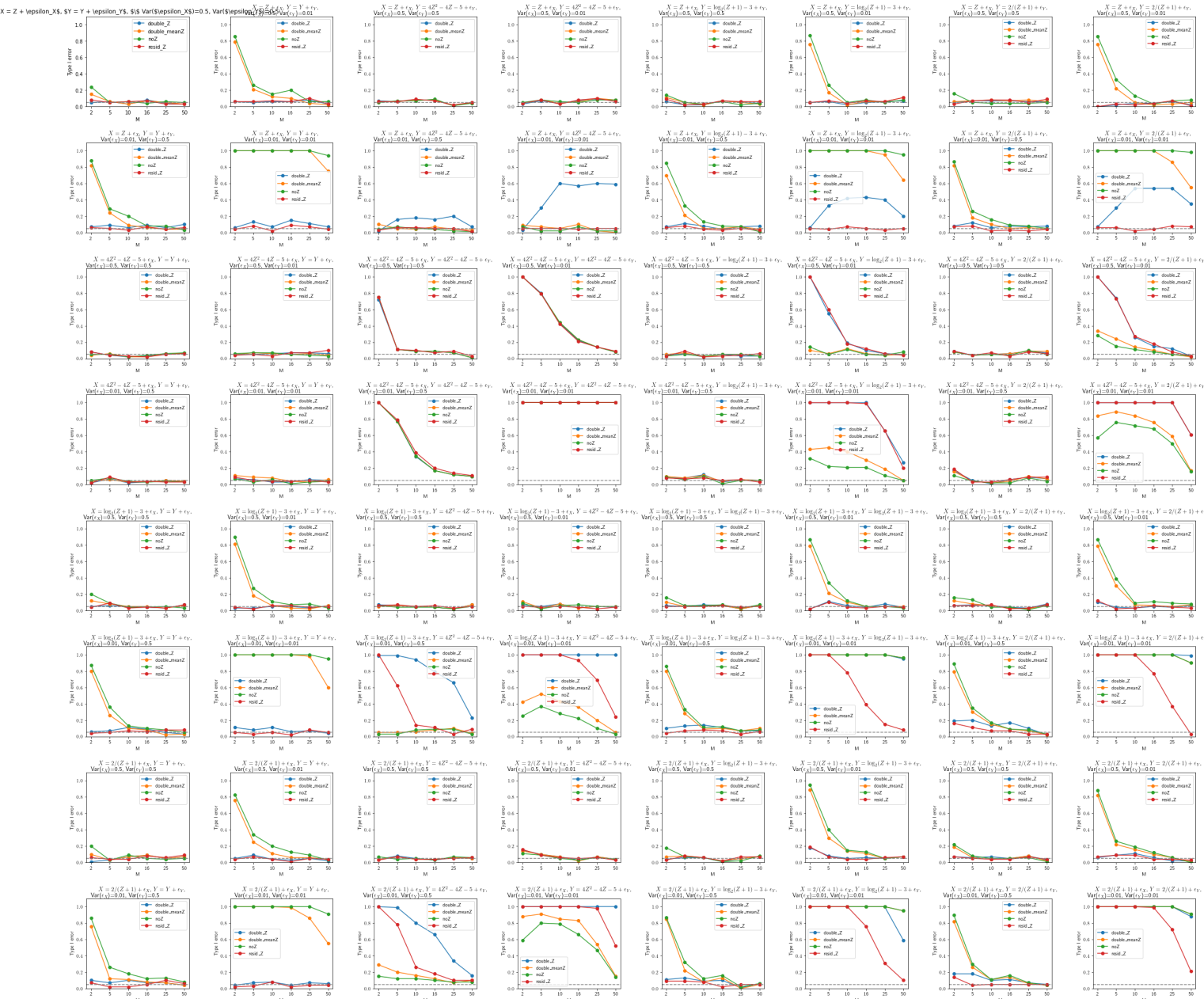
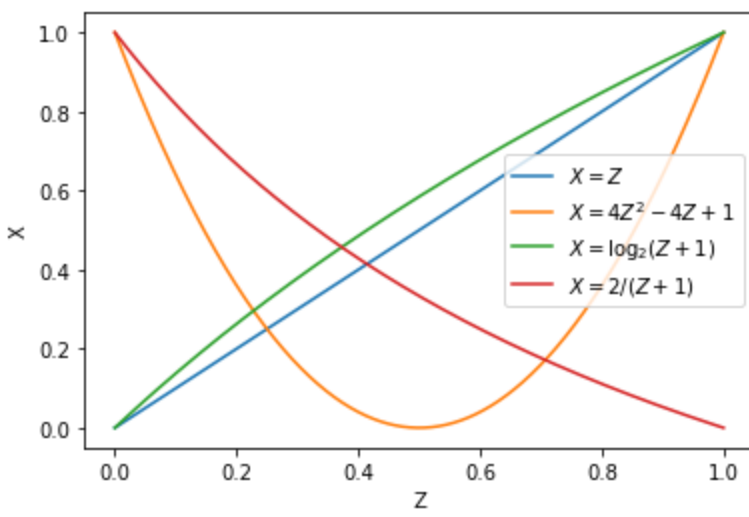


Simulation result correction

Normal Distribution





Analysis

- Methods: (\tilde{Z} is the discretized Z , and the data belonging to the same group share the same \tilde{Z} .)
 - "double_Z": permute X within each bin. At each time, regress Y on 1, Z and regress X on 1, Z separately, and take the *absolute* correlation between residuals from two linear regressions as the test statistic.

$$\text{cor}(P_{Z^\perp} X_\sigma, P_{Z^\perp} Y) = \frac{\mathbb{E}(X_\sigma^\top P_{Z^\perp} P_{Z^\perp} Y) - \mathbb{E}(P_{Z^\perp} X_\sigma) \mathbb{E}(P_{Z^\perp} Y)}{\sqrt{\text{Var}(P_{Z^\perp} X_\sigma) \text{Var}(P_{Z^\perp} Y)}} \quad (1)$$

$$\propto \frac{\mathbb{E}(X_\sigma^\top P_{Z^\perp} Y)}{\text{SD}(X_\sigma)} \quad (2)$$

- "double_meanZ": permute X within each bin. At each time, regress Y on 1, \tilde{Z} and regress X on 1, \tilde{Z} separately, and take the *absolute* correlation between residuals from two linear regressions as the test statistic.

$$\text{cor}(P_{\tilde{Z}^\perp} X_\sigma, P_{\tilde{Z}^\perp} Y) = \frac{\mathbb{E}(X_\sigma^\top P_{\tilde{Z}^\perp} P_{\tilde{Z}^\perp} Y) - \mathbb{E}(P_{\tilde{Z}^\perp} X_\sigma) \mathbb{E}(P_{\tilde{Z}^\perp} Y)}{\sqrt{\text{Var}(P_{\tilde{Z}^\perp} X_\sigma) \text{Var}(P_{\tilde{Z}^\perp} Y)}} \quad (3)$$

$$\propto \mathbb{E}(X_\sigma^\top P_{\tilde{Z}^\perp} Y) \quad (4)$$

- "noZ": use $\text{cor}(X, Y)$ as test statistic with local permutation in X with respect to Z .
- "resid_Z": regress Y on 1, Z and regress X on 1, Z separately. Permute residuals from regression on for X and take the *absolute* correlation between permuted residuals as the test statistic.

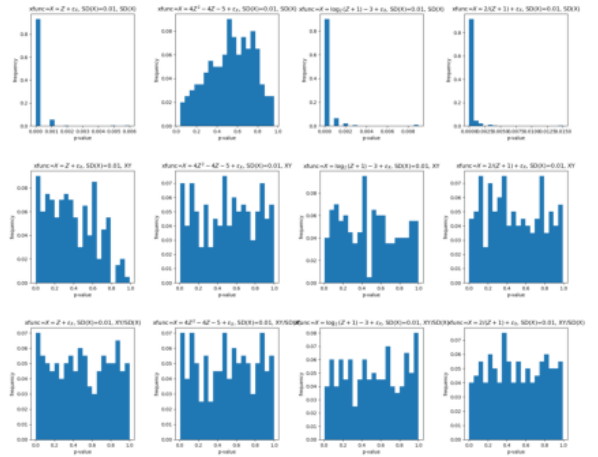
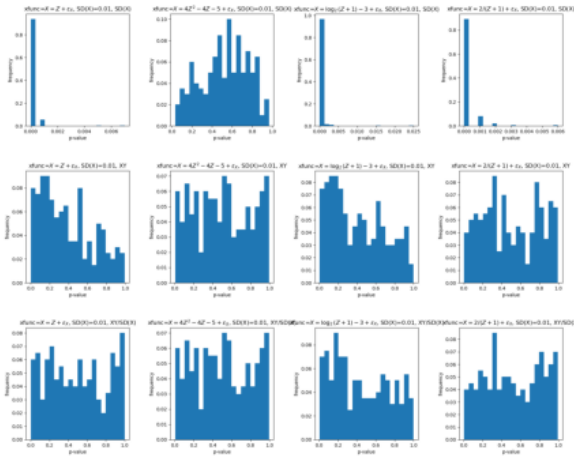
$$\text{cor}((P_{Z^\perp} X)_\sigma, P_{Z^\perp} Y) = \frac{\mathbb{E}((X^\top P_{Z^\perp})_\sigma P_{Z^\perp} Y) - \mathbb{E}((P_{Z^\perp} X)_\sigma) \mathbb{E}(P_{Z^\perp} Y)}{\sqrt{\text{Var}((P_{Z^\perp} X)_\sigma) \text{Var}(P_{Z^\perp} Y)}} \quad (5)$$

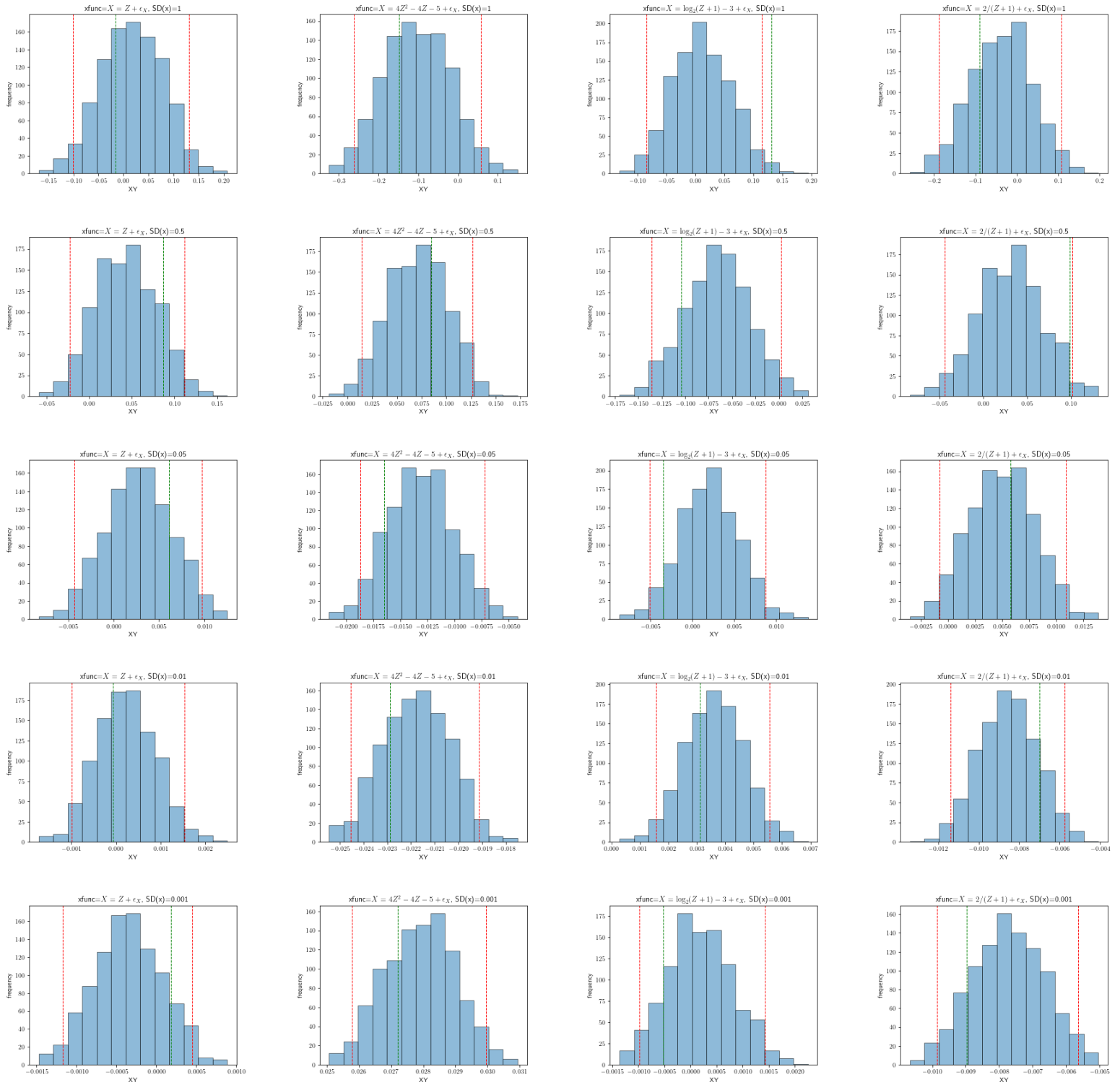
$$\propto \mathbb{E}((X^\top P_{Z^\perp})_\sigma \epsilon_Y) \quad (6)$$

- Basically "double_meanZ" and "resid_Z" are very similar - both permuting residuals after regressing $X \sim 1 + \tilde{Z}$ and $X \sim 1 + Z$ respectively.
- More generally, "no_Z" is also similar - permuting residuals after regressing $X \sim 1$ but without using information of Z .

double_Z

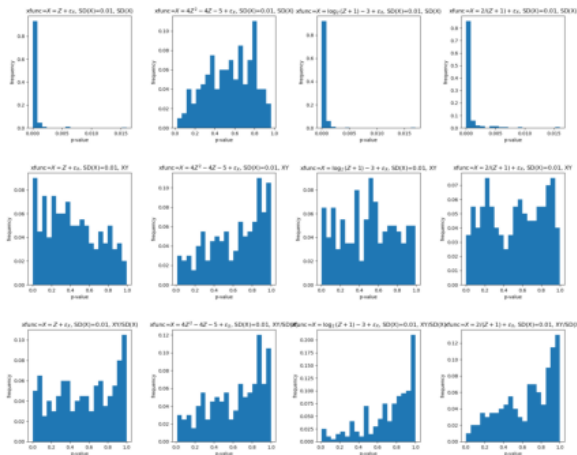
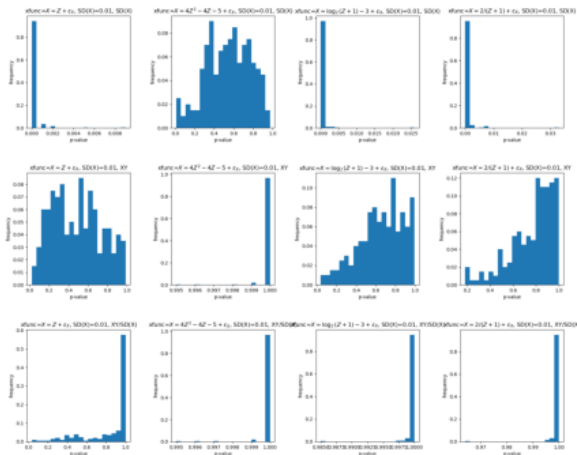
- when Y is linear in Z :
 - sometimes with uniform distribution on $X_\sigma^\top P_{Z^\perp} Y$ and $\text{SD}(X_\sigma)$ (column2)
 - sometimes with extremely small $\text{SD}(X_\sigma)$ but $X_\sigma^\top P_{Z^\perp} Y$ with mean 0 (column1,3,4) \rightarrow uniform distribution on $\frac{X_\sigma^\top P_{Z^\perp} Y}{\text{SD}(X_\sigma)}$

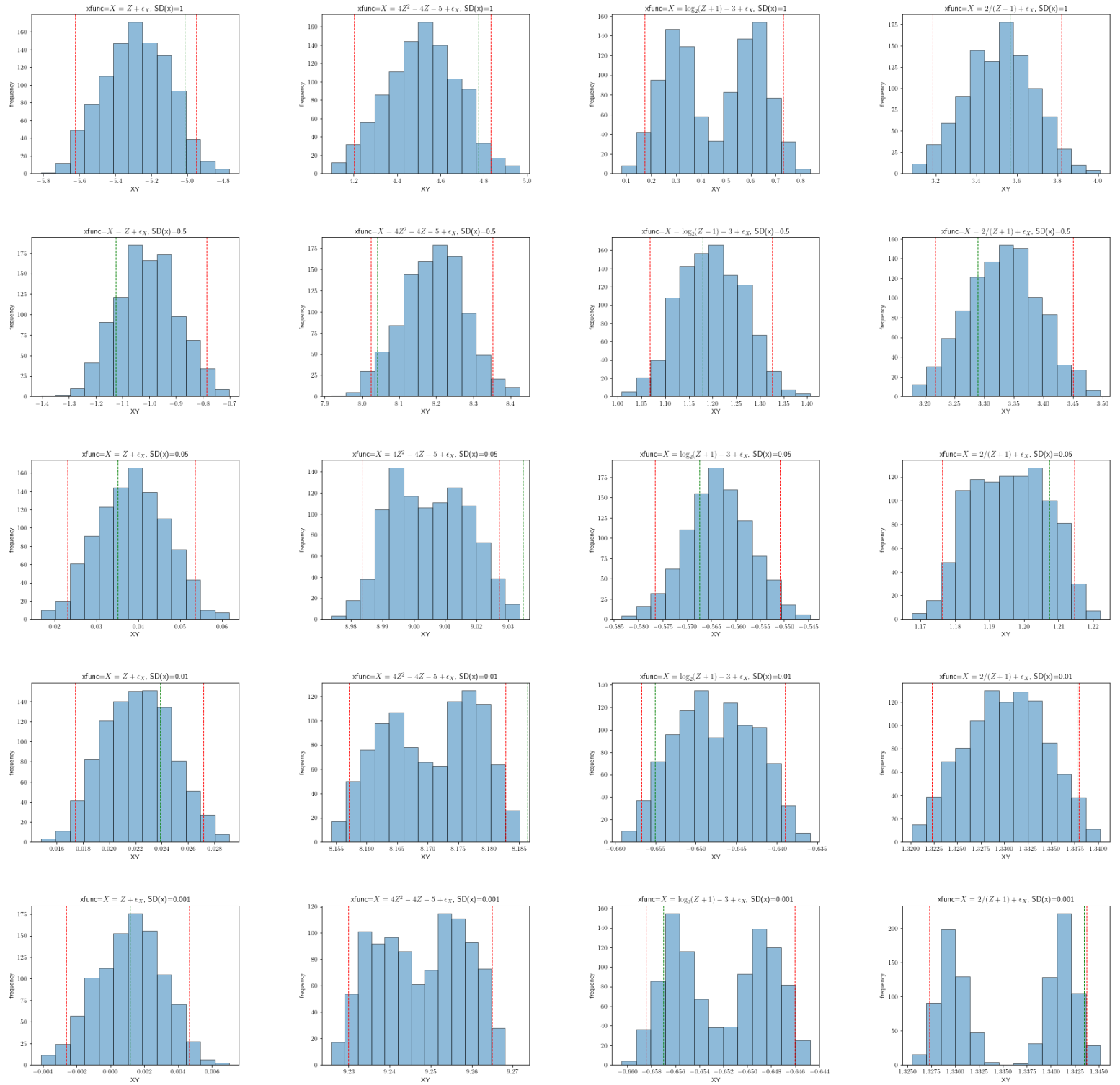




double_Z

- when Y is linear in Z :
 - sometimes with uniform distribution on $\text{SD}(X_\sigma)$ but extremely high $X_\sigma^\top P_{Z^\perp} Y$ (column2) → maybe due to the same generating function?
 - sometimes with extremely small $\text{SD}(X_\sigma)$ and $X_\sigma^\top P_{Z^\perp} Y$ with generally non-zero mean (column1,3,4) → heavy density on large percentile(>0.95) $\frac{X_\sigma^\top P_{Z^\perp} Y}{\text{SD}(X_\sigma)}$

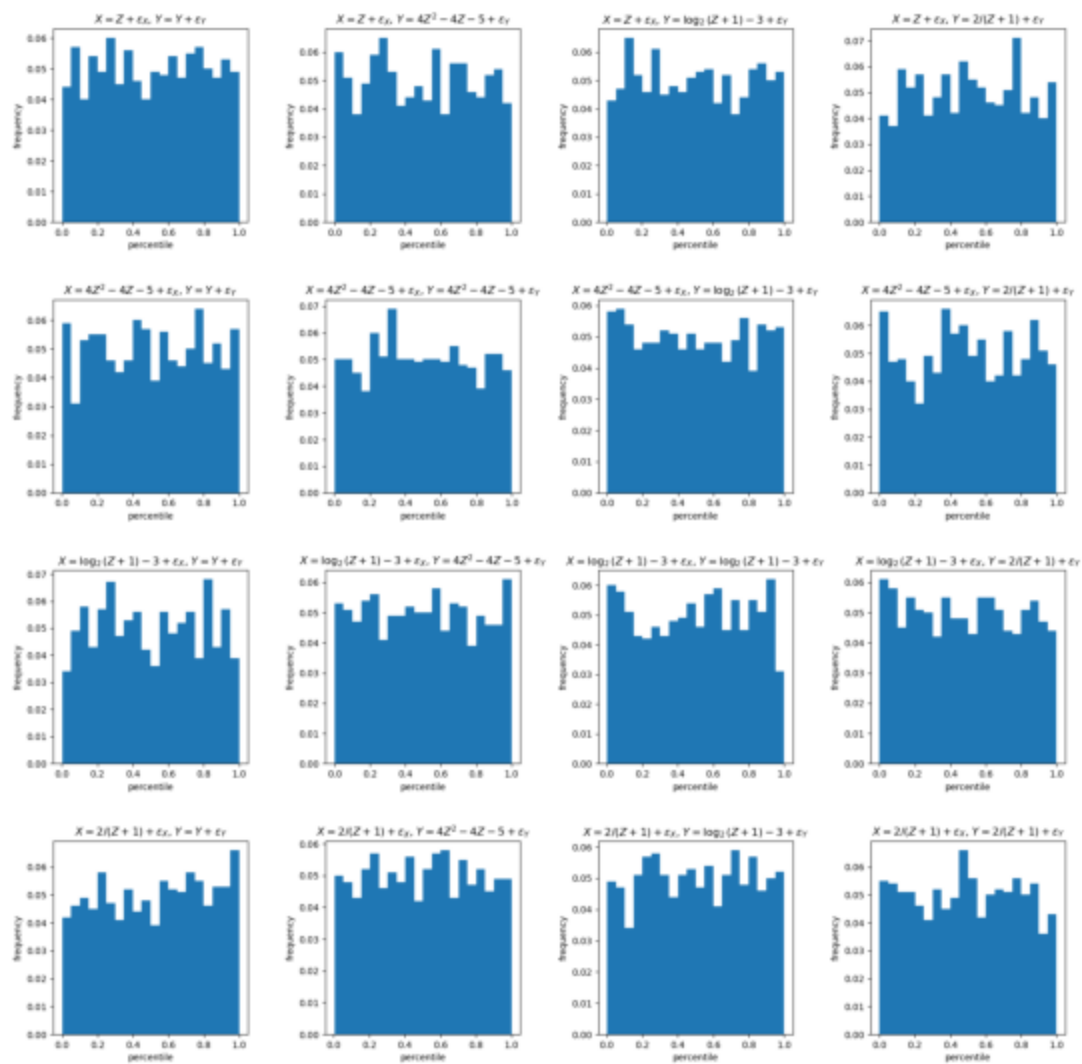




double_meanZ

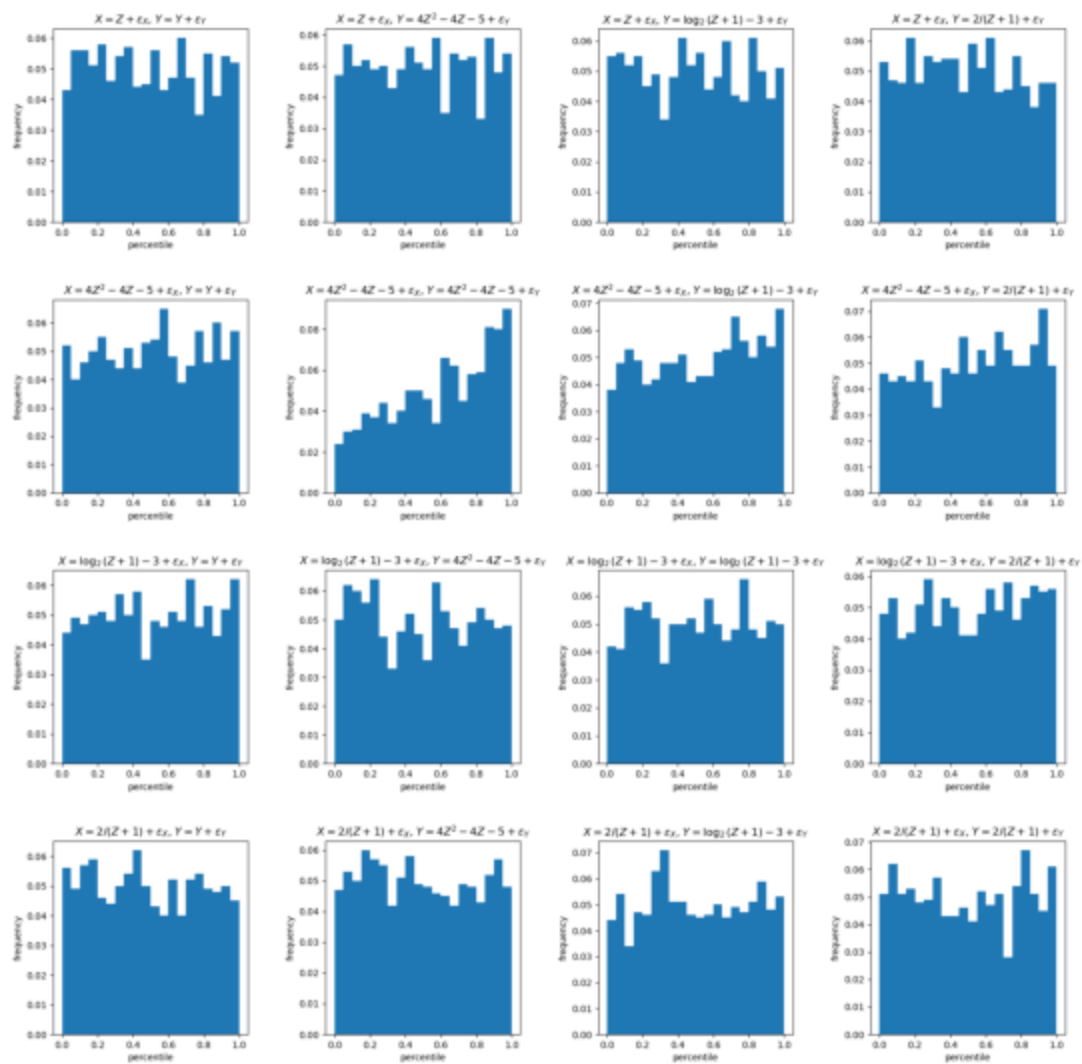
- $SD(X) = 0.5, SD(Y) = 0.5$

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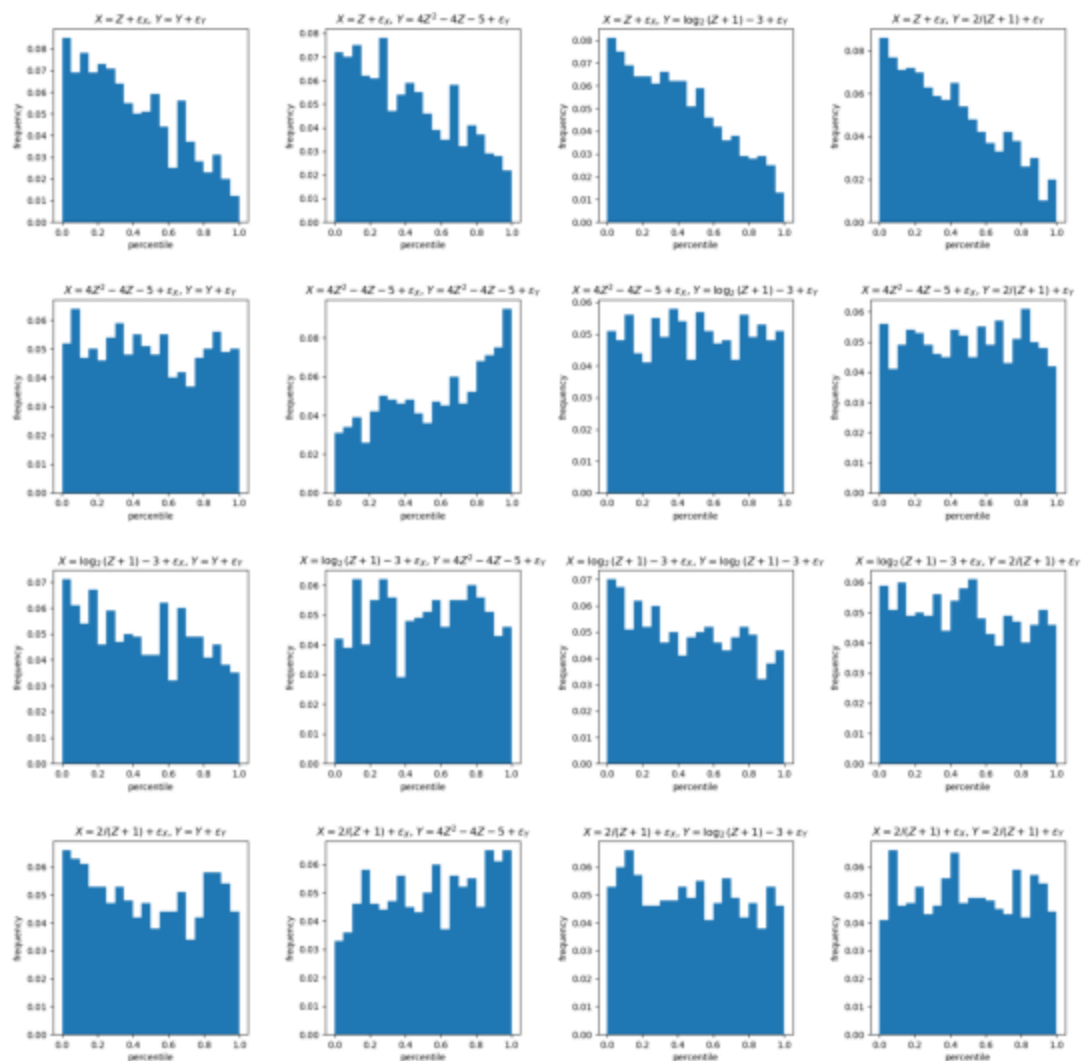
- $SD(X) = 0.5, SD(Y) = 0.01$

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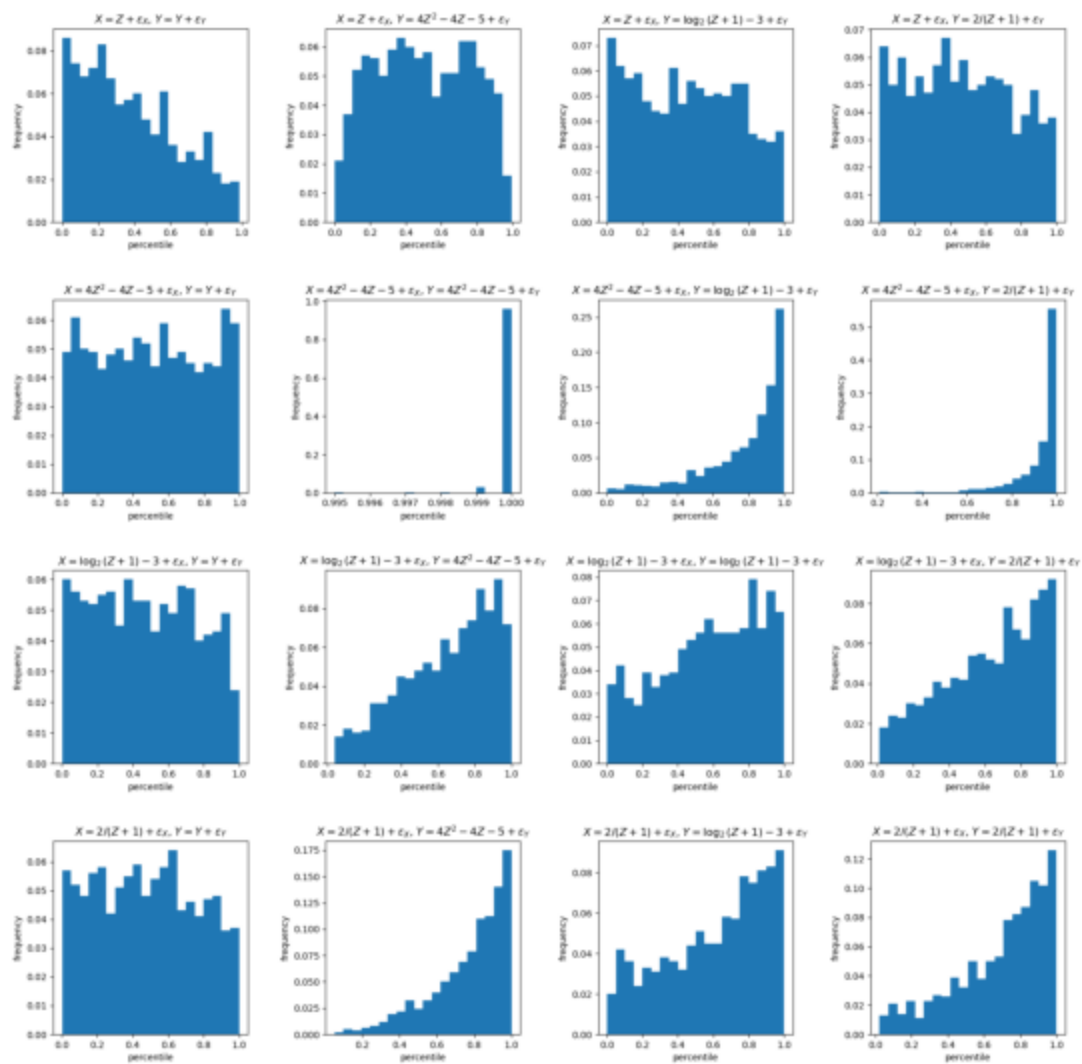
- $SD(X) = 0.01, SD(Y) = 0.5$

<matplotlib.image.AxesImage at 0x7fe3f3a7f160>



- $SD(X) = 0.01, SD(Y) = 0.01$

<matplotlib.image.AxesImage at 0x7fe3e3954490>



resid_Z

- $SD(X) = 0.5, SD(Y) = 0.5$

<matplotlib.image.AxesImage at 0x7fe3e3215f70>



- $SD(X) = 0.5, SD(Y) = 0.01$

<matplotlib.image.AxesImage at 0x7fe3d3c791c0>



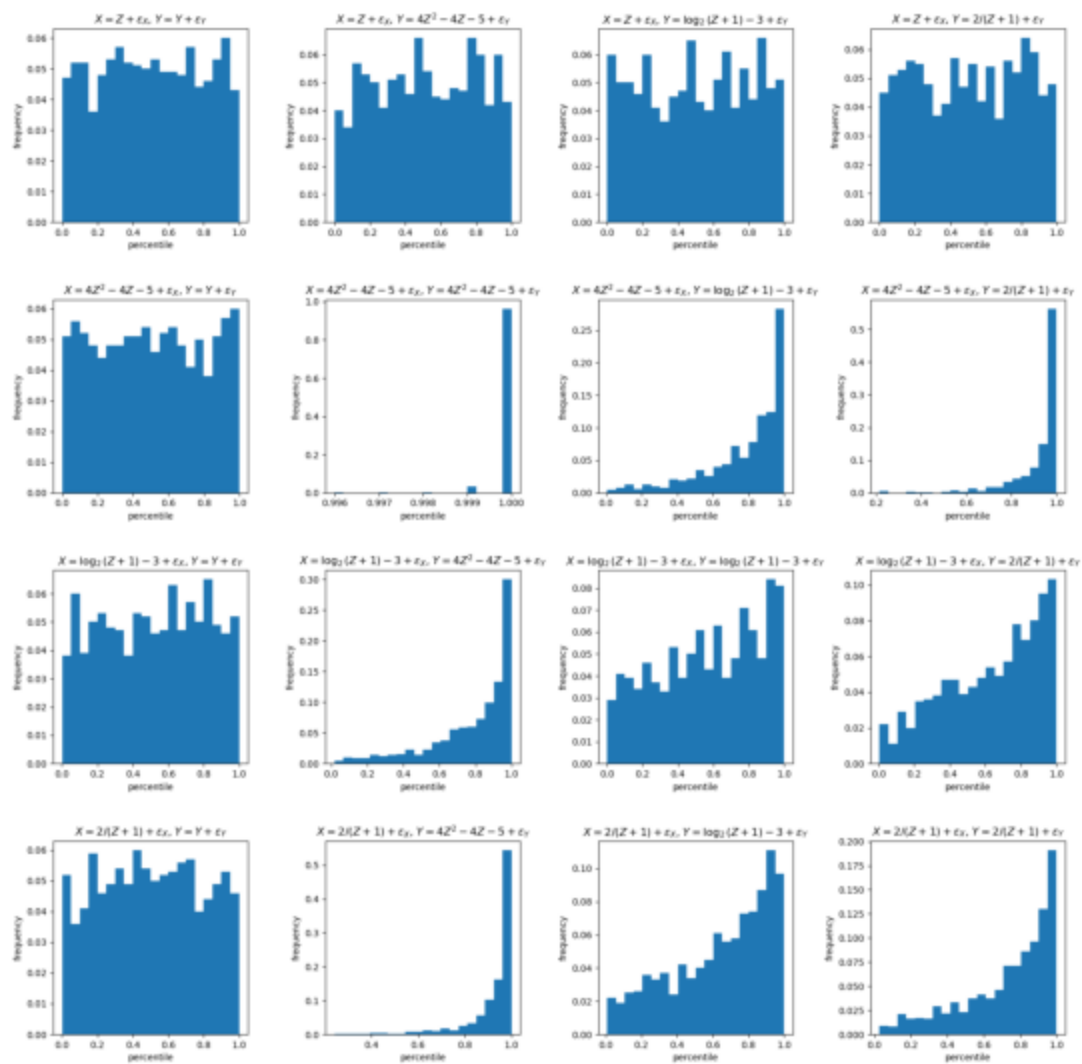
- $SD(X) = 0.01, SD(Y) = 0.5$

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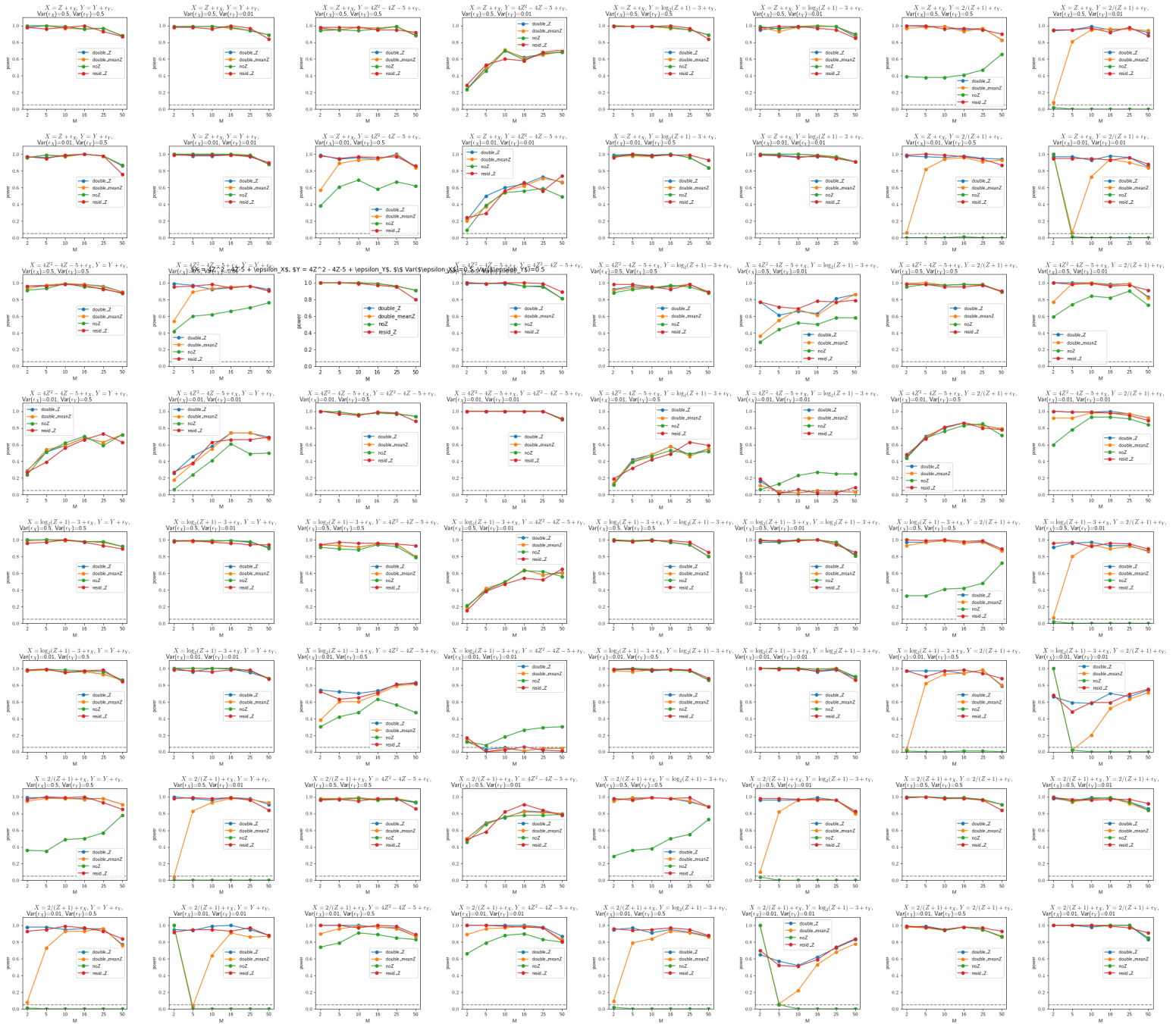
- $SD(X) = 0.01, SD(Y) = 0.01$

<matplotlib.image.AxesImage at 0x7fe3d506aa00>



Power

- $\text{cor} = 0.4$



• cor = 0.2

