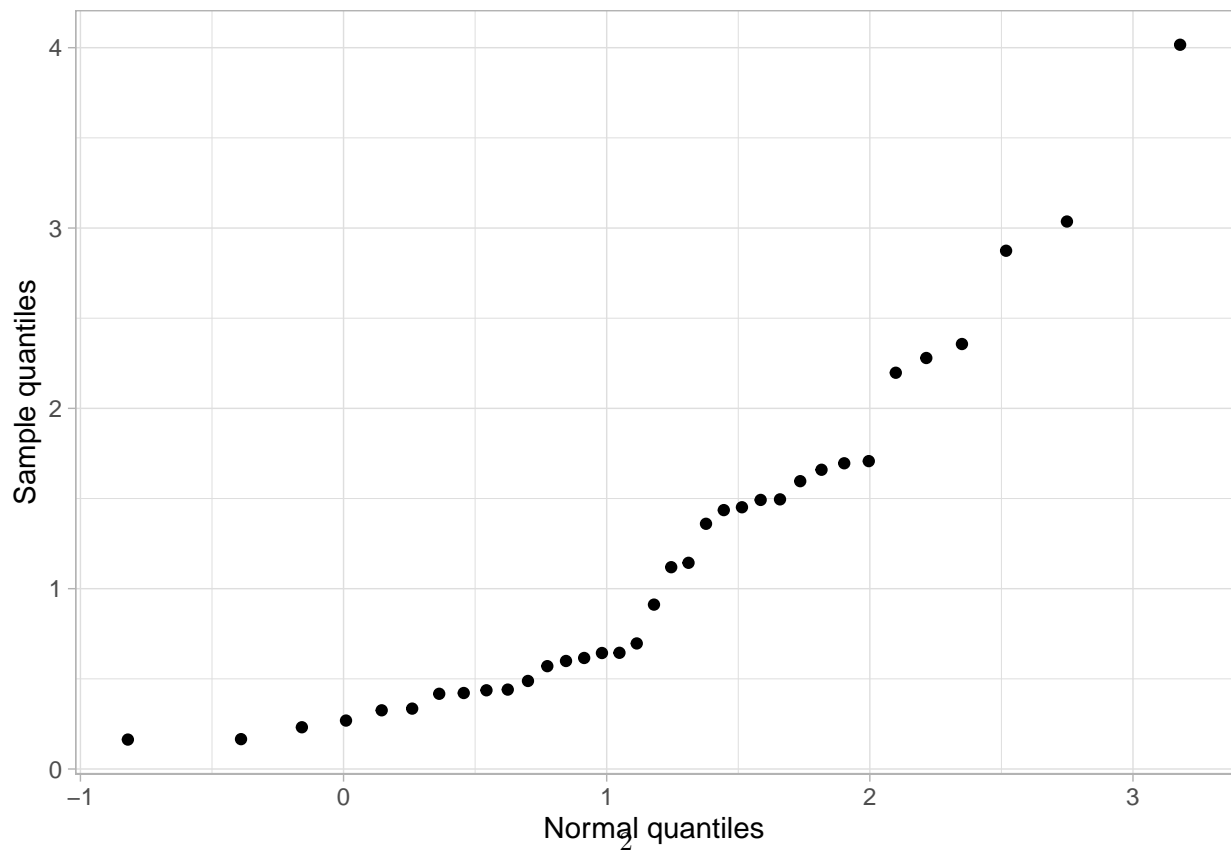
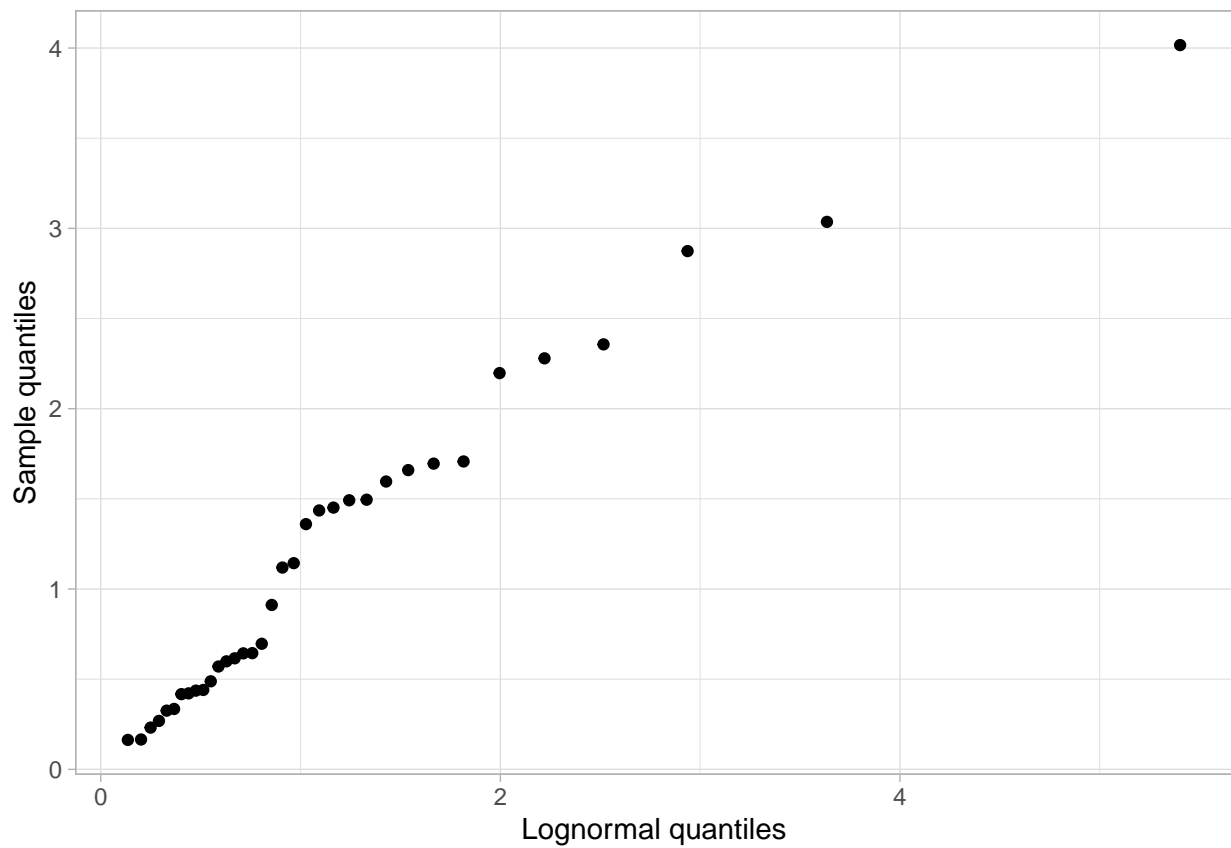
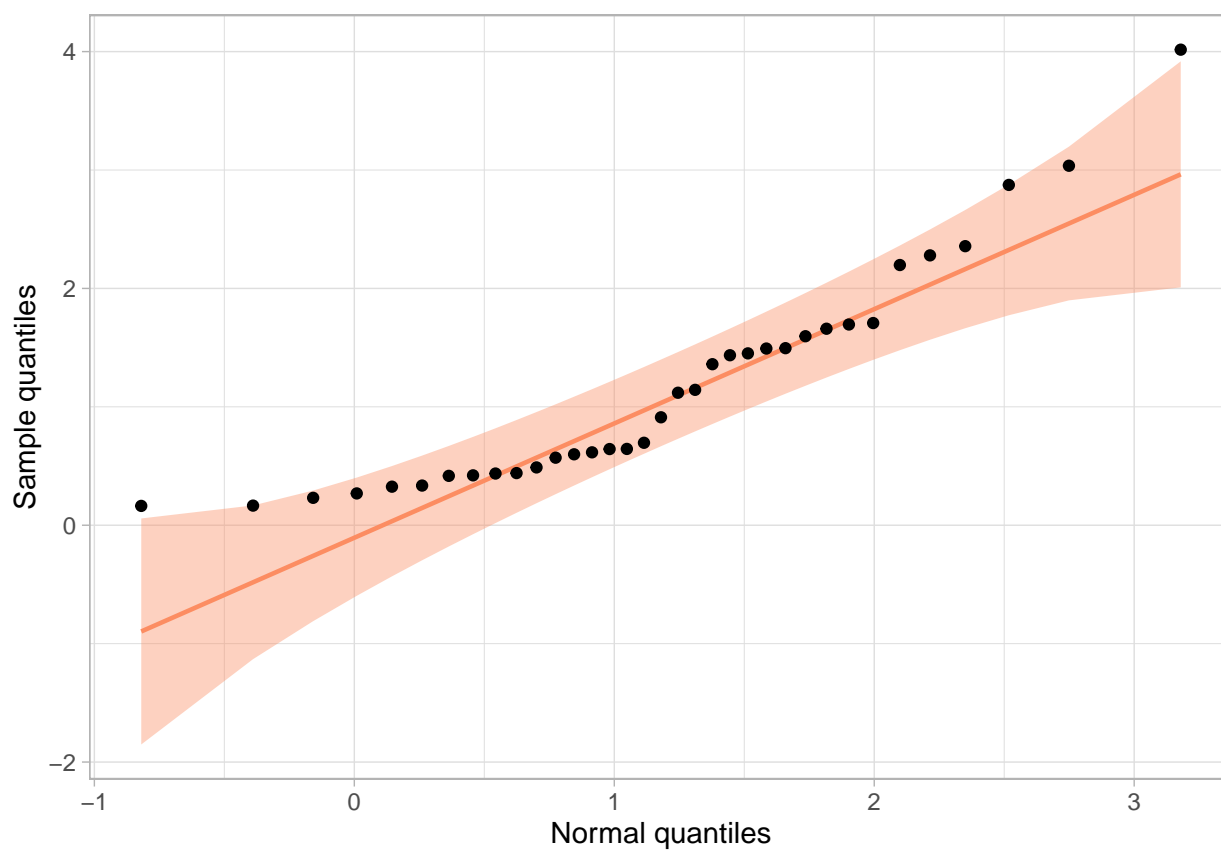
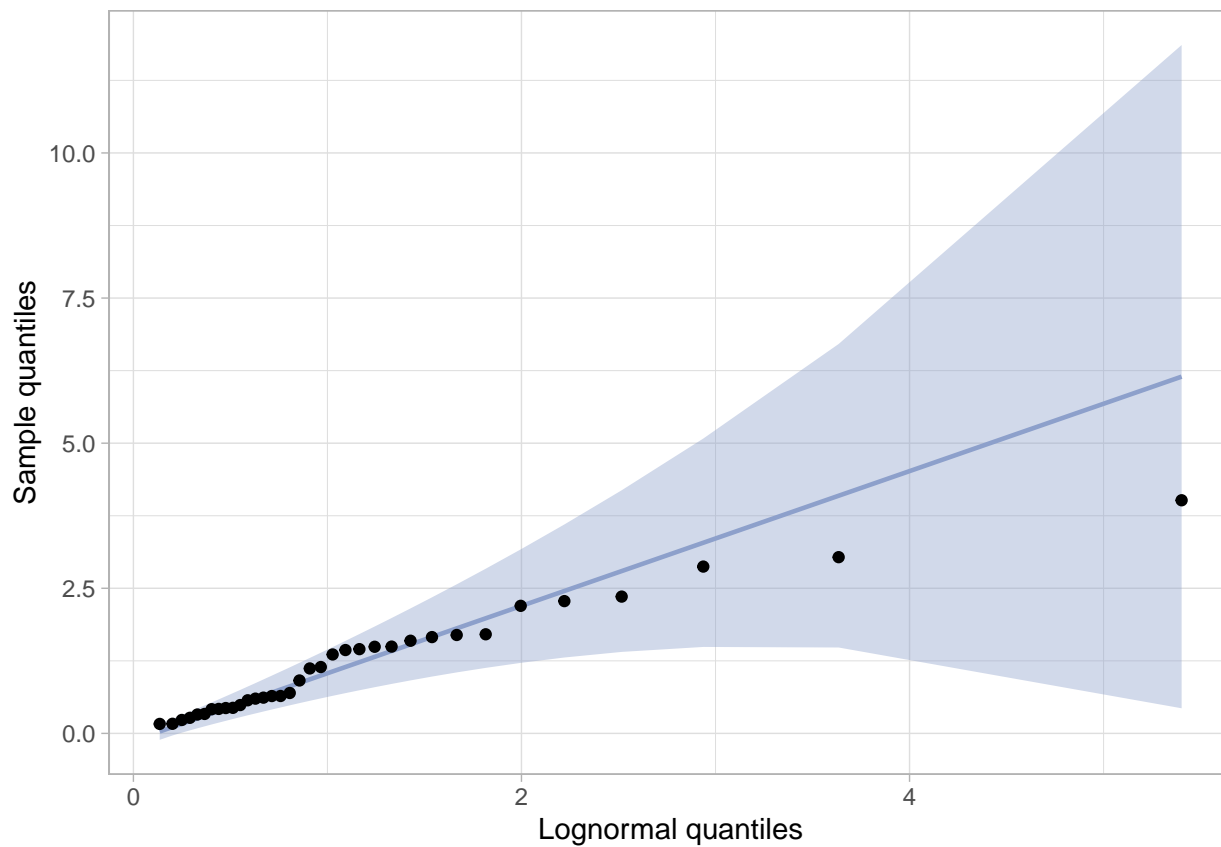


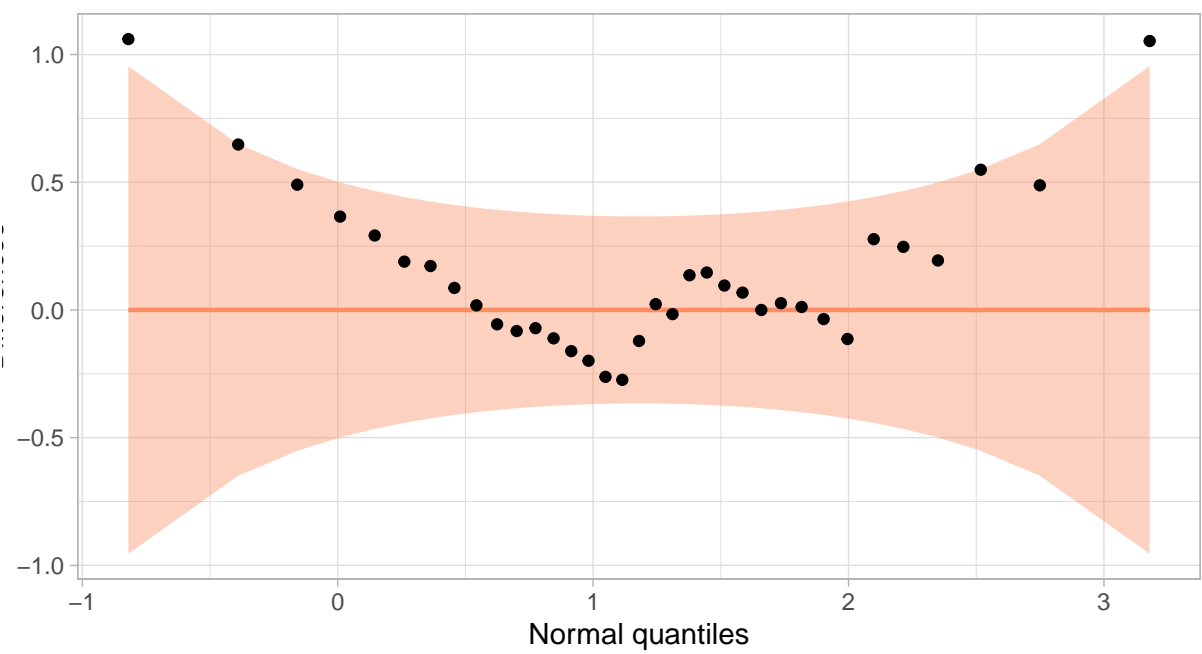
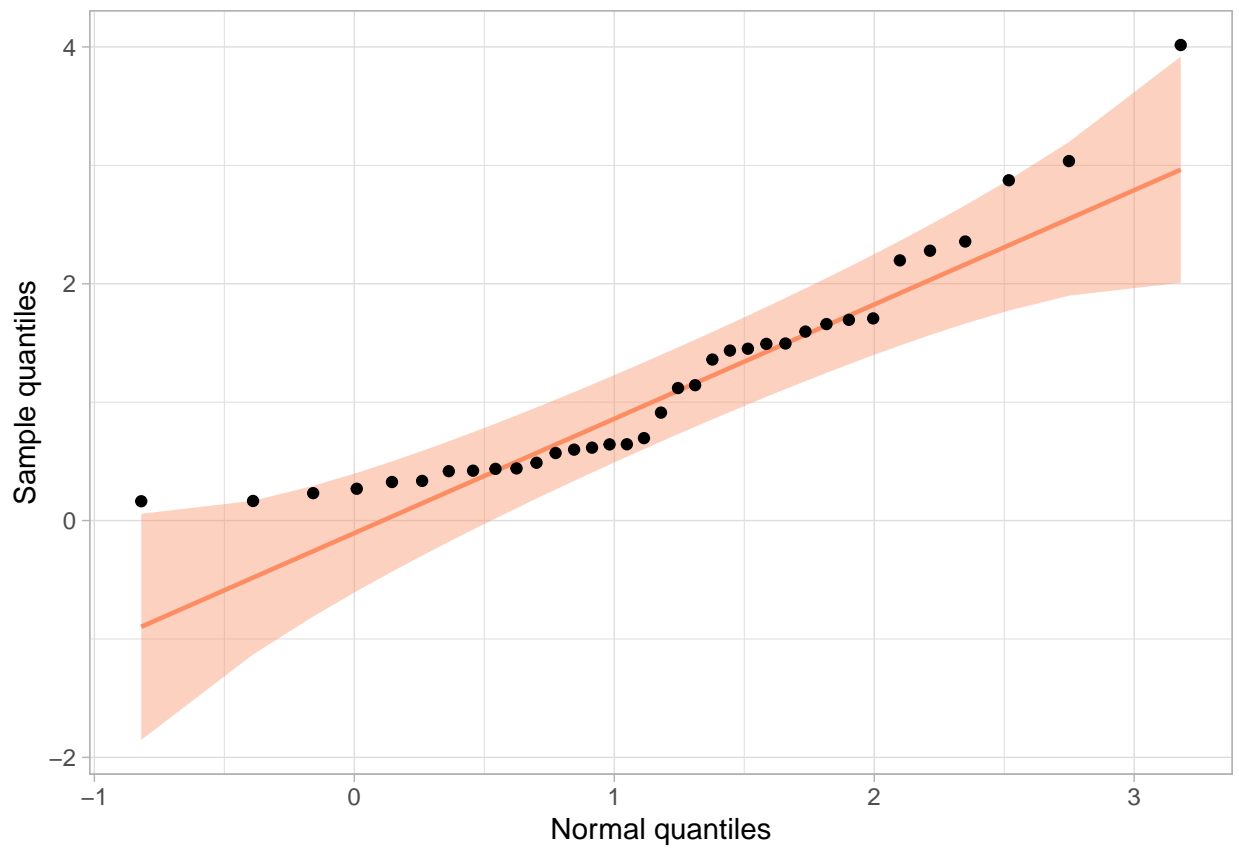
EljasiakBartlomiej

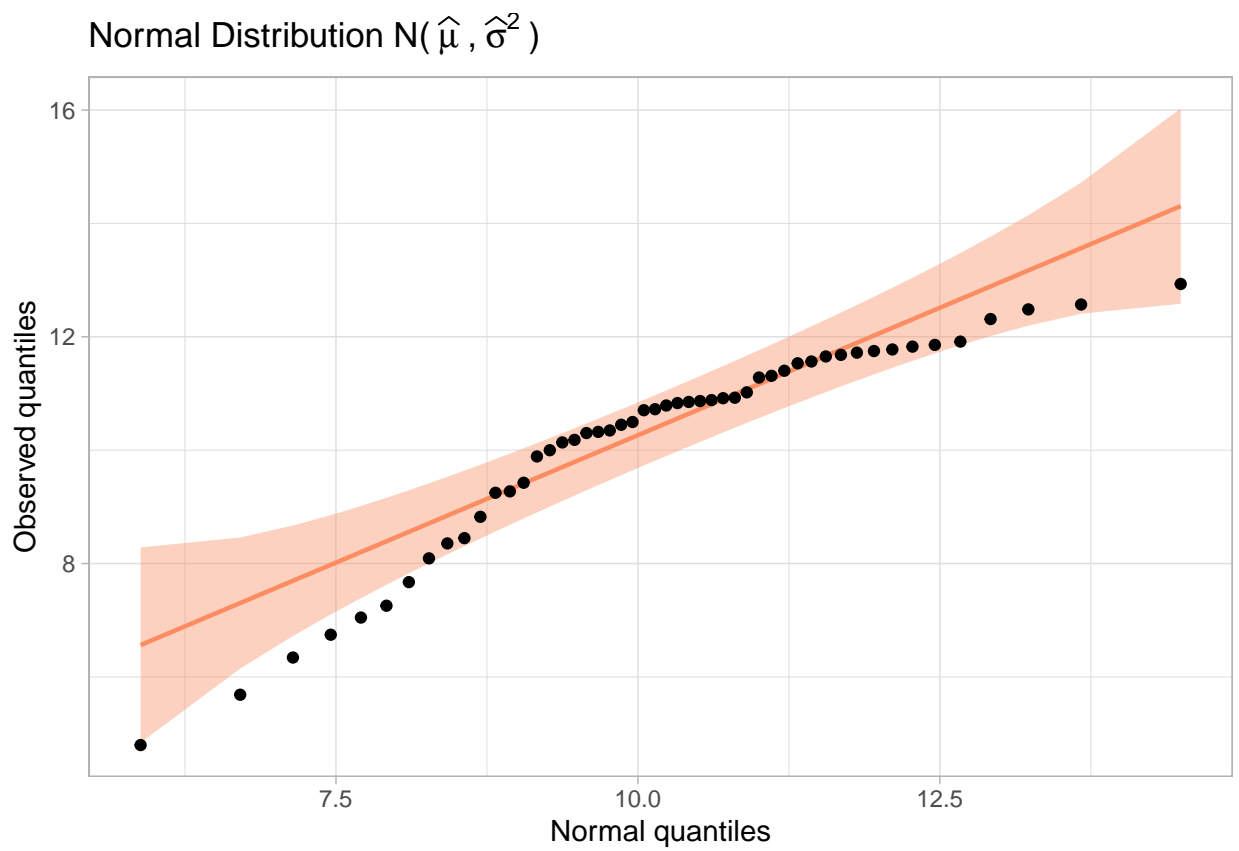
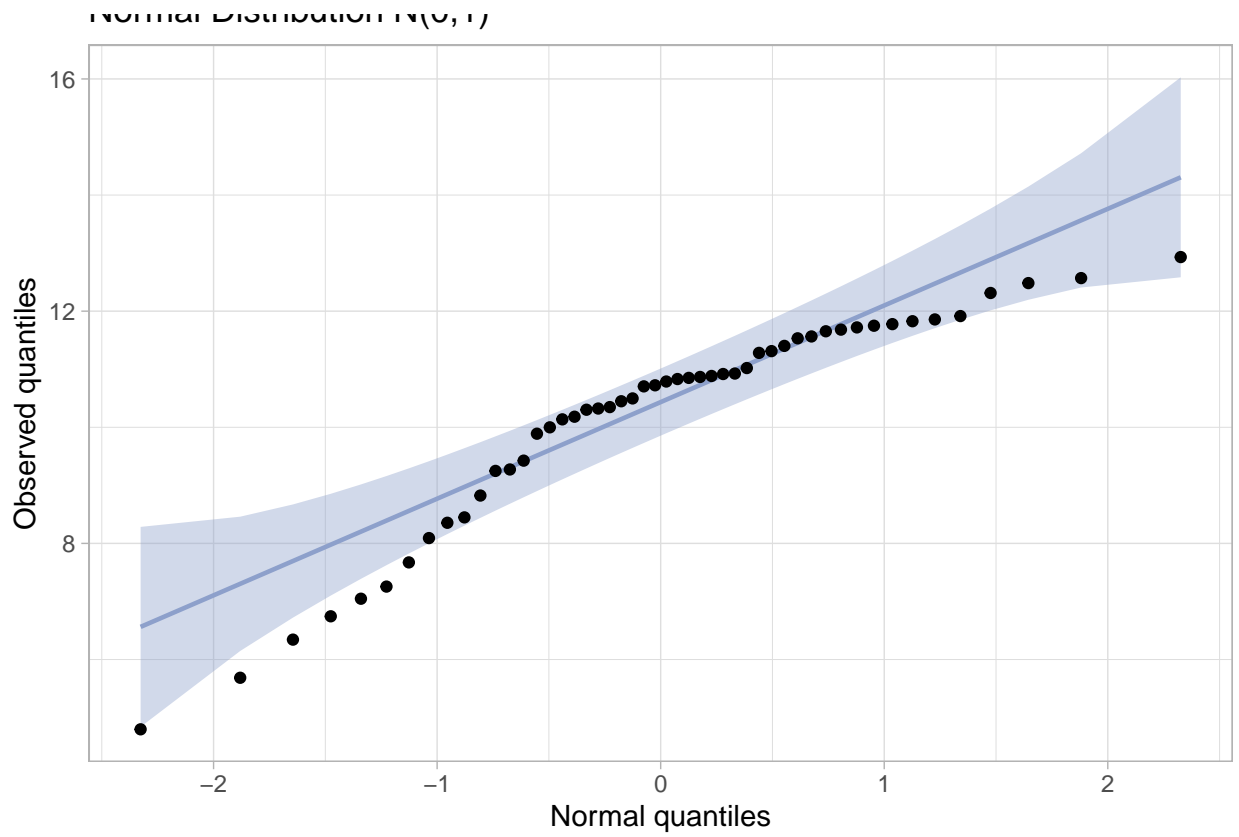
ggplot2 Compatible Quantile-Quantile Plots in R

reprodukowalny

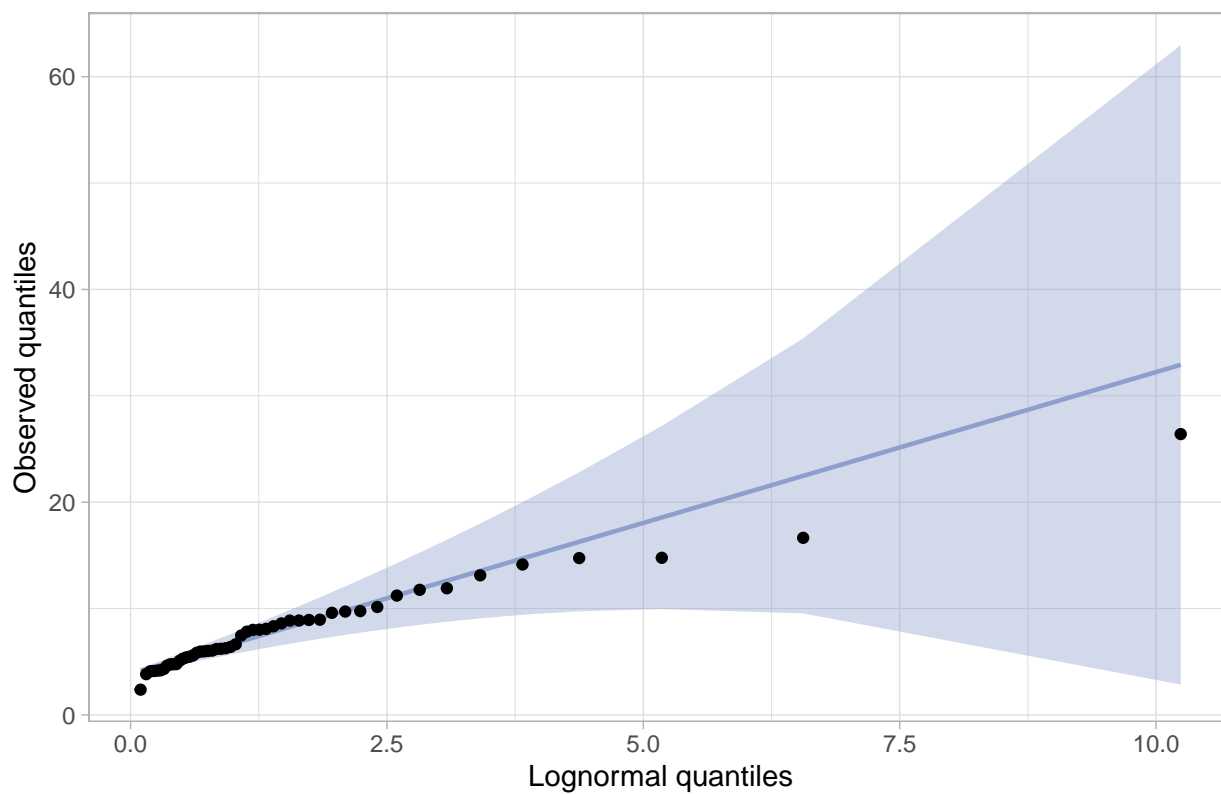




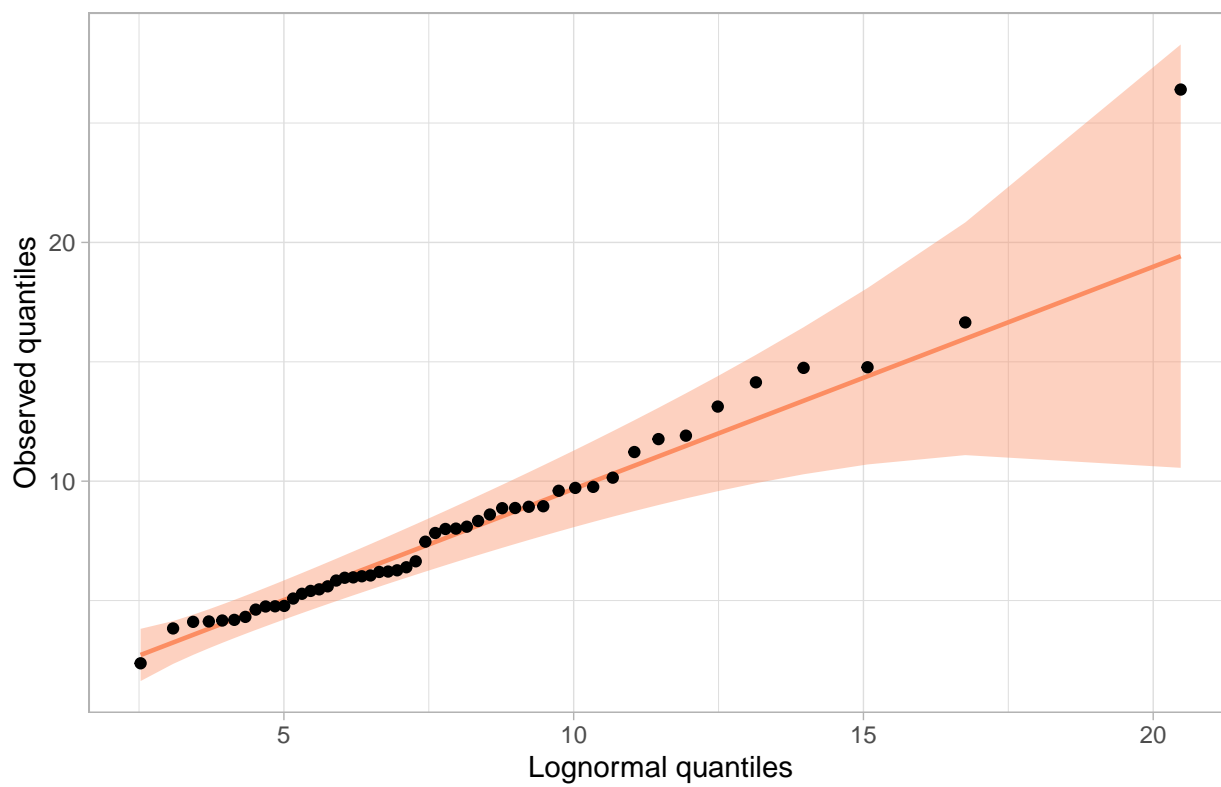


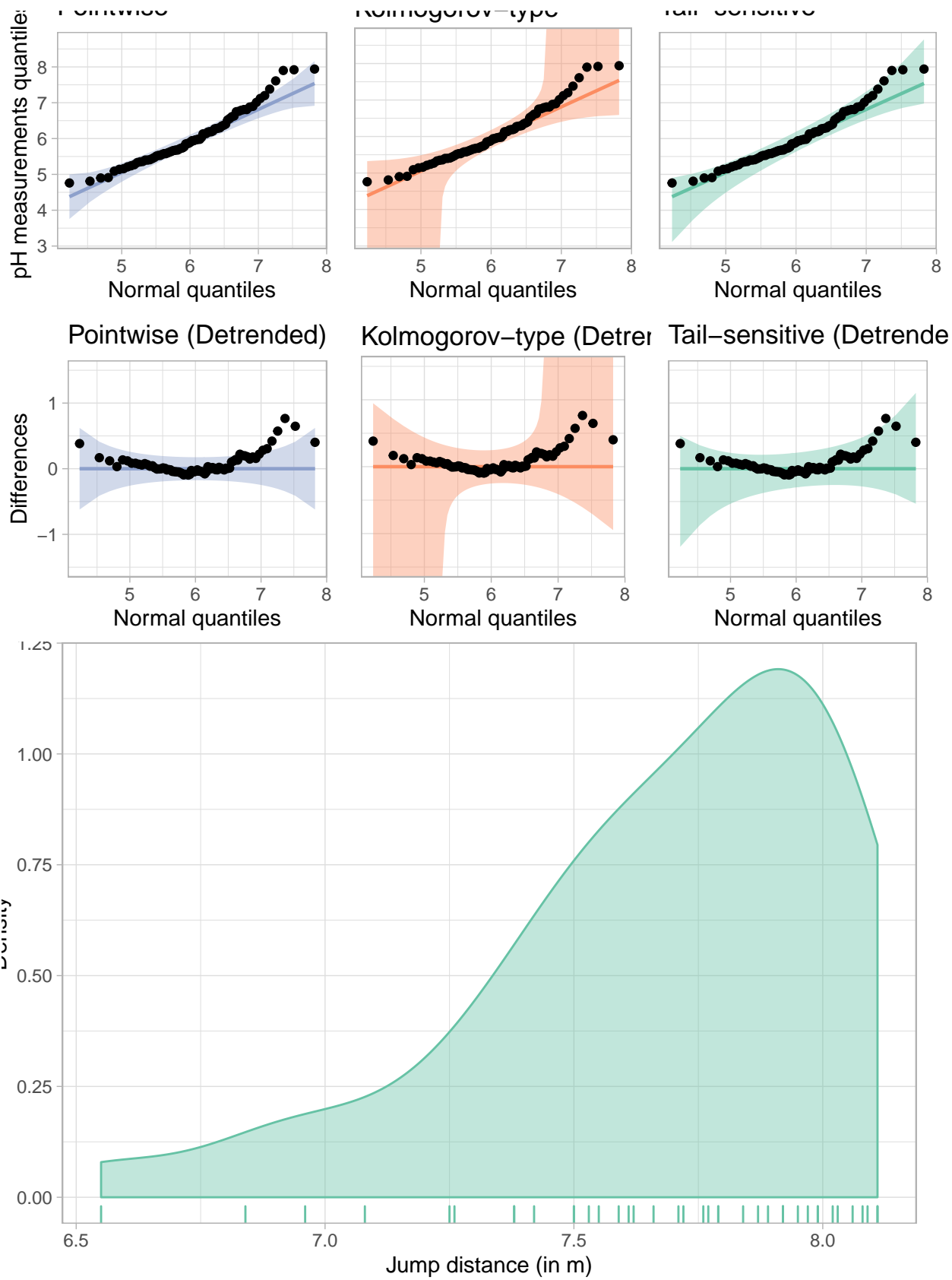


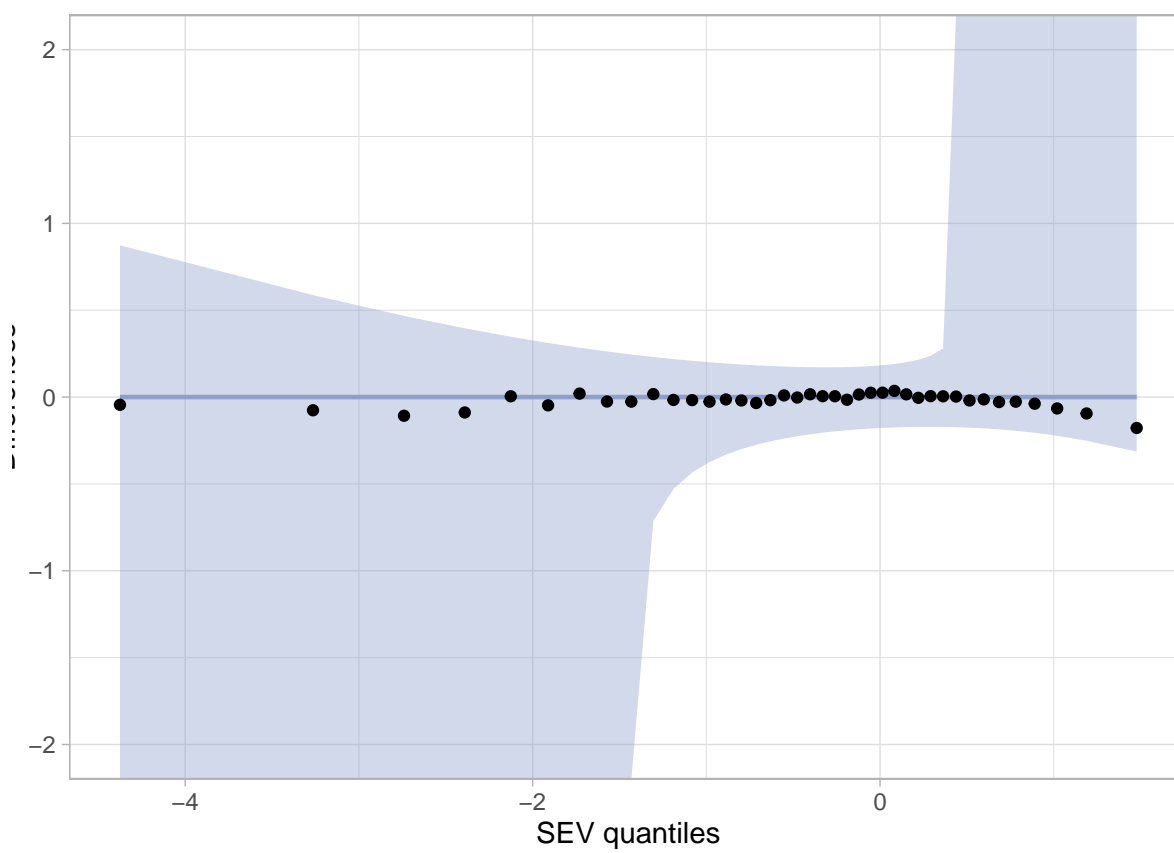
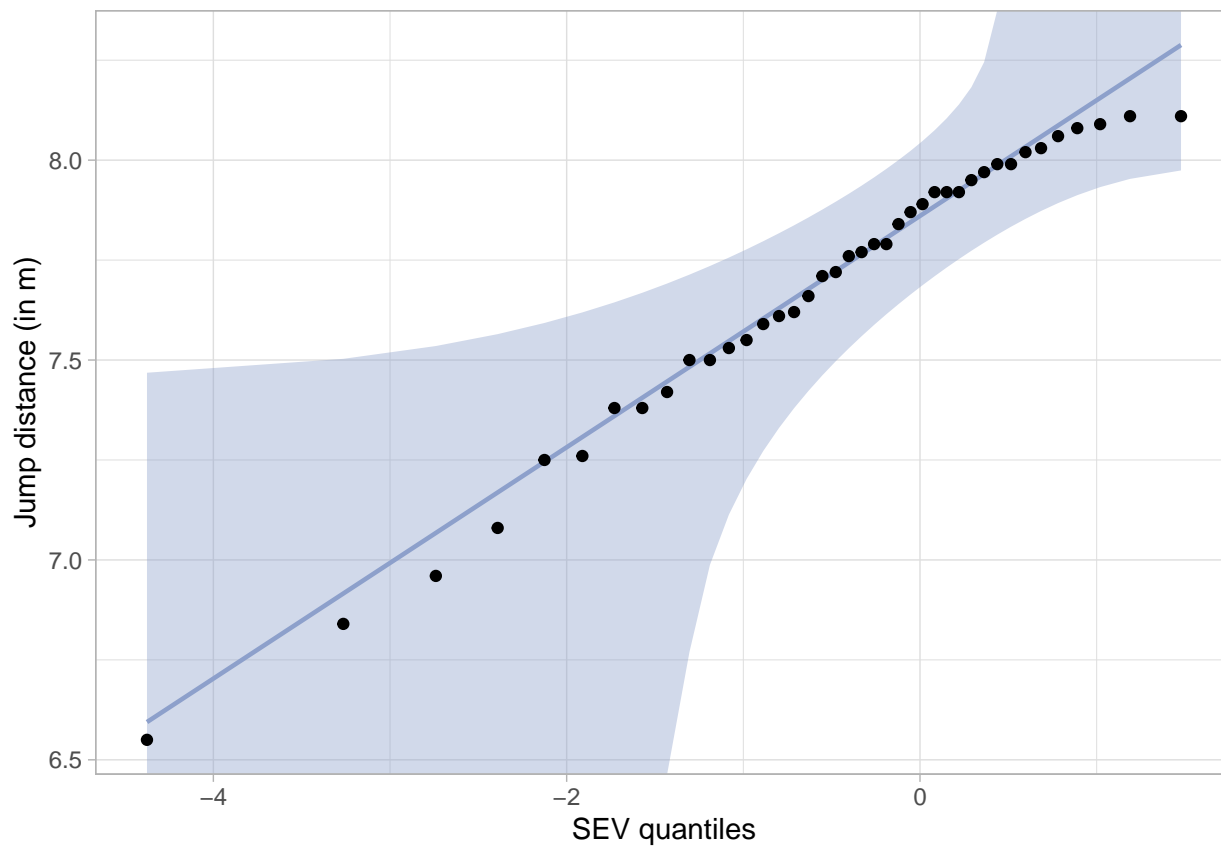
Lognormal Distribution $LN(0, 1)$

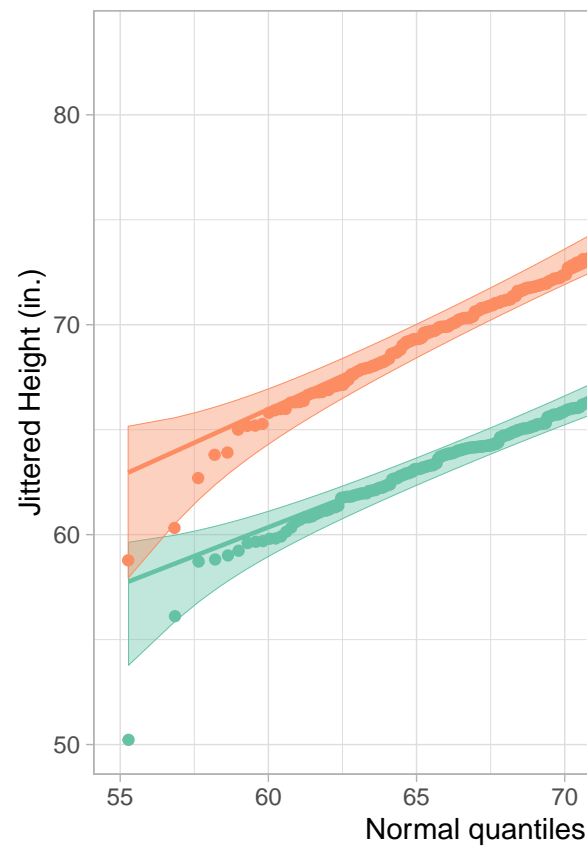
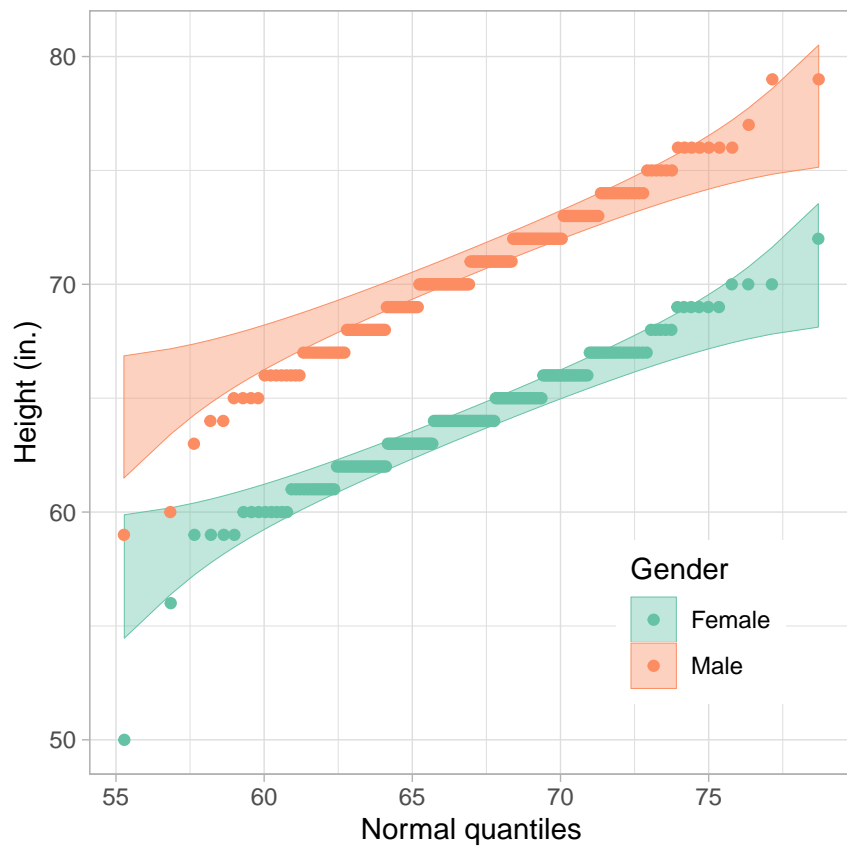


Lognormal Distribution $LN(\hat{\mu}, \hat{\sigma}^2)$

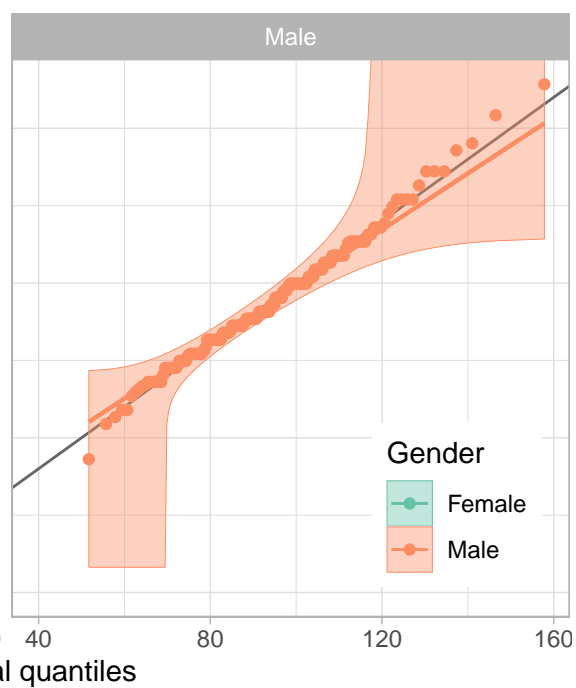
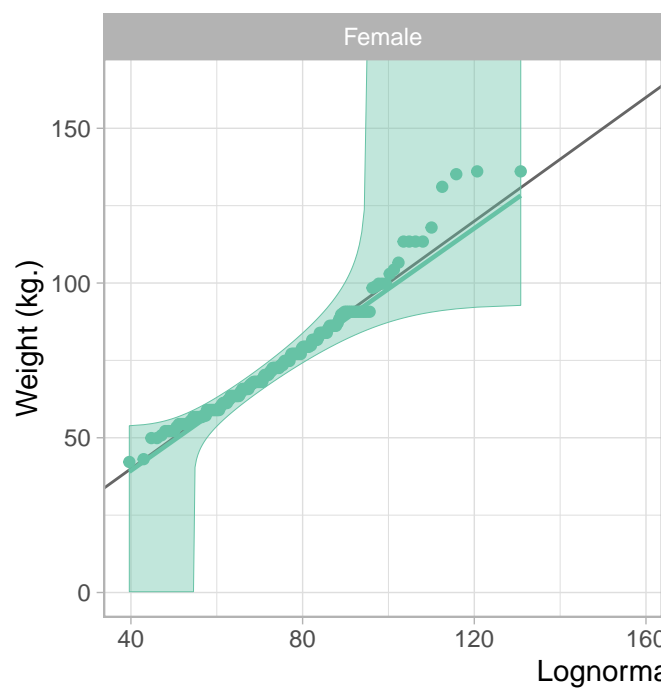
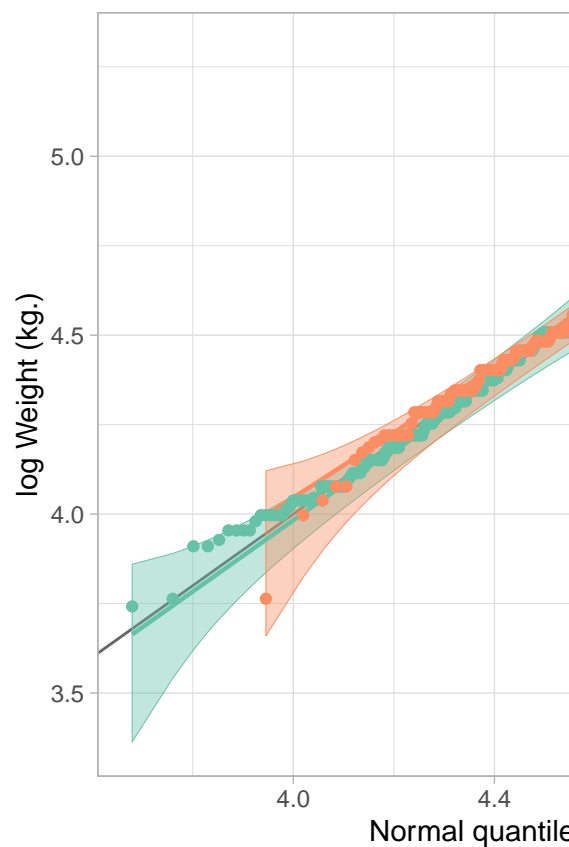
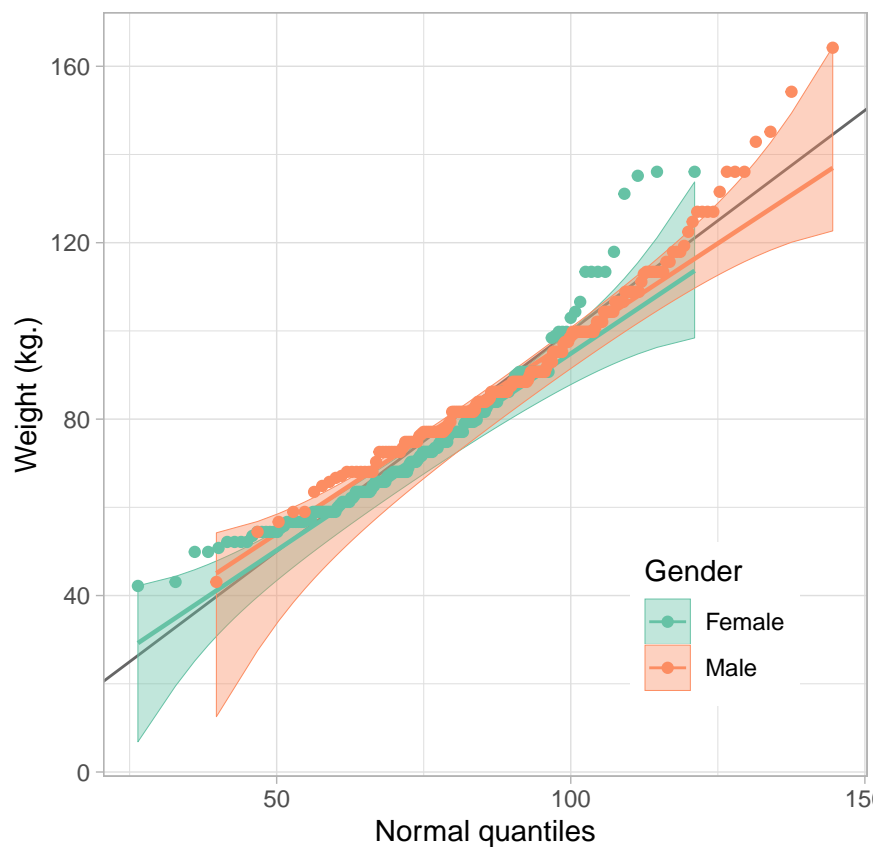








SEX	mean height (in)	sd (in)	mean log weight (kg)	sd (kg)
Male	70.55043	2.966460	9.100936	0.2011216
Female	64.50972	2.911454	8.887950	0.2254663
Total	66.99109	4.176295	8.978768	0.2397852



Dot-Pipe: an S3 Extensible Pipe for R

niereprodukowalny!!!

```
## [1] -0.9589243
```

```
## [1] 5
```

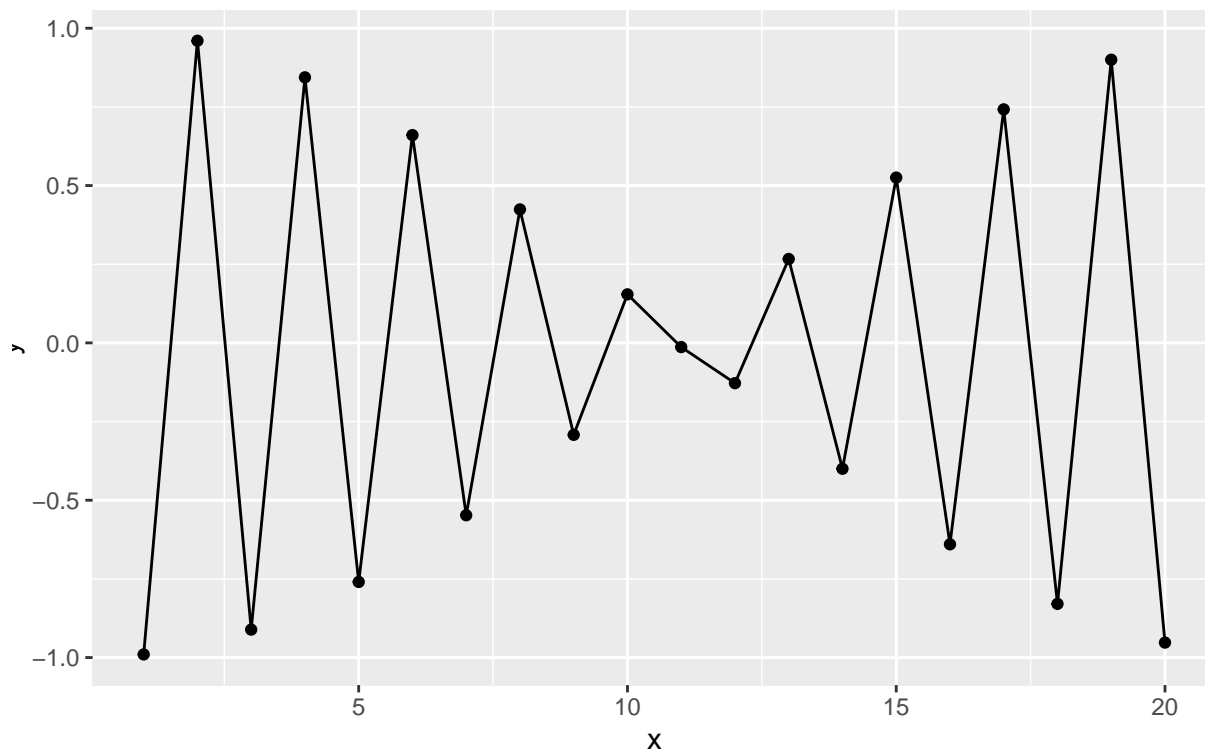
```
## [1] 6
```

```
## [1] 6
```

```
## [1] 11
```

```
  piped ggplot2
```

```
  wrapr
```



```
## [1] "relop_extend" "relop"
```

```
## [1] "mk_td(\"d\", c( \"x\")) %>% extend(., y := cos(2 * x))"
```

```
## [1] " BŁĄD !!! Powinno być 'table(d; x) %>% extend(., y := cos(2 * x))'"
```

```
## [1] "x" "y"
```

```
## $d
```

```
## [1] "x"
```

```
##   x       y
```

```
## 1 1 -0.4161468
```

```
## 2 2 -0.6536436
```

```
## 3 3  0.9601703
```

```
## wrapr::apply_right_S4 default called with classes:
```

```
## d1 data.frame
```

```
## d2 data.frame
```

```

## must have a more specific S4 method defined to dispatch
## NULL

## x
## 1 1
## 2 2

## x
## 1 2

## [1] -0.9589243
## <simpleError in pipes[[i]]: subscript out of bounds>
## [1] -0.9589243
## <simpleError in .::base: unused argument (sin)>
## [1] 5
## [1] -0.9589243

##      1      2
## 0.9428669 0.6508301

## [1] type      name      tbl_name rootpage sql
## <0 rows> (or 0-length row.names)

## [1] "abc"

## y ~ a + b + c + d + e

```