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Demo 05: noise test

A demo script to check the noise statitics. Not all spectral seperation modes and processing steps gives a proper SNR metric. This script is to verify the standard normal distribution of data, image and metabolite amplitudes after every processing step. Download noise data (800 MB "phantom-DMI-noise.tar.gz") from zenodo.org (DOI: 10.5281/zenodo.14652737) to follow this excercise.

FISP CSI dataset with acquistion weighting

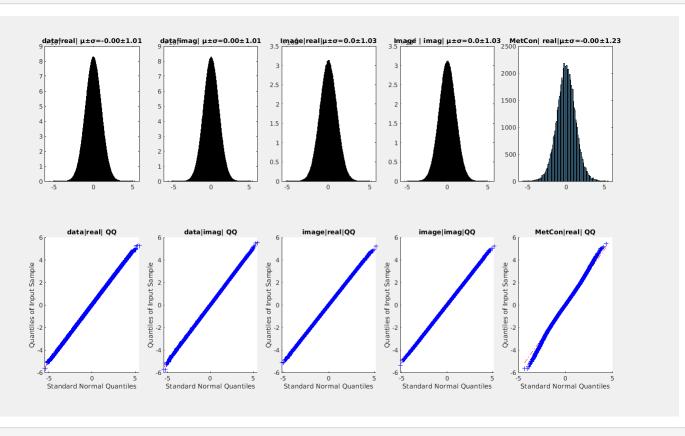
```
% add all dependencies
addpath(genpath('/ptmp/pvalsala/Packages/mapVBVD'))
addpath(genpath('/ptmp/pvalsala/Packages/DeuteMetCon'))
addpath(genpath('/ptmp/pvalsala/Packages/OXSA'))
%data path
sn='/ptmp/pvalsala/deuterium/dataForPublication/phantom-DMI-Noise';
fn=fullfile(sn,'meas_MID00750_FID11717_rpcsi_fid_Stan_res15_6_TR36_5min_0FA.dat');
metabolites=getMetaboliteStruct('phantom',0);
CSI_setting={'metabolites',metabolites,'parfor',true,...
     'doCoilCombine', 'adapt1', 'doZeropad', [0.5 0.5 0.5 0]*0, 'mask', [], 'Solver', 'IDEAL'};
mcobj_fisp=MetCon_CSI(fn,CSI_setting{:});
Software version: VD (!?)
Reader version: 1660732089 (UTC: 17-Aug-2022 10:28:09)
Scan 1/1, read all mdhs:
  324.1 MB read in 18 s
Using T2* =20 ms
starting reco
                     10 25 25
10 25 25 25 256
initial CSI data size:
                          10 25 25 25 256
final CSI data size:
```

Plot FISP noise characteristics

verify noise is uniformly distributed

reco time = 4.9 s
estimating field map(1/2)
estimating metabolities(2/2)
Metabolite mapping time = 8.0 s

```
figure,
set(gcf,'Visible','on','Position',[0,0,1400,800])
tiledlayout(2,5)
sig=permute(mcobj_fisp.twix.image{''},[2 1 3:8]);
sig=mcobj_fisp.D*sig(:,:); % noise decorrelation
sig=sig(:,abs(sum(sig,1))>0);%take all samples without zeropadding
nexttile(1), histogram(real(sig(:))), title(sprintf('data|real| \mu \pm \sigma = \%.2f \pm \%.2f', mean(real(sig(:))), std(real(sig(:)))));
nexttile(1+5),qqplot(real(siq(:))),title('data|real| 00');
\text{nexttile}(2), \text{histogram}(\text{imag}(\text{sig}(:))), \text{title}(\text{sprintf}(\text{'data}|\text{imag}|\mu\pm\sigma=\%.2\text{f}\pm\%.2\text{f}', \text{mean}(\text{imag}(\text{sig}(:))), \text{std}(\text{imag}(\text{sig}(:)))));
nexttile(2+5),qqplot(imag(sig(:))),title('data|imag| QQ');
% reconstructed images
nexttile(3), histogram(real(mcobj_fisp.img(:))), title(sprintf('Image|real|\mu \pm \sigma = \%.1f \pm \%.2f', mean(real(mcobj_fisp.img(:))), std(real(mcobj_fisp.img(:)))));
nexttile(3+5),qqplot(real(mcobj_fisp.img(:))),title('image|real|QQ');
\texttt{nexttile(4),histogram(imag(mcobj\_fisp.img(:))),title(sprintf('Image \mid imag \mid \mu \pm \sigma = \$.1f \pm \$.2f',mean(real(mcobj\_fisp.img(:))),std(real(mcobj\_fisp.img(:)))));}
nexttile(4+5),qqplot(imag(mcobj_fisp.img(:))),title('image | imag | QQ')
%Metabolite maps
[~,scl_fac]=mcobj_fisp.getNormalized();
% scl_fac is the eucledian norm of complex weights used during spectral
% seperation is used for normalization to SNR units.
Metcon_SNR=real(reshape(mcobj_fisp.Metcon,[],4)./scl_fac')/sqrt(2);
nexttile(5), histogram(Metcon_SNR(:)), title(sprintf('MetCon| real|\mu \pm \sigma = \%.2f \pm \%.2f', mean(Metcon_SNR(:)), std(Metcon_SNR(:))));
nexttile(5+5),qqplot(Metcon_SNR(:)),title('MetCon|real| QQ');
```



CSI-bSSFP dataset with acquistion weighting

```
%data path
sn='/ptmp/pvalsala/deuterium/dataForPublication/phantom-DMI-Noise';

fn=fullfile(sn,'meas_MID00749_FID11716_rpcsi_ssfp_Stan25_15_6mm_5mins_0FA.dat');
metabolites=getMetaboliteStruct('phantom',0);

CSI_setting={'metabolites',metabolites,'parfor',true,...
    'doCoilCombine','adapt1','doZeropad',[0.5 0.5 0.5 0]*0,'mask',[],'Solver','IDEAL-modes'};
mcobj_ssfp=MetCon_CSI(fn,CSI_setting{:});

Software version: VD (!?)
```

```
Software version: VD (!?)
Reader version: 1660732089 (UTC: 17-Aug-2022 10:28:09)
Scan 1/1, read all mdhs:
    163.8 MB read in 20 s
Using T2* =20 ms
starting reco
initial CSI data size: 10 25 25 25 64 4
final CSI data size: 10 25 25 25 64 4
reco time = 3.7 s
estimating field map(1/2)
estimating metabolities(2/2)
Metabolite mapping time = 5.9 s
```

Plot CSI-bSSFP noise characteristics

verify noise is uniformly distributed

```
figure, set(gcf,'Visible','on','Position',[0,0,1400,800]) tiledlayout(2,5)

%raw data sig=permute(mcobj_ssfp.twix.image{''},[2 1 3:8]); sig=mcobj_ssfp.D*sig(:,:); % noise decorrelation sig=sig(:,abs(sum(sig,1))>0):%take all samples without zeropadding

nexttile(1),histogram(real(sig(:))),title(sprintf('data|real| \pu+\sigma=\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\frac{\pi}{2}\fra
```

```
nexttile(3), histogram(real(mcobj_ssfp.img(:))), title(sprintf('Image|real|\puto=\frac{1}{2}.1ft\frac{1}{2}.2f', mean(real(mcobj_ssfp.img(:))), std(real(mcobj_ssfp.img(:))));

nexttile(3+5), qqplot(real(mcobj_ssfp.img(:))), title('image|real|\QQ');

nexttile(4), histogram(imag(mcobj_ssfp.img(:))), title(sprintf('Image|imag|\QQ'))

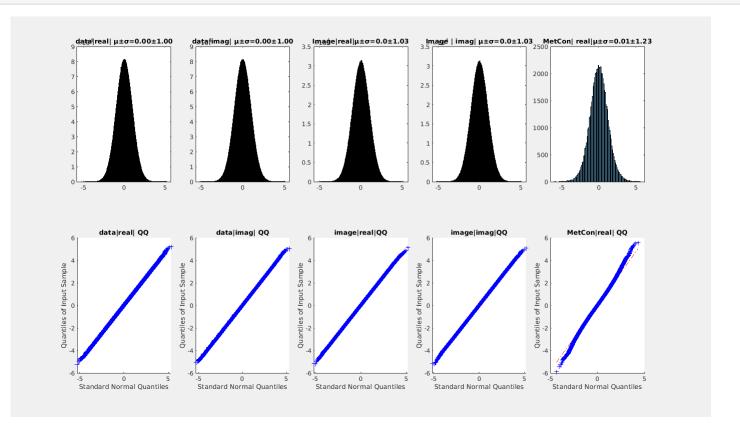
%Metabolite maps
[~,scl_fac]=mcobj_ssfp.getNormalized();

% scl_fac is the eucledian norm of complex weights used during spectral

% seperation is used for normalization to SNR units.

Metcon_SNR=real(reshape(mcobj_ssfp.Metcon,[],4)./scl_fac')/sqrt(2);

nexttile(5), histogram(Metcon_SNR(:)), title(sprintf('MetCon| real|\puto=\frac{1}{2}.2ft\frac{1}{2}.2f', mean(Metcon_SNR(:))), std(Metcon_SNR(:))));
```



multiecho-bSSFP dataset with acquistion weighting

nexttile(5+5),qqplot(Metcon_SNR(:)),title('MetCon real QQ');

assemble inputs and processing flags

%data path

Plot CSI-bSSFP noise characteristics

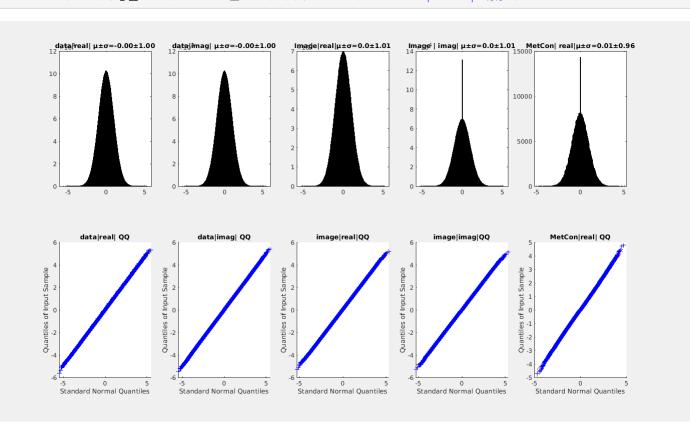
Metabolite mapping time = 22.4 s

Performed pixel shift along read: (0.0, -2.2, -5.2, -7.2) mm

estimating field map(1/2) estimating metabolities(2/2)

verify whether noise is uniformly distributed. The zeros added by the translation to compensate chemical shifts along read is visible in the metcon histogram

```
figure,
set(gcf,'Visible','on','Position',[0,0,1400,800])
tiledlayout(2,5)
%raw data
sig=permute(mcobj_ssfp.twix.image{''},[2 1 3:8]);
sig=mcobj_ssfp.D*sig(:,:); % noise decorrelation
sig=sig(:,abs(sum(sig,1))>0);%take all samples without zeropadding
nexttile(1), histogram(real(sig(:))), title(sprintf('data|real| \mu \pm \sigma = \%.2f \pm \%.2f', mean(real(sig(:)))), std(real(sig(:)))));
nexttile(1+5),qqplot(real(sig(:))),title('data|real| QQ');
nexttile(2), histogram(imag(sig(:))), title(sprintf('data|imag| \mu \pm \sigma = \%.2f \pm \%.2f', mean(imag(sig(:)))), std(imag(sig(:)))));
nexttile(2+5),qqplot(imag(sig(:))),title('data|imag| QQ');
% reconstructed images
nexttile(3), histogram(real(mcobj\_ssfp.img(:))), title(sprintf('Image|real|\mu \pm \sigma = \%.1f \pm \%.2f', mean(real(mcobj\_ssfp.img(:)))), std(real(mcobj\_ssfp.img(:)))));
nexttile(3+5),qqplot(real(mcobj_ssfp.img(:))),title('image|real|QQ');
nexttile(4), histogram(imag(mcobj\_ssfp.img(:))), title(sprintf('Image | imag | \mu \pm \sigma = \%.1f \pm \%.2f', mean(real(mcobj\_ssfp.img(:))), std(real(mcobj\_ssfp.img(:)))));
nexttile(4+5),qqplot(imag(mcobj_ssfp.img(:))),title('image imag | QQ')
%Metabolite maps
[~,scl_fac]=mcobj_ssfp.getNormalized();
% scl_fac is the eucledian norm of complex weights used during spectral
% seperation is used for normalization to SNR units.
Metcon_SNR=real(reshape(mcobj_ssfp.Metcon,[],4)./scl_fac')/sqrt(2);
nexttile(5), histogram(Metcon_SNR(:)), title(sprintf('MetCon| real|\mu \pm \sigma = \%.2f \pm \%.2f', mean(Metcon_SNR(:)), std(Metcon_SNR(:))));
nexttile(5+5),qqplot(Metcon_SNR(:)),title('MetCon|real| QQ');
```



PDF export 17 inch x 28 inch 600 dpi