# Temperature Profile Classification - 2 Class system - Full DataSet Load

GMM classification of Southern Ocean Argo float temperature profile data. This notebook uses a previously created model, PCA and sample data.

# Dask import

```
In [1]:
        #Dask server setup cell
        target version='0.19.0'
        !pip install xarray=={target version} --upgrade #--upgrade
        import logging
        import subprocess
        from dask.distributed import Client
        from dask gateway import Gateway
        from distributed import WorkerPlugin
        import dask
        dask.config.set({"array.slicing.split large chunks": True})
        class PipPlugin(WorkerPlugin):
            Install packages on a worker as it starts up.
            Parameters
            packages : List[str]
                A list of packages to install with pip on startup.
            def init (self, packages):
                self.packages = packages
            def setup(self, worker):
                logger = logging.getLogger("distributed.worker")
                subprocess.call(['python', '-m', 'pip', 'install', '--upgrade'] + self.packages)
                logger.info("Installed %s", self.packages)
        def check():
            import xarray
            return xarray. version
        gateway = Gateway()
        cluster = gateway.new cluster(worker memory=8)
        cluster.scale(20)
        client = Client(cluster)
        client
        plugin = PipPlugin([f'xarray=={target version}'])
        client.register worker plugin(plugin)
        client.run(check)
        1.1.1
        blank=1
```

## Choices for data

```
In [3]:
        #Experiment data for analysis
        dataVariableId = 'thetao'
        dataExperimentId = 'historical'
        dataSourceId = 'UKESM1-0-LL'
        dataInstitutionId = 'MOHC'
        approvedIds = ["r2i1p1f2"] #insert start of approved member ids
        #File imports
        maskName = "OceanMaskVolcello"
        modelName = "GMM UK 2Class R2"
        #Data definitions
        startDate = '1980-01'
        endDate = '2009-12'
        timeRange = slice(startDate, endDate)
        levSel = slice(0, 2000) #Selected levels to be investigated
        maxLat = -30 #Selected latitude to be investigated
        runIdSel = 0
        maskEnable = False #Decides if training data mask is applied, or if full data set is class
```

## Libaries and Modules

Importing the necessary libaries and modules for the notebook.

```
In [4]:
         #Import cell
        import calendar
         #import cartopy.crs as ccrs
         #import cartopy.feature as cfeature
        import dask.dataframe as dd
        import fsspec
        import matplotlib.dates as mdates
        import matplotlib as mpl ###
        import numpy as np
        import matplotlib.pyplot as plt
        import pandas as pd
        import pickle as pk
        import matplotlib.ticker as ticker
        import xarray as xr
        import zarr
        from dask import config
        from dask import delayed
        from joblib import dump, load
        from matplotlib.pyplot import cm
        from sklearn import mixture
        from sklearn.decomposition import PCA
        from sklearn import preprocessing
        config.set(**{'array.slicing.split large chunks': True})
        print("Imports complete")
```

Imports complete

## Importing data sets

Importing the data for the models.

#### Import sample data set and corresponding time/geo data

dfESMLevT = dfESMLev.sel(time=timeRange)

```
In [5]:
         #Importing UK ESM data cell
         #Selecting data tables
         df = pd.read csv('https://storage.googleapis.com/cmip6/cmip6-zarr-consolidated-stores.csv')
         dfFilt = df[df.variable id.eq(dataVariableId) & df.experiment id.eq(dataExperimentId) & df
         memberArr = np.empty(shape=(0), dtype=bool)
         for i in dfFilt["member id"]:
             rowSel = i[:] in approvedIds #adapt i[:] to match size of approvedIds
             memberArr = np.append(memberArr, rowSel)
         memberSer = pd.Series(memberArr, name='bools')
         dfFilt = dfFilt[memberSer.values]
         #Opening and counting number of tables
         fileSetList = []
         for i in range(len(dfFilt)):
             zstore = dfFilt.zstore.values[i]
             mapper = fsspec.get mapper(zstore)
             fileRaw = xr.open zarr(mapper, consolidated=True)
             fileSetList.append(fileRaw)
         fileCount = len(fileSetList)
         if fileCount:
             print(str(fileCount)+" "+dataSourceId+" data sets opened")
             print("No UKESM data sets opened")
         #Formatting dates into np.datetime64 format
         for i in range(fileCount):
             startDateIterate = np.datetime64(fileSetList[i]['time'].values[0],'M')
             endDateIterate = np.datetime64(fileSetList[i]['time'].values[-1],'M') + np.timedelta64
             fileSetList[i]['time']=('time', np.arange(startDateIterate, endDateIterate, dtype='dat
             fileSetList[i]['time bnds']=('time bnds', np.arange(startDateIterate, endDateIterate,
         fileSet = xr.combine nested(fileSetList, concat dim='RunId') #Combining data sets
         dataRaw = fileSet.thetao
         try: #Adjusting array names
             dataRaw = dataRaw.rename({"latitude":"lat", "longitude":"lon"})
         except:
             pass
         print ("Data sets successfully merged and renamed into dataRaw. Data dimensions are "+str(
         #dataRaw
        1 UKESM1-0-LL data sets opened
        Data sets successfully merged and renamed into dataRaw. Data dimensions are Frozen({'RunI
        d': 1, 'time': 1980, 'lev': 75, 'j': 330, 'i': 360}).
In [6]:
         dfFilt
Out[6]:
               activity_id institution_id source_id experiment_id member_id table_id variable_id grid_label
                                                                                             gs://cmip6/
                                    UKESM1-
        214481
                              MOHC
                   CMIP
                                                 historical
                                                            r2i1p1f2
                                                                    Omon
                                                                              thetao
                                        0-LL
In [7]:
         #UK ESM raw processing cell
         dfESMLev = dataRaw.sel(lev=levSel) #Selects level data down to 2k
```

```
dfESMLatLevT = dfESMLevT.where(dfESMLevT.lat < maxLat, drop=True) #Selection of latitude
dfESMLatLevT = dfESMLatLevT.squeeze()
#dfESMLatLevT = dfESMLatLevT.reset_coords(drop=True) #Removes lev if single value

globalStartDate = dfESMLatLevT["time"][0].values
globalDateInc = dfESMLatLevT["time"][1].values - globalStartDate
#np.datetime64(globalDateInc,'D')
globalEndDateIn = dfESMLatLevT["time"][-1].values
globalEndDateOut = globalEndDateIn + globalDateInc

globalStartDateStr = str(globalStartDate)[:7]
globalEndDateOutStr = str(globalEndDateIn)[:7]
print("UKESM data loaded and stored in dfESMLatLevT. Data dimensions are "+str(dfESMLatLev
#dfESMLatLevT #Uncomment to see data set</pre>
```

UKESM data loaded and stored in dfESMLatLevT. Data dimensions are Frozen({'time': 360, 'le v': 54, 'j': 139, 'i': 360}).

#### **Loading ocean Masks**

```
In [8]: #Ocean mask import cell
    maskFile = xr.open_dataset(maskName)
    oceanMask = maskFile.to_array()
    maskFile = xr.open_dataset("OceanMaskUKESM1")
    oceanMask2 = maskFile.to_array()
    print("Mask Loaded and stored in oceanMask and oceanMask2 (volcello and UKESM).")
```

Mask Loaded and stored in oceanMask and oceanMask2 (volcello and UKESM).

#### **Unpacking ocean masks**

```
In [9]: #Mask unpacking cell
    geoRange = oceanMask #copying mask
    geoRange = geoRange.rename({"variable":"cleanMe"}) #Dimension removal
    geoRange = geoRange.sel(cleanMe = geoRange.cleanMe.values[0]) #Dimension removal
    geoRange = geoRange.reset_coords("cleanMe", drop=True) #Dimension removal
    geoRangeS = geoRange.stack(ij = ("i", "j")) #Stacking
    geoRangeFilt = geoRangeS.dropna("ij")
    print("Ocean mask unpacked into geoRangeFilt.")

geoRange2 = oceanMask2 #copying mask
    geoRange2S = geoRange2.stack(ij = ("i", "j")) #Stacking
    geoRangeFilt2 = geoRange2s.dropna("ij")
    print("UKESM Ocean mask unpacked into geoRangeFilt2.")
```

Ocean mask unpacked into geoRangeFilt. UKESM Ocean mask unpacked into geoRangeFilt2.

#### **Date Calculations**

```
In [10]: #Date calculation cell
    startDateNp = np.datetime64(startDate, 'M')
    endDateNp = np.datetime64(endDate, 'M')
    timeDiff = endDateNp - startDateNp
    timeDiff = timeDiff.astype(int) + 1
    print("Calculated date range.")
```

Calculated date range.

### **Calculation functions**

#### **Functions:**

- pickRand Takes in data frame and returns sampled data frame with a randomly selected number of rows from the input data frame, controlled by the second input variable to the function.
- storeMeta Returns a np array containing the latitude and longitude data for an input xarray and associated ij.
- loadModel loadeds and returns GMM model named in input.
- saveModel saves input GMM model to provided name, if no name provided default is GMMGenerated.

```
In [11]:
         #Calculation functions cell
         def pickRand(dataArray, sampleFactor):
             '''Returns a sample of the input array, size of sampled array is based on sampleFactor
             arrLen = len(dataArray)
             if sampleFactor > 1:
                 sampleSize = int(sampleFactor)
             elif sampleFactor > 0:
                 sampleSize = int(sampleFactor*arrLen)
             else:
                 return 1
             filtArr = np.zeros(arrLen, dtype=bool) # empty mask
             sampleId = np.random.choice(arrLen, sampleSize, False) # np array of randomly generate
             for i in sampleId:
                 filtArr[i] = True # populating mask
             return dataArray[filtArr] # applies mask
         def pickRandMask(maskLen, maskQuantity, sampleFactor):
             '''Returns a linear mask for the input dimensions, size of mask is based on sampleFact
             if sampleFactor > 1:
                 sampleSize = int(sampleFactor)
             elif sampleFactor > 0:
                 sampleSize = int(sampleFactor*maskLen)
             else:
                 return 1
             globalArr = np.empty(shape=(0), dtype=bool)
             for i in range(maskQuantity):
                 filtArr = np.zeros(maskLen, dtype=bool) # empty mask
                 sampleId = np.random.choice(maskLen, sampleSize, False) # np array of randomly gen
                 for j in sampleId:
                     filtArr[j] = True # populating mask
                 globalArr = np.append(globalArr, filtArr)
             return qlobalArr
         def storeMeta(dataArray):
             "''Returns a np array containing the latitude and longitude data for the input xarray
             storeLen = len(dataArray["lat"]) # assumes each lat has a corresponding lon
             storage = np.empty(shape=(0, storeLen))
             storage = np.append(storage, [dataArray["lat"].values], axis = 0)
             storage = np.append(storage, [dataArray["lon"].values], axis = 0)
             #storage = np.append(storage, [dataArray["time"].values], axis = 0)
             #storage = np.append(storage, [dataArray["ij"].values], axis = 0)
             return storage
```

```
def loadModel(modelName:str):
    '''Loades the input GMM model named in the functions input. Returns loaded model.'''
   means = np.load(modelName + '_means.npy')
covar = np.load(modelName + '_covariances.npy')
    GMModel = mixture.GaussianMixture(n components = len(means), covariance type='full')
    GMModel.precisions cholesky = np.linalg.cholesky(np.linalg.inv(covar))
    GMModel.weights = np.load(modelName + ' weights.npy')
    GMModel.means = means
    GMModel.covariances = covar
    return GMModel
def saveModel(GMModel, modelName = "GMMGenerated"):
    '''Saves the input GMM model's weights, means and covariances. Assigns input name if
    GMModel name = str(modelName)
    np.save(modelName + ' weights', GMModel.weights , allow pickle=False)
    np.save(modelName + ' means', GMModel.means , allow pickle=False)
    np.save(modelName + ' covariances', GMModel.covariances , allow pickle=False)
    return 0
print("Calculation functions defined.")
```

Calculation functions defined.

# **Plotting functions**

#### **Functions:**

- bicPlot Plots BIC score array against component number.
- locationPlotGroup plots location and classification of data points for an input numpy array.
- locationPlotGroupDF plots location and classification of data points for an input data frame.
- locationPlotGroupDFMonthly plots location and classification of data points for an input data frame in monthly subplots.
- locationPlotTime plots locations of an input data array on a map with a colour scale for time.
- locationPlotUncertaintyDF plots uncertainty in classification on a location plot.
- tempPointPlot Plots the temperature profile of a single point against depth.
- tempGroupPlot Plots the mean/+-1std temperature profiles of all classes in input dataArrays (seperate mean and std).

```
In [12]:
         #Plotting functions Cell
         sampleDepthAxis = dfESMLatLevT["lev"]
         def bicPlot(bicArray, startNo, endNo, skipNo, title, label, plotNo):
             '''Plots input BIC score array'''
             plt.figure(plotNo, figsize=(20, 8))
             plt.style.use("seaborn-darkgrid")
             componentRange = range(startNo, endNo, skipNo)
             plt.plot(componentRange, bicArray, label = str(label))
             bicArrayMax = np.max(bicArray)
             bicArrayMin = np.min(bicArray)
             bicRange = bicArrayMax-bicArrayMin
             if bicRange == 0:
                 bicRange = 20 #provides border 1 if all bic values are identical
             plt.xticks(componentRange)
             plt.xlim([startNo-0.5, endNo+0.5])
             plt.ylim([bicArrayMin-0.05*bicRange, bicArrayMax+0.05*bicRange])
             plt.legend(bbox to anchor=(1.05, 1), loc='upper left')
```

```
plt.xlabel("Number of components")
    plt.ylabel("BIC score")
   plt.title(title)
def locationPlotGroup(metaDataArray, size, plotNo):
    '''Plots locations of numpy arrays with group colour scheme'''
    plt.figure(plotNo, figsize=size)
    ax = plt.axes(projection=ccrs.SouthPolarStereo())
    ax.add feature (cfeature.OCEAN)
    ax.add feature(cfeature.COASTLINE)
    ax.coastlines()
   ax.gridlines()
   im = ax.scatter(metaDataArray[1], metaDataArray[0], transform=ccrs.PlateCarree(), c =
   cb = plt.colorbar(im)
    plt.plot(np.arange(0,361,1),np.ones(361)*-29.5, transform=ccrs.PlateCarree(), color="E
    plt.title("Grouped Sample Locations ("+str(len(metaDataArray[0]))+")")
def locationPlotGroupDFTime(dataFrame, title, size, plotNo):
    '''Plots locations of data frame points with group colour scheme'''
    plt.figure(plotNo, figsize=size)
    ax = plt.axes(projection=ccrs.SouthPolarStereo())
    ax.add feature(cfeature.OCEAN)
   ax.add feature(cfeature.COASTLINE)
   ax.coastlines()
   ax.gridlines()
   im = ax.scatter(dataFrame["lon"], dataFrame["lat"], transform=ccrs.PlateCarree(), c =
   cb = plt.colorbar(im)
    loc = mdates.AutoDateLocator()
   cb.ax.yaxis.set major locator(loc)
    cb.ax.yaxis.set major formatter(mdates.ConciseDateFormatter(loc))
    plt.plot(np.arange(0,361,1),np.ones(361)*-29.5, transform=ccrs.PlateCarree(), color="E
    plt.title(str(title))
def locationPlotGroupDFLab(dataFrame, title, size, plotNo):
    '''Plots locations of data frame points with group colour scheme'''
    plt.figure(plotNo, figsize=size)
    ax = plt.axes(projection=ccrs.SouthPolarStereo())
    ax.add feature (cfeature.OCEAN)
   ax.add feature(cfeature.COASTLINE)
   ax.coastlines()
   ax.gridlines()
    im = ax.scatter(dataFrame["lon"], dataFrame["lat"], transform=ccrs.PlateCarree(), c =
    cb = plt.colorbar(im)
   plt.plot(np.arange(0,361,1),np.ones(361)*-29.5, transform=ccrs.PlateCarree(), color="E
    plt.title(str(title))
def locationPlotGroupDFMonthly(dataFrame, title, plotNo):
    '''Plots locations of dataframe points by monthly subplot with group colour scheme'''
    fig = plt.figure(plotNo, figsize=(30,42))
   plt.title(str(title))
    for i in range (1, 13):
        timeData = dataFrame.where(dataFrame["time"].dt.month==i)
        ax = plt.subplot(4, 3, i, projection=ccrs.SouthPolarStereo())
        ax.add feature(cfeature.OCEAN)
        ax.add feature(cfeature.COASTLINE)
        ax.coastlines()
        ax.gridlines()
        im = ax.scatter(timeData["lon"], timeData["lat"], transform=ccrs.PlateCarree(), c
        plt.plot(np.arange(0,361,1),np.ones(361)*-29.5, transform=ccrs.PlateCarree(), cold
        plt.title(calendar.month abbr[i])
    plt.subplots adjust(wspace=0, hspace=0.05)
    cb_ax = fig.add_axes([0.27, 0.1, 0.5, 0.02])
```

```
cbar = fig.colorbar(im, cax=cb ax, orientation="horizontal")
def locationPlotTime(dataArray, size, plotNo):
    '''Plots locations of numpy arrays with date colour scheme'''
    plt.figure(plotNo, figsize=size)
    ax = plt.axes(projection=ccrs.SouthPolarStereo())
    ax.add feature(cfeature.OCEAN)
    ax.add feature(cfeature.COASTLINE)
   ax.coastlines()
   ax.gridlines()
    im = ax.scatter(dataArray[1], dataArray[0], transform=ccrs.PlateCarree(), c= mdates.de
   cb = plt.colorbar(im)
   loc = mdates.AutoDateLocator()
    cb.ax.yaxis.set_major locator(loc)
    cb.ax.yaxis.set major formatter(mdates.ConciseDateFormatter(loc))
    plt.plot(np.arange(0,361,1),np.ones(361)*-29.5, transform=ccrs.PlateCarree(), color="E
   plt.title("Sample Locations ("+str(len(dataArray[0]))+")")
def locationPlotUncertaintyDF(dataFrame, title, size, plotNo):
    '''Plots input data array classification uncertainties'''
    plt.figure(plotNo, figsize=size)
    ax = plt.axes(projection=ccrs.SouthPolarStereo())
    ax.add feature(cfeature.OCEAN)
    ax.add feature(cfeature.COASTLINE)
   ax.coastlines()
   ax.gridlines()
   im = ax.scatter(dataFrame["lon"], dataFrame["lat"], transform=ccrs.PlateCarree(), c =
    cb = plt.colorbar(im)
    plt.plot(np.arange(0,361,1),np.ones(361)*-29.5, transform=ccrs.PlateCarree(), color="E
   plt.title(str(title))
def locationPlotUncertaintyDFMonthly(dataFrame, title, plotNo):
    '''Plots locations of dataframe points by monthly subplot with group colour scheme'''
    fig = plt.figure(plotNo, figsize=(30,42))
    plt.title(str(title))
    for i in range(1, 13):
        timeData = dataFrame.where(dataFrame["time"].dt.month==i)
        ax = plt.subplot(4, 3, i, projection=ccrs.SouthPolarStereo())
        ax.add feature(cfeature.OCEAN)
        ax.add feature(cfeature.COASTLINE)
        ax.coastlines()
        ax.gridlines()
        im = ax.scatter(timeData["lon"], timeData["lat"], transform=ccrs.PlateCarree(), c
        #cb = plt.colorbar(im, fraction=0.046, pad=0.04)
        plt.plot(np.arange(0,361,1),np.ones(361)*-29.5, transform=ccrs.PlateCarree(), cold
        plt.title(calendar.month abbr[i])
    plt.subplots adjust(wspace=0, hspace=0.05)
    cb ax = fig.add axes([0.27, 0.1, 0.5, 0.02])
    cbar = fig.colorbar(im, cax=cb ax, orientation="horizontal")
def locationPlotXr(dataArray, size, plotNo):
    '''Plots locations of numpy arrays with date colour scheme'''
    plt.figure(plotNo, figsize=size)
    ax = plt.axes(projection=ccrs.SouthPolarStereo())
    ax.add feature(cfeature.OCEAN)
    ax.add feature(cfeature.COASTLINE)
    ax.coastlines()
    ax.gridlines()
    im = ax.scatter(dataArray["lon"], dataArray["lat"], transform=ccrs.PlateCarree())
    plt.plot(np.arange(0,361,1),np.ones(361)*-29.5, transform=ccrs.PlateCarree(), color="E
    plt.title("Sample Locations ("+str(len(dataArray["lat"]))+")")
```

```
def surfaceTempPlot(dataArray, plotNo):
   plt.figure(plotNo, figsize=(20,20))
    ax = plt.axes(projection=ccrs.SouthPolarStereo())
    ax.add feature(cfeature.OCEAN)
    ax.add feature(cfeature.COASTLINE)
   ax.coastlines()
    ax.gridlines()
   im = ax.scatter(dataArray["lon"], dataArray["lat"], transform=ccrs.PlateCarree(), c =
    cb = plt.colorbar(im)
    plt.plot(np.arange(0,361,1),np.ones(361)*-29.5, transform=ccrs.PlateCarree(), color="E
    plt.title("Surface Temperature of Samples")
def tempPointPlot(dataArray, label, title, plotNo):
    '''Displays temperature profile plot for a given data set, singular point'''
    plt.figure(plotNo)
   plt.plot(dataArray, sampleDepthAxis, label = label)
    plt.legend(bbox to anchor=(1.05, 1), loc='upper left')
   plt.title(str(title))
    plt.gca().invert yaxis()
def tempGroupProfile(dataArrayMean, dataArrayStd, plotNo):
    '''Displays mean /+-1 std temperature profiles for classes in dataArrayMean and dataAi
    dataCompNo = len(dataArrayMean)
    columnNames = sampleDFSortMeans.columns.values
    dataStart = np.where(columnNames == sampleDepthAxis[0].values)[0][0]
    subPlotX = int(np.ceil(dataCompNo/5))
   plt.figure(plotNo, figsize=(35, 10*subPlotX))
   plt.style.use("seaborn-darkgrid")
    palette = cm.coolwarm(np.linspace(0,1, dataCompNo))
    for i in range(dataCompNo):
        meanT = dataArrayMean.iloc[i, dataStart:].to numpy()
        stdT = dataArrayStd.iloc[i, dataStart:].to numpy()
        plt.subplot(subPlotX, 5, i+1)
        plt.plot(meanT, sampleDepthAxis, marker='', linestyle="solid", color=palette[i], ]
        plt.plot(meanT+stdT, sampleDepthAxis, marker='', linestyle="dashed", color=palette
        plt.plot(meanT-stdT, sampleDepthAxis, marker='', linestyle="dashed", color=palette
       plt.xlim([-2,20])
       plt.ylim([0,1000])
        ax = plt.gca()
       ax.invert yaxis()
        ax.grid(True)
        fs = 16 #font size
        plt.xlabel("Temperature (°C)", fontsize=fs)
        plt.ylabel("Depth (m)", fontsize=fs)
        plt.title("Class = "+str(i), fontsize=fs)
        mpl.rc("xtick", labelsize=fs)
        mpl.rc("ytick", labelsize=fs)
        textstr = '\n'.join((
            r'N profs. = %i' % (nprofs[nrow], ),
            r'Mean lon = %i' % (meanLon, ),
           r'Mean lat = %i' % (meanLat, ),
           r'Post. = %i' % (meanMaxPP, )))
        props = dict(boxstyle="round", facecolor="wheat", alpha=0.8)
        ax.text(0.45, 0.25, textstr, transform=ax.transAxes, fontsize=fs, verticalalignment
```

```
print("Plotting functions defined.")
```

Plotting functions defined.

## **Plotting Ocean Mask**

```
In [13]:
#Mask plotting cell
#locationPlotXr(geoRangeFilt, (10,10), 1) #OceanMaskVolcello
#locationPlotXr(geoRangeFilt2, (10,10), 2) #OceanMaskUKESM1
plt.show()
```

## **Generating Data Samples**

#### Identifying, masking and stacking raw data

```
In [14]: #Identifying, masking and stacking raw data cell
    dfESMLatLevTStack = dfESMLatLevT.stack(ij = ("i", "j"))
    dfESMLatLevTStack = dfESMLatLevTStack.transpose('time', 'ij', 'lev')
    dfESMLatLevTStackFilt = dfESMLatLevTStack.sel(ij = geoRangeFilt.ij.values) #Produces 2219
    dfESMLatLevTStackFilt
    print("Raw data identified, stacked and stored in dfESMLatLevTStackFilt. Data dimensions:
        Raw data identified, stacked and stored in dfESMLatLevTStackFilt. Data dimensions: Frozen
        ({'time': 360, 'ij': 22194, 'lev': 54}).
In [15]: #Plotting raw data locations cell
    #locationPlotXr(dfESMLatLevTStackFilt, (10,10), 1)
    plt.show()
```

#### Selecting sample data

```
In [16]: #Mask loading cell
if maskEnable:
    importName = modelName + "_Mask.npy"
    mask = np.load(importName)
    print("Data mask loaded from "+ importName +".")
else:
    print("No mask loaded.")
```

No mask loaded.

Sample data calculated and stored in sampleData. Sample data dimensions: Frozen({'ijT': 79 89840, 'lev': 54}).

#### Placing sample data in tables

```
In [18]:
         #Location and time data to table cell
         metaData = {"lat":sampleData["lat"], "lon":sampleData["lon"], "time":sampleData["time"]}
         sampleMetaDF = pd.DataFrame(metaData, columns=["lat", "lon", "time"])
         print("Sample lat, lon and time converted to datafile (sampleMetaDF). "+str(len(sampleMeta
         sampleMetaDF.head()
```

Sample lat, lon and time converted to datafile (sampleMetaDF). 7989840 samples identified.

#### Out[18]:

	lat	lon	time
0	-66.111519	73.5	1980-01-01
1	-65.703316	73.5	1980-01-01
2	-65.288567	73.5	1980-01-01
3	-64.867195	73.5	1980-01-01

**4** -64.439102 73.5 1980-01-01

#### In [19]:

```
#Temperature data to table and table merging cell
#Generating surface temperature level value and column name
surfaceLev = sampleData["lev"][0].values
surfaceData = sampleData.sel(lev = surfaceLev)
surfaceLevName = "Surface Temp ("+str(np.round(surfaceLev,2))+")"
#Exporting sample data into pandas
if True:
    sampleDataDF = sampleData.to pandas()
    sampleDataDFClean = sampleDataDF.reset index()
    sampleDataDFClean = sampleDataDFClean.drop(columns=['ij'])
    sampleDF = pd.concat([sampleMetaDF, sampleDataDFClean.drop(columns=["time"])], axis=1)
else:
    sampleDF = sampleMetaDF
sampleDF["time"] = pd.to datetime(sampleDF["time"])
print("SampleData converted to datafile (sampleDataDF). Datafiles combined into sampleDF
sampleDF.head()
```

SampleData converted to datafile (sampleDataDF). Datafiles combined into sampleDF.

_		-	_	-	
()	1	17	a	-	0
$\cup$		1 4	ン	-	

out[19]:		lat	lon	time	0.5057600140571594	1.5558552742004395	2.6676816940307617	3.8562798500061035	į
	0	-66.111519	73.5	1980- 01-01	-1.259922	-1.262661	-1.267313	-1.274738	
	1	-65.703316	73.5	1980- 01-01	-1.240721	-1.246196	-1.254534	-1.262789	
	2	-65.288567	73.5	1980- 01-01	-1.109075	-1.123683	-1.134651	-1.145657	
	3	-64.867195	73.5	1980- 01-01	-0.735238	-0.765917	-0.792907	-0.824765	
	4	-64.439102	73.5	1980- 01-01	-0.490287	-0.573119	-0.640231	-0.684716	

# Scaling

#### **Scaling implementation**

Applying scaling to the data set, ensuring all levels have same influence over data.

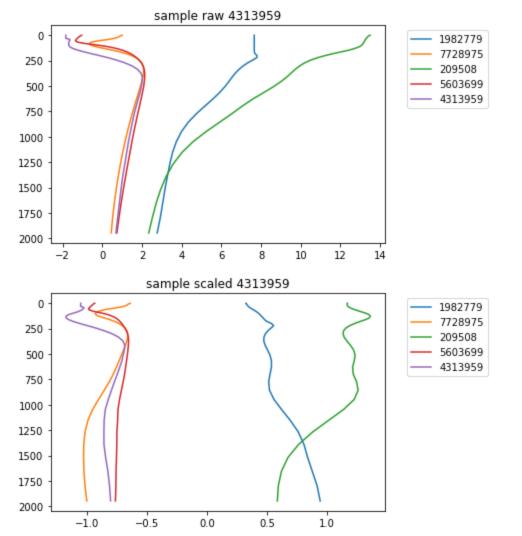
```
In [20]: #Scaler loading and transform cell
    importName = modelName + "_Scaler"
    scalerLoad = load(importName)
    sampleDataScaled = scalerLoad.transform(sampleData)
    print("Scaling of sampleData complete using "+ importName +", stored in sampleDataScaled.")
```

Scaling of sampleData complete using GMM UK 2Class R2 Scaler, stored in sampleDataScaled.

#### **Scaling comparison**

Comparing raw temperature profiles with their scaled equivalent. To show individual plots set solo to True.

```
In [21]: #Scaled temperature profile plotting cell
solo = False #Set to true for seperate plots, false for combined plots.
for i in range(5):
    x = np.random.randint(len(sampleMetaDF))
    tempPointPlot(sampleData[x], x, "sample raw "+str(x), solo*2*i)
    tempPointPlot(sampleDataScaled[x], x, "sample scaled "+str(x), solo*2*i+1)
plt.show()
```



# **Principle Component Analysis**

This process is performed to reduce the number of dimensions of the the data, as well as to improve overall model performance.

```
In [22]: #PCA importing cell
    importName = modelName + "_PCA.pkl"
    pca = pk.load(open(importName, "rb"))
    totalVarianceExplained = np.sum(pca.explained_variance_ratio_)
    print("PCA loaded into pca. Total variance explained by PCA for "+str(pca.n_components)+"

PCA loaded into pca. Total variance explained by PCA for 3 is 0.9919397558439996.

In [23]: #PCA transform cell
    sampleDataScaledPCA = pca.transform(sampleDataScaled) #converting input data into PCA representation print("Data passed through PCA to sampleDataPCA.")
```

## Model import/BIC score calculation

Data passed through PCA to sampleDataPCA.

The previously generated model is imported.

```
In [24]: #Model import and BIC + Component Calculation Cell
   bestGMModel = loadModel(modelName) #Loading model
   bicMin = bestGMModel.bic(sampleDataScaledPCA) #BIC score calculation
   bicComponentMin = bestGMModel.n_components #Identifying number of components in model (2 ;

   print("Model "+modelName+" loaded. The bicScore was "+str(np.round(bicMin, 2))+" for "+str
   print("Imported model "+modelName+" in use. No calculations necessary.")
   print("Imported model "+modelName+" in use. Model BIC score for training data: "+str(bicMinum)

Model GMM_UK_2Class_R2 loaded. The bicScore was 85523397.3 for 2.
   Imported model GMM_UK_2Class_R2 in use. No calculations necessary.
   Imported model GMM_UK_2Class_R2 in use. Model BIC score for training data: 85523397.295375
   32.
```

## Assigning class labels to each profile using the best GMM

Implementation of classification.

pass

```
In [25]: #Classification and classification probability cell
    labels = bestGMModel.predict(sampleDataScaledPCA) #Assignment of class labels from best GI
    posteriorProbs = bestGMModel.predict_proba(sampleDataScaledPCA) #Probability of profile be
    maxPosteriorProbs = np.max(posteriorProbs, axis=1) #Evaluating assigned class probability
    classUncertainty = 2 - 2*maxPosteriorProbs #I factor calculation for 2 class system (reduced try: #Removing labels to sampleDF table cell
    try: #Removing label, maxposteriorprob and classUncertainty columns from sampleDF
    sampleDF = sampleDF.drop(columns=["label", "max posterior prob", "classUncertainty"])
    except:
```

#Adding label, maxposteriorprob and classUncertainty columns to sampleDF

```
sampleDF.insert(3, "label", labels, True)
sampleDF.insert(4, "max posterior prob", maxPosteriorProbs, True)
sampleDF.insert(5, "classUncertainty", classUncertainty, True)
print("Labels identified for model ("+str(bicComponentMin)+" components) and added to same
```

Labels identified for model (2 components) and added to sampleDF with associated probability.

## Calculating class means for sorting

```
In [27]: #Class Mean Calculation Cell
    sampleDFGrouped = sampleDF.groupby("label") #group profiles according to label
    sampleDFMeans = sampleDFGrouped.mean() #calculate mean of all profiles in each class
    print("Sample dataframe grouped by label (sampleDFGrouped) and means taken (sampleDFMeans)
```

Sample dataframe grouped by label (sampleDFGrouped) and means taken (sampleDFMeans).

## Sorting the labels based on mean class temperatures

```
In [28]: #Sorted Dictionary creation cell
    surfaceMeans = sampleDFMeans[surfaceLev].to_numpy() #Takes first temperature data column
    surfaceMeansOrder = np.argsort(surfaceMeans)
    di = dict(zip(surfaceMeansOrder, range(0, bicComponentMin)))
    print("Surface temperature means taken and sorted. Label dictionary created and stored in
```

Surface temperature means taken and sorted. Label dictionary created and stored in di.

```
In [29]: #Sorted label column to tables cell
    try: #Removing labelSorted column from tables
        sampleMetaDF = sampleMetaDF.drop(columns = "labelSorted")
    except:
        pass
    try:
        sampleDF = sampleDF.drop(columns = "labelSorted")
    except:
        pass

#Adding sorted label information to sampleMetaDF and sampleDF
    sampleMetaDF.insert(3, "labelSorted", sampleDF["label"].map(di))
    sampleDF.insert(5, "labelSorted", sampleDF["label"].map(di))
    print("Sorted labels assigned to sampleDF based on surface temperature, coldest to warmest
```

Sorted labels assigned to sampleDF based on surface temperature, coldest to warmest.

```
In [30]:
#Probability data to sampleMeta table cell
try:
        sampleMetaDF = sampleMetaDF.drop(columns = ["max posterior prob", "classUncertainty"])
except:
    pass

sampleMetaDF.insert(4, "max posterior prob", maxPosteriorProbs, True)
sampleMetaDF.insert(5, "classUncertainty", classUncertainty, True)
sampleMetaDF.head()
```

	lat	lon	time	labelSorted	max posterior prob	classUncertainty
0	-66.111519	73.5	1980-01-01	0	1.0	3.899727e-09
1	-65.703316	73.5	1980-01-01	0	1.0	6.174536e-10
2	-65.288567	73.5	1980-01-01	0	1.0	1.859135e-10
3	-64.867195	73.5	1980-01-01	0	1.0	1.053628e-10
4	-64.439102	73.5	1980-01-01	0	1.0	9.551204e-11

## Use pandas to calculate the properties of the profiles by sorted label

```
In [31]: #Class temperature means and stds cell
    sampleDFSortGrouped = sampleDF.groupby("labelSorted")
    sampleDFSortMeans = sampleDFSortGrouped.mean()
    sampleDFSortStds = sampleDFSortGrouped.std()
    profileCount = sampleDFSortGrouped[sampleDF.columns[0]].count().to_numpy()
    print("sampleDF grouped by sorted label (sampleDFSortGrouped), with means and standard development ("Number of samples in each group calculated and stored in profileCount.")
```

sampleDF grouped by sorted label (sampleDFSortGrouped), with means and standard deviations calculated for each group (sampleDFSortMeans, sampleDFSortStd).

Number of samples in each group calculated and stored in profileCount.

## Confirmation of sorting

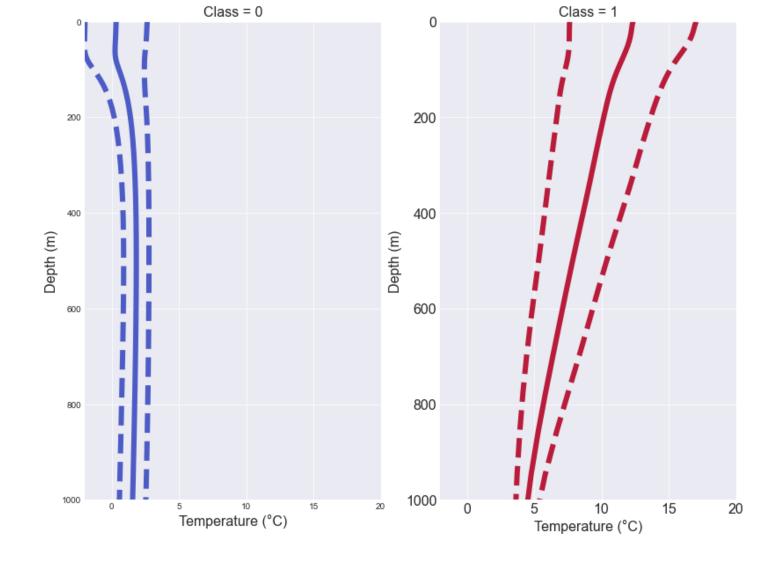
The means printed below should be ordered, going from coldest to warmest.

```
In [32]: #Temperature display cell
    print(sampleDFSortMeans[sampleDataDF.columns[0]])
```

labelSorted
0 0.339047
1 12.330616
Name: 0.5057600140571594, dtype: float32

## Plotting the means and standard deviations of the classes by profile

```
In [33]: #Plotting mean and std profiles cell
    tempGroupProfile(sampleDFSortMeans, sampleDFSortStds, 1)
    plt.show()
```



# Plotting location and cluster

```
In [34]: #surfaceTempPlot(sampleDF, 1)
plt.show()

In [35]: #locationPlotGroupDFLab(sampleDF, "Location plot of grouping", (25,25), 1)
plt.show()

In [36]: #locationPlotGroupDFMonthly(sampleDF, "Monthly summaries for training data set", 1)
print("Classifications, grouped by month.")
plt.show()

Classifications, grouped by month.

In [37]: #locationPlotUncertaintyDFMonthly(sampleDF, "Monthly uncertainty", 1)
print("Uncertainty in classifications, grouped by month.")
plt.show()
```

Uncertainty in classifications, grouped by month.

## **Exporting Meta Data**

```
Out[38]:
                  lat lon
                           time labelSorted max posterior prob classUncertainty
         0 -66.111519 73.5 1980-01-01
                                             0
                                                             1.0
                                                                    3.899727e-09
         1 -65.703316 73.5 1980-01-01
                                             0
                                                            1.0
                                                                   6.174536e-10
         2 -65.288567 73.5 1980-01-01
                                             0
                                                             1.0
                                                                 1.859135e-10
         3 -64.867195 73.5 1980-01-01
                                                             1.0 1.053628e-10
                                             0
         4 -64.439102 73.5 1980-01-01
                                                                    9.551204e-11
                                                             1.0
In [39]:
          #Meta data export cell
          exportName = modelName + " Meta Full"
          sampleMetaDF.to csv(exportName) #Exporting meta data
          print("Meta data and mask exported to "+ exportName +".")
         Meta data and mask exported to GMM UK 2Class R2 Meta Full.
In [40]:
          #Meta data reload cell
          importName = modelName + " Meta Full"
          sampleMetaReload = pd.read csv(importName)
          print("Meta data reloaded from "+ importName +". "+str(len(sampleMetaReload))+" data point
          sampleMetaReload.head()
         Meta data reloaded from GMM UK 2Class R2 Meta Full. 7989840 data points.
Out[40]:
          Unnamed: 0
                             lat lon
                                           time labelSorted max posterior prob classUncertainty
                    0 -66.111520 73.5 1980-01-01
                                                                               3.899727e-09
                                                        0
                                                                        1.0
         1
                    1 -65.703316 73.5 1980-01-01
                                                        0
                                                                        1.0
                                                                               6.174536e-10
```

0

0

0

1.0

1.0

1.0

1.859135e-10

1.053628e-10

9.551204e-11

## **End of Notebook**

2 -65.288570 73.5 1980-01-01

3 -64.867195 73.5 1980-01-01

4 -64.439100 73.5 1980-01-01

2

sampleMetaDF.head()

In [38]: