```
function [outputArg1,outputArg2] = runSPM(folderin,folderout)
% Runs processJumps using file list from getSFIfiles
% then runs SPM stats from spm1d package
%
%
%
% param.SPM = true;
% setParameters = param;
```

INPUT

```
[reslts] = processJumps(folderin,[],[],[],folderout);
SFIbench = readmatrix(fullfile("Z:\Private Student Folders\ShadanA", "wrangled_benchmark_SFI.csv"), 'Range',2, 'OutputType','string');
SFIaclr = readmatrix(fullfile("Z:\Private Student Folders\ShadanA", "wrangled ACLRmatched SFI.csv"), 'Range',2, 'OutputType','string');
MAIN
extract all force data for each jump
alliumps = [];
Dir = struct; % Dir (Directory) will summarize
col = 0;
for iFile = 1:length(reslts)
    for iJump = 1:length(reslts(iFile).jumpStats)
        if ~isempty(reslts(iFile).jumpStats(iJump).jumpData)
           col = col+1;
           iumpdata =
reslts(iFile).jumpStats(iJump).jumpData(reslts(iFile).jumpStats(iJump).phases(1,1):reslts(iFile).jumpStats(iJump).phases(4,1),2:3); % L
and R force for each jump
           normfactor = length(jumpdata)/1001;
           jumpdata = interp1([1:length(jumpdata)],jumpdata,linspace(1,length(jumpdata),101)); % normalize to 101 points
           jumpdata = sum(jumpdata,2); % sum by row to create total force
           Dir(col).Athlete = reslts(iFile).Athlete;
           Dir(col).Sex = reslts(iFile).Sex;
           Dir(col).JumpNum = reslts(iFile).jumpStats(iJump).JumpNumb; % each athlete has 3 jumps on each leg
           Dir(col).Legs = reslts(iFile).jumpStats(iJump).SL; % jumping leg
```

```
Dir(col).JumpType = reslts(iFile).JumpType; % CMJ (countermovement jump) or RHT (repeated hop test)
           Dir(col).extraload = reslts(iFile).Extra Load; % jumping with load
           Dir(col).SystemMass = reslts(iFile).System Mass; % athlete mass
           Dir(col).phases = round(reslts(iFile).jumpStats(iJump).velImpulseTimes/normfactor);
               % assign limb function catagory
               if SFIbench(find(contains(SFIbench(:,2),reslts(iFile).Athlete)),34)=="Good"
                    Dir(col).LimbFunction = 'Good';
               elseif SFIbench(find(contains(SFIbench(:,2),reslts(iFile).Athlete)),34)=="Poor"
                    Dir(col).LimbFunction = 'Poor';
               elseif SFIaclr(find(contains(SFIaclr(:,2),reslts(iFile).Athlete)),34)=="Poor" % different csv file for aclr athletes
                    Dir(col).LimbFunction = 'Poor';
               elseif SFIaclr(find(contains(SFIaclr(:,2),reslts(iFile).Athlete)),34)=="Fair" % different csv file for aclr athletes
                    Dir(col).LimbFunction = 'Fair';
                end
               % assign ACLR status
               if SFIaclr(find(contains(SFIaclr(:,2),reslts(iFile).Athlete)),24)=="Yes"
                    Dir(col).ACLstatus = 'ACLR'; % ACL injured athlete
                    Dir(col).ACLside = SFIaclr(find(contains(SFIaclr(:,2),reslts(iFile).Athlete)),26); % injured limb
               elseif SFIaclr(find(contains(SFIaclr(:,2),reslts(iFile).Athlete)),24)=="No"
                    Dir(col).ACLstatus = 'Match'; % matched control
                   Dir(col).ACLside = '':
                else
                    Dir(col).ACLstatus = 'Control'; % limb function control
                   Dir(col).ACLside = '';
                end
        alljumps(:,col) = jumpdata./(Dir(col).SystemMass*9.81); % normalize force to body mass
        end
    end
end
alljumps = alljumps'; % transpose data. SPM wants ROWS as subjects and COLS at timepoints
```

CREATE LIMB FUNCTION GROUPS FOR LEFT, RIGHT, INJURED LEG

```
% Good perceived limb function
Left good sfi = find(arrayfun(@(x) strcmp(x.Legs,'L') & strcmp(x.LimbFunction,'Good'),Dir));
Right\_good\_sfi = find(arrayfun(@(x) strcmp(x.Legs,'R') & strcmp(x.LimbFunction,'Good').Dir));
Left_good_sfi = alljumps(Left_good_sfi,:);
Right good sfi = alljumps(Right good sfi,:);
% Poor perceived limb function
Left poor sfi = find(arrayfun(@(x) strcmp(x.Legs,'L') & strcmp(x.LimbFunction,'Poor') & \simstrcmp(x.ACLstatus,'ACLR'),Dir));
Right poor sfi = find(arrayfun(@(x) strcmp(x.Legs,'R') & strcmp(x.LimbFunction,'Poor') & ~strcmp(x.ACLstatus,'ACLR'),Dir));
Left_poor_sfi = alljumps(Left_poor_sfi,:);
Right poor sfi = alljumps(Right poor sfi,:);
% ACLR between injured limbs
acl_inj = find(arrayfun(@(x) strcmp(x.ACLstatus,'ACLR') & ...
                            (strcmp(x.ACLside,'L') & strcmp(x.Legs,'L')) | ...
                            (strcmp(x.ACLside, 'R') \& strcmp(x.Legs, 'R')), Dir)); % find indicies where the jumping leg = ACLR leg
acl uninj = find(arrayfun(@(x) strcmp(x.ACLstatus, 'ACLR') \& ...
                            (strcmp(x.ACLside,'L') & strcmp(x.Legs,'R')) | ...
                            (strcmp(x.ACLside, 'R') & strcmp(x.Legs, 'L')),Dir)); % find indicies where the jumping leg is not ACLR leg
acl inj = alljumps(acl inj,:);
acl uninj = alljumps(acl uninj,:);
% ACLR matched controls
control = find(arrayfun(@(x) strcmp(x.ACLstatus, 'Match') & strcmp(x.Legs, 'R'),Dir));
control = alljumps(control,:);
```

RUN SPM ANALYSIS - spm1d program

```
% left to right
LR_good_spm = spm1d.stats.ttest_paired(Left_good_sfi, Right_good_sfi);
LR_good_spmi = LR_good_spm.inference(0.05, 'two_tailed', true);

LR_poor_spm = spm1d.stats.ttest_paired(Left_poor_sfi, Right_poor_sfi);
LR_poor_spmi = LR_poor_spm.inference(0.05, 'two_tailed', true);
```

```
% ACLR between injured limb groups
aclr_spm = spm1d.stats.ttest_paired(acl_inj, acl_uninj);
aclr_spmi = aclr_spm.inference(0.05, 'two_tailed', true);
```

CREATE FIGURE

```
fig12 = tiledlayout(2,3);
    fig12.Title.String = 'Between Limb Compairison of Single Leg CMJ';
    fig12.Title.FontWeight = 'bold';
    fig12.Title.FontSize = 14;
    fig12.TileSpacing = 'compact';
   % fig12.Position = [1000 300 650 1000];
    poorspm = nexttile; % SPM - Poor SFI
           LR poor spmi.plot()
           LR_poor_spmi.plot_p_values()
           title('Poor SFI Group', "FontWeight", "normal")
    goodspm = nexttile; % SPM - Good SFI
           LR_good_spmi.plot()
           LR good spmi.plot p values()
           title('Good SFI Group', "FontWeight", "normal")
    aclrspm = nexttile; % SPM - ACLR SFI
            aclr_spmi.plot()
           aclr spmi.plot p values()
           title('ACLR Group', "FontWeight", "normal")
    poorfz = nexttile; % Fz-time curve - Poor SFI (Left is blue, Right is grey)
            spm1d.plot.plot meanSD(Left poor sfi, 'color', '#457DEE')
           hold on
            spm1d.plot.plot meanSD(Right poor sfi, 'color', '#8D8E90')
           hold off
```

```
goodfz = nexttile; % Fz-time curve - Good SFI (Left is blue, Right is grey)
            spm1d.plot.plot meanSD(Left good sfi, 'color', '#457DEE')
           hold on
            spm1d.plot.plot_meanSD(Right_good_sfi, 'color', '#8D8E90')
            hold off
    aclrfz = nexttile; % Fz-time curve - injured is blue, uninjured is grey
            spm1d.plot.plot meanSD(acl inj, 'color', '#457DEE')
           hold on
            spm1d.plot.plot_meanSD(acl_uninj, 'color', '#8D8E90')
            hold off
    linkaxes ([poorspm, poorfz], 'x');
    linkaxes ([goodspm, goodfz], 'x');
    linkaxes ([aclrspm, aclrfz], 'x');
    linkaxes ([poorspm, goodspm, aclrspm], 'y');
   xlabel ([poorfz, goodfz, aclrfz], 'Time (%)')
    ylabel ([goodspm, aclrspm], '')
   ylabel (poorfz, 'Force (N/BW)')
   xticks ([poorspm, poorfz, goodspm, goodfz, aclrspm, aclrfz], [25 50 75 100])
    xticklabels ([poorfz, goodfz, aclrfz], {'25' '50' '75' '100'})
    xticklabels ([poorspm, goodspm, aclrspm], {})
    legend(poorfz, {'Left Limb', '', 'Right Limb'});
    legend(goodfz, {'Left Limb', '', 'Right Limb'});
    legend(aclrfz, {'Injured Limb', '', 'Uninjured Limb'});
    saveas(fig12,'Z:\Private Student Folders\ShadanA\Figures\BetweenLimbs ACL Good Poor.jpg')
outputArg1 = fig12;
outputArg2 = fig13;
end
```