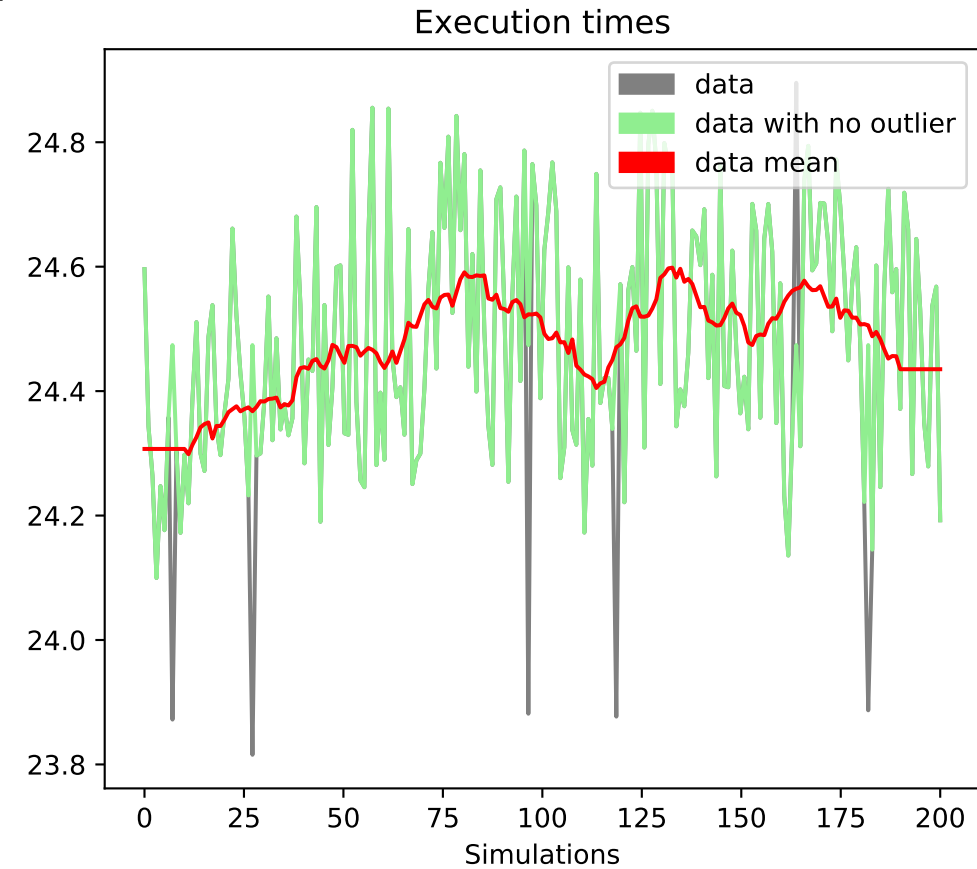
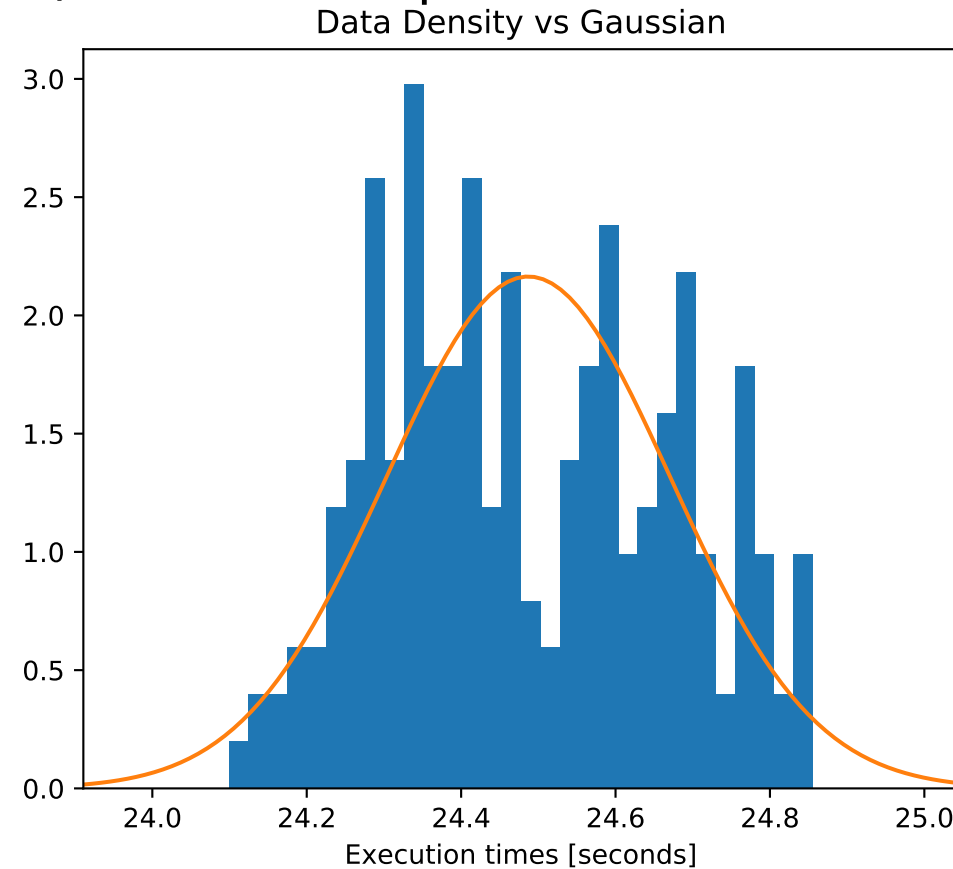
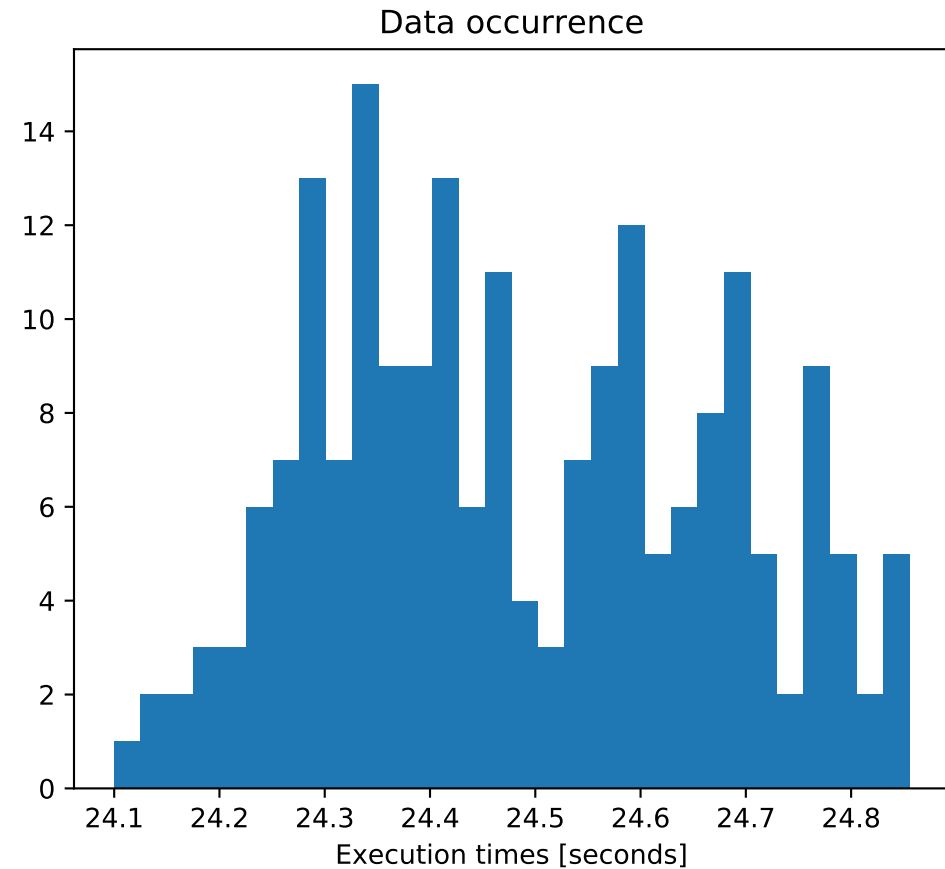
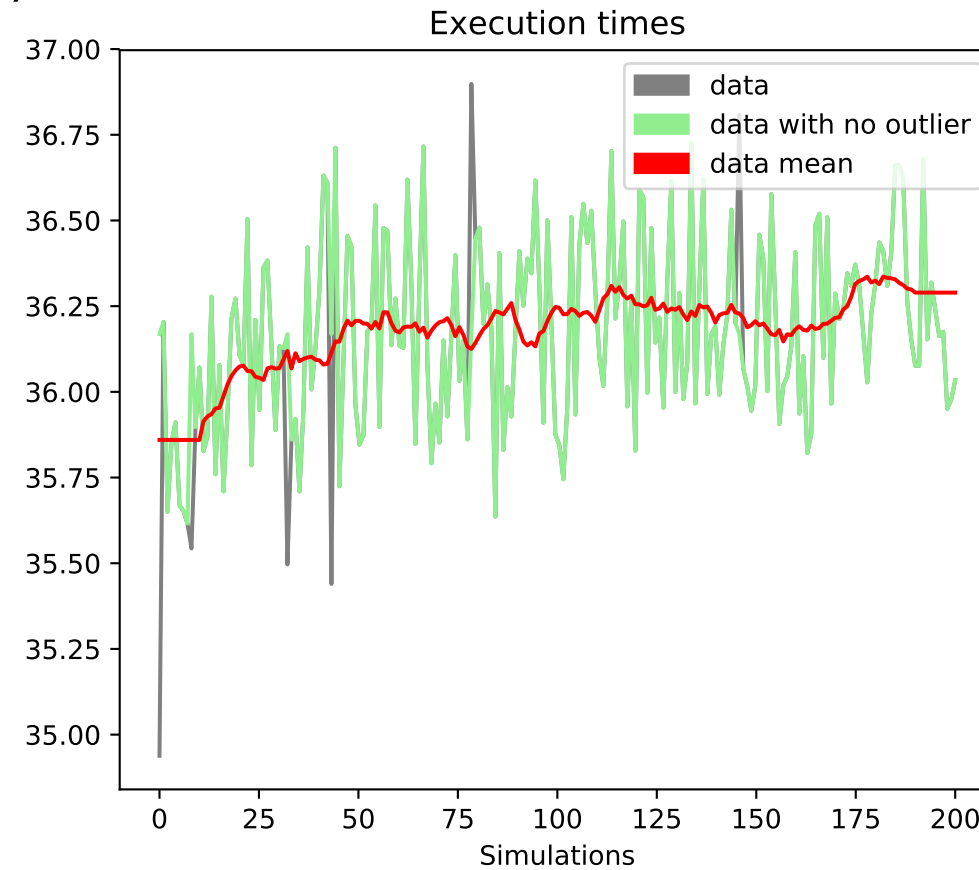
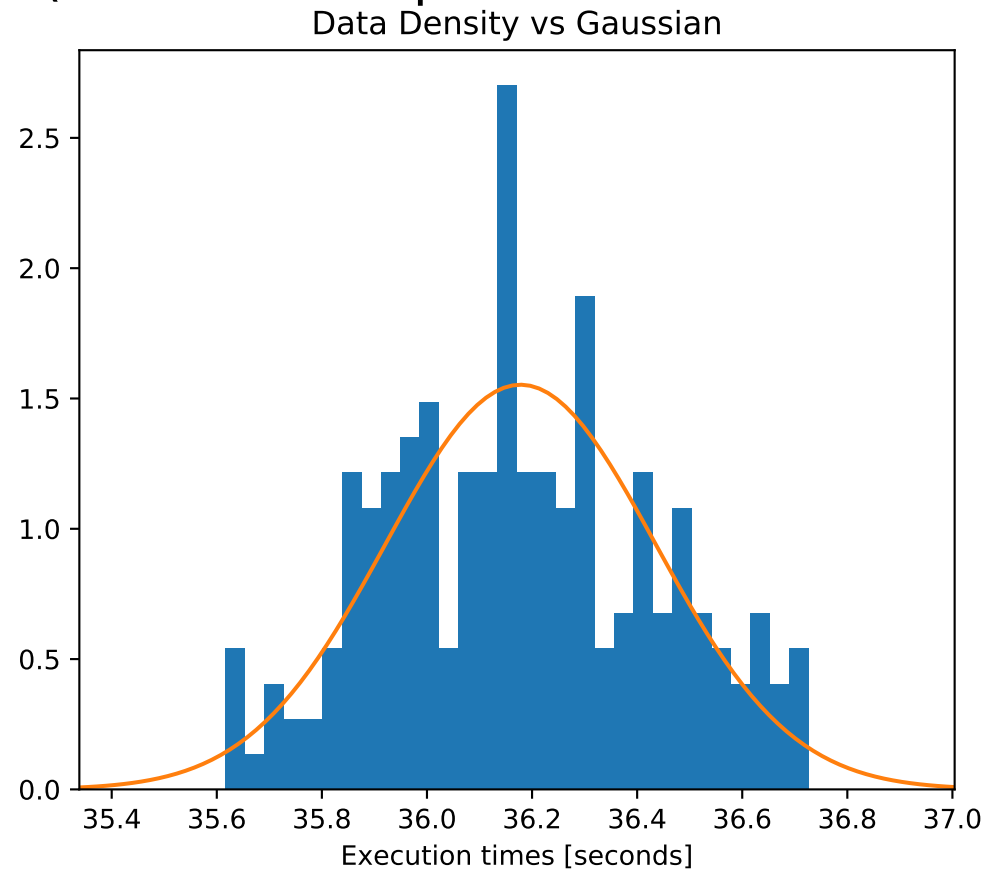
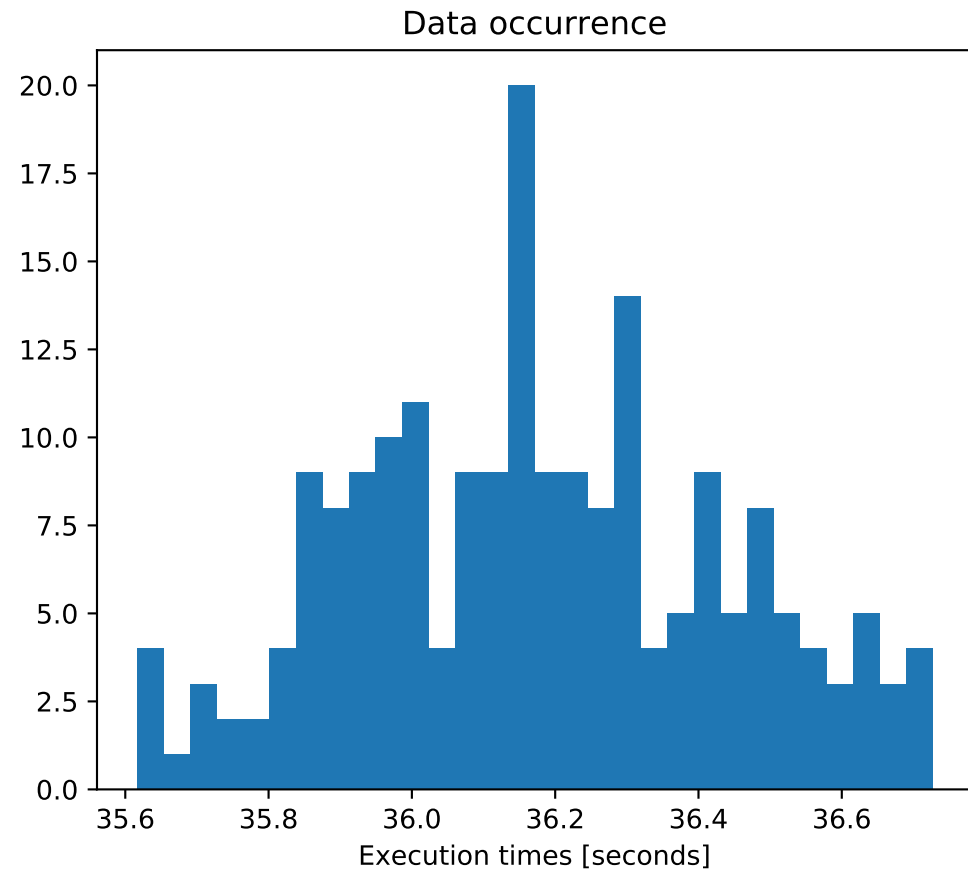


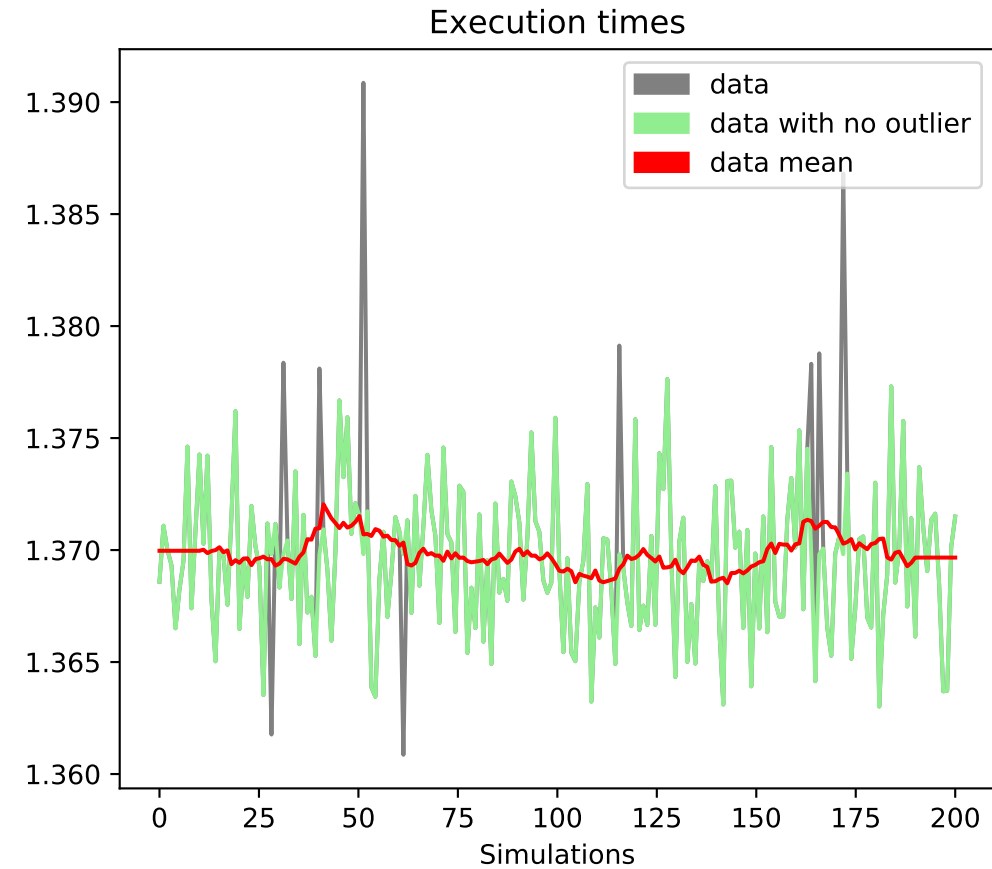
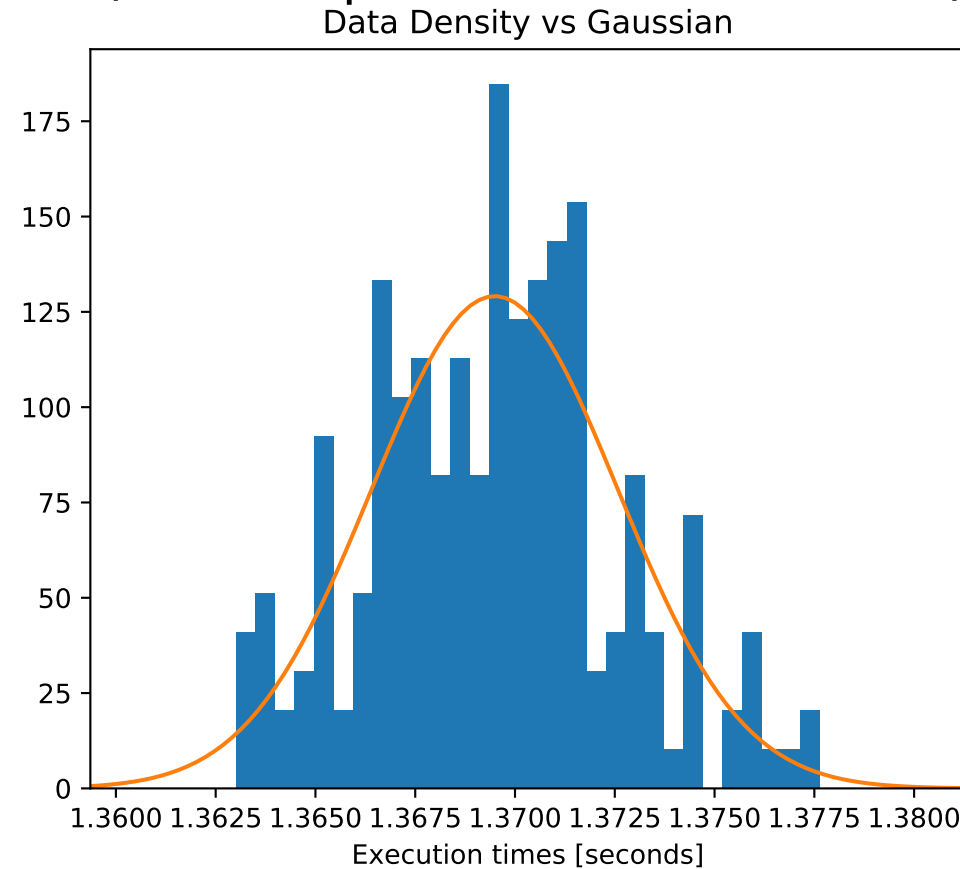
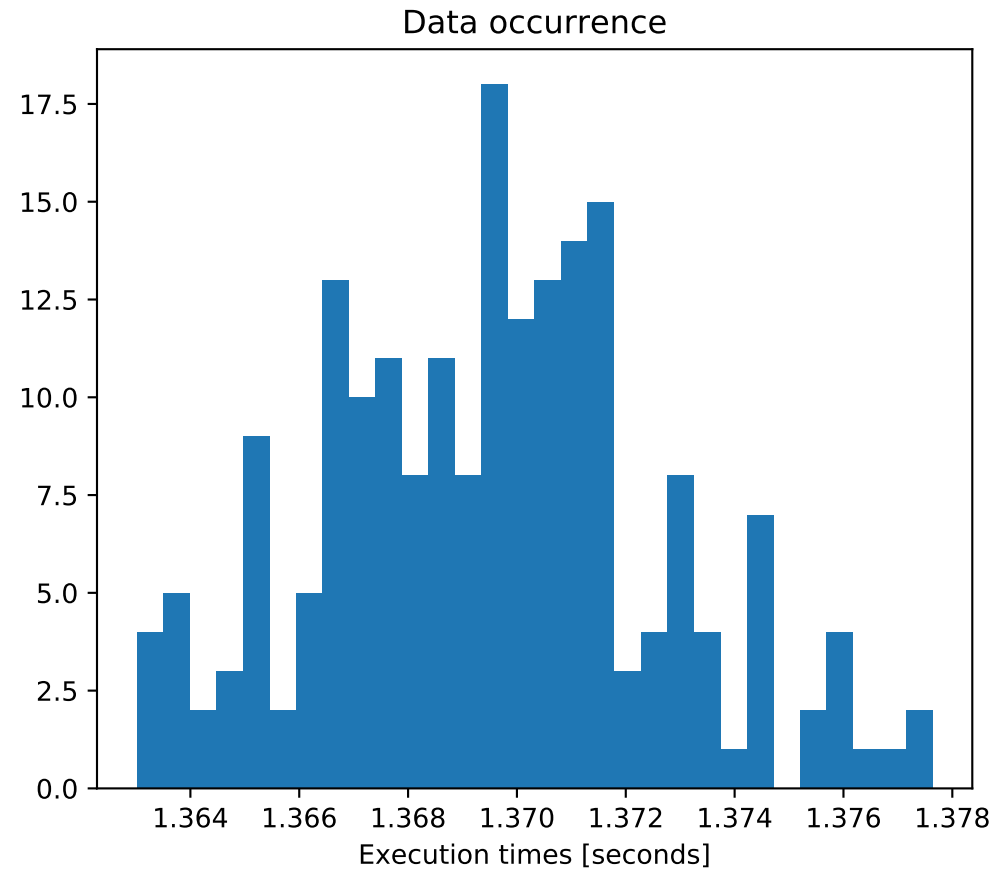
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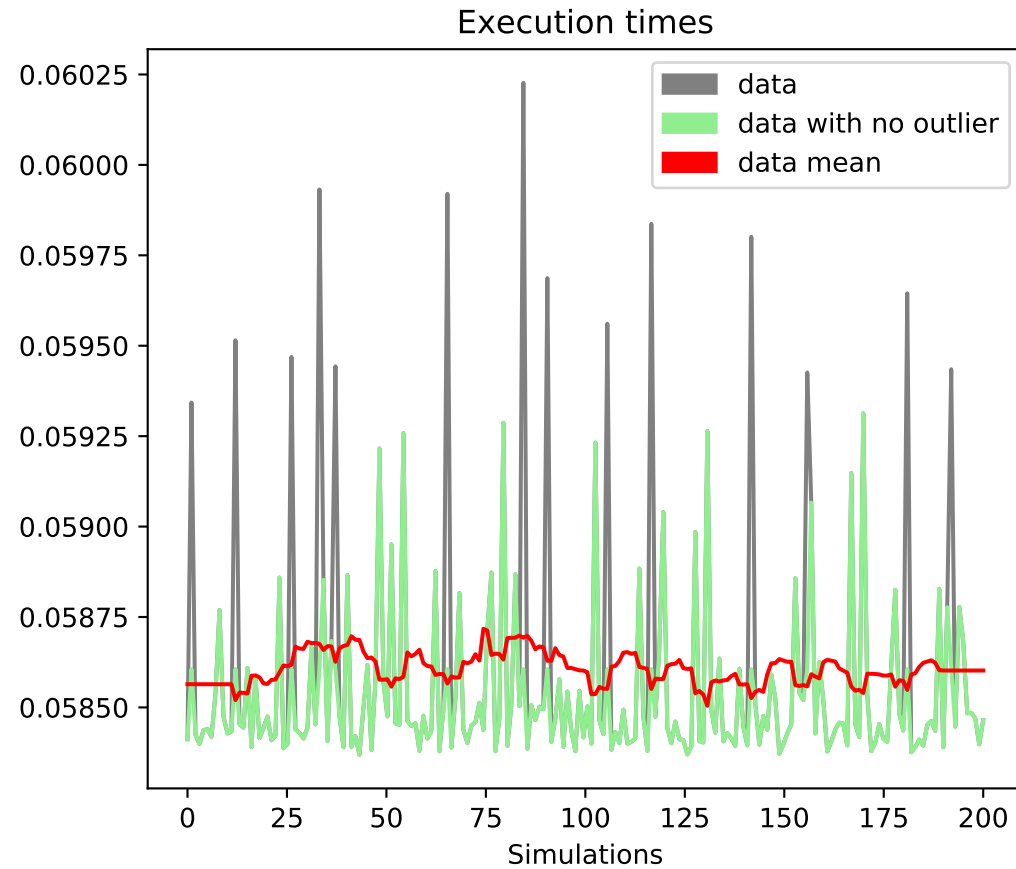
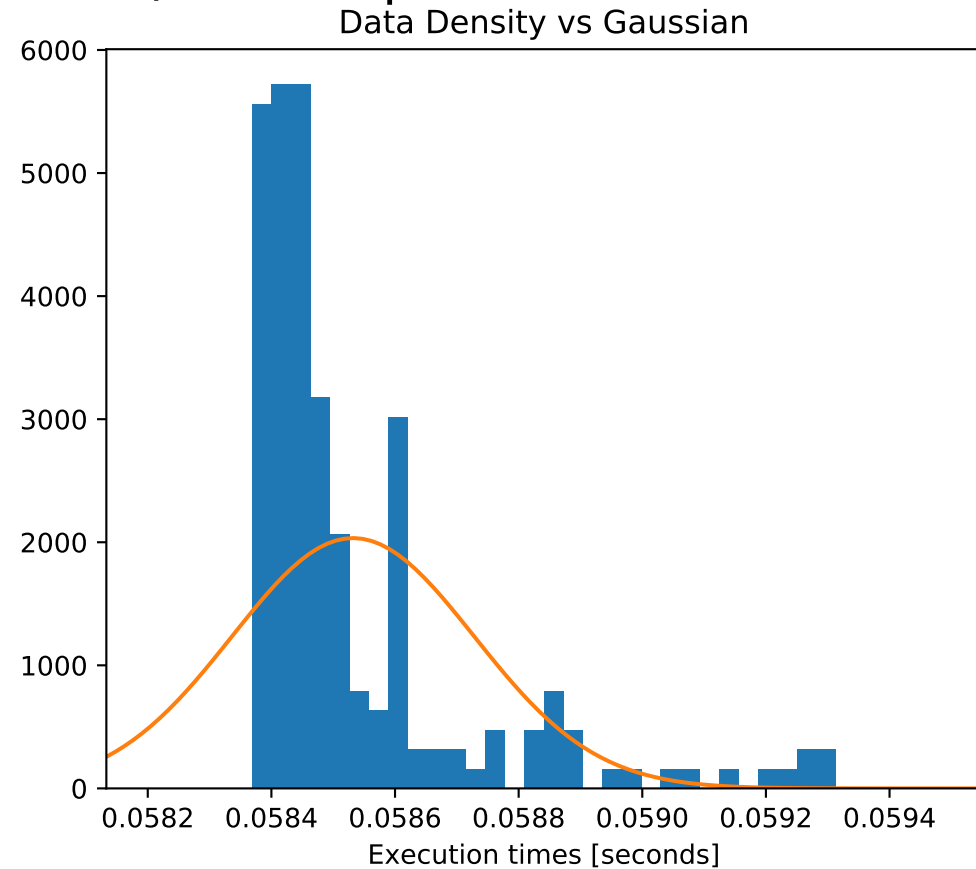
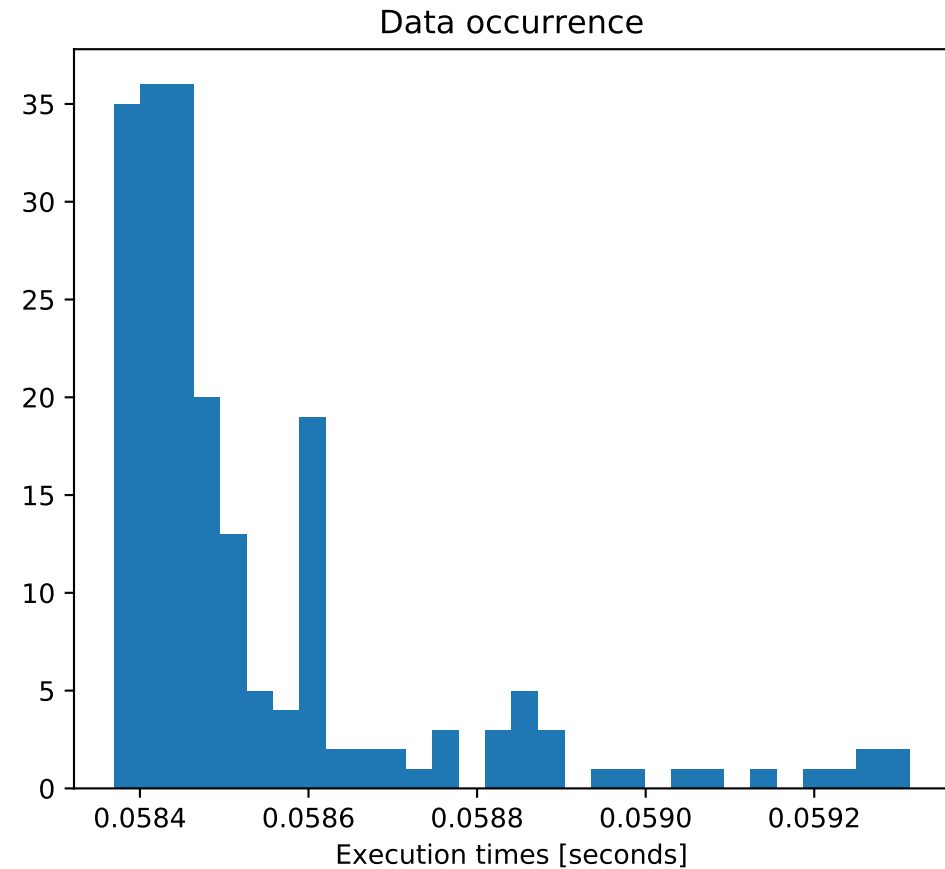
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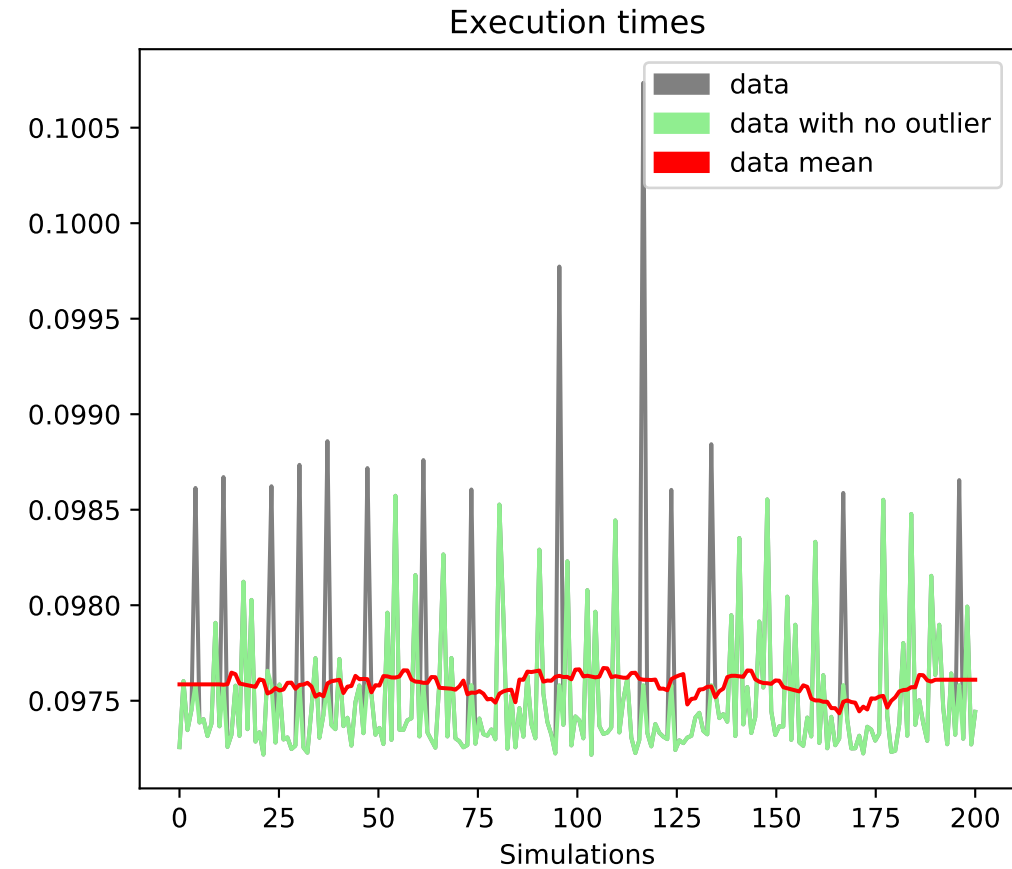
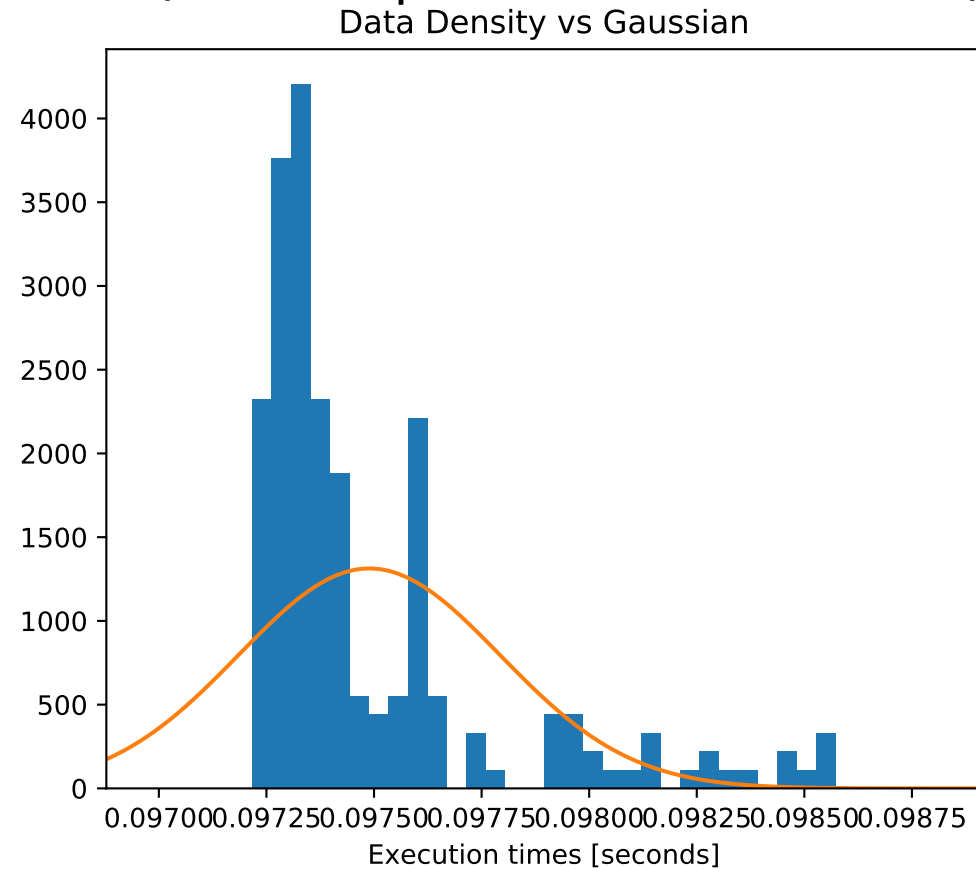
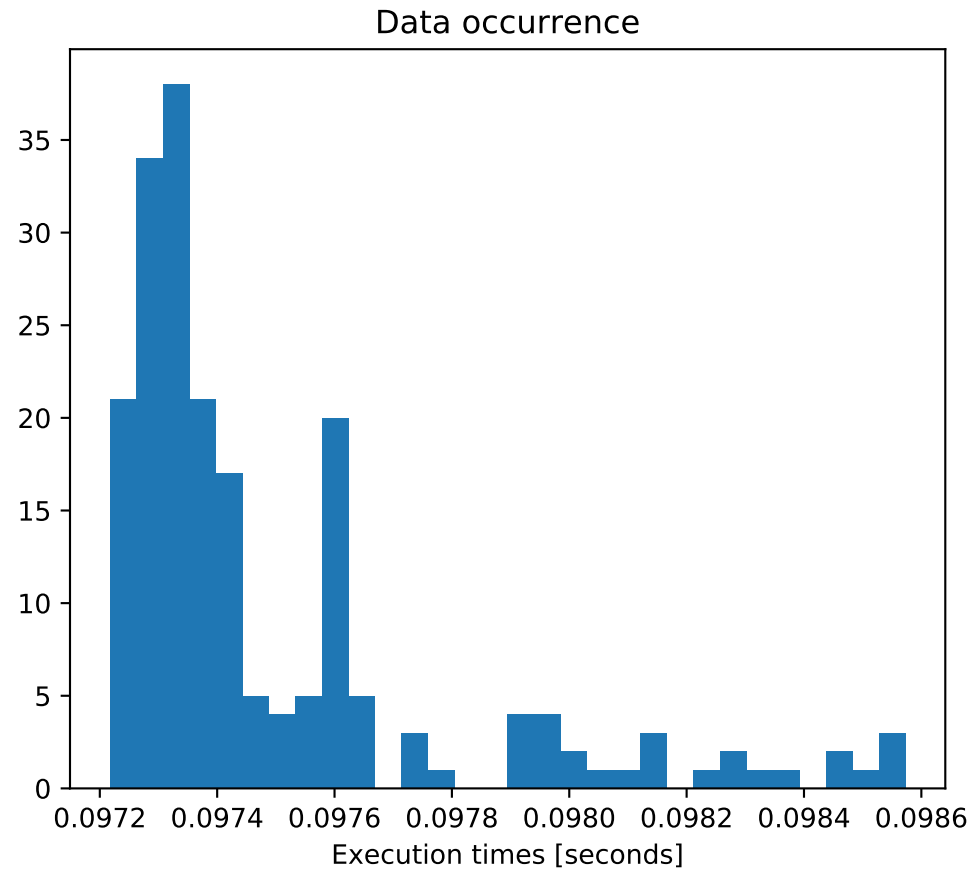
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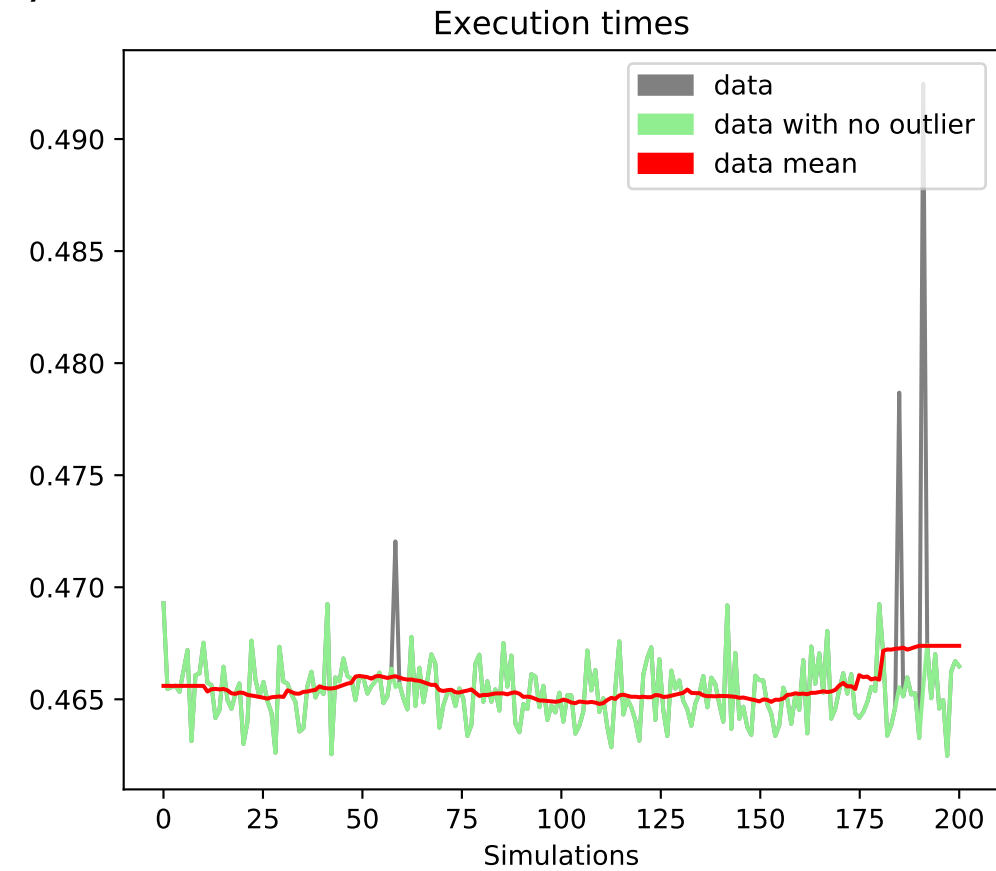
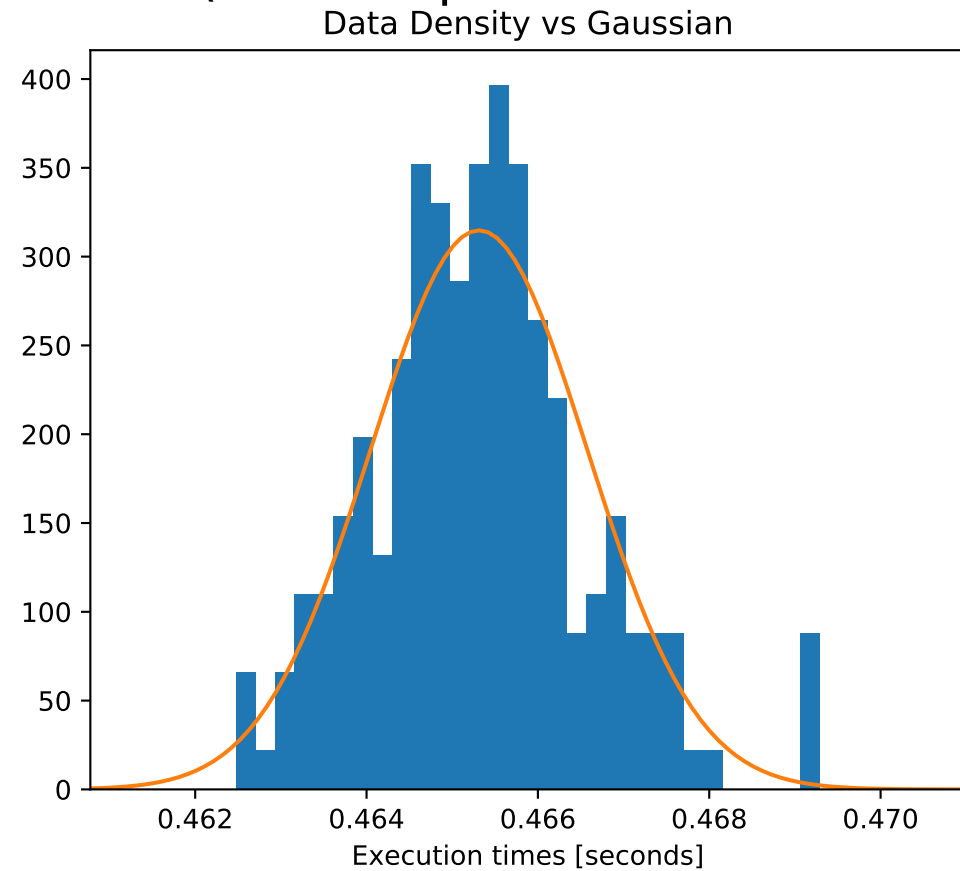
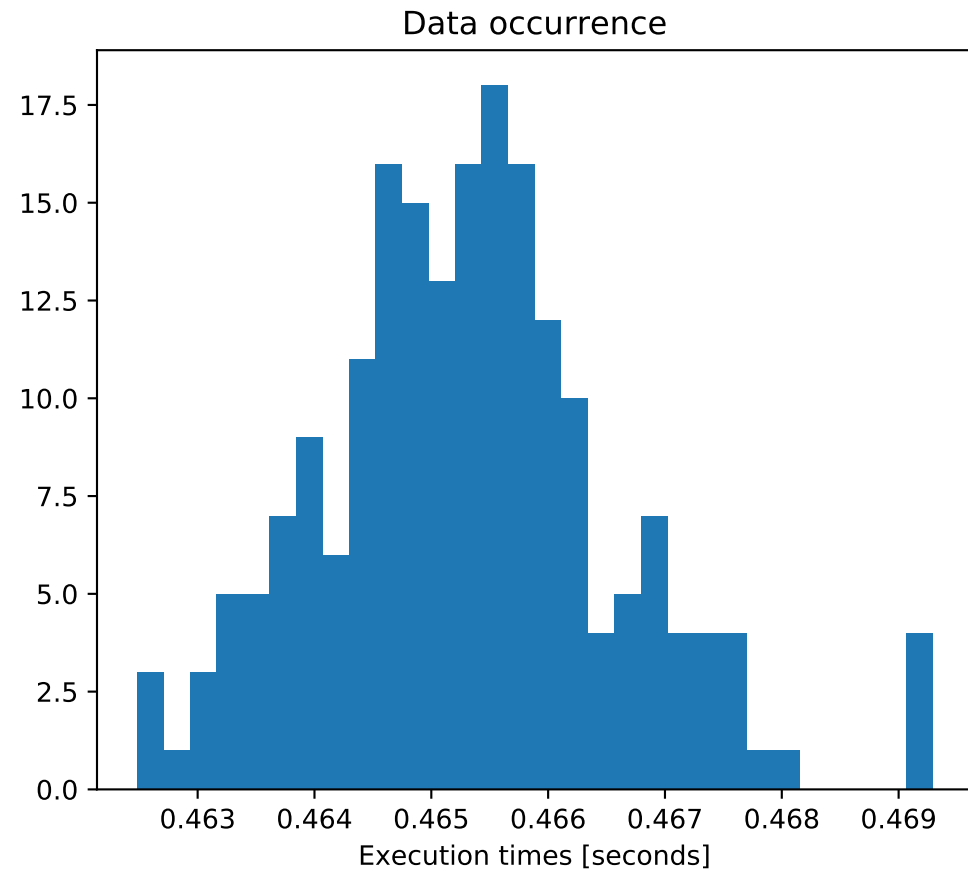
ATAX (IID with p-value 0.9897329091)



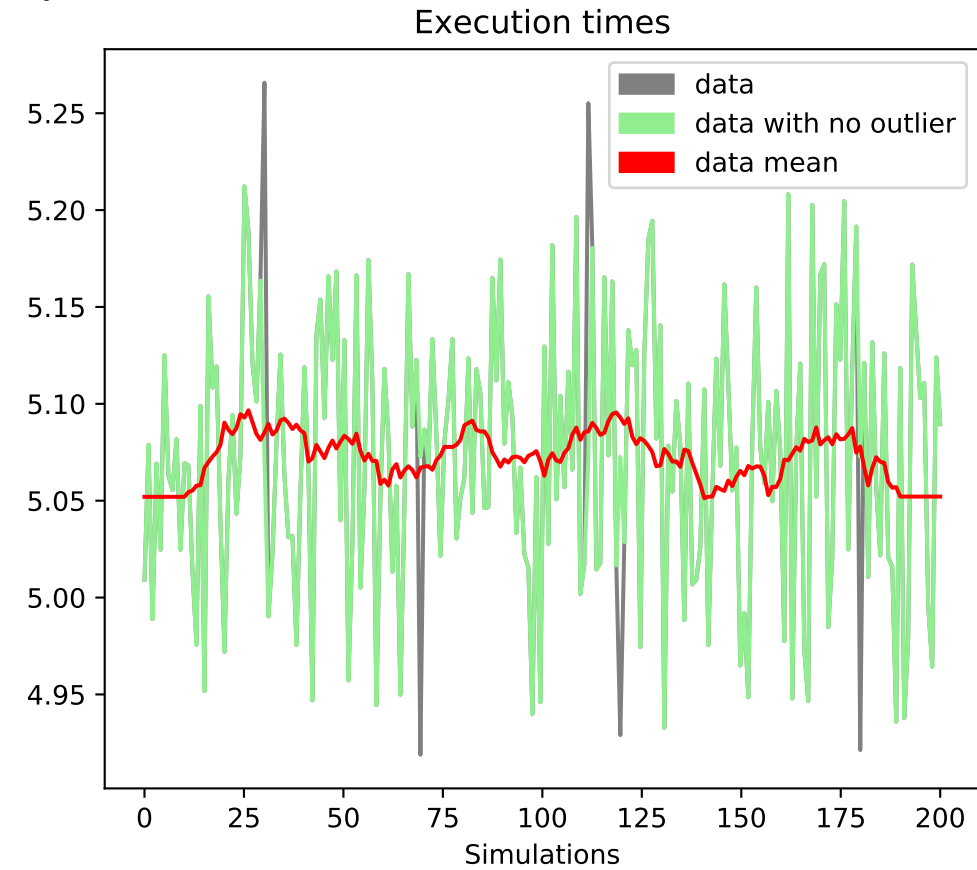
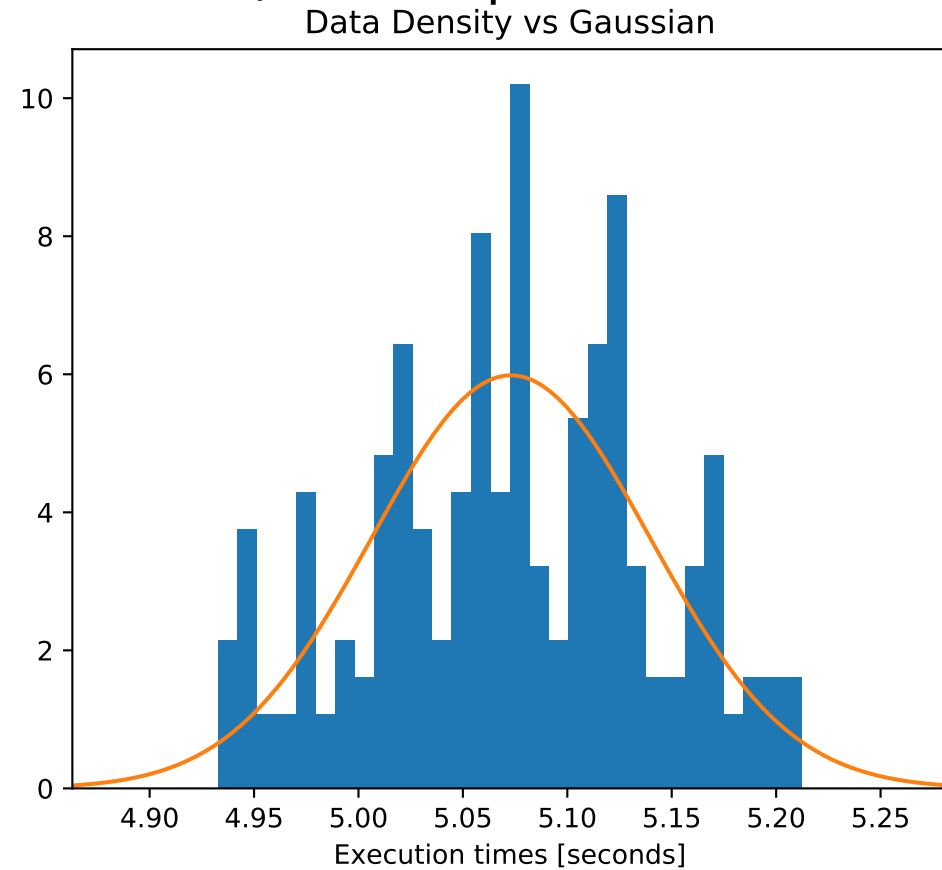
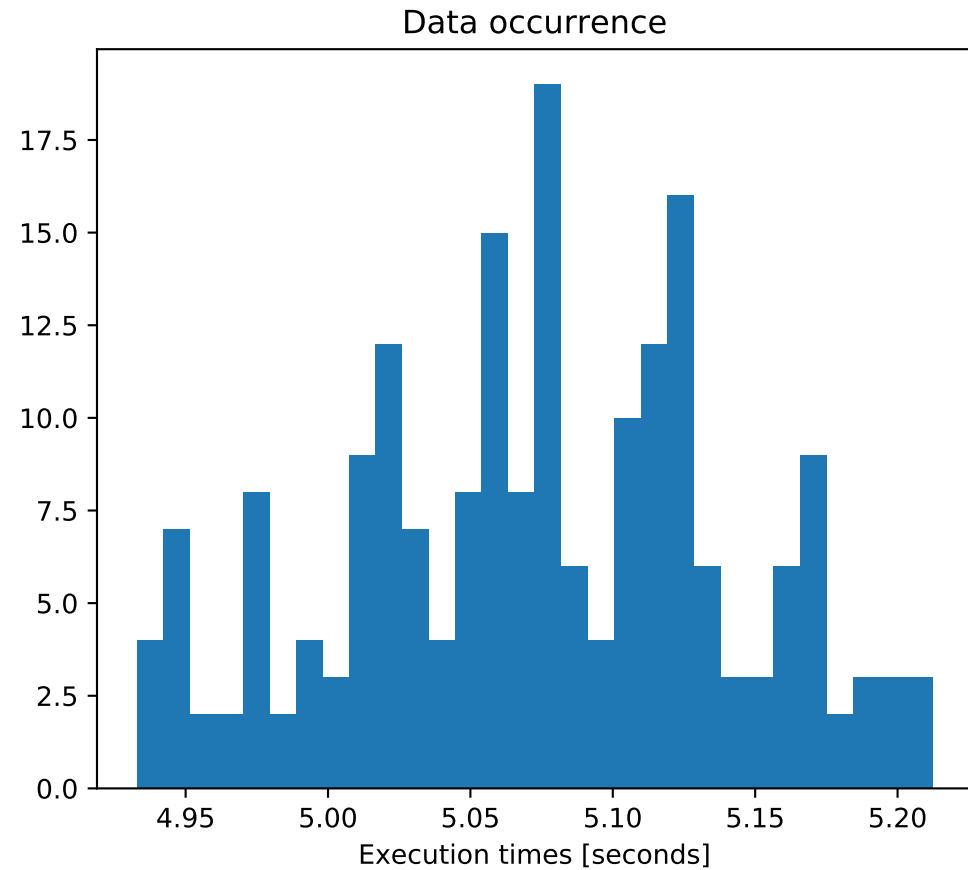
BICG (IID with p-value 0.747418002)



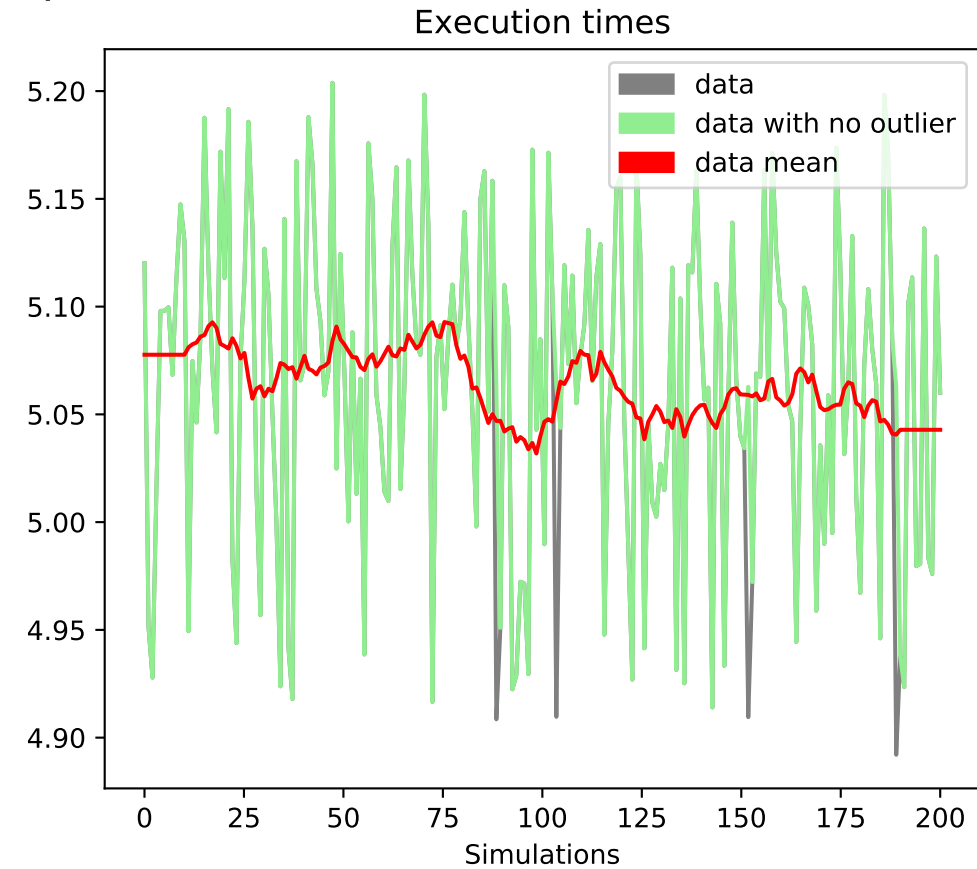
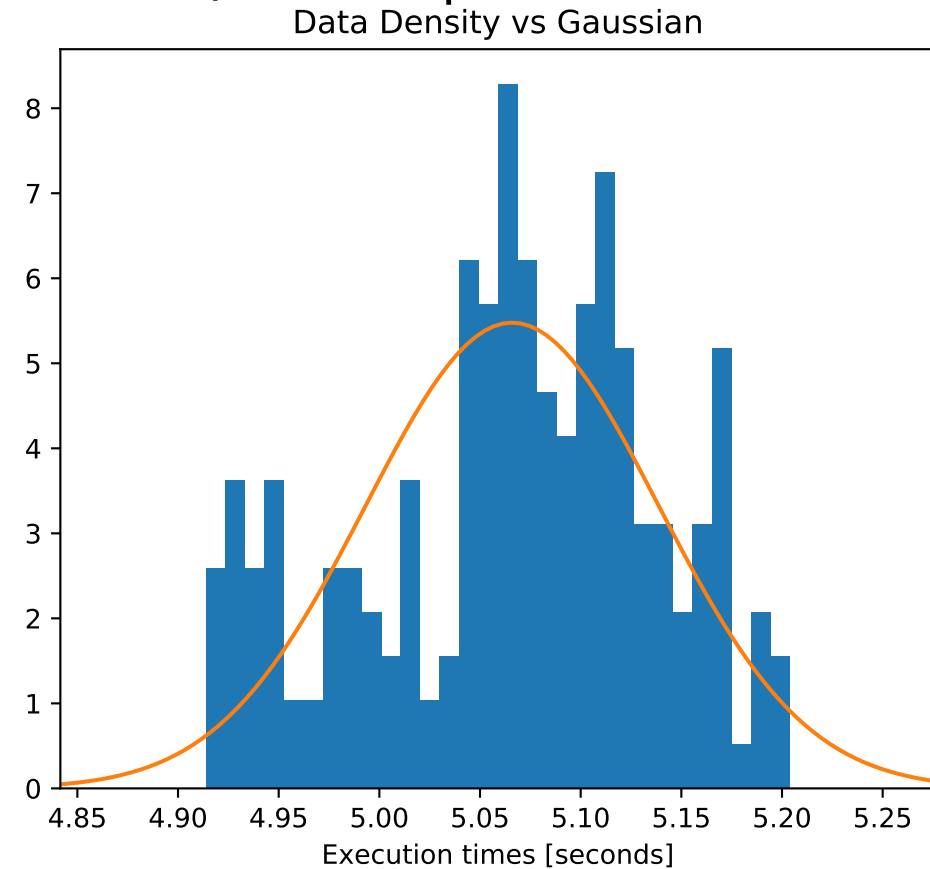
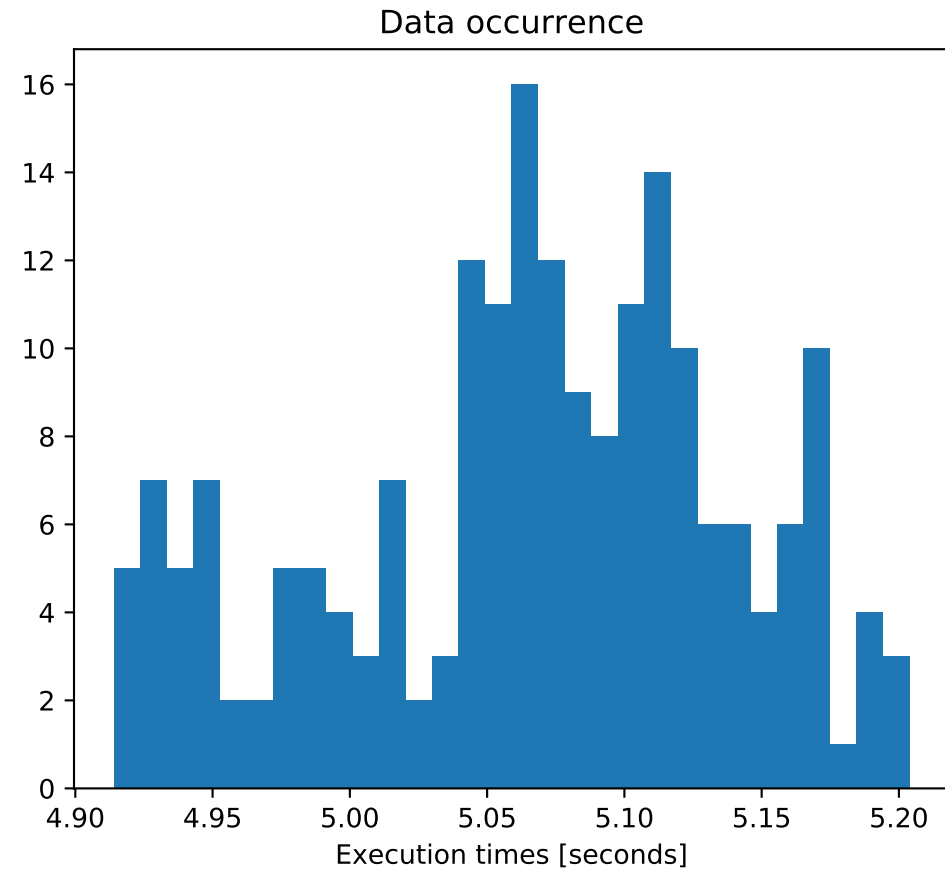
CHOLSKY (IID with p-value 0.2032544257)



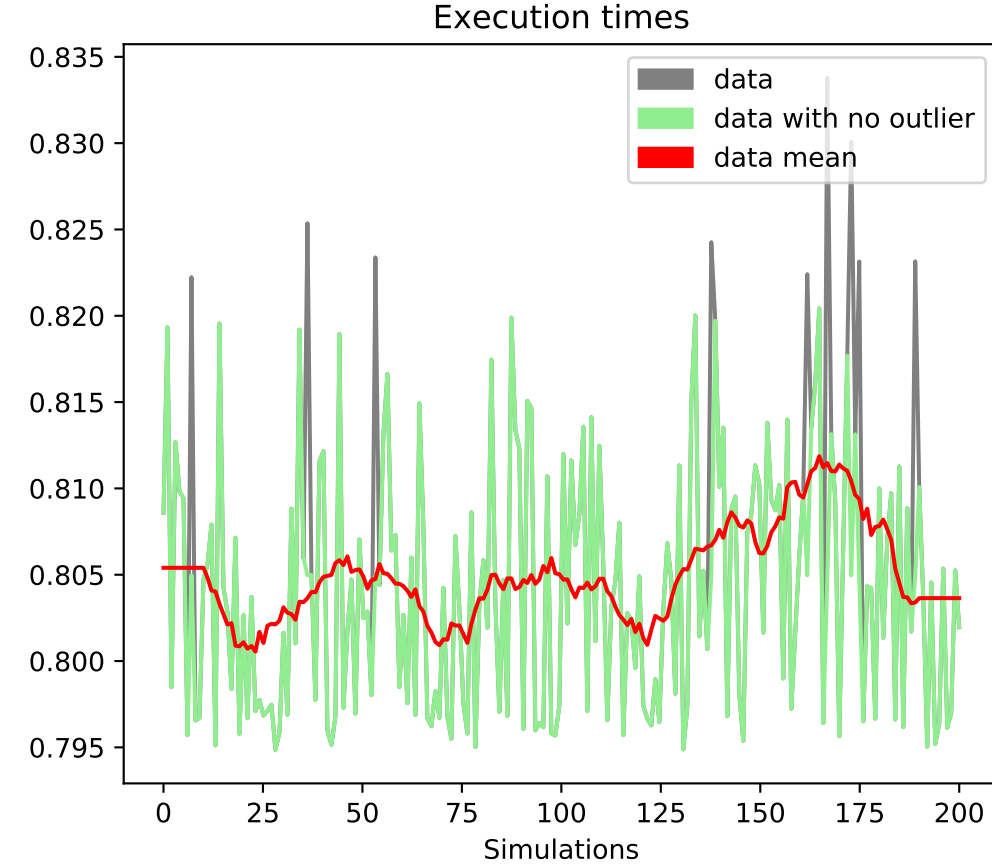
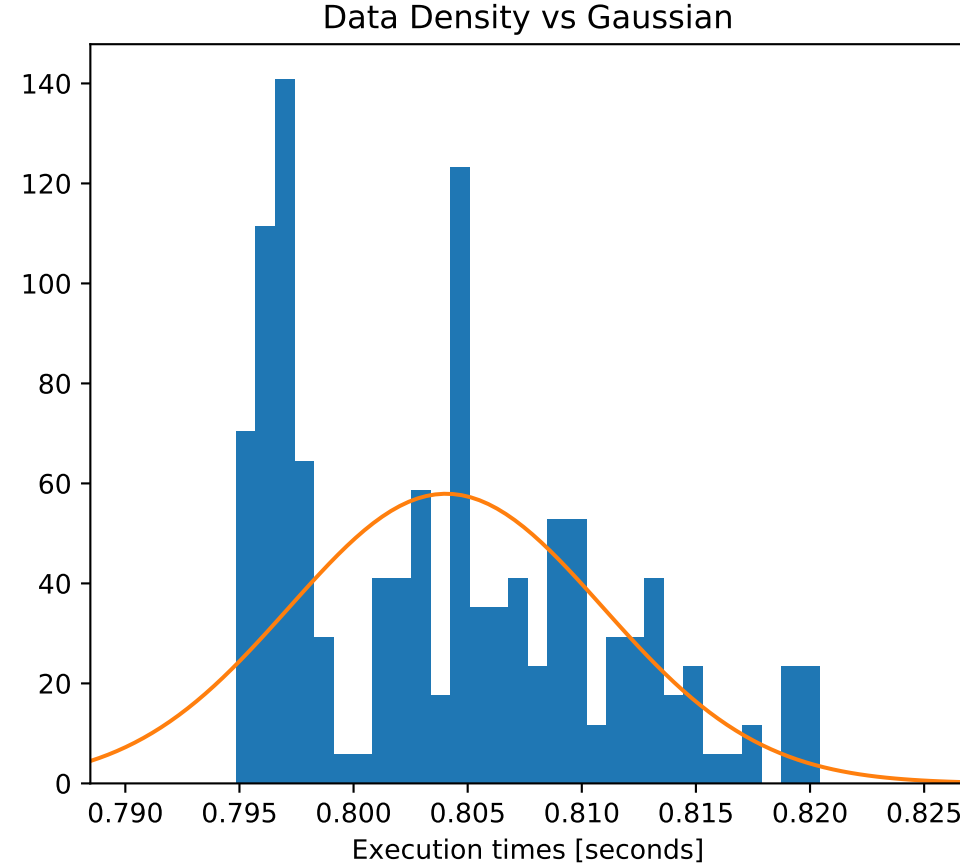
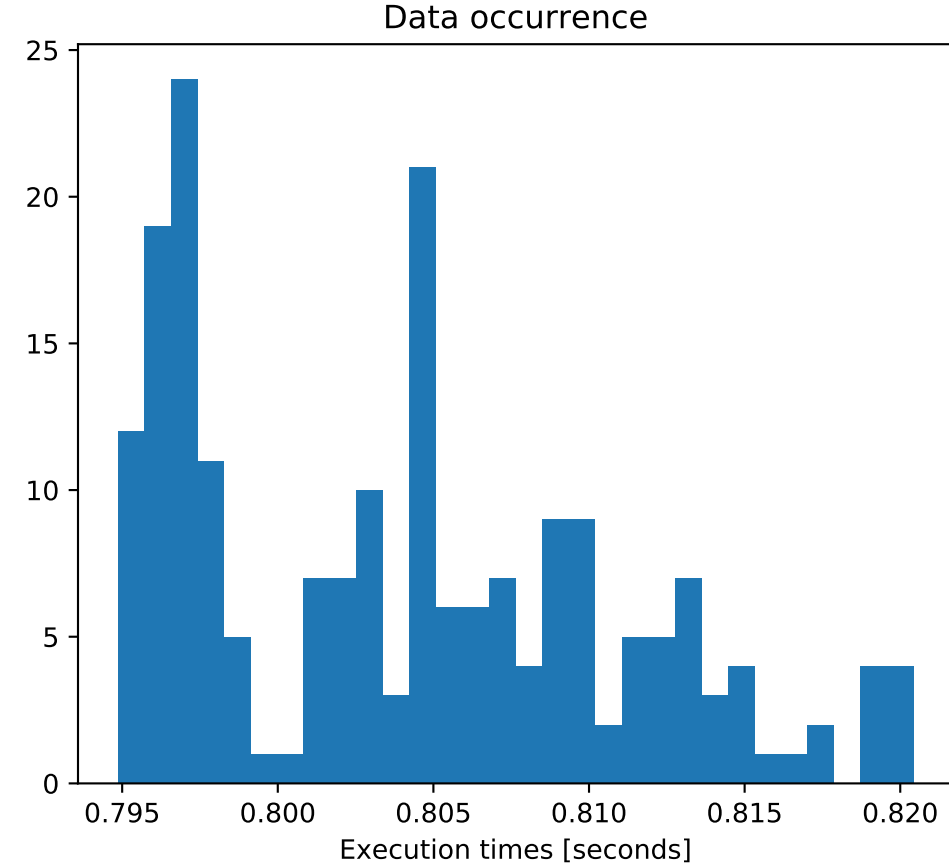
CORRELATION (IID with p-value 0.693079555)



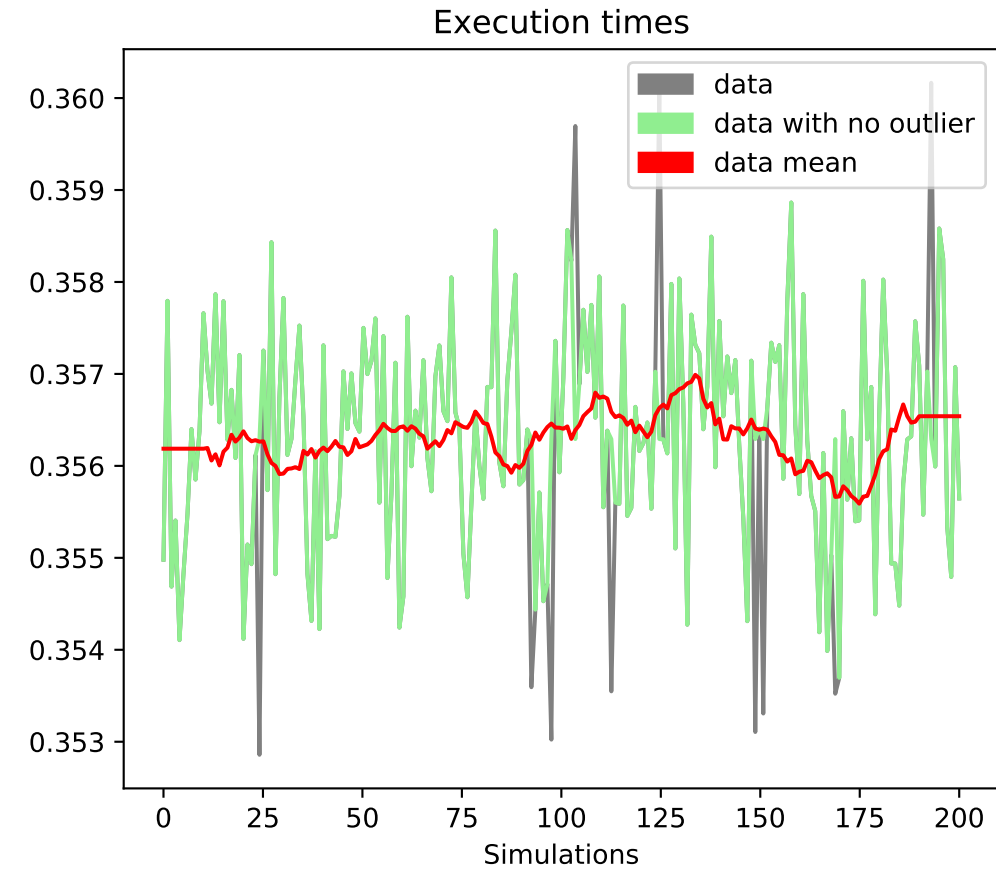
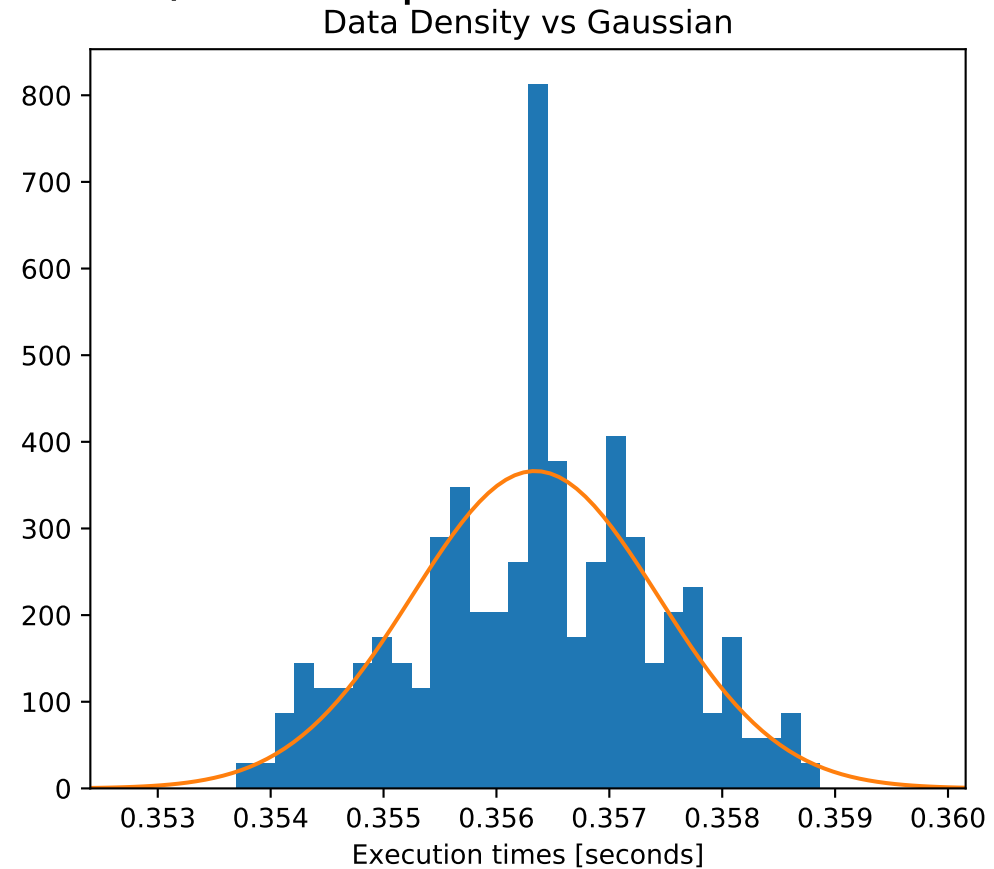
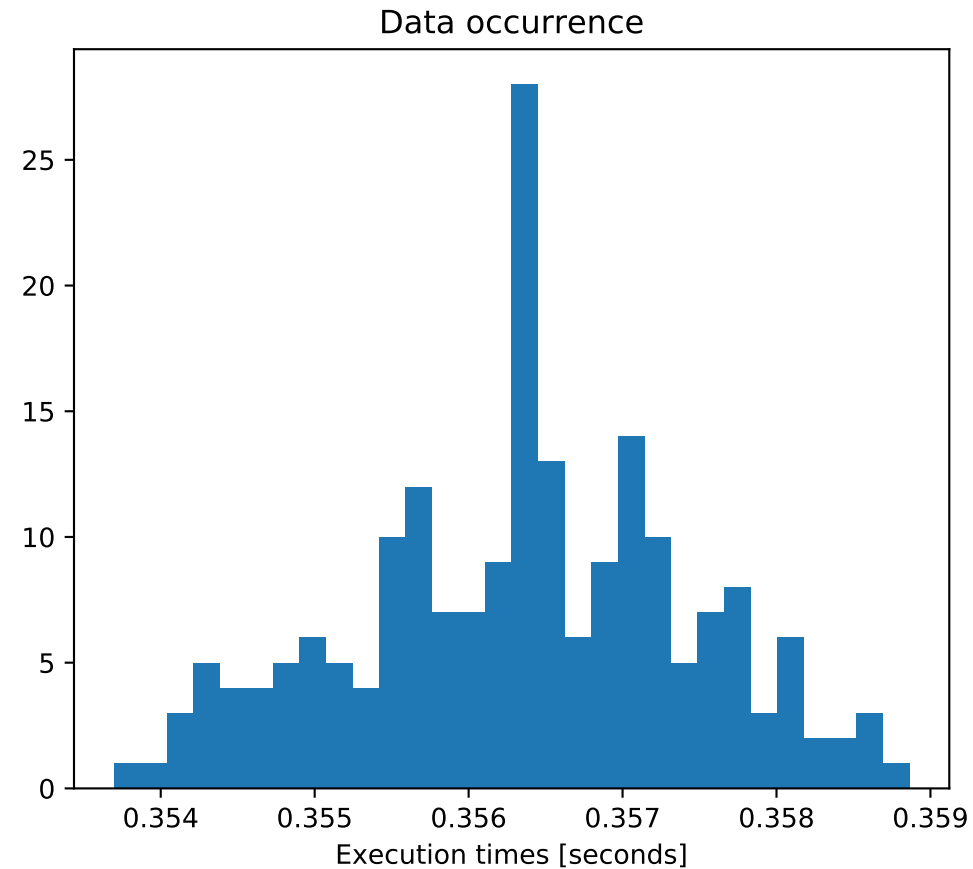
COVARIANCE (IID with p-value 0.3115718757)



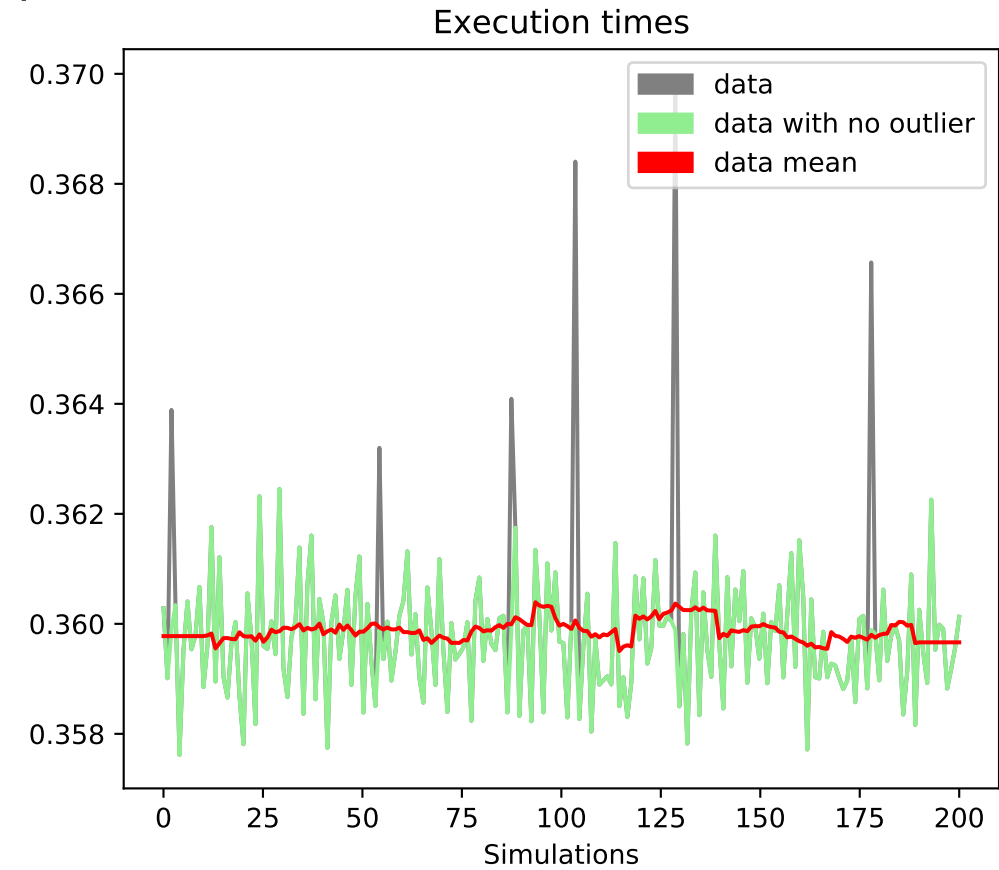
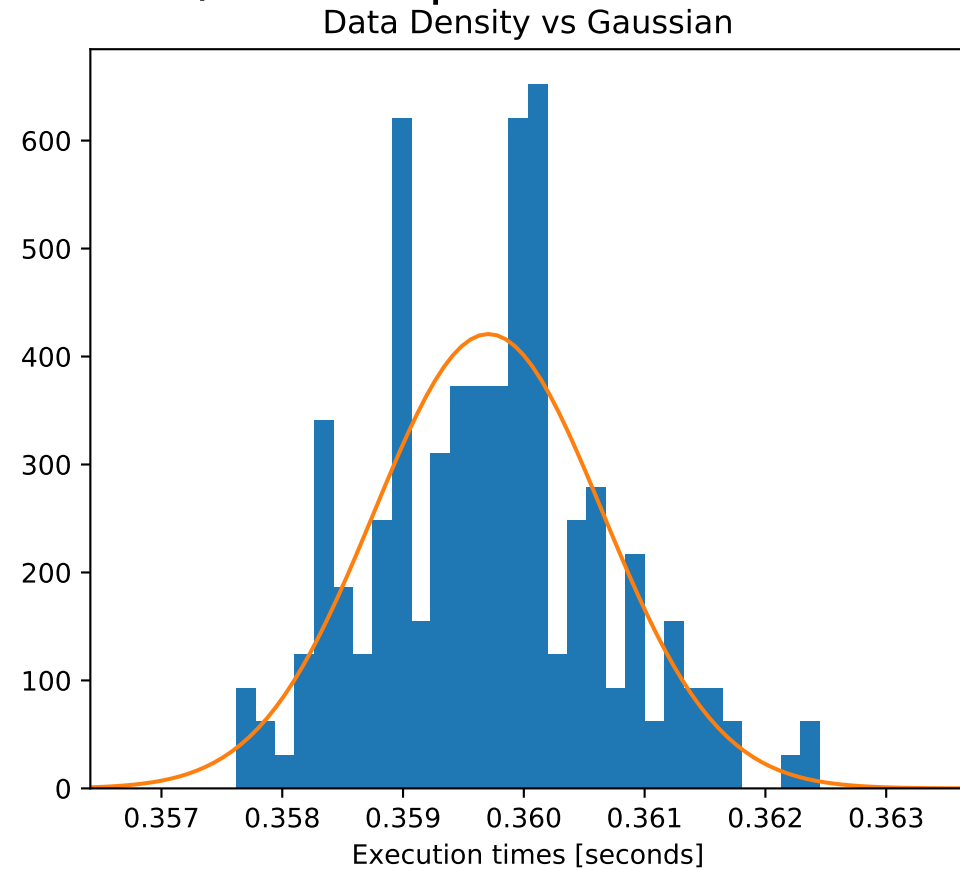
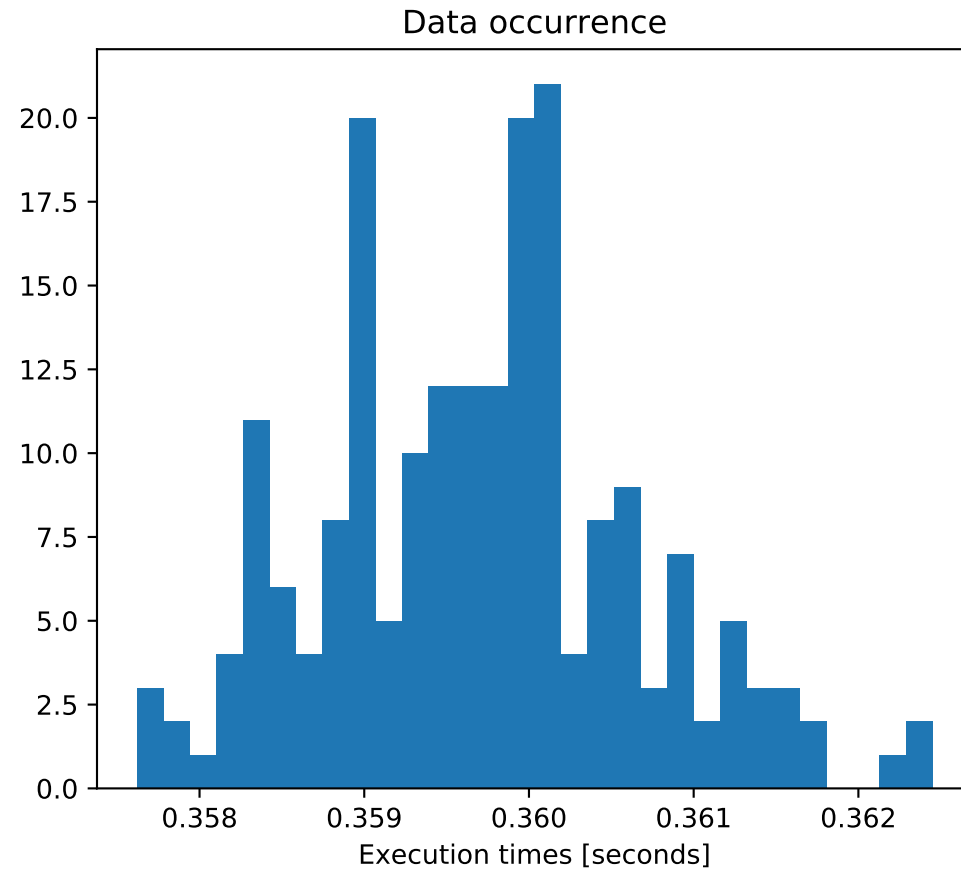
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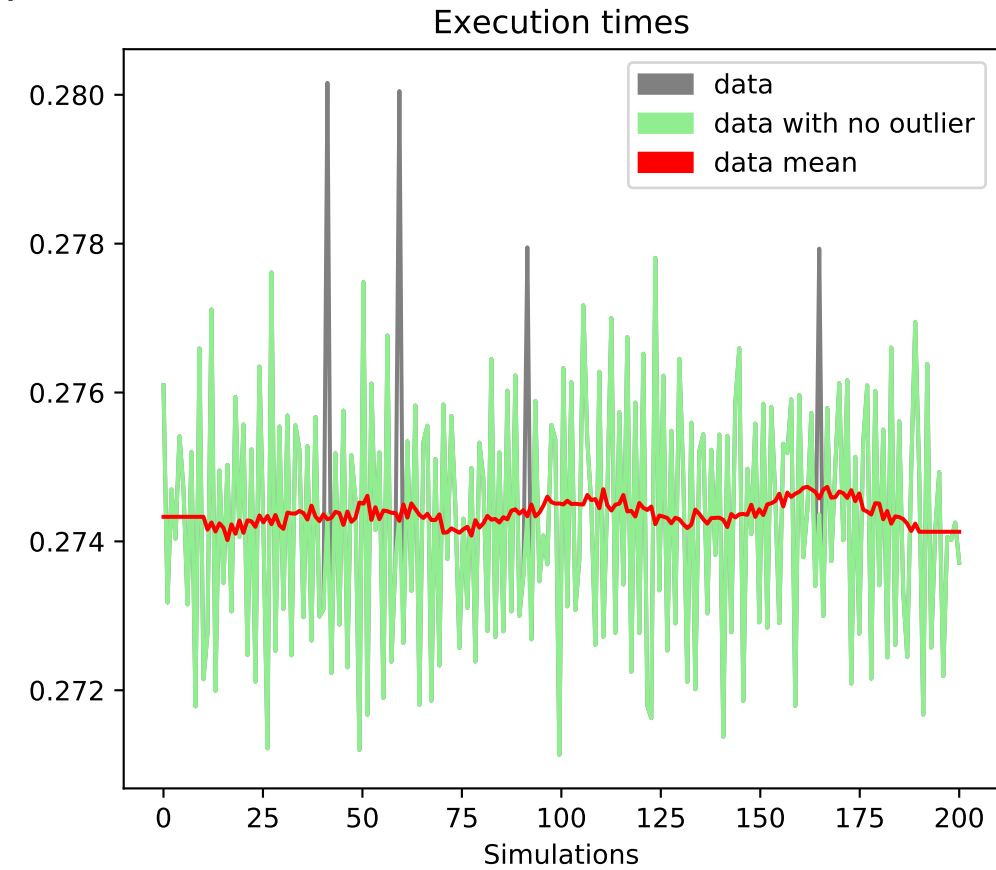
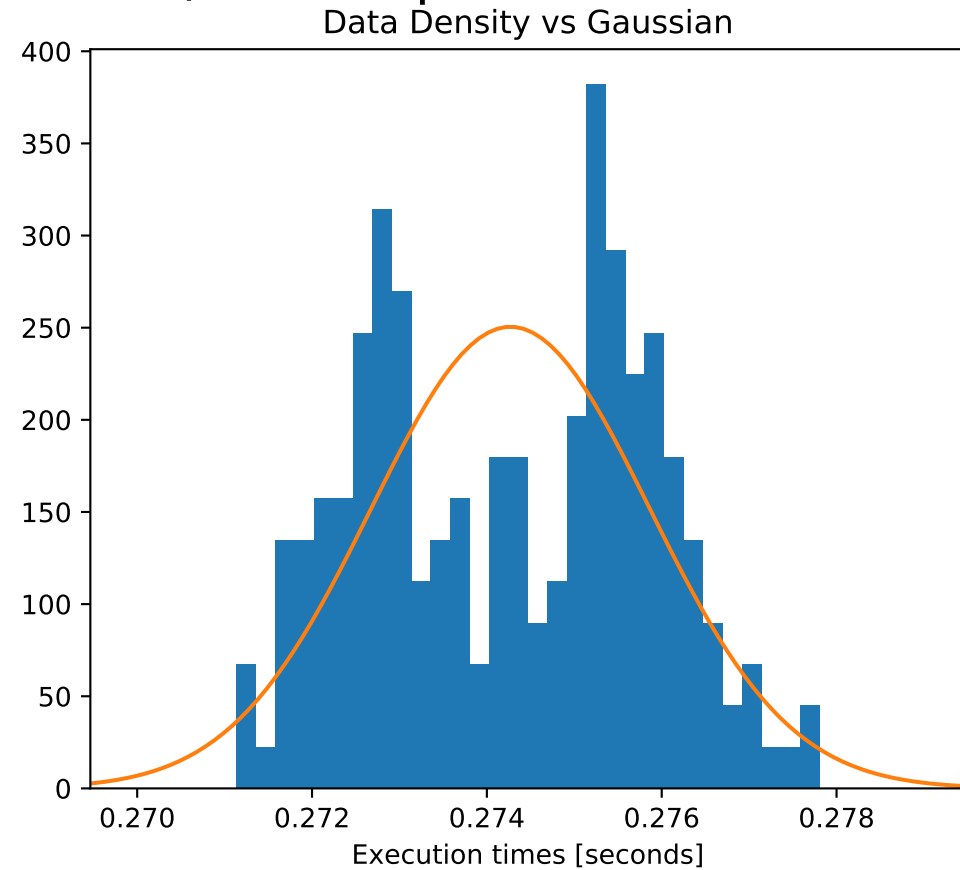
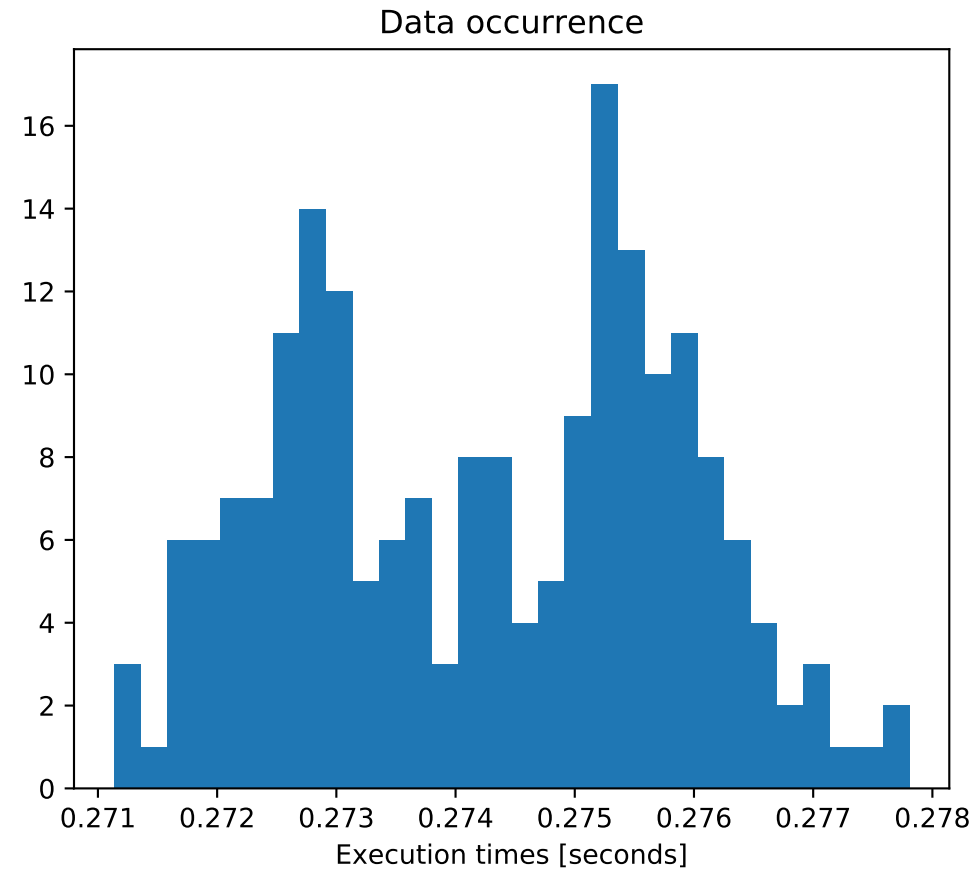
DURBIN (IID with p-value 0.1862154808)



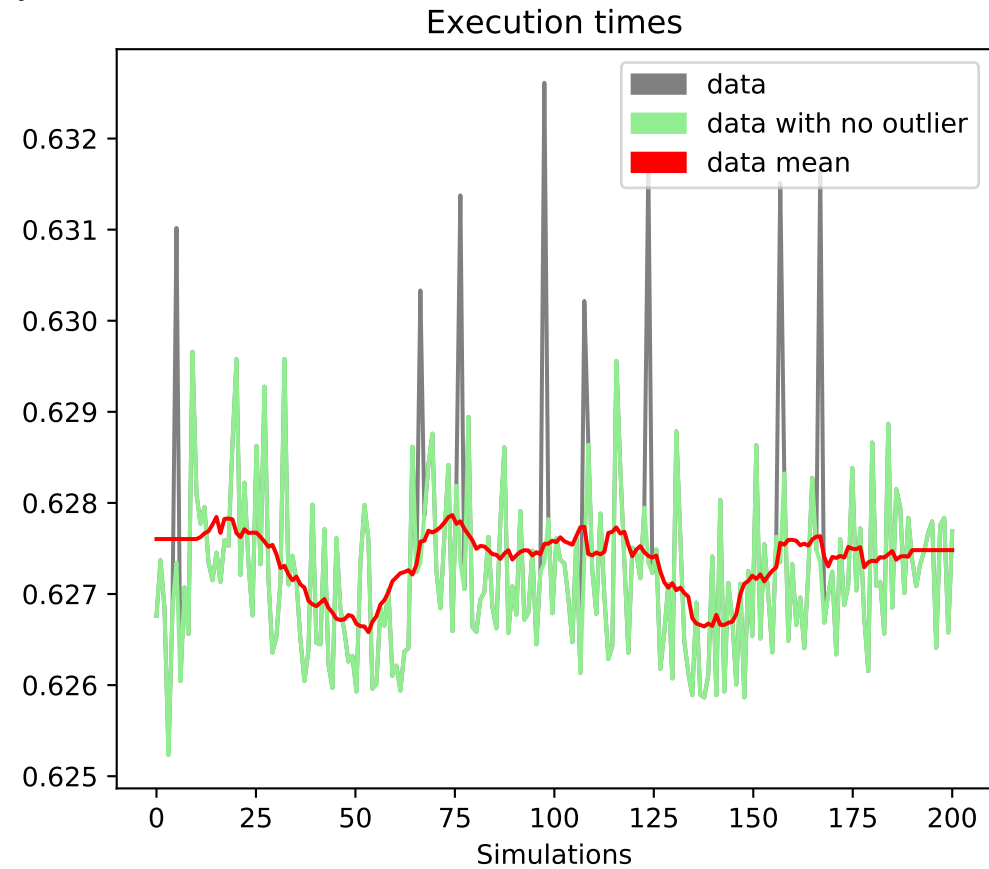
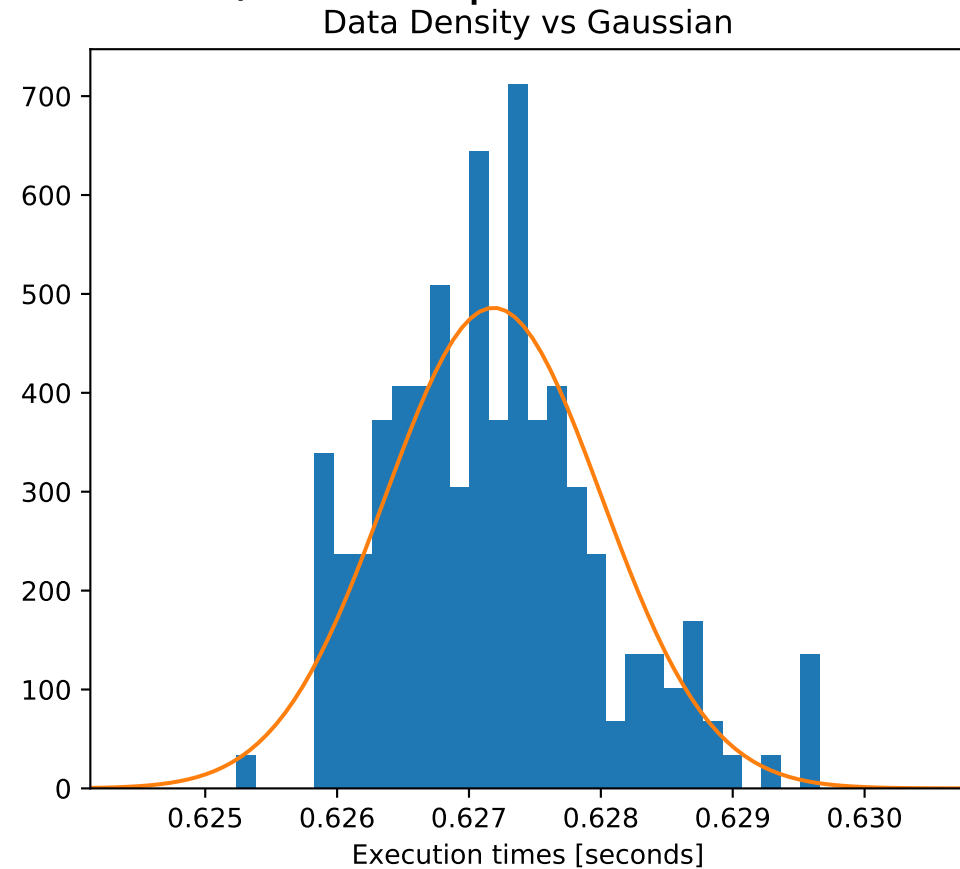
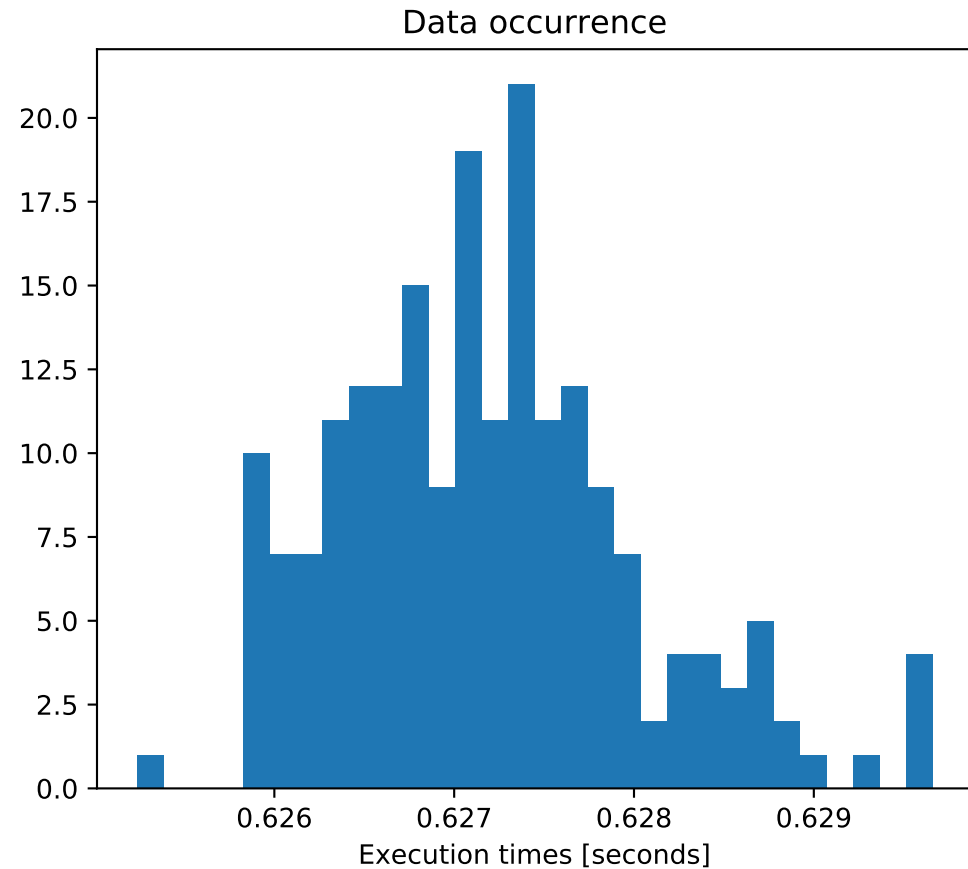
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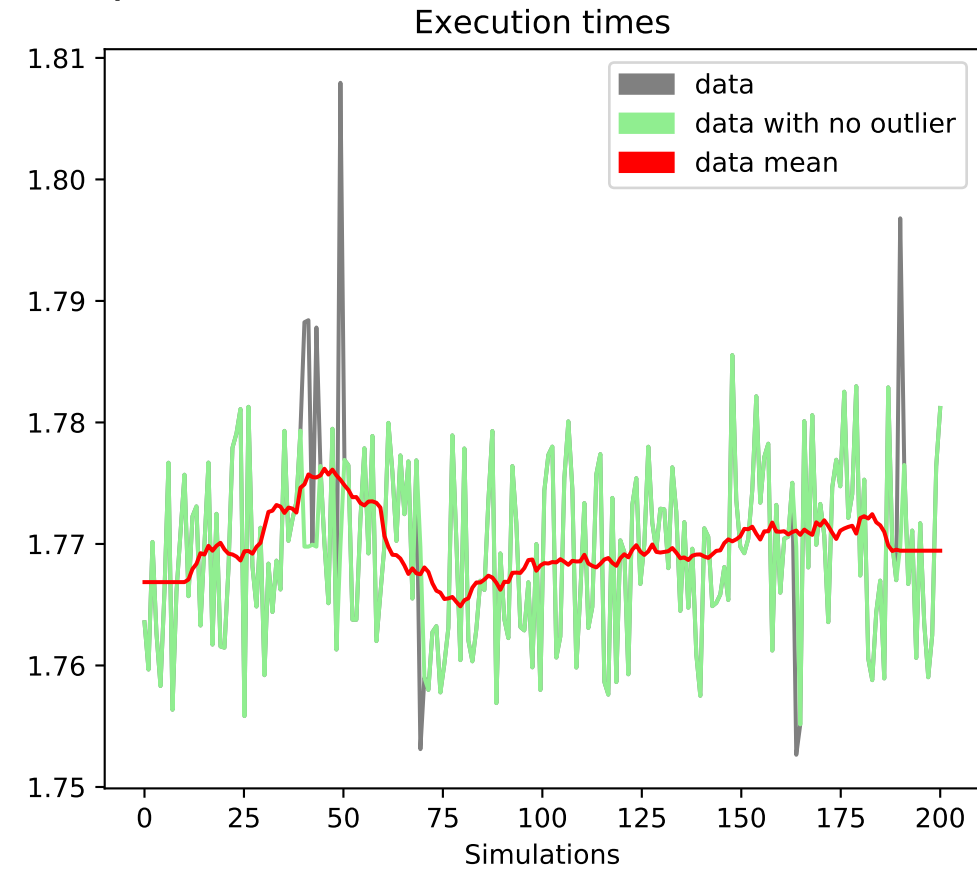
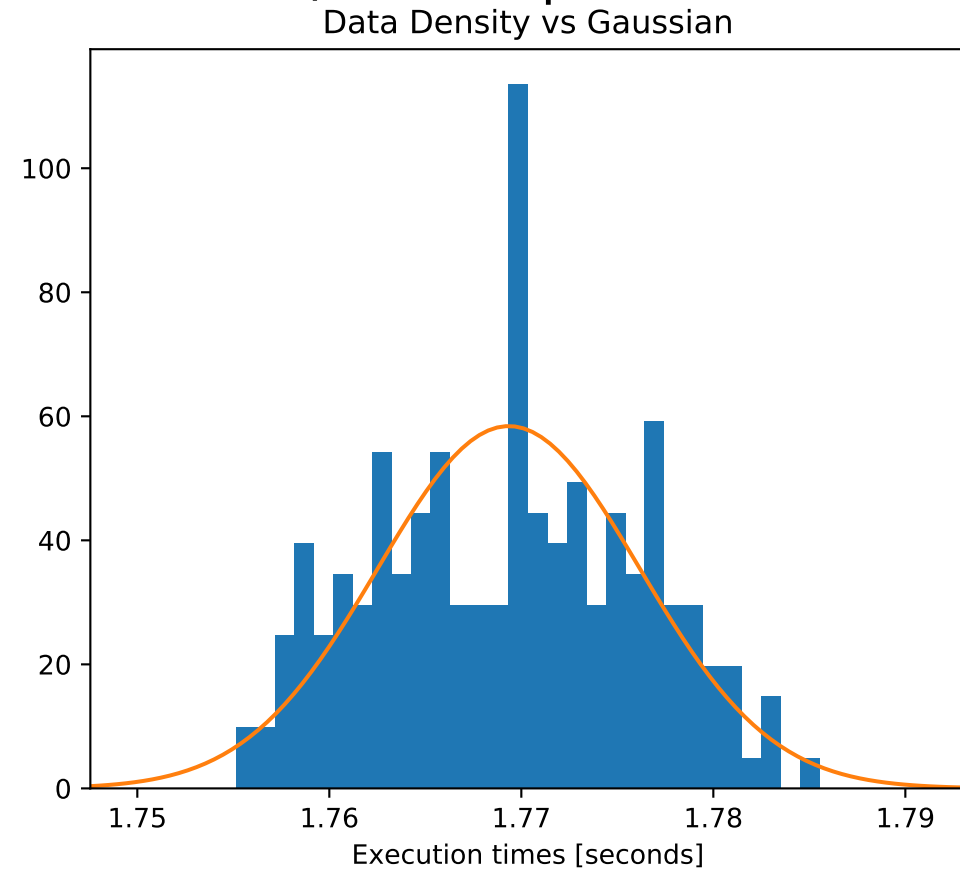
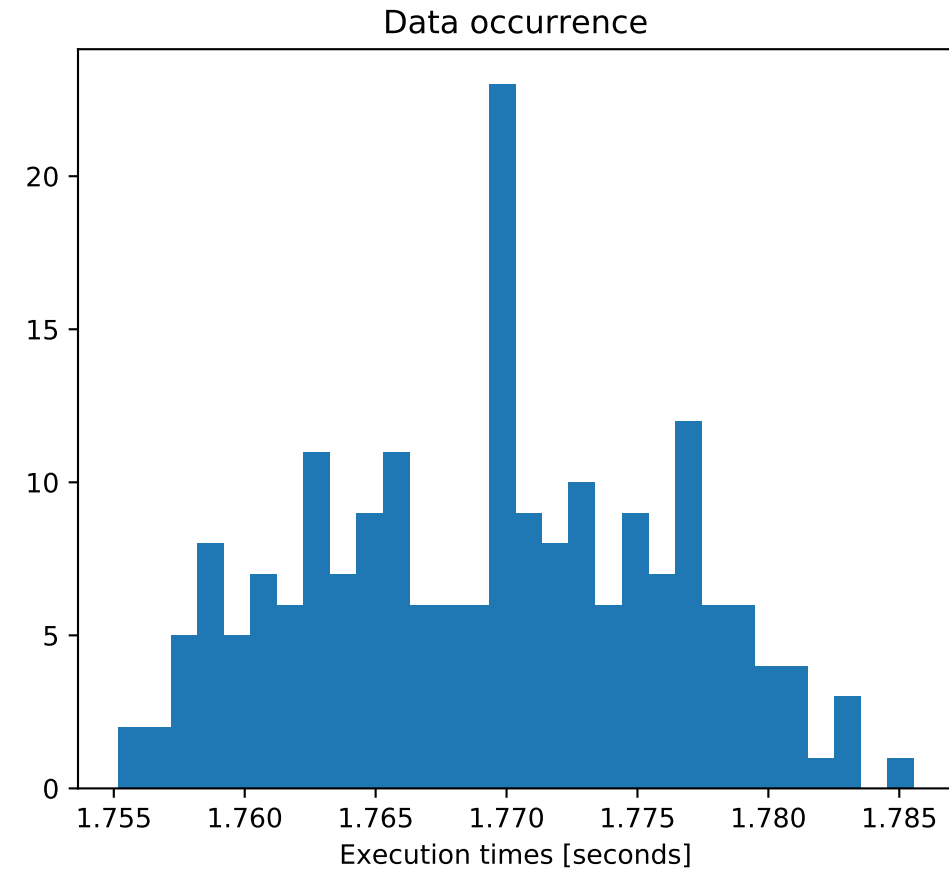
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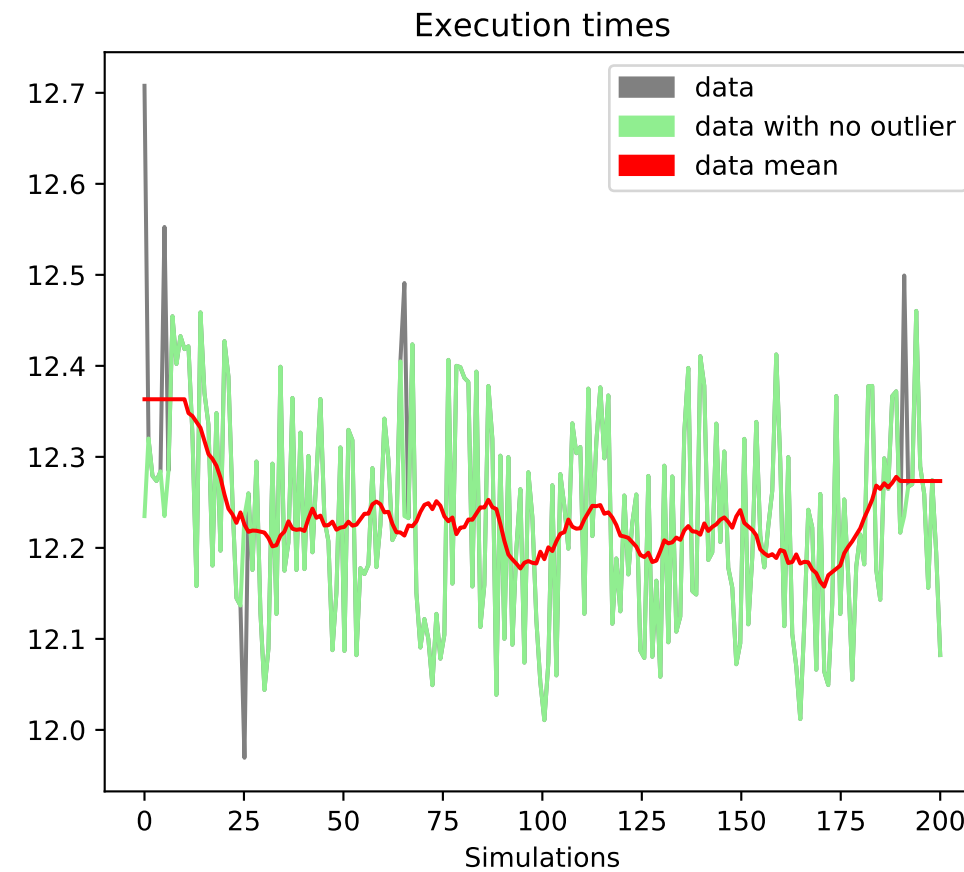
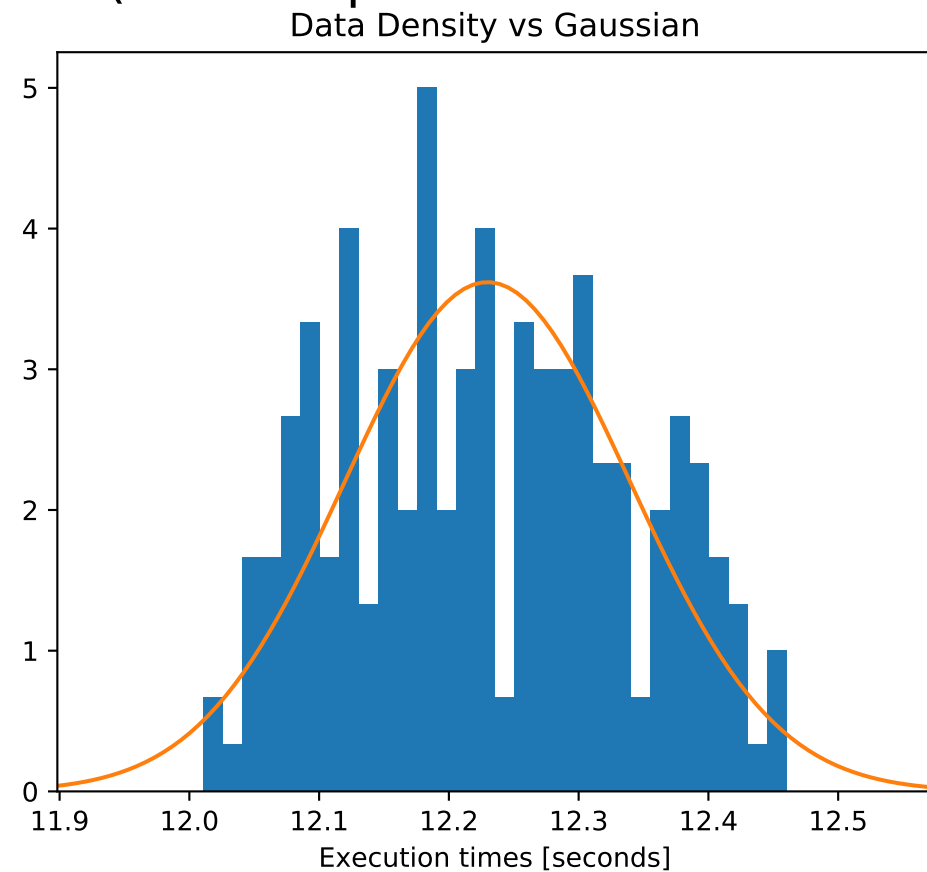
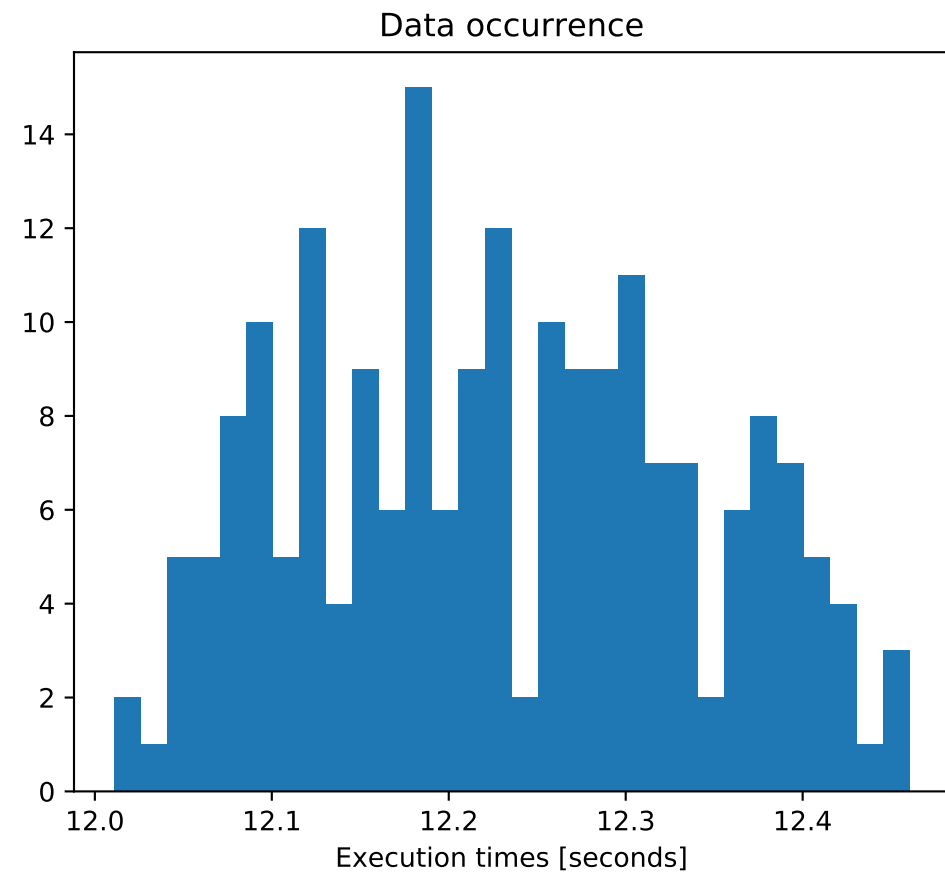
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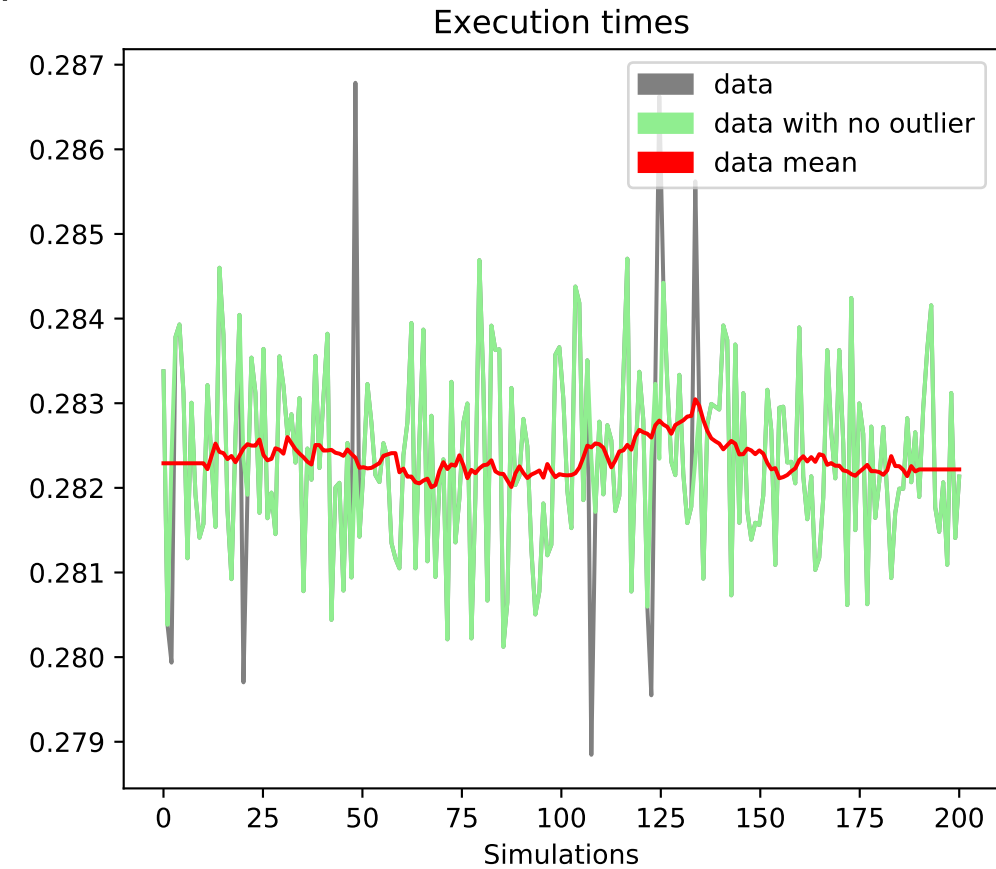
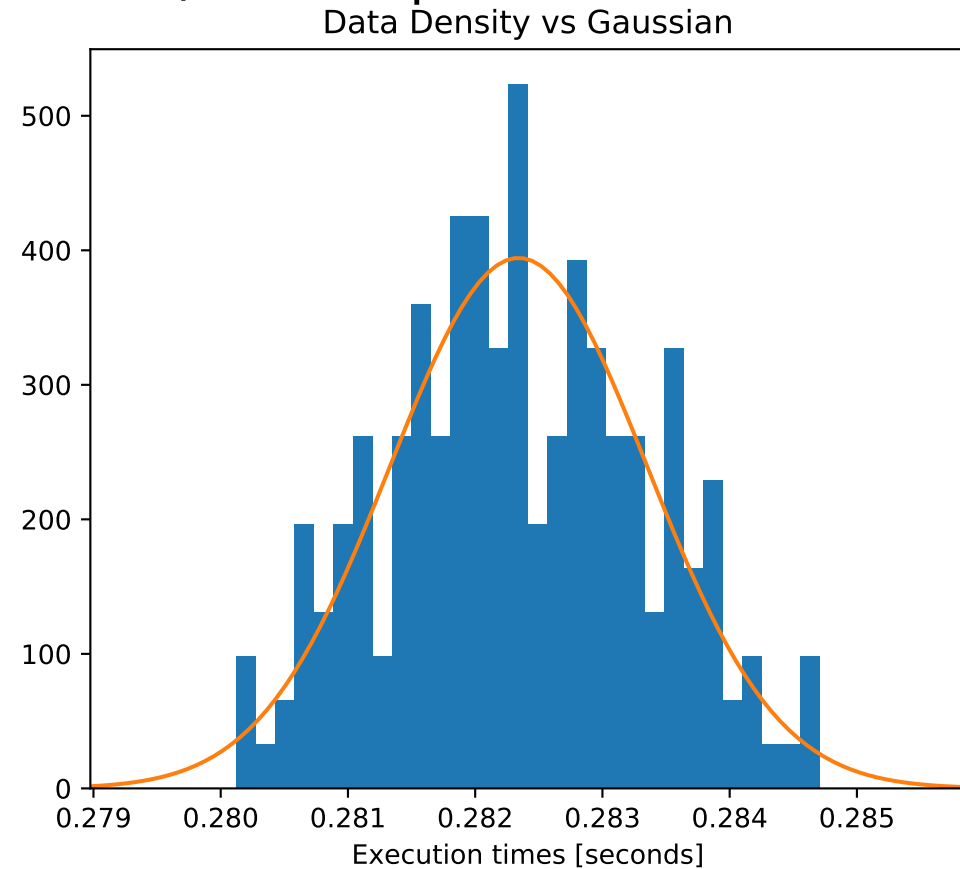
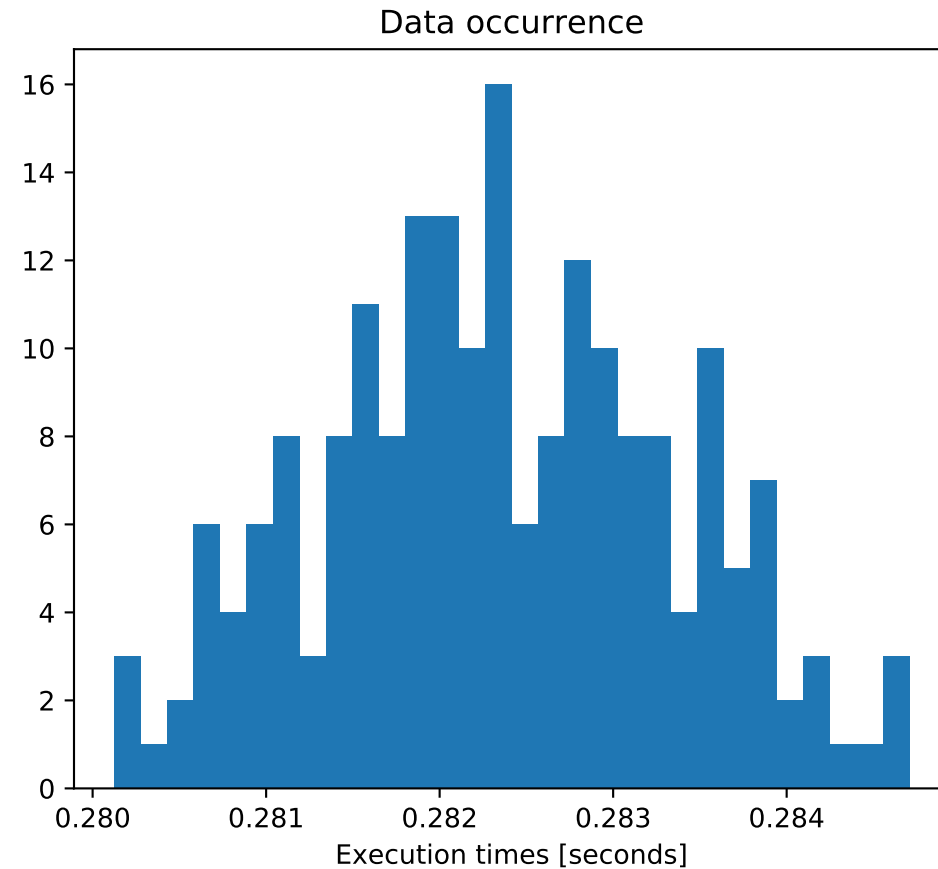
FLOYD-WARSHALL (IID with p-value 0.1221796539)



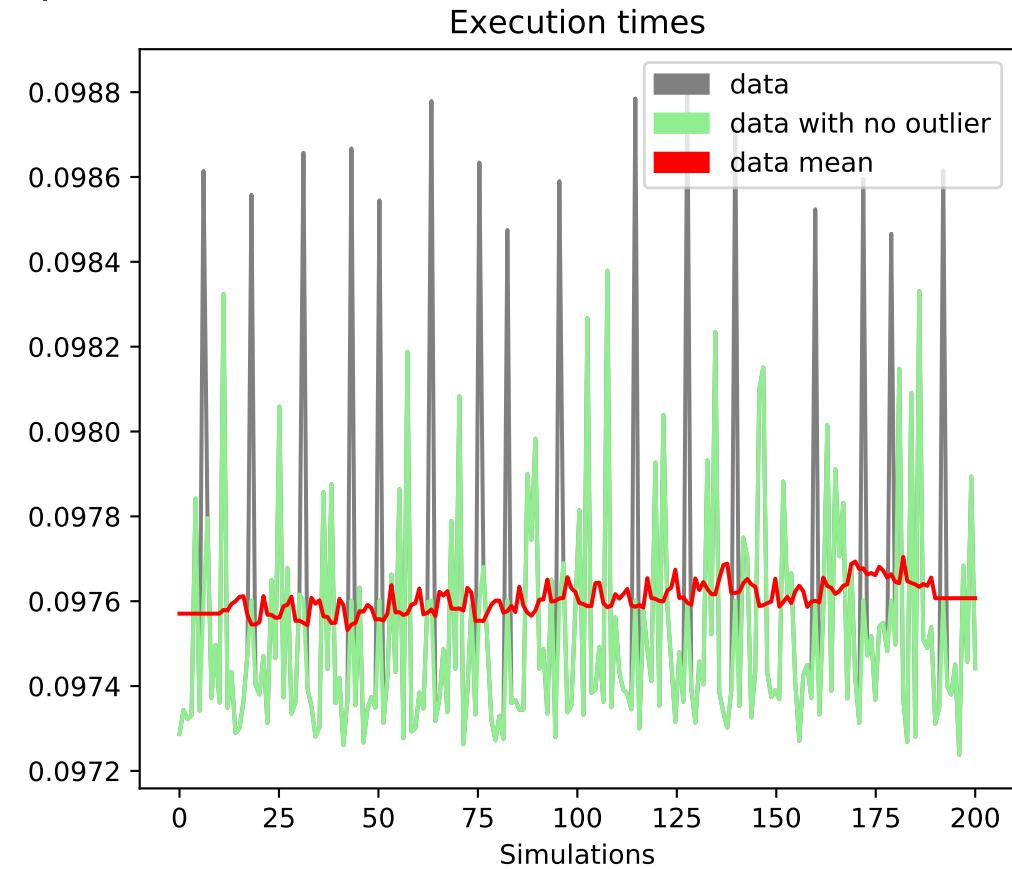
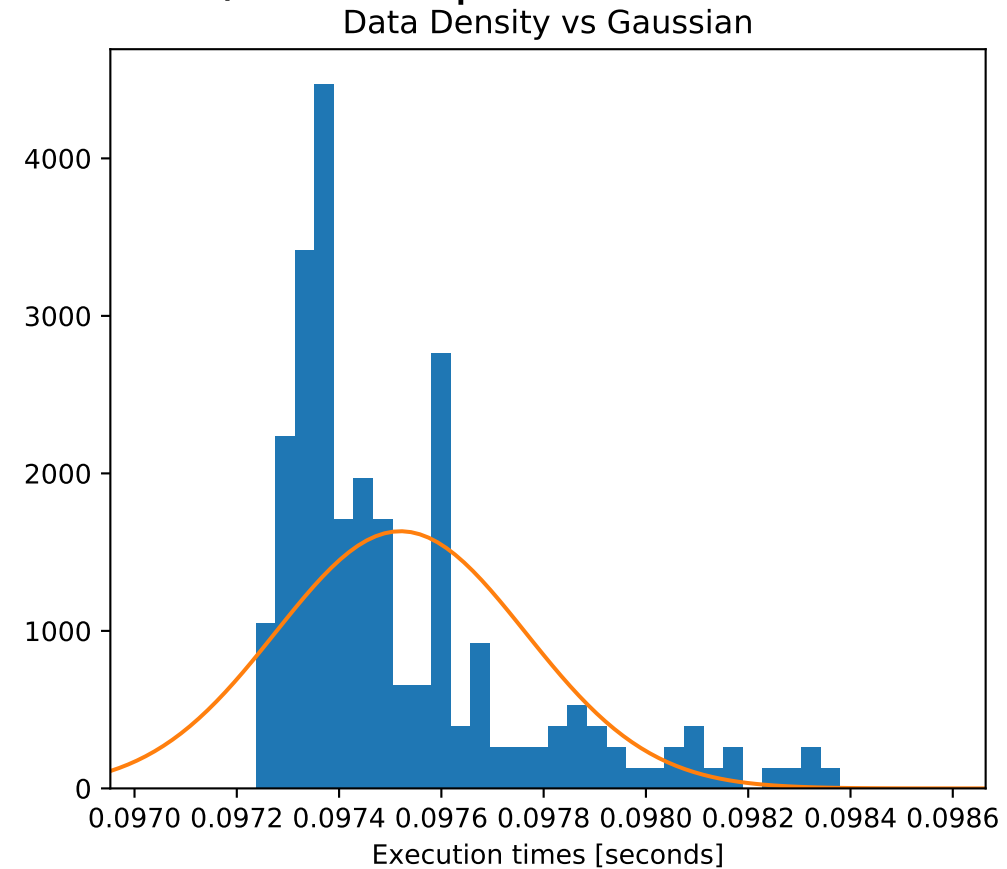
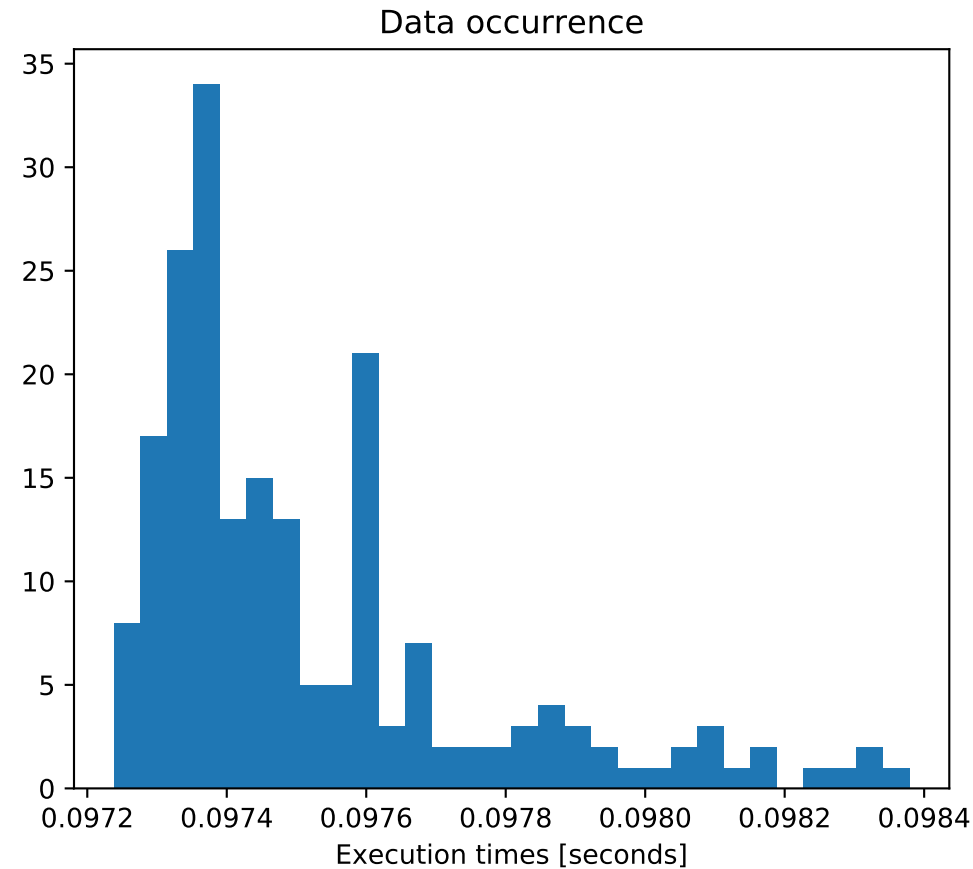
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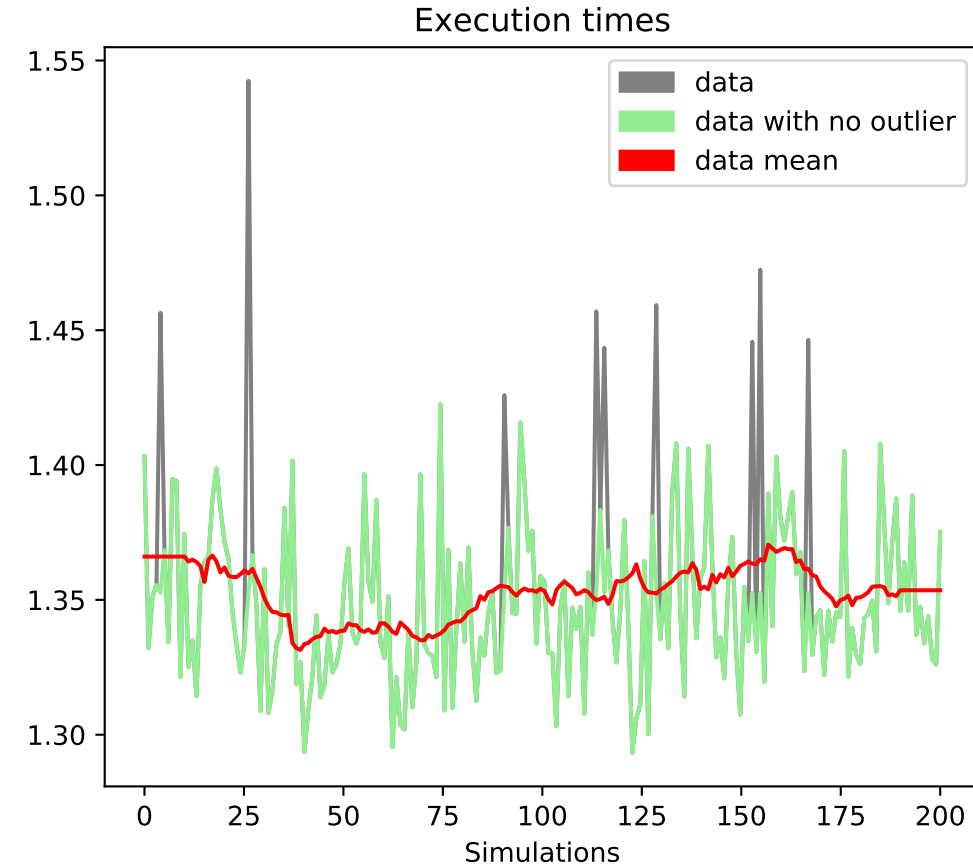
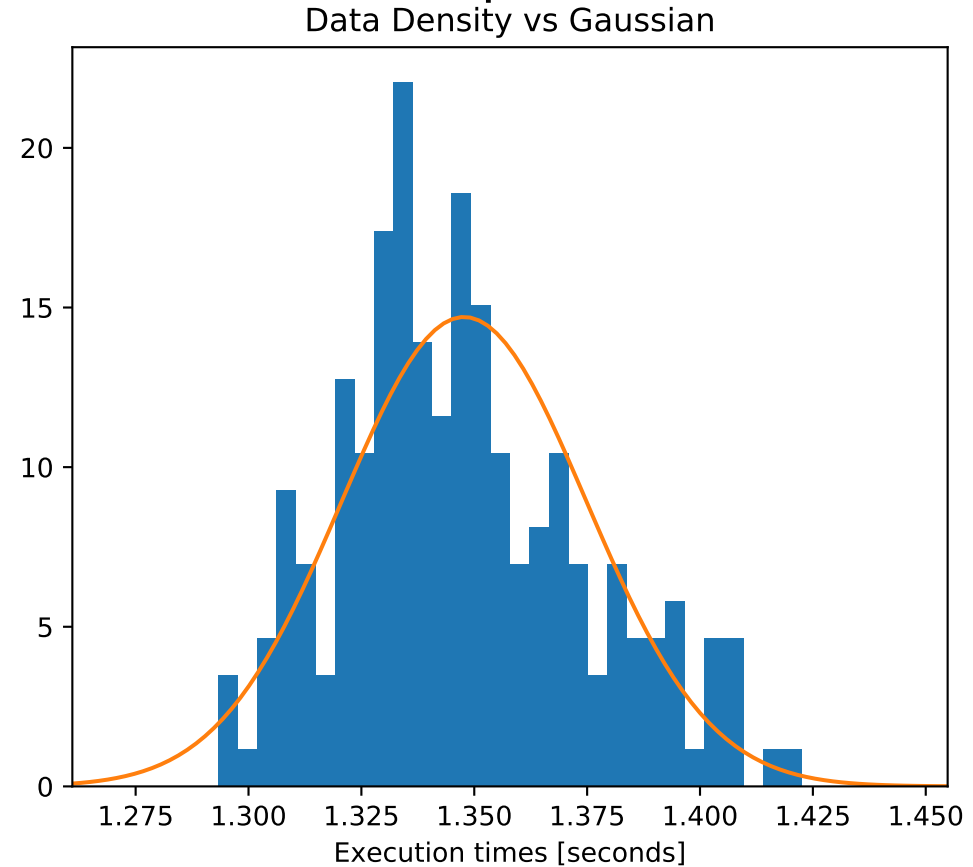
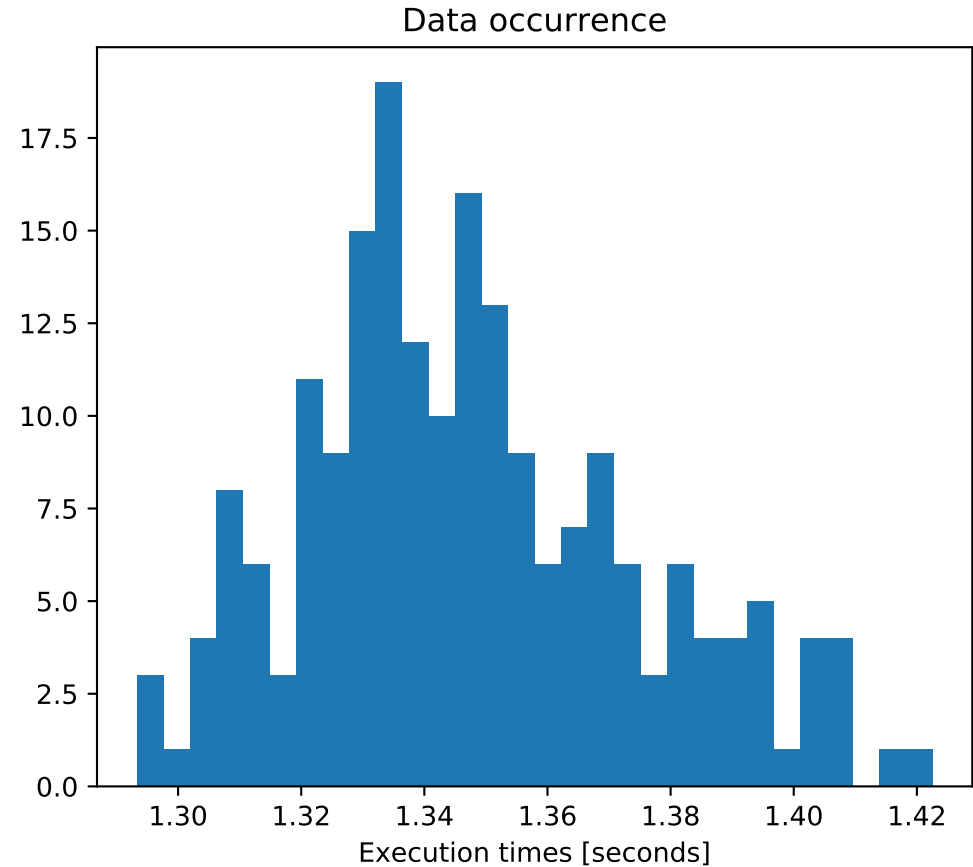
GEMVER (IID with p-value 0.4750853029)



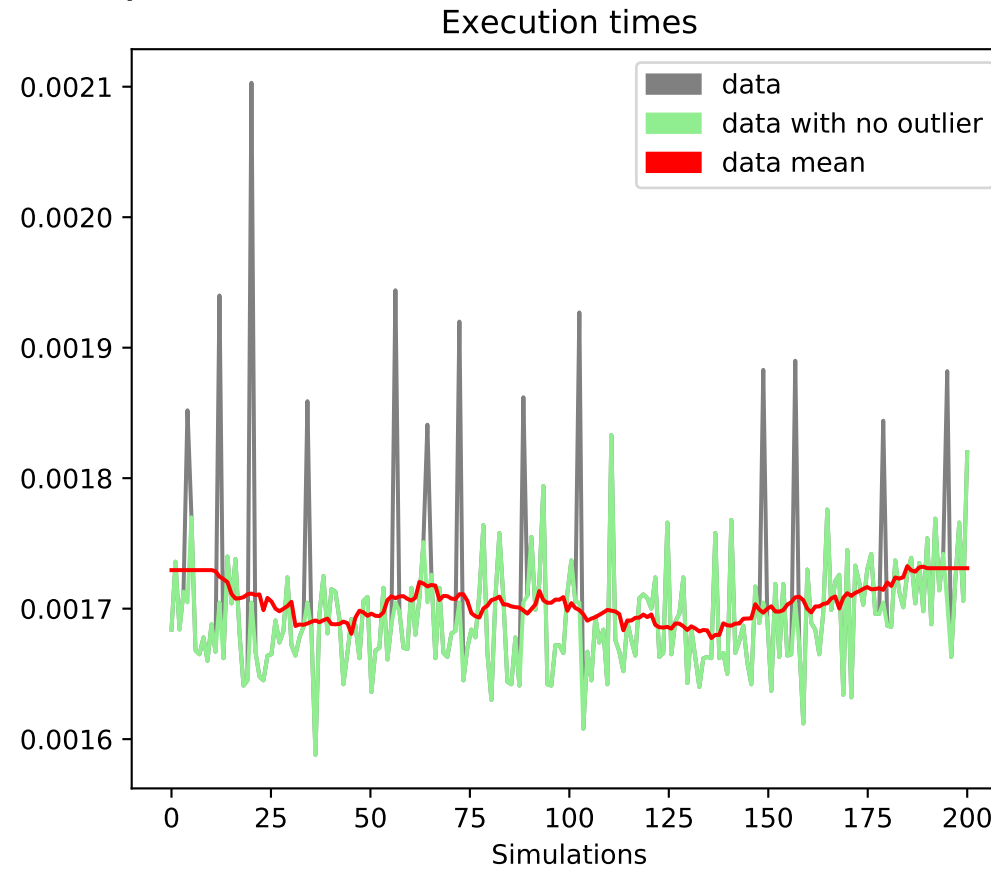
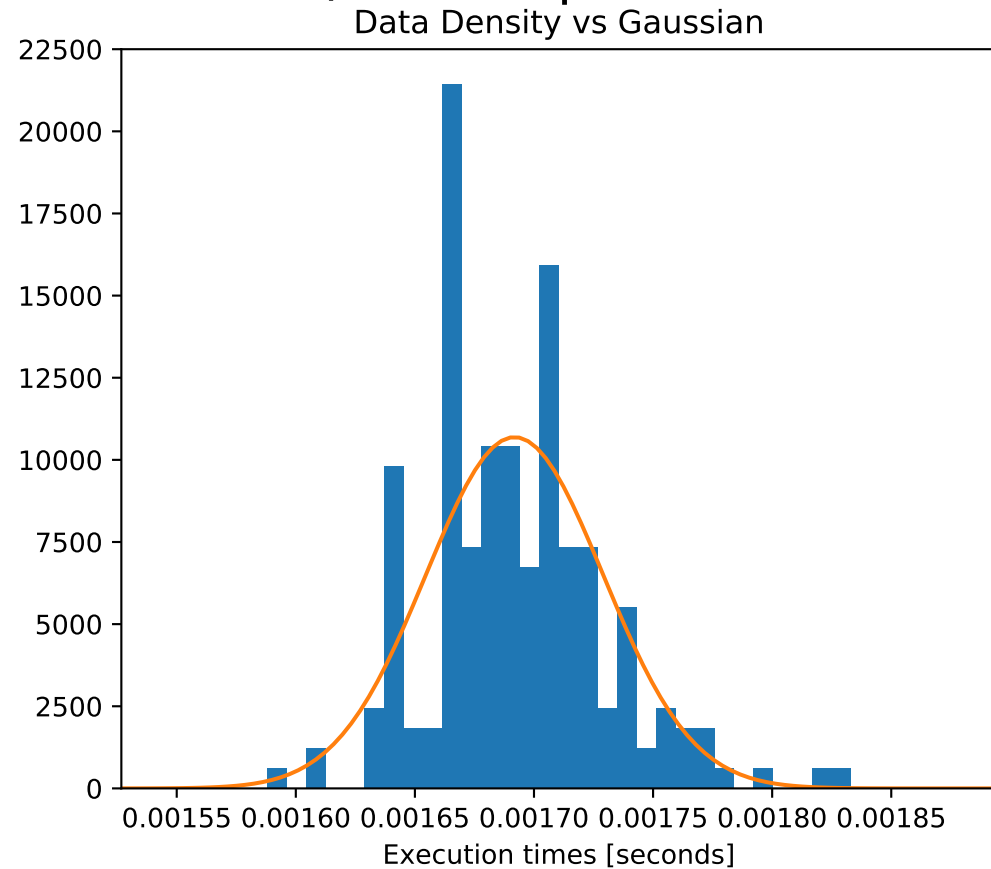
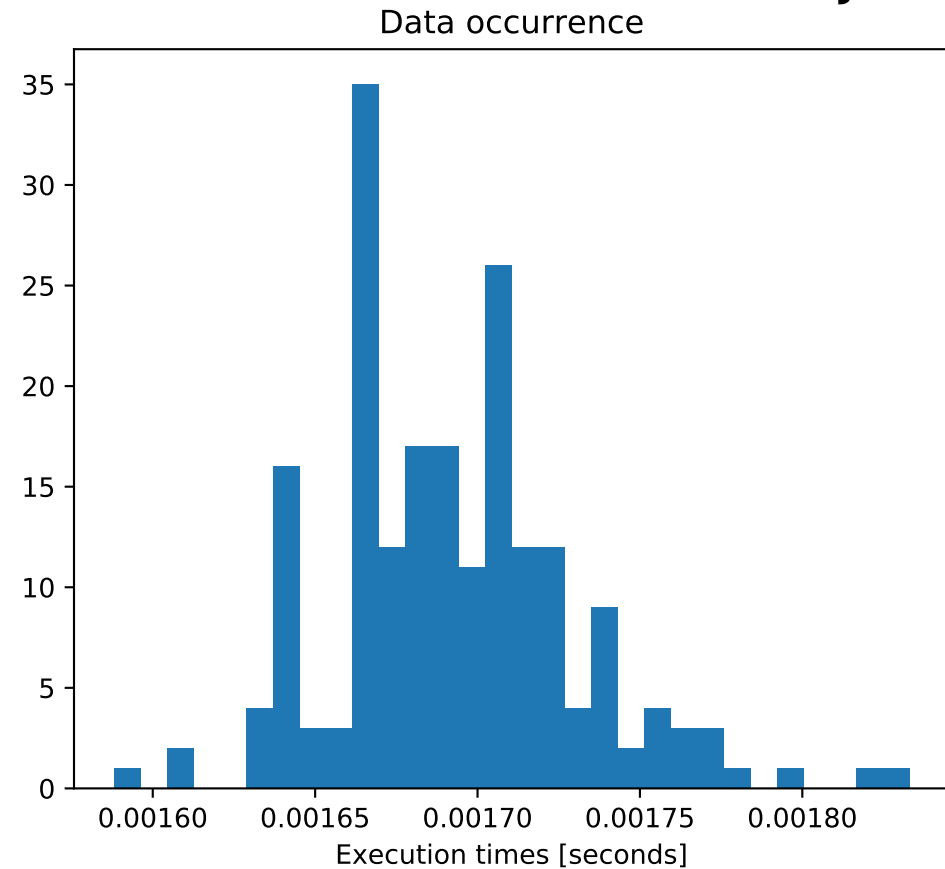
GESUMMV (IID with p-value 0.0674368936)



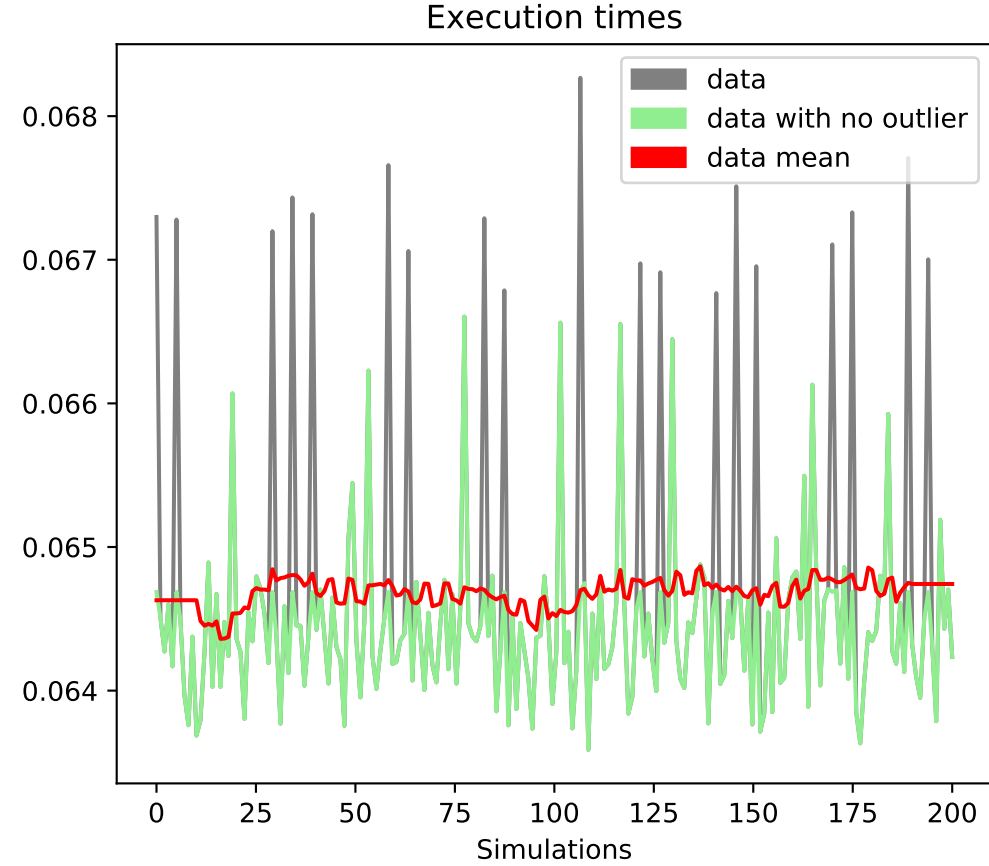
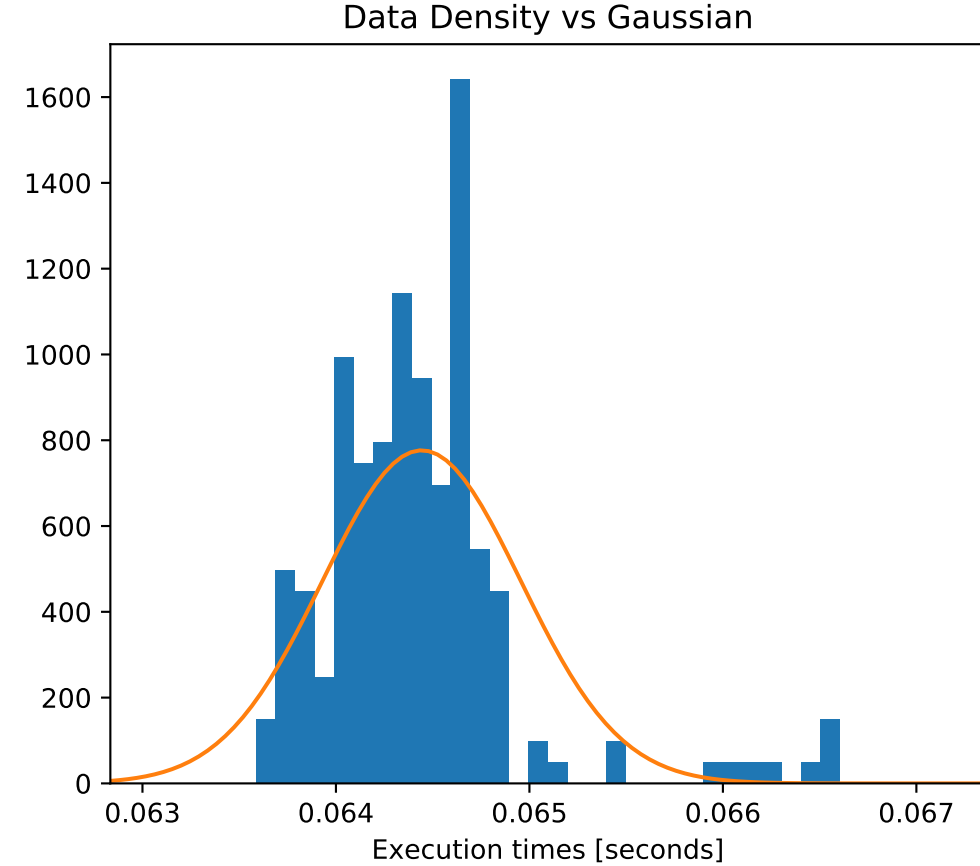
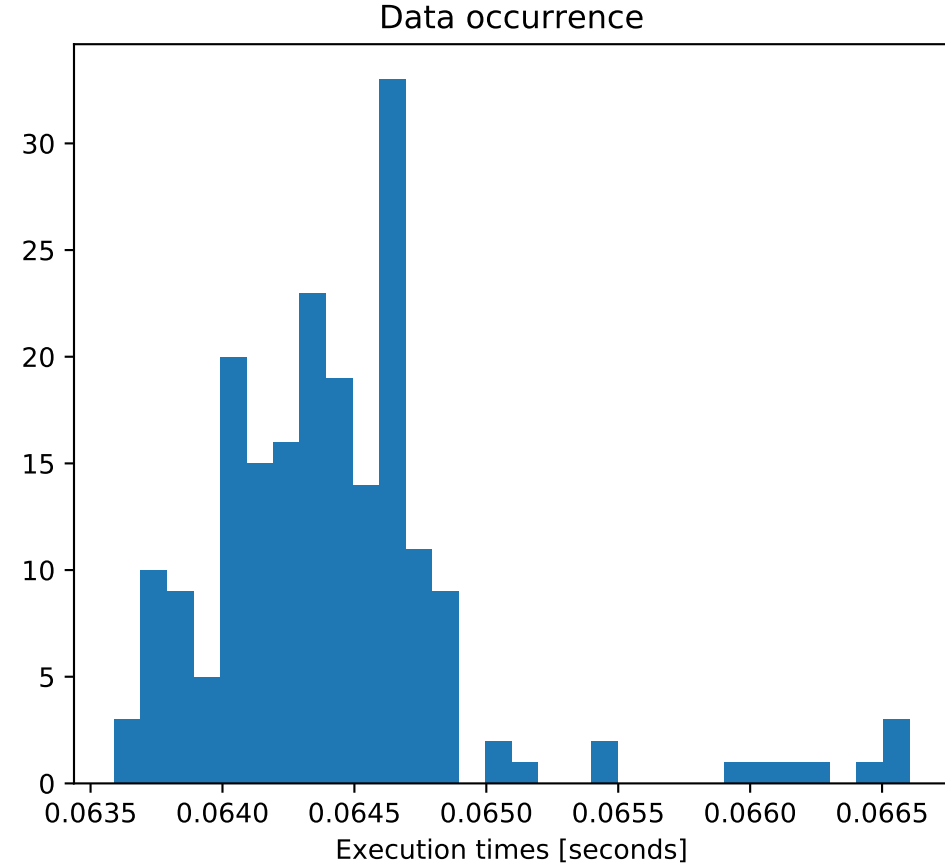
GRAMSCHMIDT (IID with p-value 0.3549676307)



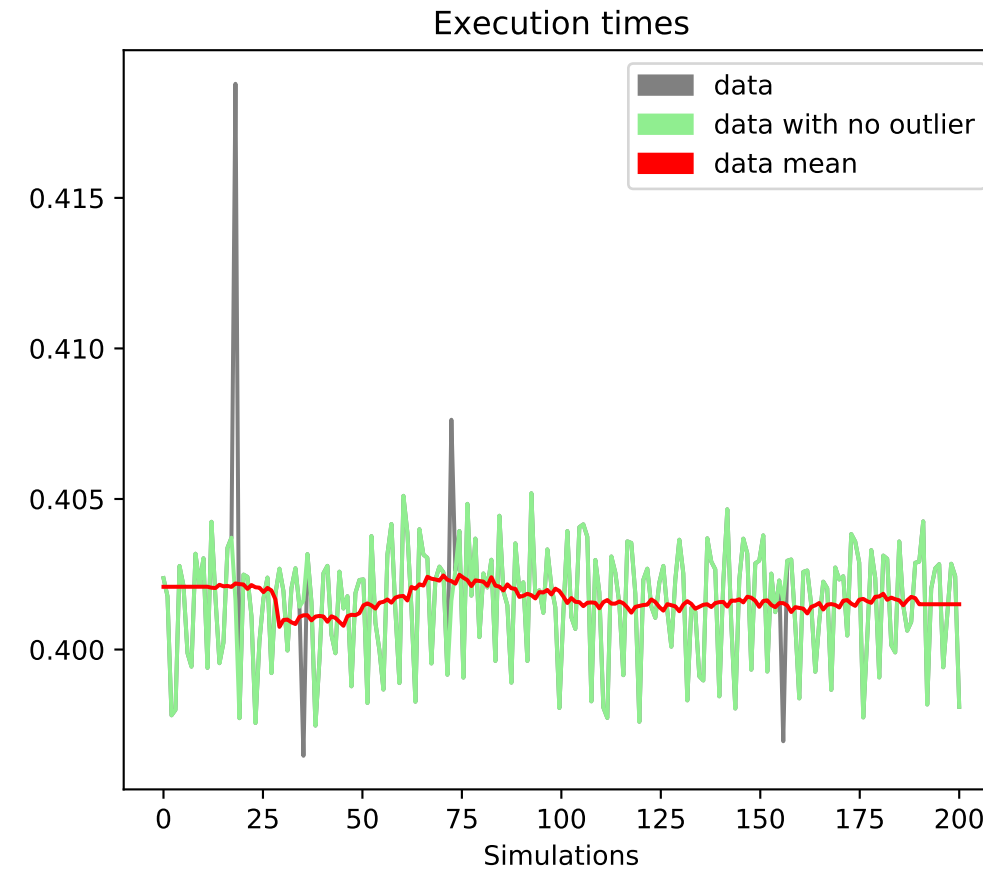
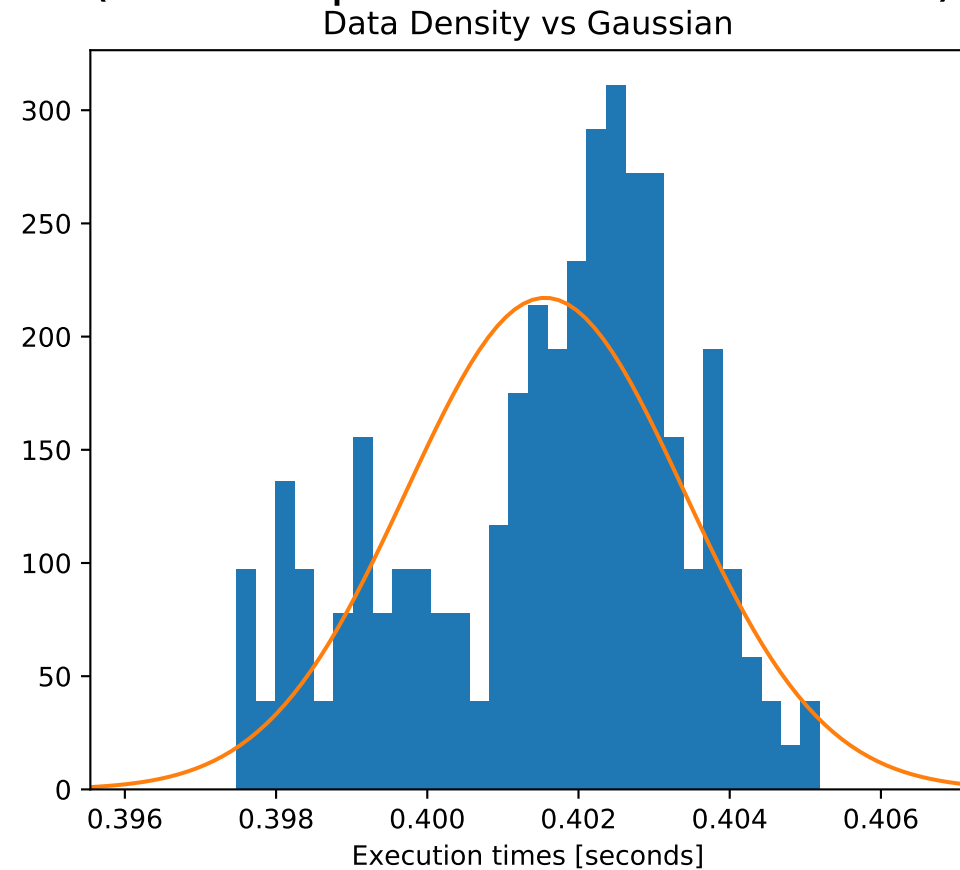
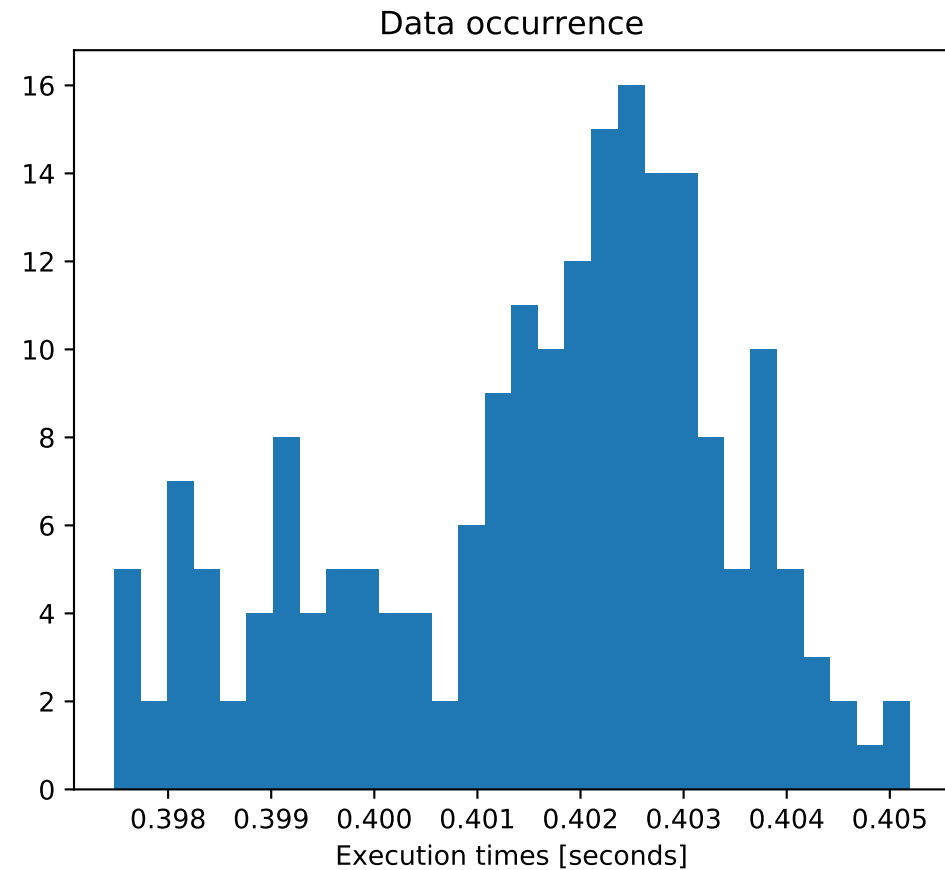
JACOBI-1D-IMPER (IID with p-value 0.0793504386)



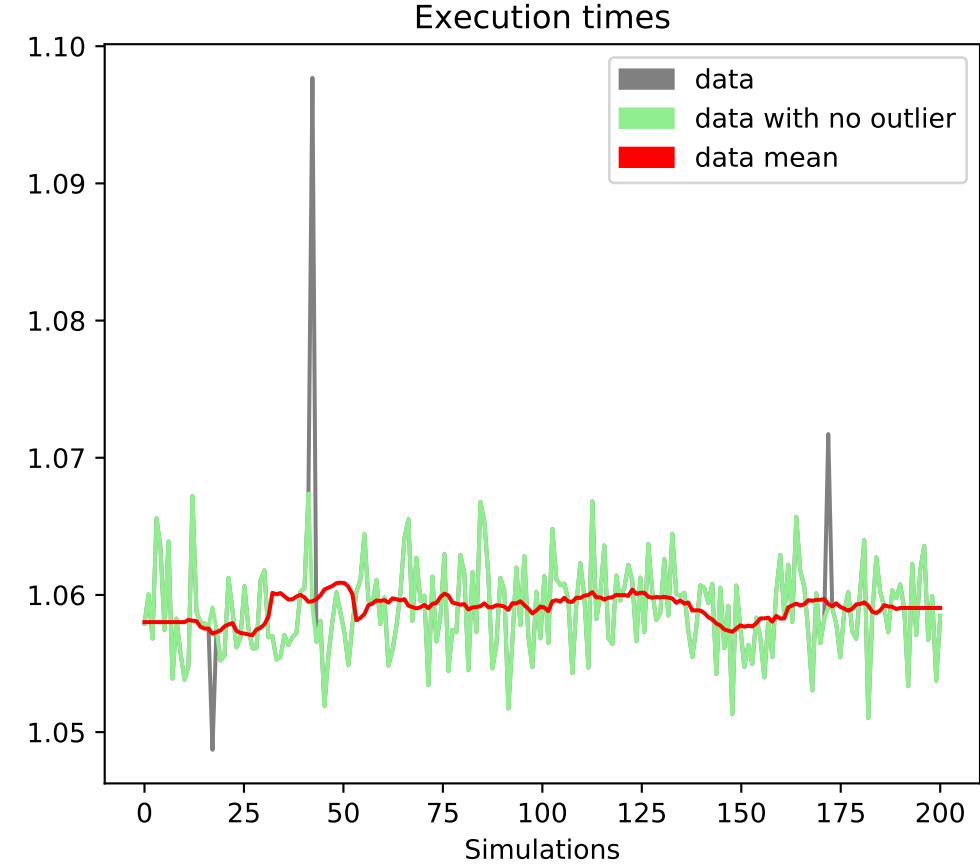
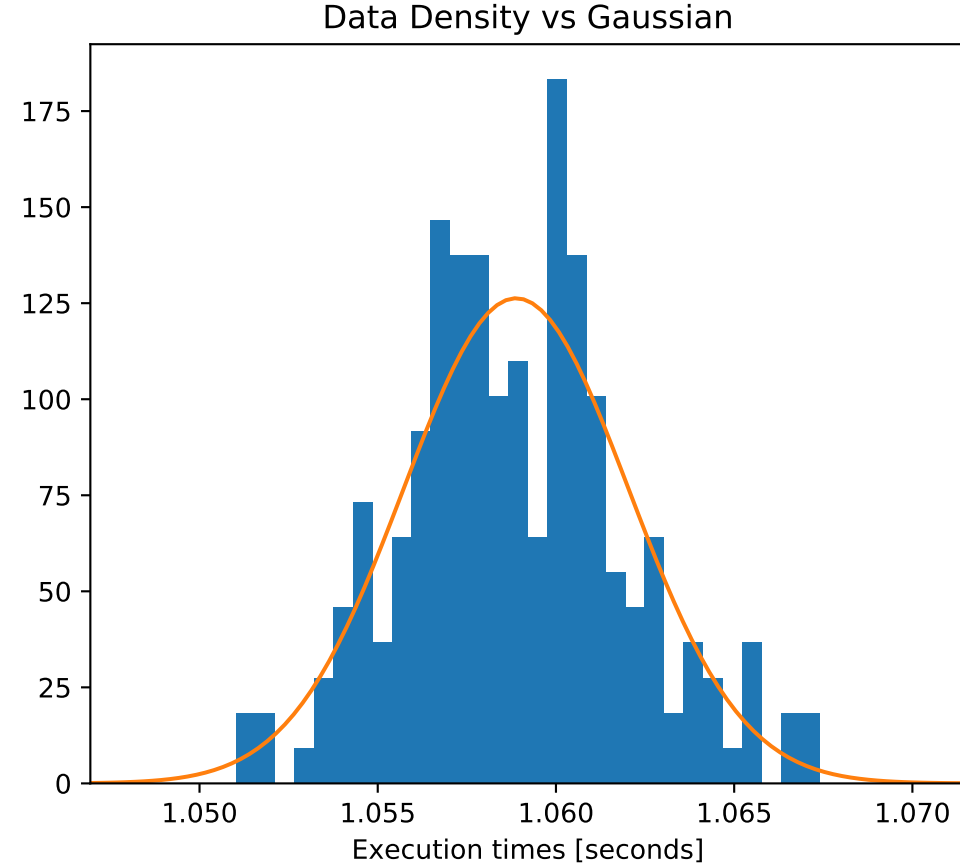
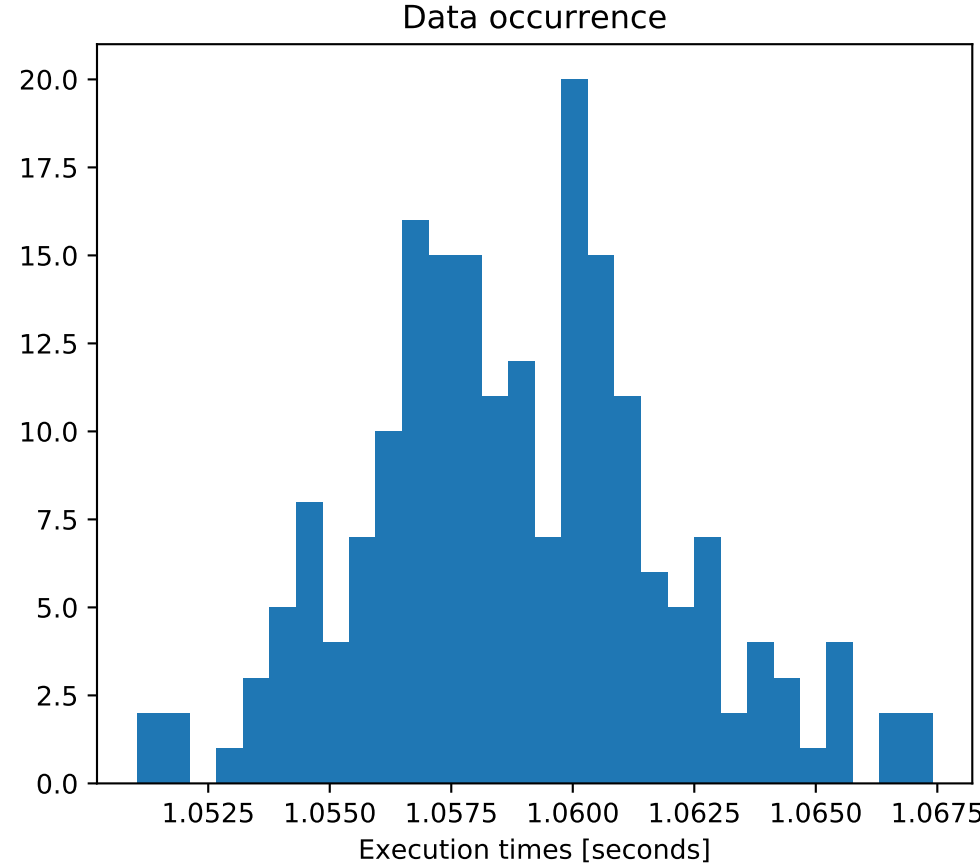
JACOBI-2D-IMPER (IID with p-value 0.4050985017)



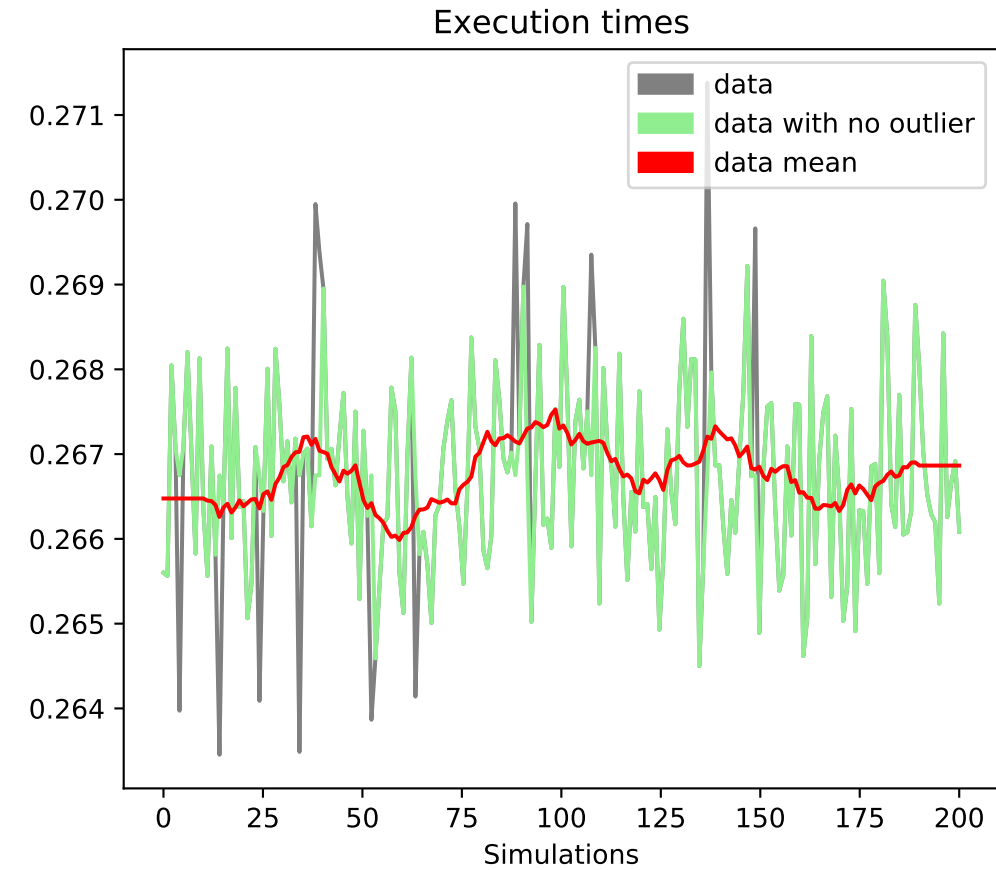
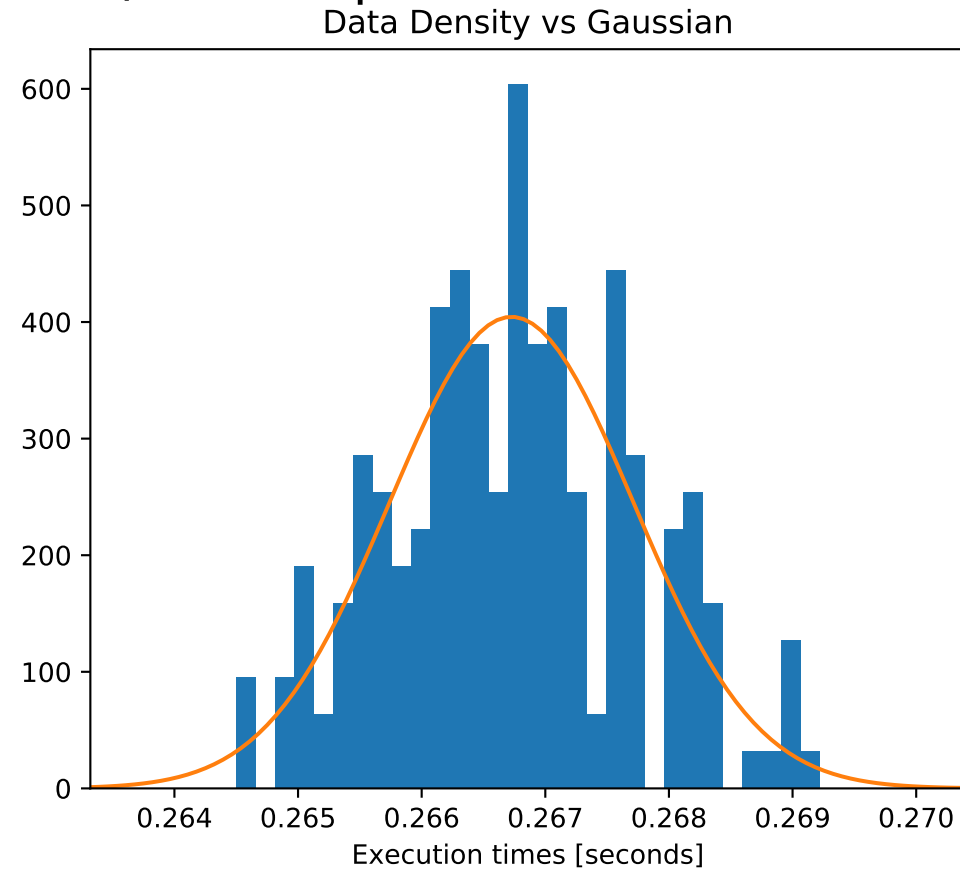
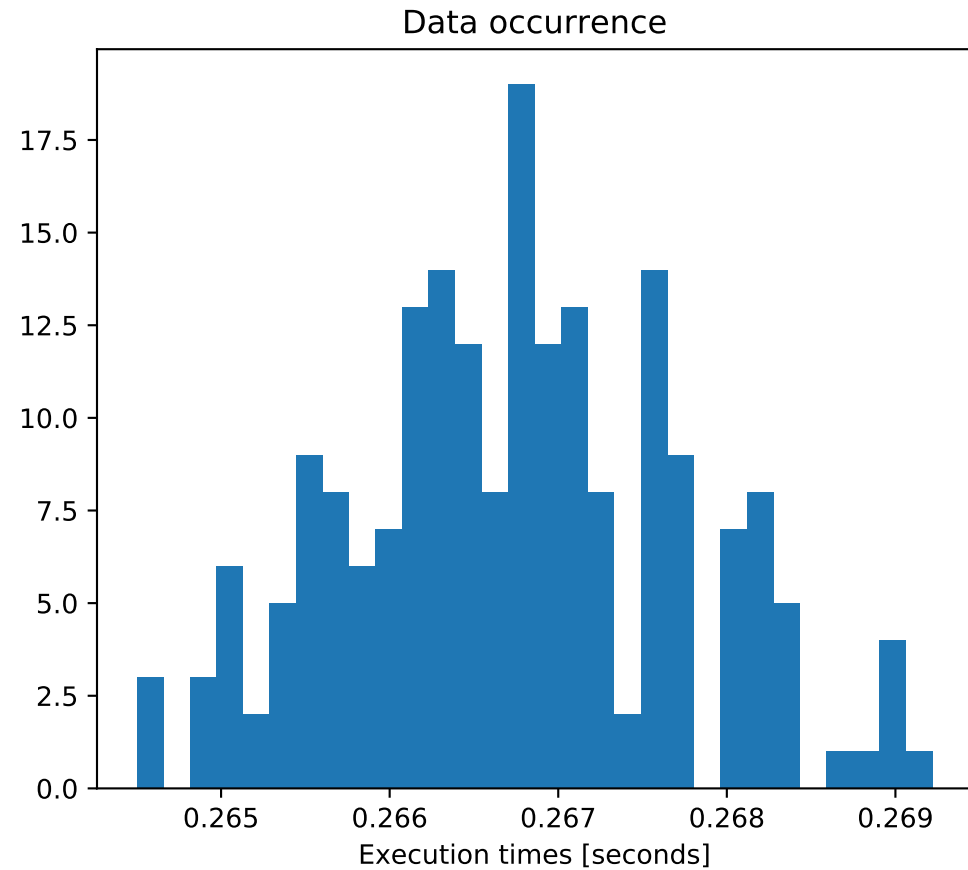
LU (IID with p-value 0.7940641206)



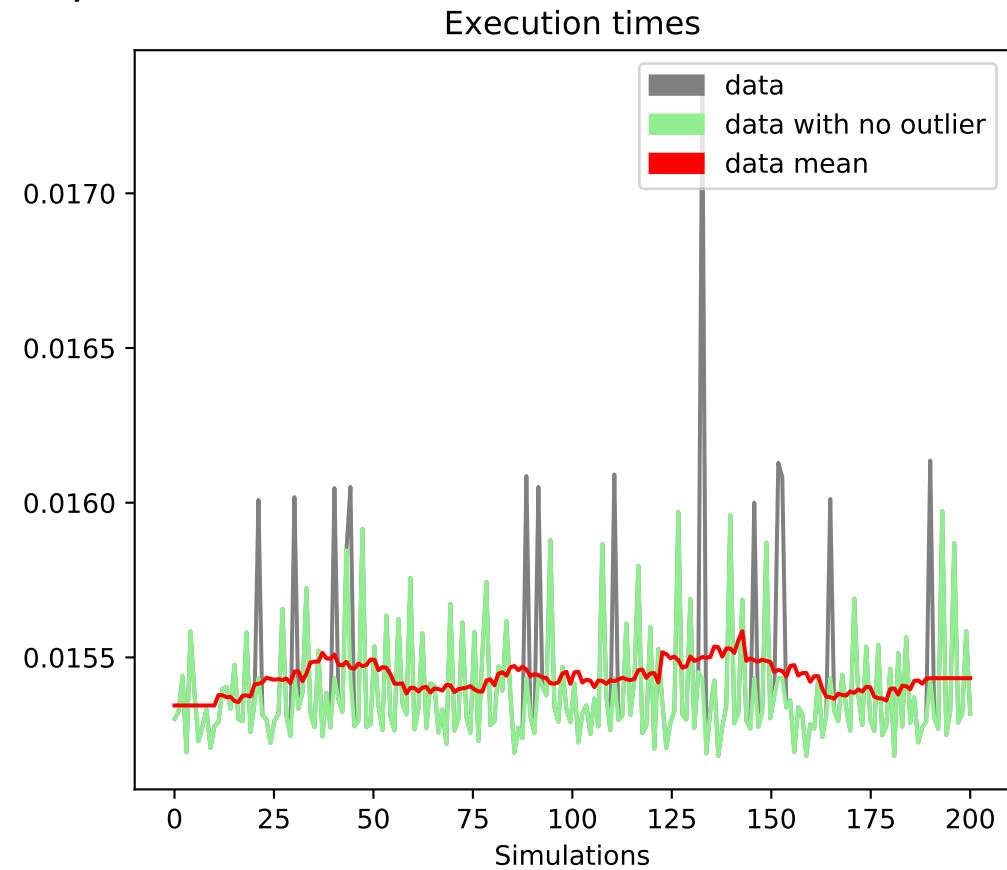
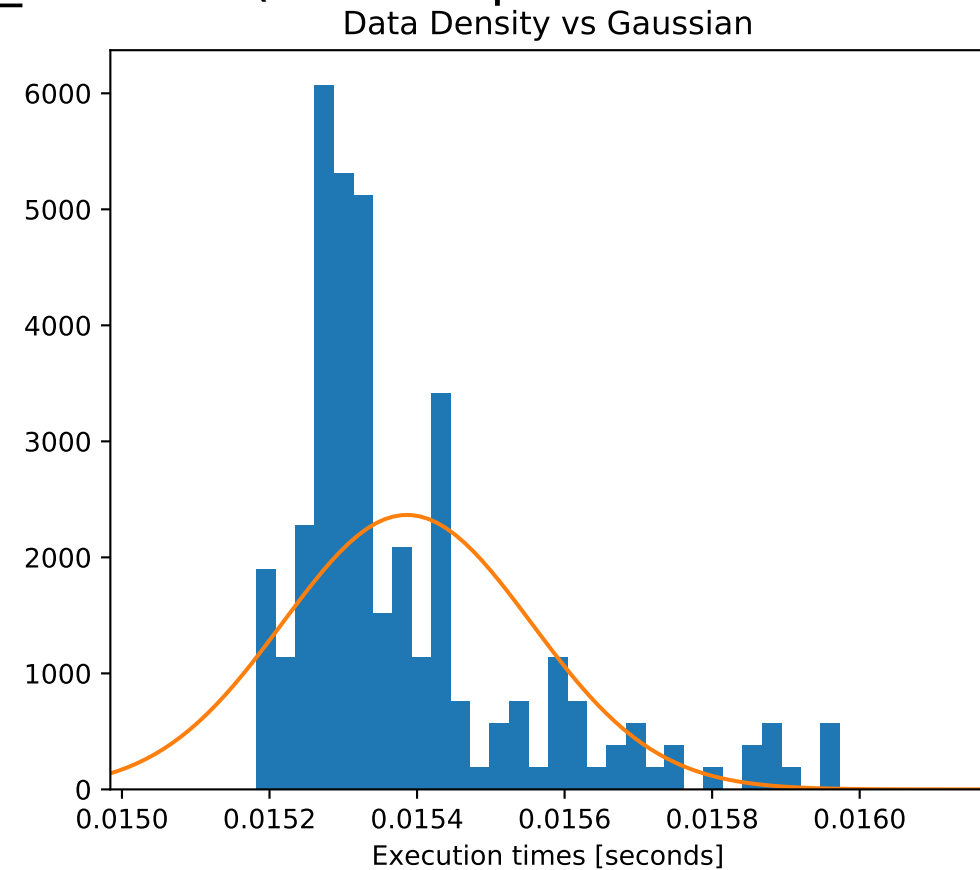
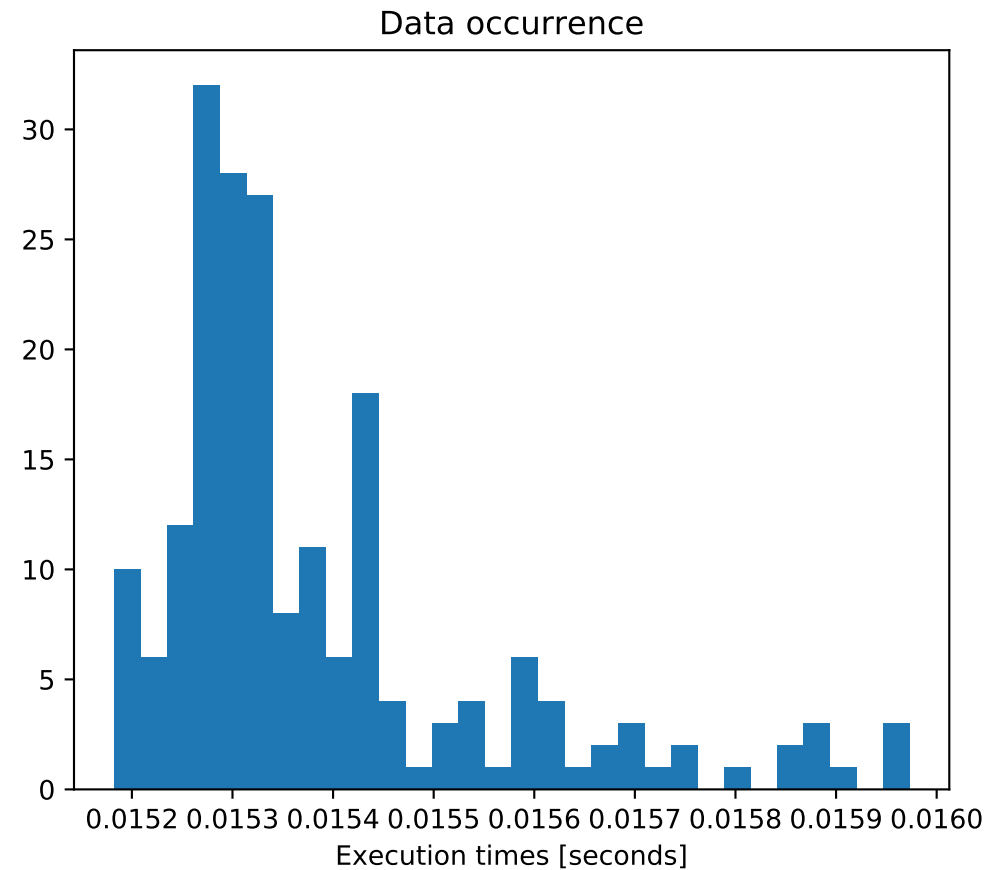
LUDCMP (IID with p-value 0.4623868991)



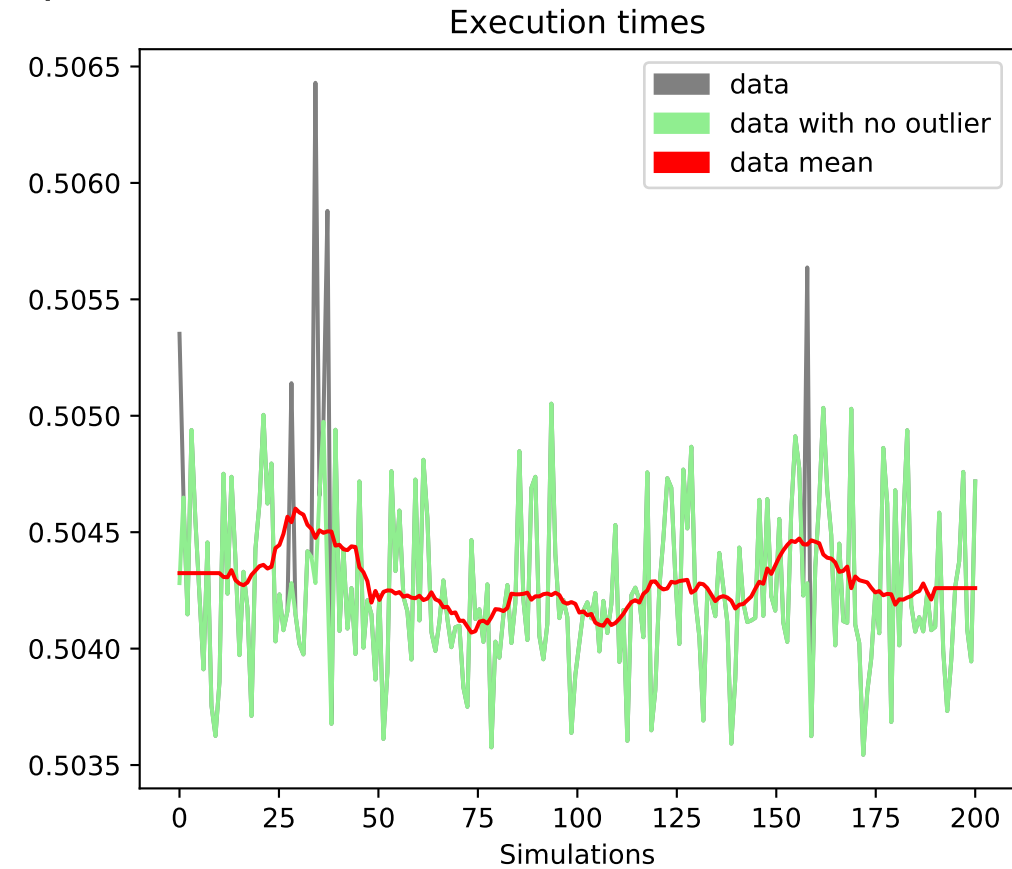
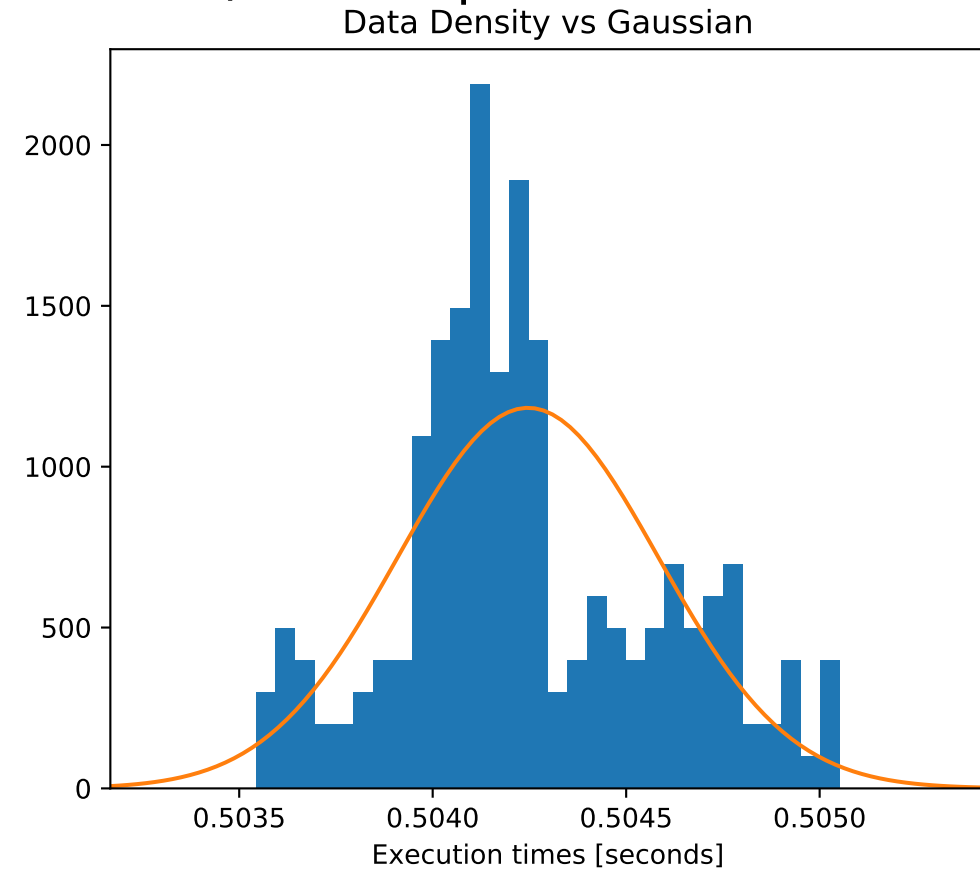
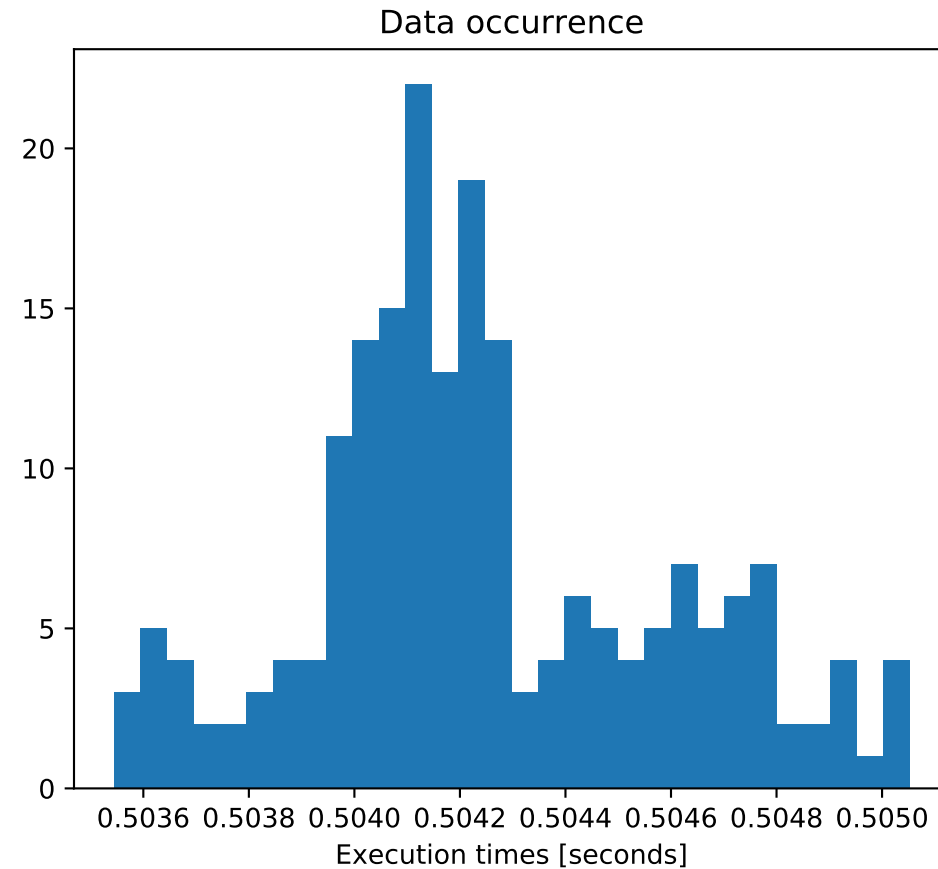
MVT (IID with p-value 0.9594452095)



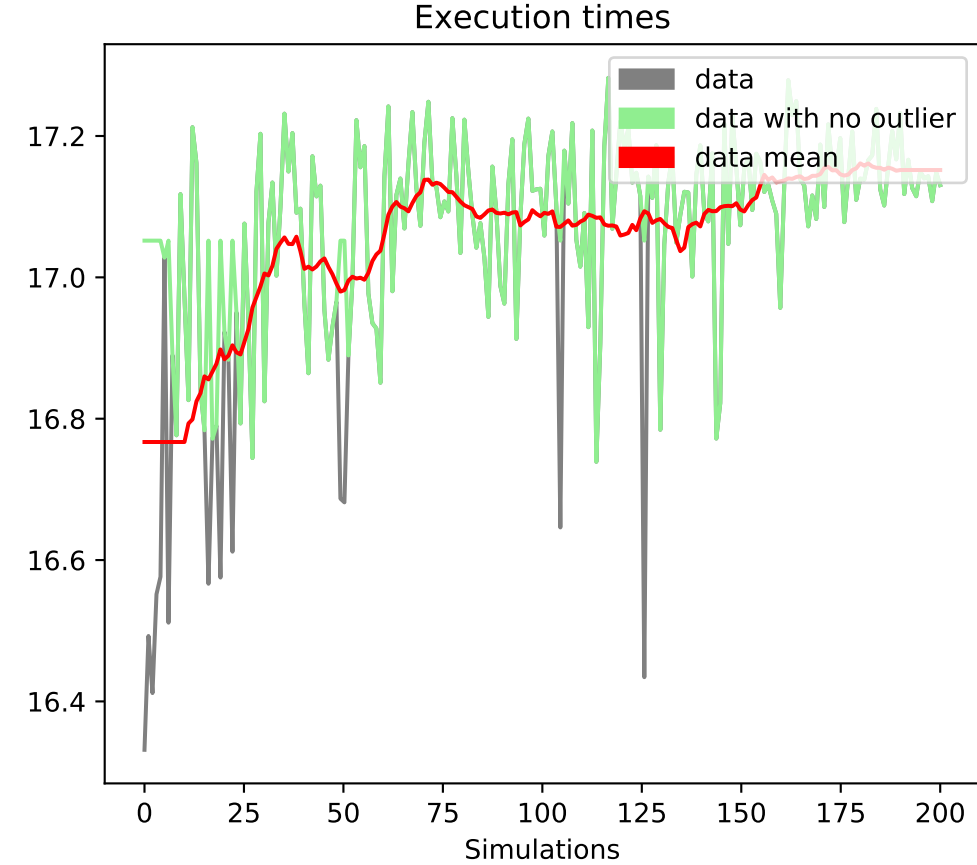
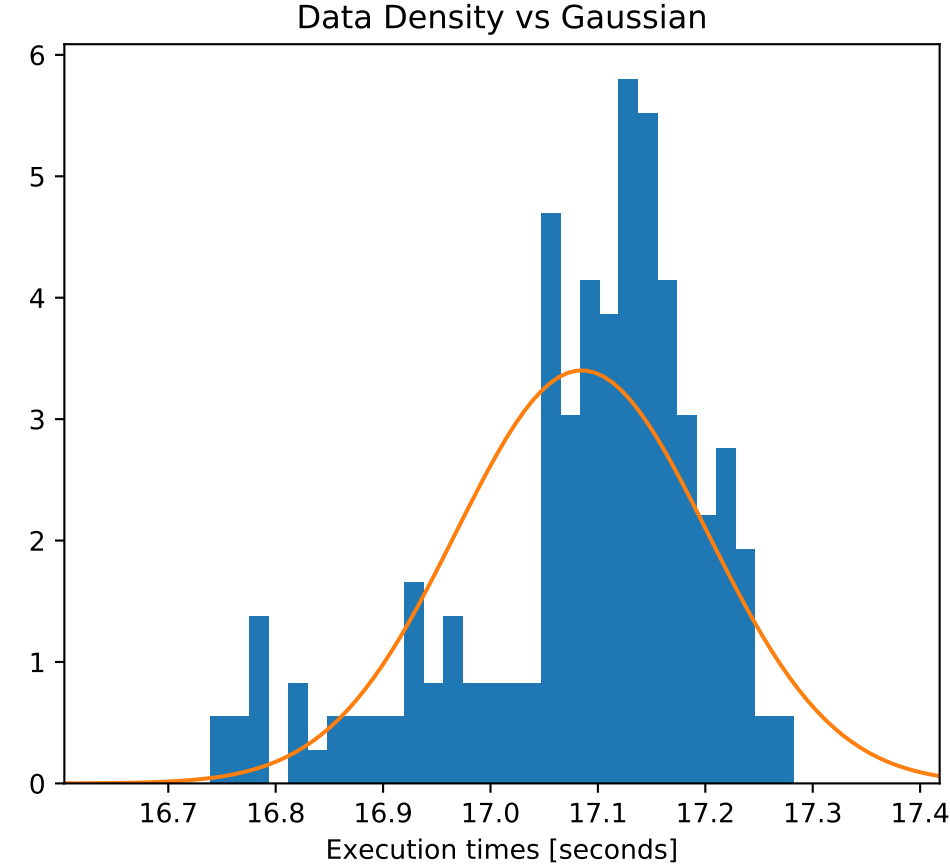
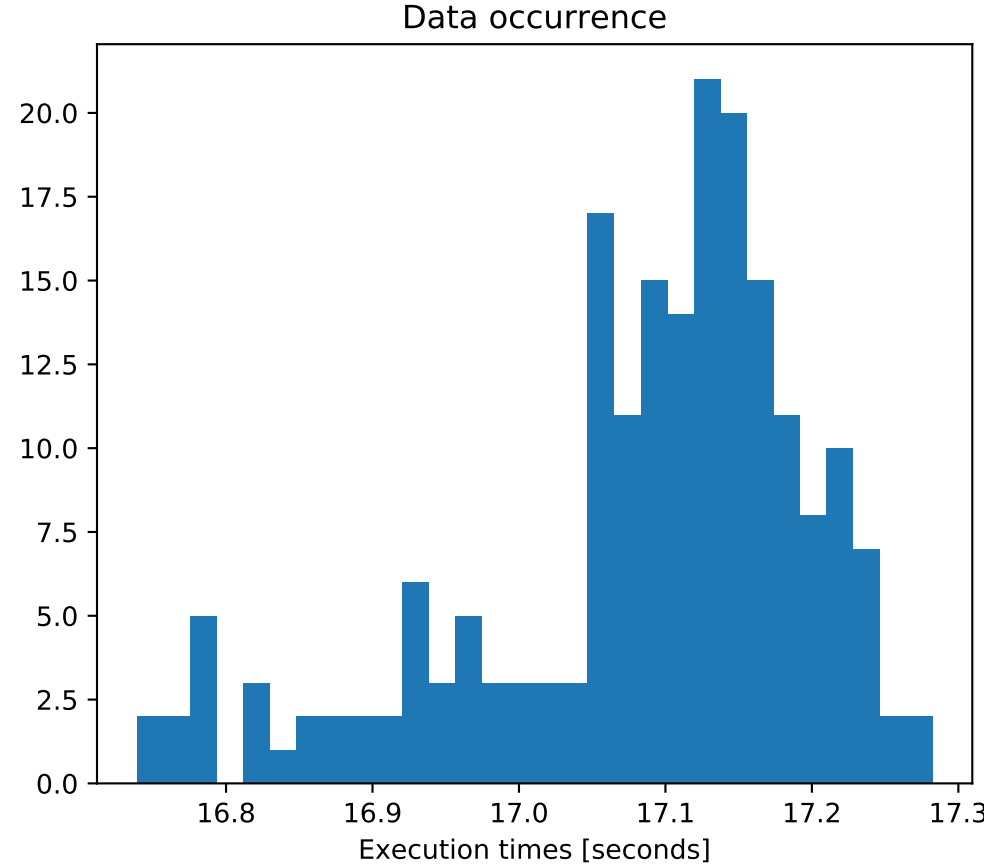
REG_DETECT (IID with p-value 0.9848576449)



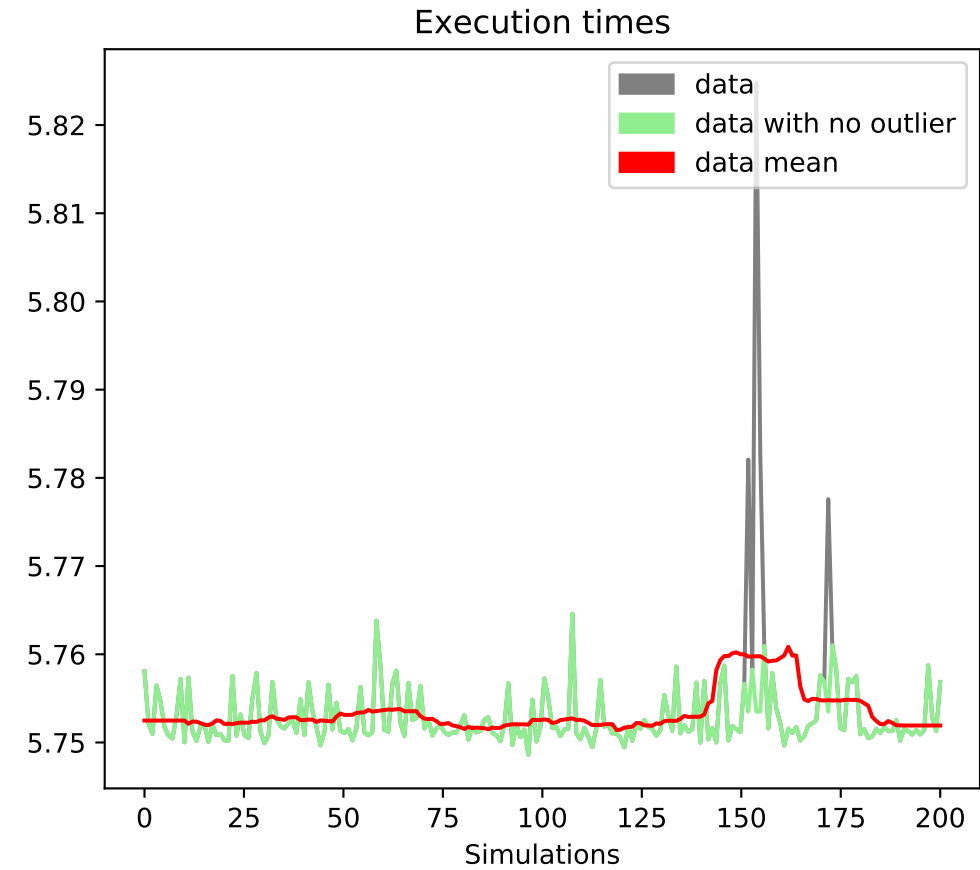
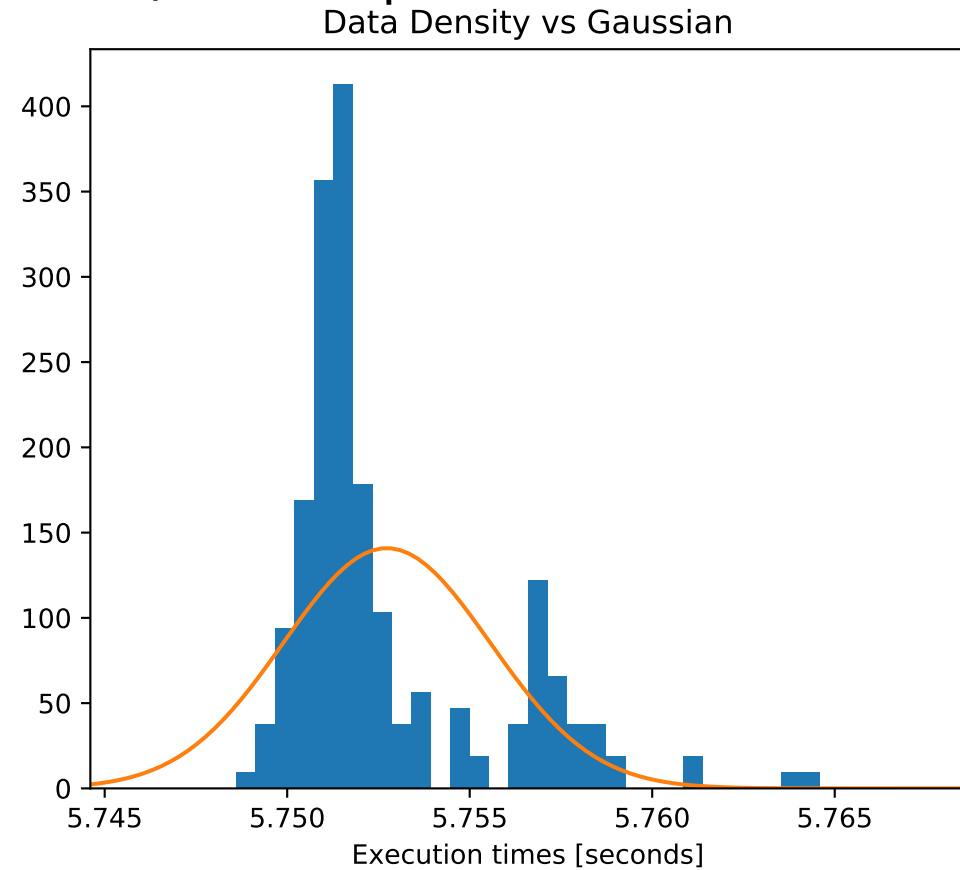
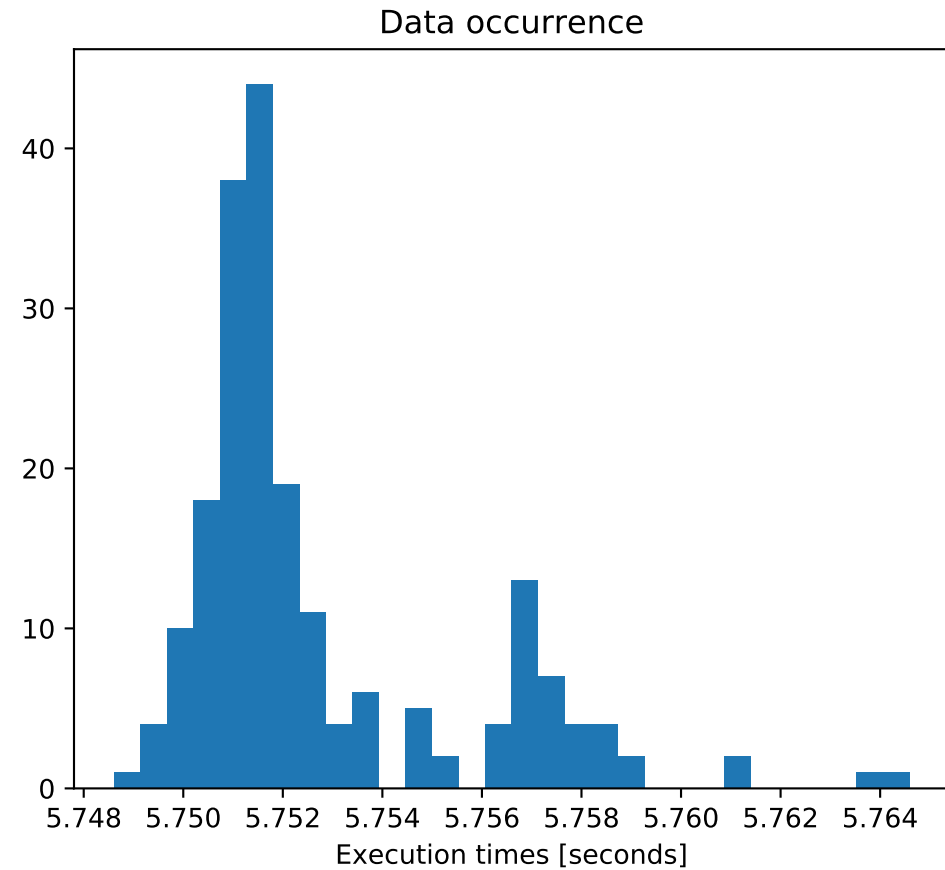
SEIDEL-2D (IID with p-value 0.8573258743)



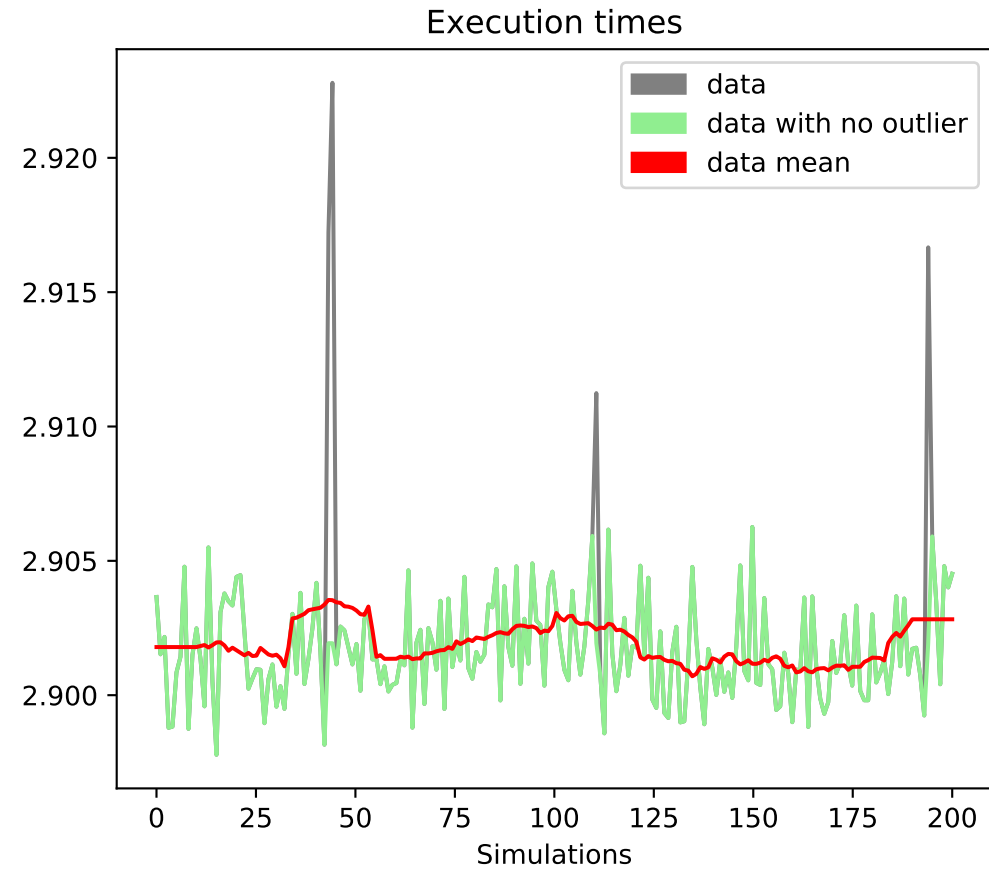
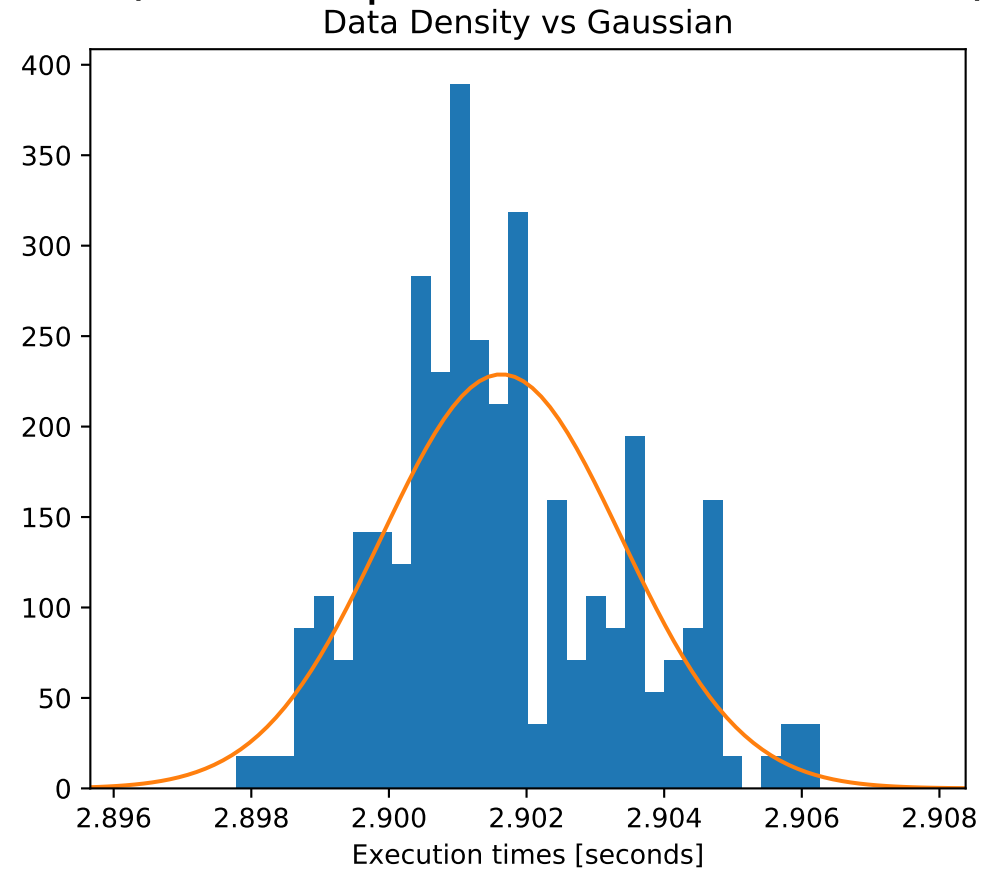
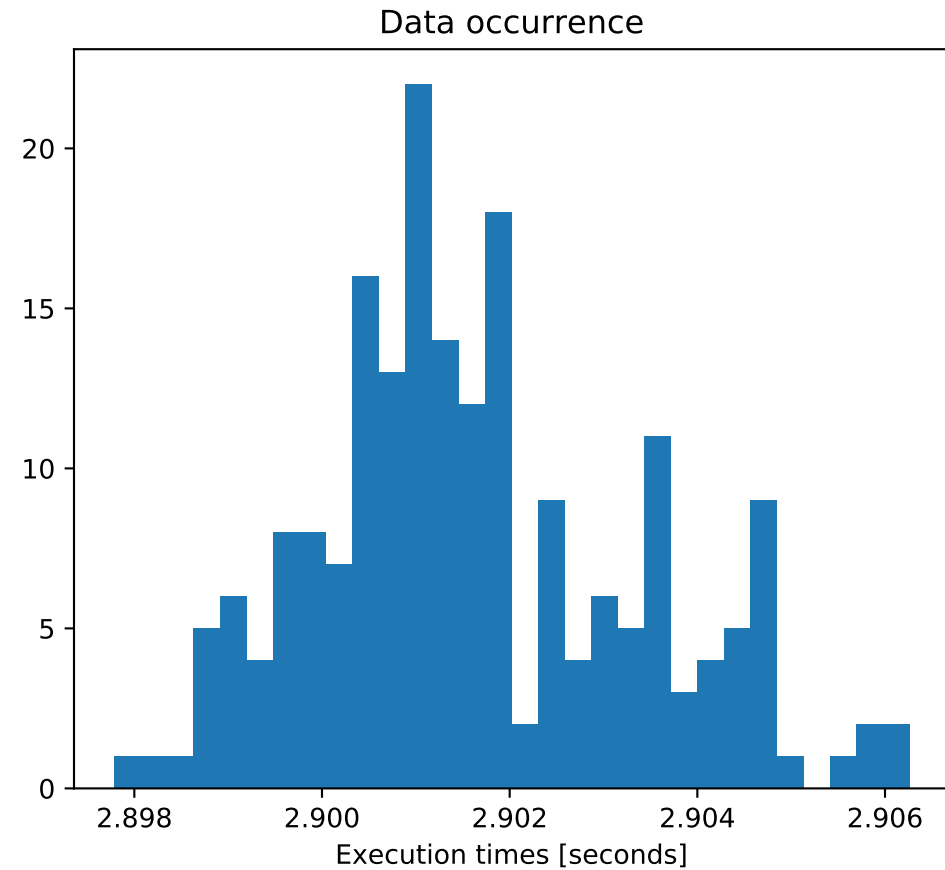
SYMM (NOT IID with p-value 1.19615e-05)



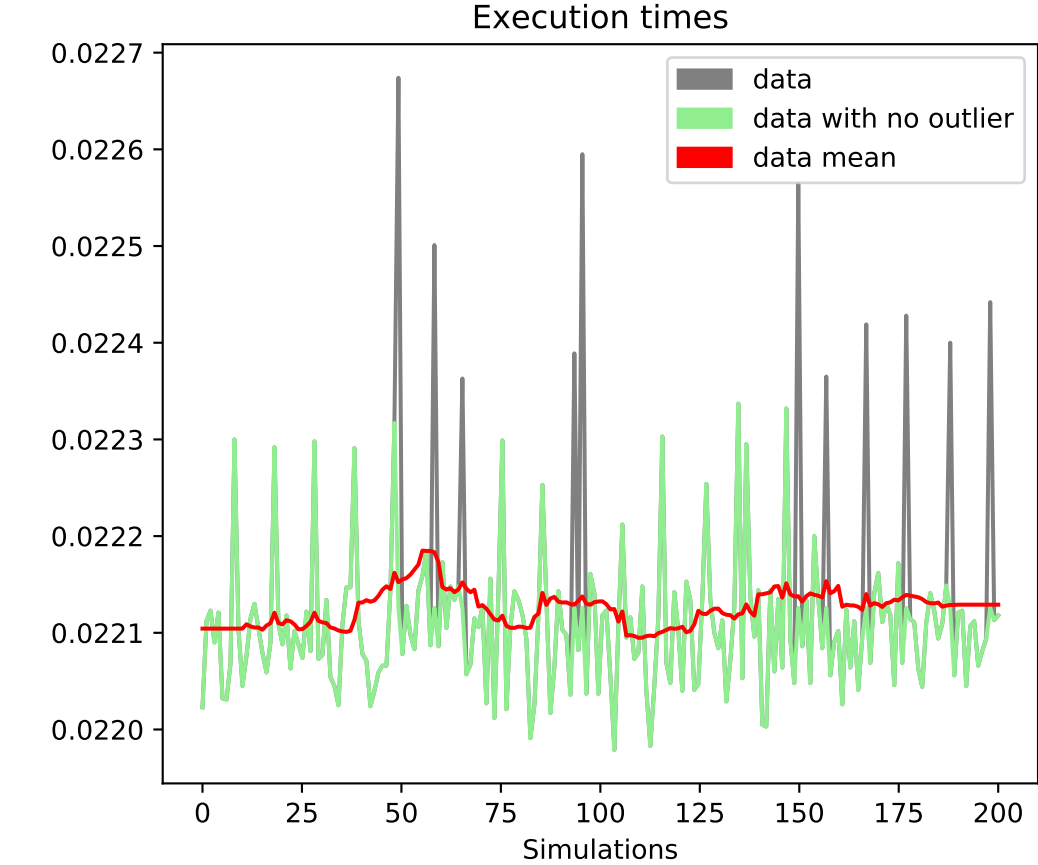
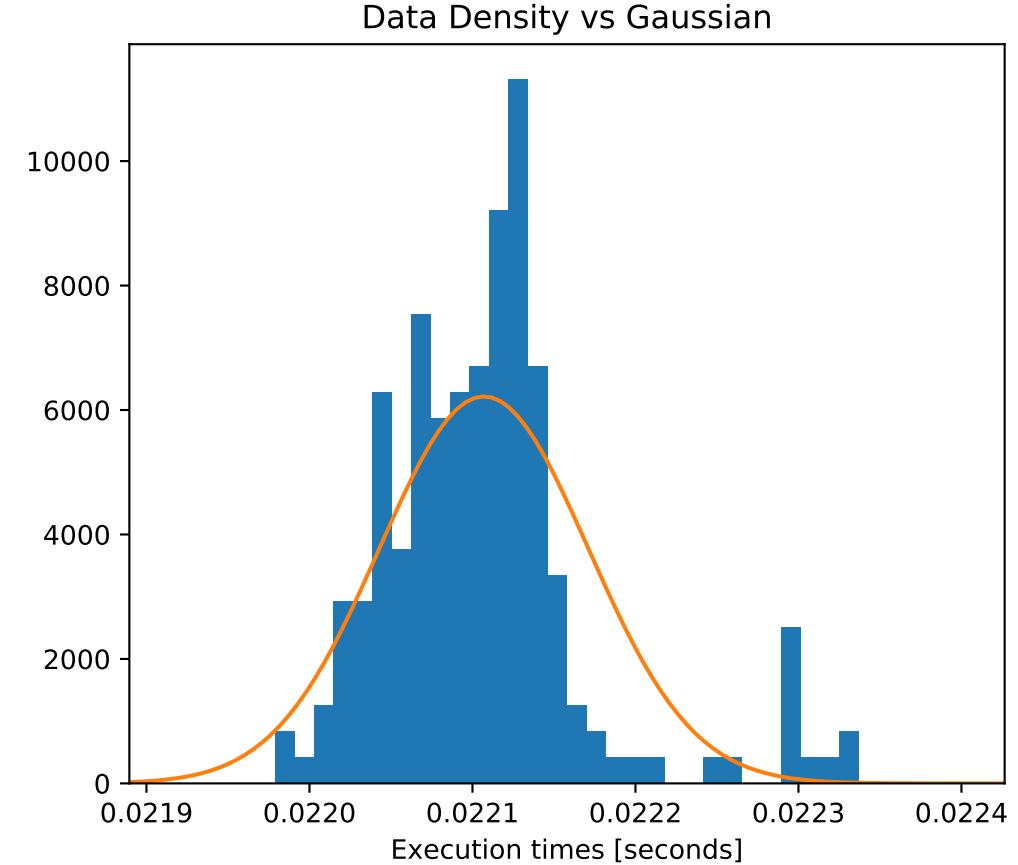
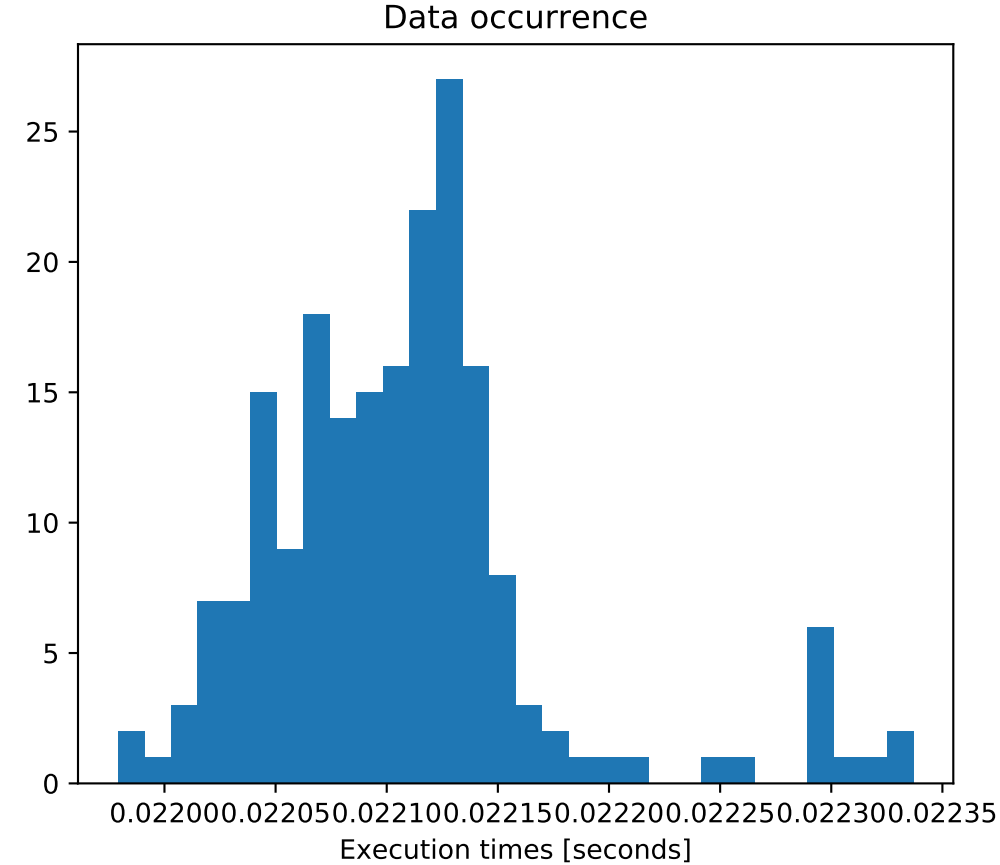
SYR2K (IID with p-value 0.3658504096)



SYRK (IID with p-value 0.3953499915)



TRISOLV (IID with p-value 0.8784999741)



TRMM (IID with p-value 0.0638468825)

