

```
In [9]: resnet_results = {

1: {"coverage": 0.899, "avg_size": 0.904,
    "class_cov": {
        "esantema-maculo-papuloso": 0.877,
        "esantema-morbilliforme": 0.920,
        "esantema-polimorfo-like": 0.848,
        "esantema-virale": 0.892,
        "esantema_iatrogeno_farmaco_indotta": 0.928,
        "orticaria": 0.904,
        "pediculosi": 0.930,
        "scabbia": 0.901,
        "varicella": 0.836
    }},

2: {"coverage": 0.898, "avg_size": 0.902,
    "class_cov": {
        "esantema-maculo-papuloso": 0.866,
        "esantema-morbilliforme": 0.926,
        "esantema-polimorfo-like": 0.825,
        "esantema-virale": 0.887,
        "esantema_iatrogeno_farmaco_indotta": 0.922,
        "orticaria": 0.915,
        "pediculosi": 0.891,
        "scabbia": 0.898,
        "varicella": 0.811
    }},

3: {"coverage": 0.900, "avg_size": 0.904,
    "class_cov": {
        "esantema-maculo-papuloso": 0.825,
        "esantema-morbilliforme": 0.915,
        "esantema-polimorfo-like": 0.846,
        "esantema-virale": 0.876,
        "esantema_iatrogeno_farmaco_indotta": 0.915,
        "orticaria": 0.927,
        "pediculosi": 0.915,
        "scabbia": 0.899,
        "varicella": 0.819
    }},

4: {"coverage": 0.899, "avg_size": 0.905,
    "class_cov": {
        "esantema-maculo-papuloso": 0.851,
        "esantema-morbilliforme": 0.936,
        "esantema-polimorfo-like": 0.862,
        "esantema-virale": 0.885,
        "esantema_iatrogeno_farmaco_indotta": 0.929,
        "orticaria": 0.908,
        "pediculosi": 0.930,
        "scabbia": 0.907,
        "varicella": 0.843
    }},
}
```

```
5: {"coverage": 0.900, "avg_size": 0.906,
    "class_cov": {
      "esantema-maculo-papuloso": 0.866,
      "esantema-morbilliforme": 0.936,
      "esantema-polimorfo-like": 0.853,
      "esantema-virale": 0.886,
      "esantema_iatrogeno_farmaco_indotta": 0.923,
      "orticaria": 0.911,
      "pediculosi": 0.903,
      "scabbia": 0.917,
      "varicella": 0.841
    }
  },

6: {"coverage": 0.902, "avg_size": 0.908,
    "class_cov": {
      "esantema-maculo-papuloso": 0.877,
      "esantema-morbilliforme": 0.925,
      "esantema-polimorfo-like": 0.865,
      "esantema-virale": 0.904,
      "esantema_iatrogeno_farmaco_indotta": 0.929,
      "orticaria": 0.899,
      "pediculosi": 0.937,
      "scabbia": 0.921,
      "varicella": 0.827
    }
  },

7: {"coverage": 0.900, "avg_size": 0.905,
    "class_cov": {
      "esantema-maculo-papuloso": 0.876,
      "esantema-morbilliforme": 0.944,
      "esantema-polimorfo-like": 0.864,
      "esantema-virale": 0.860,
      "esantema_iatrogeno_farmaco_indotta": 0.933,
      "orticaria": 0.919,
      "pediculosi": 0.927,
      "scabbia": 0.924,
      "varicella": 0.832
    }
  },

8: {"coverage": 0.900, "avg_size": 0.905,
    "class_cov": {
      "esantema-maculo-papuloso": 0.863,
      "esantema-morbilliforme": 0.955,
      "esantema-polimorfo-like": 0.860,
      "esantema-virale": 0.892,
      "esantema_iatrogeno_farmaco_indotta": 0.931,
      "orticaria": 0.906,
      "pediculosi": 0.914,
      "scabbia": 0.907,
      "varicella": 0.814
    }
  },

9: {"coverage": 0.900, "avg_size": 0.906,
    "class_cov": {
      "esantema-maculo-papuloso": 0.854,
      "esantema-morbilliforme": 0.925,
```

```
    "esantema-polimorfo-like": 0.843,  
    "esantema-virale": 0.877,  
    "esantema_iatrogeno_farmaco_indotta": 0.937,  
    "orticaria": 0.917,  
    "pediculosi": 0.916,  
    "scabbia": 0.913,  
    "varicella": 0.827  
  }},  
  
10: {"coverage": 0.899, "avg_size": 0.903,  
    "class_cov": {  
      "esantema-maculo-papuloso": 0.890,  
      "esantema-morbilliforme": 0.952,  
      "esantema-polimorfo-like": 0.840,  
      "esantema-virale": 0.884,  
      "esantema_iatrogeno_farmaco_indotta": 0.927,  
      "orticaria": 0.904,  
      "pediculosi": 0.924,  
      "scabbia": 0.920,  
      "varicella": 0.822  
    }},  
  
11: {"coverage": 0.900, "avg_size": 0.905,  
    "class_cov": {  
      "esantema-maculo-papuloso": 0.861,  
      "esantema-morbilliforme": 0.935,  
      "esantema-polimorfo-like": 0.845,  
      "esantema-virale": 0.880,  
      "esantema_iatrogeno_farmaco_indotta": 0.924,  
      "orticaria": 0.915,  
      "pediculosi": 0.915,  
      "scabbia": 0.910,  
      "varicella": 0.855  
    }},  
  
12: {"coverage": 0.900, "avg_size": 0.905,  
    "class_cov": {  
      "esantema-maculo-papuloso": 0.874,  
      "esantema-morbilliforme": 0.937,  
      "esantema-polimorfo-like": 0.857,  
      "esantema-virale": 0.898,  
      "esantema_iatrogeno_farmaco_indotta": 0.920,  
      "orticaria": 0.904,  
      "pediculosi": 0.914,  
      "scabbia": 0.907,  
      "varicella": 0.816  
    }},  
  
13: {"coverage": 0.899, "avg_size": 0.905,  
    "class_cov": {  
      "esantema-maculo-papuloso": 0.880,  
      "esantema-morbilliforme": 0.931,  
      "esantema-polimorfo-like": 0.849,  
      "esantema-virale": 0.887,  
      "esantema_iatrogeno_farmaco_indotta": 0.934,  
      "orticaria": 0.905,
```

```
        "pediculosi": 0.935,  
        "scabbia": 0.907,  
        "varicella": 0.829  
    }},  
  
14: {"coverage": 0.901, "avg_size": 0.906,  
    "class_cov": {  
        "esantema-maculo-papuloso": 0.864,  
        "esantema-morbilliforme": 0.948,  
        "esantema-polimorfo-like": 0.837,  
        "esantema-virale": 0.899,  
        "esantema_iatrogeno_farmaco_indotta": 0.918,  
        "orticaria": 0.905,  
        "pediculosi": 0.916,  
        "scabbia": 0.913,  
        "varicella": 0.851  
    }},  
  
15: {"coverage": 0.900, "avg_size": 0.904,  
    "class_cov": {  
        "esantema-maculo-papuloso": 0.851,  
        "esantema-morbilliforme": 0.931,  
        "esantema-polimorfo-like": 0.816,  
        "esantema-virale": 0.904,  
        "esantema_iatrogeno_farmaco_indotta": 0.944,  
        "orticaria": 0.903,  
        "pediculosi": 0.929,  
        "scabbia": 0.894,  
        "varicella": 0.819  
    }},  
  
16: {"coverage": 0.902, "avg_size": 0.908,  
    "class_cov": {  
        "esantema-maculo-papuloso": 0.872,  
        "esantema-morbilliforme": 0.951,  
        "esantema-polimorfo-like": 0.840,  
        "esantema-virale": 0.888,  
        "esantema_iatrogeno_farmaco_indotta": 0.928,  
        "orticaria": 0.914,  
        "pediculosi": 0.931,  
        "scabbia": 0.910,  
        "varicella": 0.826  
    }},  
  
17: {"coverage": 0.900, "avg_size": 0.905,  
    "class_cov": {  
        "esantema-maculo-papuloso": 0.868,  
        "esantema-morbilliforme": 0.943,  
        "esantema-polimorfo-like": 0.870,  
        "esantema-virale": 0.870,  
        "esantema_iatrogeno_farmaco_indotta": 0.939,  
        "orticaria": 0.912,  
        "pediculosi": 0.930,  
        "scabbia": 0.916,  
        "varicella": 0.855  
    }},
```

```

18: {"coverage": 0.901, "avg_size": 0.907,
    "class_cov": {
      "esantema-maculo-papuloso": 0.843,
      "esantema-morbilloforme": 0.939,
      "esantema-polimorfo-like": 0.810,
      "esantema-virale": 0.886,
      "esantema_iatrogeno_farmaco_indotta": 0.932,
      "orticaria": 0.917,
      "pediculosi": 0.912,
      "scabbia": 0.908,
      "varicella": 0.847
    }
  },

19: {"coverage": 0.901, "avg_size": 0.906,
    "class_cov": {
      "esantema-maculo-papuloso": 0.882,
      "esantema-morbilloforme": 0.952,
      "esantema-polimorfo-like": 0.862,
      "esantema-virale": 0.880,
      "esantema_iatrogeno_farmaco_indotta": 0.927,
      "orticaria": 0.914,
      "pediculosi": 0.905,
      "scabbia": 0.909,
      "varicella": 0.824
    }
  },

20: {"coverage": 0.899, "avg_size": 0.905,
    "class_cov": {
      "esantema-maculo-papuloso": 0.881,
      "esantema-morbilloforme": 0.926,
      "esantema-polimorfo-like": 0.850,
      "esantema-virale": 0.873,
      "esantema_iatrogeno_farmaco_indotta": 0.930,
      "orticaria": 0.916,
      "pediculosi": 0.905,
      "scabbia": 0.907,
      "varicella": 0.842
    }
  },
}

```

```

vit_results = {

  1: {"coverage": 0.900, "avg_size": 0.901,
    "class_cov": {
      "esantema-maculo-papuloso": 0.872,
      "esantema-morbilloforme": 0.931,
      "esantema-polimorfo-like": 0.780,
      "esantema-virale": 0.879,
      "esantema_iatrogeno_farmaco_indotta": 0.951,
      "orticaria": 0.923,
      "pediculosi": 0.887,
      "scabbia": 0.887,
      "varicella": 0.838
    }
  }
}

```

```
    }},  
  
3: {"coverage": 0.900, "avg_size": 0.901,  
    "class_cov": {  
        "esantema-maculo-papuloso": 0.868,  
        "esantema-morbilliforme": 0.951,  
        "esantema-polimorfo-like": 0.859,  
        "esantema-virale": 0.884,  
        "esantema_iatrogeno_farmaco_indotta": 0.935,  
        "orticaria": 0.917,  
        "pediculosi": 0.913,  
        "scabbia": 0.870,  
        "varicella": 0.854  
    }},  
  
2: {"coverage": 0.900, "avg_size": 0.901,  
    "class_cov": {  
        "esantema-maculo-papuloso": 0.873,  
        "esantema-morbilliforme": 0.916,  
        "esantema-polimorfo-like": 0.859,  
        "esantema-virale": 0.899,  
        "esantema_iatrogeno_farmaco_indotta": 0.926,  
        "orticaria": 0.900,  
        "pediculosi": 0.898,  
        "scabbia": 0.918,  
        "varicella": 0.852  
    }},  
  
4: {"coverage": 0.902, "avg_size": 0.903,  
    "class_cov": {  
        "esantema-maculo-papuloso": 0.899,  
        "esantema-morbilliforme": 0.936,  
        "esantema-polimorfo-like": 0.873,  
        "esantema-virale": 0.864,  
        "esantema_iatrogeno_farmaco_indotta": 0.957,  
        "orticaria": 0.925,  
        "pediculosi": 0.896,  
        "scabbia": 0.882,  
        "varicella": 0.844  
    }},  
  
5: {"coverage": 0.899, "avg_size": 0.900,  
    "class_cov": {  
        "esantema-maculo-papuloso": 0.862,  
        "esantema-morbilliforme": 0.911,  
        "esantema-polimorfo-like": 0.815,  
        "esantema-virale": 0.889,  
        "esantema_iatrogeno_farmaco_indotta": 0.920,  
        "orticaria": 0.920,  
        "pediculosi": 0.901,  
        "scabbia": 0.872,  
        "varicella": 0.863  
    }},  
  
6: {"coverage": 0.900, "avg_size": 0.901,  
    "class_cov": {
```

```
    "esantema-maculo-papuloso": 0.862,  
    "esantema-morbilliforme": 0.927,  
    "esantema-polimorfo-like": 0.834,  
    "esantema-virale": 0.879,  
    "esantema_iatrogeno_farmaco_indotta": 0.942,  
    "orticaria": 0.927,  
    "pediculosi": 0.910,  
    "scabbia": 0.851,  
    "varicella": 0.844  
  }},  
  
7: {"coverage": 0.900, "avg_size": 0.902,  
   "class_cov": {  
     "esantema-maculo-papuloso": 0.901,  
     "esantema-morbilliforme": 0.961,  
     "esantema-polimorfo-like": 0.886,  
     "esantema-virale": 0.880,  
     "esantema_iatrogeno_farmaco_indotta": 0.936,  
     "orticaria": 0.894,  
     "pediculosi": 0.939,  
     "scabbia": 0.938,  
     "varicella": 0.879  
   }},  
  
8: {"coverage": 0.900, "avg_size": 0.902,  
   "class_cov": {  
     "esantema-maculo-papuloso": 0.926,  
     "esantema-morbilliforme": 0.958,  
     "esantema-polimorfo-like": 0.812,  
     "esantema-virale": 0.892,  
     "esantema_iatrogeno_farmaco_indotta": 0.941,  
     "orticaria": 0.893,  
     "pediculosi": 0.895,  
     "scabbia": 0.933,  
     "varicella": 0.894  
   }},  
  
9: {"coverage": 0.899, "avg_size": 0.900,  
   "class_cov": {  
     "esantema-maculo-papuloso": 0.874,  
     "esantema-morbilliforme": 0.921,  
     "esantema-polimorfo-like": 0.855,  
     "esantema-virale": 0.900,  
     "esantema_iatrogeno_farmaco_indotta": 0.942,  
     "orticaria": 0.896,  
     "pediculosi": 0.915,  
     "scabbia": 0.900,  
     "varicella": 0.859  
   }},  
  
10: {"coverage": 0.900, "avg_size": 0.901,  
     "class_cov": {  
       "esantema-maculo-papuloso": 0.860,  
       "esantema-morbilliforme": 0.964,  
       "esantema-polimorfo-like": 0.861,
```

```
        "esantema-virale": 0.903,  
        "esantema_iatrogeno_farmaco_indotta": 0.918,  
        "orticaria": 0.903,  
        "pediculosi": 0.865,  
        "scabbia": 0.898,  
        "varicella": 0.877  
    }},  
  
11: {"coverage": 0.899, "avg_size": 0.901,  
    "class_cov": {  
        "esantema-maculo-papuloso": 0.925,  
        "esantema-morbilliforme": 0.950,  
        "esantema-polimorfo-like": 0.882,  
        "esantema-virale": 0.877,  
        "esantema_iatrogeno_farmaco_indotta": 0.951,  
        "orticaria": 0.891,  
        "pediculosi": 0.919,  
        "scabbia": 0.939,  
        "varicella": 0.871  
    }},  
  
12: {"coverage": 0.900, "avg_size": 0.901,  
    "class_cov": {  
        "esantema-maculo-papuloso": 0.881,  
        "esantema-morbilliforme": 0.939,  
        "esantema-polimorfo-like": 0.804,  
        "esantema-virale": 0.907,  
        "esantema_iatrogeno_farmaco_indotta": 0.938,  
        "orticaria": 0.895,  
        "pediculosi": 0.905,  
        "scabbia": 0.927,  
        "varicella": 0.827  
    }},  
  
13: {"coverage": 0.900, "avg_size": 0.901,  
    "class_cov": {  
        "esantema-maculo-papuloso": 0.869,  
        "esantema-morbilliforme": 0.946,  
        "esantema-polimorfo-like": 0.873,  
        "esantema-virale": 0.882,  
        "esantema_iatrogeno_farmaco_indotta": 0.939,  
        "orticaria": 0.906,  
        "pediculosi": 0.937,  
        "scabbia": 0.897,  
        "varicella": 0.869  
    }},  
  
14: {"coverage": 0.900, "avg_size": 0.901,  
    "class_cov": {  
        "esantema-maculo-papuloso": 0.891,  
        "esantema-morbilliforme": 0.970,  
        "esantema-polimorfo-like": 0.867,  
        "esantema-virale": 0.899,  
        "esantema_iatrogeno_farmaco_indotta": 0.943,  
        "orticaria": 0.891,  
        "pediculosi": 0.904,
```



```
      "scabbia": 0.913,  
      "varicella": 0.863  
    }},  
  
15: {"coverage": 0.902, "avg_size": 0.904,  
     "class_cov": {  
       "esantema-maculo-papuloso": 0.914,  
       "esantema-morbilliforme": 0.955,  
       "esantema-polimorfo-like": 0.862,  
       "esantema-virale": 0.883,  
       "esantema_iatrogeno_farmaco_indotta": 0.935,  
       "orticaria": 0.916,  
       "pediculosi": 0.851,  
       "scabbia": 0.898,  
       "varicella": 0.865  
     }},  
  
16: {"coverage": 0.899, "avg_size": 0.900,  
     "class_cov": {  
       "esantema-maculo-papuloso": 0.908,  
       "esantema-morbilliforme": 0.926,  
       "esantema-polimorfo-like": 0.897,  
       "esantema-virale": 0.913,  
       "esantema_iatrogeno_farmaco_indotta": 0.954,  
       "orticaria": 0.890,  
       "pediculosi": 0.740,  
       "scabbia": 0.909,  
       "varicella": 0.852  
     }},  
  
17: {"coverage": 0.899, "avg_size": 0.900,  
     "class_cov": {  
       "esantema-maculo-papuloso": 0.857,  
       "esantema-morbilliforme": 0.916,  
       "esantema-polimorfo-like": 0.864,  
       "esantema-virale": 0.885,  
       "esantema_iatrogeno_farmaco_indotta": 0.948,  
       "orticaria": 0.909,  
       "pediculosi": 0.909,  
       "scabbia": 0.907,  
       "varicella": 0.774  
     }},  
  
18: {"coverage": 0.897, "avg_size": 0.899,  
     "class_cov": {  
       "esantema-maculo-papuloso": 0.871,  
       "esantema-morbilliforme": 0.950,  
       "esantema-polimorfo-like": 0.863,  
       "esantema-virale": 0.882,  
       "esantema_iatrogeno_farmaco_indotta": 0.923,  
       "orticaria": 0.910,  
       "pediculosi": 0.845,  
       "scabbia": 0.900,  
       "varicella": 0.864  
     }},
```

```

19: {"coverage": 0.898, "avg_size": 0.900,
    "class_cov": {
        "esantema-maculo-papuloso": 0.869,
        "esantema-morbilloforme": 0.925,
        "esantema-polimorfo-like": 0.884,
        "esantema-virale": 0.877,
        "esantema_iatrogeno_farmaco_indotta": 0.927,
        "orticaria": 0.927,
        "pediculosi": 0.875,
        "scabbia": 0.843,
        "varicella": 0.839
    }},

20: {"coverage": 0.898, "avg_size": 0.899,
    "class_cov": {
        "esantema-maculo-papuloso": 0.893,
        "esantema-morbilloforme": 0.934,
        "esantema-polimorfo-like": 0.902,
        "esantema-virale": 0.899,
        "esantema_iatrogeno_farmaco_indotta": 0.930,
        "orticaria": 0.893,
        "pediculosi": 0.913,
        "scabbia": 0.891,
        "varicella": 0.861
    }},

}

```

```

In [10]: import numpy as np
import matplotlib.pyplot as plt

```

```

In [ ]: # disease_name_map = {
#         "esantema-maculo-papuloso": "Maculopapular exanthema",
#         "esantema-morbilloforme": "Morbilloform exanthema",
#         "esantema-polimorfo-like": "Polymorphous exanthema",
#         "esantema-virale": "Viral exanthema",
#         "esantema_iatrogeno_farmaco_indotta": "Iatrogenic drug-induced exanthema",
#         "orticaria": "Urticaria",
#         "pediculosi": "Pediculosis",
#         "scabbia": "Scabies",
#         "varicella": "Chickenpox",
#     }

# def rename_class_names(results_dict, name_map):
#     new_results = {}
#     for k, v in results_dict.items():
#         new_class_cov = {
#             name_map.get(class_name, class_name): value
#             for class_name, value in v["class_cov"].items()
#         }
#         new_results[k] = {
#             "coverage": v["coverage"],
#             "avg_size": v["avg_size"],
#             "class_cov": new_class_cov
#         }

```

```
#     return new_results

# resnet_results_en = rename_class_names(resnet_results, disease_name_map)
# vit_results_en = rename_class_names(vit_results, disease_name_map)
```

```
In [21]: disease_name_map = {
    "esantema-maculo-papuloso": "Maculopapular exanthema",
    "esantema-morbilliforme": "Morbilliform exanthema",
    "esantema-polimorfo-like": "Polymorphous exanthema",
    "esantema-virale": "Viral exanthema",
    "esantema_iatrogeno_farmaco_indotta": "Iatrogenic drug-induced exanthema",
    "orticaria": "Urticaria",
    "pediculosi": "Pediculosis",
    "scabbia": "Scabies",
    "varicella": "Chickenpox",
}

def rename_class_names_inplace(results_dict, name_map):
    for _, v in results_dict.items():
        class_cov = v["class_cov"]

        for old_name in list(class_cov.keys()):
            if old_name in name_map:
                class_cov[name_map[old_name]] = class_cov.pop(old_name)

rename_class_names_inplace(resnet_results, disease_name_map)
rename_class_names_inplace(vit_results, disease_name_map)
```

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```
In [22]: import numpy as np

def avg_metric(results_dict, metric):
    return np.mean([results_dict[r][metric] for r in results_dict])

avg_resnet_coverage = avg_metric(resnet_results, "coverage")
avg_resnet_size     = avg_metric(resnet_results, "avg_size")

avg_vit_coverage = avg_metric(vit_results, "coverage")
```

```

avg_vit_size      = avg_metric(vit_results, "avg_size")

print("ResNet  AVG Coverage:", round(avg_resnet_coverage,3))
print("ResNet  AVG Set Size:", round(avg_resnet_size,3))
print("ViT     AVG Coverage:", round(avg_vit_coverage,3))
print("ViT     AVG Set Size:", round(avg_vit_size,3))

```

```

ResNet  AVG Coverage: 0.9
ResNet  AVG Set Size: 0.905
ViT     AVG Coverage: 0.9
ViT     AVG Set Size: 0.901

```

```

In [23]: # ALL class names
class_names = list(resnet_results[9]["class_cov"].keys())

def avg_class_cov(results_dict):
    class_covs = {cls: [] for cls in class_names}
    for r in results_dict:
        for cls in class_names:
            class_covs[cls].append(results_dict[r]["class_cov"][cls])
    return {cls: np.mean(class_covs[cls]) for cls in class_names}

resnet_class_avg = avg_class_cov(resnet_results)
vit_class_avg    = avg_class_cov(vit_results)

print("\nAvg class-wise coverage (ResNet):")
print(resnet_class_avg)
print("\nAvg class-wise coverage (ViT):")
print(vit_class_avg)

```

Avg class-wise coverage (ResNet):

```

{'Maculopapular exanthema': np.float64(0.8660500000000001), 'Morbilliform exanthema': np.float64(0.93635), 'Polymorphous exanthema': np.float64(0.8471), 'Viral exanthema': np.float64(0.8854), 'Iatrogenic drug-induced exanthema': np.float64(0.9285), 'Urticaria': np.float64(0.9107500000000002), 'Pediculosis': np.float64(0.9189500000000003), 'Scabies': np.float64(0.9094000000000001), 'Chickenpox': np.float64(0.8318)}

```

Avg class-wise coverage (ViT):

```

{'Maculopapular exanthema': np.float64(0.88375), 'Morbilliform exanthema': np.float64(0.9393500000000001), 'Polymorphous exanthema': np.float64(0.8566), 'Viral exanthema': np.float64(0.8886499999999999), 'Iatrogenic drug-induced exanthema': np.float64(0.9378), 'Urticaria': np.float64(0.9063000000000001), 'Pediculosis': np.float64(0.89085), 'Scabies': np.float64(0.89865), 'Chickenpox': np.float64(0.8544499999999999)}

```

```

In [24]: x = np.arange(len(class_names))
width = 0.38
target = 0.90

resnet_vals = [resnet_class_avg[c] for c in class_names]
vit_vals    = [vit_class_avg.get(c, np.nan) for c in class_names]

plt.figure(figsize=(14,6))

resnet_colors = ["cornflowerblue" if v >= target else "royalblue" for v in resnet_vals]
vit_colors    = ["lightcoral" if v >= target else "firebrick" for v in vit_vals]

bars_res = plt.bar(x - width/2, resnet_vals, width, label="ResNet", color=resnet_co

```

```

bars_vit = plt.bar(x + width/2, vit_vals, width, label="ViT", color=vit_color)

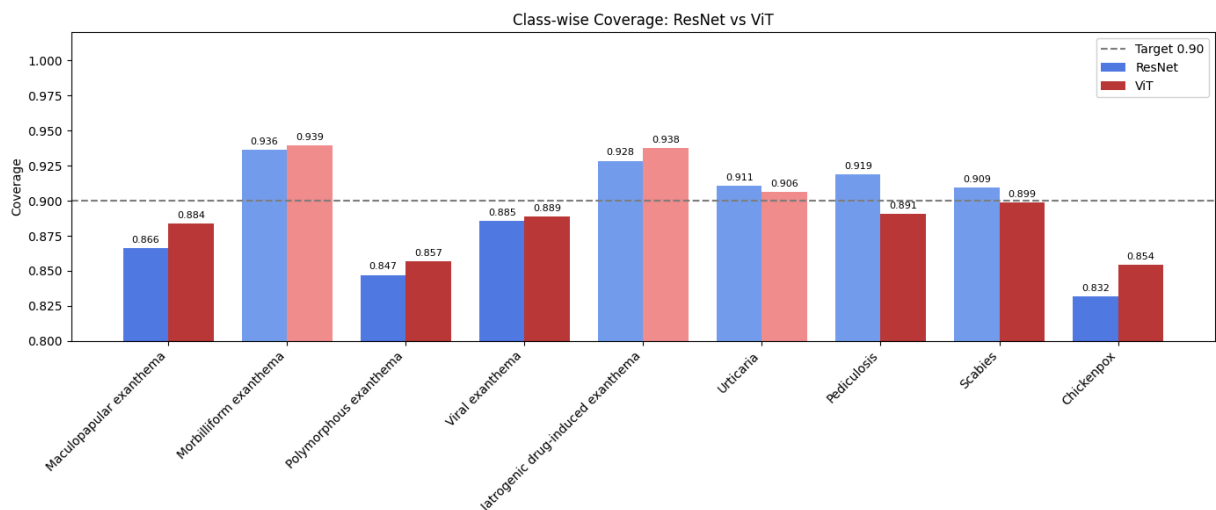
plt.axhline(target, linestyle="--", color="gray", label=f"Target {target:.2f}")

for bar, val in zip(bars_res, resnet_vals):
    height = bar.get_height()
    plt.annotate(f"{val:.3f}",
                 xy=(bar.get_x() + bar.get_width()/2, height),
                 xytext=(0, 3),
                 textcoords="offset points",
                 ha="center", va="bottom", fontsize=8)

for bar, val in zip(bars_vit, vit_vals):
    height = bar.get_height()
    plt.annotate(f"{val:.3f}",
                 xy=(bar.get_x() + bar.get_width()/2, height),
                 xytext=(0, 3),
                 textcoords="offset points",
                 ha="center", va="bottom", fontsize=8)

plt.xticks(x, class_names, rotation=45, ha="right")
plt.ylabel("Coverage")
plt.title("Class-wise Coverage: ResNet vs ViT")
plt.ylim(0.80, 1.02)
plt.legend()
plt.tight_layout()
plt.show()

```



```

In [25]: import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
from typing import Optional, Tuple

def plot_avg_coverage_at_nominal(
    df: pd.DataFrame,
    nominal: float = 0.90,
    model_order: Tuple[str, str] = ("ResNet50", "ViT"),

```

```

model_labels: Tuple[str, str] = ("ResNet-50", "Vision Transformer (ViT)"),
dpi: int = 300,
out_path: Optional[str] = None,
):
    required = {"model", "run", "target_coverage", "coverage"}
    if not required.issubset(df.columns):
        raise ValueError(f"df must contain columns: {required}")

    t = round(float(nominal), 2)

    d = df.copy()
    d["target_coverage"] = d["target_coverage"].astype(float).round(2)
    d = d[d["target_coverage"] == t].copy()

    if d.empty:
        raise ValueError(
            "No rows found for target_coverage={}. Available targets: {}".format(
                t, sorted(df["target_coverage"].astype(float).round(2).unique())
            )
        )

    agg = (
        d.groupby("model")["coverage"]
        .agg(["mean", "std", "count"])
        .reindex(list(model_order))
    )
    agg["sem"] = agg["std"] / np.sqrt(agg["count"])

    means = agg["mean"].values

    plt.figure(figsize=(6.2, 4.6), dpi=dpi)

    bars = plt.bar(
        list(model_labels),
        means,
        color=["#4C72B0", "#DD8452"],
        alpha=0.85
    )

    for x in range(len(means)):
        plt.axvline(
            x=x,
            color="gray",
            linewidth=2.0,
            alpha=0.35,
            zorder=0
        )

    plt.axhline(
        y=t,
        linestyle="--",
        linewidth=1.8,
        color="black",

```

```

        alpha=0.7,
        zorder=5
    )

    for bar in bars:
        h = bar.get_height()
        plt.annotate(
            "{:.4f}".format(h),
            xy=(bar.get_x() + bar.get_width() / 2, h),
            xytext=(0, 6),
            textcoords="offset points",
            ha="center",
            va="bottom",
            fontsize=9
        )

    plt.ylabel("Empirical coverage")
    plt.title("Average Empirical Coverage at Nominal = {:.2f} ( $\alpha$  = {:.2f})"
              .format(t, 1.0 - t),
              fontweight="normal")

    ymin = float(np.min(means)) - 0.002
    ymax = max(float(np.max(means)) + 0.002, t + 0.002)
    plt.ylim(ymin, ymax)

    plt.grid(True, axis="y", linewidth=0.6, alpha=0.25)
    plt.tight_layout()

    if out_path is not None:
        plt.savefig(out_path, dpi=dpi, bbox_inches="tight")

    plt.show()
    return agg

```

```

In [26]: import pandas as pd

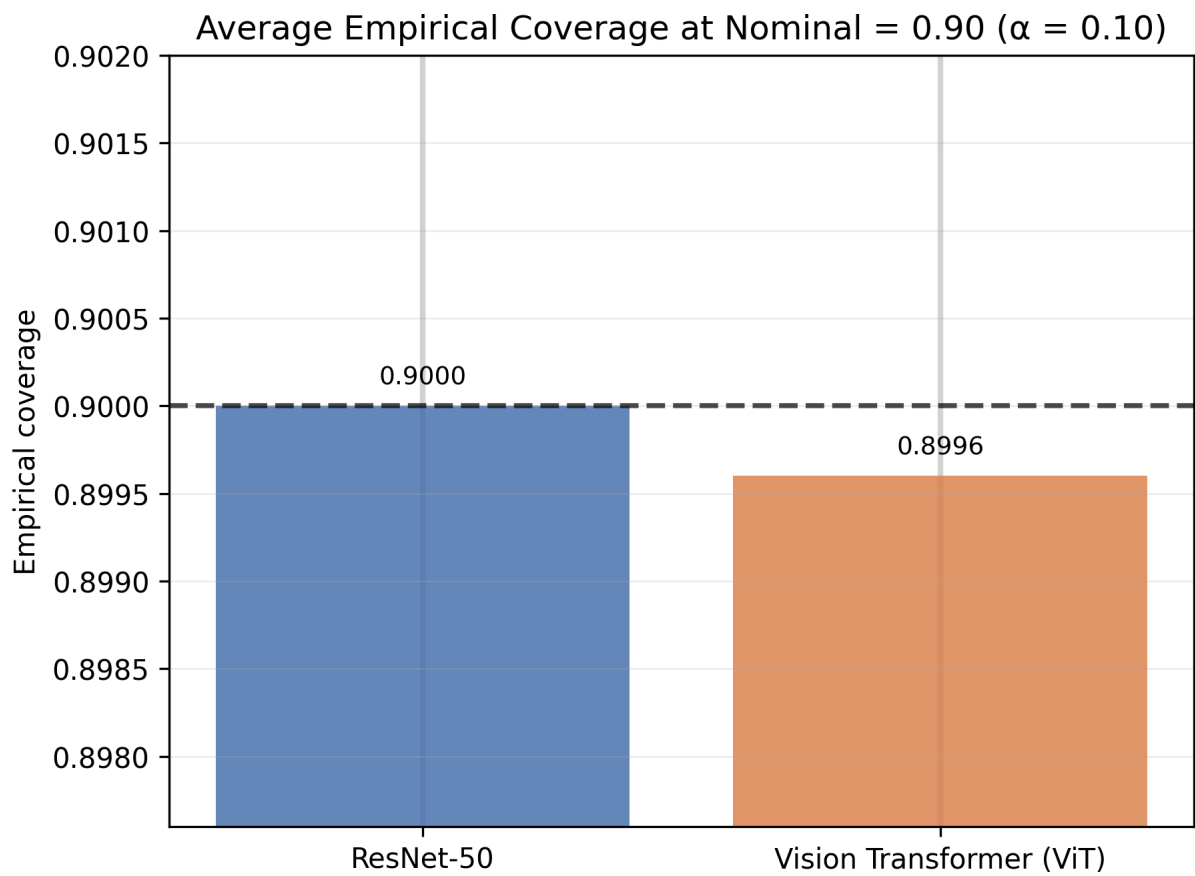
rows = []

for run, v in resnet_results.items():
    rows.append({
        "model": "ResNet50",
        "run": run,
        "target_coverage": 0.90,
        "coverage": v["coverage"]
    })

for run, v in vit_results.items():
    rows.append({
        "model": "ViT",
        "run": run,
        "target_coverage": 0.90,
        "coverage": v["coverage"]
    })

```

```
df_coverage = pd.DataFrame(rows)
plot_avg_coverage_at_nominal(
    df_coverage,
    nominal=0.90,
    out_path="avg_coverage_nominal_0.90.png"
)
```



Out[26]:

	mean	std	count	sem
model				
ResNet50	0.9000	0.001026	20	0.000229
ViT	0.8996	0.001188	20	0.000266

In []:

```
In [27]: import numpy as np
import matplotlib.pyplot as plt

def plot_classwise_coverage_comparison(
    class_names,
    resnet_class_avg,
    vit_class_avg,
    target: float = 0.90,

    figsize=(14, 6),
    dpi=300,
```



```

class_fontsize: int = 10,          # x-tick labels
y_label_fontsize: int = 12,
title_fontsize: int = 14,
legend_fontsize: int = 10,
annotate_fontsize: int = 9,

bar_width: float = 0.38,
resnet_color_pass: str = "cornflowerblue",
resnet_color_fail: str = "royalblue",
vit_color_pass: str = "lightcoral",
vit_color_fail: str = "firebrick",
bar_alpha: float = 0.9,

target_line_style: str = "--",
target_line_color: str = "gray",
target_line_width: float = 1.2,

grid: bool = True,
grid_axis: str = "y",
grid_linewidth: float = 0.6,
grid_alpha: float = 0.35,

ylim: tuple = (0.80, None),
headroom_frac: float = 0.06,
):
    x = np.arange(len(class_names))

    resnet_vals = [resnet_class_avg[c] for c in class_names]
    vit_vals     = [vit_class_avg.get(c, np.nan) for c in class_names]

    plt.figure(figsize=figsize, dpi=dpi)

    resnet_colors = [
        resnet_color_pass if v >= target else resnet_color_fail
        for v in resnet_vals
    ]
    vit_colors = [
        vit_color_pass if v >= target else vit_color_fail
        for v in vit_vals
    ]

    bars_res = plt.bar(
        x - bar_width / 2,
        resnet_vals,
        bar_width,
        label="ResNet",
        color=resnet_colors,
        alpha=bar_alpha,

```

```

)

bars_vit = plt.bar(
    x + bar_width / 2,
    vit_vals,
    bar_width,
    label="ViT",
    color=vit_colors,
    alpha=bar_alpha,
)

y_max = np.nanmax(resnet_vals + vit_vals)
plt.ylim(
    bottom=ylim[0],
    top=y_max * (1.0 + headroom_frac) if ylim[1] is None else ylim[1],
)

plt.axhline(
    target,
    linestyle=target_line_style,
    color=target_line_color,
    linewidth=target_line_width,
    label=f"Target {target:.2f}",
)

def _annotate(bars, values):
    for bar, val in zip(bars, values):
        if np.isnan(val):
            continue
        h = bar.get_height()
        plt.annotate(
            f"{val:.3f}",
            xy=(bar.get_x() + bar.get_width() / 2, h),
            xytext=(0, 3),
            textcoords="offset points",
            ha="center",
            va="bottom",
            fontsize=annotate_fontsize,
        )

_annotate(bars_res, resnet_vals)
_annotate(bars_vit, vit_vals)

plt.xticks(x, class_names, rotation=45, ha="right", fontsize=class_fontsize)
plt.ylabel("Coverage", fontsize=y_label_fontsize)
plt.title("Class-wise Coverage: ResNet vs ViT", fontsize=title_fontsize)

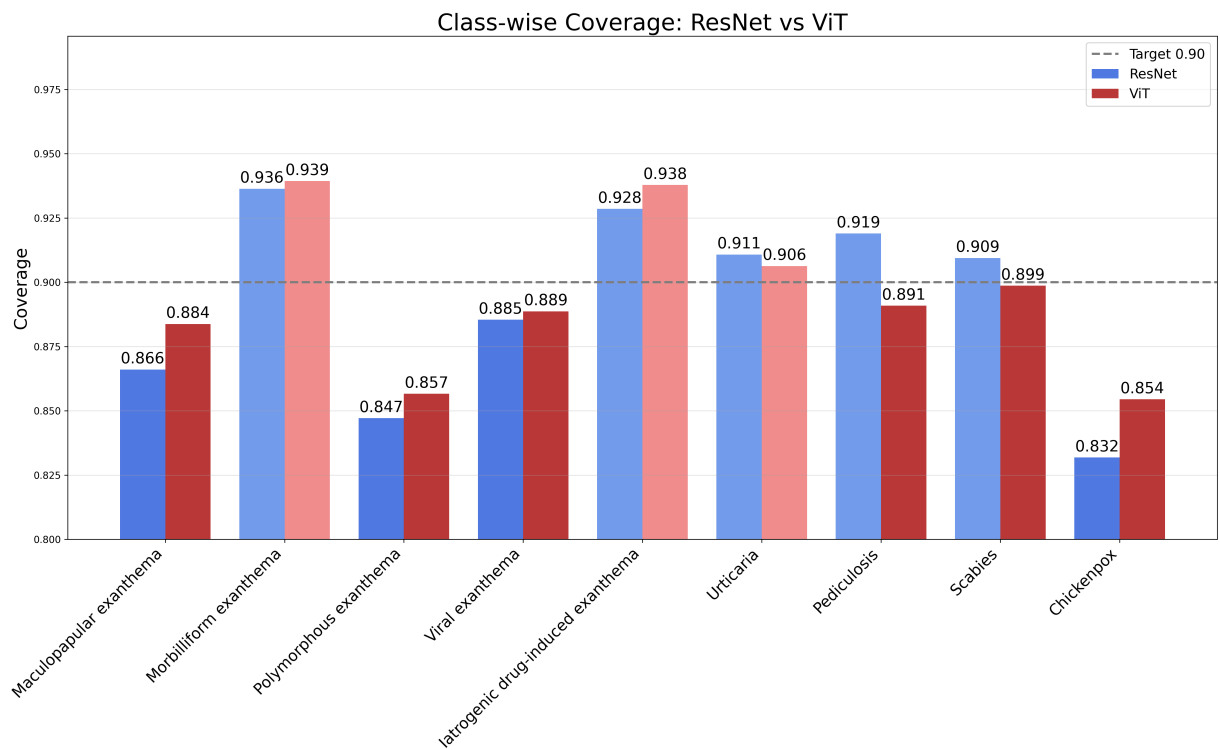
if grid:
    plt.grid(True, axis=grid_axis, linewidth=grid_linewidth, alpha=grid_alpha)

plt.legend(fontsize=legend_fontsize)

```

```
plt.tight_layout()
plt.show()
```

```
In [30]: plot_classwise_coverage_comparison(
    class_names,
    resnet_class_avg,
    vit_class_avg,
    target=0.90,
    figsize=(18, 11),
    class_fontsize=16,
    annotate_fontsize=16,
    title_fontsize=24,
    legend_fontsize=14,
    y_label_fontsize=18,
    grid=True,
    grid_linewidth=0.8,
    grid_alpha=0.35,
    target_line_width=2.3,
)
```



In []:

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