```
classdef TopoDatastore < matlab.io.Datastore & ...</pre>
       matlab.io.datastore.MiniBatchable & ...
       matlab.io.datastore.Shuffleable
    %TOPODATASTORE Custom datastore to read in data from file structure for TMN
       To be used as a datastore to a local directory of TUEG files.
    응
       Authors:
    응
           Michael Caiola (Michael.Caiola@fda.hhs.gov)
    응
           Meijun Ye (Meijun.Ye@fda.hhs.gov)
    응
    응
       Disclaimer: This software and documentation (the "Software") were
    응
       developed at the Food and Drug Administration (FDA) by employees of
       the Federal Government in the course of their official duties.
    응
       Pursuant to Title 17, Section 105 of the United States Code,
    응
    응
       this work is not subject to copyright protection and is in the
       public domain. Permission is hereby granted, free of charge, to any
       person obtaining a copy of the Software, to deal in the Software
    응
       without restriction, including without limitation the rights to
       use, copy, modify, merge, publish, distribute, sublicense, or sell
    응
       copies of the Software or derivatives, and to permit persons to
    응
       whom the Software is furnished to do so. FDA assumes no
    응
       responsibility whatsoever for use by other parties of the Software,
       its source code, documentation or compiled executables, and makes
    응
       no guarantees, expressed or implied, about its quality,
    응
    응
       reliability, or any other characteristic. Further, use of this code
       in no way implies endorsement by the FDA or confers any advantage
    응
       in regulatory decisions. Although this software can be
       redistributed and/or modified freely, we ask that any derivative
       works bear some notice that they are derived from it, and any
       modified versions bear some notice that they have been modified.
   properties
       Datastore
       Labels
       NumClasses
       SequenceDimension
       MiniBatchSize
       Augmented
       TestSet
        newHz
        fband=[1 4; 4 8; 8 12; 12 16; 16 20; 25 40];
       chLocations
   end
   properties(SetAccess = protected)
       NumObservations
   end
   properties (Access = private)
       CurrentFileIndex
```

```
%FileSet matlab.io.datastore.DsFileSet
end
methods
    function ds = TopoDatastore(folder,newHz,chLocations,varargin)
        % ds = NewDatastore(folder,newHz) creates a custom datastore
        % from the data in folder with new sampling sampling rate newHz
        % chLocations are needed for topoplot, use chlocs2.mat
        % Optional arguments:
          "DataAugmentation" (default = false)
        엉
                Augments the data in random shuffles of 90 seconds
        응
            "TestSet (default = false)
        응
                Creates a testset of 90 second data, to be used when
                DataAugmentation is being used
        if isempty(newHz)
            ds.newHz = 250;
        else
            ds.newHz = newHz;
        end
        inputs = ds.parseInputs(varargin{:});
        if inputs.DataAugmentation
            ds.Augmented = true;
        else
            ds.Augmented = false;
        end
        if inputs.TestSet
            ds.TestSet = true;
            ds.Augmented = false;
        else
            ds.TestSet = false;
        end
        if exist("topoplot", 'file') ~=2
            %open eeglab
            warning ("Need EEGLAB! Trying to open...")
            try
                path = pwd;
                eeglabpath = uigetdir([], "Locate EEGLAB Directory");
                cd(eeglabpath)
                eeglab
                close
                cd(path)
            catch
                error ("Could not find EEGLAB. Open prior to use to add to path.")
            end
```

```
end
```

```
% Save chLocations
    ds.chLocations = chLocations;
    % Create file datastore.
    fds = fileDatastore(folder, ...
        'ReadFcn',@(x) readSequence(x,ds.newHz,ds.Augmented,ds.TestSet,...
        ds.fband, ds.chLocations), ...
        'IncludeSubfolders', true);
    ds.Datastore = fds;
    % Read labels from folder names.
    numObservations = numel(fds.Files);
    for i = 1:numObservations
        file = fds.Files{i};
        filepath = fileparts(file);
        [~,label] = fileparts(filepath);
        labels{i,1} = label;
    end
    ds.Labels = categorical(labels);
    ds.NumClasses = numel(unique(labels));
    % Determine sequence dimension.
   X = preview(fds);
   ds.SequenceDimension = size(X,1);
    % Initialize datastore properties.
    ds.MiniBatchSize = 128;
    ds.NumObservations = numObservations;
    ds.CurrentFileIndex = 1;
end
function tf = hasdata(ds)
    % tf = hasdata(ds) returns true if more data is available.
    %tf = hasdata(ds.Datastore);
    tf = ds.CurrentFileIndex + ds.MiniBatchSize - 1 ...
        <= ds.NumObservations;
end
function [data,info] = read(ds)
    % [data,info] = read(ds) read one mini-batch of data.
   miniBatchSize = ds.MiniBatchSize;
    if ds.NumObservations < ds.MiniBatchSize</pre>
        ds.MiniBatchSize = ds.NumObservations;
    end
    i = 0;
```

```
while i < miniBatchSize && hasdata(ds) %hasdata(ds.Datastore)
        i = i + 1;
        Predictors{i,1} = read(ds.Datastore);
        %temppred = read(ds.Datastore);
        %predictors{i,1} = spect(temppred);
        Response(i,1) = ds.Labels(ds.CurrentFileIndex);
        ds.CurrentFileIndex = ds.CurrentFileIndex + 1;
    end
    data = preprocessData(Predictors, Response);
    info.Size = size(data);
end
function reset(ds)
    % reset(ds) resets the datastore to the start of the data.
    reset (ds.Datastore);
    ds.CurrentFileIndex = 1;
end
function dsNew = subset(ds,in)
    dsNew = copy(ds);
   dsNew.Datastore=subset(ds.Datastore,in);
    dsNew.NumObservations = numel(dsNew.Datastore.Files);
    for i = 1:dsNew.NumObservations
        file = dsNew.Datastore.Files{i};
        filepath = fileparts(file);
        [~,label] = fileparts(filepath);
        labels{i,1} = label;
    end
    dsNew.Labels = categorical(labels);
    dsNew.NumClasses = numel(unique(labels));
end
function varargout = grabEachLabel(ds,N,varargin)
    % pulls a certain number of each individual label
    opt random=0;
    numvarargin=length(varargin);
   k=1;
    while k<=numvarargin
        switch varargin(k)
            case "random"
                opt random=1;
        end
        k=k+1;
    end
    for i=1:length(N)
        in{i}=[];
    labels=unique(ds.Labels);
```

```
for i=1:ds.NumClasses
            labs=ds.Labels==labels(i);
            numLabs=sum(labs);
            x=find(labs);
            if opt_random
                x=x (randperm (length (x)));
            end
            if sum(N)>numLabs
                n=floor(numLabs.*N/sum(N));
                warning("Not enough " + string(labels(i))+" Labels")
            else
                n=N;
            end
            in{1}=[in{1}; x(1:n(1))];
            for j=2:length(n)
                in{j}=[in{j};x(n(j-1)+1:n(j-1)+n(j))];
            end
        end
        for i=1:length(in)
            varargout{i} = subset(ds,in{i});
        end
    end
    function dsNew = shuffle(ds)
        % dsNew = shuffle(ds) shuffles the files and the corresponding
        % labels in the datastore.
        % Create copy of datastore.
        dsNew = copy(ds);
        dsNew.Datastore = copy(ds.Datastore);
        fds = dsNew.Datastore;
        % Shuffle files and corresponding labels.
        numObservations = dsNew.NumObservations;
        idx = randperm(numObservations);
        fds.Files = fds.Files(idx);
        dsNew.Labels = dsNew.Labels(idx);
    end
end
methods (Hidden = true)
    function frac = progress(ds)
        % frac = progress(ds) returns the percentage of observations
        % read in the datastore.
        frac = (ds.CurrentFileIndex - 1) / ds.NumObservations;
    end
end
```

```
methods (Access = 'private')
        function inputStruct = parseInputs(ds, varargin)
            p = inputParser();
            p.addParameter('DataAugmentation', false, @augmentationValidator);
            p.addParameter('TestSet', false, @augmentationValidator);
            p.parse(varargin(:));
            inputStruct = p.Results;
        end
    end
end
function data = preprocessData(Predictors, Response)
% data = preprocessData(predictors, responses) preprocesses
% the data in predictors and responses and returns the table
% data
miniBatchSize = size(Predictors,1);
% Pad data to length of longest sequence.
sequenceLengths = cellfun(@(X) size(X,2), Predictors);
maxSequenceLength = max(sequenceLengths);
for i = 1:miniBatchSize
    X = Predictors{i};
    % Pad sequence with zeros.
    if size(X,2) < maxSequenceLength</pre>
        X(:, maxSequenceLength) = 0;
    end
    Predictors{i} = X;
end
% Return data as a table.
data = table(Predictors, Response);
end
function data = readSequence(filename, newHz, aug, ts, fband, chlocs)
            % data = readSequence(filename) reads the sequence y from the MAT file
            % filename
            oldHz = 250;
            S = load(filename);
            if newHz == oldHz
                data = S.y;
            else
                data = single(resample(double(S.y),newHz,oldHz,'Dimension',2)); % ✓
resampled
            end
            downsample = 90; %sec
            if aug
```

```
r = randi(downsample*newHz-2)+1;
                data = data(:,r:r+(downsample*newHz-1));
            end
            if ts
                data = data(:,1:1+(downsample*newHz-1));
            end
            %Topo
            data = data';
            tot=bandpower(data,newHz,[1 newHz/2]);
            n = size(fband, 1);
            abs psd = zeros(size(data,2),n);
            for i=1:n
                abs_psd(:,i)=bandpower(data,newHz,fband(i,:));
            end
            rel psd = abs psd./tot';
            gscale = 67*2; %interpoltation value
            y = zeros(gscale, gscale, n);
            for i = 1:n
                [~,y(:,:,i)] = topoplot(rel psd(:,i),chlocs,'noplot','on','gridscale', ✓
gscale);
            end
            data = y;
            data = fillmissing(data, 'constant', 0);
end
function TF = augmentationValidator(valIn)
% if ischar(valIn) || isstring(valIn)
     TF = string('none').contains(lower(valIn)); %#ok<STRQUOT>
% elseif isa(valIn,'imageDataAugmenter') && isscalar(valIn)
     TF = true;
if islogical(valIn)
   TF = true;
else
    TF = false;
end
end
```