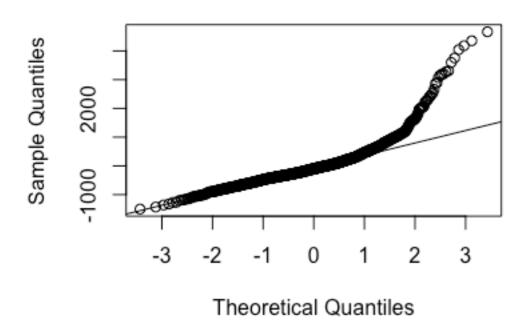
• Also need to check the residuals at the random effects level using the ranef() function to extract the random effects parameters.

```
model.full <- lmer(RT ~ Context * Sentence + (1 + Context + Sentence | Subject) +
                      (1 + Context + Sentence | Item), data = DV, REML = TRUE)
qqnorm(residuals(model.full))
summary(model.full)
# checking residuals
r int <- lme4::ranef(model.full)$Subject$`(Intercept)`</pre>
qqnorm(r int)
qqline(r int)
r slope <- lme4::ranef(model.full)$Subject$Context1</pre>
qqnorm(r slope)
qqline(r slope)
r slope <- lme4::ranef(model.full)$Subject$Sentence1
qqnorm(r slope)
qqline(r slope)
```

• This is for a crossed design - for nested random effects the above code will need a little modification...

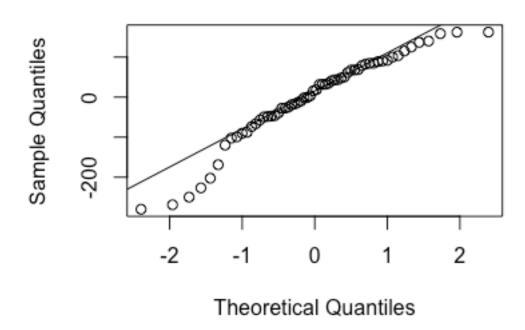
## Overall residuals

## **Normal Q-Q Plot**



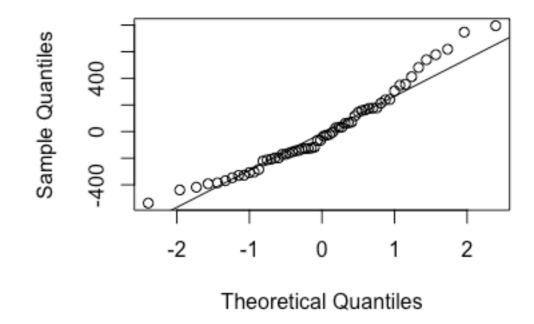
Factor 1 residuals

## Normal Q-Q Plot



# Intercept residuals

## **Normal Q-Q Plot**



Factor 2 residuals

#### Normal Q-Q Plot

