## Code Examples

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## Chapter 2

## Section 1

```
set.seed(23)
P \leftarrow RM_norm(N = 5, mean = 0, sd = 1)
spectrum_P <- spectrum(P)</pre>
# Outputs the following
spectrum_P
##
          Re
                   Im
                         Norm Order
## 1 -0.5434 1.3539 1.4589
## 2 -0.5434 -1.3539 1.4589
                                  2
## 3 0.2255 1.4250 1.4427
                                  3
## 4 0.2255 -1.4250 1.4427
                                  4
## 5 -0.8678 0.0000 0.8678
\begin{lstlisting}[language=R]
library(RMAT)
P \leftarrow RM_{norm}(N = 5, mean = 0, sd = 1)
spectrum_P <- spectrum(P)</pre>
# Outputs the following
spectrum_P
\end{lstlisting}
```

Spectrum Dataframe			
Re	Im	Norm	Order
-0.5434	1.3539	1.4589	1
-0.5434	-1.3539	1.4589	2
0.2255	1.4250	1.4427	3
0.2255	-1.4250	1.4427	4
-0.8678	0.0000	0.8678	5

## Section 2

```
set.seed(23)
ens <- RME_norm(N = 3, mean = 0, sd = 1, size = 3)
spectrum_ens <- spectrum(ens)
# Outputs the following
spectrum_ens</pre>
```

```
## Re Im Norm Order
```

```
## 1 1.7581 0.0000 1.7581
## 2 -0.2614 1.0012 1.0347
                                2
## 3 -0.2614 -1.0012 1.0347
                                3
## 4 1.2327 0.4227 1.3032
                                1
## 5  1.2327 -0.4227 1.3032
                                2
## 6 -0.8504 0.0000 0.8504
                                3
## 7 -0.5296 1.0508 1.1767
                                1
## 8 -0.5296 -1.0508 1.1767
                                2
## 9 0.7357 0.0000 0.7357
\begin{lstlisting}[language=R]
library(RMAT)
P \leftarrow RM_norm(N = 5, mean = 0, sd = 1)
spectrum_P <- spectrum(P)</pre>
# Outputs the following
spectrum_P
\end{lstlisting}
```