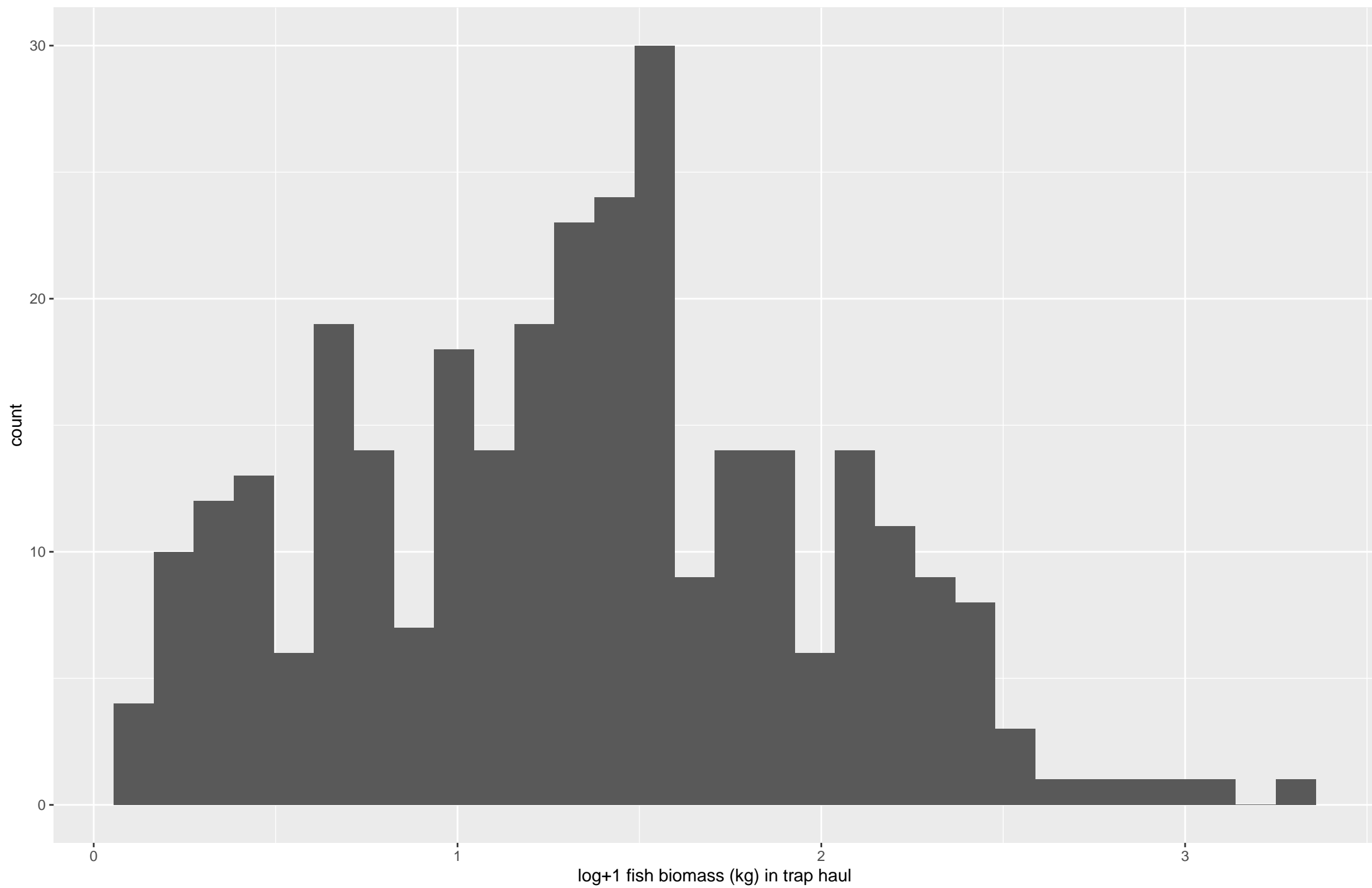
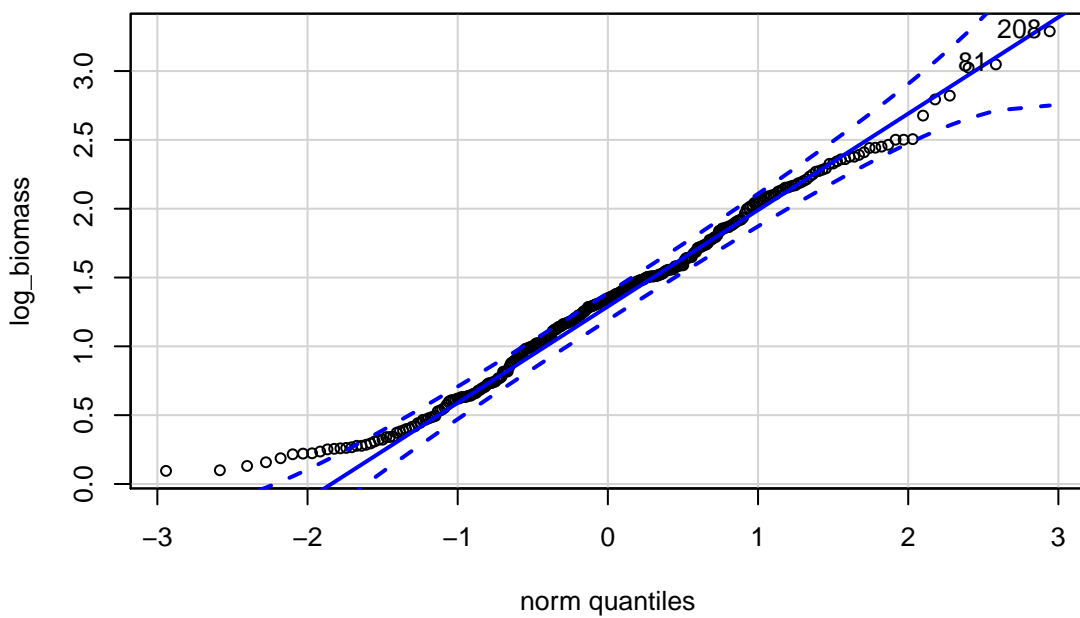


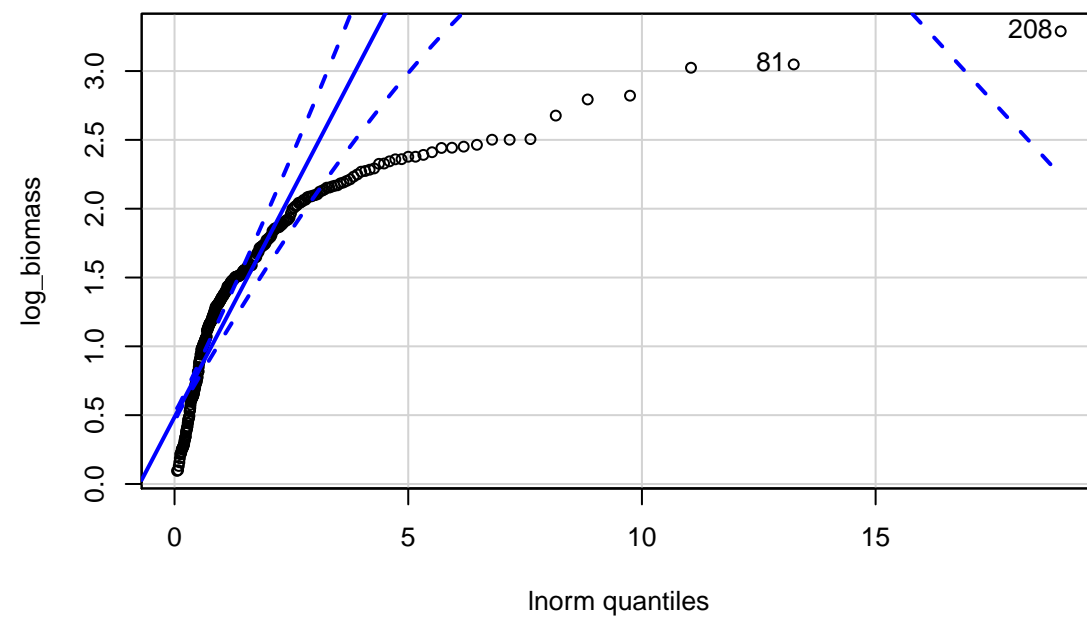
Histogram of log+1 fish biomass (kg)



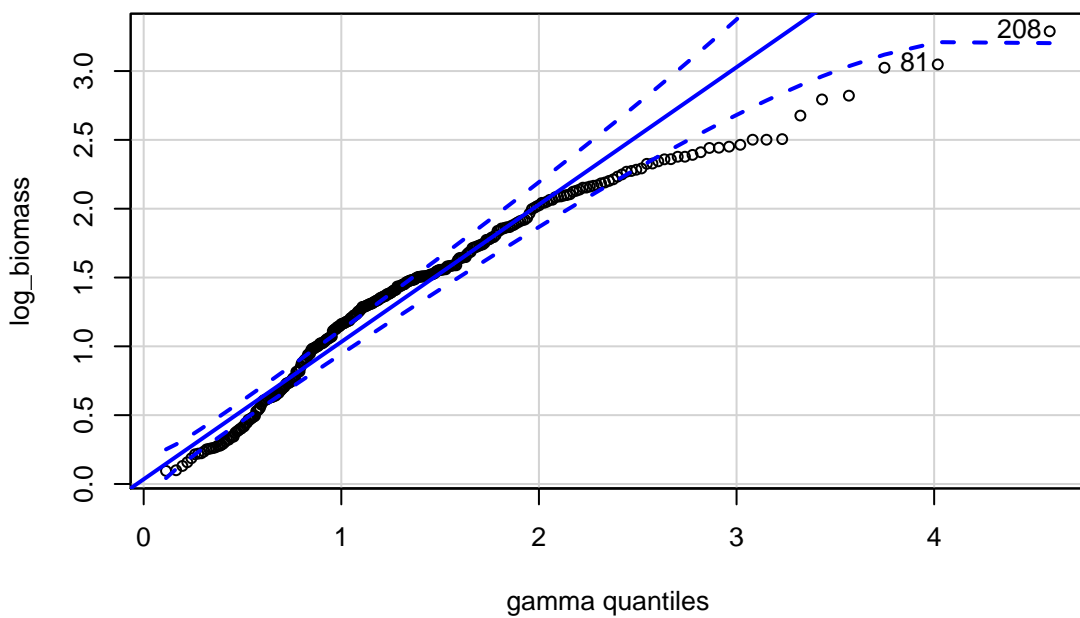
**Normal distribution**



**Log-normal distribution**



**Gamma distribution**



```
Global model call: lmer(formula = log_biomass ~ design + log_days_since_last_haul +
  location_exposure + Exp_or_Cont + (1 | TrapID) + (1 | Date_YMD),
  data = trap_haul_no_zero, REML = FALSE, na.action = "na.fail")
```

```
---
```

```
Model selection table
```

	(Int)	dsg	Exp_or_Cont	lct_exp	log_dys_snc_lst_hal	df	logLik	AICc	delta	weight
9	0.5791					0.3165	5 -273.856	557.9	0.00	0.244
13	0.6096			+		0.3173	6 -272.893	558.1	0.16	0.226
10	0.4430	+				0.3339	6 -273.164	558.6	0.70	0.172
14	0.5414	+		+		0.3249	7 -272.806	560.0	2.07	0.087
11	0.5777		+			0.3165	6 -273.856	560.0	2.08	0.086
15	0.6089		+	+		0.3173	7 -272.893	560.2	2.25	0.079
12	0.4432	+	+			0.3339	7 -273.164	560.7	2.79	0.060
16	0.5414	+	+	+		0.3249	8 -272.806	562.1	4.18	0.030
1	1.2930					4	-278.982	566.1	8.19	0.004
5	1.3240			+		5	-278.041	566.3	8.37	0.004
2	1.2430	+				5	-278.804	567.8	9.90	0.002
3	1.2910		+			5	-278.982	568.2	10.25	0.001
6	1.3550	+		+		6	-278.008	568.3	10.39	0.001
7	1.3230		+	+		6	-278.041	568.4	10.45	0.001
4	1.2420	+	+			6	-278.804	569.9	11.98	0.001
8	1.3540	+	+	+		7	-278.008	570.4	12.48	0.000

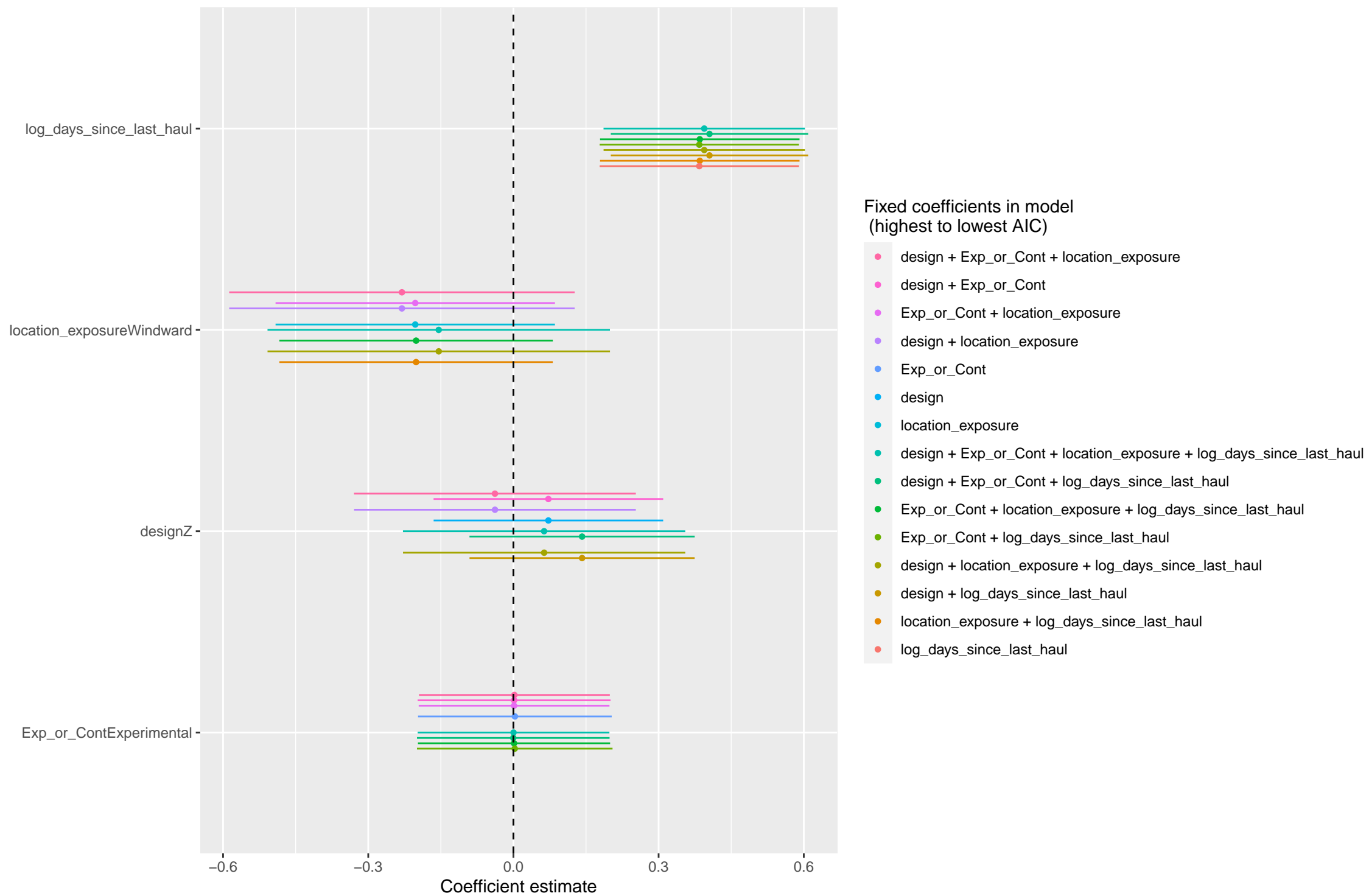
```
Models ranked by AICc(x)
```

```
Random terms (all models):
```

```
...1 | TrapID..., ...1 | Date_YMD...
```

	<b>model</b>	<b>sigma</b>	<b>logLik</b>	<b>AIC</b>	<b>BIC</b>	<b>deviance</b>	<b>df.residual</b>
	log_days_since_last_haul	0.54	-273.86	557.71	576.35	547.71	302
	location_exposure + log_days_since_last_haul	0.54	-272.89	557.79	580.15	545.79	301
	design + log_days_since_last_haul	0.55	-273.16	558.33	580.69	546.33	301
	design + location_exposure + log_days_since_last_haul	0.54	-272.81	559.61	585.70	545.61	300
	Exp_or_Cont + log_days_since_last_haul	0.54	-273.86	559.71	582.07	547.71	301
	Exp_or_Cont + location_exposure + log_days_since_last_haul	0.54	-272.89	559.79	585.87	545.79	300
	design + Exp_or_Cont + log_days_since_last_haul	0.55	-273.16	560.33	586.42	546.33	300
	design + Exp_or_Cont + location_exposure + log_days_since_last_haul	0.54	-272.81	561.61	591.43	545.61	299
	none	0.54	-278.98	565.96	580.87	557.96	303
	location_exposure	0.54	-278.04	566.08	584.72	556.08	302
	design	0.54	-278.80	567.61	586.24	557.61	302
	Exp_or_Cont	0.54	-278.98	567.96	586.60	557.96	302
	design + location_exposure	0.54	-278.01	568.02	590.38	556.02	301
	Exp_or_Cont + location_exposure	0.54	-278.04	568.08	590.44	556.08	301
	design + Exp_or_Cont	0.54	-278.80	569.61	591.97	557.61	301
	design + Exp_or_Cont + location_exposure	0.54	-278.01	570.02	596.10	556.02	300

# Predicting log+1 fish biomass (kg) in trap haul



Backward reduced random-effect table:

	Eliminated	npar	logLik	AIC	LRT	Df	Pr(>Chisq)
<none>		8	-272.81	561.61			
(1   TrapID)	0	7	-281.77	577.54	17.929	1	2.293e-05 ***
(1   Date_YMD)	0	7	-278.97	571.95	12.337	1	0.0004442 ***

---  
 Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

Backward reduced fixed-effect table:

Degrees of freedom method: Satterthwaite

	Eliminated	Sum Sq	Mean Sq	NumDF	DenDF	F value	Pr(>F)
Exp_or_Cont	1	0.0000	0.0000	1	34.841	0.0000	0.998222
design	2	0.0538	0.0538	1	79.167	0.1817	0.671040
location_exposure	3	0.5756	0.5756	1	90.986	1.9476	0.166240
log_days_since_last_haul	0	3.9617	3.9617	1	23.238	13.3568	0.001304 **

---  
 Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

Model found:

log\_biomass ~ log\_days\_since\_last\_haul + (1 | TrapID) + (1 |  
 Date\_YMD)

## Optimal model found using 'step' function which performs backward elimination of fixed-effect terms

```
Linear mixed model fit by maximum likelihood . t-tests use Satterthwaite's method ['lmerModLmerTest']
Formula: log_biomass ~ log_days_since_last_haul + (1 | TrapID) + (1 | Date_YMD)
Data: trap_haul_no_zero
```

AIC	BIC	logLik	deviance	df.resid
557.7	576.3	-273.9	547.7	302

Scaled residuals:

Min	1Q	Median	3Q	Max
-2.19806	-0.73098	-0.03229	0.65968	2.82404

Random effects:

Groups	Name	Variance	Std.Dev.
TrapID	(Intercept)	0.04657	0.2158
Date_YMD	(Intercept)	0.04001	0.2000
Residual		0.29660	0.5446

Number of obs: 307, groups: TrapID, 37; Date\_YMD, 23

Fixed effects:

	Estimate	Std. Error	df	t value	Pr(> t )
(Intercept)	0.5791	0.2067	26.4505	2.802	0.00937 **
log_days_since_last_haul	0.3165	0.0866	23.2383	3.655	0.00130 **

---

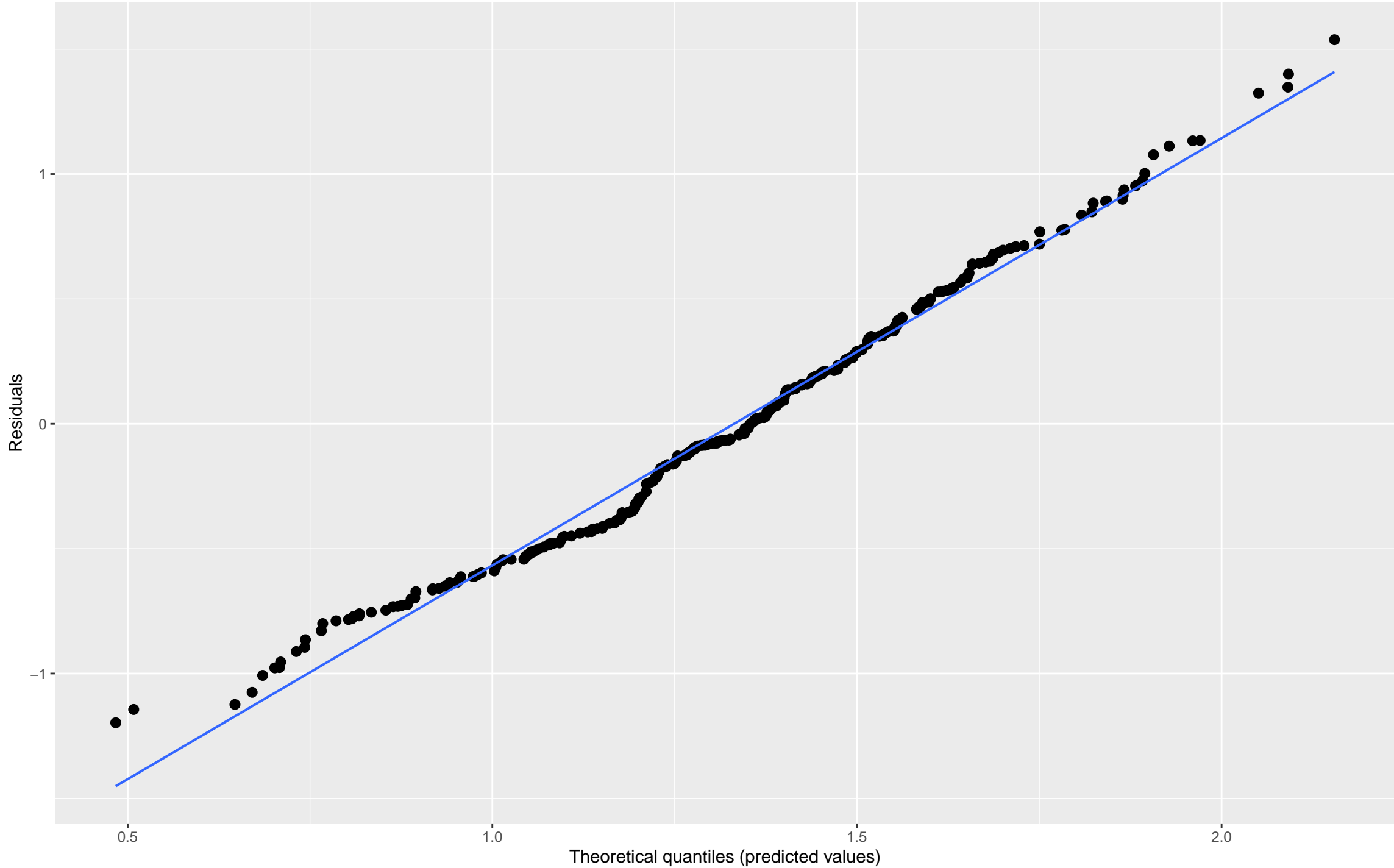
Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

Correlation of Fixed Effects:

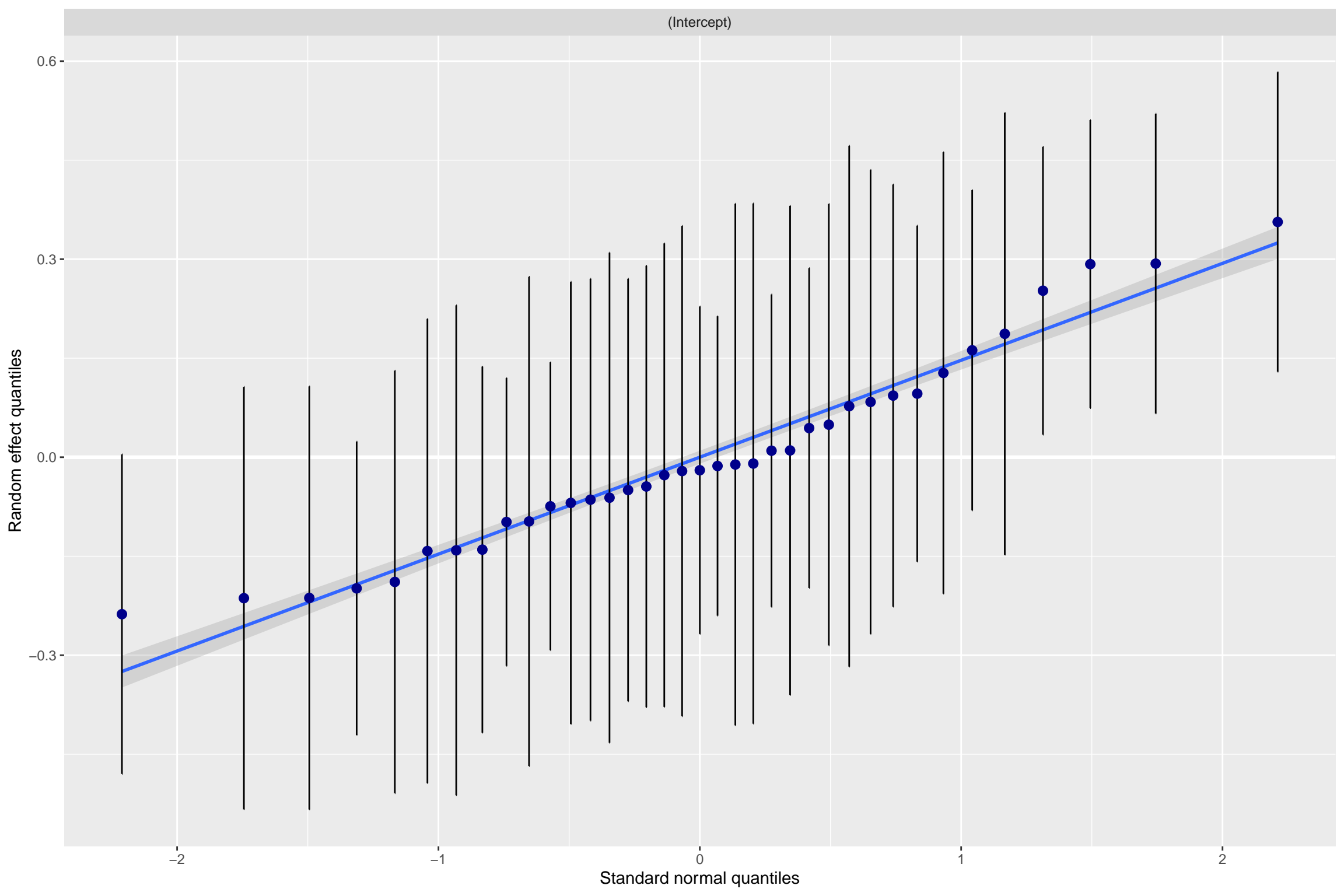
	(Intr)
lg_dys_sn__	-0.946

Non-normality of residuals and outliers

Dots should be plotted along the line







(Intercept)

Random effect quantiles

0.6  
0.3  
0.0  
-0.3

-2

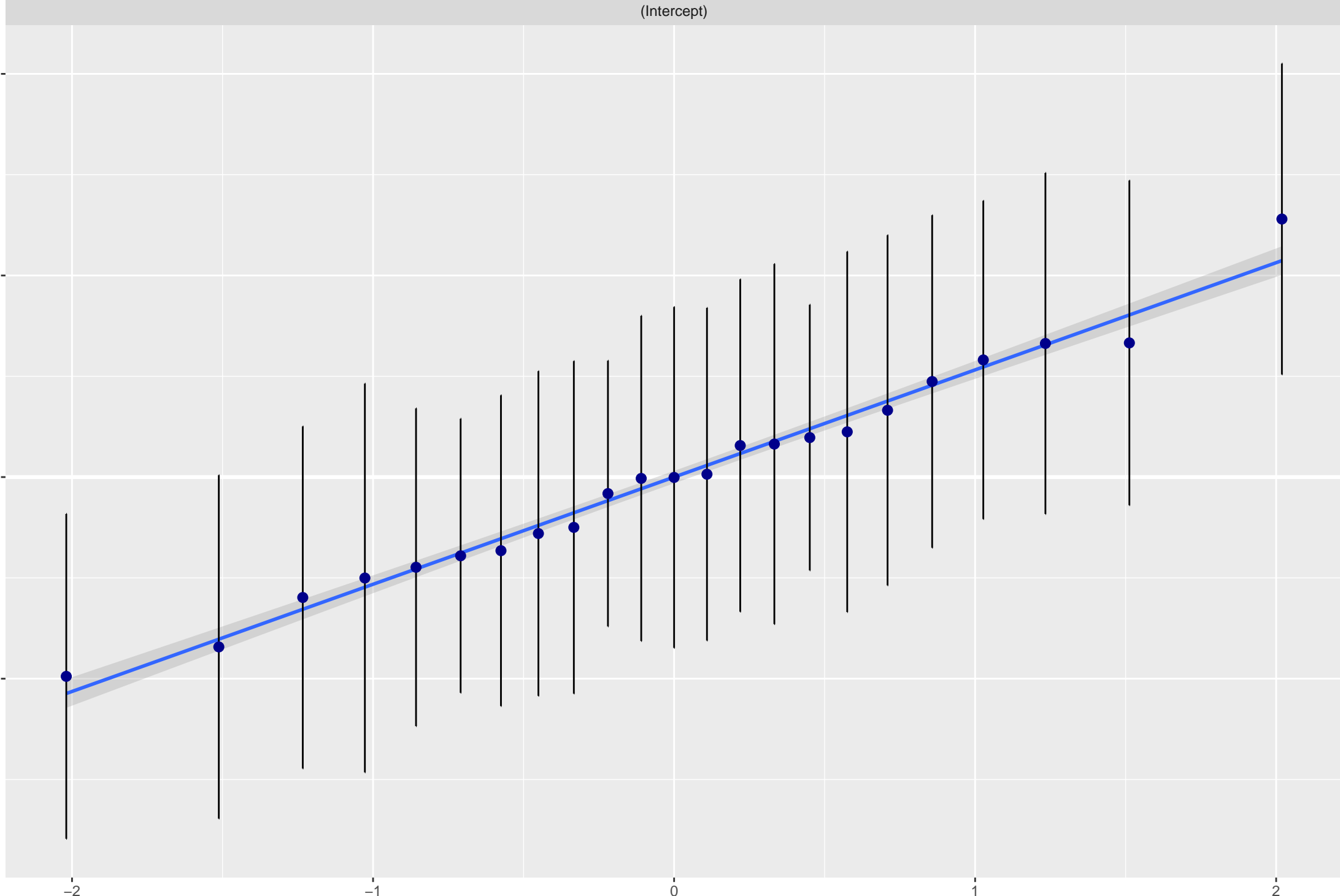
-1

0

1

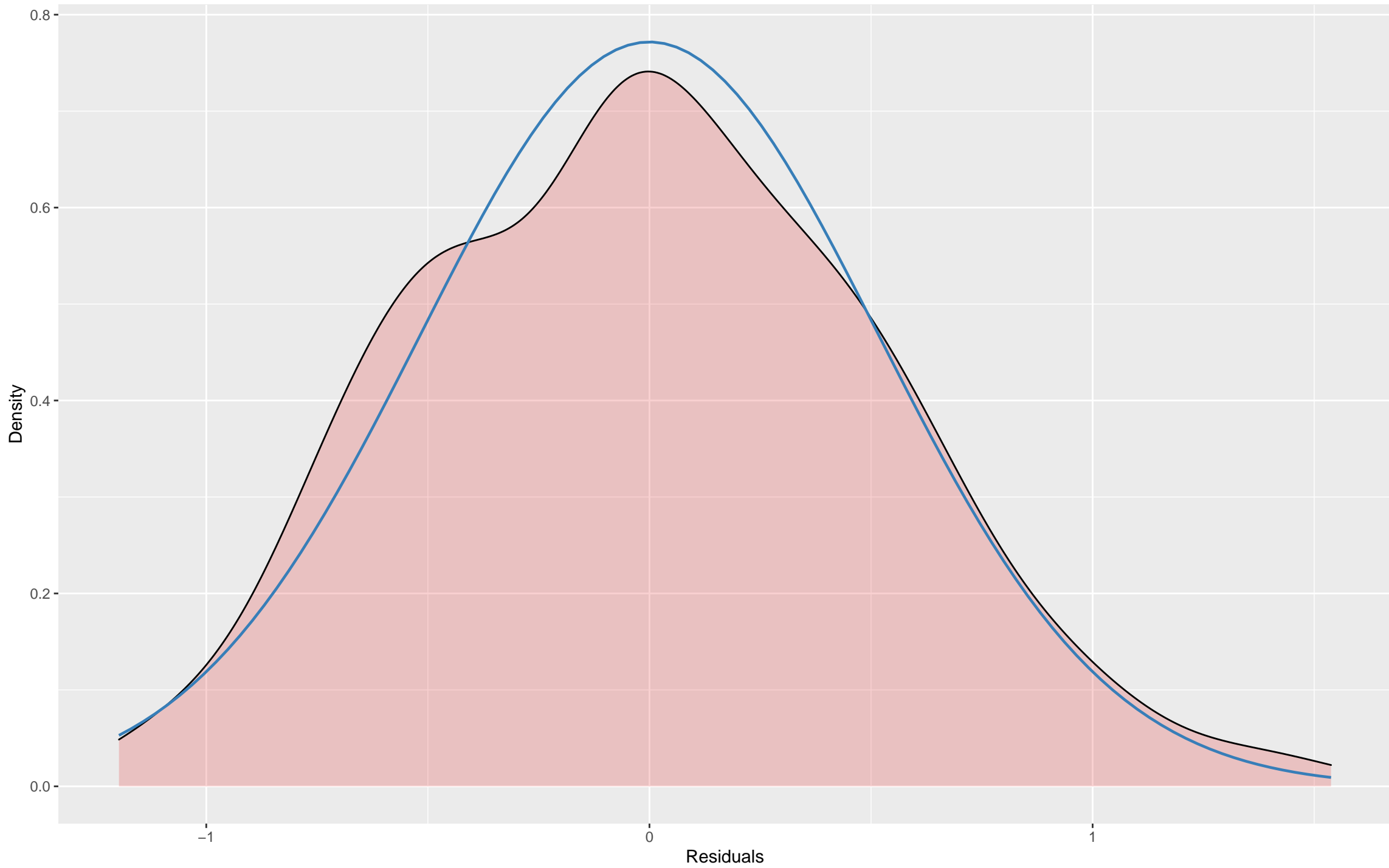
2

Standard normal quantiles



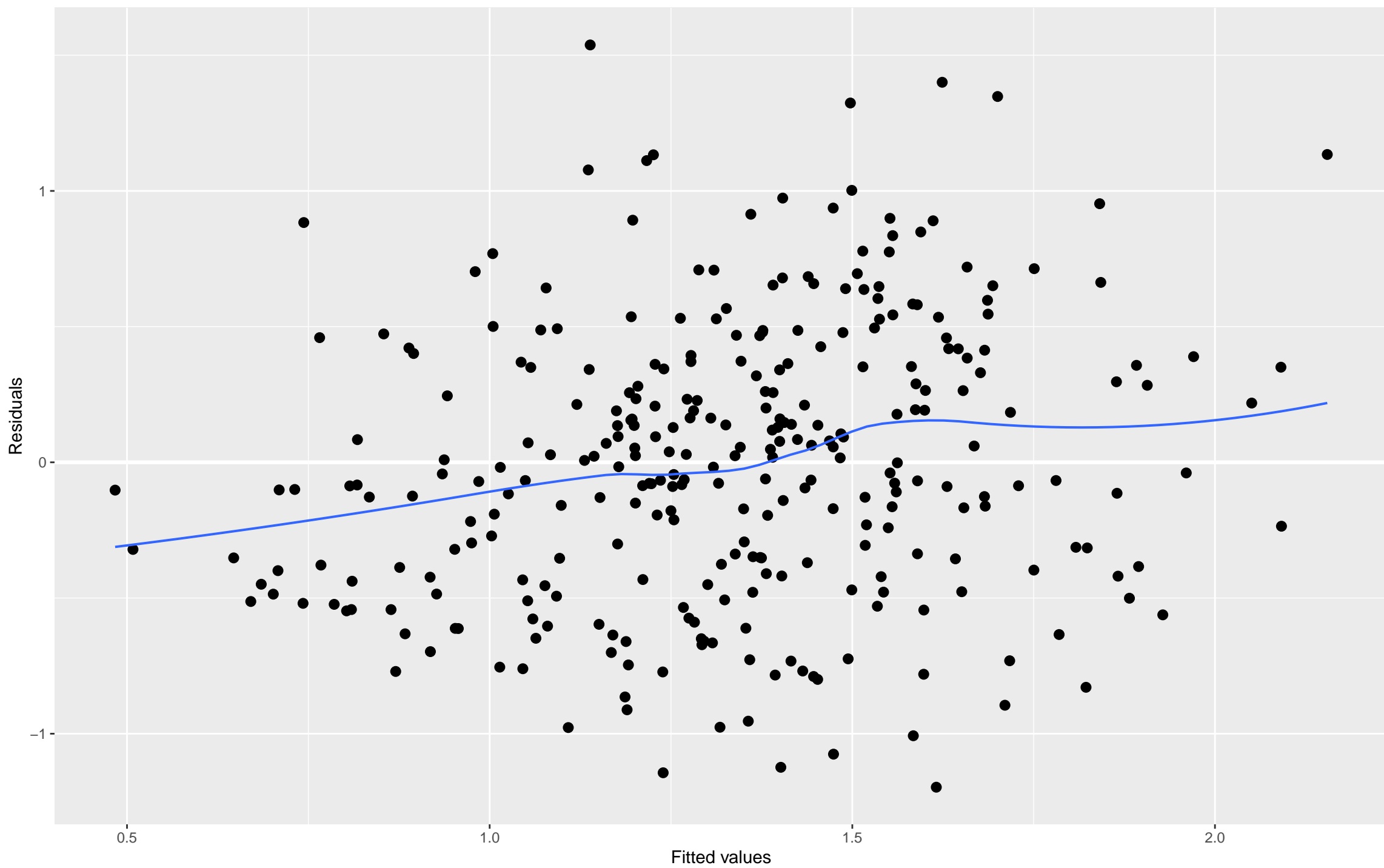
# Non-normality of residuals

Distribution should look like normal curve



# Homoscedasticity (constant variance of residuals)

Amount and distance of points scattered above/below line is equal or randomly spread



# Model summaries for all models with delta AIC < 2

```
$log_days_since_last_haul
Linear mixed model fit by maximum likelihood . t-tests use Satterthwaite's method ['lmerModLmerTest']
Formula: log_biomass ~ log_days_since_last_haul + (1 | TrapID) + (1 | Date_YMD)
Data: trap_haul_no_zero

      AIC      BIC   logLik deviance df.resid
557.7      576.3   -273.9    547.7      302

Scaled residuals:
    Min       1Q   Median       3Q      Max
-2.19806 -0.73098 -0.03229  0.65968  2.82404

Random effects:
Groups   Name              Variance Std.Dev.
TrapID   (Intercept)      0.04657    0.2158
Date_YMD (Intercept)      0.04001    0.2000
Residual                    0.29660    0.5446
Number of obs: 307, groups: TrapID, 37; Date_YMD, 23

Fixed effects:
              Estimate Std. Error    df t value Pr(>|t|)
(Intercept)      0.5791      0.2067 26.4505   2.802  0.00937 **
log_days_since_last_haul  0.3165      0.0866 23.2383   3.655  0.00130 **
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Correlation of Fixed Effects:
      (Intr)
lg_dys_sn_ -0.946

$'location_exposure + log_days_since_last_haul'
Linear mixed model fit by maximum likelihood . t-tests use Satterthwaite's method ['lmerModLmerTest']
Formula: log_biomass ~ location_exposure + log_days_since_last_haul + (1 | TrapID) + (1 | Date_YMD)
Data: trap_haul_no_zero

      AIC      BIC   logLik deviance df.resid
557.8      580.1   -272.9    545.8      301

Scaled residuals:
    Min       1Q   Median       3Q      Max
-2.20897 -0.73974 -0.01921  0.64412  2.81684

Random effects:
Groups   Name              Variance Std.Dev.
TrapID   (Intercept)      0.04385    0.2094
Date_YMD (Intercept)      0.04024    0.2006
Residual                    0.29555    0.5436
Number of obs: 307, groups: TrapID, 37; Date_YMD, 23

Fixed effects:
              Estimate Std. Error    df t value Pr(>|t|)
(Intercept)      0.60960      0.20745 26.99061   2.939  0.00668 **
location_exposureWindward -0.20115      0.14414 90.98576  -1.396  0.16624
log_days_since_last_haul  0.31733      0.08665 23.59684   3.662  0.00126 **
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Correlation of Fixed Effects:
      (Intr) lctn_W
lctn_xparWn -0.097
lg_dys_sn_ -0.941 -0.012

$'design + log_days_since_last_haul'
Linear mixed model fit by maximum likelihood . t-tests use Satterthwaite's method ['lmerModLmerTest']
Formula: log_biomass ~ design + log_days_since_last_haul + (1 | TrapID) + (1 | Date_YMD)
Data: trap_haul_no_zero

      AIC      BIC   logLik deviance df.resid
558.3      580.7   -273.2    546.3      301

Scaled residuals:
    Min       1Q   Median       3Q      Max
-2.22809 -0.74814 -0.01972  0.64751  2.81760

Random effects:
Groups   Name              Variance Std.Dev.
TrapID   (Intercept)      0.04394    0.2096
Date_YMD (Intercept)      0.03694    0.1922
Residual                    0.29737    0.5453
Number of obs: 307, groups: TrapID, 37; Date_YMD, 23

Fixed effects:
              Estimate Std. Error    df t value Pr(>|t|)
(Intercept)      0.44301      0.23187 41.03757   1.911  0.06306 .
designZ           0.14185      0.11868 70.68139   1.195  0.23602
log_days_since_last_haul  0.33386      0.08575 24.49901   3.893  0.00067 ***
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Correlation of Fixed Effects:
      (Intr) desgnZ
designZ      -0.492
lg_dys_sn_ -0.894  0.167
```

# Full model summary

```
Linear mixed model fit by maximum likelihood . t-tests use Satterthwaite's method ['lmerModLmerTest']
Formula: log_biomass ~ design + log_days_since_last_haul + location_exposure + Exp_or_Cont + (1 | TrapID) + (1 | Date_YMD)
Data: trap_haul_no_zero

      AIC      BIC    logLik deviance df.resid
561.6    591.4    -272.8    545.6      299

Scaled residuals:
    Min       1Q   Median       3Q      Max
-2.21977 -0.74005 -0.02548  0.64625  2.81568

Random effects:
Groups   Name              Variance Std.Dev.
TrapID   (Intercept)  0.04338  0.2083
Date_YMD (Intercept)  0.03889  0.1972
Residual                    0.29606  0.5441
Number of obs: 307, groups:  TrapID, 37; Date_YMD, 23

Fixed effects:
              Estimate Std. Error    df t value Pr(>|t|)
(Intercept)  5.414e-01  2.642e-01  5.977e+01  2.049  0.04482 *
designZ       6.337e-02  1.488e-01  7.909e+01  0.426  0.67141
log_days_since_last_haul  3.249e-01  8.749e-02  2.468e+01  3.714  0.00105 **
location_exposureWindward -1.544e-01  1.805e-01  1.001e+02 -0.855  0.39448
Exp_or_ContExperimental  2.268e-04  1.011e-01  3.484e+01  0.002  0.99822
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Correlation of Fixed Effects:
      (Intr) desgnZ lg_____ lctn_W
designZ      -0.595
lg_dya_sn_   -0.834  0.197
lctn_xparWn  -0.422  0.605  0.110
Exp_r_CntEx -0.187 -0.025 -0.007 -0.006
```