Examples

October 30, 2025

1 Simulations for $EJAB_{10}$

Desired graphs (from talk w Dr. Nathoo): 1. Sensitivity, precision wrt n

2. ROC curve (Sensitivity vs 1-specificity as effect size varies)

3. Collect all type 1 errors from fixed sample size and fixed q; plot density: Should be uniform - This will be discussed later (i.e. during Friday's meeting)

To Do this week (due Friday): (1) and (2)

Note: Initially should be a simple t-test, we will then move to more complex models.

1.1 Notes and definitions

1.1.1 Sensitivity

$$sensitivity = \frac{True\ Positives}{True\ Positives + False\ Negatives} \in [0,1]$$

- The proportion actual positives correctly identified by the test
- E.g. if the a medical test for a disease detects 90/100 people who actually have the disease then the test has a sensitivity of 90%
- Interpretation: Higher sensitivity means fewer false negatives (type 2); the test rarely misses the positive cases

1.1.2 Precision (positive predictive value)

$$\label{eq:Precision} \begin{aligned} & \text{Precision} = \frac{\text{True Positives}}{\text{True Positives} + \text{False Positives}} \in [0, 1] \end{aligned}$$

- The proportion of positive predictions that are actually correct
- E.g. if a medical test for a disease identifies 100 people as having the disease, but only 80 of those people actually have the disease, then the test has a precision of 80%
- Interpretation: Higher precision means fewer false positives (type 1); the test rarely mislabels negative cases as positive

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1.1.3 ROC curve

(Note: Defined differently from discussed; we will plot both)

• TPR =
$$\frac{\text{TP}}{\text{TP+FN}}$$

• $FPR = 1 - Specificity = 1 - \frac{TN}{TN+FP} = \frac{FP}{FP+TN}$

ROC curve w effect size - TPR vs FPR as effect size varies - We will use Cohen's d: $d = \frac{\bar{X}_1 - \bar{X}_2}{S_{\text{pooled}}}$

1.2 Summary for plots:

- 1. Sensitivity vs sample size n
- 2. Precision vs sample size n
- 3. ROC curve (TPR vs FPR) as effect size d varies

1.3 Code

```
[1]: import numpy as np
     import scipy.stats as st
     rng = np.random.default_rng(277)
     # ---- settings ----
     n_grid = np.linspace(10, 100, 100, dtype=int) # sample sizes for the x-axis
     R = 5000
                                                        # replicates per (n)
     d0 = 0.5
                                                        # effect size for H1
                                                        # t-test threshold
     alpha = 0.05
     # ---- helpers ----
     def simulate_pvals(n, R, d_scalar, rng=None):
        if rng is None:
            rng = np.random.default_rng()
         # R experiments of size n from N(d, 1)
        X = rng.normal(loc=d_scalar, scale=1.0, size=(R, n))
         # two-sided one-sample t-test per experiment (axis=1)
        res = st.ttest_1samp(X, popmean=0.0, axis=1, alternative='two-sided')
        return res.pvalue # shape (R,)
     def ejab01(n, pvals):
         # safe chi-square quantile to avoid inf at 0/1
        eps = 1e-12
        q = st.chi2.ppf(np.clip(1 - pvals, eps, 1 - eps), df=1)
        return np.sqrt(n) * np.exp(-0.5 * q * (n - 1) / n)
     # ---- outputs to plot ----
     sens_ejab = np.empty(len(n_grid))
     prec_ejab = np.empty(len(n_grid))
     sens_t = np.empty(len(n_grid))
     prec_t = np.empty(len(n_grid))
     \# ---- main loop over n ----
```

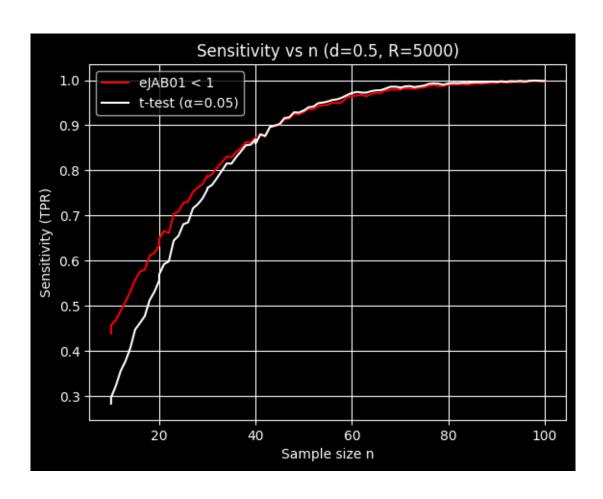
```
for i, n in enumerate(n_grid):
    # H1 (positives in truth) and H0 (negatives in truth)
    p_H1 = simulate_pvals(int(n), R, d0, rng=rng)
    p_H0 = simulate_pvals(int(n), R, 0.0, rng=rng)
    # eJAB01 decisions
    dec_ejab_H1 = ejab01(int(n), p_H1) < 1
    dec_ejab_H0 = ejab01(int(n), p_H0) < 1
    TP = dec ejab H1.sum()
    FN = (~dec ejab H1).sum()
    FP = dec_ejab_H0.sum()
    TN = (~dec_ejab_H0).sum()
    sens_ejab[i] = TP / (TP + FN) if (TP + FN) else np.nan
    prec_ejab[i] = TP / (TP + FP) if (TP + FP) else np.nan
    # classical t-test decisions at alpha
    dec_t_H1 = p_H1 \le alpha
    dec_t_H0 = p_H0 \le alpha
    TP_t = dec_t_H1.sum()
    FN_t = (\sim dec_t_{H1}).sum()
    FP t = dec t H0.sum()
    TN_t = (\sim dec_t_{H0}).sum()
    sens_t[i] = TP_t / (TP_t + FN_t) if (TP_t + FN_t) else np.nan
    prec_t[i] = TP_t / (TP_t + FP_t) if (TP_t + FP_t) else np.nan
```

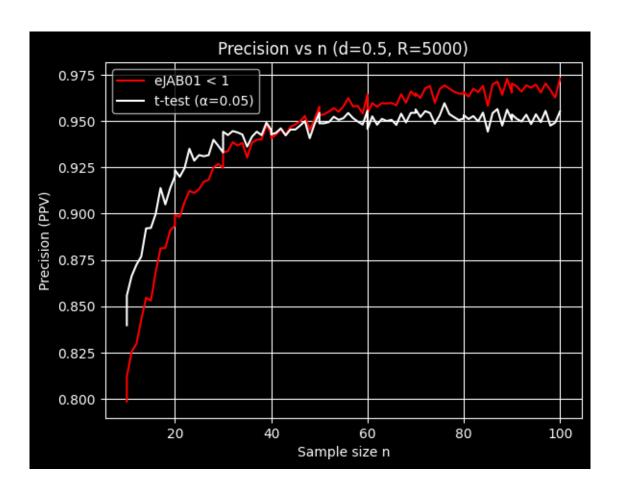
```
[2]: import matplotlib.pyplot as plt

plt.style.use('dark_background')

plt.figure()
plt.plot(n_grid, sens_ejab, color='red', label='eJABO1 < 1')
plt.plot(n_grid, sens_t, color='white', label=f't-test (={alpha})')
plt.xlabel('Sample size n'); plt.ylabel('Sensitivity (TPR)')
plt.title(f'Sensitivity vs n (d={d0}, R={R})'); plt.grid(True); plt.legend()

plt.figure()
plt.plot(n_grid, prec_ejab, color='red', label='eJABO1 < 1')
plt.plot(n_grid, prec_t, color='white', label=f't-test (={alpha})')
plt.xlabel('Sample size n'); plt.ylabel('Precision (PPV)')
plt.title(f'Precision vs n (d={d0}, R={R})'); plt.grid(True); plt.legend()
plt.show()</pre>
```

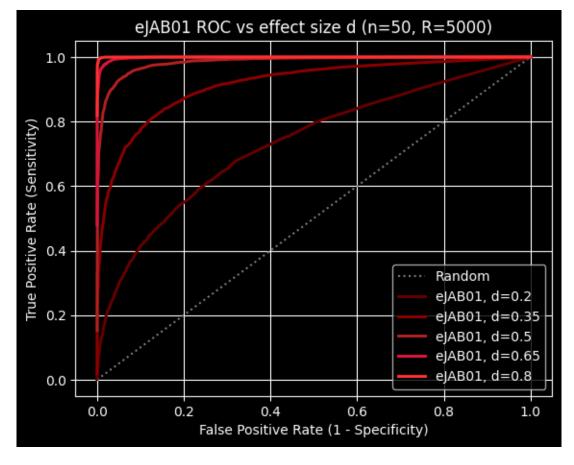




```
[]: # eJAB-only ROC across effect sizes
     def roc_from_ejab(p_H1, p_H0, n, thr_grid):
         e1 = ejab01(int(n), p_H1)
         e0 = ejab01(int(n), p_H0)
         # decision: reject HO if eJAB <
         tpr = np.array([np.mean(e1 < t) for t in thr_grid])</pre>
         fpr = np.array([np.mean(e0 < t) for t in thr_grid])</pre>
         return fpr, tpr
     n_fixed = 50
     d_{grid} = [0.2, 0.35, 0.5, 0.65, 0.8]
     # alpha_grid = np.r_[0.0, np.logspace(-6, -1, 200), np.linspace(0.1, 1.0, 200)]_{\sqcup}
     → # (unused here; keep if needed elsewhere)
     thr_grid = np.logspace(-3, 1, 400)
     colors = ['#660000', '#8B0000', '#B22222', '#DC143C', '#FF3333']
     plt.figure()
     plt.plot([0,1], [0,1], ':', color='gray', label='Random')
```

```
for d, c in zip(d_grid, colors):
    p_H1 = simulate_pvals(int(n_fixed), R, d, rng=rng)
    p_H0 = simulate_pvals(int(n_fixed), R, 0.0, rng=rng)
    fpr_e, tpr_e = roc_from_ejab(p_H1, p_H0, n_fixed, thr_grid)
    plt.plot(fpr_e, tpr_e, lw=2.2, color=c, label=f'eJAB01, d={d}')

plt.xlabel('False Positive Rate (1 - Specificity)')
plt.ylabel('True Positive Rate (Sensitivity)')
plt.title(f'eJAB01 ROC vs effect size d (n={n_fixed}, R={R})')
plt.grid(True); plt.legend(); plt.show()
```



Here's some clarification, I will add these to the graphs as I see now that the labeling is unclear: - EJAB01 Sensitivity/precision vs n (Red lines) - We first define the ejab01 decisions: - $dec_ejab_H1 = ejab01(int(n), p_H1) < 1 \# true effects - <math>dec_ejab_H0 = ejab01(int(n), p_H0) < 1 \# false positives - We sum them to get totals to use in the sensitivity and precision plots: - <math>TP = dec_ejab_H1.sum() \# Num true positives as predicted by ejab - FN = (~dec_ejab_H1).sum() \# False negatives ... ejab - FP = <math>dec_ejab_H0.sum() \# False positives ... - TN = (~dec_ejab_H0).sum() \# True negatives ... - We use these to$

compute sensitivity, precision: - sens_t[i] = TP_t / (TP_t + FN_t) if (TP_t + FN_t) else np.nan - prec_t[i] = TP_t / (TP_t + FP_t) if (TP_t + FP_t) else np.nan - We compare with the actual results of the t-test alpha=0.05 (White lines) - dec_t_H1 = p_H1 <= alpha - dec_t_H0 = p_H0 <= alpha

e.g. plt.plot(n_grid, sens_ejab, color='red', label='eJAB01 < 1') plt.plot(n_grid, sens_t, color='white', label=f't-test (={alpha})')

- For the ROC curves:
 - The roc_from_ejabfunciton takes as input:
 - * Simulated p values with no/an effect (p_H0, p_H1)
 - * n (sample size)
 - * thr_grid: A log spaced array of possible decision thresholds t; "reject H0 if eJAB01 < t"
 - It uses these to calculate the TPR/FPR (True/false pos rates) (mean gives proportion of values)
 - * tpr = np.array([np.mean(e1 < t) for t in thr_grid])
 - * fpr = np.array([np.mean(e0 < t) for t in thr_grid])
 - As requested, we plot far various effect sizes using cohen's d:
 - $* d_grid = [0.2, 0.35, 0.5, 0.65, 0.8]$
 - The darker lines in the plot correspond to smaller effect sizes. We can see that, as expected, for smaller effect sizes they approach x=y; truly random data.
 - For larger effect size (lighter lines) the curves hug the y-axis, reflecting higher true positive rates (greater sensitivity) for the same false positive rate.