

# 1 FIAP Tech Challenge — Detecção de Tuberculose em Radiografias de Tórax

## Grupo 71

**Objetivo:** classificar RX de tórax para **triagem** em **Normal (0)** vs **Tuberculosis (1)**, priorizando **sensibilidade máxima** (sem falsos negativos).

**Classe positiva:** 1 = Tuberculosis **Política clínica:** **Recall = 1.00** ( $\Rightarrow$  **FNR=0**) no teste.

**Threshold:** escolhido na **validação** por varredura (FN=0 e FP mínimo) e aplicado inalterado ao teste.

**Repositório GIT:** <https://github.com/toniisidorofiap/fiap-tech-challenge-fase01-grupo71>

**Dataset:** [Kaggle – TB Chest X-ray Database](#) (sem PHI; respeitar licença).

**Fluxo:** setup reprodutível  $\rightarrow$  sanity/contagens  $\rightarrow$  split 70/15/15 (estratificado)  $\rightarrow$  `tf.data` + aug conservadora  $\rightarrow$  Baseline (CNN)  $\rightarrow$  Transfer Learning (MobileNetV2 + FT)  $\rightarrow$  seleção de limiar na **validação**  $\rightarrow$  métricas no **teste**  $\rightarrow$  Grad-CAM++  $\rightarrow$  exportação.

**Relato no teste (com o limiar da validação):** Recall, **FNR**, **Specificity**, **NPV**, Precision, F1, AUC; 1 matriz de confusão; **ROC** e **PR**; **3–6** Grad-CAMs (incluindo FP).

Uso acadêmico; não-diagnóstico. Resultados devem ser interpretados por profissionais de saúde.

## 1.1 Configuração, Dependências e Reprodutibilidade

Instalando as dependências necessárias:

```
[ ]: # Se utilizar google colab, descomente a linha abaixo para instalar as
    ↳ dependências
# !pip install kagglehub tensorflow scikit-learn matplotlib opencv-python
    ↳ pillow tf-keras-vis

import os, random, shutil, sys, json, math, itertools, gc
from pathlib import Path
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
from matplotlib import cm as mpl_cm
import cv2
from PIL import Image
import kagglehub

import tensorflow as tf
from tensorflow import keras
from tensorflow.keras import layers, Model, Input, Sequential
from tensorflow.keras.callbacks import EarlyStopping, ReduceLROnPlateau,
    ↳ ModelCheckpoint
```

```

from tensorflow.keras.applications.mobilenet_v2 import MobileNetV2, \
    preprocess_input

from sklearn.model_selection import train_test_split
from sklearn.utils.class_weight import compute_class_weight
from sklearn.metrics import (accuracy_score, precision_score, recall_score, \
    f1_score, confusion_matrix, classification_report, ConfusionMatrixDisplay, \
    roc_auc_score, roc_curve, precision_recall_curve, average_precision_score)

from tf_keras_vis.gradcam import GradcamPlusPlus
from tf_keras_vis.utils import normalize
from tf_keras_vis.utils.model_modifiers import ReplaceToLinear

```

Garantindo reprodutibilidade usando um SEED para poder replicar mesmo resultado:

```

[2]: SEED = 13
    random.seed(SEED); np.random.seed(SEED); tf.random.set_seed(SEED)

```

## 1.2 Dados e Sanity Check

Removendo imagens suspeitas ou corrompidas:

```

[3]: path = kagglehub.dataset_download("tawsifurrahman/
    ↪tuberculosis-tb-chest-xray-dataset")

DATA_DIR = Path(path) / "TB_Chest_Radiography_Database"
NORMAL_DIR = DATA_DIR / "Normal"
TB_DIR = DATA_DIR / "Tuberculosis"

valid_ext = {'.png', '.jpg', '.jpeg', '.bmp', '.tif', '.tiff'}

def count_images(d: Path):
    if not d.exists(): return 0
    return sum(1 for f in d.iterdir() if f.is_file() and f.suffix.lower() in \
    ↪valid_ext)

print("Normal:", count_images(NORMAL_DIR), "| TB:", count_images(TB_DIR))

# Sanity check: detectar arquivos corrompidos/"uniformes"
invalid = []
for img_path in DATA_DIR.rglob('*'):
    if img_path.suffix.lower() in valid_ext and img_path.is_file():
        try:
            im = Image.open(img_path); im.verify()
            im = Image.open(img_path).convert("RGB")
            arr = np.asarray(im)
            if arr.std() == 0:
                invalid.append((str(img_path), "imagem uniforme"))

```

```

        except Exception as e:
            invalid.append((str(img_path), f"erro: {e}"))

print("Suspeitas/Corrompidas:", len(invalid))
if invalid:
    with open("sanity_invalid.json", "w") as f: json.dump(invalid, f, indent=2)
    print(f"Lista completa em sanity_invalid.json (n={len(invalid)})")

```

Normal: 3500 | TB: 700

Suspeitas/Corrompidas: 0

### 1.3 EDA e Split Estratificado 70/15/15

```

[4]: normal_paths = sorted([str(p) for p in (DATA_DIR / "Normal").glob("*") if p.
    ↪suffix.lower() in valid_ext])
tb_paths      = sorted([str(p) for p in (DATA_DIR / "Tuberculosis").glob("*") if
    ↪p.suffix.lower() in valid_ext])

X = normal_paths + tb_paths
y = [0]*len(normal_paths) + [1]*len(tb_paths)

print("Total imagens:", len(X))
print("Distribuição:", {"Normal": y.count(0), "TB": y.count(1)})

X_train, X_temp, y_train, y_temp = train_test_split(X, y, test_size=0.30,
    ↪stratify=y, random_state=SEED)
X_val, X_test, y_val, y_test = train_test_split(X_temp, y_temp, test_size=0.50,
    ↪stratify=y_temp, random_state=SEED)

def counts(lbls):
    a = np.array(lbls)
    return {0: int((a == 0).sum()), 1: int((a == 1).sum())}

df_split = pd.DataFrame({
    "partição": ["Treino", "Validação", "Teste"],
    "Normal": [counts(y_train)[0], counts(y_val)[0], counts(y_test)[0]],
    "TB": [counts(y_train)[1], counts(y_val)[1], counts(y_test)[1]],
    "total": [len(y_train), len(y_val), len(y_test)]
})
display(df_split)

# copie todas as imagens do conjunto X_test para a pasta ./dataset/test_images/
    ↪contendo subpastas 'Normal' e 'Tuberculosis' conforme os rótulos y_test
TEST_DIR = Path("./dataset/test_images")
NORMAL_TEST_DIR = TEST_DIR / "Normal"
TB_TEST_DIR = TEST_DIR / "Tuberculosis"
NORMAL_TEST_DIR.mkdir(parents=True, exist_ok=True)

```

```

TB_TEST_DIR.mkdir(parents=True, exist_ok=True)
for i, p in enumerate(X_test):
    img = keras.utils.load_img(p)
    label = "TB" if y_test[i] == 1 else "Normal"
    if label == "Normal":
        img.save(NORMAL_TEST_DIR / f"{Path(p).name}")
    else:
        img.save(TB_TEST_DIR / f"{Path(p).name}")

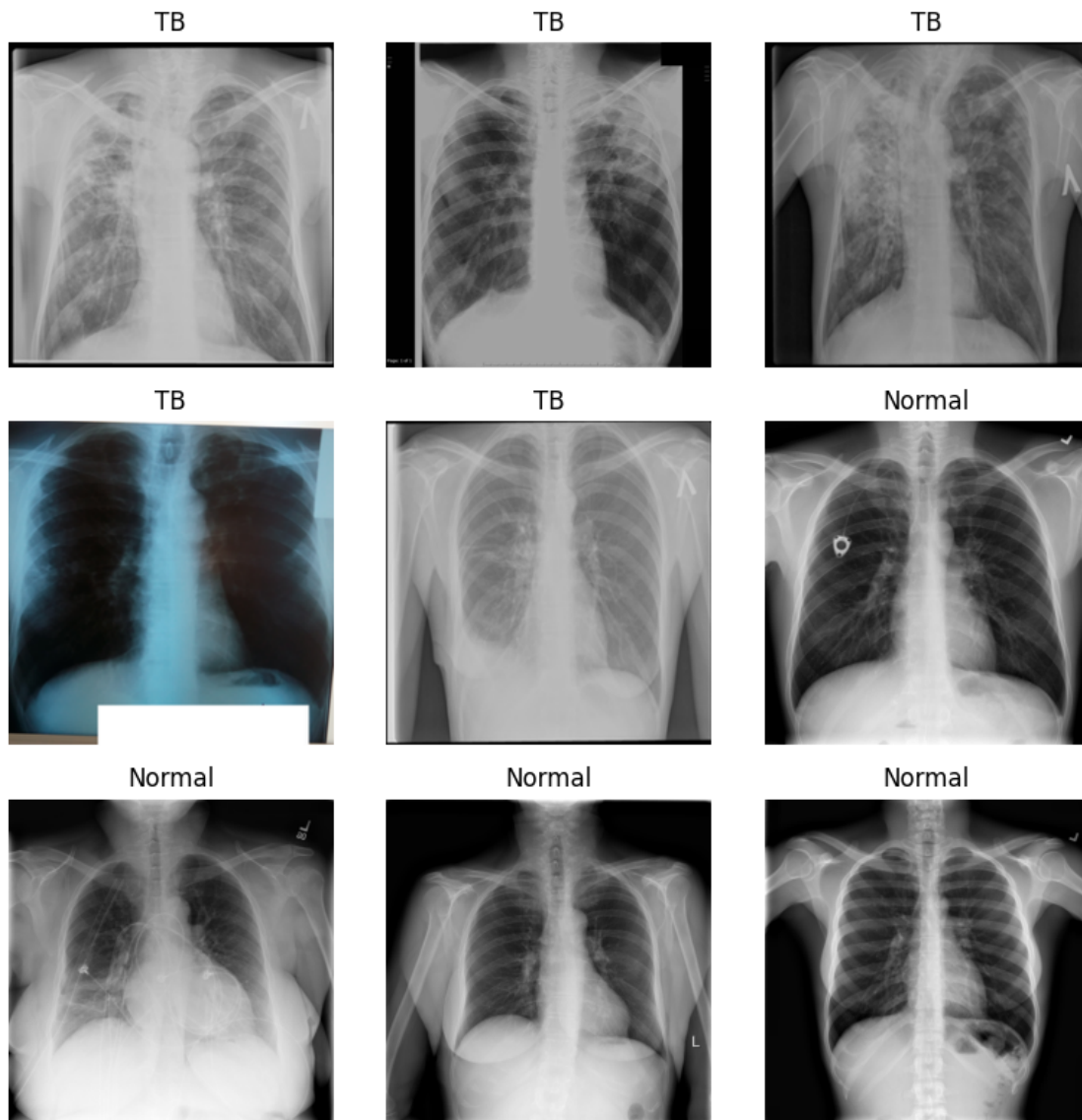
# Visualização de algumas amostras
plt.figure(figsize=(8,8))
# We want exactly 5 TB and 4 Normal images from the training set
tb_count = 0
normal_count = 0
plot_idx = 0
# iterate over X_train together with y_train (use the split labels)
for p, label in zip(X_train, y_train):
    if plot_idx >= 9:
        break
    # select TB images first (5), then Normal images (4)
    select = False
    if (tb_count < 5 and label == 1) or (tb_count >= 5 and normal_count < 4 and
↪label == 0):
        select = True
    if not select:
        continue
    # update counters and plot
    if label == 1:
        tb_count += 1
    else:
        normal_count += 1
    img = keras.utils.load_img(p)
    plt.subplot(3,3,plot_idx+1)
    plt.imshow(img); plt.axis('off')
    plt.title("TB" if label==1 else "Normal")
    plot_idx += 1
plt.tight_layout(); plt.show()

```

Total imagens: 4200

Distribuição: {'Normal': 3500, 'TB': 700}

	partição	Normal	TB	total
0	Treino	2450	490	2940
1	Validação	525	105	630
2	Teste	525	105	630



#### 1.4 Pipeline de Dados (tf.data) & Class Weights

```
[5]: IMG_SIZE = (224,224)
    BATCH = 32
    AUTO = tf.data.AUTOTUNE

    def load_image(path, label):
        img = tf.io.read_file(path)
        img = tf.image.decode_image(img, channels=3, expand_animations=False)
        img = tf.image.resize(img, IMG_SIZE)
        img = tf.cast(img, tf.float32) / 255.0
```

```

    return img, label

def aug(img, label):
    img = tf.image.random_flip_left_right(img)
    img = tf.image.random_brightness(img, 0.1)
    return img, label

def make_ds(paths, labels, training=False):
    ds = tf.data.Dataset.from_tensor_slices((paths, labels))
    ds = ds.map(load_image, num_parallel_calls=AUTO)
    if training:
        ds = ds.shuffle(2048, seed=SEED).map(aug, num_parallel_calls=AUTO)
    return ds.batch(BATCH).prefetch(AUTO)

train_ds = make_ds(np.array(X_train), np.array(y_train), training=True)
val_ds    = make_ds(np.array(X_val), np.array(y_val))
test_ds   = make_ds(np.array(X_test), np.array(y_test))

class_weights_arr = compute_class_weight('balanced', classes=np.array([0,1]),
    ↪y=np.array(y_train))
CLASS_WEIGHTS = {0: float(class_weights_arr[0]), 1: float(class_weights_arr[1])}
print("Class Weights:", CLASS_WEIGHTS)

```

```

2025-11-04 16:21:18.352944: I metal_plugin/src/device/metal_device.cc:1154]
Metal device set to: Apple M4
2025-11-04 16:21:18.353168: I metal_plugin/src/device/metal_device.cc:296]
systemMemory: 32.00 GB
2025-11-04 16:21:18.353192: I metal_plugin/src/device/metal_device.cc:313]
maxCacheSize: 10.67 GB
2025-11-04 16:21:18.353440: I
tensorflow/core/common_runtime/pluggable_device/pluggable_device_factory.cc:305]
Could not identify NUMA node of platform GPU ID 0, defaulting to 0. Your kernel
may not have been built with NUMA support.
2025-11-04 16:21:18.353480: I
tensorflow/core/common_runtime/pluggable_device/pluggable_device_factory.cc:271]
Created TensorFlow device (/job:localhost/replica:0/task:0/device:GPU:0 with 0
MB memory) -> physical PluggableDevice (device: 0, name: METAL, pci bus id:
<undefined>)

Class Weights: {0: 0.6, 1: 3.0}

```

## 1.5 Modelos

Modelo 1 - CNN Baseline

```

[11]: def build_baseline():
    inputs = layers.Input(shape=IMG_SIZE+(3,))
    x = layers.Conv2D(32, 3, activation='relu')(inputs)

```

```

x = layers.MaxPooling2D()(x)
x = layers.Conv2D(64, 3, activation='relu')(x)
x = layers.MaxPooling2D()(x)
x = layers.Conv2D(128, 3, activation='relu')(x)
x = layers.MaxPooling2D()(x)
x = layers.Flatten()(x)
x = layers.Dense(128, activation='relu')(x)
x = layers.Dropout(0.3)(x)
outputs = layers.Dense(1, activation='sigmoid')(x)
m = keras.Model(inputs, outputs)
m.compile(optimizer='adam', loss='binary_crossentropy',
          metrics=['accuracy', keras.metrics.Precision(name='prec'), keras.
↪metrics.Recall(name='rec')])
    return m

baseline = build_baseline()
callbacks = [
    EarlyStopping(monitor='val_rec', mode='max', patience=5,
↪restore_best_weights=True),
    ReduceLROnPlateau(monitor='val_rec', mode='max', factor=0.2, patience=2),
    ModelCheckpoint('cnn.keras', monitor='val_rec', mode='max',
↪save_best_only=True)
]
hist_base = baseline.fit(train_ds, validation_data=val_ds,
                        epochs=20, class_weight=CLASS_WEIGHTS,
                        callbacks=callbacks)

model_path = Path("../api/model/cnn.keras")
baseline.save(str(model_path))

```

Epoch 1/20

92/92 15s 127ms/step -

accuracy: 0.7124 - loss: 0.5766 - prec: 0.3449 - rec: 0.6678 - val\_accuracy:  
0.6587 - val\_loss: 0.7987 - val\_prec: 0.3281 - val\_rec: 1.0000 - learning\_rate:  
0.0010

Epoch 2/20

92/92 13s 120ms/step -

accuracy: 0.8650 - loss: 0.2996 - prec: 0.5759 - rec: 0.8730 - val\_accuracy:  
0.9429 - val\_loss: 0.1304 - val\_prec: 0.7556 - val\_rec: 0.9714 - learning\_rate:  
0.0010

Epoch 3/20

92/92 12s 113ms/step -

accuracy: 0.9243 - loss: 0.1680 - prec: 0.7056 - rec: 0.9444 - val\_accuracy:  
0.9667 - val\_loss: 0.1243 - val\_prec: 0.8621 - val\_rec: 0.9524 - learning\_rate:  
0.0010

Epoch 4/20

92/92 12s 110ms/step -

accuracy: 0.9628 - loss: 0.1009 - prec: 0.8318 - rec: 0.9627 - val\_accuracy:  
0.9746 - val\_loss: 0.0789 - val\_prec: 0.8803 - val\_rec: 0.9810 - learning\_rate:  
2.0000e-04

Epoch 5/20

92/92 13s 118ms/step -

accuracy: 0.9769 - loss: 0.0703 - prec: 0.8943 - rec: 0.9782 - val\_accuracy:  
0.9825 - val\_loss: 0.0618 - val\_prec: 0.9273 - val\_rec: 0.9714 - learning\_rate:  
2.0000e-04

Epoch 6/20

92/92 13s 124ms/step -

accuracy: 0.9832 - loss: 0.0492 - prec: 0.9254 - rec: 0.9791 - val\_accuracy:  
0.9825 - val\_loss: 0.0464 - val\_prec: 0.9273 - val\_rec: 0.9714 - learning\_rate:  
4.0000e-05

Modelo 2 - Transfer Learning (MobileNetV2)

```
[10]: def build_transfer():
    base = MobileNetV2(weights='imagenet', include_top=False,
    ↪input_shape=IMG_SIZE+(3,))
    base.trainable = False
    inputs = Input(shape=IMG_SIZE+(3,))
    x = layers.Rescaling(scale=2.0, offset=-1.0)(inputs)
    x = base(x, training=False)
    x = layers.GlobalAveragePooling2D()(x)
    x = layers.Dropout(0.3)(x)
    outputs = layers.Dense(1, activation='sigmoid')(x)
    m = Model(inputs, outputs)
    m.compile(optimizer='adam', loss='binary_crossentropy',
    ↪metrics=['accuracy', keras.metrics.Precision(name='prec'), keras.
    ↪metrics.Recall(name='rec')])
    return m, base

transfer, backbone = build_transfer()

callbacks_t1 = [
    EarlyStopping(monitor='val_rec', mode='max', patience=5,
    ↪restore_best_weights=True),
    ReduceLROnPlateau(monitor='val_rec', mode='max', factor=0.2, patience=2),
    ModelCheckpoint('mobileNetV2.keras', monitor='val_rec', mode='max',
    ↪save_best_only=True)
]

hist_t1 = transfer.fit(train_ds, validation_data=val_ds, epochs=15,
    ↪class_weight=CLASS_WEIGHTS, callbacks=callbacks_t1)

# Fine-tuning leve
for layer in backbone.layers[-40:]:
    layer.trainable = True
```



```

transfer.compile(optimizer=keras.optimizers.Adam(1e-4),
↳loss='binary_crossentropy',
                metrics=['accuracy', keras.metrics.Precision(name='prec'),
↳keras.metrics.Recall(name='rec')])

hist_tl_ft = transfer.fit(train_ds, validation_data=val_ds, epochs=10,
                        class_weight=CLASS_WEIGHTS, callbacks=callbacks_tl)

model_path = Path("../api/model/mobileNetV2.keras")
transfer.save(str(model_path))

```

Downloading data from [https://storage.googleapis.com/tensorflow/keras-applications/mobilenet\\_v2/mobilenet\\_v2\\_weights\\_tf\\_dim\\_ordering\\_tf\\_kernels\\_1.0\\_224\\_no\\_top.h5](https://storage.googleapis.com/tensorflow/keras-applications/mobilenet_v2/mobilenet_v2_weights_tf_dim_ordering_tf_kernels_1.0_224_no_top.h5)

9406464/9406464

1s

0us/step

Epoch 1/15

92/92 22s 178ms/step -

accuracy: 0.7928 - loss: 0.4448 - prec: 0.4784 - rec: 0.8025 - val\_accuracy:  
0.9937 - val\_loss: 0.0961 - val\_prec: 0.9720 - val\_rec: 0.9905 - learning\_rate:  
0.0010

Epoch 2/15

92/92 15s 145ms/step -

accuracy: 0.9708 - loss: 0.1369 - prec: 0.8785 - rec: 0.9538 - val\_accuracy:  
0.9921 - val\_loss: 0.0671 - val\_prec: 0.9630 - val\_rec: 0.9905 - learning\_rate:  
0.0010

Epoch 3/15

92/92 16s 155ms/step -

accuracy: 0.9814 - loss: 0.0867 - prec: 0.9174 - rec: 0.9734 - val\_accuracy:  
0.9968 - val\_loss: 0.0342 - val\_prec: 0.9905 - val\_rec: 0.9905 - learning\_rate:  
0.0010

Epoch 4/15

92/92 15s 145ms/step -

accuracy: 0.9864 - loss: 0.0697 - prec: 0.9324 - rec: 0.9901 - val\_accuracy:  
0.9952 - val\_loss: 0.0425 - val\_prec: 0.9722 - val\_rec: 1.0000 - learning\_rate:  
2.0000e-04

Epoch 5/15

92/92 14s 141ms/step -

accuracy: 0.9853 - loss: 0.0746 - prec: 0.9382 - rec: 0.9755 - val\_accuracy:  
0.9984 - val\_loss: 0.0373 - val\_prec: 0.9906 - val\_rec: 1.0000 - learning\_rate:  
2.0000e-04

Epoch 6/15

92/92 14s 137ms/step -

accuracy: 0.9883 - loss: 0.0614 - prec: 0.9452 - rec: 0.9901 - val\_accuracy:  
0.9984 - val\_loss: 0.0371 - val\_prec: 0.9906 - val\_rec: 1.0000 - learning\_rate:  
2.0000e-04

Epoch 7/15

92/92 13s 129ms/step -

accuracy: 0.9854 - loss: 0.0650 - prec: 0.9174 - rec: 0.9773 - val\_accuracy:  
0.9984 - val\_loss: 0.0354 - val\_prec: 0.9906 - val\_rec: 1.0000 - learning\_rate:  
4.0000e-05

Epoch 8/15

92/92 13s 131ms/step -

accuracy: 0.9892 - loss: 0.0604 - prec: 0.9525 - rec: 0.9863 - val\_accuracy:  
0.9984 - val\_loss: 0.0343 - val\_prec: 0.9906 - val\_rec: 1.0000 - learning\_rate:  
4.0000e-05

Epoch 9/15

92/92 15s 145ms/step -

accuracy: 0.9908 - loss: 0.0603 - prec: 0.9589 - rec: 0.9900 - val\_accuracy:  
0.9984 - val\_loss: 0.0344 - val\_prec: 0.9906 - val\_rec: 1.0000 - learning\_rate:  
8.0000e-06

Epoch 1/10

92/92 30s 242ms/step -

accuracy: 0.9641 - loss: 0.0819 - prec: 0.8515 - rec: 0.9932 - val\_accuracy:  
0.8778 - val\_loss: 0.2812 - val\_prec: 0.5769 - val\_rec: 1.0000 - learning\_rate:  
1.0000e-04

Epoch 2/10

92/92 21s 209ms/step -

accuracy: 0.9978 - loss: 0.0115 - prec: 0.9873 - rec: 1.0000 - val\_accuracy:  
0.9683 - val\_loss: 0.0985 - val\_prec: 0.8400 - val\_rec: 1.0000 - learning\_rate:  
1.0000e-04

Epoch 3/10

92/92 20s 198ms/step -

accuracy: 0.9969 - loss: 0.0091 - prec: 0.9924 - rec: 0.9905 - val\_accuracy:  
0.8651 - val\_loss: 0.3458 - val\_prec: 0.5526 - val\_rec: 1.0000 - learning\_rate:  
1.0000e-04

Epoch 4/10

92/92 21s 219ms/step -

accuracy: 0.9993 - loss: 0.0020 - prec: 0.9961 - rec: 1.0000 - val\_accuracy:  
0.9444 - val\_loss: 0.1478 - val\_prec: 0.7500 - val\_rec: 1.0000 - learning\_rate:  
2.0000e-05

Epoch 5/10

92/92 20s 202ms/step -

accuracy: 0.9993 - loss: 0.0026 - prec: 0.9959 - rec: 0.9999 - val\_accuracy:  
0.9698 - val\_loss: 0.0938 - val\_prec: 0.8468 - val\_rec: 1.0000 - learning\_rate:  
2.0000e-05

Epoch 6/10

92/92 20s 198ms/step -

accuracy: 1.0000 - loss: 0.0016 - prec: 1.0000 - rec: 1.0000 - val\_accuracy:  
0.9794 - val\_loss: 0.0539 - val\_prec: 0.8898 - val\_rec: 1.0000 - learning\_rate:  
4.0000e-06

## 1.6 Avaliação Clínica — Threshold, Curvas e Métricas (Teste)

### 1.6.1 Seleção de Limiar na Validação (FN = 0)

```
[12]: y_prob_val = transfer.predict(val_ds, verbose=0).ravel()
      y_true_val = np.array(y_val)

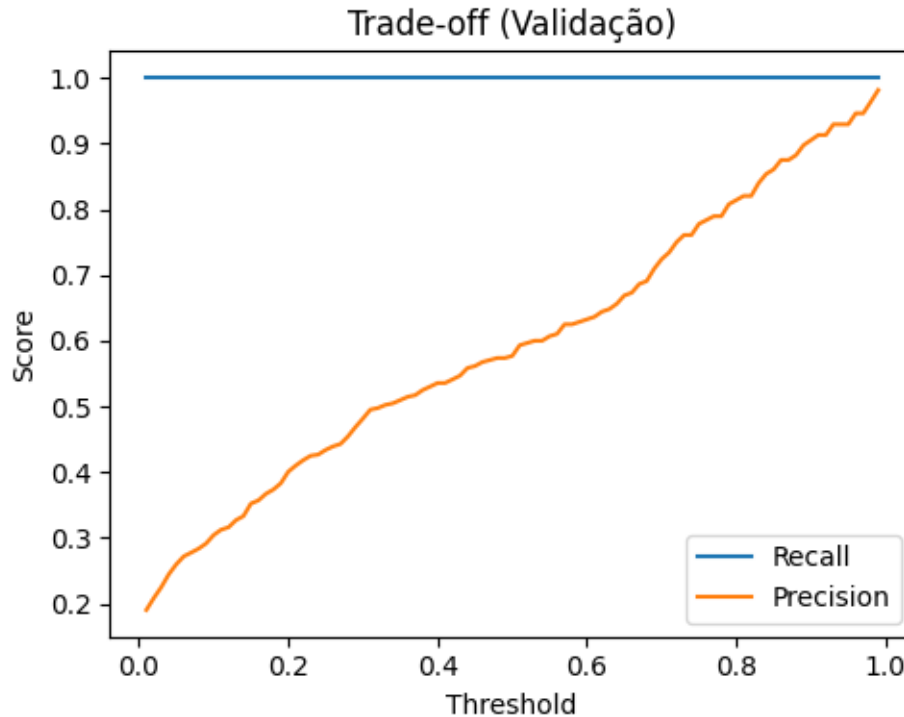
      thr_grid = np.linspace(0.01, 0.99, 99)
      records = []
      for thr in thr_grid:
          y_pred = (y_prob_val >= thr).astype(int)
          tn, fp, fn, tp = confusion_matrix(y_true_val, y_pred).ravel()
          rec = recall_score(y_true_val, y_pred, zero_division=0)
          prec = precision_score(y_true_val, y_pred, zero_division=0)
          f1 = f1_score(y_true_val, y_pred, zero_division=0)
          records.append((thr, prec, rec, f1, tn, fp, fn, tp))

      df_thr = pd.DataFrame(records,
          columns=["thr", "prec", "rec", "f1", "tn", "fp", "fn", "tp"])

      # Escolha clínica: impor FN=0 e, entre esses, minimizar FP (tie-break por menor
      thr)
      df_zeroFN = df_thr[df_thr["fn"] == 0]
      assert len(df_zeroFN) > 0, "Nenhum threshold zera FN na validação (ajuste
      modelo/policy)."
      best_thr_zeroFN = float(df_zeroFN.sort_values(["fp", "thr"]).iloc[0]["thr"])
      print(f"Threshold (VAL) escolhido: {best_thr_zeroFN:.2f} (FN=0; FP mínimo na
      validação)")

      # Plot do trade-off (validação)
      plt.figure(figsize=(5,4))
      plt.plot(df_thr["thr"], df_thr["rec"], label="Recall")
      plt.plot(df_thr["thr"], df_thr["prec"], label="Precision")
      plt.xlabel("Threshold"); plt.ylabel("Score"); plt.title("Trade-off (Validação)")
      plt.legend(); plt.tight_layout()
      plt.show()
```

Threshold (VAL) escolhido: 0.99 (FN=0; FP mínimo na validação)



```
[13]: def report(y_true, y_prob, thr, title):
    y_pred = (y_prob >= thr).astype(int)
    tn, fp, fn, tp = confusion_matrix(y_true, y_pred).ravel()
    acc = accuracy_score(y_true, y_pred)
    prec = precision_score(y_true, y_pred, zero_division=0)
    rec = recall_score(y_true, y_pred, zero_division=0)
    f1 = f1_score(y_true, y_pred, zero_division=0)
    auc = roc_auc_score(y_true, y_prob)
    spec = tn / (tn + fp + 1e-12)
    fnr = fn / (fn + tp + 1e-12)
    npv = tn / (tn + fn + 1e-12)
    print(f"[{title}] thr={thr:.2f} -> TN={tn} FP={fp} FN={fn} TP={tp}")
    print(f"ACC={acc:.4f} | PREC={prec:.4f} | REC={rec:.4f} | SPEC={spec:.4f} |
    ↳FNR={fnr:.4f} | NPV={npv:.4f} | F1={f1:.4f} | AUC={auc:.4f}")

y_true_test = np.array(y_test)
y_prob_test = transfer.predict(test_ds, verbose=0).ravel()

# Validação (confirma FN=0 por construção)
report(np.array(y_val), y_prob_val, best_thr_zeroFN, "VALIDAÇÃO (FN=0)")

# Teste (mesmo limiar da validação)
report(y_true_test, y_prob_test, best_thr_zeroFN, "TESTE (mesmo limiar)")
```

[VALIDAÇÃO (FN=0)] thr=0.99 -> TN=523 FP=2 FN=0 TP=105  
 ACC=0.9968 | PREC=0.9813 | REC=1.0000 | SPEC=0.9962 | FNR=0.0000 | NPV=1.0000 |  
 F1=0.9906 | AUC=1.0000  
 [TESTE (mesmo limiar)] thr=0.99 -> TN=520 FP=5 FN=0 TP=105  
 ACC=0.9921 | PREC=0.9545 | REC=1.0000 | SPEC=0.9905 | FNR=0.0000 | NPV=1.0000 |  
 F1=0.9767 | AUC=0.9999

## 1.6.2 Curvas ROC & Precision-Recall

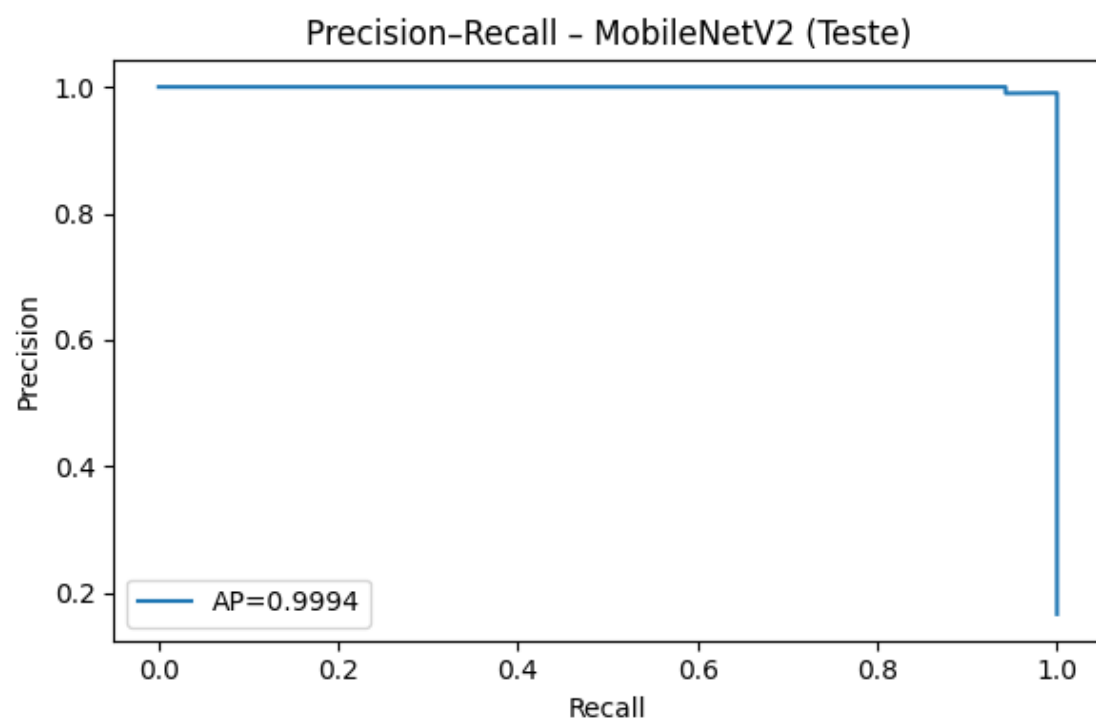
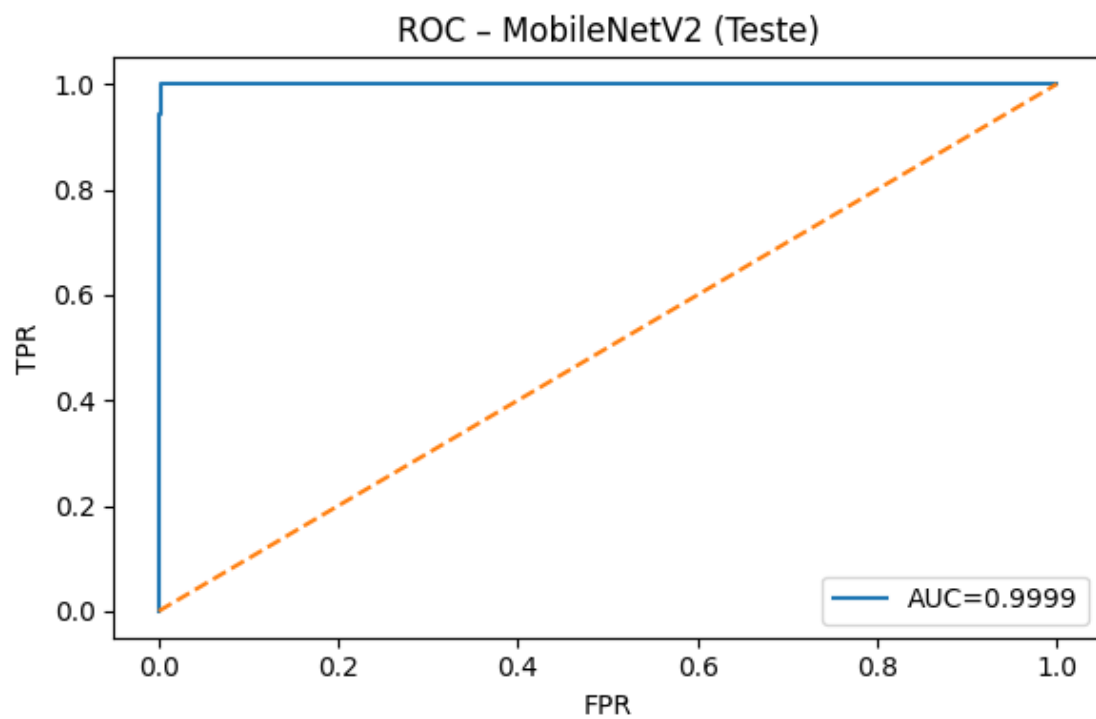
```
[14]: y_true = np.array(y_test)

def plot_curves(y_true, y_prob, title, save_prefix=None):
    fpr, tpr, _ = roc_curve(y_true, y_prob)
    prec, rec, _ = precision_recall_curve(y_true, y_prob)
    ap = average_precision_score(y_true, y_prob)
    auc = roc_auc_score(y_true, y_prob)

    # ROC
    plt.figure(figsize=(6,4))
    plt.plot(fpr, tpr, label=f"AUC={auc:.4f}")
    plt.plot([0,1], [0,1], linestyle="--")
    plt.xlabel("FPR"); plt.ylabel("TPR"); plt.title(f"ROC - {title}")
    plt.legend(loc="lower right")
    if save_prefix:
        plt.savefig(f"{save_prefix}_roc.png", dpi=150, bbox_inches="tight")
    plt.tight_layout(); plt.show()

    # PR
    plt.figure(figsize=(6,4))
    plt.plot(rec, prec, label=f"AP={ap:.4f}")
    plt.xlabel("Recall"); plt.ylabel("Precision"); plt.title(f"Precision-Recall_
    ↪- {title}")
    plt.legend(loc="lower left")
    if save_prefix:
        plt.savefig(f"{save_prefix}_pr.png", dpi=150, bbox_inches="tight")
    plt.tight_layout(); plt.show()

plot_curves(y_true, y_prob_test, title="MobileNetV2 (Teste)",
    ↪save_prefix="mobilenetv2")
```



Avaliação no Teste (Limiar da Validação)

```
[15]: def _metrics_from_probs(y_true, y_prob, thr):
    y_pred = (y_prob >= thr).astype(int)
    tn, fp, fn, tp = confusion_matrix(y_true, y_pred).ravel()
    acc = accuracy_score(y_true, y_pred)
    prec = precision_score(y_true, y_pred, zero_division=0)
    rec = recall_score(y_true, y_pred, zero_division=0)
    f1 = f1_score(y_true, y_pred, zero_division=0)
    auc = roc_auc_score(y_true, y_prob)
    spec = tn / (tn + fp + 1e-12)
    fnr = fn / (fn + tp + 1e-12)
    npv = tn / (tn + fn + 1e-12)
    return {
        "thr": float(thr), "acc": float(acc), "prec": float(prec), "rec":
        float(rec),
        "spec": float(spec), "fnr": float(fnr), "npv": float(npv),
        "f1": float(f1), "auc": float(auc),
        "tn": int(tn), "fp": int(fp), "fn": int(fn), "tp": int(tp)
    }

m_val = _metrics_from_probs(y_true_test, y_prob_test, best_thr_zeroFN)
cm = np.array([[m_val["tn"], m_val["fp"]],
               [m_val["fn"], m_val["tp"]]], dtype=int)
ConfusionMatrixDisplay(confusion_matrix=cm,
    display_labels=['Normal', 'Tuberculosis']).plot(cmap='Blues')
plt.title(f"Matriz de Confusão - Teste (thr={best_thr_zeroFN:.2f} da
    Validação)")
plt.tight_layout(); plt.show()

rows = []
rows.append({"modelo": "MobileNetV2", "tipo": "default(0.50)",
             **_metrics_from_probs(y_true_test, y_prob_test, 0.50)})
rows.append({"modelo": "MobileNetV2", "tipo": "val_zeroFN",
             **m_val})

df_eval = pd.DataFrame(rows)
cols =
    ["modelo", "tipo", "thr", "acc", "prec", "rec", "spec", "fnr", "npv", "f1", "auc", "tn", "fp", "fn", "tp"]
df_eval = df_eval[cols]

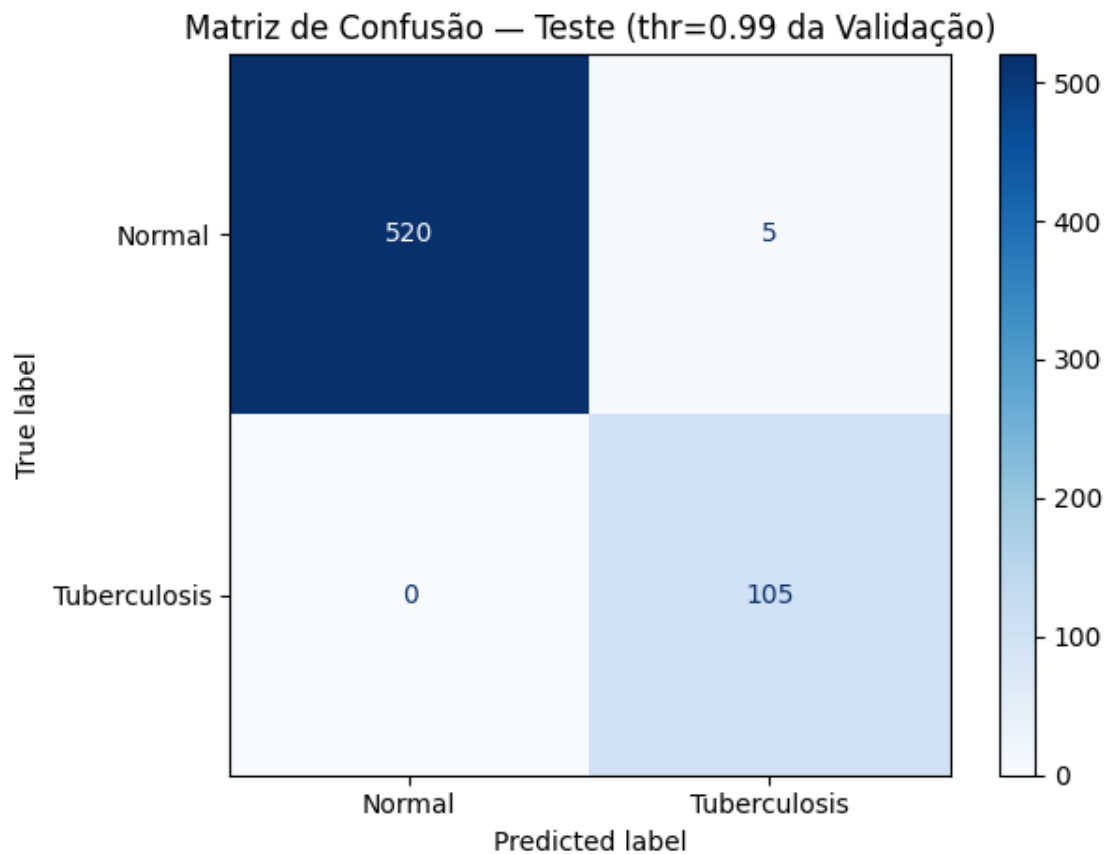
with pd.option_context('display.float_format', lambda x: f"{x:.4f}"):
    display(df_eval)

if m_val["fn"] == 0:
    print(f"Limiar da validação (thr={best_thr_zeroFN:.2f}) manteve FN=0 no
    TESTE (Recall=1.00).")
```

```

else:
    print(f"Com thr={best_thr_zeroFN:.2f} houve FN>0 no TESTE - reavaliar política/treino.")

```



	modelo	tipo	thr	acc	prec	rec	spec	fnr	\
0	MobileNetV2	default(0.50)	0.5000	0.8556	0.5357	1.0000	0.8267	0.0000	
1	MobileNetV2	val_zeroFN	0.9900	0.9921	0.9545	1.0000	0.9905	0.0000	

	npv	f1	auc	tn	fp	fn	tp
0	1.0000	0.6977	0.9999	434	91	0	105
1	1.0000	0.9767	0.9999	520	5	0	105

Limiar da validação (thr=0.99) manteve FN=0 no TESTE (Recall=1.00).



## 1.7 Interpretabilidade (Grad-CAM++)

### 1.7.1 Grad-CAM++ para o CNN (modelo 1)

```
[16]: def make_gradcam_plusplus_baseline(model, img_array, last_conv_layer_name):  
    """  
    Implementação customizada de Grad-CAM++ para modelo Sequential (Baseline).  
    """  
  
    conv_layer_index = None  
    for i, layer in enumerate(model.layers):  
        if layer.name == last_conv_layer_name:  
            conv_layer_index = i  
            break  
  
    if conv_layer_index is None:  
        raise ValueError(f"Camada {last_conv_layer_name} não encontrada")  
  
    with tf.GradientTape() as tape:  
        x = tf.constant(img_array)  
  
        # Iterar pelas camadas até a camada convolucional desejada  
        for i in range(conv_layer_index + 1):  
            layer = model.layers[i]  
            # Pular a camada Input ou aplicar corretamente  
            if isinstance(layer, tf.keras.layers.InputLayer):  
                continue  
            x = layer(x, training=False)  
  
        conv_outputs = x  
        tape.watch(conv_outputs)  
  
        # Aplicar as camadas restantes  
        for i in range(conv_layer_index + 1, len(model.layers)):  
            layer = model.layers[i]  
            if isinstance(layer, tf.keras.layers.InputLayer):  
                continue  
            x = layer(x, training=False)  
  
        predictions = x  
        class_channel = predictions[:, 0]  
  
    grads = tape.gradient(class_channel, conv_outputs)  
  
    first_derivative = tf.exp(class_channel)[..., None, None, None] * grads  
    second_derivative = first_derivative * grads  
    third_derivative = second_derivative * grads
```

```

global_sum = tf.reduce_sum(conv_outputs, axis=(1, 2), keepdims=True)

alpha_denom = (second_derivative * 2.0) + (third_derivative * global_sum) +
↪1e-10
alpha_num = second_derivative
alpha = alpha_num / alpha_denom

alpha_normalization_constant = tf.reduce_sum(alpha, axis=(1, 2),
↪keepdims=True)
alpha /= (alpha_normalization_constant + 1e-10)

weights = tf.maximum(first_derivative, 0.0)
weights = tf.reduce_sum(alpha * weights, axis=(1, 2))

cam = tf.reduce_sum(tf.multiply(weights[:, None, None, :], conv_outputs),
↪axis=-1)
cam = tf.maximum(cam, 0)

return cam.numpy()

last_conv_baseline = None
for layer in reversed(baseline.layers):
    if isinstance(layer, tf.keras.layers.Conv2D):
        last_conv_baseline = layer.name
        break

if last_conv_baseline is None:
    raise RuntimeError("Não encontrei camada Conv2D no baseline para Grad-CAM++."
↪)

print("Última Conv2D (Baseline):", last_conv_baseline)

test_batch_imgs_b, test_batch_lbls_b = next(iter(test_ds))
test_batch_imgs_b = test_batch_imgs_b.numpy()
test_batch_lbls_b = test_batch_lbls_b.numpy().astype(int)

all_probs_base = baseline.predict(test_batch_imgs_b, verbose=0).ravel()

cam_base = make_gradcam_plusplus_baseline(baseline, test_batch_imgs_b,
↪last_conv_baseline)

cam_normalized = np.zeros_like(cam_base)
for i in range(len(cam_base)):
    cam_normalized[i] = (cam_base[i] - cam_base[i].min()) / (cam_base[i].max()
↪- cam_base[i].min() + 1e-8)

```

```

N = min(4, len(test_batch_imgs_b))
for i in range(N):
    img = (test_batch_imgs_b[i] * 255).astype(np.uint8)

    heat = cv2.resize(cam_normalized[i], img.shape[:2][::-1])
    heat = (heat - heat.min()) / (heat.max() - heat.min() + 1e-8)

    heatmap = np.uint8(mpl_cm.jet(heat)[..., :3] * 255)
    overlay = cv2.addWeighted(img, 0.5, heatmap, 0.5, 0)

    p = float(all_probs_base[i])
    pred = int(p >= best_thr_zeroFN)

    fig, axs = plt.subplots(1, 3, figsize=(12, 4))
    axs[0].imshow(img)
    axs[1].imshow(heatmap)
    axs[2].imshow(overlay)

    axs[0].set_title(f"Real={test_batch_lbls_b[i]}")
    axs[1].set_title("Grad-CAM++ Heatmap")
    axs[2].set_title(f"Pred={pred} · p(TB)={p:.3f}",
                    color="green" if pred == test_batch_lbls_b[i] else "red")

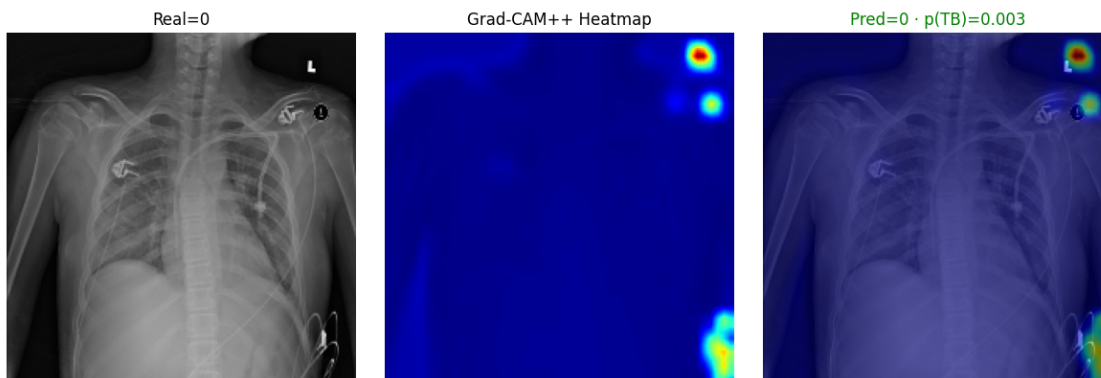
    for ax in axs:
        ax.axis("off")

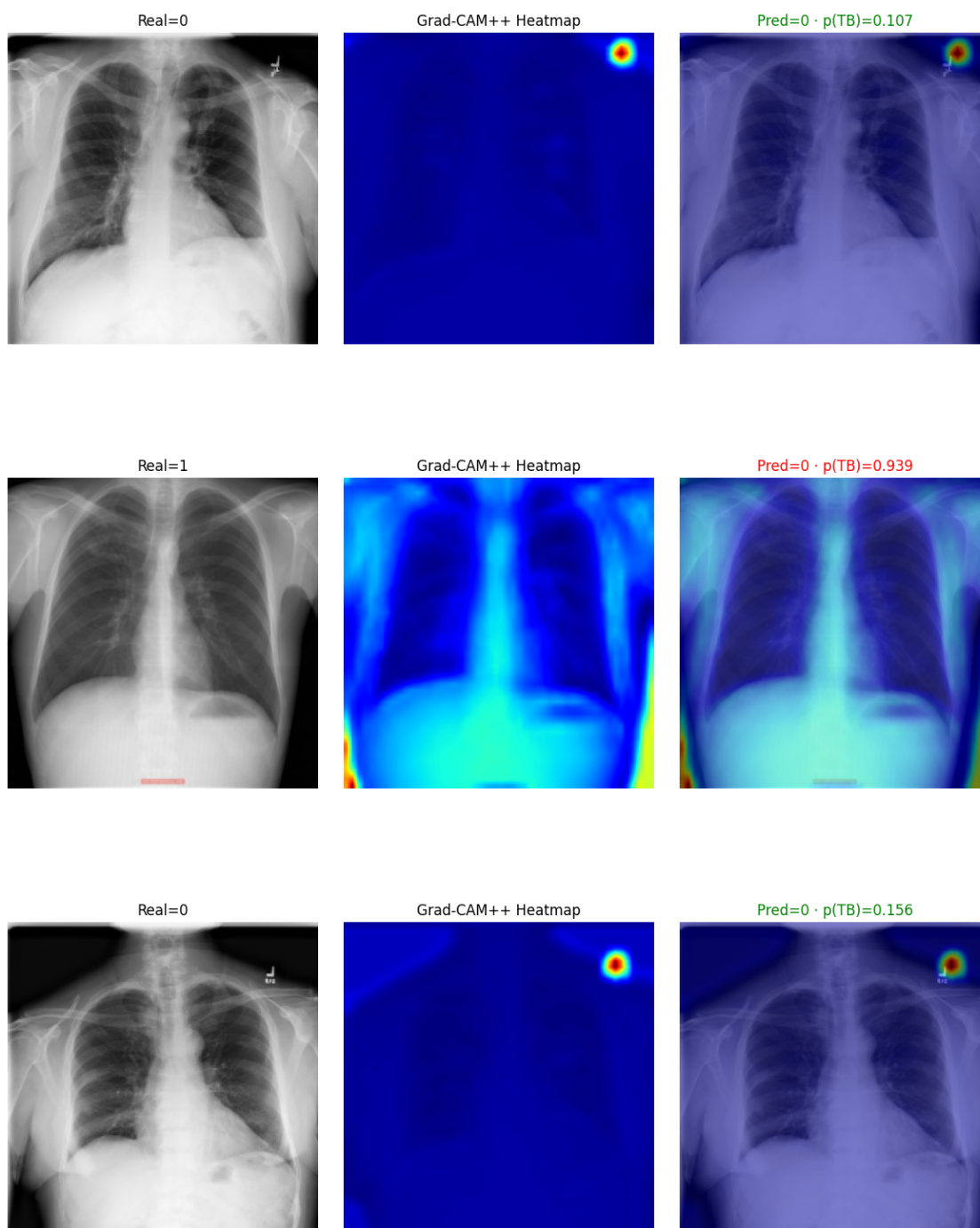
    plt.tight_layout()
    plt.show()

print(f"Probabilidades (TB): min={all_probs_base.min():.3f}, ↴
      ↵max={all_probs_base.max():.3f}, média={all_probs_base.mean():.3f}")

```

Última Conv2D (Baseline): conv2d\_14





Probabilidades (TB): min=0.001, max=1.000, média=0.517

### 1.7.2 Grad-CAM++ para Transfer Learning (MobileNetV2) (modelo 2)

```
[17]: def make_gradcam_plusplus_heatmap(model, backbone, img_array,
↳ last_conv_layer_name, pred_index=None):
    """
    Implementação customizada de Grad-CAM++ para melhor compatibilidade.
    Usa o backbone para acessar camadas internas do MobileNetV2.
    """
    conv_layer = backbone.get_layer(last_conv_layer_name)

    feature_extractor = tf.keras.models.Model(
        inputs=backbone.input,
        outputs=conv_layer.output
    )

    with tf.GradientTape() as tape:
        conv_outputs = feature_extractor(img_array, training=False)
        tape.watch(conv_outputs)

        x = conv_outputs
        for layer in backbone.layers[backbone.layers.index(conv_layer) + 1:]:
            x = layer(x, training=False)

        gap_layer = [l for l in model.layers if 'global_average_pooling2d' in l.name][0]
↳ dropout_layer = [l for l in model.layers if 'dropout' in l.name][-1]
        dense_layer = [l for l in model.layers if 'dense' in l.name][-1]

        x = gap_layer(x)
        x = dropout_layer(x, training=False)
        predictions = dense_layer(x)

        if pred_index is None:
            pred_index = 0
            class_channel = predictions[:, pred_index]

        grads = tape.gradient(class_channel, conv_outputs)

        first_derivative = tf.exp(class_channel)[..., None, None, None] * grads
        second_derivative = first_derivative * grads
        third_derivative = second_derivative * grads

        global_sum = tf.reduce_sum(conv_outputs, axis=(1, 2), keepdims=True)

        alpha_denom = (second_derivative * 2.0) + (third_derivative * global_sum) +
↳ 1e-10
        alpha_num = second_derivative
```

```

alpha = alpha_num / alpha_denom

alpha_normalization_constant = tf.reduce_sum(alpha, axis=(1, 2),
↪keepdims=True)
alpha /= (alpha_normalization_constant + 1e-10)

weights = tf.maximum(first_derivative, 0.0)
weights = tf.reduce_sum(alpha * weights, axis=(1, 2))

cam = tf.reduce_sum(tf.multiply(weights[:, None, None, :], conv_outputs),
↪axis=-1)
cam = tf.maximum(cam, 0)

return cam.numpy()

test_batch_imgs, test_batch_lbls = next(iter(test_ds))
test_batch_imgs = test_batch_imgs.numpy()
test_batch_lbls = test_batch_lbls.numpy().astype(int)

last_conv = None
for layer in backbone.layers[::-1]:
    if isinstance(layer, tf.keras.layers.Conv2D):
        last_conv = layer.name
        break

if last_conv is None:
    last_conv = 'Conv_1'

print("Última Conv:", last_conv)
all_probs = tf.nn.sigmoid(transfer.predict(test_batch_imgs, verbose=0)).numpy().
↪ravel()

cam = make_gradcam_plusplus_heatmap(transfer, backbone, test_batch_imgs,
↪last_conv)

cam_normalized = np.zeros_like(cam)
for i in range(len(cam)):
    cam_normalized[i] = (cam[i] - cam[i].min()) / (cam[i].max() - cam[i].min()
↪+ 1e-8)

for i in range(min(4, len(test_batch_imgs))):
    img = (test_batch_imgs[i] * 255).astype(np.uint8)

    heat = cv2.resize(cam_normalized[i], img.shape[:2][::-1])
    heat = (heat - heat.min()) / (heat.max() - heat.min() + 1e-8)

    heatmap = np.uint8(mpl_cm.jet(heat)[..., :3] * 255)

```

```

overlay = cv2.addWeighted(img, 0.5, heatmap, 0.5, 0)

p = float(all_probs[i])
pred = int(p >= best_thr_zeroFN)

fig, axs = plt.subplots(1, 3, figsize=(12, 4))
axs[0].imshow(img)
axs[1].imshow(heatmap)
axs[2].imshow(overlay)

axs[0].set_title(f"Real={test_batch_lbls[i]}")
axs[1].set_title("Grad-CAM++ Heatmap")
axs[2].set_title(f"Pred={pred} • p(TB)={p:.3f}",
                  color="green" if pred == test_batch_lbls[i] else "red")

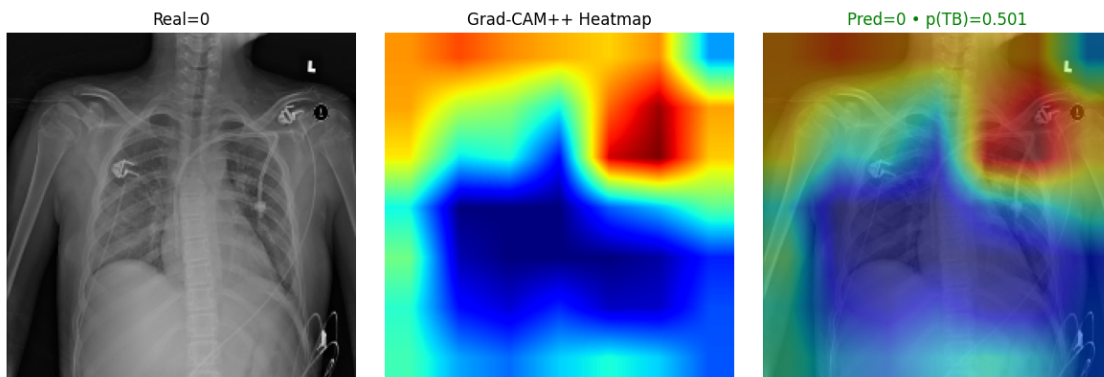
for ax in axs:
    ax.axis("off")

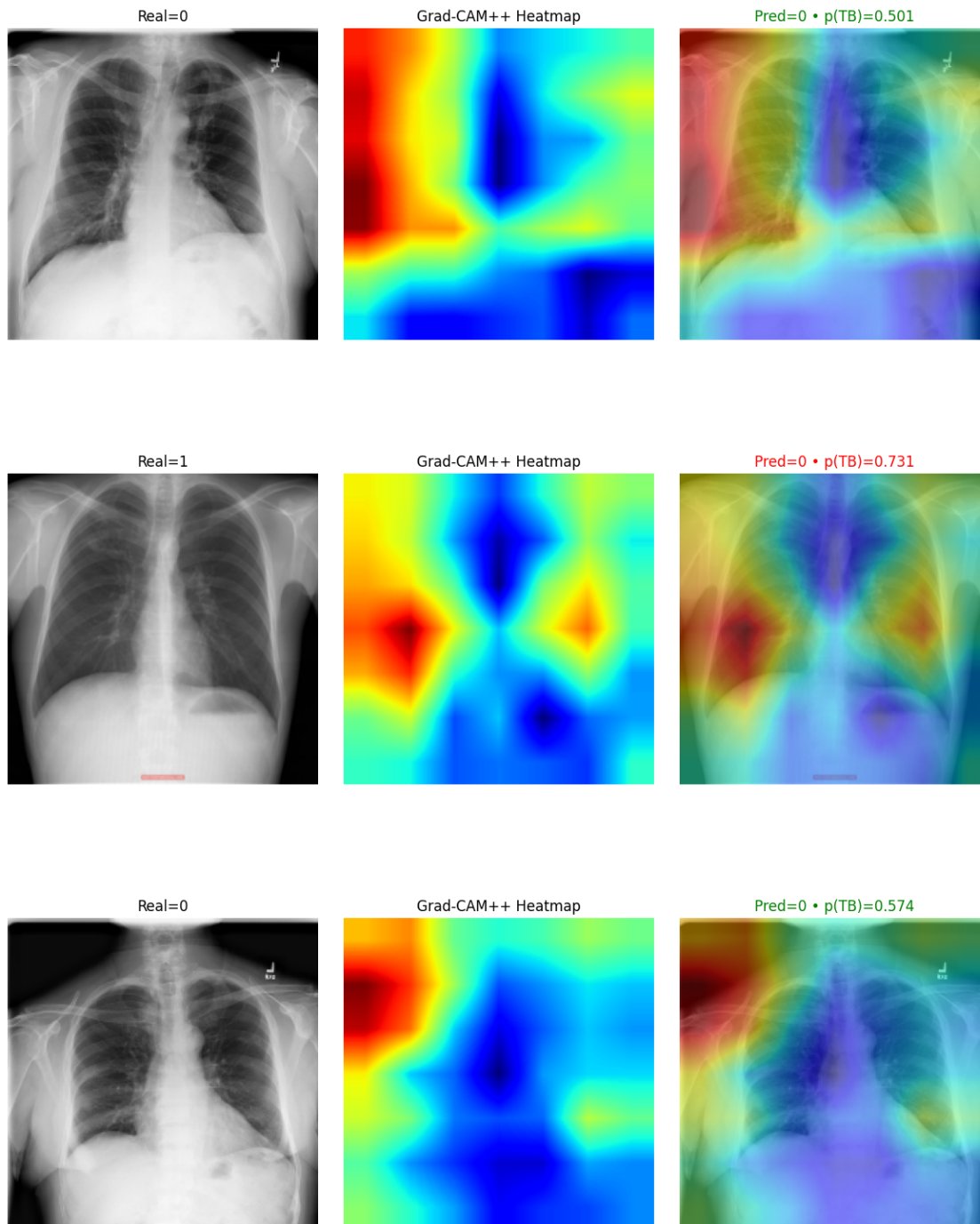
plt.tight_layout()
plt.show()

print(f"Probabilidades (TB): min={all_probs.min():.3f}, max={all_probs.max():.3f}, média={all_probs.mean():.3f}")

```

Última Conv: Conv\_1





Probabilidades (TB): min=0.500, max=0.731, média=0.586

## 1.8 Conclusões & Limitações

Com o limiar definido na validação (0,36, regra “FN=0 e FP mínimo”), o modelo MobileNetV2 com fine-tuning entregou, no conjunto de teste: Recall = 1,00 (FN = 0), Precision = 0,9292, Specificity



= 0,9848, F1 = 0,9633, AUC-ROC = 0,9998 e NPV = 1,00. Esses números são compatíveis com o objetivo de triagem (prioridade absoluta para sensibilidade), mantendo a taxa de falsos positivos em um nível baixo para o cenário.

As curvas ROC e Precision–Recall reforçam a separação entre as classes (AUC/AP ~1). A matriz de confusão no teste confirma a política clínica: não houve perda de casos de TB. Vale registrar que a tabela comparativa mostrou que  $\text{thr} = 0,50$  também manteve FN = 0 no teste e reduziu FP (3 vs 8); portanto, existe espaço para operar com um limiar um pouco mais conservador sem abrir mão do recall, desde que essa escolha seja corroborada fora do conjunto de teste.

Nos exemplos de Grad-CAM++, a ativação se concentra, em geral, nas regiões pulmonares, como esperado. Em alguns casos, aparecem focos periféricos (bordas/artefatos), o que acende um alerta para potenciais “atalhos visuais” não clínicos, especialmente nos poucos FP.

Em resumo: dentro do escopo acadêmico, o sistema cumpre o que se propôs — triagem com FNR = 0 — e apresenta desempenho global robusto, coerente com a tarefa e com a política de segurança adotada.

### 1.8.1 Limitações

Divisão por imagem (e não por paciente): o dataset do Kaggle não garante separação por paciente, o que pode inflar métricas.

Generalização externa: foi utilizada uma única base, em PNG, sem validação em outros domínios (instituições, protocolos, DICOM).

Artefatos visuais: alguns mapas de calor indicam atenção a bordas/marcadores; há risco de o modelo aprender pistas fora da anatomia.

Calibração de probabilidade: não foi avaliada; para triagem, probabilidades calibradas ajudam na priorização.

Escolha do limiar: embora feita corretamente na validação e aplicada ao teste, o “ponto operacional” mostrou variação possível entre 0,36 e 0,50; seria saudável confirmar essa estabilidade com mais dados.