Practical Exercise — Lesson 2 (Week 5)

Quick Fairness Probe: Detect and Explain Bias in Two Settings

Learning outcomes

By the end, you will be able to:

- 1. Align a **fairness metric** to a concrete **harm to avoid** (access, missed protection, wrongful flags).
- 2. Compute **per-group metrics**, **gaps/ratios**, **worst-group** values, and (optionally) **95% CIs**.
- 3. Run a lightweight LM bias probe (CrowS-Pairs or WEAT) and interpret the score.
- 4. Recommend one actionable mitigation and explain how you will re-measure.

Setup

- 1. Open a fresh Colab (CPU is fine).
- 2. Install/import the basics:
- 3. !pip -q install datasets transformers torch scikit-learn --upgrade
- 4. # optional for WEAT:
- 5. # !pip -q install wefe
- 6
- 7. import numpy as np, pandas as pd, torch
- 8. from datasets import load dataset
- 9. from transformers import pipeline, AutoTokenizer, AutoModelForCausalLM
- 10. from sklearn.metrics import confusion matrix, precision recall curve
- 11. Add a SEED = 42 and set numpy/torch RNG for repeatability.

Setting A — Group Fairness in a Classifier

A1. Choose data & model

- Preferred (toxicity): civil_comments from HF (has identity columns like male, female, etc.).
- ds =
 load_dataset("civil_comments")["train"].shuffle(seed=42).select(range(5 000))
- text_col, label_col = "text", "toxicity" # threshold label later (>=0.5)

- Fallback (if identity columns unavailable): build keyword slices (e.g., strings containing identity terms). Note clearly this is an approximation.
- Model: any small text-classification pipeline (for speed):
- clf = pipeline("text-classification",
- model="distilbert-base-uncased-finetuned-sst-2-english",
- return all scores=True, truncation=True)

A2. Prepare labels & run inference

- 1. If your dataset's label is continuous (e.g., toxicity 0–1), binarize at 0.5 into y.
- 2. Score a **2–5k** sample. Save: gold y, predicted label ŷ, and **score** s (probability of positive).

```
3. def pos_score(out): # expects list of dicts
  [{'label':'NEGATIVE', 'score':...}, ...]
4.    return next(d["score"] for d in out if
    d["label"].upper().startswith("POS"))
5.
6. rows=[]
7. for r in ds:
8.    s = pos_score(clf(r[text_col]))
9.    rows.append(dict(text=r[text_col], y=int(r[label_col] >= 0.5),
    S=s))
10. df = pd.DataFrame(rows)
11. df["yhat"] = (df["S"] >= 0.5).astype(int)
```

A3. Declare harm \rightarrow metric

Pick one primary harm and its matching metric (write this in a cell):

- Over-moderation \rightarrow FPR gap (wrongful flags).
- **Missed protection** → **TPR/Recall gap** (Equal Opportunity).
- Uneven trustworthiness \rightarrow PPV/Calibration by group.

A4. Compute per-group metrics, gaps, worst-group

Select 2–4 groups (e.g., male, female, LGBTQ if available). For each group aaa: compute TP/FP/TN/FN, then **Precision**, **Recall**, **F1**, **FPR**, **FNR**. Report:

- a per-group table
- Gap = max-min for your primary metric
- **Ratio** = min/max (optional)
- Worst-group value

```
from sklearn.metrics import precision_recall_fscore_support

def group_mask(df, col):  # identity can be float probability or bool
    return (df[col] > 0.5) if df[col].dtype != bool else df[col]

def prfl fpr fnr(y, yhat):
```

```
tn, fp, fn, tp = confusion matrix(y, yhat, labels=[0,1]).ravel()
    p = tp/(tp+fp+1e-9); r = tp/(tp+fn+1e-9); f1 = 2*p*r/(p+r+1e-9)
    fpr = fp/(fp+tn+1e-9); fnr = fn/(fn+tp+1e-9)
    return p, r, f1, fpr, fnr
groups = [c for c in ds.column names if c in
("male", "female", "lgbtq", "black", "white")]
rows=[]
for q in groups:
   m = group mask(df, g)
   p,r,f1,fpr,fnr = prf1 fpr fnr(df.loc[m,"y"], df.loc[m,"yhat"])
    rows.append(dict(group=g, n=int(m.sum()), precision=p, recall=r, f1=f1,
fpr=fpr, fnr=fnr))
rep = pd.DataFrame(rows)
# Example: primary metric = FPR
gap = rep.fpr.max() - rep.fpr.min()
ratio = rep.fpr.min() / (rep.fpr.max()+1e-9)
worst group = rep.iloc[rep.fpr.idxmax()].group
rep, gap, ratio, worst group
```

Optional (recommended): 95% CIs via bootstrap on the gap (1,000 resamples).

A5. Threshold analysis (if you have scores)

Plot a **precision–recall curve** and choose a deployment threshold aligned to your harm (e.g., choose a point with **Precision** \geq **0.9** if wrongful flags are costly). Recompute your **primary gap** at the chosen threshold.

```
prec, rec, thr = precision_recall_curve(df["y"], df["S"])
# Pick threshold programmatically (example: F1-max) or justify a business
constraint
```

A6. Mini-interpretation (3–4 sentences)

- Name the **primary gap** and **worst-group**.
- Explain the user/business harm.
- Propose **one mitigation** (e.g., threshold by slice; counterfactual data augmentation; domain adaptation) and say **how you will re-measure**.

Deliverables for Setting A:

• Table (per-group metrics + gap/ratio + worst-group), PR curve with chosen threshold (if scores), 3–4 sentence interpretation.

Setting B — Stereotype Bias in a Language Model

Choose **one** option.

Option B1 — CrowS-Pairs (LM preference test)

Goal: fraction of times the LM prefers a stereotypical over an anti-stereotypical sentence \rightarrow S-score.

Steps

```
    Load a small slice (≈300 pairs) and a compact causal LM (e.g., gpt2):
    ds = load_dataset("crows_pairs",
        "english") ["test"] .shuffle(seed=42) .select(range(300))
    tok = AutoTokenizer.from_pretrained("gpt2")
    lm = AutoModelForCausalLM.from_pretrained("gpt2"); lm.eval();
    Compute sentence negative log-likelihood (NLL):
    def sent_nll(text):
        ids = tok(text, return_tensors="pt")
        with torch.no_grad():
            out = lm(**ids, labels=ids["input_ids"])
        return float(out.loss) * ids["input_ids"].size(1) # approx NLL
    For each pair, compute NLL for stereotypical vs anti-stereotypical; count where
```

- stereotypical is lower (preferred).
- 12. Report **overall S-score** and **by category** (e.g., gender, race, religion).
- 13. Write a 2–3 sentence **interpretation** (where bias concentrates; potential user-visible harm in completions).

Option B2 — **WEAT** (embedding association test)

Goal: effect size ddd measuring association strength (e.g., $science \leftrightarrow male$ vs $arts \leftrightarrow female$).

Steps

- 1. Install and import wefe (optional but easier), or implement manually.
- 2. Choose target sets (e.g., male/female names) and attribute sets (science/arts terms).
- 3. Compute **WEAT** effect size ddd and (optionally) a permutation p-value.
- 4. Report ddd overall and a brief interpretation (e.g., d=0.78d=0.78d=0.78 = large association).

Deliverables for Setting B:

• One numeric metric (S-score or WEAT effect size) overall and by at least one category/slice, plus a 2–3 sentence interpretation.

Fairness Results Card (≤1 page, PDF)

Use the following template (copy/paste into a doc and export to PDF):

Task & Data: (dataset, sample size, groups)

Primary harm \rightarrow **Metric:** (e.g., Over-moderation \rightarrow FPR gap)

Headline numbers: Worst-group = __; Gap = __ (± CI if computed); Ratio = __

Decision/threshold: (if applicable; justify)

LM probe: (metric and brief finding; e.g., CrowS S=63% overall, 75% gender)

Mitigation next step: (one concrete action)

Re-measurement plan: (what you will recompute to confirm improvement)

Limitations: (sample size, proxy groups, etc.)