

degeneracyCount

December 10, 2025

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[31]: import numpy as np
import matplotlib.pyplot as plt
import tqdm
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[32]: import sys, pathlib
sys.path.insert(0, str(pathlib.Path.cwd().parent))
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[33]: from decoding.beliefPropagation import performBeliefPropagationFast
from decoding.OSD import performOSD
```

```
[34]: codes = [
    "[[72, 12, 6]]",
    "[[90, 8, 10]]",
    "[[108, 8, 10]]",
    "[[144, 12, 12]]",
    "[[288, 12, 18]]",
]

trials = 1000
physicalErrorRates = np.logspace(-3.2, -1.3, 8)
results_BP = {}
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[35]: np.random.seed(0)
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[36]: for code in tqdm.tqdm(codes):
    oc = np.load(f'../codes/{code}.npz')
    name = code
    results_BP[name] = {}
    code = oc['Hx']
    Lx = oc['Lx']
    distance = oc['distance']
    n = len(code[0])
    logicalErrorRates = []
    BPs_fault_rates = []
    BPs_miscorrected_rates = []
    incorrectable_rates = []
    degeneracies = []
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for errorRate in physicalErrorRates:

    initialBeliefs = [np.log((1 - errorRate) / errorRate)] * n
    logical_error = 0
    BPs_fault = 0
    BPs_miscorrected = 0
    incorruptable = 0
    degenerateErrors = 0

    for _ in range(trials):

        ##### CODE CAPACITY ERROR MODEL #####
        # non-trivial pythonic way to generate random bitstring with given
        ↪ error rate
        error = (np.random.random(n) < errorRate).astype(int)

        syndrome = (error @ code.T) % 2

        ##### SIMPLE PHENOMENOLOGICAL ERROR MODEL #####
        # measurementError = (np.random.random(len(syndrome)) < errorRate).
        ↪ astype(int)
        # syndrome = (syndrome + measurementError) % 2

        detection, isSyndromeFound, llrs =
        ↪ performBeliefPropagationFast(code, syndrome, initialBeliefs, verbose=False)

        # if not isSyndromeFound:
        #     logical_error += 1
        #     BPs_fault += 1

        # detection = performOSD(code, syndrome, llrs, detection)

        # This is the XOR, between the actual error and the detected error.
        ↪ We are simulating the correction of the error
        residual = (detection + error) % 2

        syndromeLogic = (Lx @ residual) % 2

        if isSyndromeFound and not np.any(syndromeLogic) and (np.
        ↪ array_equal(detection, error) == False):
            degenerateErrors += 1

        if np.any(syndromeLogic):
            logical_error += 1

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        error_weight = np.sum(error)
        if error_weight < (distance // 2):
            BPs_miscorrected += 1
        else:
            incorrectable += 1

    ler = logical_error / trials
    logicalErrorRates.append(ler)
    BPs_fault_rates.append(BPs_fault)
    BPs_miscorrected_rates.append(BPs_miscorrected)
    incorrectable_rates.append(incorrectable)
    degeneracies.append(degenerateErrors)

results_BP[name]['ler'] = logicalErrorRates
results_BP[name]['BPs_fault'] = BPs_fault_rates
results_BP[name]['BPs_miscorrected'] = BPs_miscorrected_rates
results_BP[name]['incorrectable'] = incorrectable_rates
results_BP[name]['degeneracies'] = degeneracies

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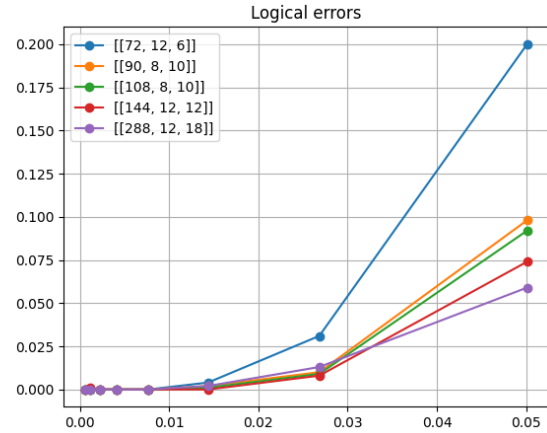
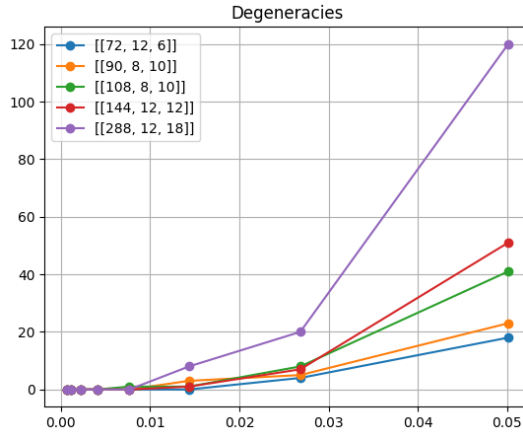
```

[37]: fig, axes = plt.subplots(1, 2, figsize=(14, 5))

for name in results_BP:
    axes[0].plot(physicalErrorRates, results_BP[name]['degeneracies'],
    ↪label=name, marker='o')
    axes[0].grid(True)
    axes[0].legend()
    axes[0].set_title('Degeneracies')

for name in results_BP:
    axes[1].plot(physicalErrorRates, results_BP[name]['ler'], label=name,
    ↪marker='o')
    axes[1].grid(True)
    axes[1].legend()
    axes[1].set_title('Logical errors')

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[38]: results_OSD = {}

for code in tqdm.tqdm(codes):
    oc = np.load(f'../codes/{code}.npz')
    name = code
    results_OSD[name] = {}
    code = oc['Hx']
    Lx = oc['Lx']
    distance = oc['distance']
    n = len(code[0])
    logicalErrorRates = []
    BPs_fault_rates = []
    BPs_miscorrected_rates = []
    incorrectable_rates = []
    degeneracies = []

    for errorRate in physicalErrorRates:

        initialBeliefs = [np.log((1 - errorRate) / errorRate)] * n
        logical_error = 0
        BPs_fault = 0
        BPs_miscorrected = 0
        incorrectable = 0
        degenerateErrors = 0

        for _ in range(trials):

            ##### CODE CAPACITY ERROR MODEL #####
            # non-trivial pythonic way to generate random bitstring with given
            ↪ error rate
```

```

error = (np.random.random(n) < errorRate).astype(int)

syndrome = (error @ code.T) % 2

#### SIMPLE PHENOMENOLOGICAL ERROR MODEL ####
# measurementError = (np.random.random(len(syndrome)) < errorRate).
→astype(int)
# syndrome = (syndrome + measurementError) % 2

detection, isSyndromeFound, llrs =
→performBeliefPropagationFast(code, syndrome, initialBeliefs, verbose=False)

if not isSyndromeFound:
    # logical_error += 1
    # BPs_fault += 1

    detection = performOSD(code, syndrome, llrs, detection)

    # This is the XOR, between the actual error and the detected error.
→We are simulating the correction of the error
    residual = (detection + error) % 2

    syndromeLogic = (Lx @ residual) % 2

    osd_syndrome_check = (detection @ code.T) % 2
    is_valid_osd = np.array_equal(osd_syndrome_check, syndrome)

    if is_valid_osd and not np.any(syndromeLogic) and (np.
→array_equal(detection, error) == False):
        degenerateErrors += 1

    if np.any(syndromeLogic):
        logical_error += 1

        error_weight = np.sum(error)
        if error_weight < (distance // 2):
            BPs_miscorrected += 1
        else:
            incorrectable += 1

ler = logical_error / trials
logicalErrorRates.append(ler)
BPs_fault_rates.append(BPs_fault)
BPs_miscorrected_rates.append(BPs_miscorrected)

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incorrectable_rates.append(incorrectable)
degeneracies.append(degenerateErrors)

results_OSD[name]['ler'] = logicalErrorRates
results_OSD[name]['BPs_fault'] = BPs_fault_rates
results_OSD[name]['BPs_miscorrected'] = BPs_miscorrected_rates
results_OSD[name]['incorrectable'] = incorrectable_rates
results_OSD[name]['degeneracies'] = degeneracies

```

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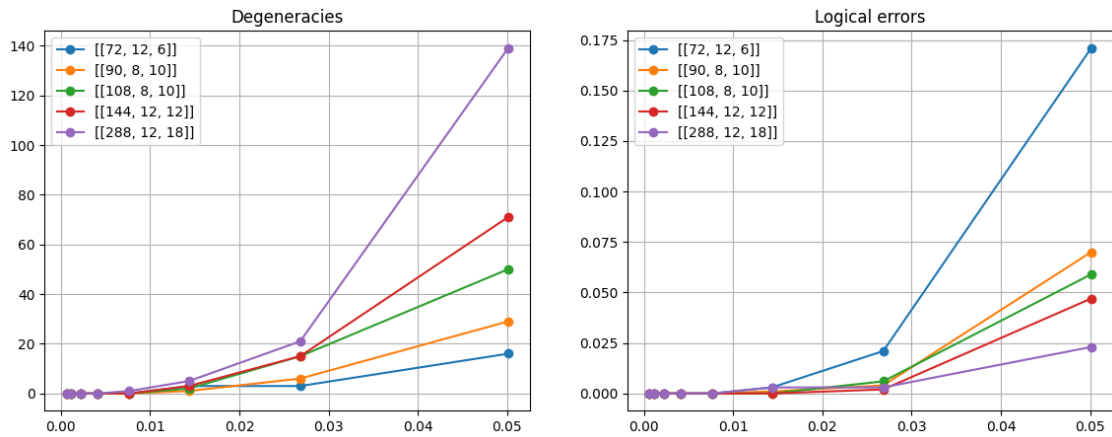
```

[39]: fig, axes = plt.subplots(1, 2, figsize=(14, 5))

for name in results_OSD:
    axes[0].plot(physicalErrorRates, results_OSD[name]['degeneracies'],
        label=name, marker='o')
    axes[0].grid(True)
    axes[0].legend()
    axes[0].set_title('Degeneracies')

for name in results_OSD:
    axes[1].plot(physicalErrorRates, results_OSD[name]['ler'], label=name,
        marker='o')
    axes[1].grid(True)
    axes[1].legend()
    axes[1].set_title('Logical errors')

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[40]: fig, axes = plt.subplots(2, 2, figsize=(14, 5))

# Left plots: BP results
for name in results_BP:

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    axes[0][0].plot(physicalErrorRates, results_BP[name]['degeneracies'],
        ↪label=name, marker='o')
axes[0][0].grid(True)
axes[0][0].legend()
# axes[0][0].set_xlabel('Physical error rate')
axes[0][0].set_xscale('log')
axes[0][0].set_ylabel('Degeneracies')
axes[0][0].set_title('BP degeneracy')

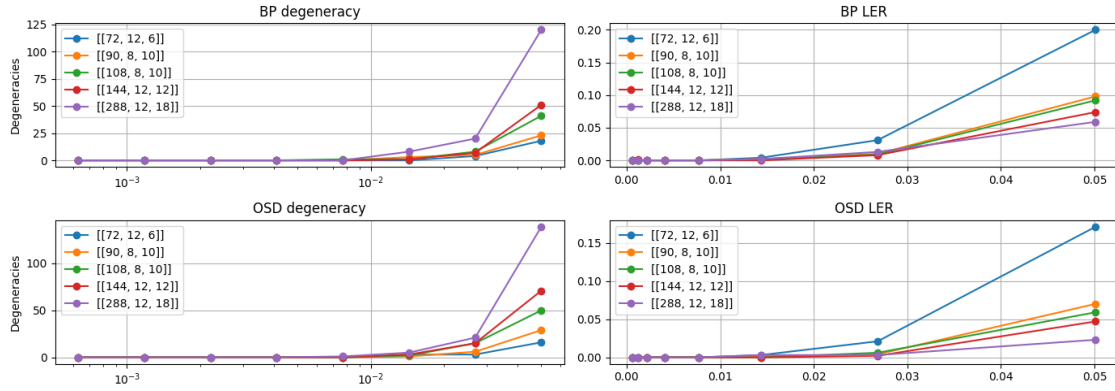
# Right plot: OSD results
for name in results_OSD:
    axes[1][0].plot(physicalErrorRates, results_OSD[name]['degeneracies'],
        ↪label=name, marker='o')
axes[1][0].grid(True)
axes[1][0].legend()
# axes[1][0].set_xlabel('Physical error rate')
axes[1][0].set_xscale('log')
axes[1][0].set_ylabel('Degeneracies')
axes[1][0].set_title('OSD degeneracy')

for name in results_BP:
    axes[0][1].plot(physicalErrorRates, results_BP[name]['ler'], label=name,
        ↪marker="o")
axes[0][1].grid(True)
axes[0][1].set_title("BP LER")
axes[0][1].legend()

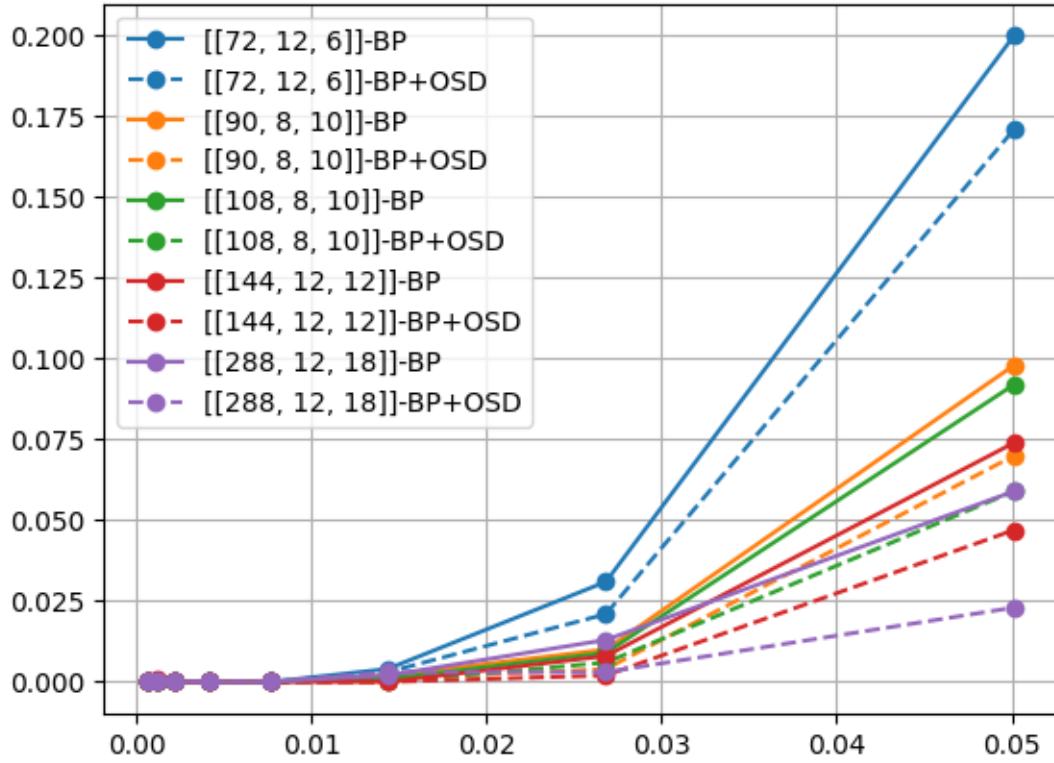
for name in results_OSD:
    axes[1][1].plot(physicalErrorRates, results_OSD[name]['ler'], label=name,
        ↪marker="o")
axes[1][1].grid(True)
axes[1][1].set_title("OSD LER")
axes[1][1].legend()

plt.tight_layout()
plt.show()

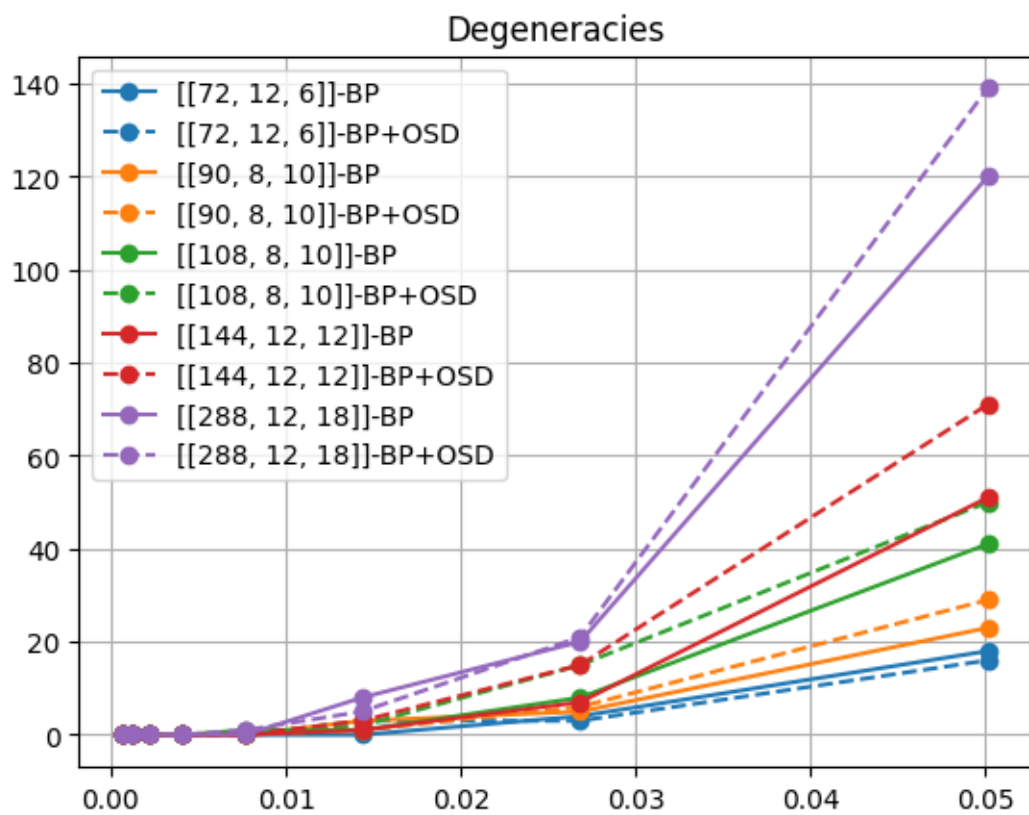
```



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[ ]: colors = plt.rcParams['axes.prop_cycle'].by_key()['color']
for i, name in enumerate(results_BP):
    c = colors[i % len(colors)]
    plt.plot(physicalErrorRates, results_BP[name]['ler'], label=f"{name}-BP",
    ↪marker='o', color=c)
    plt.plot(physicalErrorRates, results_OSD[name]['ler'],
    ↪label=f"{name}-BP+OSD", marker='o', linestyle='dashed', color=c)
plt.legend()
plt.grid(True)
plt.title("Logical Error Rate")
plt.show()
```

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[43]: colors = plt.rcParams['axes.prop_cycle'].by_key()['color']
for i, name in enumerate(results_BP):
    c = colors[i % len(colors)]
    plt.plot(physicalErrorRates, results_BP[name]['degeneracies'],
             label=f"{name}-BP", marker='o', color=c)
    plt.plot(physicalErrorRates, results_OSD[name]['degeneracies'],
             label=f"{name}-BP+OSD", marker='o', linestyle='dashed', color=c)
plt.legend()
plt.grid(True)
plt.title("Degeneracies")
plt.show()
```



[]: