**Gram matrix reconstruction error
USPST dataset for the angular kernel**

Legend:
- G
- $G_{circ}K_2K_1$
- $G_{Toeplitz}D_2HD_1$
- $G_{skew-circ}D_2HD_1$
- $HD_{g_1, g_2, ..., g_n}HD_2HD_1$
- $HD_3HD_2HD_1$

Y-axis: Gram matrix reconstruction error (0.00, 0.01, 0.02, 0.03, 0.04, 0.05)

X-axis: Number of random features (2^8, 2^9, 2^{10}, 2^{11}, 2^{12}, 2^{13})