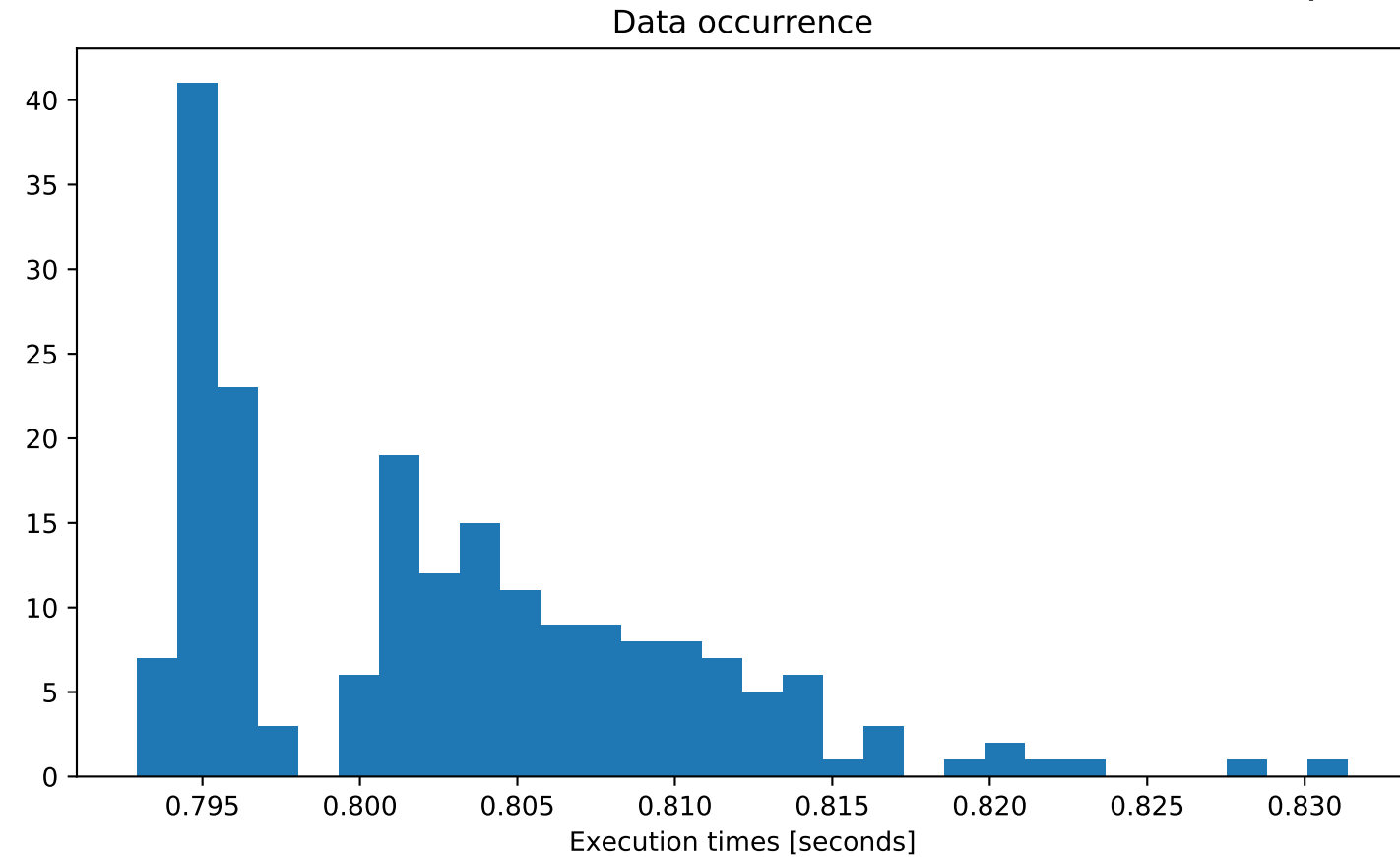


# COVARIANCE (IID with p-value 0.4988)

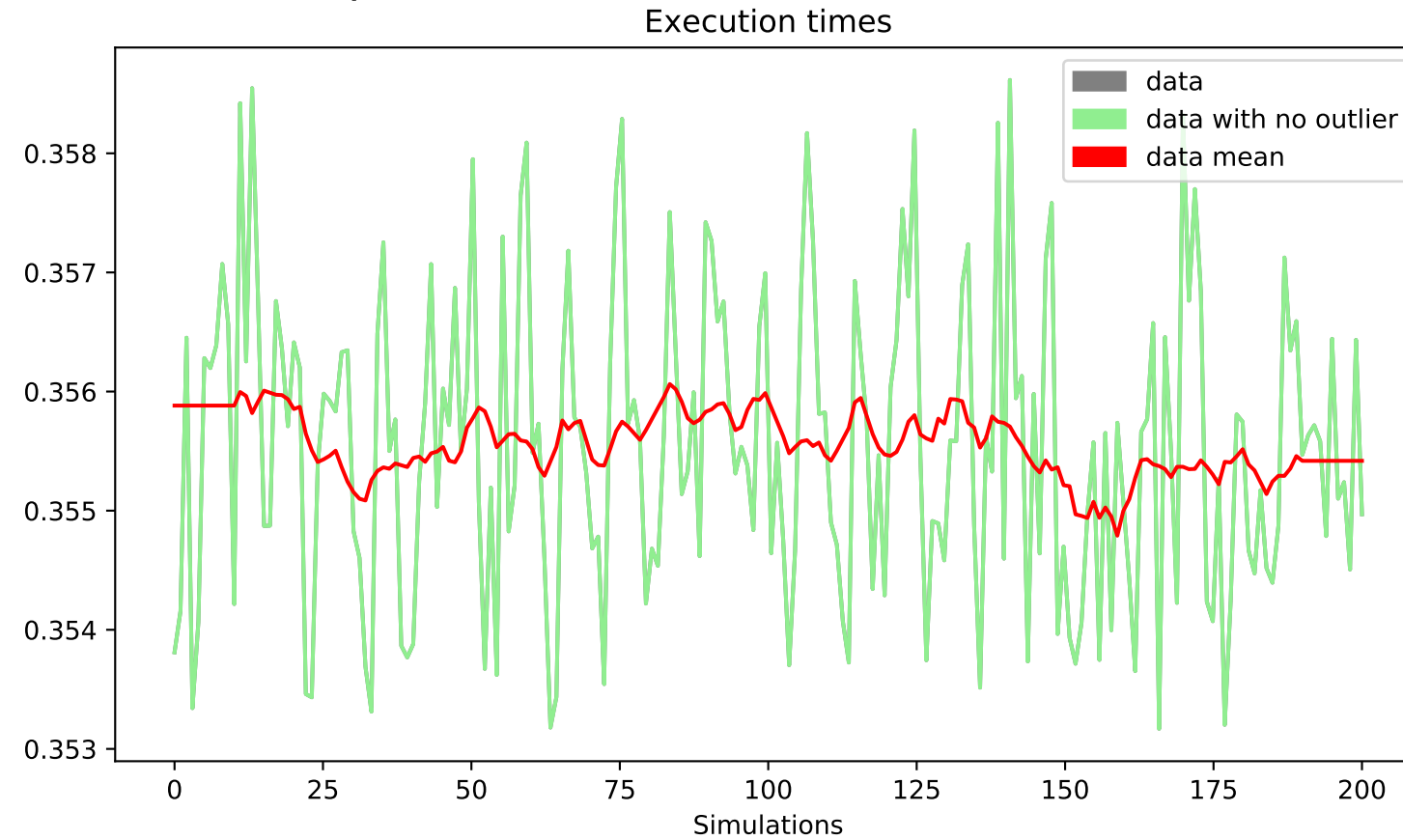
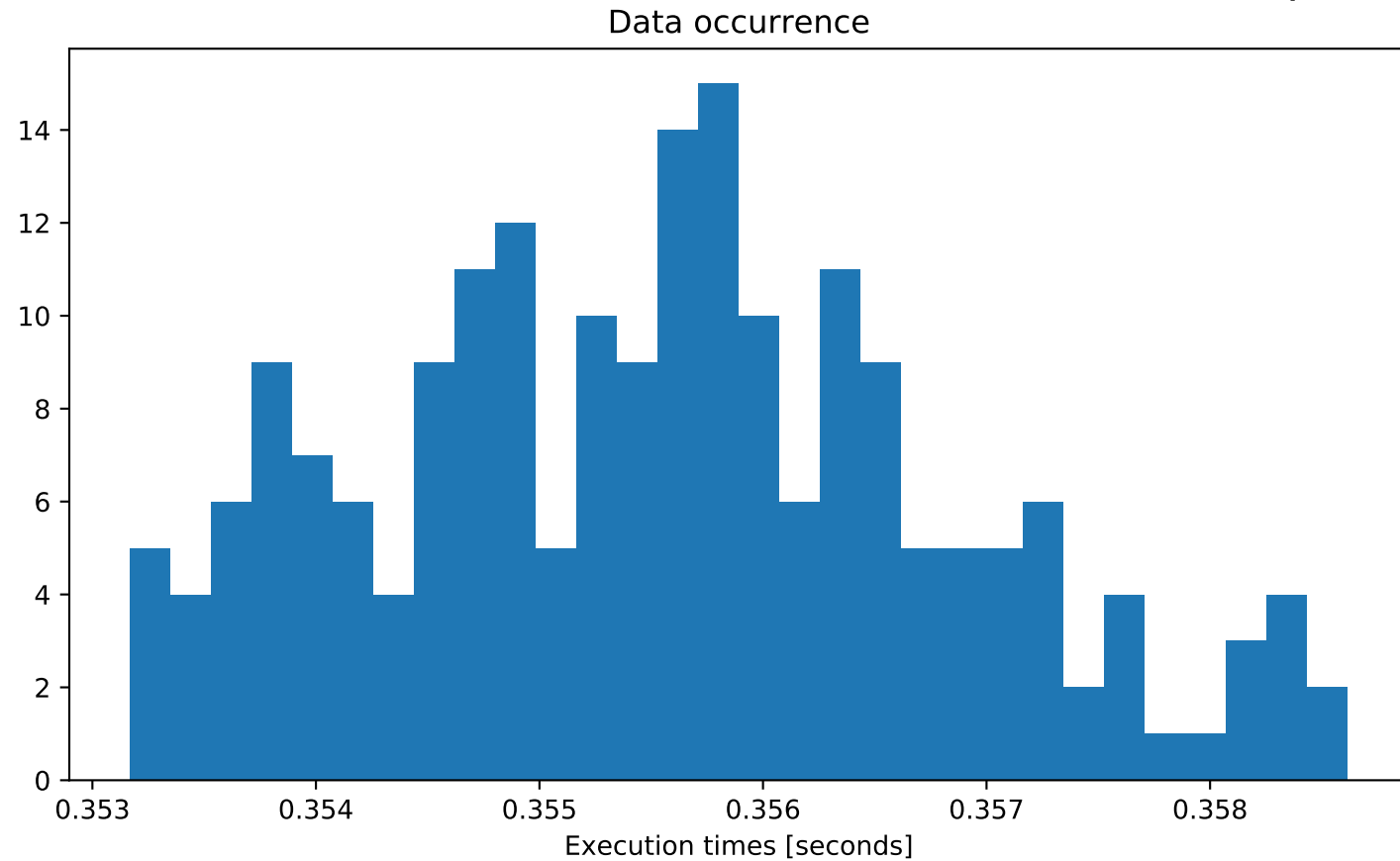




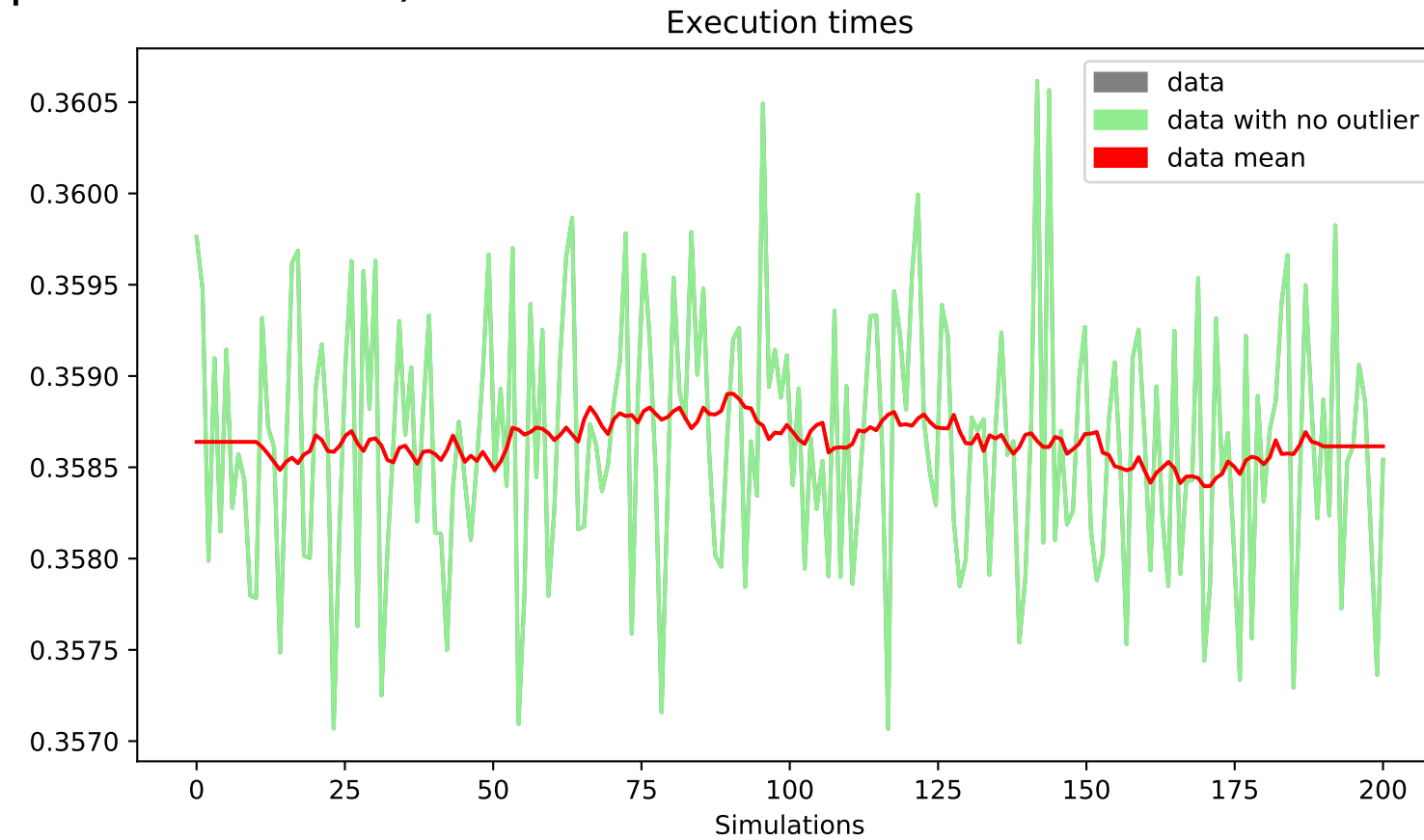
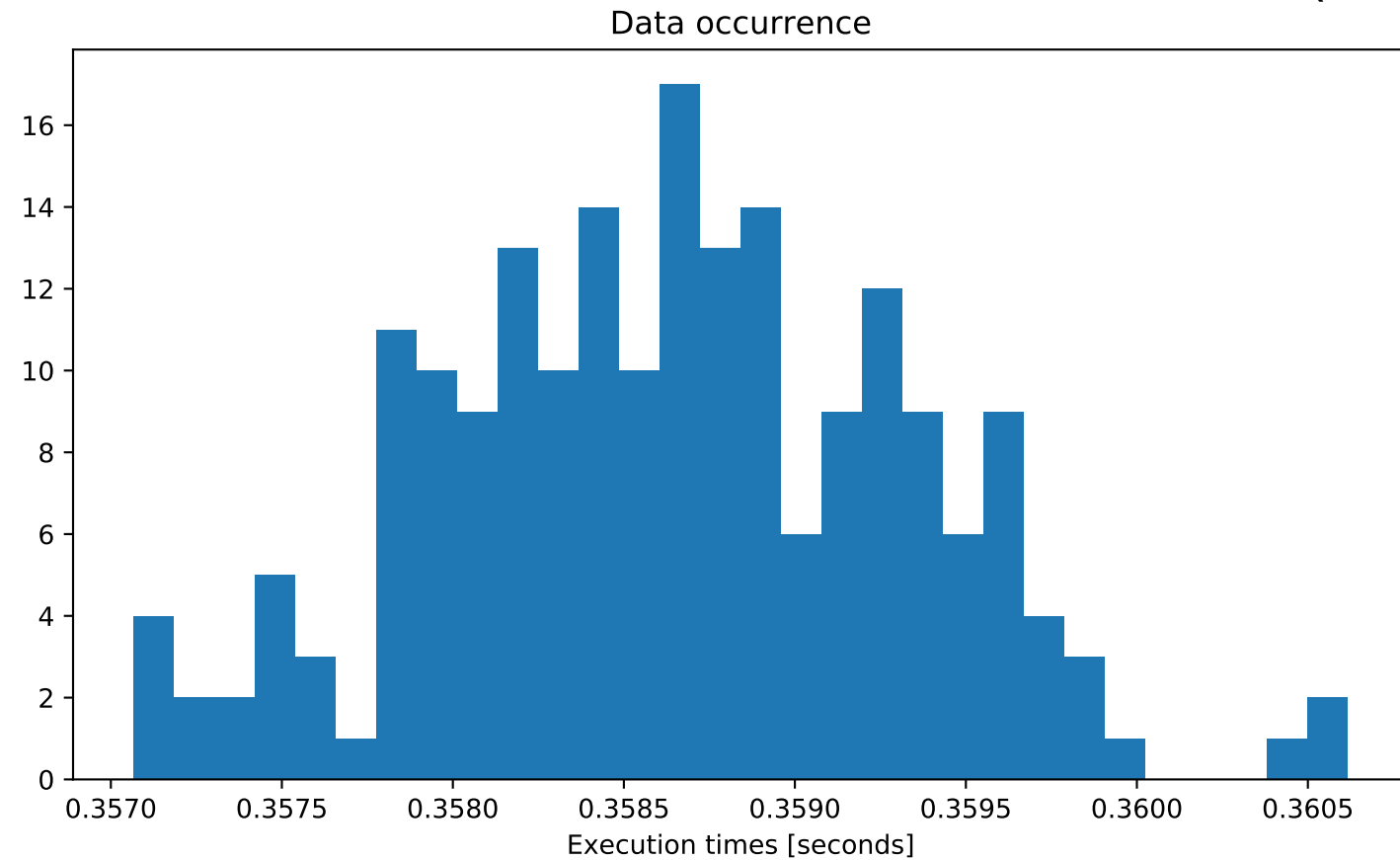
# DOITGEN (IID with p-value 0.5938)



# DURBIN (IID with p-value 0.6962)

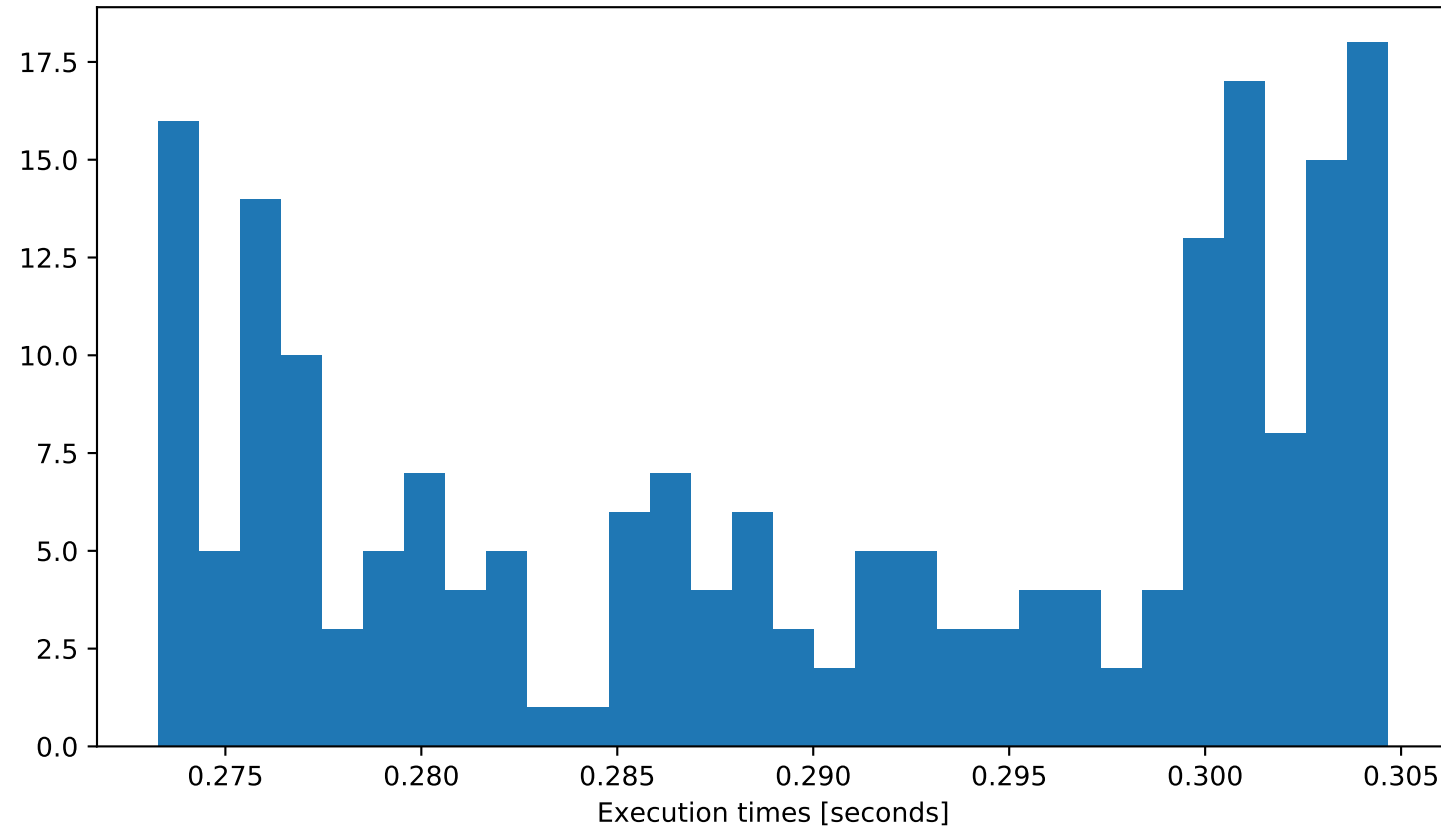


# DYNPROG (IID with p-value 0.5702)

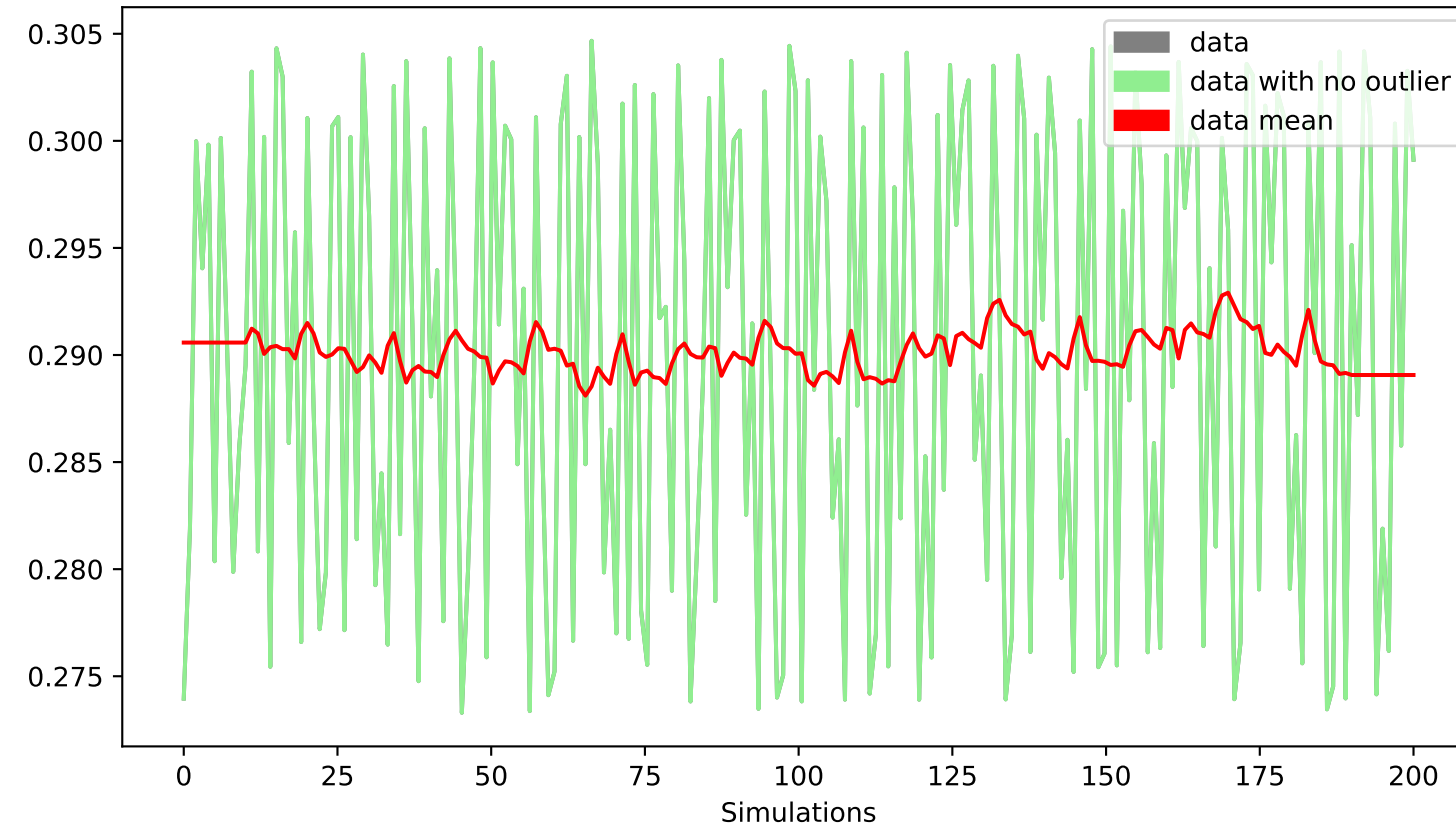


# FDTD-2D (IID with p-value 0.78)

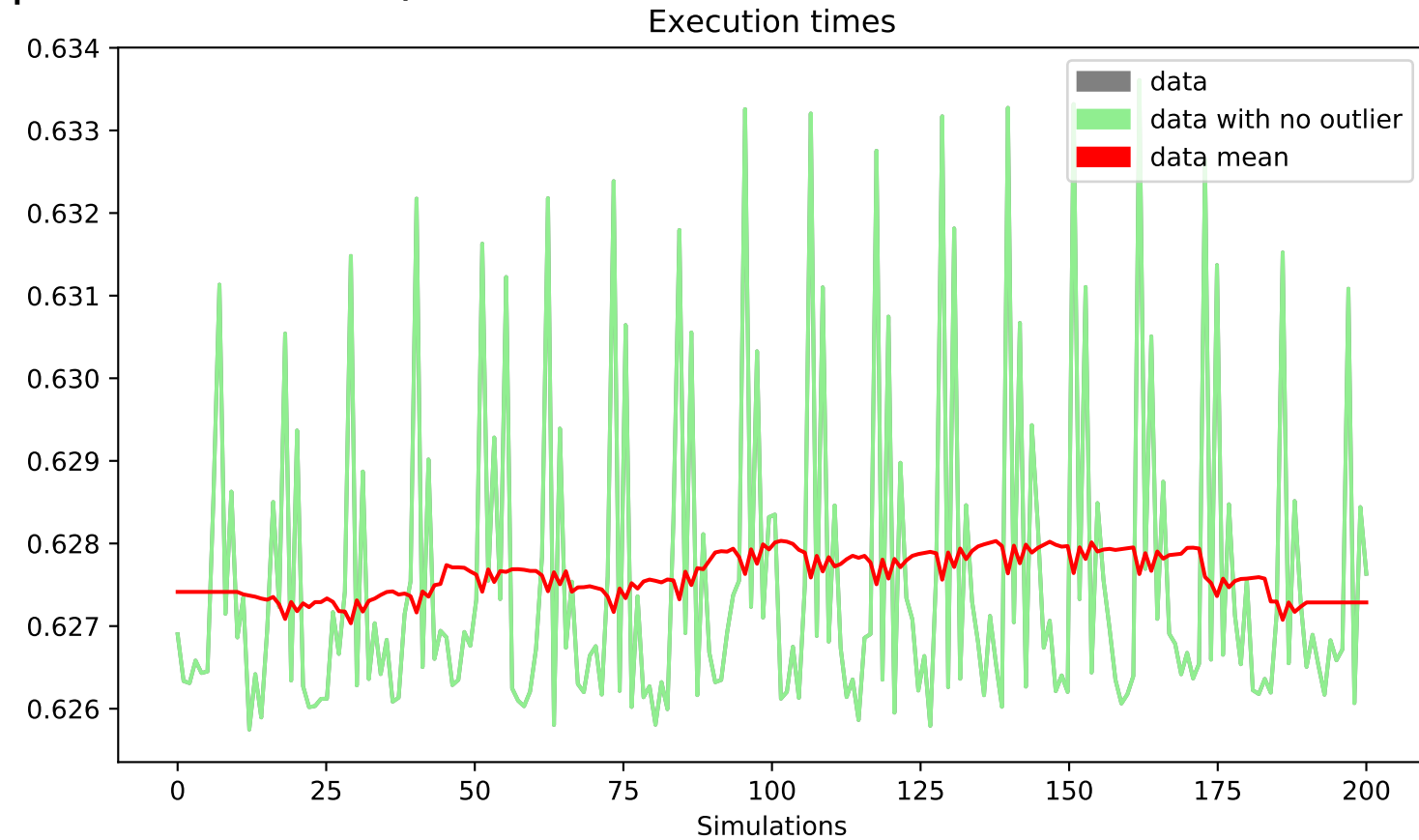
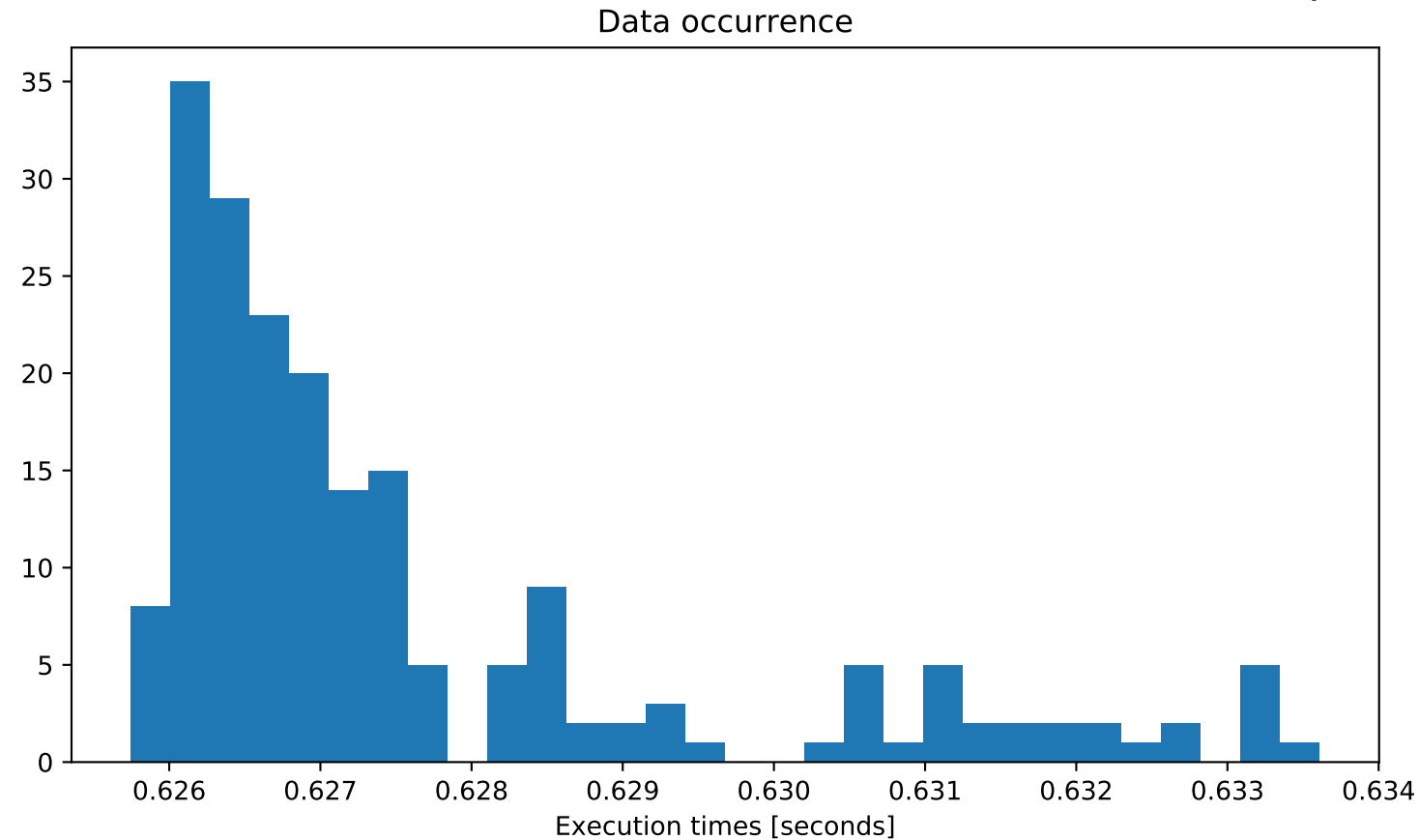
Data occurrence



Execution times

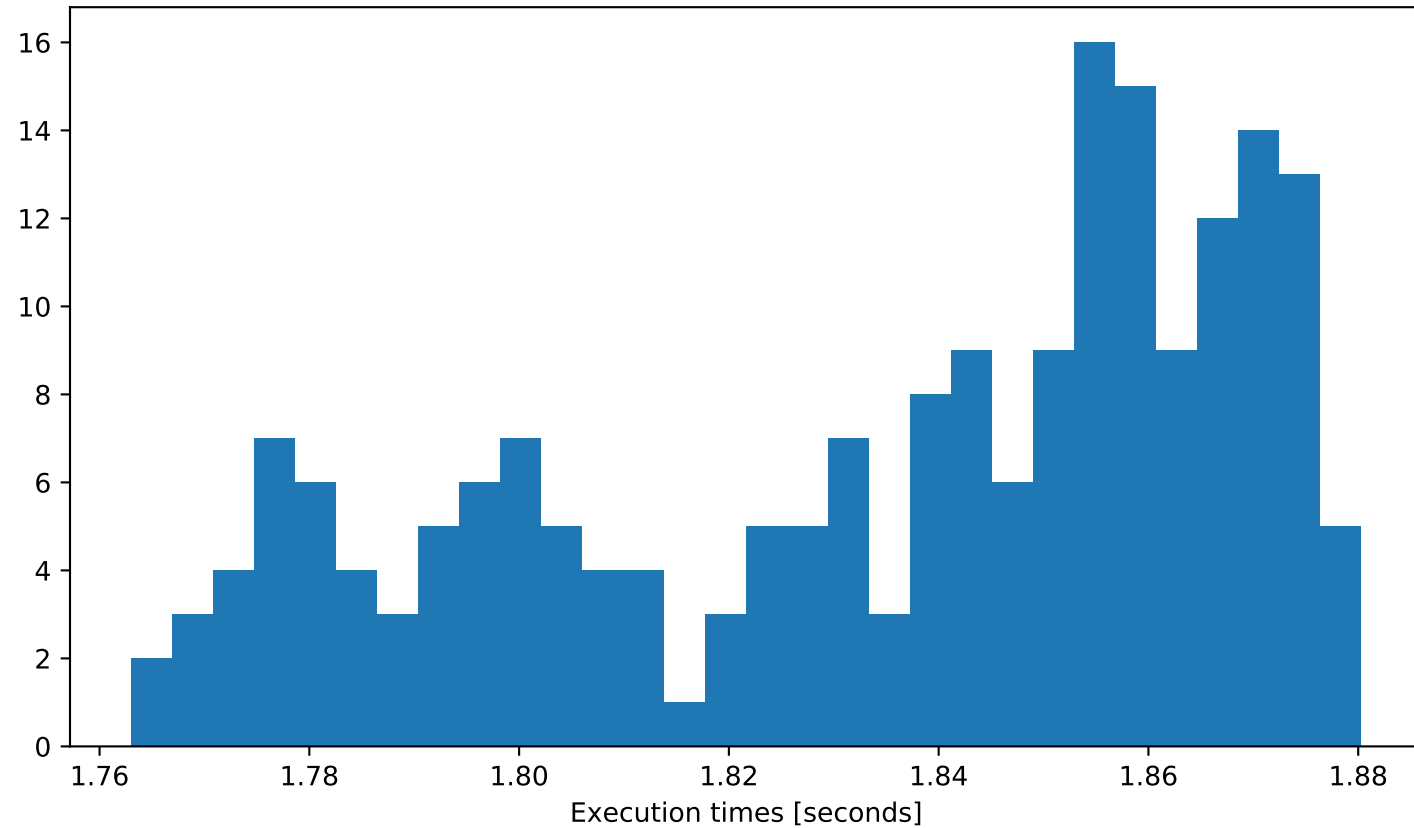


# FDTD-APML (IID with p-value 0.6076)

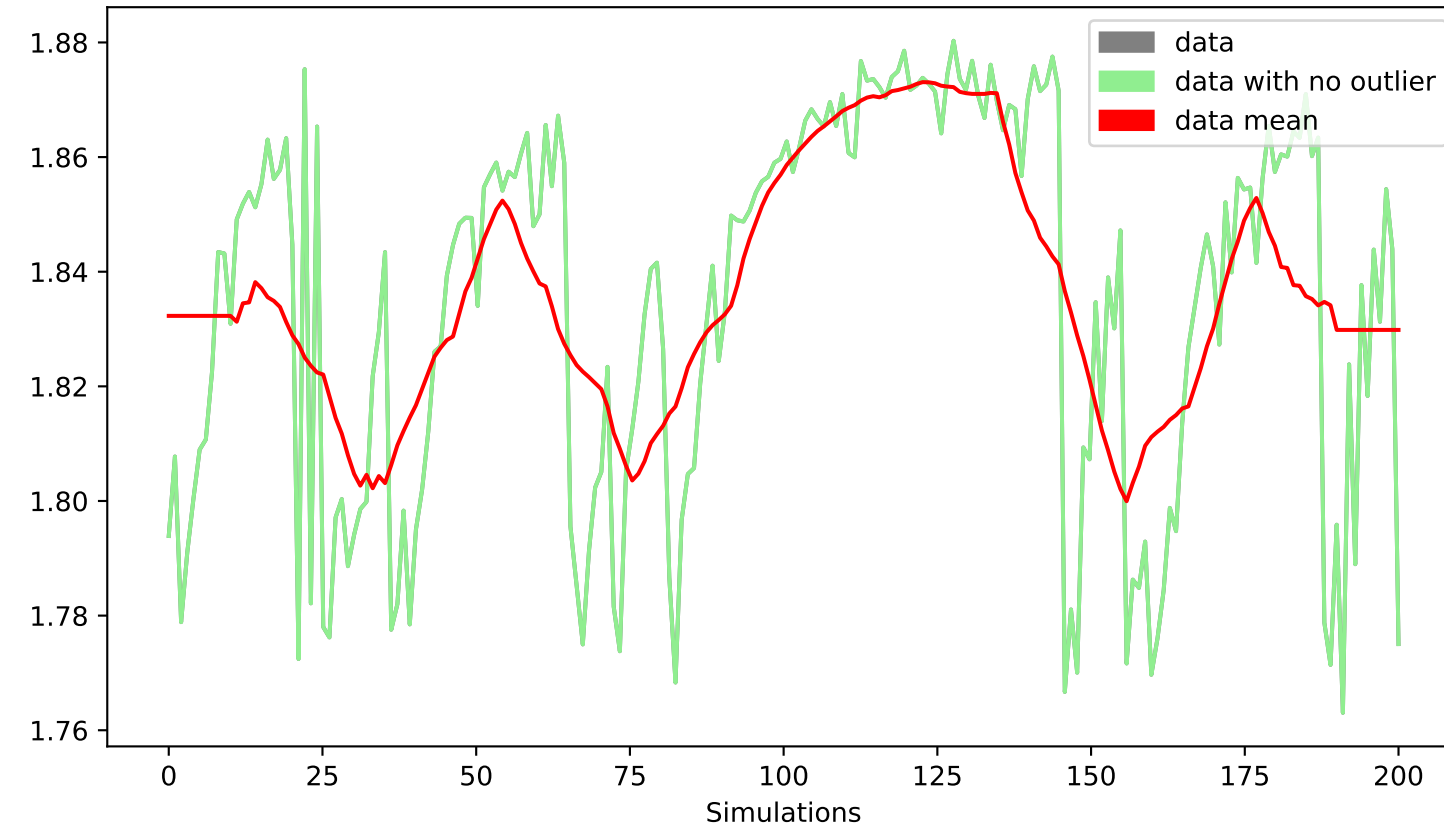


# FLOYD-WARSHALL (NOT IID with p-value 0.0113333333)

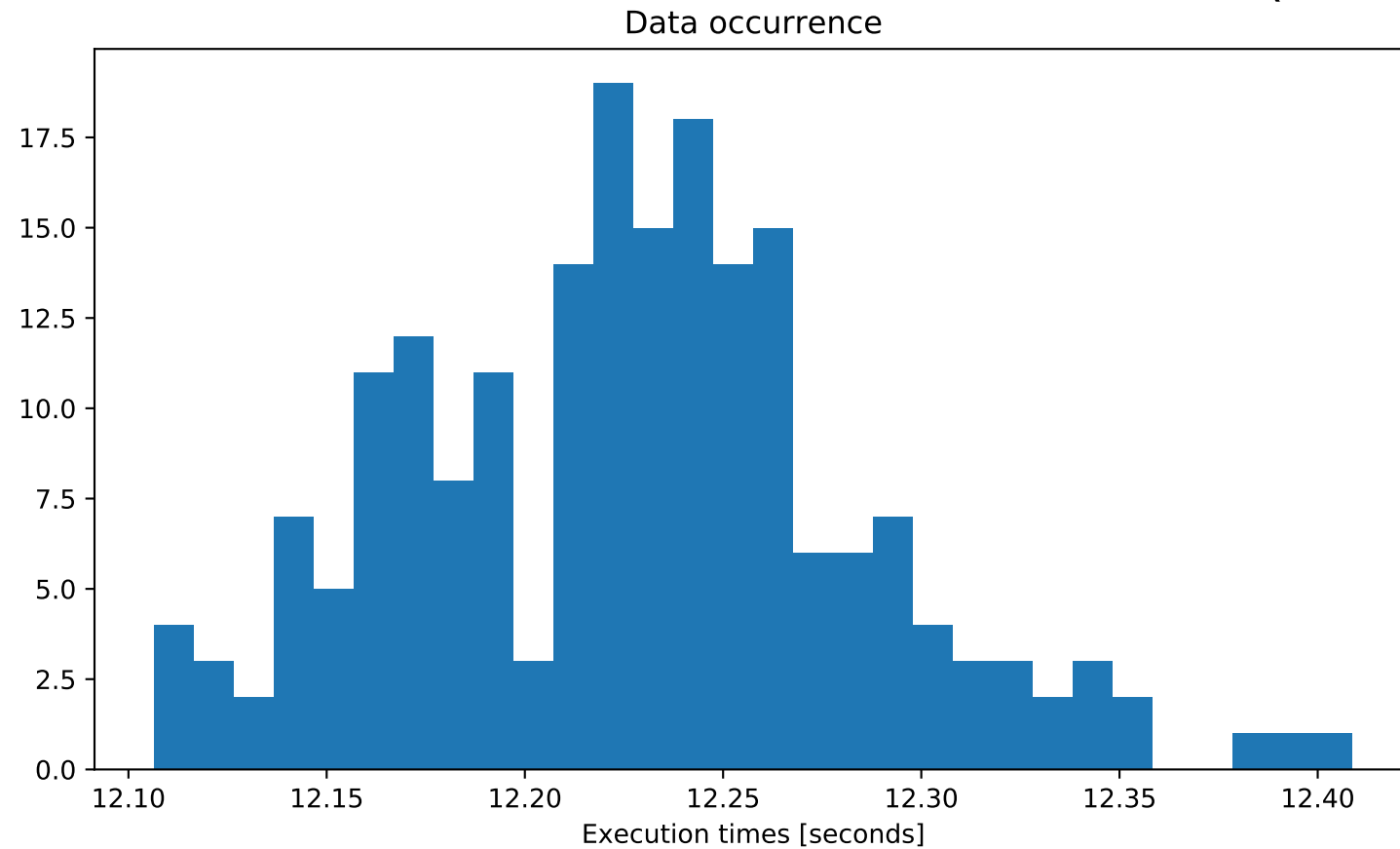
Data occurrence



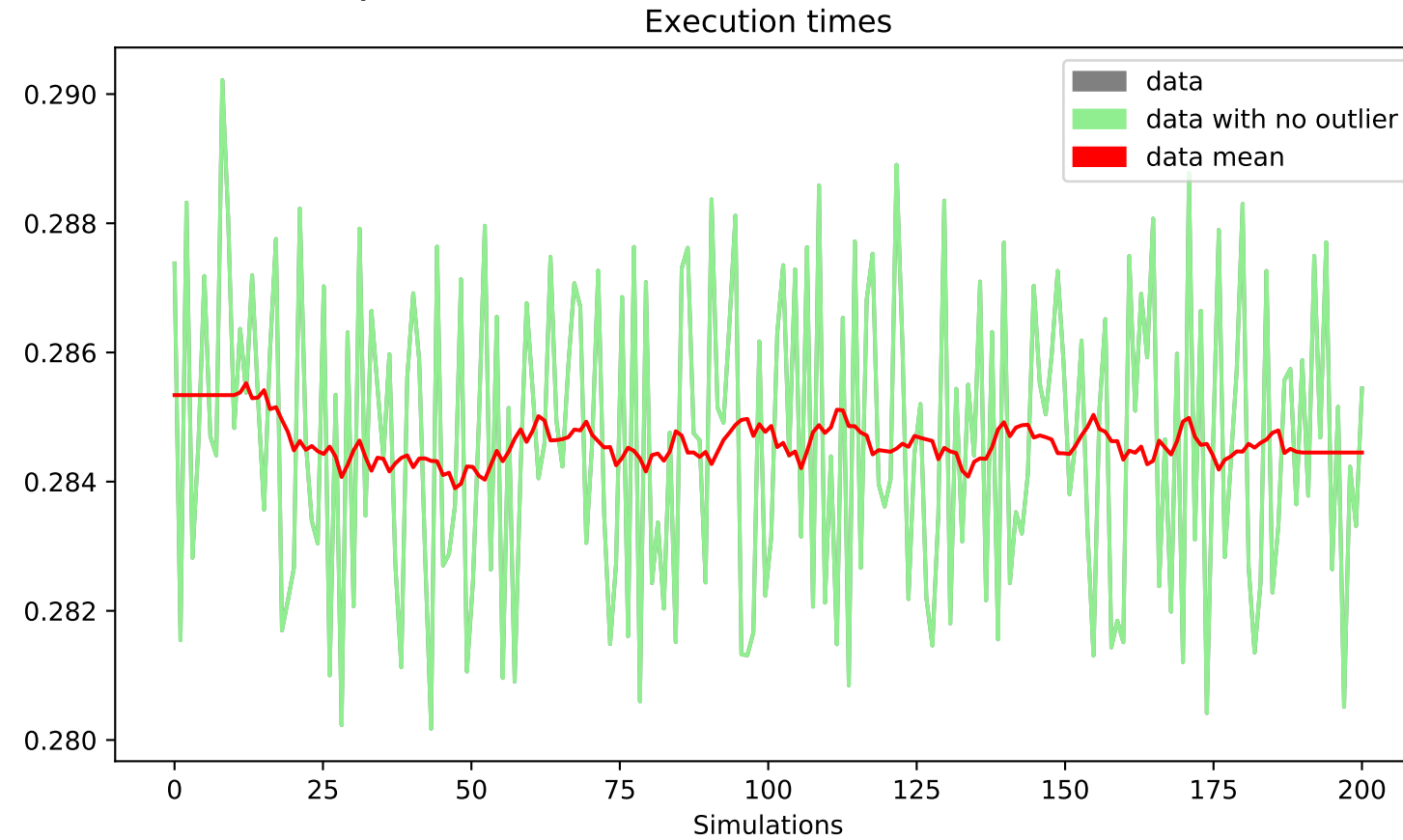
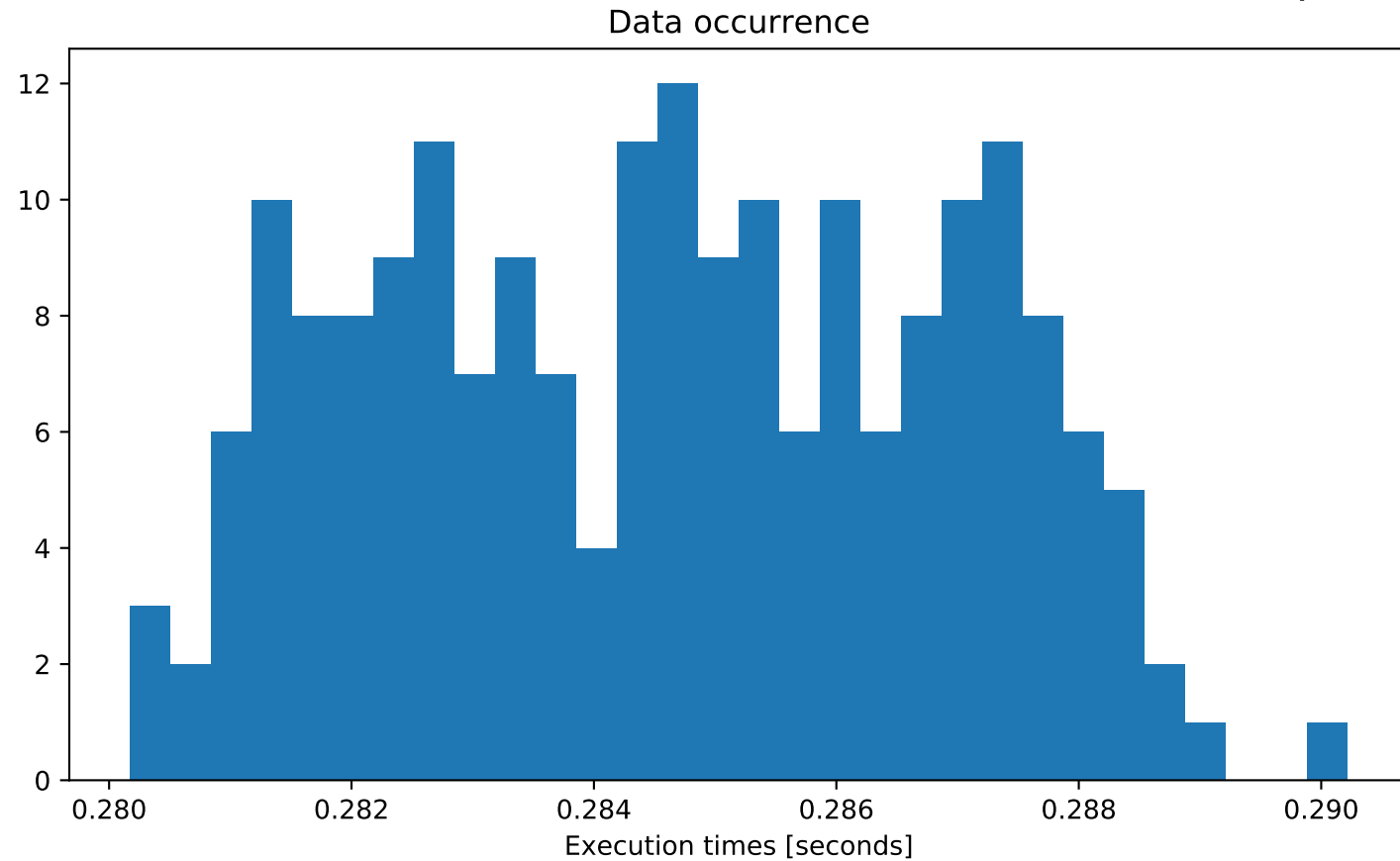
Execution times



# GEMM (IID with p-value 0.4248)

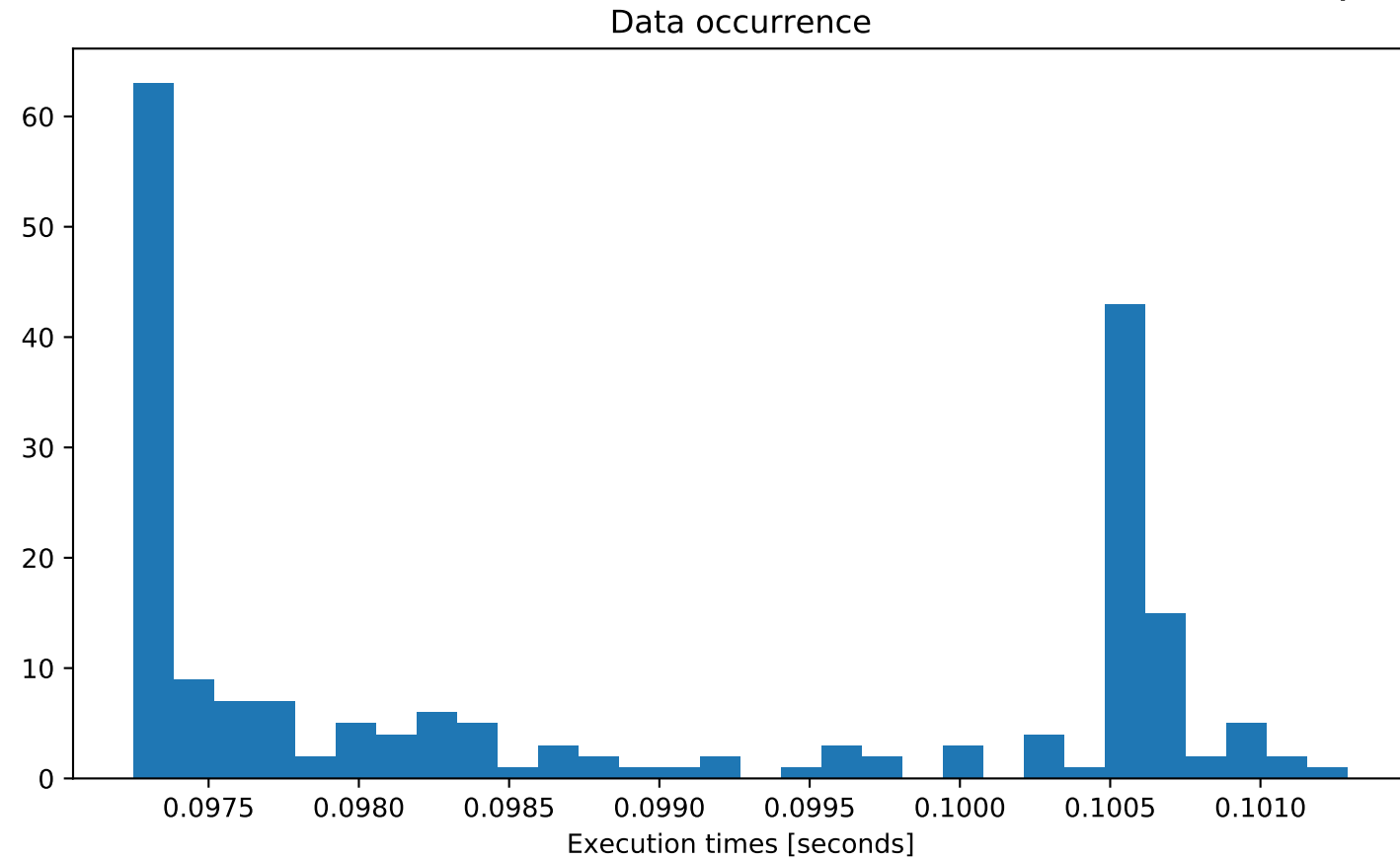


# GEMVER (IID with p-value 0.6864)

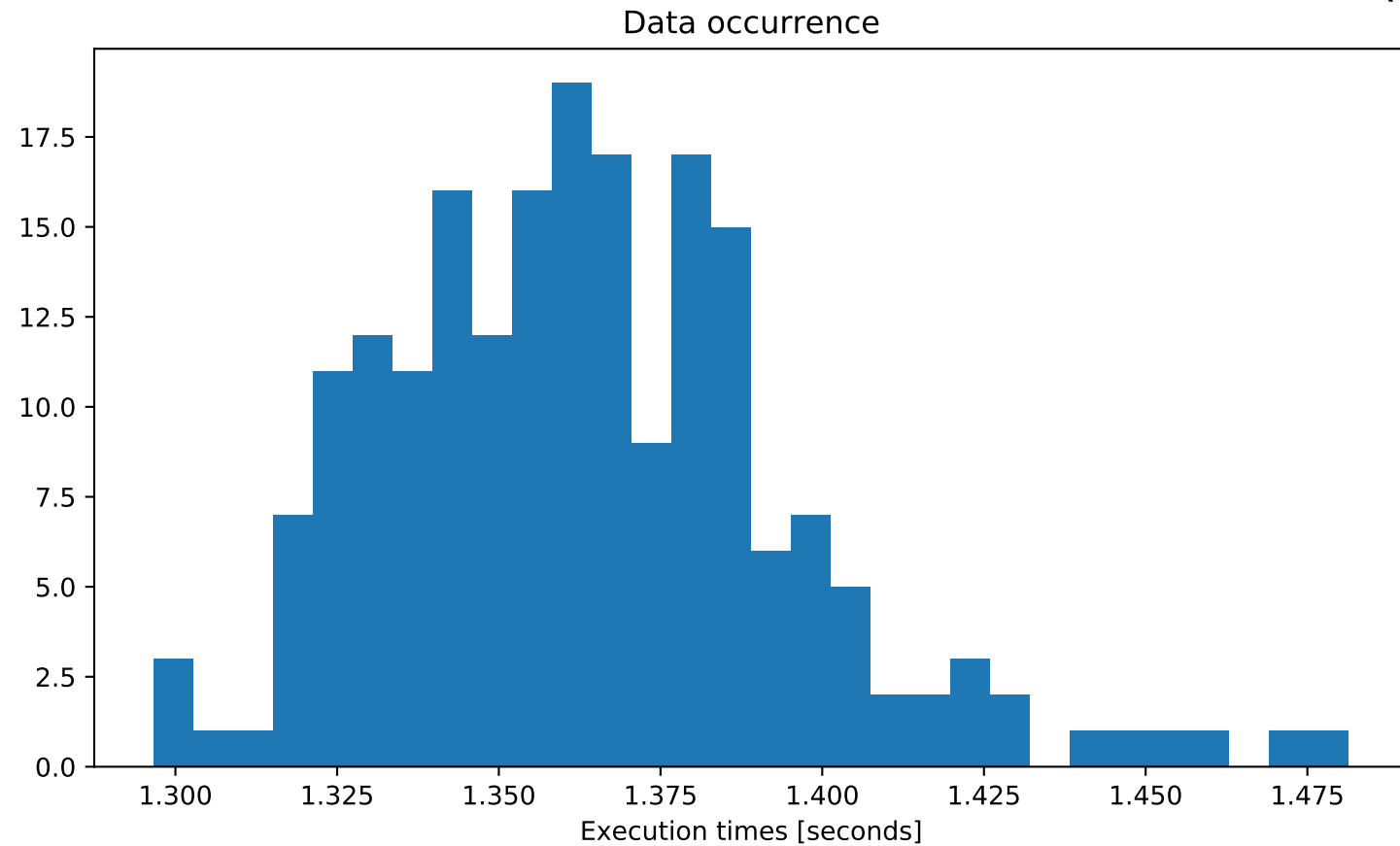




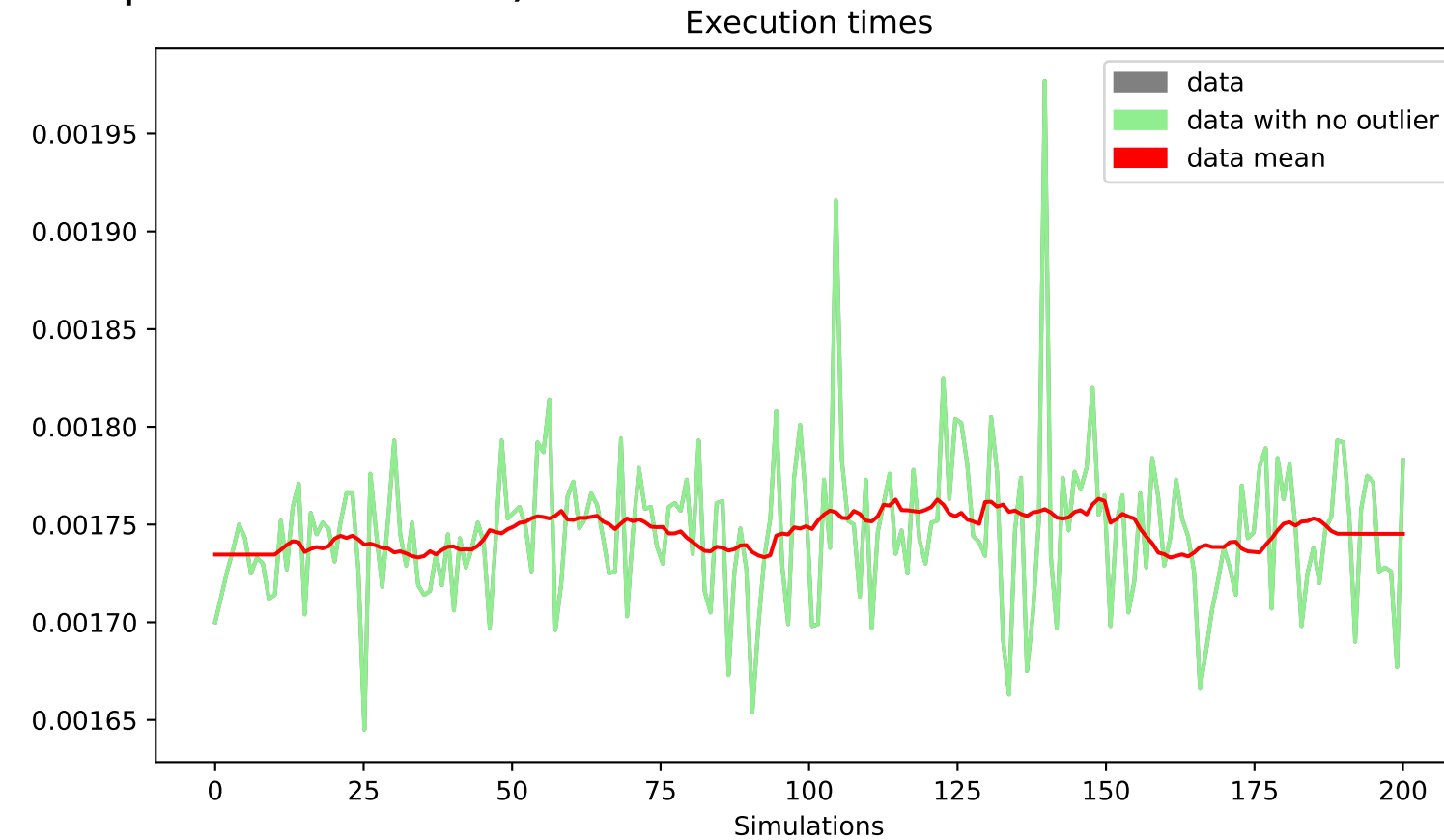
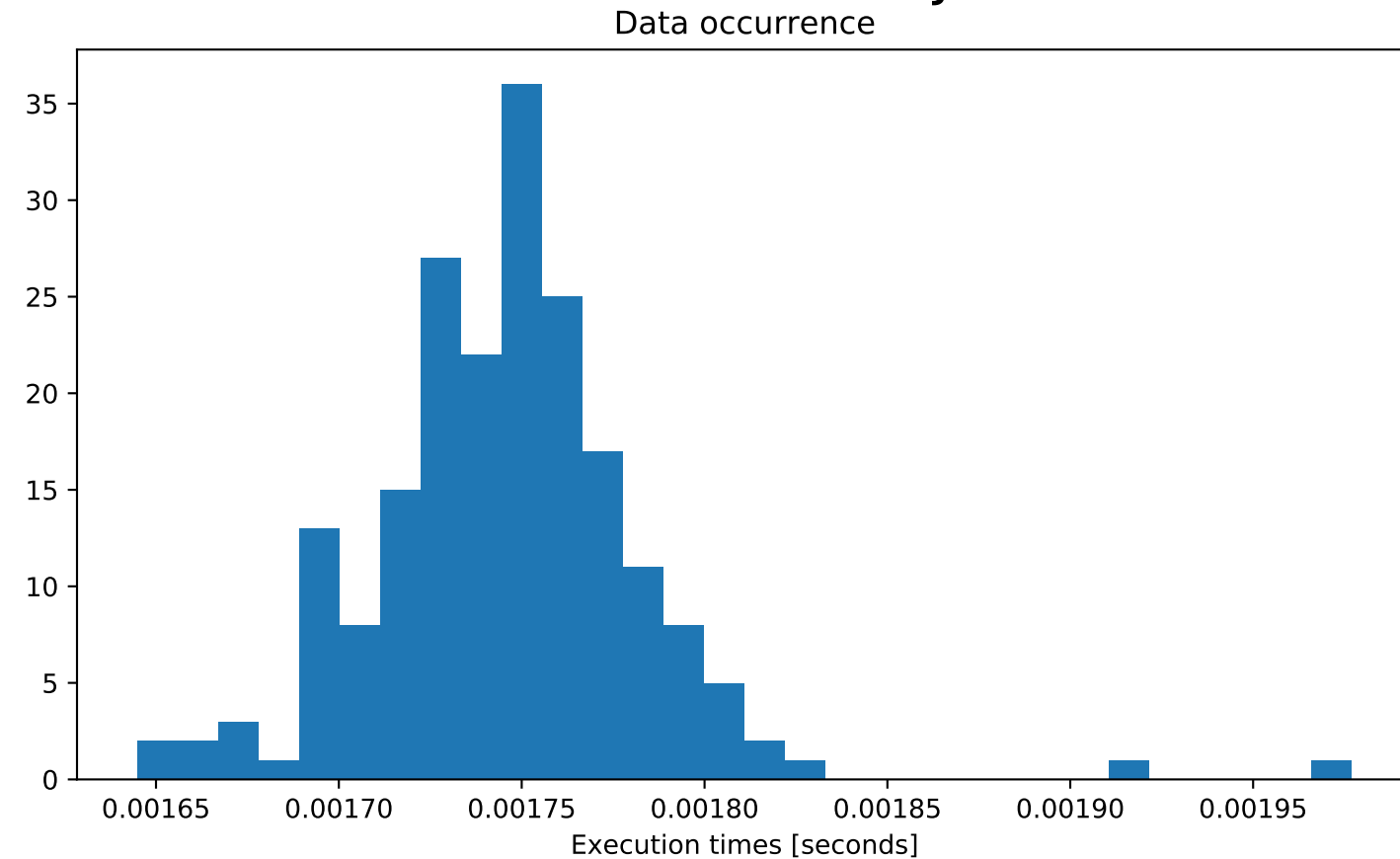
# GESUMMV (IID with p-value 0.768)



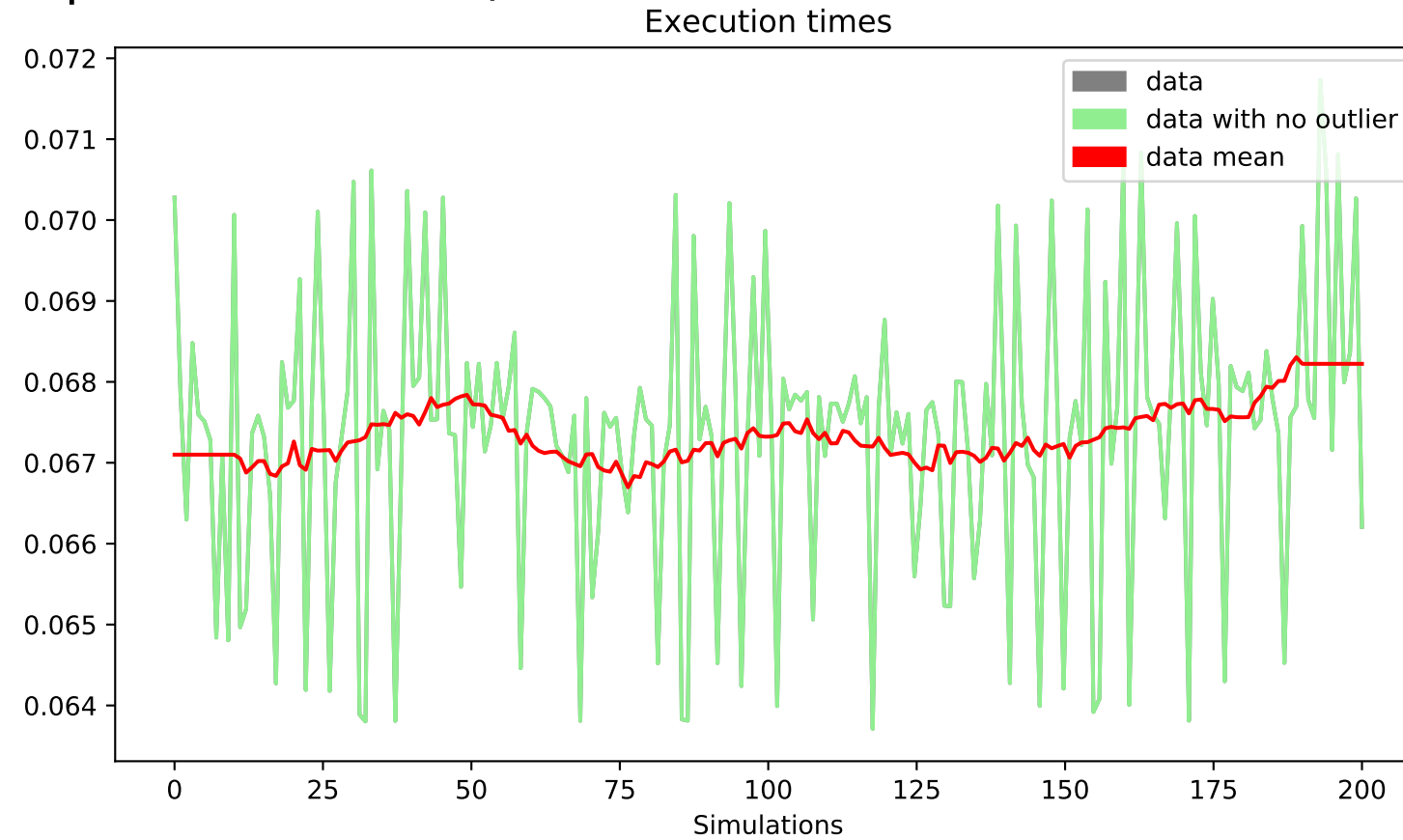
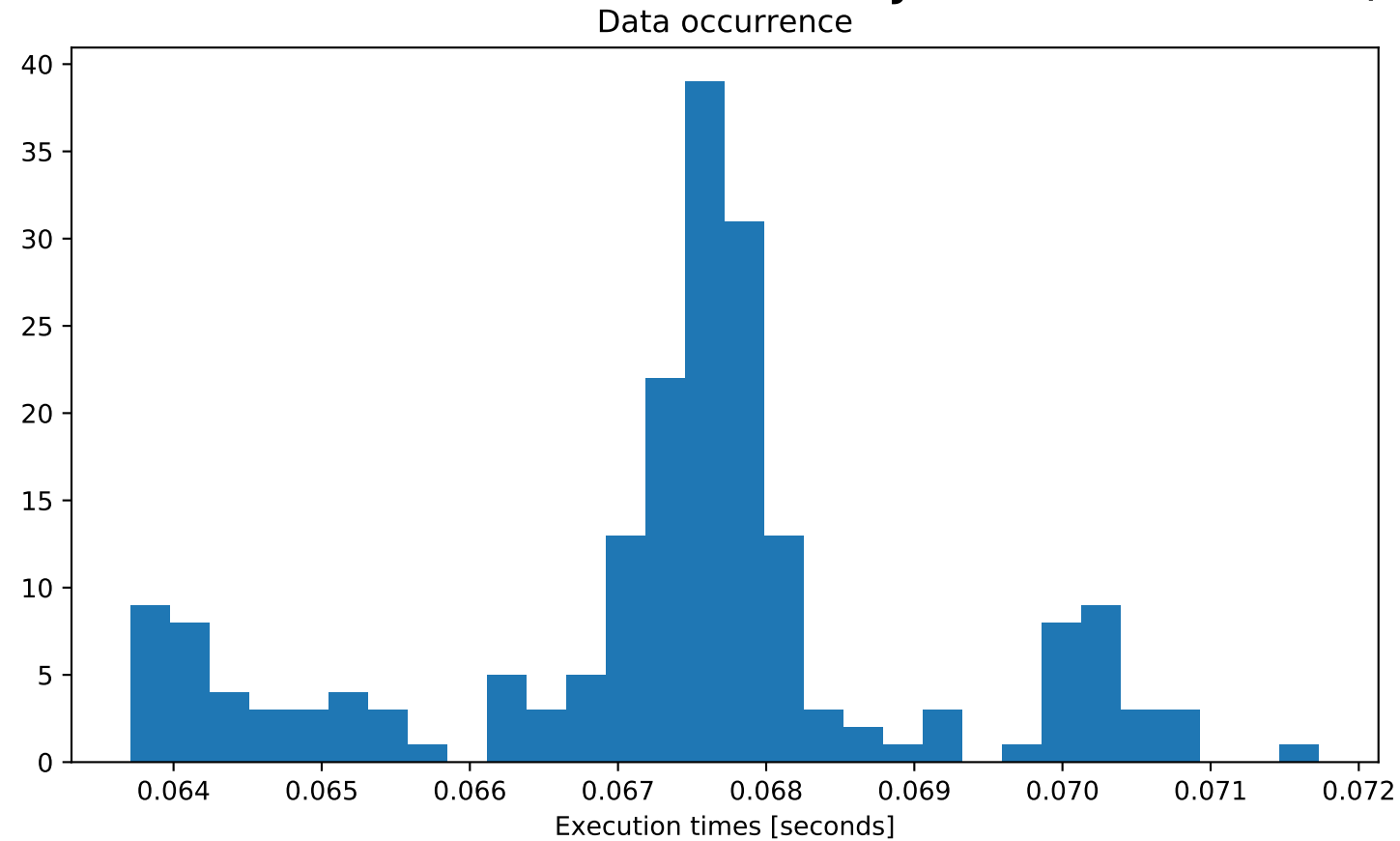
# GRAMSCHMIDT (IID with p-value 0.2632)



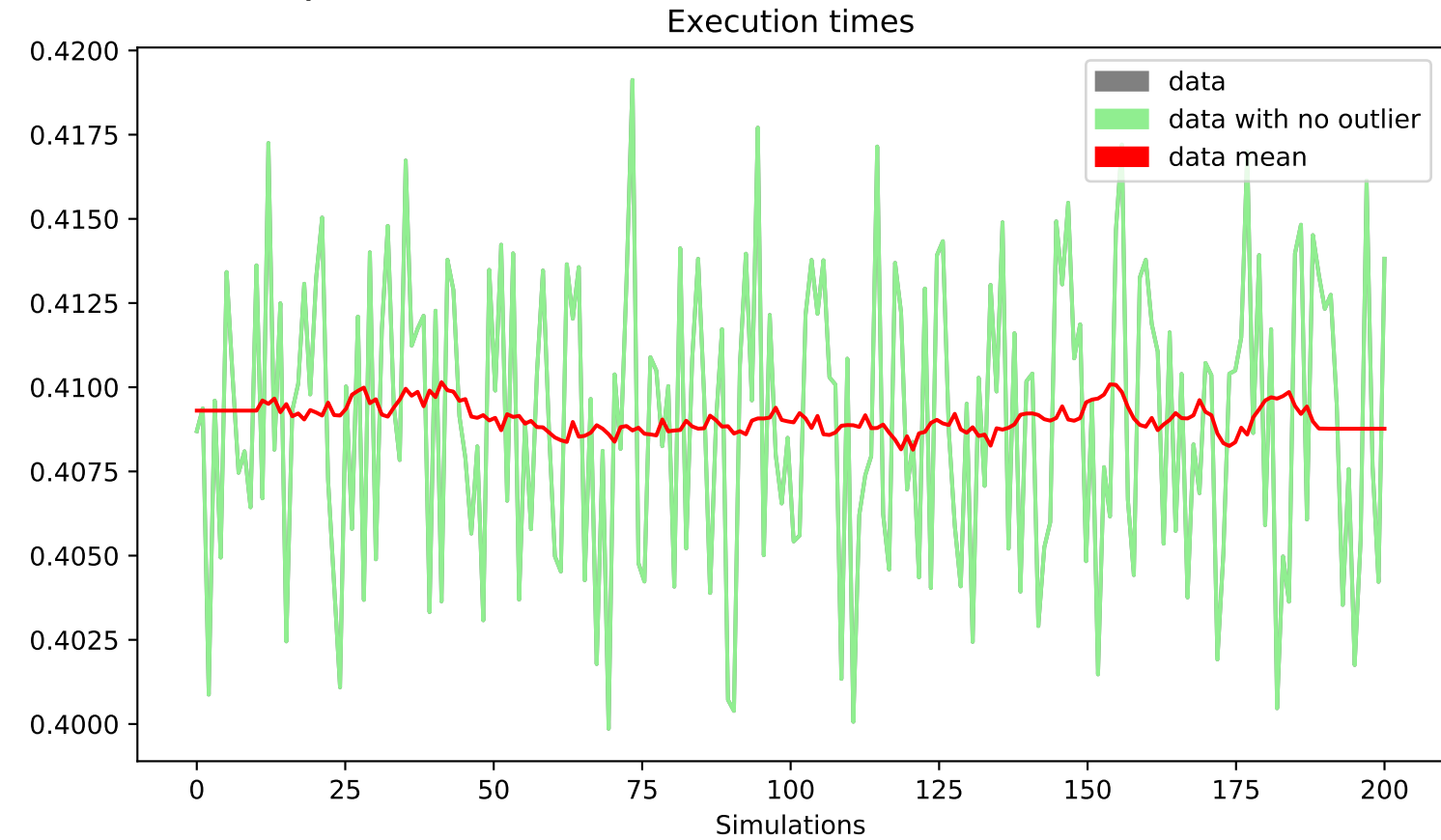
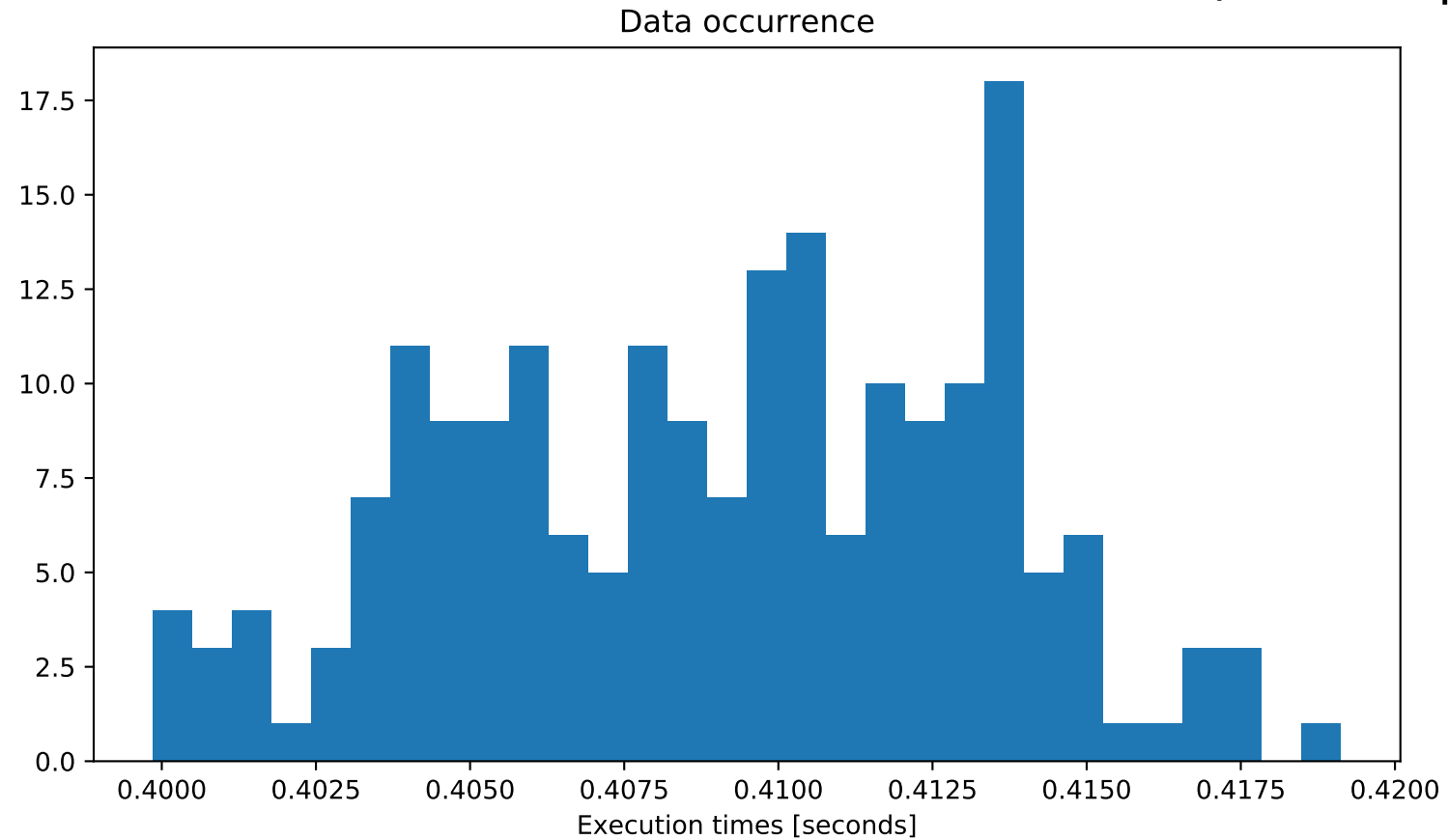
# JACOBI-1D-IMPER (IID with p-value 0.4365)



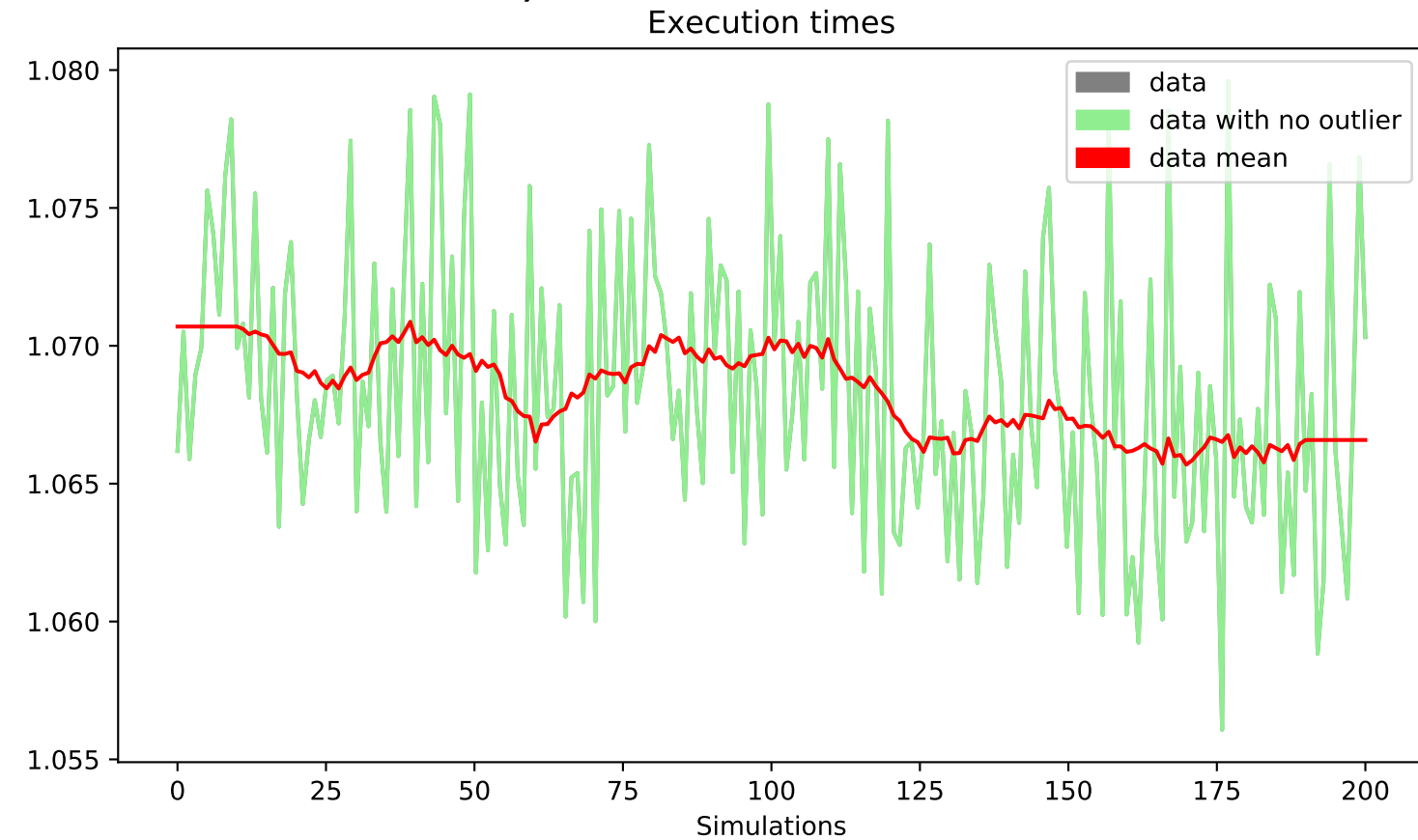
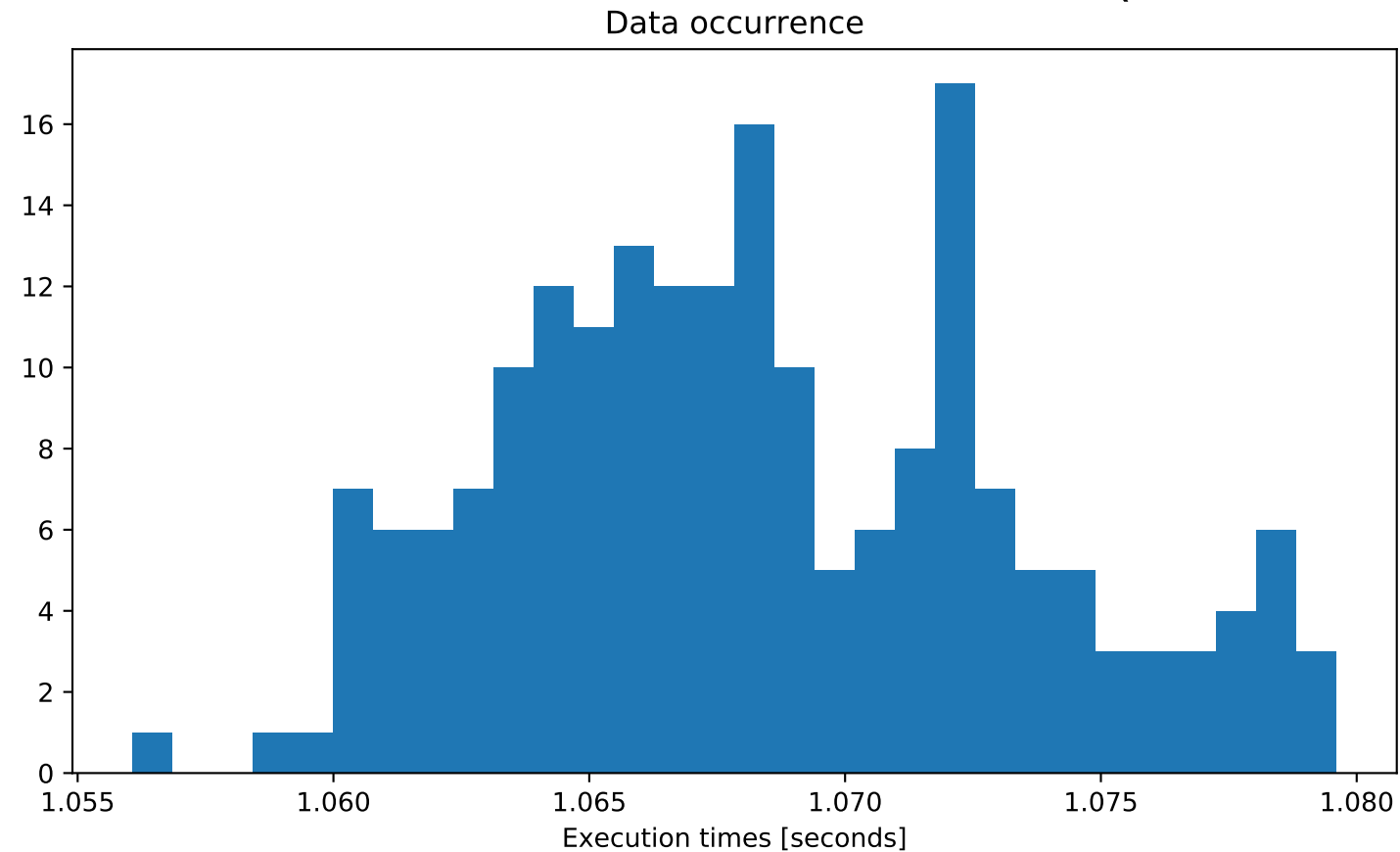
# JACOBI-2D-IMPER (IID with p-value 0.55625)



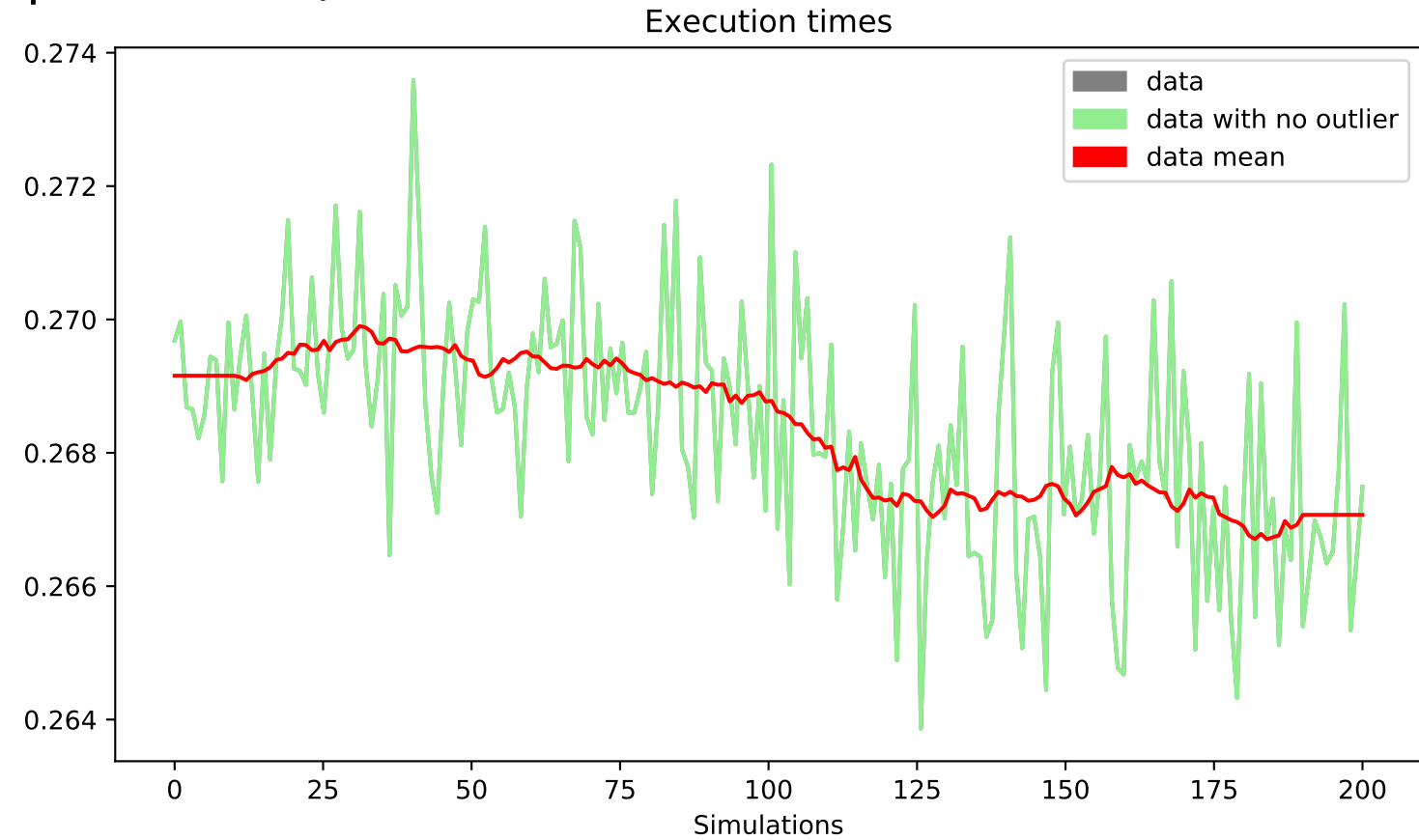
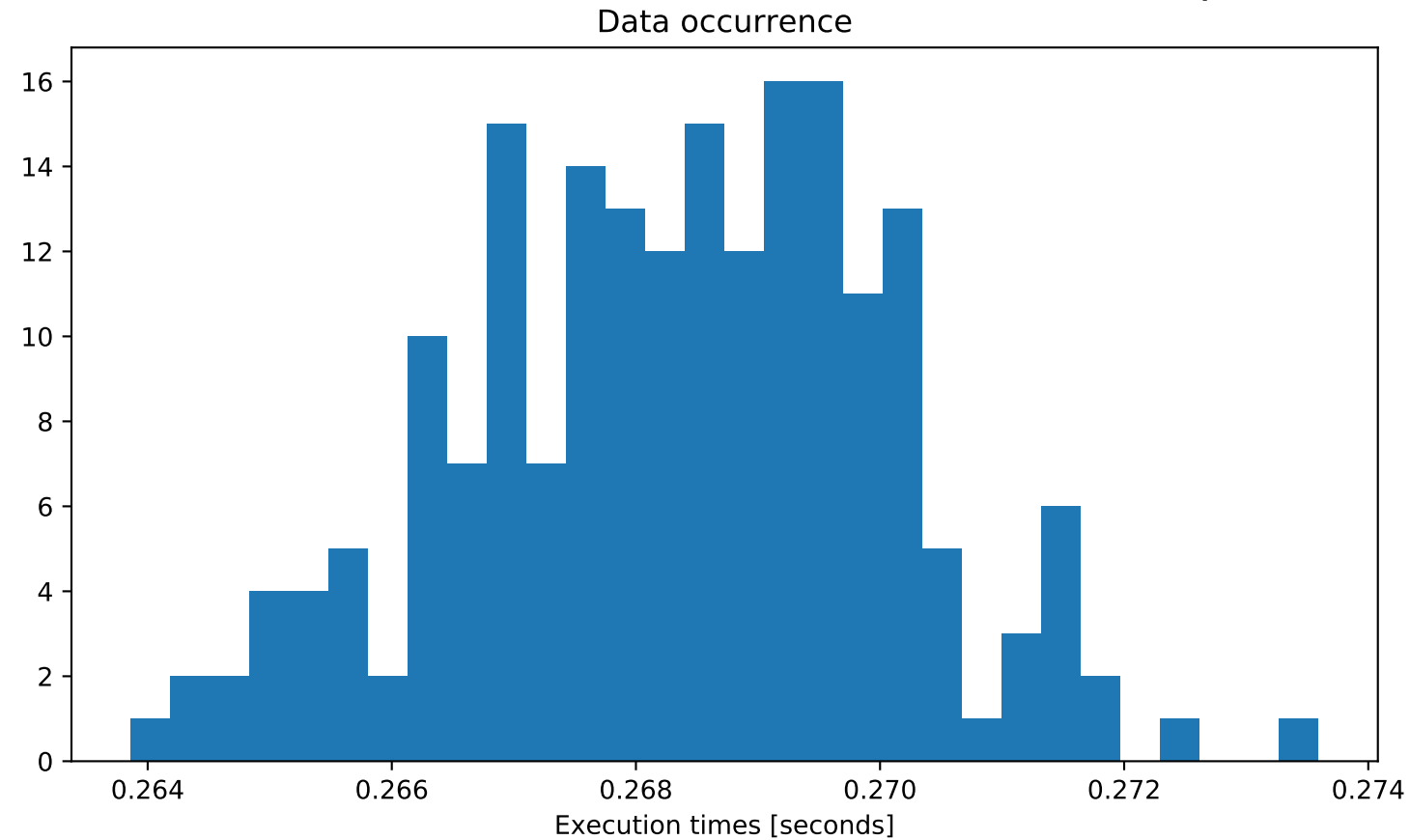
# LU (IID with p-value 0.8186)



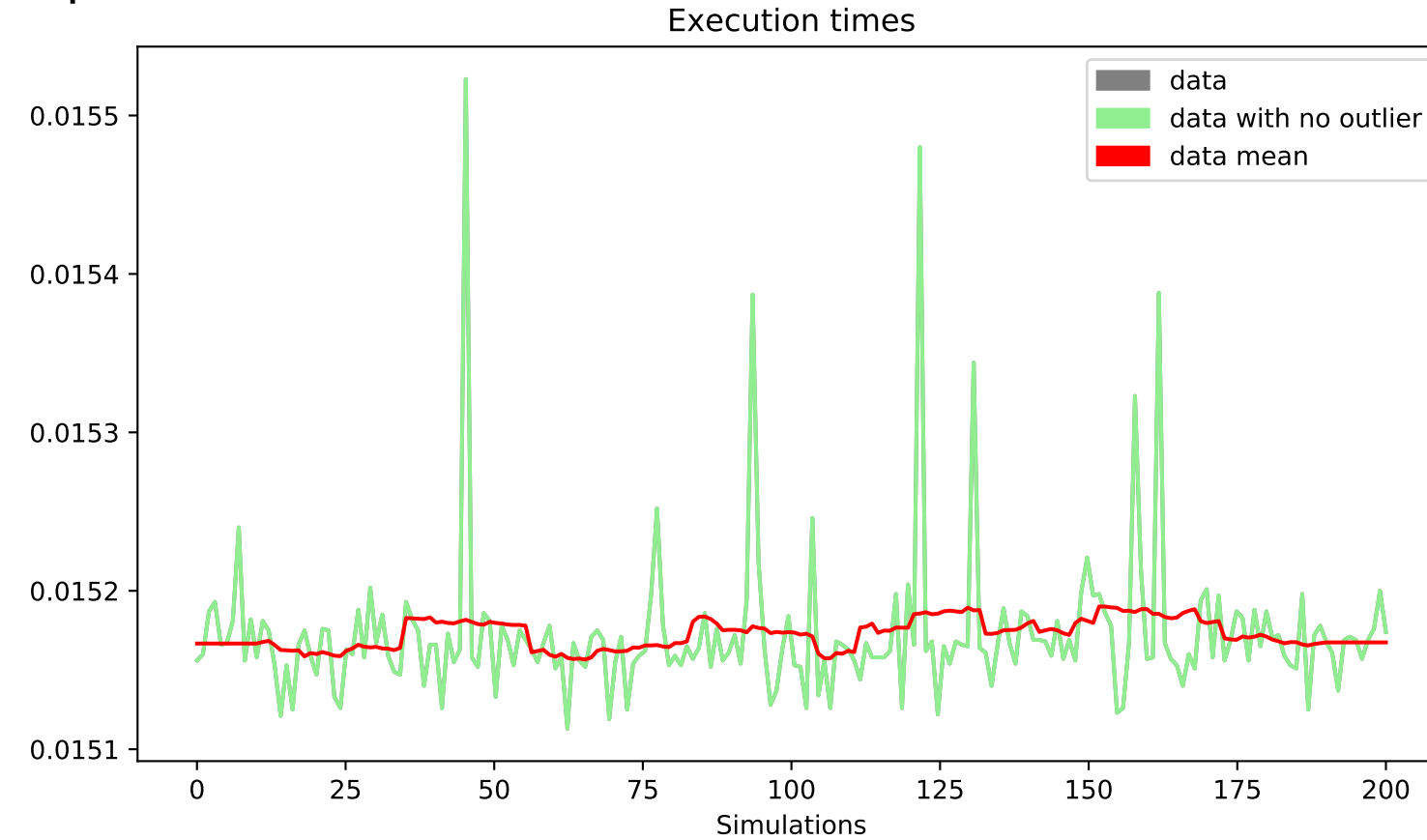
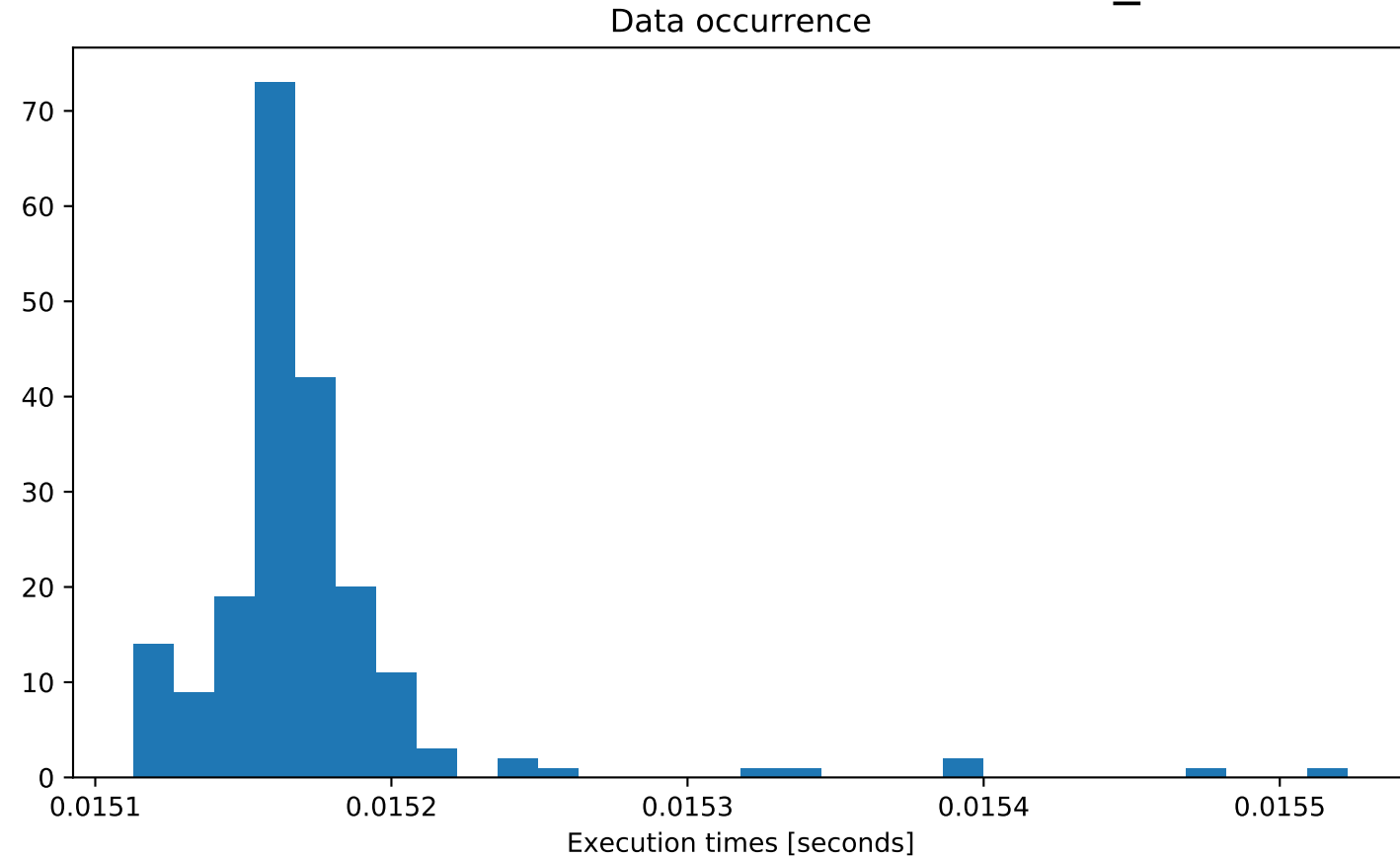
# LUDCMP (NOT IID with p-value 0.0243333333)



# MVT (NOT IID with p-value 0.0)

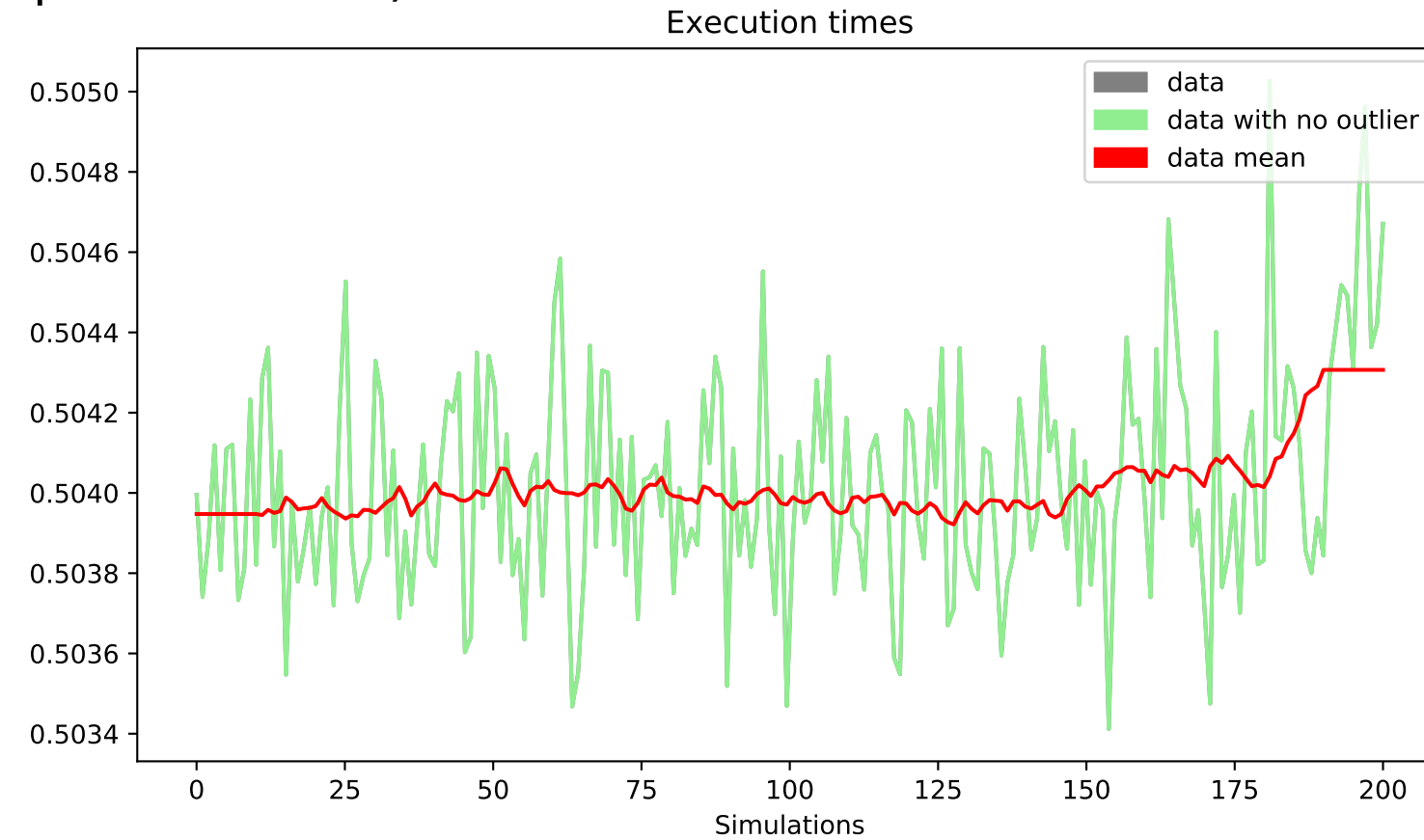
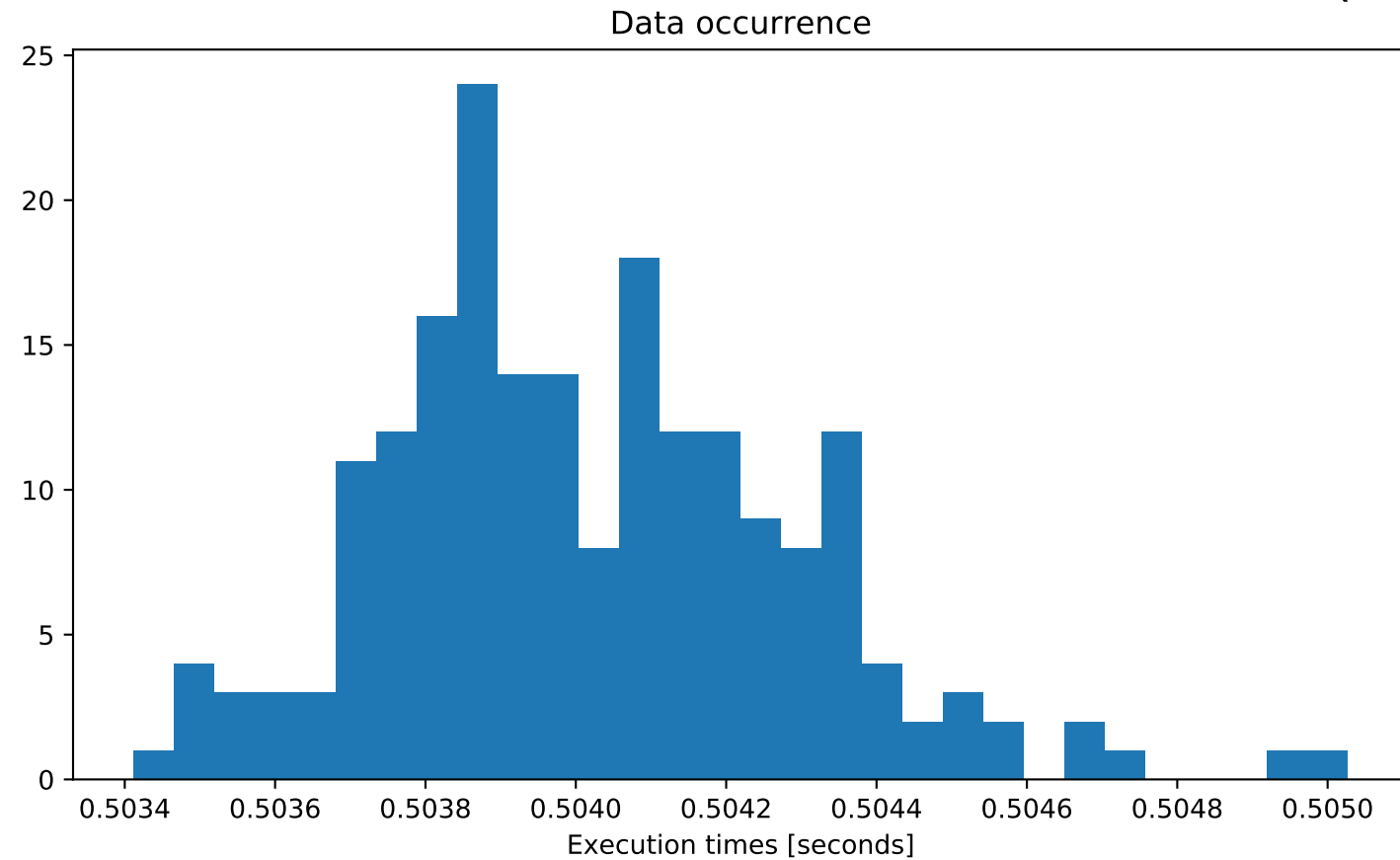


# REG\_DETECT (IID with p-value 0.533)

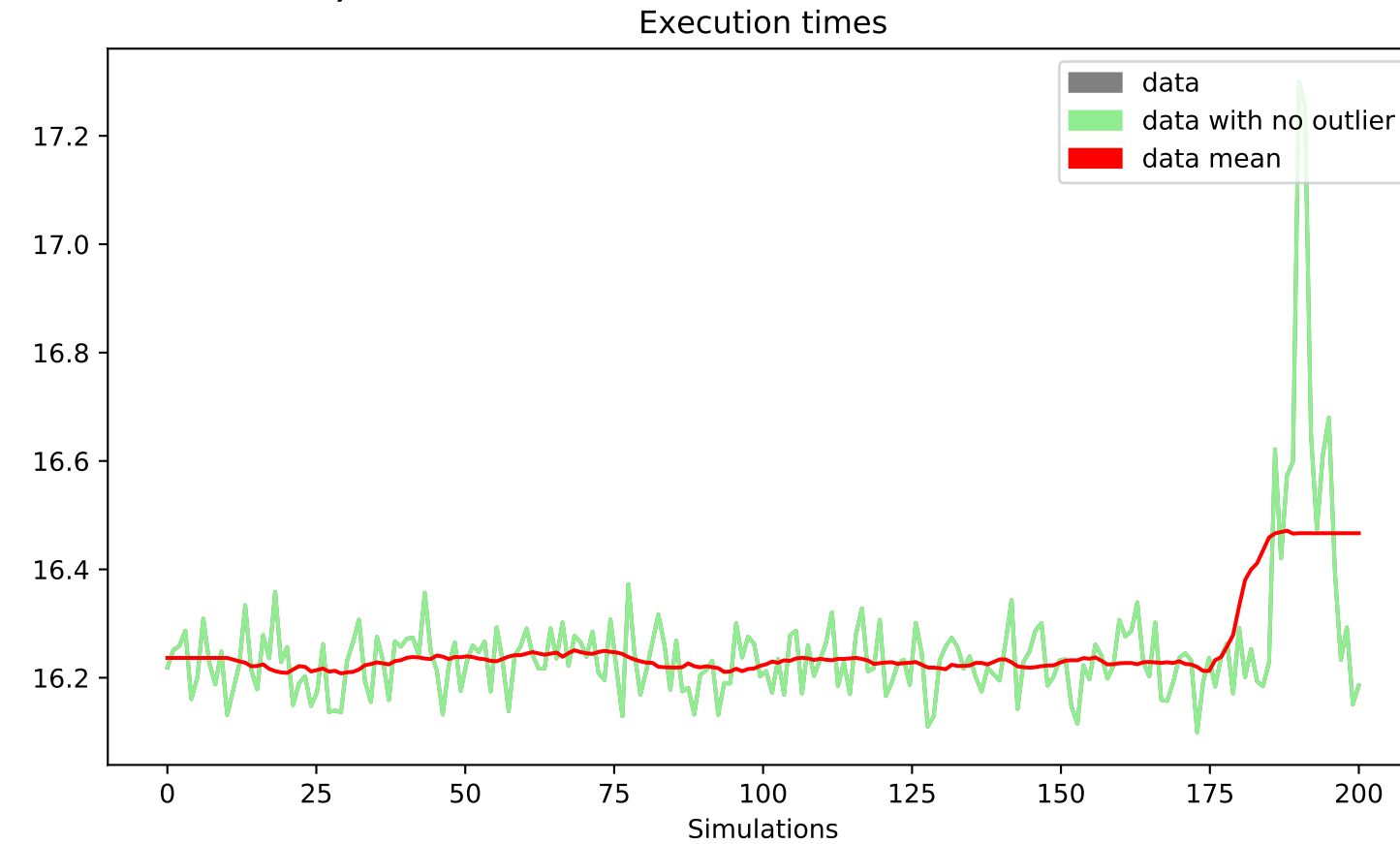
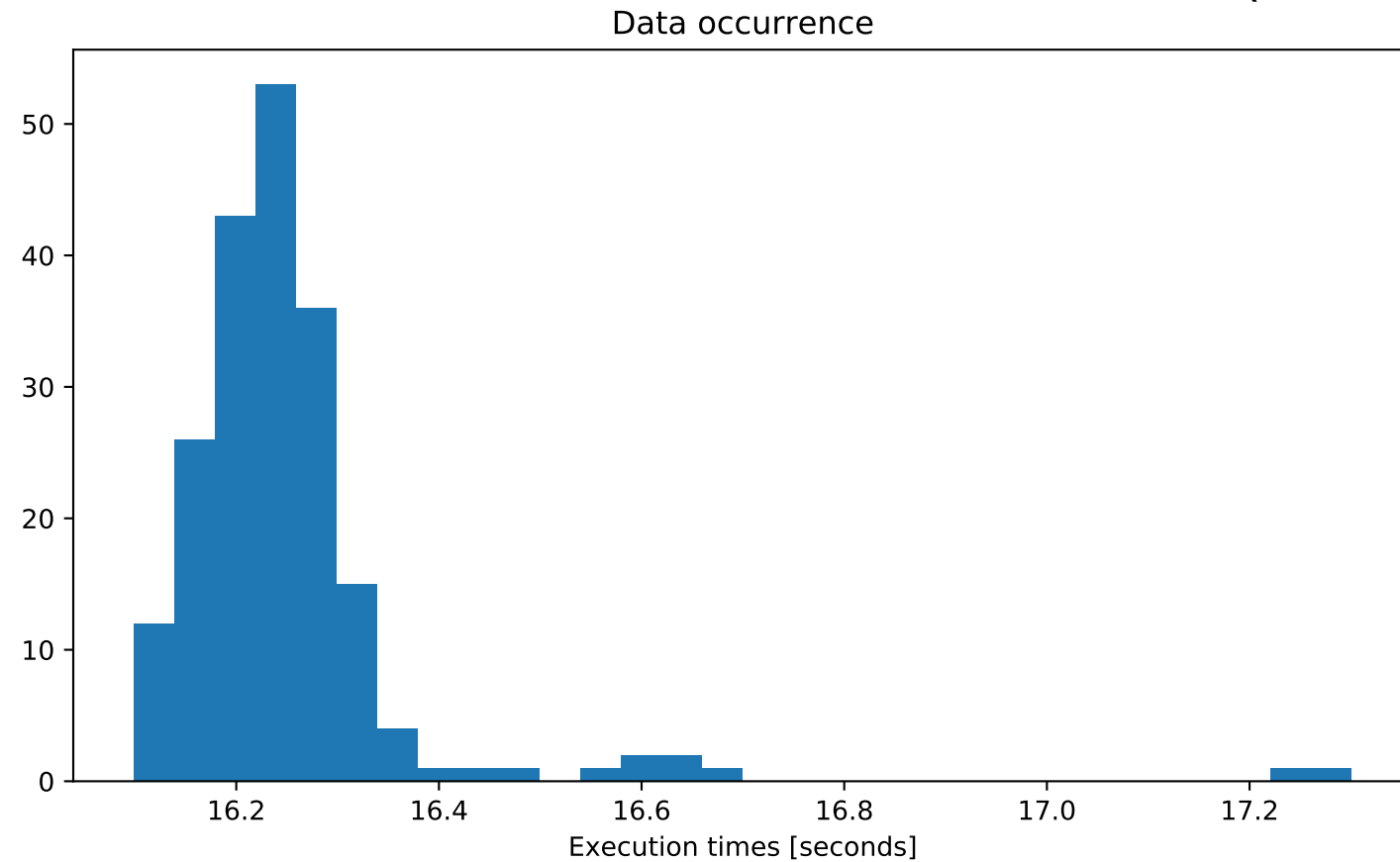




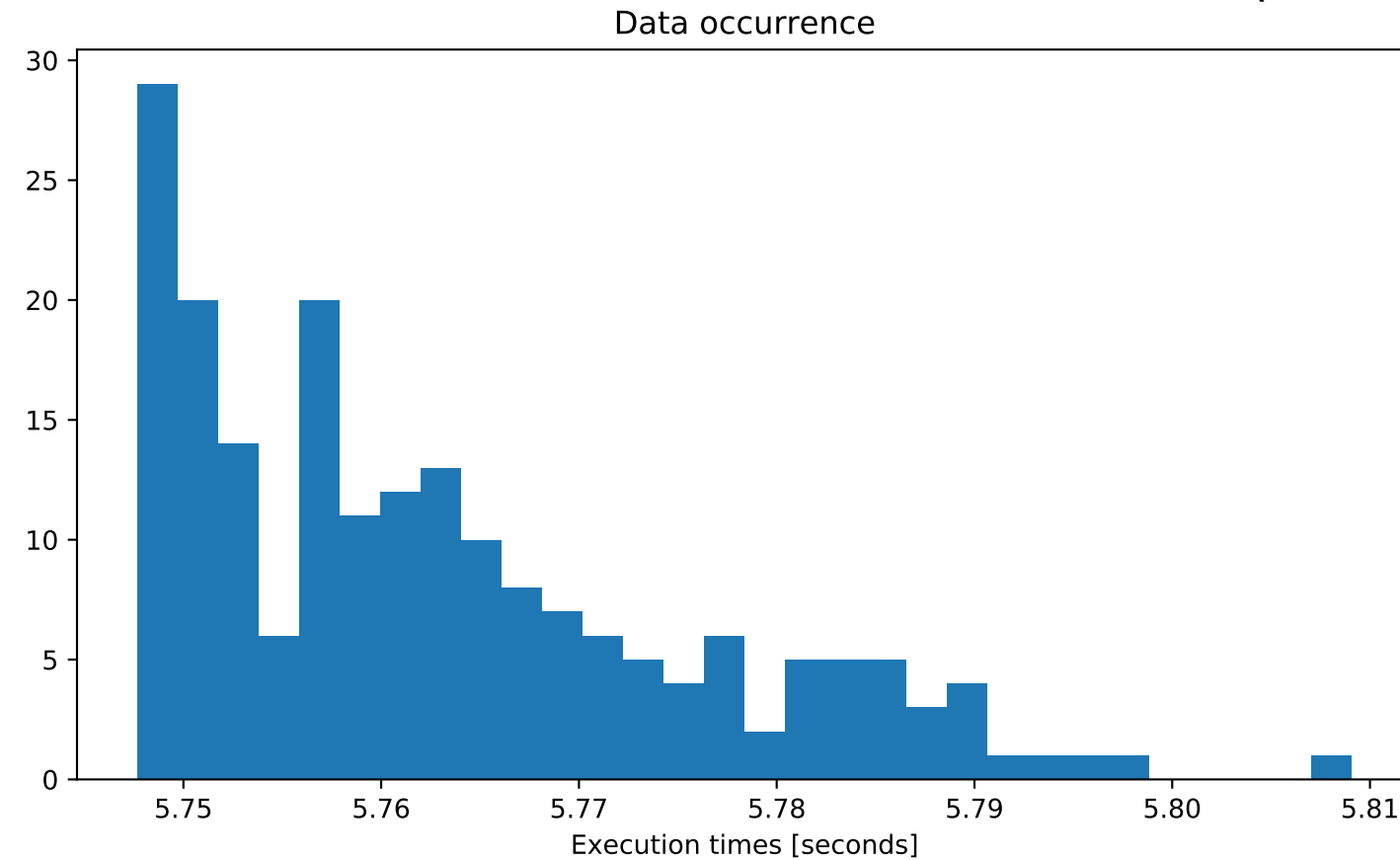
# SEIDEL-2D (IID with p-value 0.399)



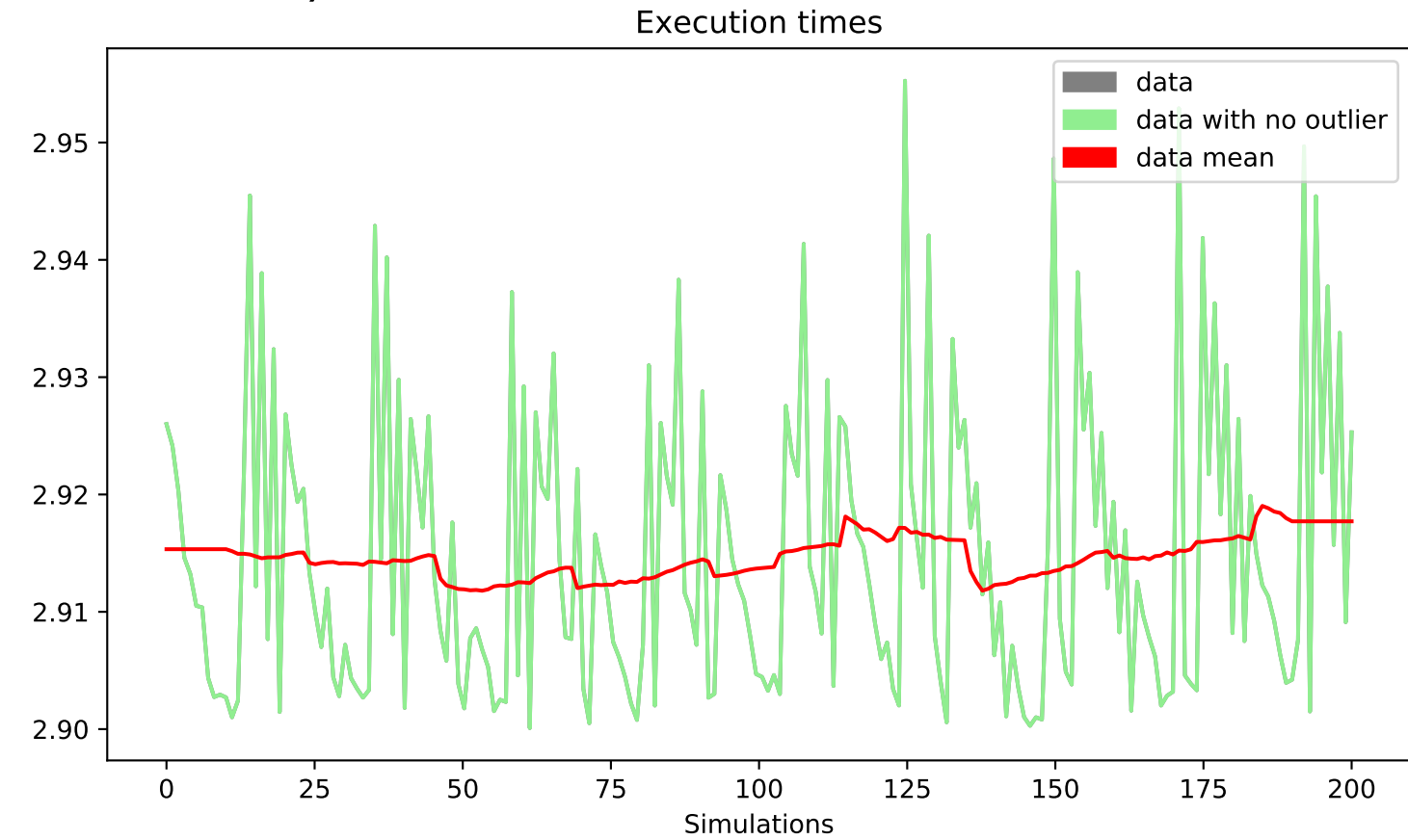
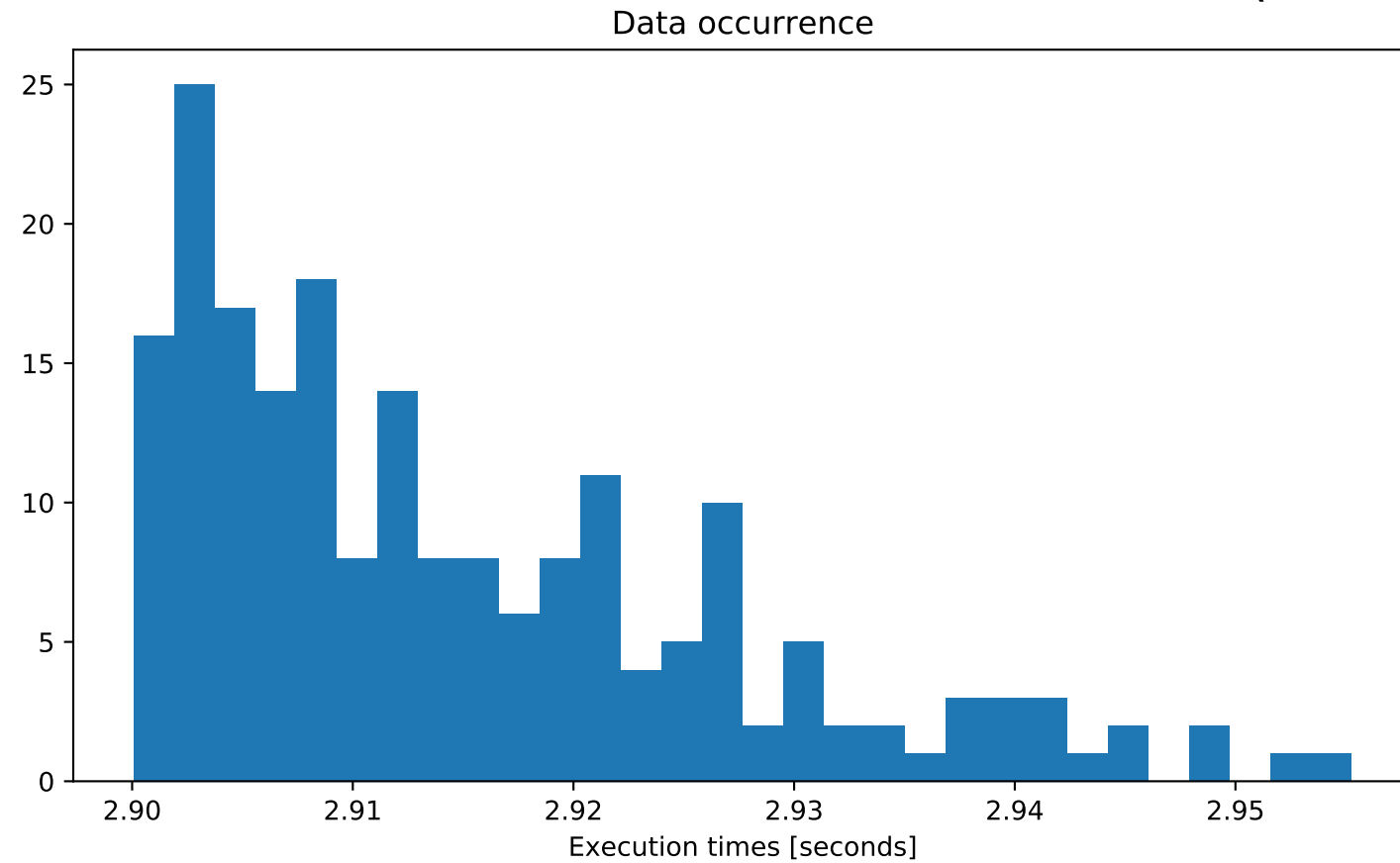
# SYMM (IID with p-value 0.25525)



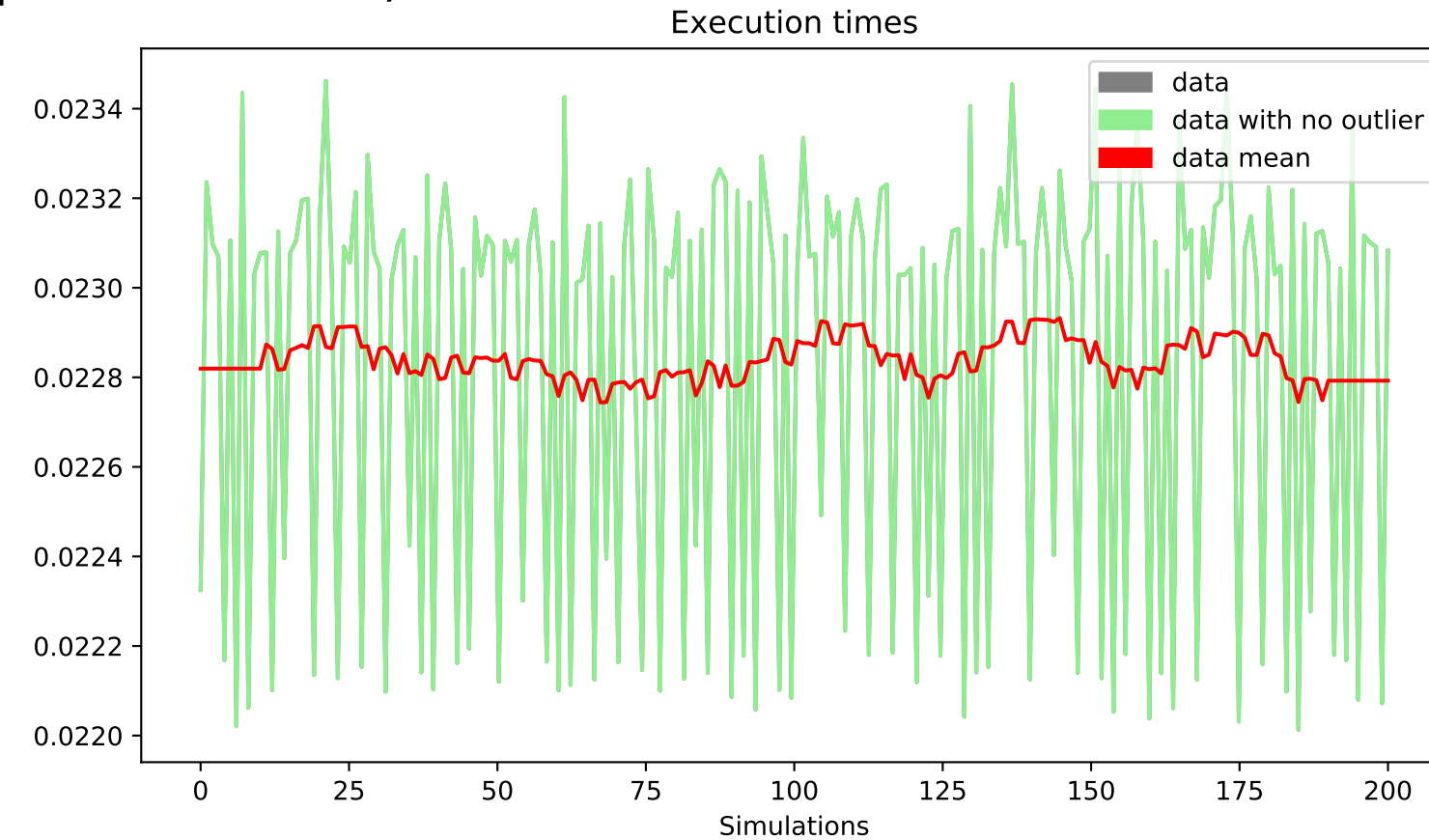
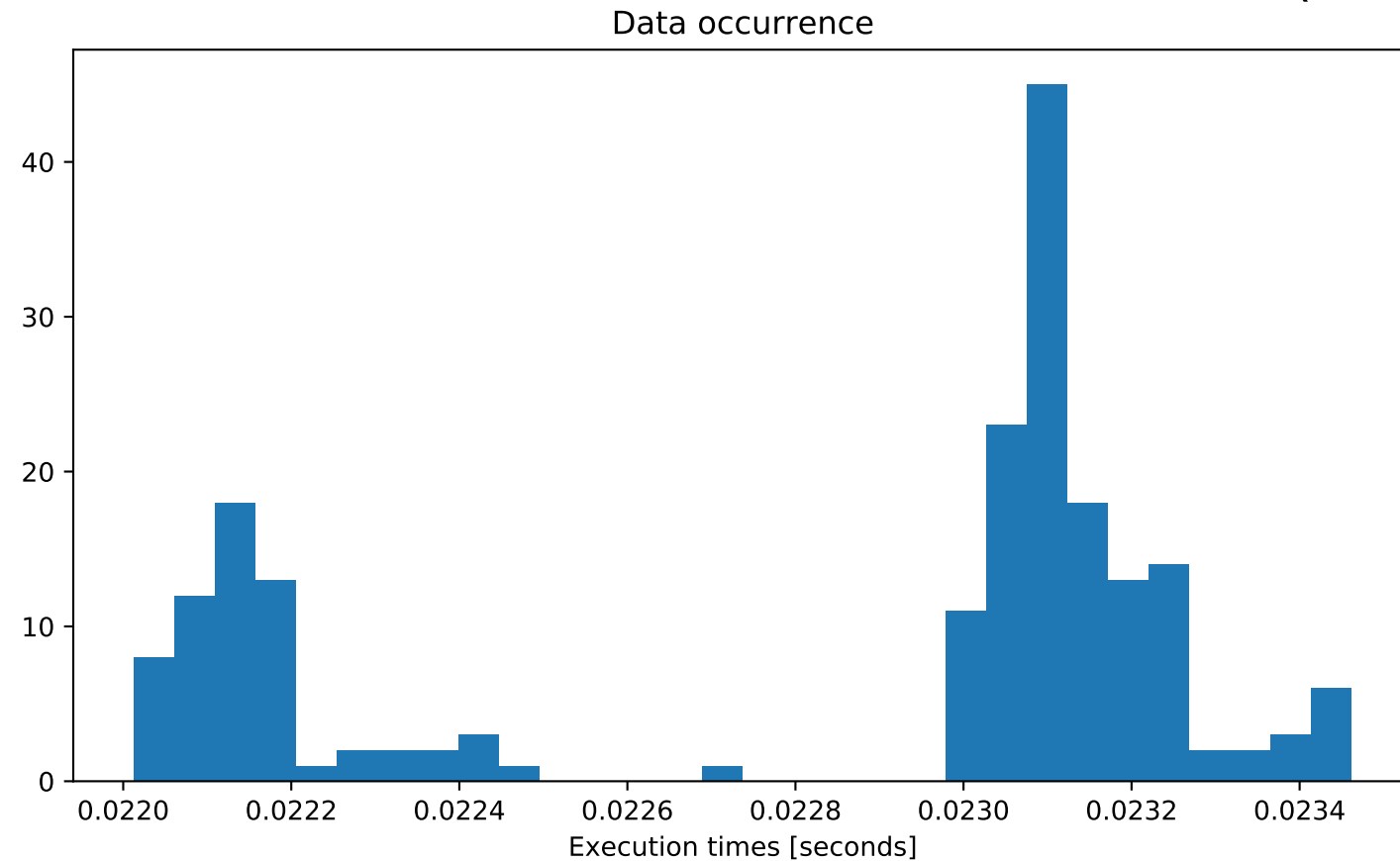
# SYR2K (IID with p-value 0.47825)



# SYRK (IID with p-value 0.5562)



# TRISOLV (IID with p-value 0.8834)



# TRMM (IID with p-value 0.5598)

