

```
% For Fig 4:
% adult+juven lob 6, juven crus 1-> YM PC5, PC6
% juven crus 1, juven crus 2, adult+juven lob 7 -> SC PC3

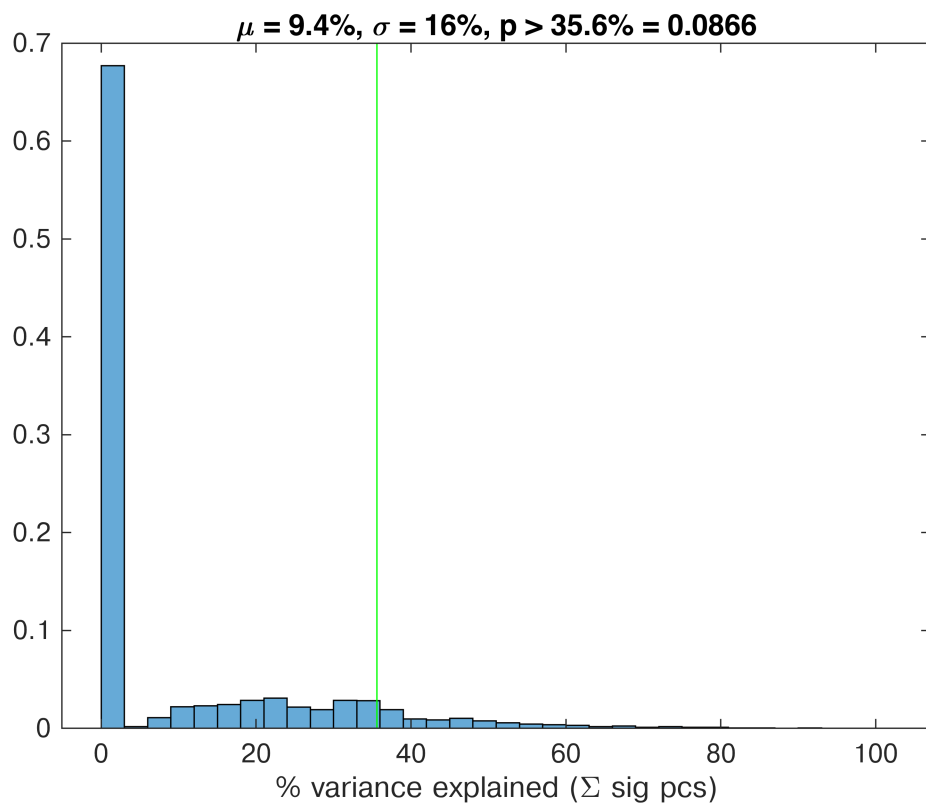
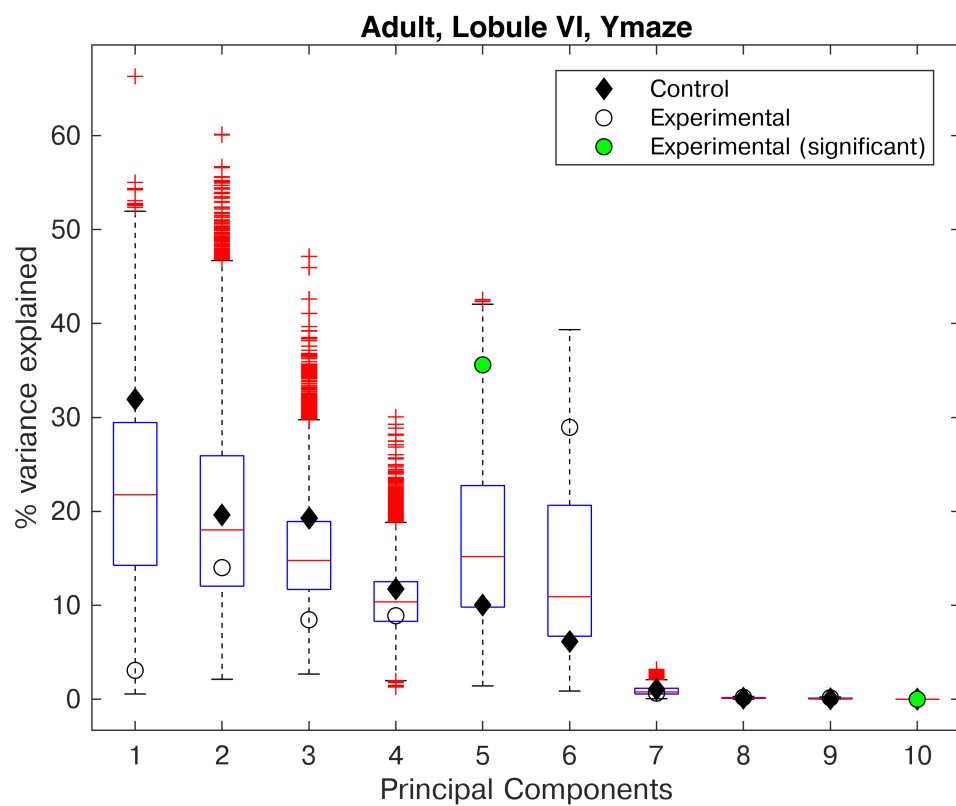
clear all, close all, clc
```

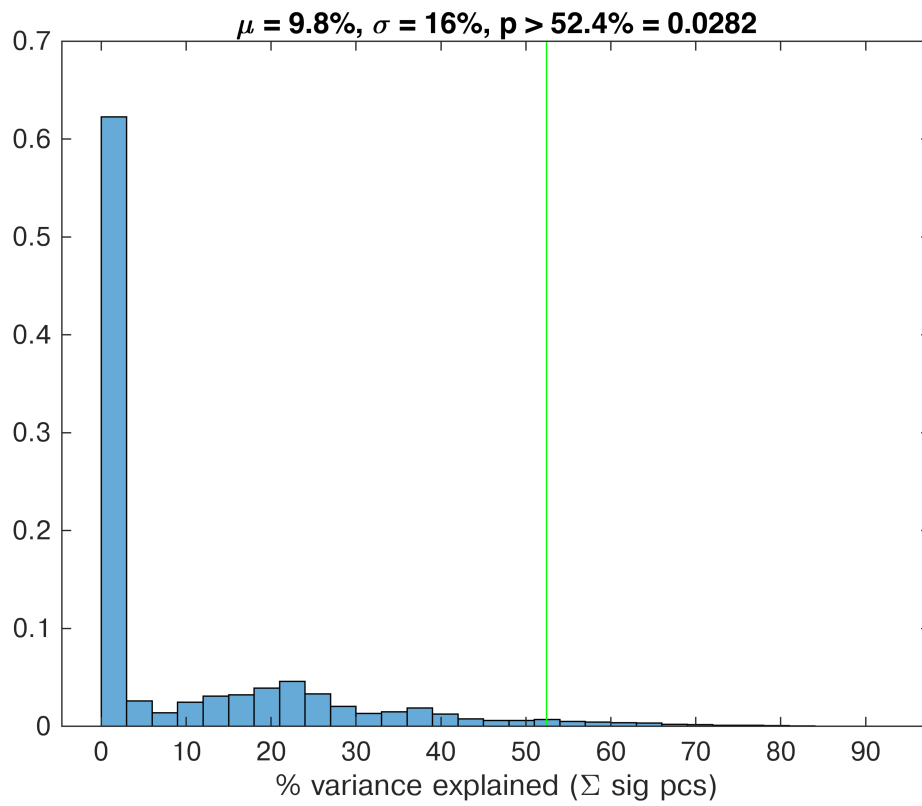
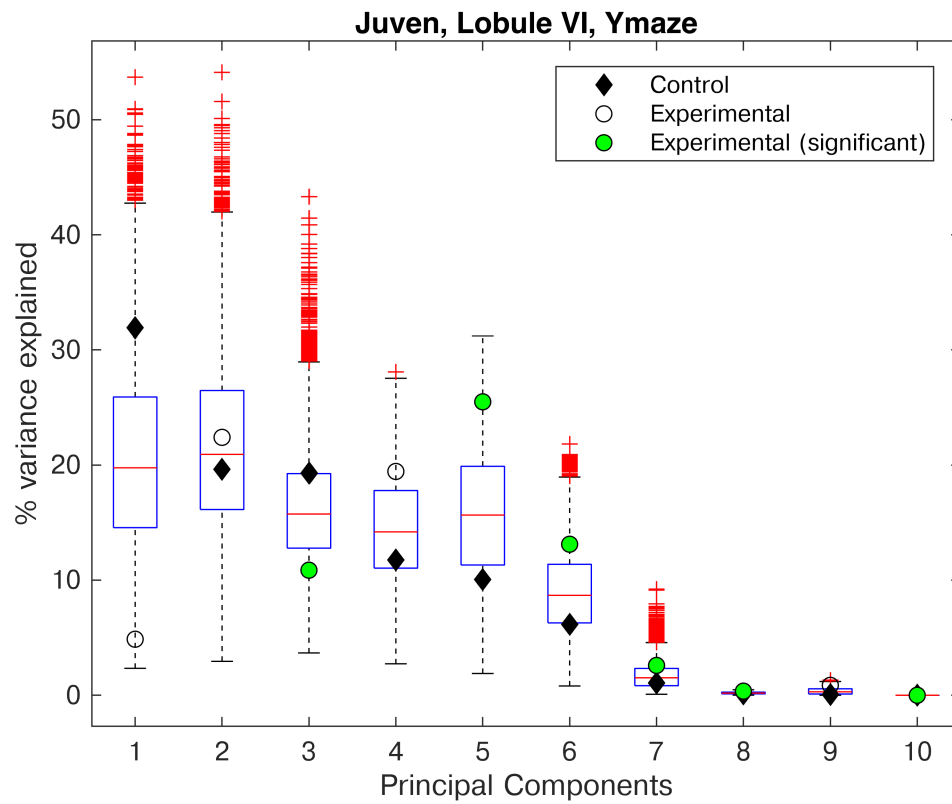
```
load('data/2017-08-31-ks_analysis.mat')
whos
```

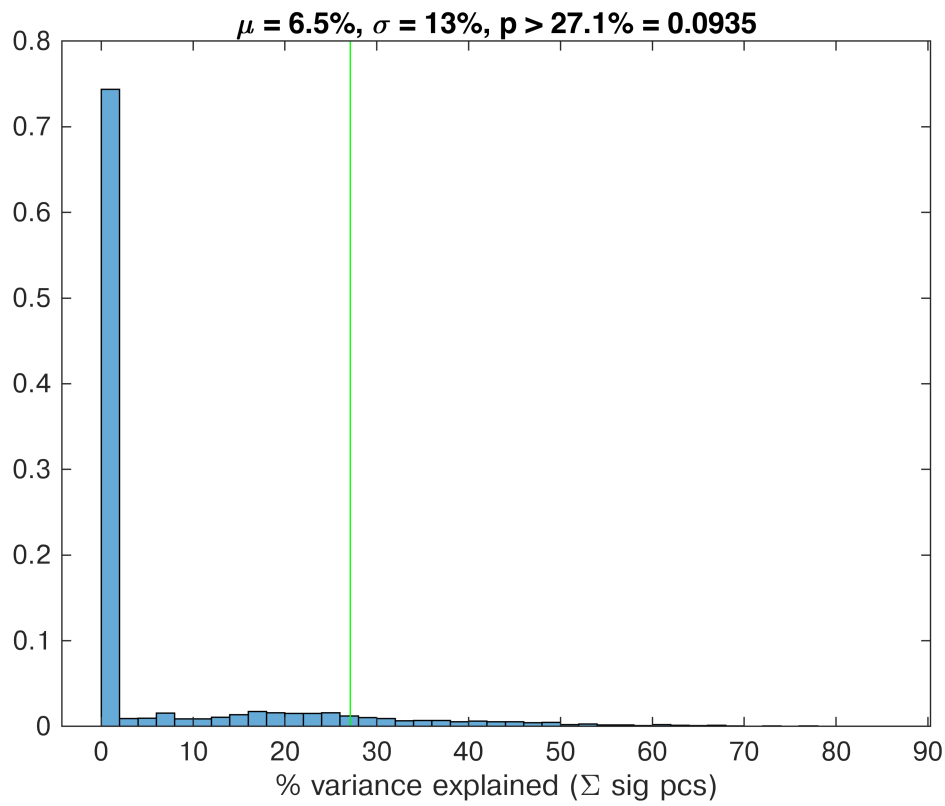
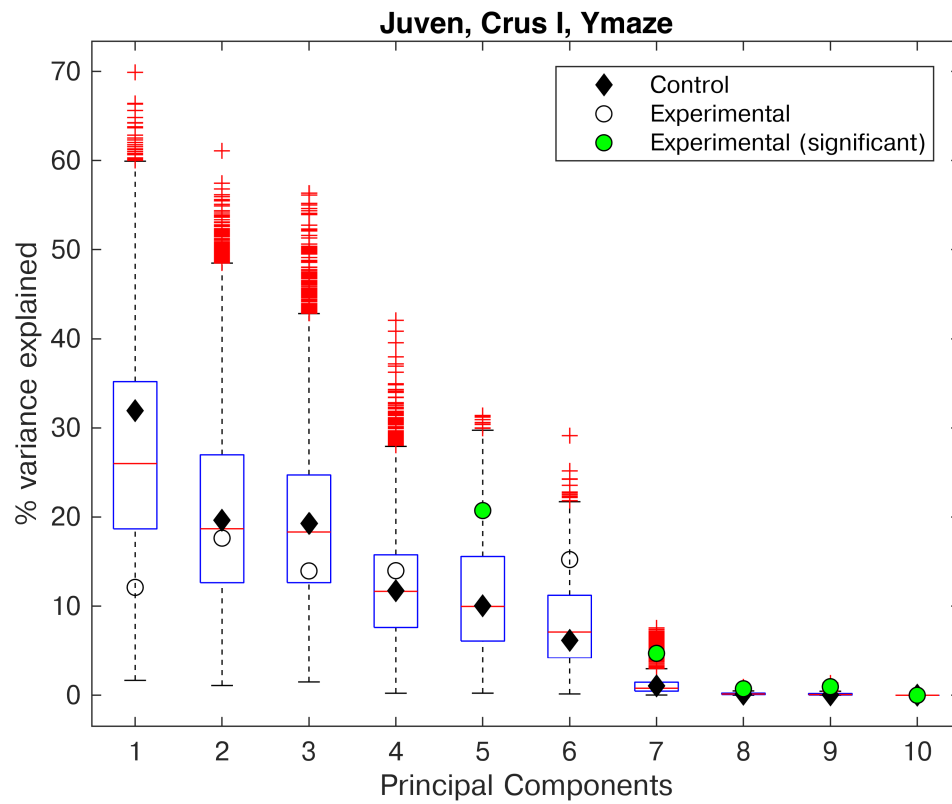
Name	Size	Bytes	Class	Attributes
epm	1x1	841024	struct	
exptAges	2x1	244	cell	
exptRegions	4x1	512	cell	
gr	1x1	422265	struct	
kstable	336x9	53448809	table	
sc	1x1	998745	struct	
timestamp	1x20	40	char	
ym	1x1	1731200	struct	

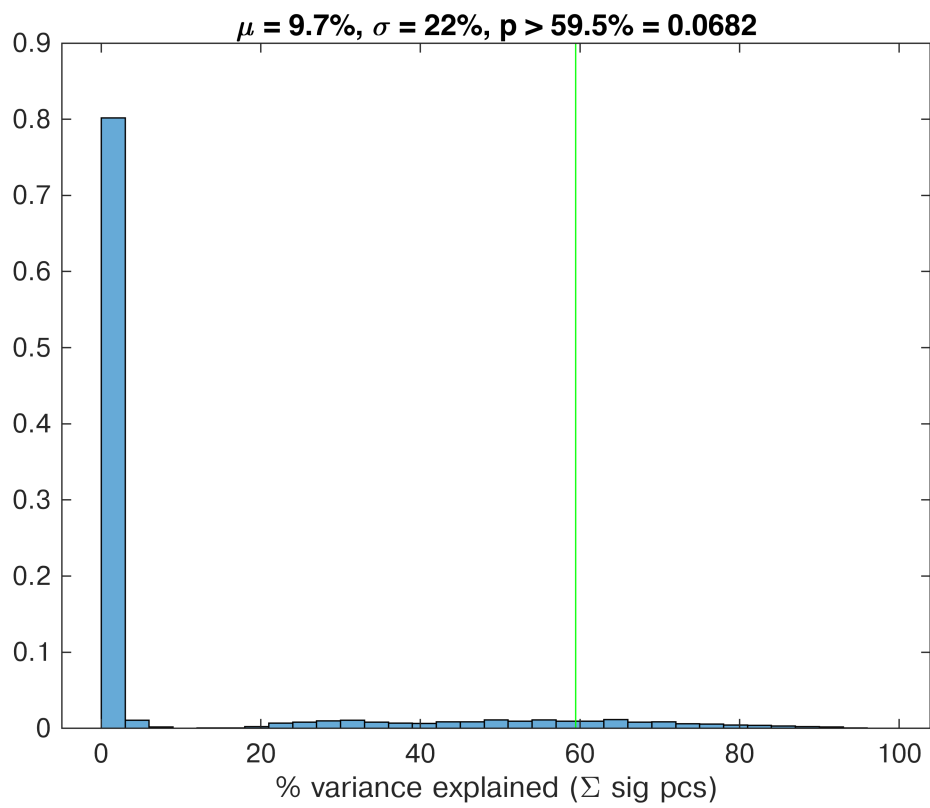
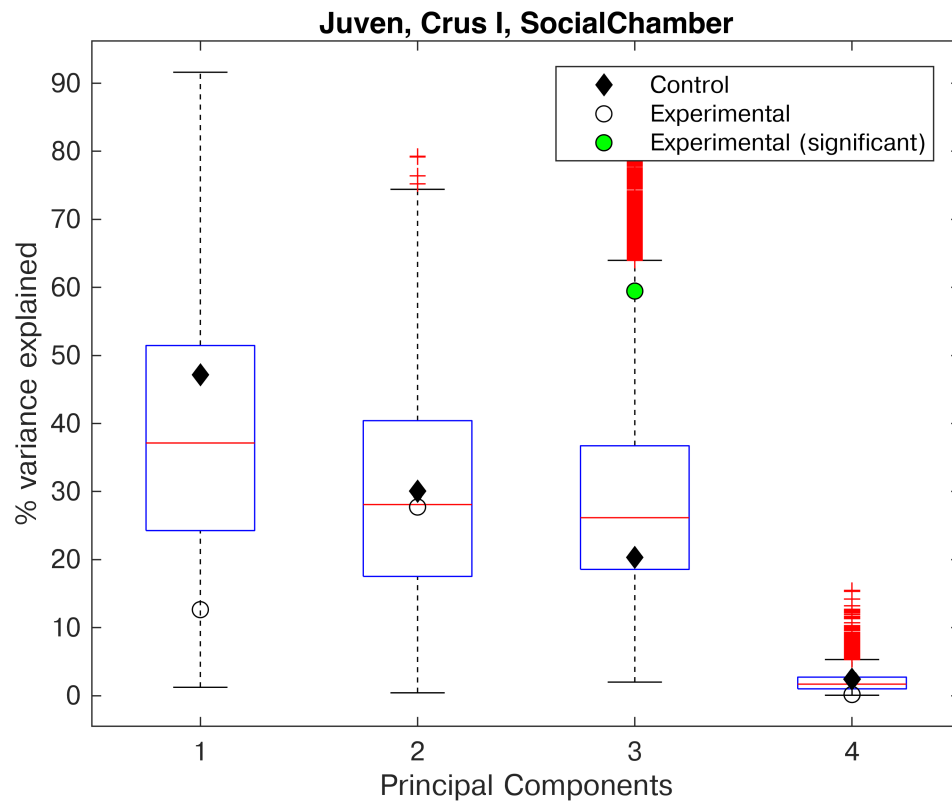
```
Bs = {
    ym.LobuleVI.Adult
    ym.LobuleVI.Juven
    ym.CrusI.Juven
    sc.CrusI.Juven
    sc.CrusII.Juven
    sc.LobuleVII.Adult
    sc.LobuleVII.Juven
};

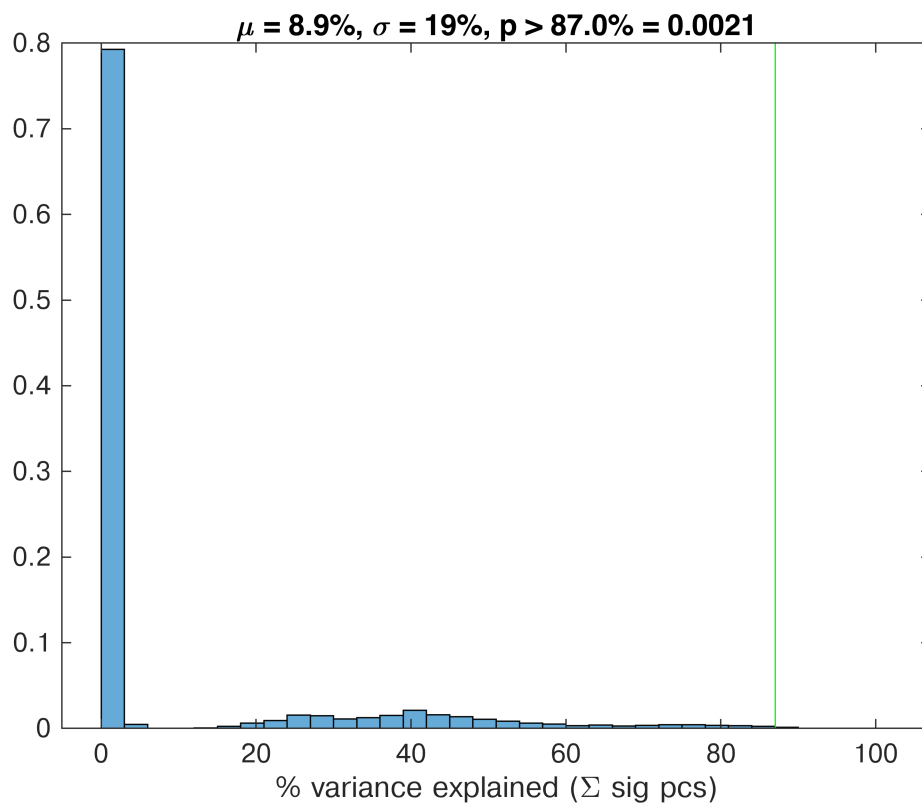
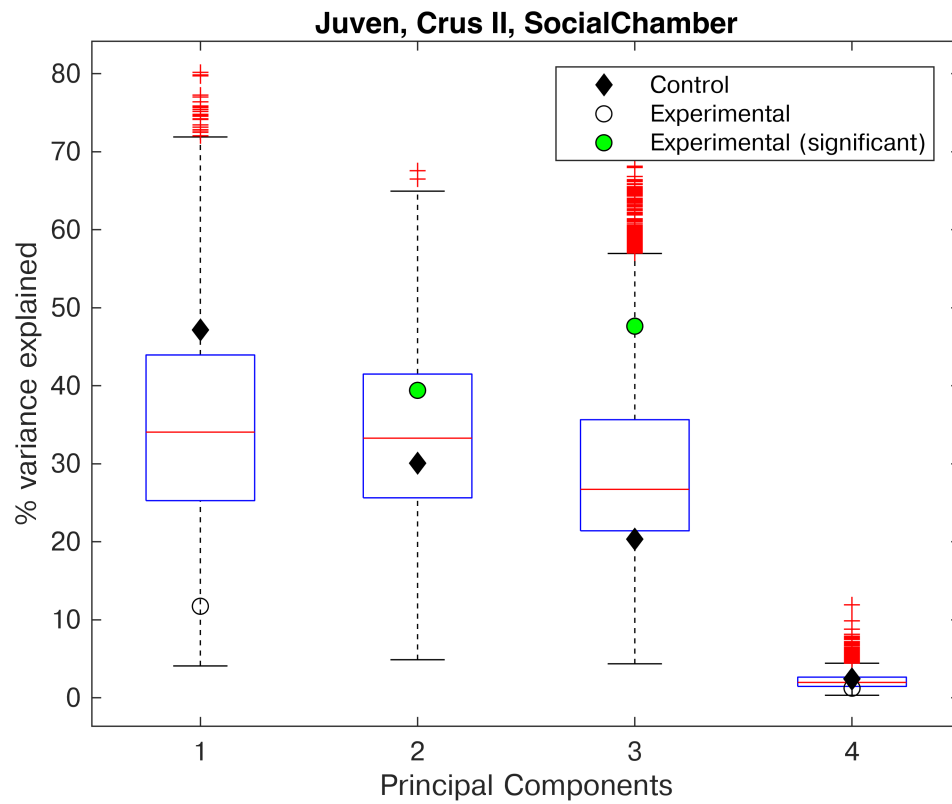
for i = 1:numel(Bs)
    run_shuffle(Bs{i})
end
```

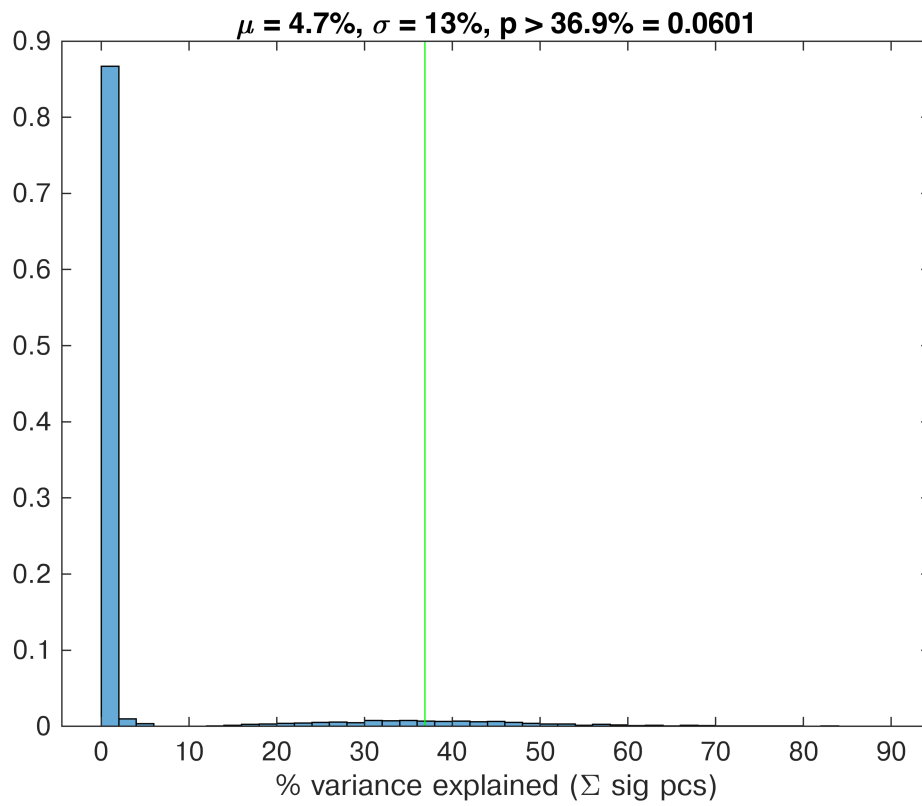
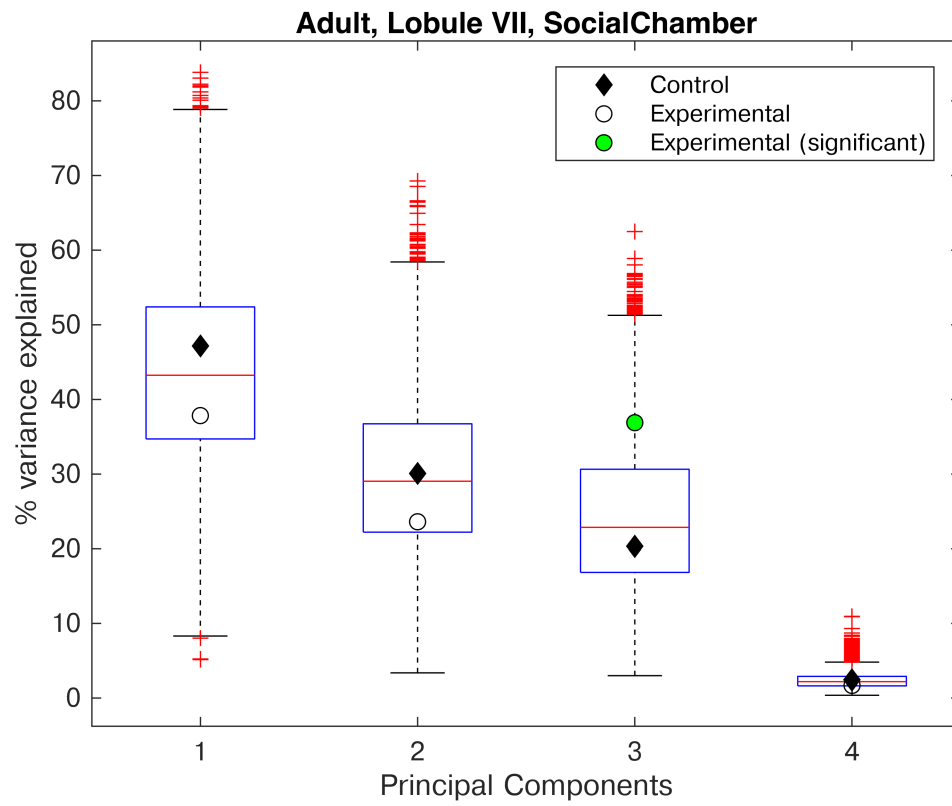


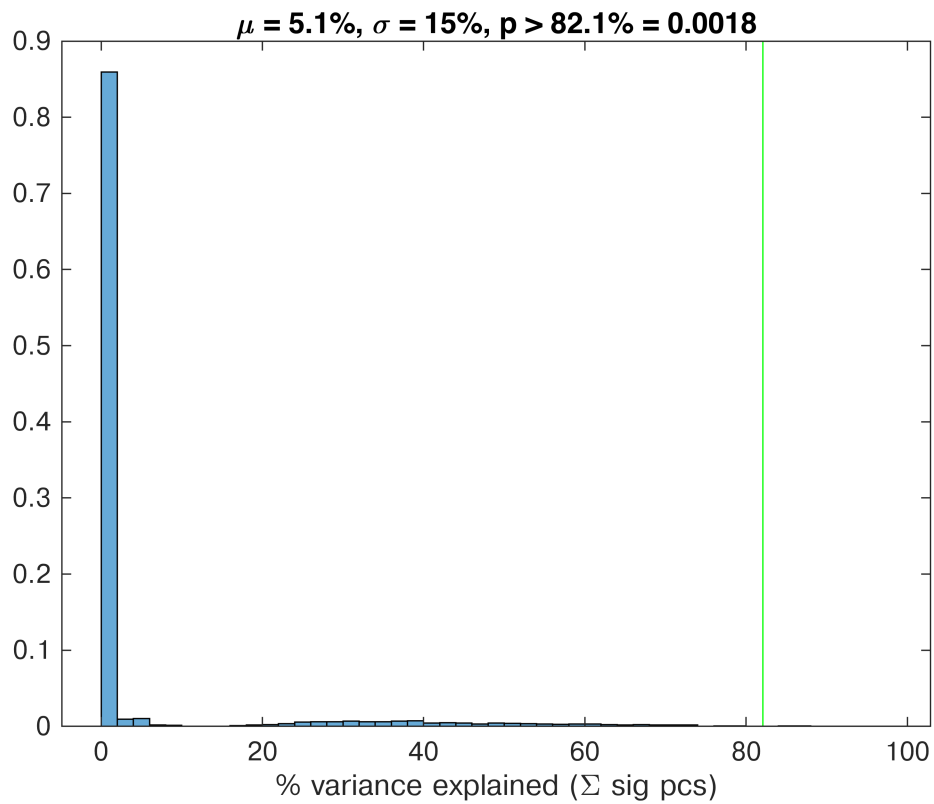
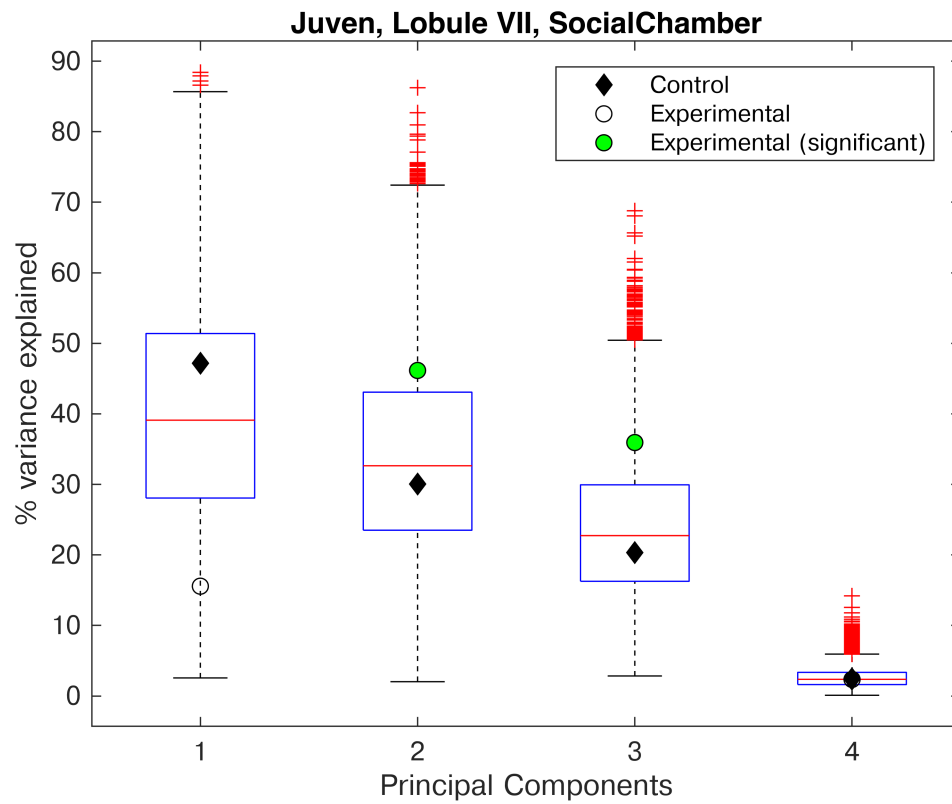












```
function run_shuffle(B, sig_var_perc_thresh)
if nargin < 2; sig_var_perc_thresh = []; end
```



```

% already normalized/projected:
allmice_amps = B.pca{:,,:};
untreat_amps = B.pca{B.isCtrl,:}; % controls
treated_amps = B.pca{~B.isCtrl,:}; % experimentals

num_untreat = size(untreat_amps,1);
num_treated = size(treated_amps,1);
d = size(allmice_amps,2); % number of pcs

% latent (eigenvalues of covariance)
untreat_latent = sum(untreat_amps.^2) ./ size(untreat_amps,1);
treated_latent = sum(treated_amps.^2) ./ size(treated_amps,1);

% percent variance explained
untreat_var_perc = untreat_latent ./ sum(untreat_latent) .* 100;
treated_var_perc = treated_latent ./ sum(treated_latent) .* 100;

N = 10000; % number of bootstrap samples
% k = size(untreat_amps,1); % number of control mice per sample
k = size(treated_amps,1); % number of experimental mice per sample
alpha = 0.05; % significance level

% repeat procedure with data for reference
[~,treated_pval] = ttest2(untreat_amps,treated_amps); % significance test
treated_is_sig = treated_pval < alpha;
treated_sig_var = sum(treated_latent(treated_is_sig));
treated_sig_var_perc = treated_sig_var / sum(treated_latent) * 100;

% bootstrap
% tic;
sample_pval = NaN(N,d);
sample_var = NaN(N,d);
sample_sig_var = NaN(N,1);
sample_num_sig = NaN(N,1);
for i = 1:N
    sample_amps = datasample(allmice_amps,k); % random subsample
    sample_latent = sum(sample_amps.^2) ./ size(sample_amps,1); % compute latents (eigenvalues)

    [~,sample_pval(i,:)] = ttest2(untreat_amps,sample_amps); % significance test
    is_sig = sample_pval(i,:) < alpha;

    sample_var(i,:) = sample_latent; % variance per pc
    sample_sig_var(i) = sum(sample_latent(is_sig)); % sum of variance for significant pcs
    sample_num_sig(i) = sum(is_sig); % count of significant pcs
end
% toc

% compute percent variance
sample_var_perc = sample_var ./ sum(sample_var,2) * 100;
sample_sig_var_perc = sample_sig_var ./ sum(sample_var,2) * 100;

% Plot

```

```

figure,boxplot(sample_var_perc), hold on
h1 = plot(1:d, untreat_var_perc, 'dk', 'MarkerSize',6, 'MarkerFaceColor','k');
h2 = plot(1:d, treated_var_perc, 'ok', 'MarkerSize',6);
h3 = plot(find(treated_is_sig), treated_var_perc(treated_is_sig), 'ok', 'MarkerSize',6, 'MarkerFaceColor','k');
xlabel('Principal Components'),ylabel('% variance explained')
legend([h1 h2 h3],{'Control', 'Experimental', 'Experimental (significant)'})
title(sprintf('%s, %s, %s', B.age_name, B.region_desc, B.assay_name))

% default to empirically computed
if isempty(sig_var_perc_thresh); sig_var_perc_thresh = treated_sig_var_perc; end

figure,histogram(sample_sig_var_perc, 'Normalization','probability'),hold on
plot([1 1] * sig_var_perc_thresh, ylim(), 'g-')
xlabel('% variance explained (\Sigma sig pcs)')
title(sprintf('\mu = %.1f%%, \sigma = %1.1f%%, p > %.1f%% = %g', ...
    mean(sample_sig_var_perc), std(sample_sig_var_perc), ...
    sig_var_perc_thresh, mean(sample_sig_var_perc > sig_var_perc_thresh)))

end

```