

Hierarchical models with different priors for between groups standard deviation

Laura Balasso

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Data

```
regions <- c('Lazio', 'Lombardia', 'Abruzzo', 'Veneto', 'Emilia-Romagna', 'Toscana', 'Campania', 'Friuli-Venezia Giulia', 'Calabria')

regions
```

```
## [1] "Lazio"          "Lombardia"       "Abruzzo"
## [4] "Veneto"         "Emilia-Romagna" "Toscana"
## [7] "Campania"       "Friuli Venezia Giulia" "Sicilia"
## [10] "Calabria"
```

```
hier_data <- get_hier_data(data_it, regions, initial_date = as.Date('2020-08-30'))

p_delay <- get_delay_distribution()

stan_data_hier <- list(J = length(regions),
                      N = nrow(hier_data$exposures),
                      N_nonzero = length(hier_data$nonzero_days),
                      nonzero_days = hier_data$nonzero_days,
                      conv_gt = get_gt_convolution_ln2(nrow(hier_data$exposures)),
                      length_delay = length(p_delay),
                      p_delay = p_delay,
                      exposures = hier_data$exposures,
                      nonzero_positives = hier_data$positives[hier_data$nonzero_days ,]
)
```

Models

Three hierarchical models with different priors on tau are compared to the simple model (that fits a different model for each region).

Hierarchical model 1

$$\log(r_t) \sim N(\mu_t, \tau^2)$$
$$\tau \sim \text{HalfCauchy}(0, 2.5)$$

Hierarchical model 2

$$\log(r_t) \sim N(\mu_t, \tau^2)$$

$$\tau \sim \text{HalfCauchy}(0, 1)$$

Hierarchical model 3

$$\log(r_t) \sim N(\mu_t, \tau^2)$$

$$\tau \sim \text{HalfCauchy}(0, 0.5)$$

```
## Simple model
compiled_grouped <- stan_model('../stan/grouped_model.stan')

## Trying to compile a simple C file

## Running /Library/Frameworks/R.framework/Resources/bin/R CMD SHLIB foo.c
## clang -mmacosx-version-min=10.13 -I"/Library/Frameworks/R.framework/Resources/include" -DNDEBUG -I
## In file included from <built-in>:1:
## In file included from /Library/Frameworks/R.framework/Versions/4.0/Resources/library/StanHeaders/inc
## In file included from /Library/Frameworks/R.framework/Versions/4.0/Resources/library/RcppEigen/inclu
## In file included from /Library/Frameworks/R.framework/Versions/4.0/Resources/library/RcppEigen/inclu
## /Library/Frameworks/R.framework/Versions/4.0/Resources/library/RcppEigen/include/Eigen/src/Core/util
## namespace Eigen {
## ~
## /Library/Frameworks/R.framework/Versions/4.0/Resources/library/RcppEigen/include/Eigen/src/Core/util
## namespace Eigen {
## ~
## ;
## In file included from <built-in>:1:
## In file included from /Library/Frameworks/R.framework/Versions/4.0/Resources/library/StanHeaders/inc
## In file included from /Library/Frameworks/R.framework/Versions/4.0/Resources/library/RcppEigen/inclu
## /Library/Frameworks/R.framework/Versions/4.0/Resources/library/RcppEigen/include/Eigen/Core:96:10: f
## #include <complex>
## ~~~~~
## 3 errors generated.
## make: *** [foo.o] Error 1

fit_grouped <- sampling(compiled_grouped, data= stan_data_hier, iter=2000, cores=getOption("mc.cores", 1))

##
## SAMPLING FOR MODEL 'grouped_model' NOW (CHAIN 1).
## Chain 1:
## Chain 1: Gradient evaluation took 0.022425 seconds
## Chain 1: 1000 transitions using 10 leapfrog steps per transition would take 224.25 seconds.
## Chain 1: Adjust your expectations accordingly!
## Chain 1:
## Chain 1:
## Chain 1: Iteration:    1 / 2000 [  0%] (Warmup)
## Chain 1: Iteration:   200 / 2000 [ 10%] (Warmup)
## Chain 1: Iteration:   400 / 2000 [ 20%] (Warmup)
## Chain 1: Iteration:   600 / 2000 [ 30%] (Warmup)
## Chain 1: Iteration:   800 / 2000 [ 40%] (Warmup)
## Chain 1: Iteration:  1000 / 2000 [ 50%] (Warmup)
## Chain 1: Iteration:  1001 / 2000 [ 50%] (Sampling)
## Chain 1: Iteration:  1200 / 2000 [ 60%] (Sampling)
```

```

## Chain 1: Iteration: 1400 / 2000 [ 70%] (Sampling)
## Chain 1: Iteration: 1600 / 2000 [ 80%] (Sampling)
## Chain 1: Iteration: 1800 / 2000 [ 90%] (Sampling)
## Chain 1: Iteration: 2000 / 2000 [100%] (Sampling)
## Chain 1:
## Chain 1: Elapsed Time: 1351.81 seconds (Warm-up)
## Chain 1: 1075.06 seconds (Sampling)
## Chain 1: 2426.88 seconds (Total)
## Chain 1:
##
## SAMPLING FOR MODEL 'grouped_model' NOW (CHAIN 2).
## Chain 2:
## Chain 2: Gradient evaluation took 0.007134 seconds
## Chain 2: 1000 transitions using 10 leapfrog steps per transition would take 71.34 seconds.
## Chain 2: Adjust your expectations accordingly!
## Chain 2:
## Chain 2:
## Chain 2: Iteration: 1 / 2000 [ 0%] (Warmup)
## Chain 2: Iteration: 200 / 2000 [ 10%] (Warmup)
## Chain 2: Iteration: 400 / 2000 [ 20%] (Warmup)
## Chain 2: Iteration: 600 / 2000 [ 30%] (Warmup)
## Chain 2: Iteration: 800 / 2000 [ 40%] (Warmup)
## Chain 2: Iteration: 1000 / 2000 [ 50%] (Warmup)
## Chain 2: Iteration: 1001 / 2000 [ 50%] (Sampling)
## Chain 2: Iteration: 1200 / 2000 [ 60%] (Sampling)
## Chain 2: Iteration: 1400 / 2000 [ 70%] (Sampling)
## Chain 2: Iteration: 1600 / 2000 [ 80%] (Sampling)
## Chain 2: Iteration: 1800 / 2000 [ 90%] (Sampling)
## Chain 2: Iteration: 2000 / 2000 [100%] (Sampling)
## Chain 2:
## Chain 2: Elapsed Time: 1268.62 seconds (Warm-up)
## Chain 2: 1154.82 seconds (Sampling)
## Chain 2: 2423.45 seconds (Total)
## Chain 2:
##
## SAMPLING FOR MODEL 'grouped_model' NOW (CHAIN 3).
## Chain 3:
## Chain 3: Gradient evaluation took 0.012382 seconds
## Chain 3: 1000 transitions using 10 leapfrog steps per transition would take 123.82 seconds.
## Chain 3: Adjust your expectations accordingly!
## Chain 3:
## Chain 3:
## Chain 3: Iteration: 1 / 2000 [ 0%] (Warmup)
## Chain 3: Iteration: 200 / 2000 [ 10%] (Warmup)
## Chain 3: Iteration: 400 / 2000 [ 20%] (Warmup)
## Chain 3: Iteration: 600 / 2000 [ 30%] (Warmup)
## Chain 3: Iteration: 800 / 2000 [ 40%] (Warmup)
## Chain 3: Iteration: 1000 / 2000 [ 50%] (Warmup)
## Chain 3: Iteration: 1001 / 2000 [ 50%] (Sampling)
## Chain 3: Iteration: 1200 / 2000 [ 60%] (Sampling)
## Chain 3: Iteration: 1400 / 2000 [ 70%] (Sampling)
## Chain 3: Iteration: 1600 / 2000 [ 80%] (Sampling)
## Chain 3: Iteration: 1800 / 2000 [ 90%] (Sampling)
## Chain 3: Iteration: 2000 / 2000 [100%] (Sampling)

```

```

## Chain 3:
## Chain 3: Elapsed Time: 1189.94 seconds (Warm-up)
## Chain 3: 1118.24 seconds (Sampling)
## Chain 3: 2308.19 seconds (Total)
## Chain 3:
##
## SAMPLING FOR MODEL 'grouped_model' NOW (CHAIN 4).
## Chain 4:
## Chain 4: Gradient evaluation took 0.008676 seconds
## Chain 4: 1000 transitions using 10 leapfrog steps per transition would take 86.76 seconds.
## Chain 4: Adjust your expectations accordingly!
## Chain 4:
## Chain 4:
## Chain 4: Iteration: 1 / 2000 [ 0%] (Warmup)
## Chain 4: Iteration: 200 / 2000 [ 10%] (Warmup)
## Chain 4: Iteration: 400 / 2000 [ 20%] (Warmup)
## Chain 4: Iteration: 600 / 2000 [ 30%] (Warmup)
## Chain 4: Iteration: 800 / 2000 [ 40%] (Warmup)
## Chain 4: Iteration: 1000 / 2000 [ 50%] (Warmup)
## Chain 4: Iteration: 1001 / 2000 [ 50%] (Sampling)
## Chain 4: Iteration: 1200 / 2000 [ 60%] (Sampling)
## Chain 4: Iteration: 1400 / 2000 [ 70%] (Sampling)
## Chain 4: Iteration: 1600 / 2000 [ 80%] (Sampling)
## Chain 4: Iteration: 1800 / 2000 [ 90%] (Sampling)
## Chain 4: Iteration: 2000 / 2000 [100%] (Sampling)
## Chain 4:
## Chain 4: Elapsed Time: 1285.77 seconds (Warm-up)
## Chain 4: 1075.81 seconds (Sampling)
## Chain 4: 2361.58 seconds (Total)
## Chain 4:
## Hierarchical model 1
compiled_hier <- stan_model('../stan/hier_rt_model.stan')

## Trying to compile a simple C file

## Running /Library/Frameworks/R.framework/Resources/bin/R CMD SHLIB foo.c
## clang -mmacosx-version-min=10.13 -I"/Library/Frameworks/R.framework/Resources/include" -DNDEBUG -I
## In file included from <built-in>:1:
## In file included from /Library/Frameworks/R.framework/Versions/4.0/Resources/library/StanHeaders/inc
## In file included from /Library/Frameworks/R.framework/Versions/4.0/Resources/library/RcppEigen/inclu
## In file included from /Library/Frameworks/R.framework/Versions/4.0/Resources/library/RcppEigen/inclu
## /Library/Frameworks/R.framework/Versions/4.0/Resources/library/RcppEigen/include/Eigen/src/Core/util
## namespace Eigen {
## ~
## /Library/Frameworks/R.framework/Versions/4.0/Resources/library/RcppEigen/include/Eigen/src/Core/util
## namespace Eigen {
## ~
## ;
## In file included from <built-in>:1:
## In file included from /Library/Frameworks/R.framework/Versions/4.0/Resources/library/StanHeaders/inc
## In file included from /Library/Frameworks/R.framework/Versions/4.0/Resources/library/RcppEigen/inclu
## /Library/Frameworks/R.framework/Versions/4.0/Resources/library/RcppEigen/include/Eigen/Core:96:10: f
## #include <complex>
## ~~~~~~

```

```

## 3 errors generated.
## make: *** [foo.o] Error 1

fit_hier <- sampling(compiled_hier, data = stan_data_hier, iter= 2000, cores=getOption("mc.cores", 1L))

##
## SAMPLING FOR MODEL 'hier_rt_model' NOW (CHAIN 1).
## Chain 1:
## Chain 1: Gradient evaluation took 0.013848 seconds
## Chain 1: 1000 transitions using 10 leapfrog steps per transition would take 138.48 seconds.
## Chain 1: Adjust your expectations accordingly!
## Chain 1:
## Chain 1:
## Chain 1: Iteration:    1 / 2000 [  0%] (Warmup)
## Chain 1: Iteration:   200 / 2000 [ 10%] (Warmup)
## Chain 1: Iteration:   400 / 2000 [ 20%] (Warmup)
## Chain 1: Iteration:   600 / 2000 [ 30%] (Warmup)
## Chain 1: Iteration:   800 / 2000 [ 40%] (Warmup)
## Chain 1: Iteration:  1000 / 2000 [ 50%] (Warmup)
## Chain 1: Iteration: 1001 / 2000 [ 50%] (Sampling)
## Chain 1: Iteration: 1200 / 2000 [ 60%] (Sampling)
## Chain 1: Iteration: 1400 / 2000 [ 70%] (Sampling)
## Chain 1: Iteration: 1600 / 2000 [ 80%] (Sampling)
## Chain 1: Iteration: 1800 / 2000 [ 90%] (Sampling)
## Chain 1: Iteration: 2000 / 2000 [100%] (Sampling)
## Chain 1:
## Chain 1: Elapsed Time: 1274.6 seconds (Warm-up)
## Chain 1:                948.957 seconds (Sampling)
## Chain 1:                2223.56 seconds (Total)
## Chain 1:
##
## SAMPLING FOR MODEL 'hier_rt_model' NOW (CHAIN 2).
## Chain 2:
## Chain 2: Gradient evaluation took 0.008572 seconds
## Chain 2: 1000 transitions using 10 leapfrog steps per transition would take 85.72 seconds.
## Chain 2: Adjust your expectations accordingly!
## Chain 2:
## Chain 2:
## Chain 2: Iteration:    1 / 2000 [  0%] (Warmup)
## Chain 2: Iteration:   200 / 2000 [ 10%] (Warmup)
## Chain 2: Iteration:   400 / 2000 [ 20%] (Warmup)
## Chain 2: Iteration:   600 / 2000 [ 30%] (Warmup)
## Chain 2: Iteration:   800 / 2000 [ 40%] (Warmup)
## Chain 2: Iteration:  1000 / 2000 [ 50%] (Warmup)
## Chain 2: Iteration: 1001 / 2000 [ 50%] (Sampling)
## Chain 2: Iteration: 1200 / 2000 [ 60%] (Sampling)
## Chain 2: Iteration: 1400 / 2000 [ 70%] (Sampling)
## Chain 2: Iteration: 1600 / 2000 [ 80%] (Sampling)
## Chain 2: Iteration: 1800 / 2000 [ 90%] (Sampling)
## Chain 2: Iteration: 2000 / 2000 [100%] (Sampling)
## Chain 2:
## Chain 2: Elapsed Time: 1315.35 seconds (Warm-up)
## Chain 2:                938.585 seconds (Sampling)
## Chain 2:                2253.94 seconds (Total)
## Chain 2:

```

```

##
## SAMPLING FOR MODEL 'hier_rt_model' NOW (CHAIN 3).
## Chain 3:
## Chain 3: Gradient evaluation took 0.008591 seconds
## Chain 3: 1000 transitions using 10 leapfrog steps per transition would take 85.91 seconds.
## Chain 3: Adjust your expectations accordingly!
## Chain 3:
## Chain 3:
## Chain 3: Iteration:    1 / 2000 [  0%] (Warmup)
## Chain 3: Iteration:   200 / 2000 [ 10%] (Warmup)
## Chain 3: Iteration:   400 / 2000 [ 20%] (Warmup)
## Chain 3: Iteration:   600 / 2000 [ 30%] (Warmup)
## Chain 3: Iteration:   800 / 2000 [ 40%] (Warmup)
## Chain 3: Iteration:  1000 / 2000 [ 50%] (Warmup)
## Chain 3: Iteration: 1001 / 2000 [ 50%] (Sampling)
## Chain 3: Iteration: 1200 / 2000 [ 60%] (Sampling)
## Chain 3: Iteration: 1400 / 2000 [ 70%] (Sampling)
## Chain 3: Iteration: 1600 / 2000 [ 80%] (Sampling)
## Chain 3: Iteration: 1800 / 2000 [ 90%] (Sampling)
## Chain 3: Iteration: 2000 / 2000 [100%] (Sampling)
## Chain 3:
## Chain 3: Elapsed Time: 1260.59 seconds (Warm-up)
## Chain 3:                944.517 seconds (Sampling)
## Chain 3:                2205.11 seconds (Total)
## Chain 3:
##
## SAMPLING FOR MODEL 'hier_rt_model' NOW (CHAIN 4).
## Chain 4:
## Chain 4: Gradient evaluation took 0.007199 seconds
## Chain 4: 1000 transitions using 10 leapfrog steps per transition would take 71.99 seconds.
## Chain 4: Adjust your expectations accordingly!
## Chain 4:
## Chain 4:
## Chain 4: Iteration:    1 / 2000 [  0%] (Warmup)
## Chain 4: Iteration:   200 / 2000 [ 10%] (Warmup)
## Chain 4: Iteration:   400 / 2000 [ 20%] (Warmup)
## Chain 4: Iteration:   600 / 2000 [ 30%] (Warmup)
## Chain 4: Iteration:   800 / 2000 [ 40%] (Warmup)
## Chain 4: Iteration:  1000 / 2000 [ 50%] (Warmup)
## Chain 4: Iteration: 1001 / 2000 [ 50%] (Sampling)
## Chain 4: Iteration: 1200 / 2000 [ 60%] (Sampling)
## Chain 4: Iteration: 1400 / 2000 [ 70%] (Sampling)
## Chain 4: Iteration: 1600 / 2000 [ 80%] (Sampling)
## Chain 4: Iteration: 1800 / 2000 [ 90%] (Sampling)
## Chain 4: Iteration: 2000 / 2000 [100%] (Sampling)
## Chain 4:
## Chain 4: Elapsed Time: 1285.94 seconds (Warm-up)
## Chain 4:                978.256 seconds (Sampling)
## Chain 4:                2264.19 seconds (Total)
## Chain 4:
## Hierarchical model 2
compiled_hierC1 <- stan_model('../stan/hier_model_C1.stan')

## Trying to compile a simple C file

```

```

## Running /Library/Frameworks/R.framework/Resources/bin/R CMD SHLIB foo.c
## clang -mmacosx-version-min=10.13 -I"/Library/Frameworks/R.framework/Resources/include" -DNDEBUG -I
## In file included from <built-in>:1:
## In file included from /Library/Frameworks/R.framework/Versions/4.0/Resources/library/StanHeaders/inc
## In file included from /Library/Frameworks/R.framework/Versions/4.0/Resources/library/RcppEigen/inclu
## In file included from /Library/Frameworks/R.framework/Versions/4.0/Resources/library/RcppEigen/inclu
## /Library/Frameworks/R.framework/Versions/4.0/Resources/library/RcppEigen/include/Eigen/src/Core/util
## namespace Eigen {
## ~
## /Library/Frameworks/R.framework/Versions/4.0/Resources/library/RcppEigen/include/Eigen/src/Core/util
## namespace Eigen {
## ~
## ;
## In file included from <built-in>:1:
## In file included from /Library/Frameworks/R.framework/Versions/4.0/Resources/library/StanHeaders/inc
## In file included from /Library/Frameworks/R.framework/Versions/4.0/Resources/library/RcppEigen/inclu
## /Library/Frameworks/R.framework/Versions/4.0/Resources/library/RcppEigen/include/Eigen/Core:96:10: f
## #include <complex>
## ~~~~~
## 3 errors generated.
## make: *** [foo.o] Error 1

```

```

fit_C1 <- sampling(compiled_hierC1, data = stan_data_hier, iter=2000, cores=getOption("mc.cores", 1L))

```

```

##
## SAMPLING FOR MODEL 'hier_model_C1' NOW (CHAIN 1).
## Chain 1:
## Chain 1: Gradient evaluation took 0.014197 seconds
## Chain 1: 1000 transitions using 10 leapfrog steps per transition would take 141.97 seconds.
## Chain 1: Adjust your expectations accordingly!
## Chain 1:
## Chain 1:
## Chain 1: Iteration:    1 / 2000 [ 0%] (Warmup)
## Chain 1: Iteration:   200 / 2000 [ 10%] (Warmup)
## Chain 1: Iteration:   400 / 2000 [ 20%] (Warmup)
## Chain 1: Iteration:   600 / 2000 [ 30%] (Warmup)
## Chain 1: Iteration:   800 / 2000 [ 40%] (Warmup)
## Chain 1: Iteration:  1000 / 2000 [ 50%] (Warmup)
## Chain 1: Iteration: 1001 / 2000 [ 50%] (Sampling)
## Chain 1: Iteration: 1200 / 2000 [ 60%] (Sampling)
## Chain 1: Iteration: 1400 / 2000 [ 70%] (Sampling)
## Chain 1: Iteration: 1600 / 2000 [ 80%] (Sampling)
## Chain 1: Iteration: 1800 / 2000 [ 90%] (Sampling)
## Chain 1: Iteration: 2000 / 2000 [100%] (Sampling)
## Chain 1:
## Chain 1: Elapsed Time: 1257.01 seconds (Warm-up)
## Chain 1:           951.29 seconds (Sampling)
## Chain 1:           2208.3 seconds (Total)
## Chain 1:
##
## SAMPLING FOR MODEL 'hier_model_C1' NOW (CHAIN 2).
## Chain 2:
## Chain 2: Gradient evaluation took 0.007642 seconds
## Chain 2: 1000 transitions using 10 leapfrog steps per transition would take 76.42 seconds.
## Chain 2: Adjust your expectations accordingly!

```

```

## Chain 2:
## Chain 2:
## Chain 2: Iteration:    1 / 2000 [  0%] (Warmup)
## Chain 2: Iteration:   200 / 2000 [ 10%] (Warmup)
## Chain 2: Iteration:   400 / 2000 [ 20%] (Warmup)
## Chain 2: Iteration:   600 / 2000 [ 30%] (Warmup)
## Chain 2: Iteration:   800 / 2000 [ 40%] (Warmup)
## Chain 2: Iteration:  1000 / 2000 [ 50%] (Warmup)
## Chain 2: Iteration: 1001 / 2000 [ 50%] (Sampling)
## Chain 2: Iteration: 1200 / 2000 [ 60%] (Sampling)
## Chain 2: Iteration: 1400 / 2000 [ 70%] (Sampling)
## Chain 2: Iteration: 1600 / 2000 [ 80%] (Sampling)
## Chain 2: Iteration: 1800 / 2000 [ 90%] (Sampling)
## Chain 2: Iteration: 2000 / 2000 [100%] (Sampling)
## Chain 2:
## Chain 2: Elapsed Time: 1377.47 seconds (Warm-up)
## Chain 2:           963.79 seconds (Sampling)
## Chain 2:           2341.26 seconds (Total)
## Chain 2:
##
## SAMPLING FOR MODEL 'hier_model_C1' NOW (CHAIN 3).
## Chain 3:
## Chain 3: Gradient evaluation took 0.007546 seconds
## Chain 3: 1000 transitions using 10 leapfrog steps per transition would take 75.46 seconds.
## Chain 3: Adjust your expectations accordingly!
## Chain 3:
## Chain 3:
## Chain 3: Iteration:    1 / 2000 [  0%] (Warmup)
## Chain 3: Exception: Exception: poisson_rng: Rate parameter is 0, but must be > 0! (in 'modeld77356b
## (in 'modeld77356b4a1c0_hier_model_C1' at line 148)
##
## Chain 3: Exception: Exception: poisson_rng: Rate parameter is 0, but must be > 0! (in 'modeld77356b
## (in 'modeld77356b4a1c0_hier_model_C1' at line 148)
##
## Chain 3: Exception: Exception: poisson_rng: Rate parameter is 0, but must be > 0! (in 'modeld77356b
## (in 'modeld77356b4a1c0_hier_model_C1' at line 148)
##
## Chain 3: Exception: Exception: poisson_rng: Rate parameter is 0, but must be > 0! (in 'modeld77356b
## (in 'modeld77356b4a1c0_hier_model_C1' at line 148)
##
## Chain 3: Iteration:   200 / 2000 [ 10%] (Warmup)
## Chain 3: Iteration:   400 / 2000 [ 20%] (Warmup)
## Chain 3: Iteration:   600 / 2000 [ 30%] (Warmup)
## Chain 3: Iteration:   800 / 2000 [ 40%] (Warmup)
## Chain 3: Iteration:  1000 / 2000 [ 50%] (Warmup)
## Chain 3: Iteration: 1001 / 2000 [ 50%] (Sampling)
## Chain 3: Iteration: 1200 / 2000 [ 60%] (Sampling)
## Chain 3: Iteration: 1400 / 2000 [ 70%] (Sampling)
## Chain 3: Iteration: 1600 / 2000 [ 80%] (Sampling)
## Chain 3: Iteration: 1800 / 2000 [ 90%] (Sampling)
## Chain 3: Iteration: 2000 / 2000 [100%] (Sampling)
## Chain 3:
## Chain 3: Elapsed Time: 1289.43 seconds (Warm-up)
## Chain 3:           978.512 seconds (Sampling)

```



```

## Chain 3:                2267.94 seconds (Total)
## Chain 3:
##
## SAMPLING FOR MODEL 'hier_model_C1' NOW (CHAIN 4).
## Chain 4:
## Chain 4: Gradient evaluation took 0.007417 seconds
## Chain 4: 1000 transitions using 10 leapfrog steps per transition would take 74.17 seconds.
## Chain 4: Adjust your expectations accordingly!
## Chain 4:
## Chain 4:
## Chain 4: Iteration:      1 / 2000 [  0%] (Warmup)
## Chain 4: Iteration:    200 / 2000 [ 10%] (Warmup)
## Chain 4: Iteration:    400 / 2000 [ 20%] (Warmup)
## Chain 4: Iteration:    600 / 2000 [ 30%] (Warmup)
## Chain 4: Iteration:    800 / 2000 [ 40%] (Warmup)
## Chain 4: Iteration:   1000 / 2000 [ 50%] (Warmup)
## Chain 4: Iteration:   1001 / 2000 [ 50%] (Sampling)
## Chain 4: Iteration:   1200 / 2000 [ 60%] (Sampling)
## Chain 4: Iteration:   1400 / 2000 [ 70%] (Sampling)
## Chain 4: Iteration:   1600 / 2000 [ 80%] (Sampling)
## Chain 4: Iteration:   1800 / 2000 [ 90%] (Sampling)
## Chain 4: Iteration:   2000 / 2000 [100%] (Sampling)
## Chain 4:
## Chain 4: Elapsed Time: 1423.53 seconds (Warm-up)
## Chain 4:                1016.24 seconds (Sampling)
## Chain 4:                2439.77 seconds (Total)
## Chain 4:

## Warning in validityMethod(object): The following variables have undefined
## values: log_lik[1,1],The following variables have undefined values:
## log_lik[2,1],The following variables have undefined values: log_lik[3,1],The
## following variables have undefined values: log_lik[4,1],The following
## variables have undefined values: log_lik[5,1],The following variables have
## undefined values: log_lik[6,1],The following variables have undefined values:
## log_lik[7,1],The following variables have undefined values: log_lik[8,1],The
## following variables have undefined values: log_lik[9,1],The following
## variables have undefined values: log_lik[10,1],The following variables have
## undefined values: log_lik[11,1],The following variables have undefined values:
## log_lik[12,1],The following variables have undefined values: log_lik[13,1],The
## following variables have undefined values: log_lik[14,1],The following
## variables have undefined values: log_lik[15,1],The following variables have
## undefined values: log_lik[16,1],The following variables have undefined values:
## log_lik[17,1],The following variables have undefined values: log_lik[18,1],The
## following variables have undefined values: log_lik[19,1],The following
## variables have undefined values: log_lik[20,1],The following variables have
## undefined values: log_lik[21,1],The following variables have undefined values:
## log_lik[22,1],The following variables have undefined values: log_lik[23,1],The
## following variables have undefined values: log_lik[24,1],The following
## variables have undefined values: log_lik[25,1],The following variables have
## undefined values: log_lik[26,1],The following variables have undefined values:
## log_lik[27,1],The following variables have undefined values: log_lik[28,1],The
## following variables have undefined values: log_lik[29,1],The following
## variables have undefined values: log_lik[30,1],The following variables have
## undefined values: log_lik[31,1],The following variables have undefined values:

```

[illegible]

```
## variables have undefined values: log_lik[20,2],The following variables have
## undefined values: log_lik[21,2],The following variables have undefined values:
## log_lik[22,2],The following variables have undefined values: log_lik[23,2],The
## following variables have undefined values: log_lik[24,2],The following
## variables have undefined values: log_lik[25,2],The following variables have
## undefined values: log_lik[26,2],The following variables have undefined values:
## log_lik[27,2],The following variables have undefined values: log_lik[28,2],The
## following variables have undefined values: log_lik[29,2],The following
## variables have undefined values: log_lik[30,2],The following variables have
## undefined values: log_lik[31,2],The following variables have undefined values:
## log_lik[32,2],The following variables have undefined values: log_lik[33,2],The
## following variables have undefined values: log_lik[34,2],The following
## variables have undefined values: log_lik[35,2],The following variables have
## undefined values: log_lik[36,2],The following variables have undefined values:
## log_lik[37,2],The following variables have undefined values: log_lik[38,2],The
## following variables have undefined values: log_lik[39,2],The following
## variables have undefined values: log_lik[40,2],The following variables have
## undefined values: log_lik[41,2],The following variables have undefined values:
## log_lik[42,2],The following variables have undefined values: log_lik[43,2],The
## following variables have undefined values: log_lik[44,2],The following
## variables have undefined values: log_lik[45,2],The following variables have
## undefined values: log_lik[46,2],The following variables have undefined values:
## log_lik[47,2],The following variables have undefined values: log_lik[48,2],The
## following variables have undefined values: log_lik[49,2],The following
## variables have undefined values: log_lik[50,2],The following variables have
## undefined values: log_lik[51,2],The following variables have undefined values:
## log_lik[52,2],The following variables
```

```
## Hierarchical model 3
```

```
compiled_hierC05 <- stan_model(' ../stan/hier_model_C05.stan')
```

```
## Trying to compile a simple C file
```

```
## Running /Library/Frameworks/R.framework/Resources/bin/R CMD SHLIB foo.c
## clang -mmacosx-version-min=10.13 -I"/Library/Frameworks/R.framework/Resources/include" -DNDEBUG -I
## In file included from <built-in>:1:
## In file included from /Library/Frameworks/R.framework/Versions/4.0/Resources/library/StanHeaders/inc
## In file included from /Library/Frameworks/R.framework/Versions/4.0/Resources/library/RcppEigen/inclu
## In file included from /Library/Frameworks/R.framework/Versions/4.0/Resources/library/RcppEigen/inclu
## /Library/Frameworks/R.framework/Versions/4.0/Resources/library/RcppEigen/include/Eigen/src/Core/util
## namespace Eigen {
## ~
## /Library/Frameworks/R.framework/Versions/4.0/Resources/library/RcppEigen/include/Eigen/src/Core/util
## namespace Eigen {
## ~
## ;
## In file included from <built-in>:1:
## In file included from /Library/Frameworks/R.framework/Versions/4.0/Resources/library/StanHeaders/inc
## In file included from /Library/Frameworks/R.framework/Versions/4.0/Resources/library/RcppEigen/inclu
## /Library/Frameworks/R.framework/Versions/4.0/Resources/library/RcppEigen/include/Eigen/Core:96:10: f
## #include <complex>
## ~~~~~
## 3 errors generated.
## make: *** [foo.o] Error 1
```

```

fit_C05 <- sampling(compiled_hierC05, data= stan_data_hier, iter=2000,cores=getOption("mc.cores", 1L))

##
## SAMPLING FOR MODEL 'hier_model_C05' NOW (CHAIN 1).
## Chain 1:
## Chain 1: Gradient evaluation took 0.015648 seconds
## Chain 1: 1000 transitions using 10 leapfrog steps per transition would take 156.48 seconds.
## Chain 1: Adjust your expectations accordingly!
## Chain 1:
## Chain 1:
## Chain 1: Iteration:    1 / 2000 [  0%] (Warmup)
## Chain 1: Iteration:   200 / 2000 [ 10%] (Warmup)
## Chain 1: Iteration:   400 / 2000 [ 20%] (Warmup)
## Chain 1: Iteration:   600 / 2000 [ 30%] (Warmup)
## Chain 1: Iteration:   800 / 2000 [ 40%] (Warmup)
## Chain 1: Iteration:  1000 / 2000 [ 50%] (Warmup)
## Chain 1: Iteration:  1001 / 2000 [ 50%] (Sampling)
## Chain 1: Iteration:  1200 / 2000 [ 60%] (Sampling)
## Chain 1: Iteration:  1400 / 2000 [ 70%] (Sampling)
## Chain 1: Iteration:  1600 / 2000 [ 80%] (Sampling)
## Chain 1: Iteration:  1800 / 2000 [ 90%] (Sampling)
## Chain 1: Iteration:  2000 / 2000 [100%] (Sampling)
## Chain 1:
## Chain 1: Elapsed Time: 1340.02 seconds (Warm-up)
## Chain 1:                972.676 seconds (Sampling)
## Chain 1:                2312.69 seconds (Total)
## Chain 1:
##
## SAMPLING FOR MODEL 'hier_model_C05' NOW (CHAIN 2).
## Chain 2:
## Chain 2: Gradient evaluation took 0.009088 seconds
## Chain 2: 1000 transitions using 10 leapfrog steps per transition would take 90.88 seconds.
## Chain 2: Adjust your expectations accordingly!
## Chain 2:
## Chain 2:
## Chain 2: Iteration:    1 / 2000 [  0%] (Warmup)
## Chain 2: Iteration:   200 / 2000 [ 10%] (Warmup)
## Chain 2: Iteration:   400 / 2000 [ 20%] (Warmup)
## Chain 2: Iteration:   600 / 2000 [ 30%] (Warmup)
## Chain 2: Iteration:   800 / 2000 [ 40%] (Warmup)
## Chain 2: Iteration:  1000 / 2000 [ 50%] (Warmup)
## Chain 2: Iteration:  1001 / 2000 [ 50%] (Sampling)
## Chain 2: Iteration:  1200 / 2000 [ 60%] (Sampling)
## Chain 2: Iteration:  1400 / 2000 [ 70%] (Sampling)
## Chain 2: Iteration:  1600 / 2000 [ 80%] (Sampling)
## Chain 2: Iteration:  1800 / 2000 [ 90%] (Sampling)
## Chain 2: Iteration:  2000 / 2000 [100%] (Sampling)
## Chain 2:
## Chain 2: Elapsed Time: 1300.29 seconds (Warm-up)
## Chain 2:                936.442 seconds (Sampling)
## Chain 2:                2236.73 seconds (Total)
## Chain 2:
##
## SAMPLING FOR MODEL 'hier_model_C05' NOW (CHAIN 3).

```

```

## Chain 3:
## Chain 3: Gradient evaluation took 0.009058 seconds
## Chain 3: 1000 transitions using 10 leapfrog steps per transition would take 90.58 seconds.
## Chain 3: Adjust your expectations accordingly!
## Chain 3:
## Chain 3:
## Chain 3: Iteration:    1 / 2000 [  0%] (Warmup)
## Chain 3: Iteration:   200 / 2000 [ 10%] (Warmup)
## Chain 3: Iteration:   400 / 2000 [ 20%] (Warmup)
## Chain 3: Iteration:   600 / 2000 [ 30%] (Warmup)
## Chain 3: Iteration:   800 / 2000 [ 40%] (Warmup)
## Chain 3: Iteration:  1000 / 2000 [ 50%] (Warmup)
## Chain 3: Iteration: 1001 / 2000 [ 50%] (Sampling)
## Chain 3: Iteration: 1200 / 2000 [ 60%] (Sampling)
## Chain 3: Iteration: 1400 / 2000 [ 70%] (Sampling)
## Chain 3: Iteration: 1600 / 2000 [ 80%] (Sampling)
## Chain 3: Iteration: 1800 / 2000 [ 90%] (Sampling)
## Chain 3: Iteration: 2000 / 2000 [100%] (Sampling)
## Chain 3:
## Chain 3: Elapsed Time: 1282.25 seconds (Warm-up)
## Chain 3:                938.121 seconds (Sampling)
## Chain 3:                2220.37 seconds (Total)
## Chain 3:
##
## SAMPLING FOR MODEL 'hier_model_C05' NOW (CHAIN 4).
## Chain 4:
## Chain 4: Gradient evaluation took 0.008213 seconds
## Chain 4: 1000 transitions using 10 leapfrog steps per transition would take 82.13 seconds.
## Chain 4: Adjust your expectations accordingly!
## Chain 4:
## Chain 4:
## Chain 4: Iteration:    1 / 2000 [  0%] (Warmup)
## Chain 4: Iteration:   200 / 2000 [ 10%] (Warmup)
## Chain 4: Iteration:   400 / 2000 [ 20%] (Warmup)
## Chain 4: Iteration:   600 / 2000 [ 30%] (Warmup)
## Chain 4: Iteration:   800 / 2000 [ 40%] (Warmup)
## Chain 4: Iteration:  1000 / 2000 [ 50%] (Warmup)
## Chain 4: Iteration: 1001 / 2000 [ 50%] (Sampling)
## Chain 4: Iteration: 1200 / 2000 [ 60%] (Sampling)
## Chain 4: Iteration: 1400 / 2000 [ 70%] (Sampling)
## Chain 4: Iteration: 1600 / 2000 [ 80%] (Sampling)
## Chain 4: Iteration: 1800 / 2000 [ 90%] (Sampling)
## Chain 4: Iteration: 2000 / 2000 [100%] (Sampling)
## Chain 4:
## Chain 4: Elapsed Time: 1247.98 seconds (Warm-up)
## Chain 4:                893.731 seconds (Sampling)
## Chain 4:                2141.71 seconds (Total)
## Chain 4:

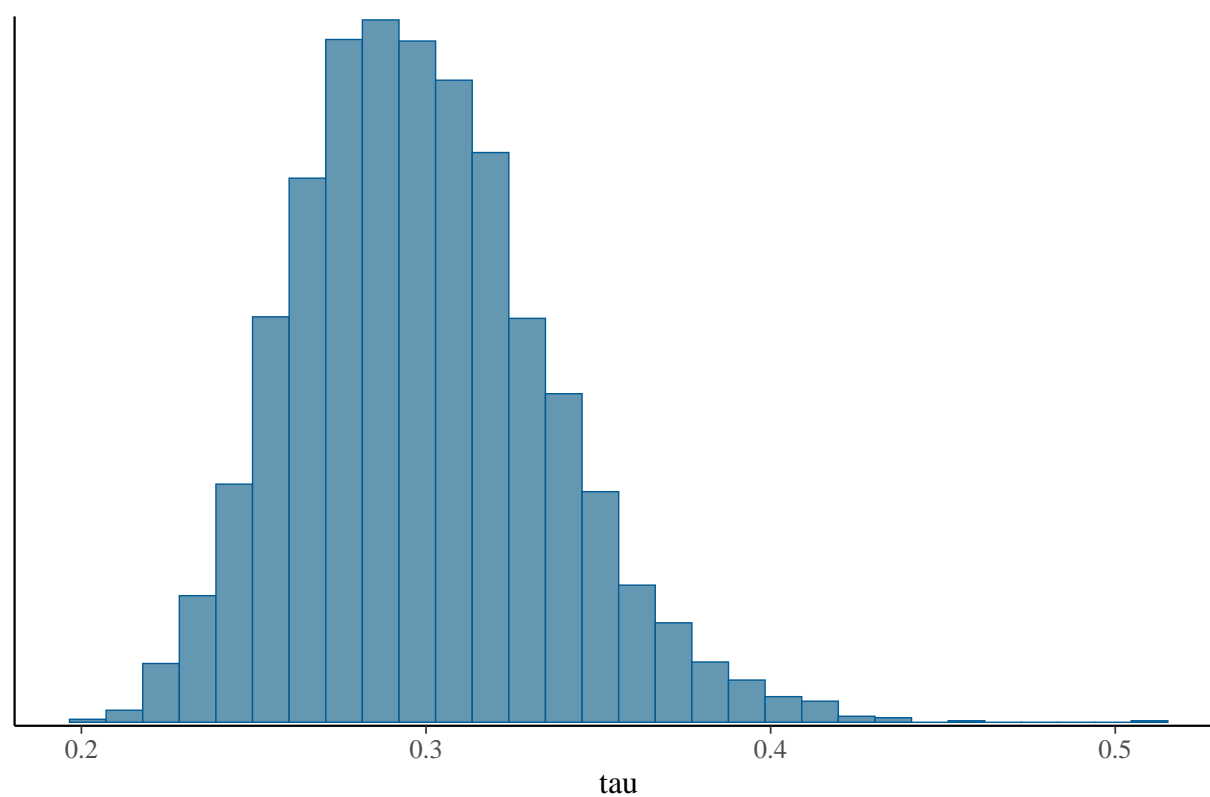
```

Posterior histograms for tau

```
mcmc_hist(fit_hier, pars = 'tau') + ggtitle('Model 1')
```

```
## `stat_bin()` using `bins = 30`. Pick better value with `binwidth`.
```

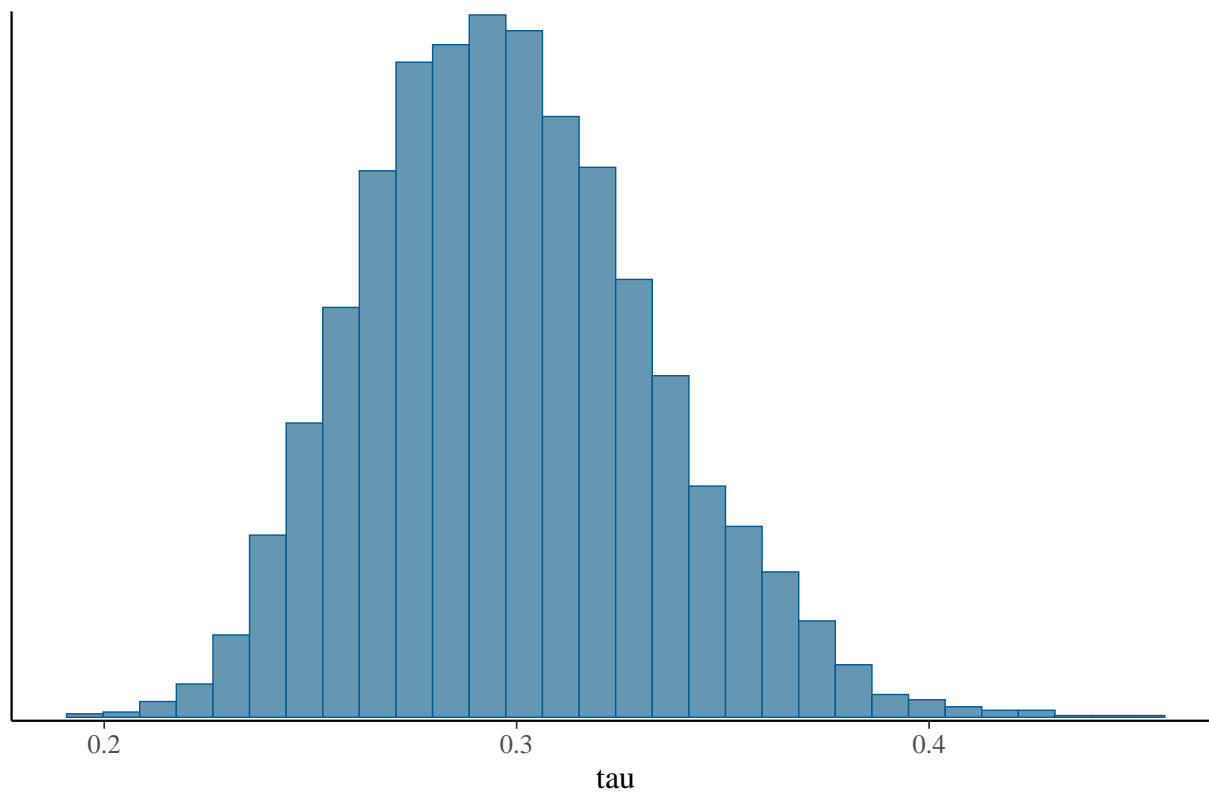
Model 1



```
mcmc_hist(fit_C1, pars = 'tau') + ggtitle('Model 2')
```

```
## `stat_bin()` using `bins = 30`. Pick better value with `binwidth`.
```

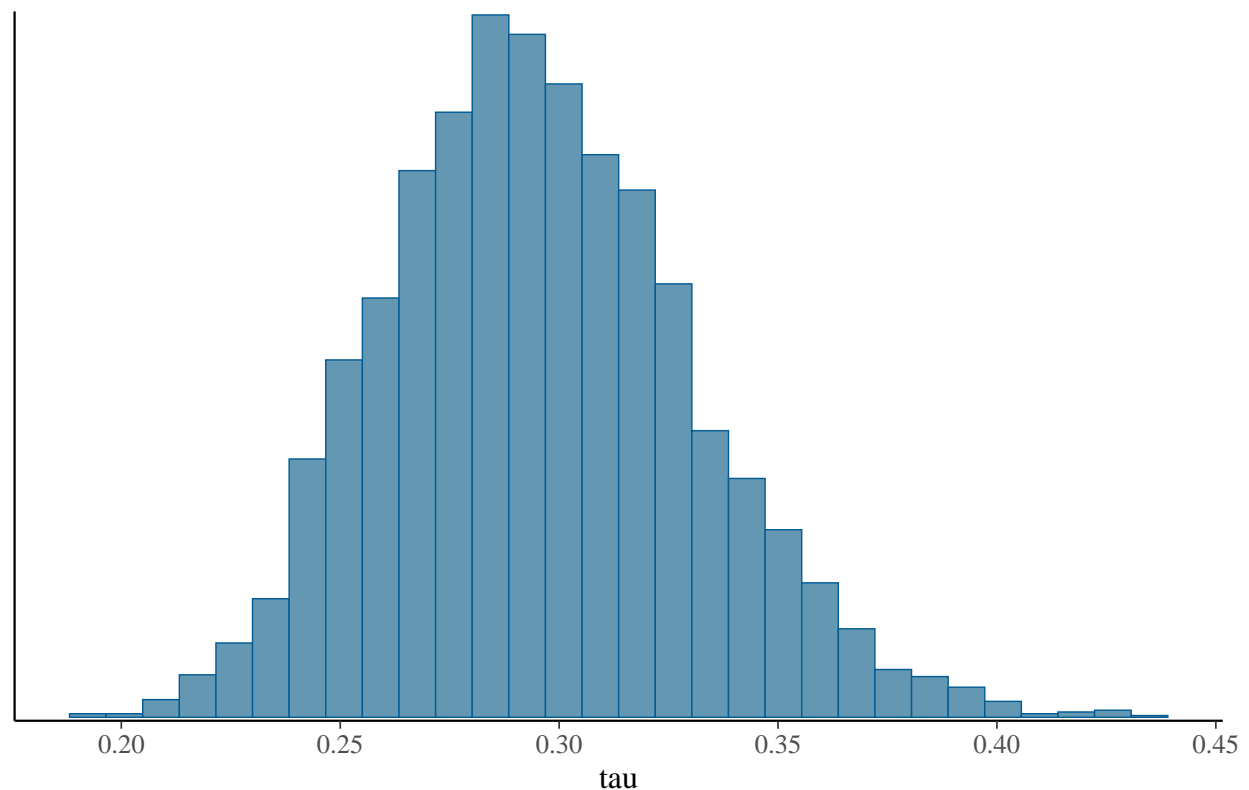
Model 2



```
mcmc_hist(fit_C05, pars = 'tau') + ggtitle('Model 3')
```

```
## `stat_bin()` using `bins = 30`. Pick better value with `binwidth`.
```

Model 3



Rt

For each of the three hierarchical models, the resulting Rt curve is overlapped to the Rt curve produced by the single region model, both with 50% and 95% credibility intervals.

The same plots are produced for the regions Lazio, Lombardia, Veneto, Friuli Venezia Giulia and Abruzzo.

```
### Retrieve rt medians and intervals for a given model and region
rt_intervals <- function(data, fit, regions_vector, region){
  summary <- summary(fit)
  region_idx = which(regions_vector == region)
  rt_1 <- which(rownames(summary$summary) == paste('r_t[1,', region_idx, ']', sep=''))
  rt_index <- seq(rt_1, rt_1 + length(data$dates) * length(regions_vector) - 1, by=length(regions_vector))
  rt_median <- summary$summary[rt_index, '50%']
  min_rt_50_interval <- summary$summary[rt_index, '25%']
  max_rt_50_interval <- summary$summary[rt_index, '75%']
  min_rt_95_interval <- summary$summary[rt_index, '2.5%']
  max_rt_95_interval <- summary$summary[rt_index, '97.5%']

  return(list(median = rt_median, min_50 = min_rt_50_interval, max_50 = max_rt_50_interval, min_95 = min_rt_95_interval, max_95 = max_rt_95_interval))
}

### Plot rt curve for a single region
plot_rt <- function(data, fit, regions_vector, region){

  intervals <- rt_intervals(data, fit, regions_vector, region)

  p <- ggplot( ) +
```



```

    geom_line(aes(x = data$dates, y=intervals$median ), color='darkred')+
    xlab('Date') +
    ylab('') +
    geom_hline(yintercept=1, linetype="dashed", color = "red") +
    geom_vline(xintercept = data$dates[1]) +
    geom_ribbon(aes(x = data$dates, ymin = intervals$min_50, ymax = intervals$max_50), alpha= 0.5, fill="red")+
    geom_ribbon(aes(x = data$dates,ymin =intervals$min_95, ymax = intervals$max_95), alpha= 0.1, fill="red")+

    return(p)
}

### overlay with rt plot of the single region model
plot_overlay <- function(plot, region){

  simple_intervals <- rt_intervals(hier_data, fit_grouped, regions, region)

  p <- plot + geom_line(aes(x=hier_data$dates, y = simple_intervals$median), color = 'navyblue') +
  geom_ribbon(aes(x = hier_data$dates,ymin = simple_intervals$min_50, ymax = simple_intervals$max_50), alpha= 0.5, fill="red")+
  geom_ribbon(aes(x = hier_data$dates,ymin = simple_intervals$min_95, ymax = simple_intervals$max_95), alpha= 0.1, fill="red")+
  ggtitle(paste('Rt', region , sep = ' '))

  return(p)
}

get_plot_limits <- function(plot) {
  gb = ggplot_build(plot)
  xmin = gb$layout$panel_params[[1]]$x.range[1]
  xmax = gb$layout$panel_params[[1]]$x.range[2]
  ymin = gb$layout$panel_params[[1]]$y.range[1]
  ymax = gb$layout$panel_params[[1]]$y.range[2]
  list(xmin = xmin, xmax = xmax, ymin = ymin, ymax = ymax)
}

```

Lazio

```

p1 <- plot_rt(hier_data, fit_hier, regions, 'Lazio')
p1 <- plot_overlay(p1, 'Lazio') + labs(subtitle = 'hierarchical model n.1 (red) and single region model')

p2 <- plot_rt(hier_data, fit_C05, regions, 'Lazio')
p2 <- plot_overlay(p2, 'Lazio') + labs(subtitle = 'hierarchical model n.2 (red) and single region model')

p3 <- plot_rt(hier_data, fit_C1, regions, 'Lazio')
p3 <- plot_overlay(p3, 'Lazio') + labs(subtitle = 'hierarchical model n.3 (red) and single region model')

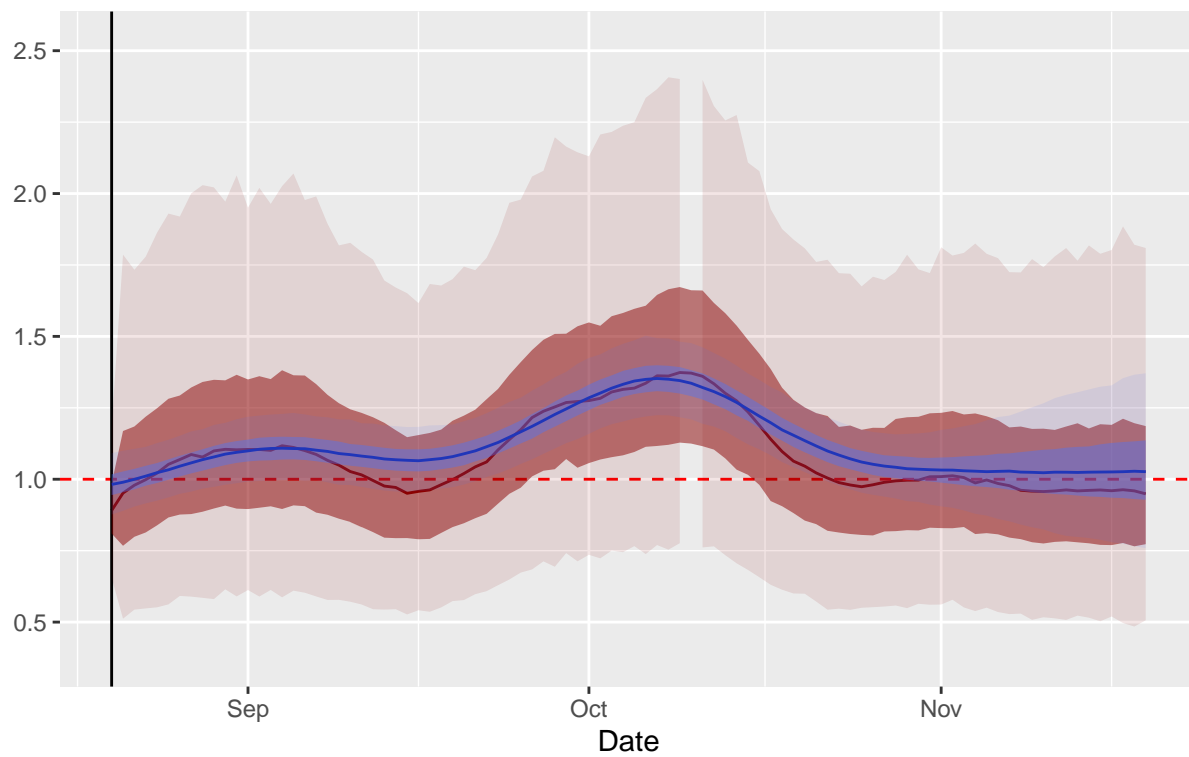
ymin <- min(get_plot_limits(p1)$ymin, get_plot_limits(p2)$ymin, get_plot_limits(p3)$ymin)
ymax <- min(get_plot_limits(p1)$ymax, get_plot_limits(p2)$ymax, get_plot_limits(p3)$ymax)

p1 + ylim(c(ymin, ymax))

```

Rt Lazio

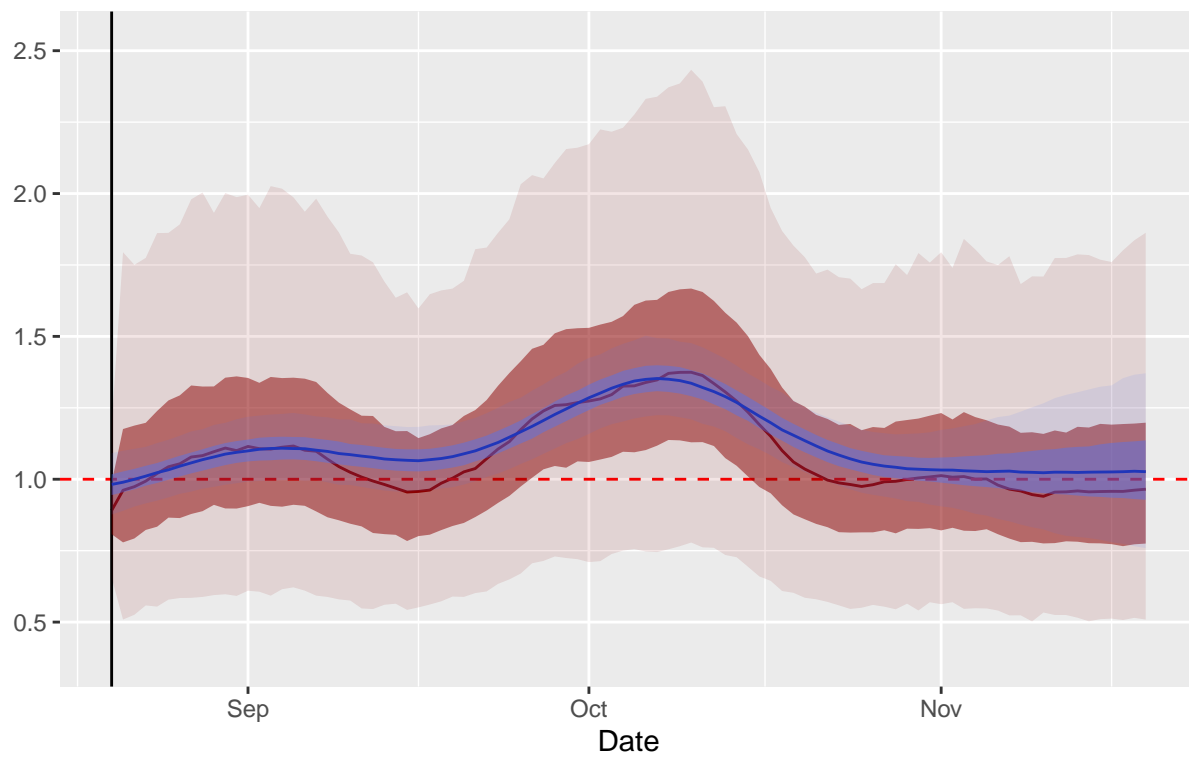
hierarchical model n.1 (red) and single region model (blue)



```
p2 + ylim(c(ymin, ymax))
```

Rt Lazio

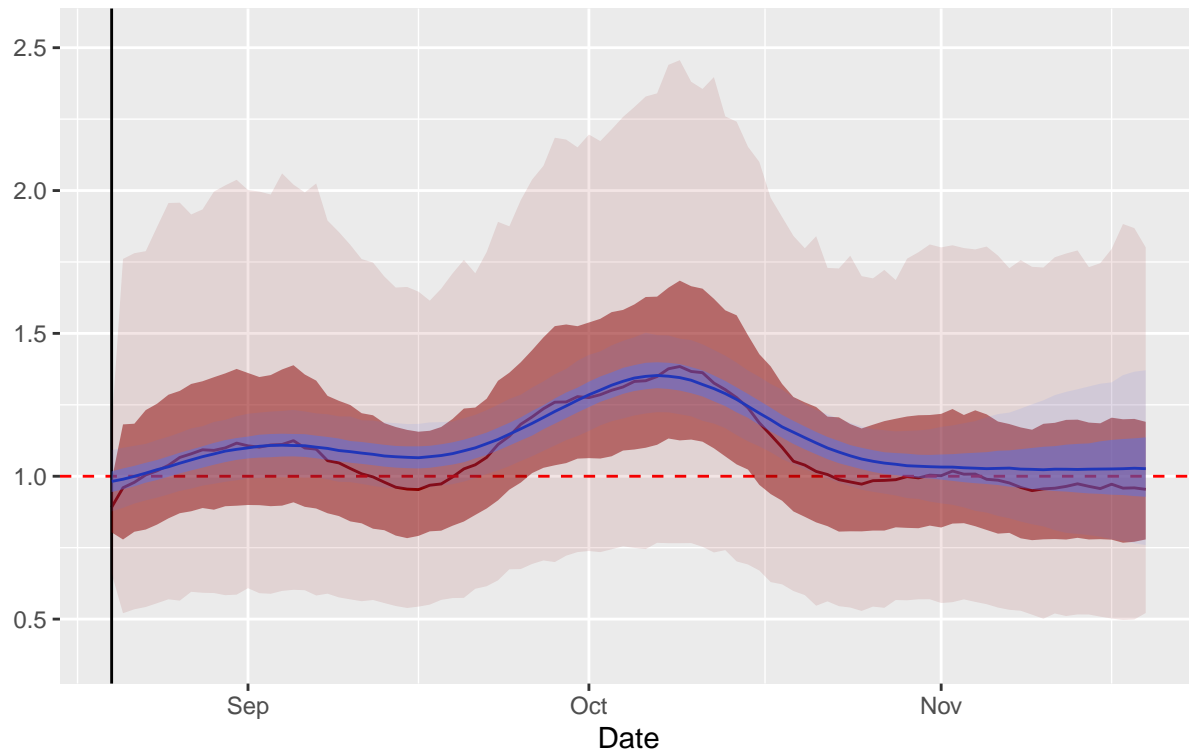
hierarchical model n.2 (red) and single region model (blue)



```
p3 + ylim(c(ymin, ymax))
```

Rt Lazio

hierarchical model n.3 (red) and single region model (blue)



Lombardia

```
p1 <- plot_rt(hier_data, fit_hier, regions, 'Lombardia')
p1 <- plot_overlay(p1, 'Lombardia') + labs(subtitle = 'hierarchical model n.1 (red) and single region model (blue)')

p2 <- plot_rt(hier_data, fit_C05, regions, 'Lazio')
p2 <- plot_overlay(p2, 'Lombardia') + labs(subtitle = 'hierarchical model n.2 (red) and single region model (blue)')

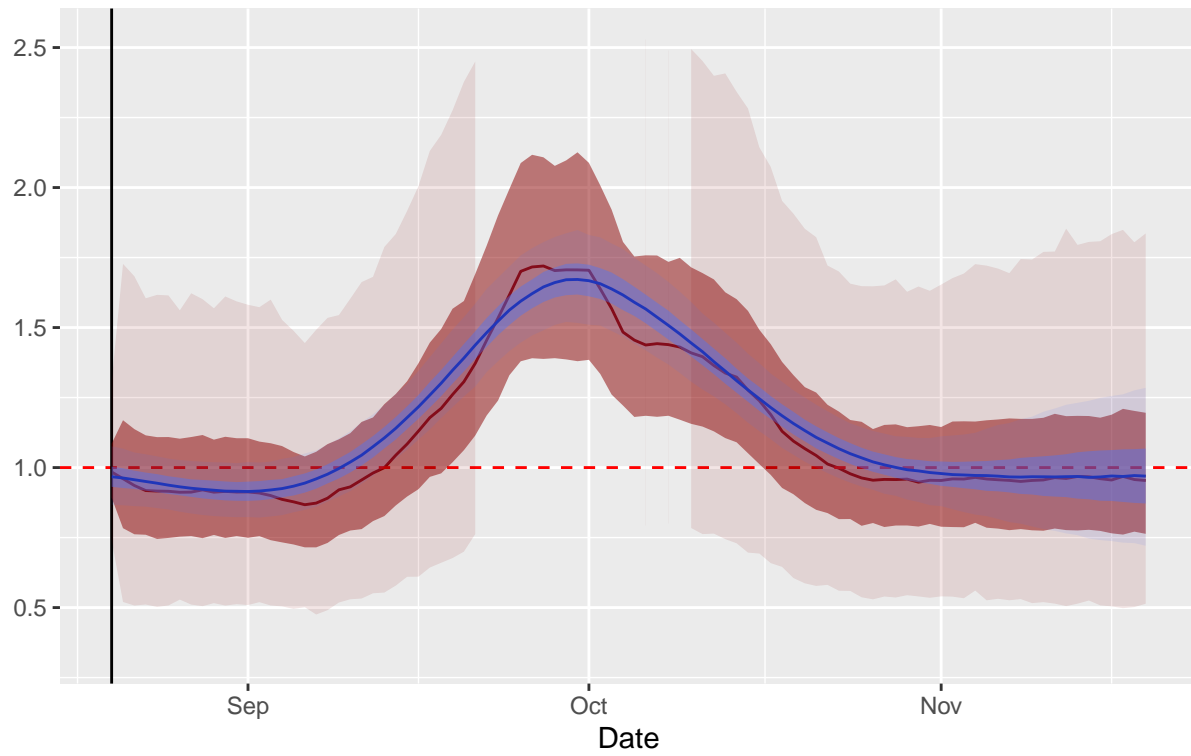
p3 <- plot_rt(hier_data, fit_C1, regions, 'Lazio')
p3 <- plot_overlay(p3, 'Lombardia') + labs(subtitle = 'hierarchical model n.3 (red) and single region model (blue)')

ymin <- min(get_plot_limits(p1)$ymin, get_plot_limits(p2)$ymin, get_plot_limits(p3)$ymin)
ymax <- min(get_plot_limits(p1)$ymax, get_plot_limits(p2)$ymax, get_plot_limits(p3)$ymax)

p1 + ylim(c(ymin, ymax))
```

Rt Lombardia

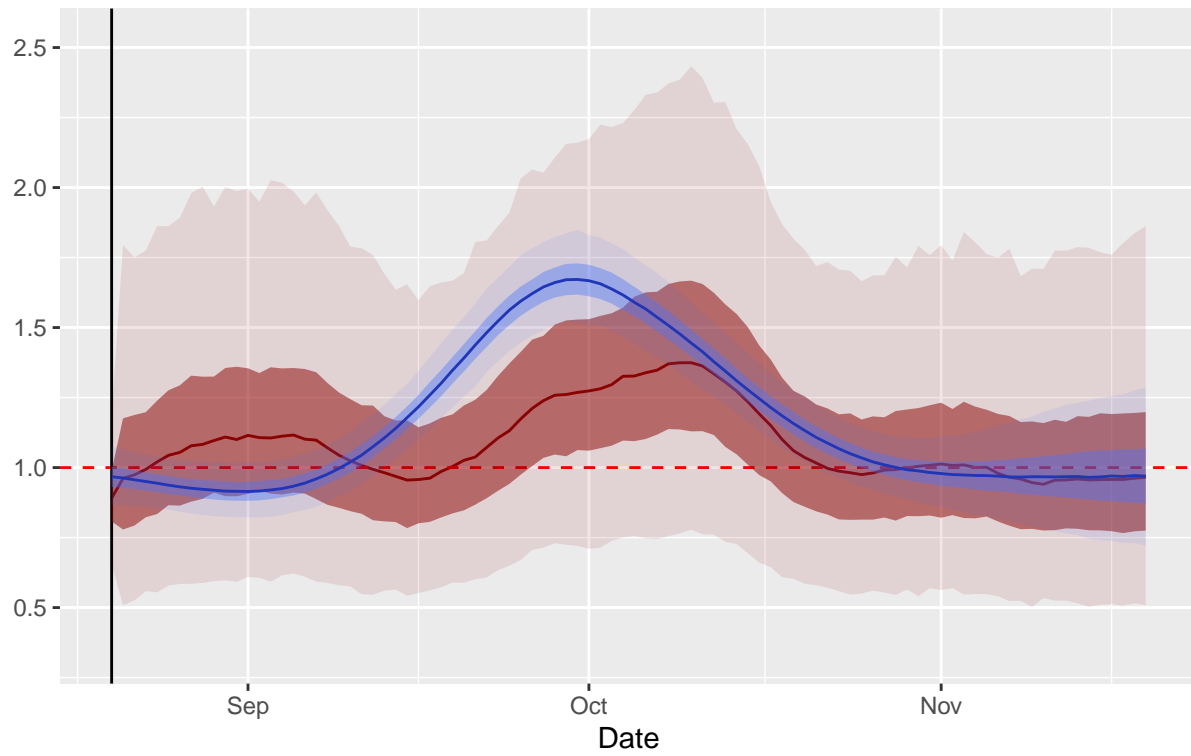
hierarchical model n.1 (red) and single region model (blue)



```
p2 + ylim(c(ymin, ymax))
```

Rt Lombardia

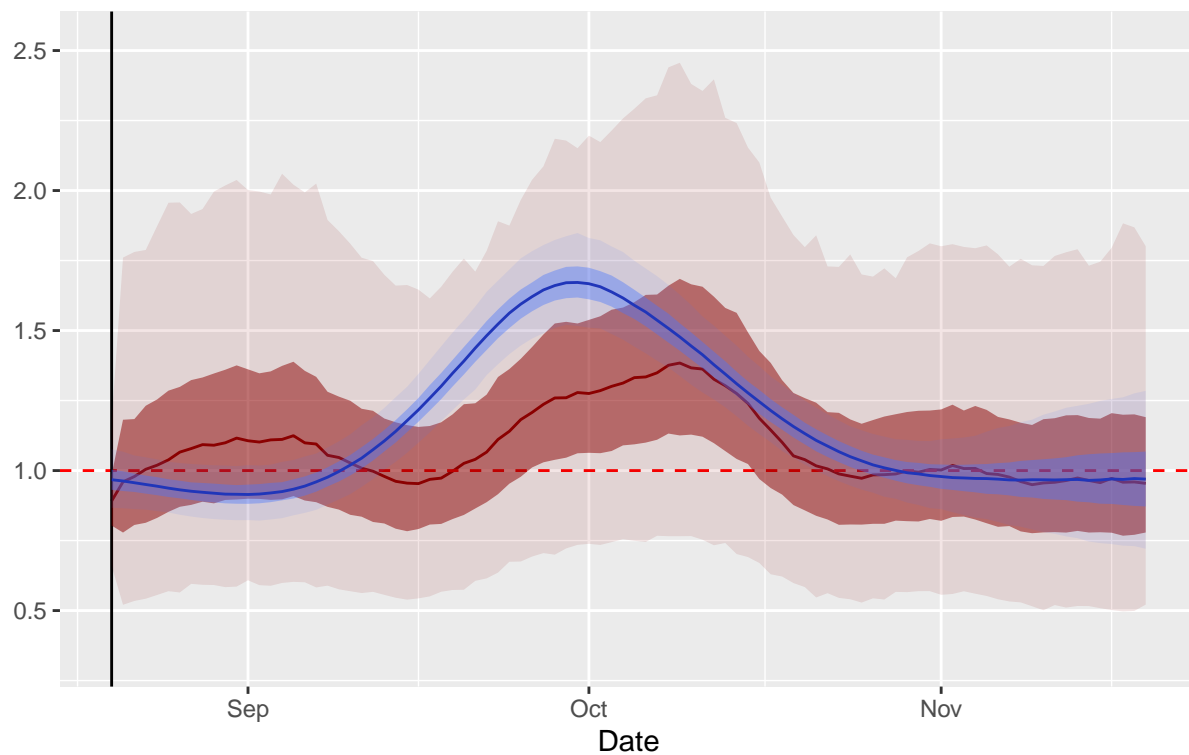
hierarchical model n.2 (red) and single region model (blue)



```
p3 + ylim(c(ymin, ymax))
```

Rt Lombardia

hierarchical model n.3 (red) and single region model (blue)



Veneto

```
p1 <- plot_rt(hier_data, fit_hier, regions, 'Veneto')
p1 <- plot_overlay(p1, 'Veneto') + labs(subtitle = 'hierarchical model n.1 (red) and single region model')

p2 <- plot_rt(hier_data, fit_C05, regions, 'Lazio')
p2 <- plot_overlay(p2, 'Veneto') + labs(subtitle = 'hierarchical model n.2 (red) and single region model')

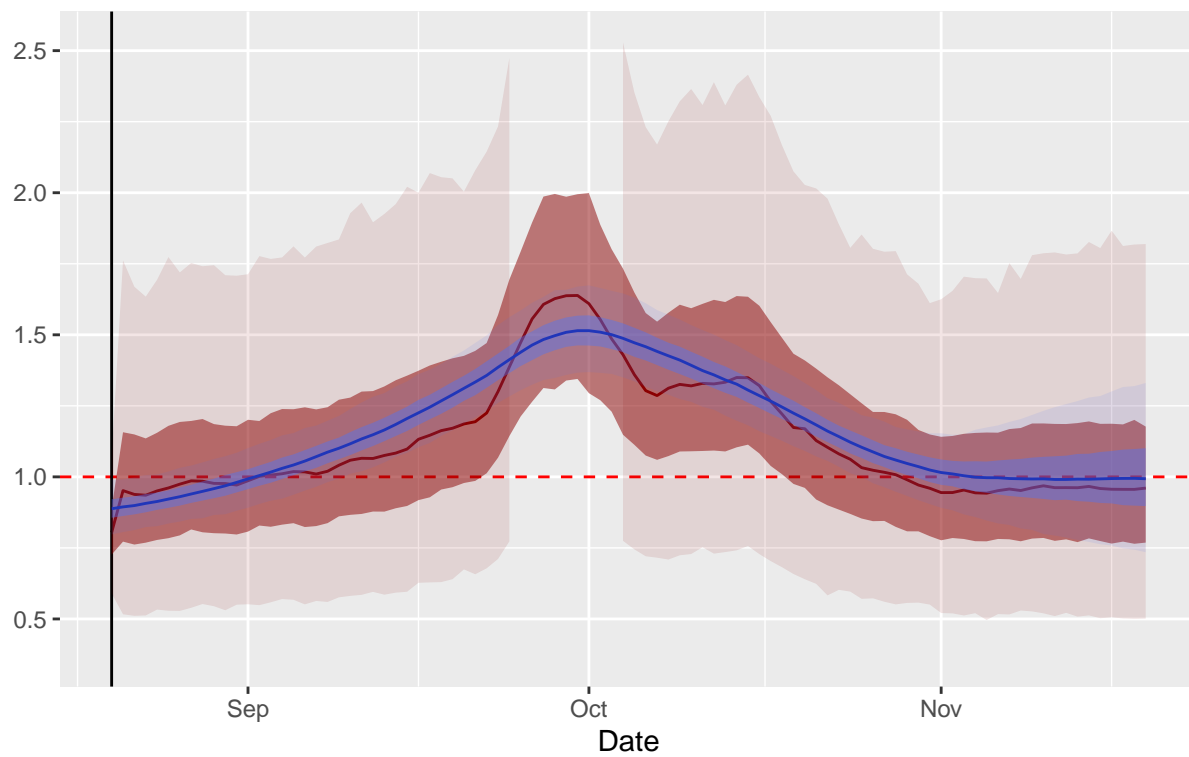
p3 <- plot_rt(hier_data, fit_C1, regions, 'Lazio')
p3 <- plot_overlay(p3, 'Veneto') + labs(subtitle = 'hierarchical model n.3 (red) and single region model')

ymin <- min(get_plot_limits(p1)$ymin, get_plot_limits(p2)$ymin, get_plot_limits(p3)$ymin)
ymax <- min(get_plot_limits(p1)$ymax, get_plot_limits(p2)$ymax, get_plot_limits(p3)$ymax)

p1 + ylim(c(ymin, ymax))
```

Rt Veneto

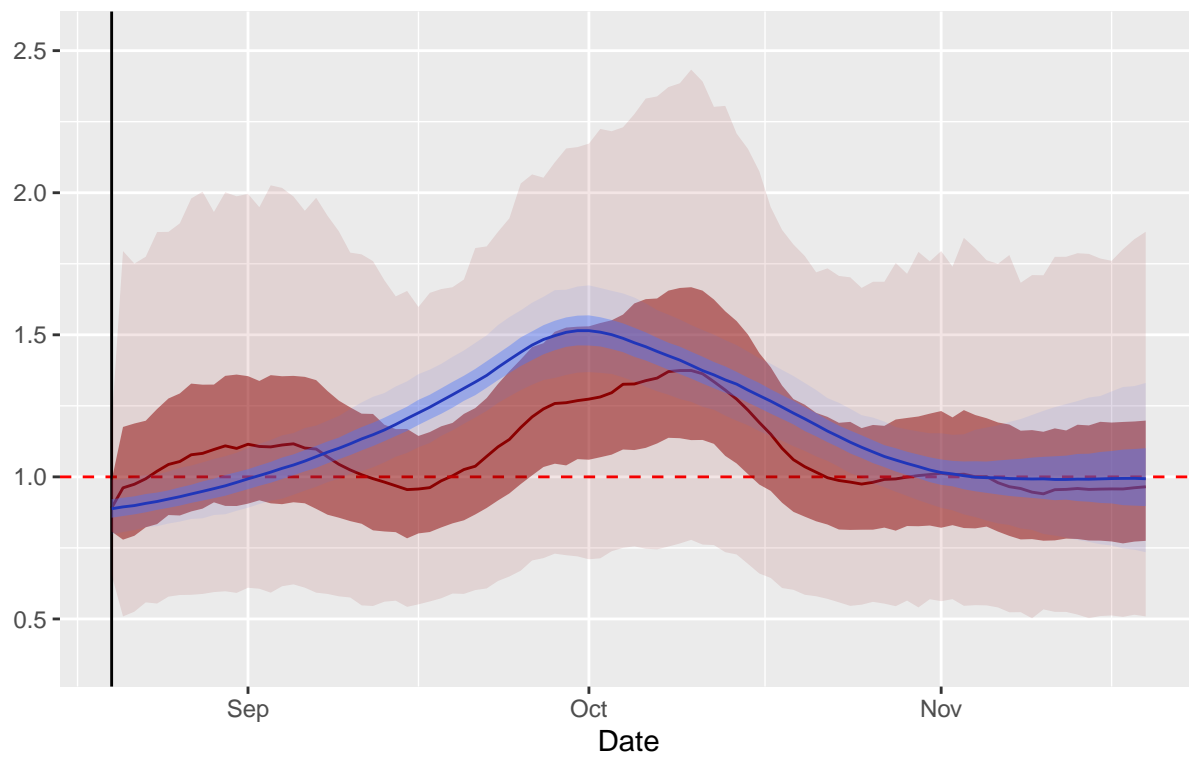
hierarchical model n.1 (red) and single region model (blue)



```
p2 + ylim(c(ymin, ymax))
```


Rt Veneto

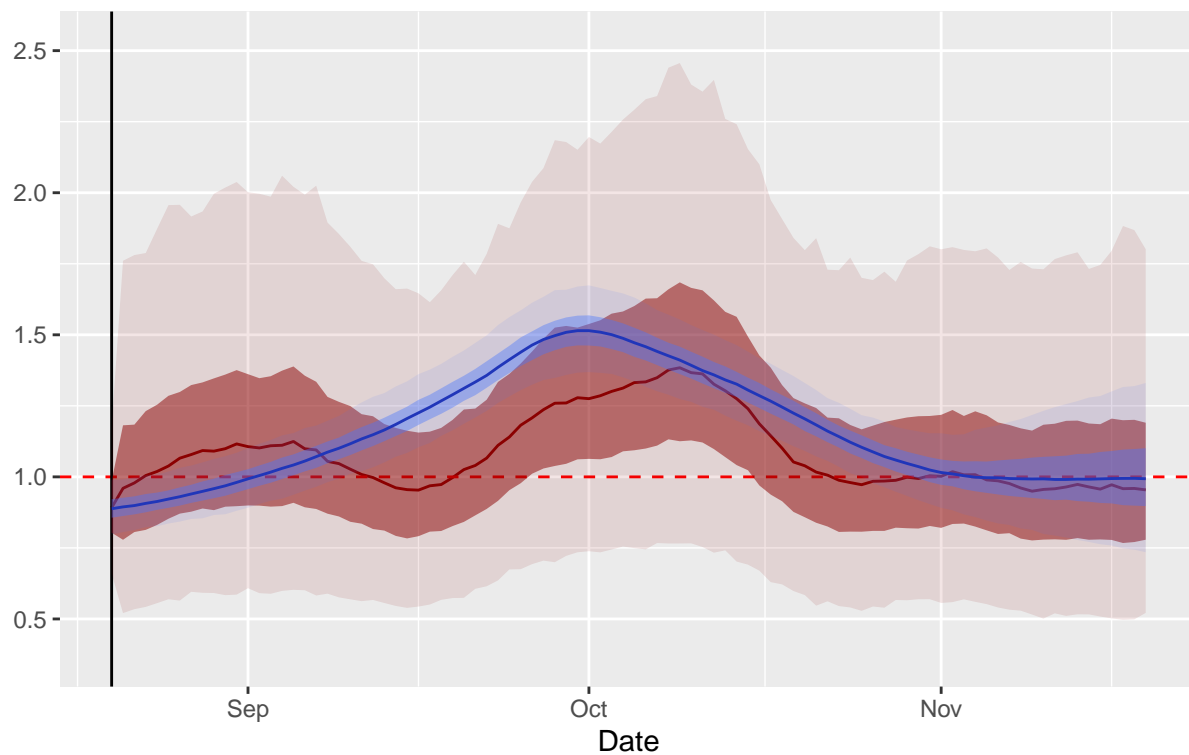
hierarchical model n.2 (red) and single region model (blue)



```
p3 + ylim(c(ymin, ymax))
```

Rt Veneto

hierarchical model n.3 (red) and single region model (blue)



FVG

```
p1 <- plot_rt(hier_data, fit_hier, regions, 'Friuli Venezia Giulia')
p1 <- plot_overlay(p1, 'Friuli Venezia Giulia') + labs(subtitle = 'hierarchical model n.1 (red) and single region model (blue)')

p2 <- plot_rt(hier_data, fit_C05, regions, 'Lazio')
p2 <- plot_overlay(p2, 'Friuli Venezia Giulia') + labs(subtitle = 'hierarchical model n.2 (red) and single region model (blue)')

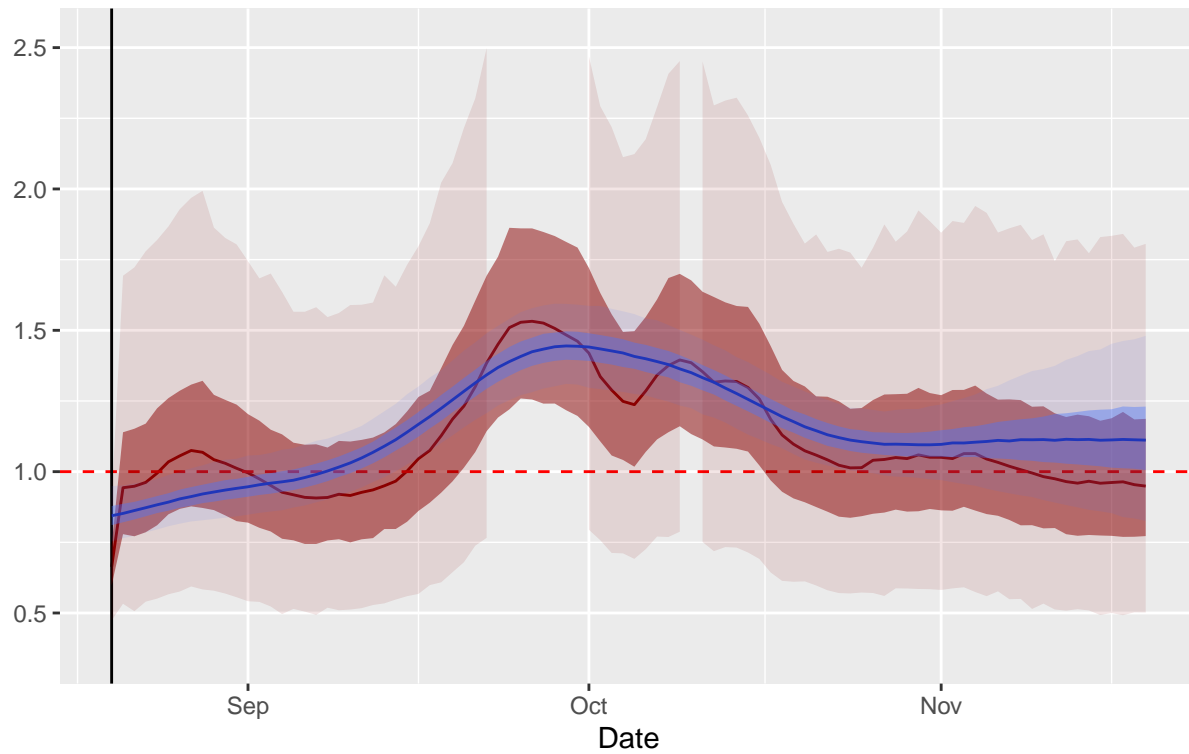
p3 <- plot_rt(hier_data, fit_C1, regions, 'Lazio')
p3 <- plot_overlay(p3, 'Friuli Venezia Giulia') + labs(subtitle = 'hierarchical model n.3 (red) and single region model (blue)')

ymin <- min(get_plot_limits(p1)$ymin, get_plot_limits(p2)$ymin, get_plot_limits(p3)$ymin)
ymax <- min(get_plot_limits(p1)$ymax, get_plot_limits(p2)$ymax, get_plot_limits(p3)$ymax)

p1 + ylim(c(ymin, ymax))
```

Rt Friuli Venezia Giulia

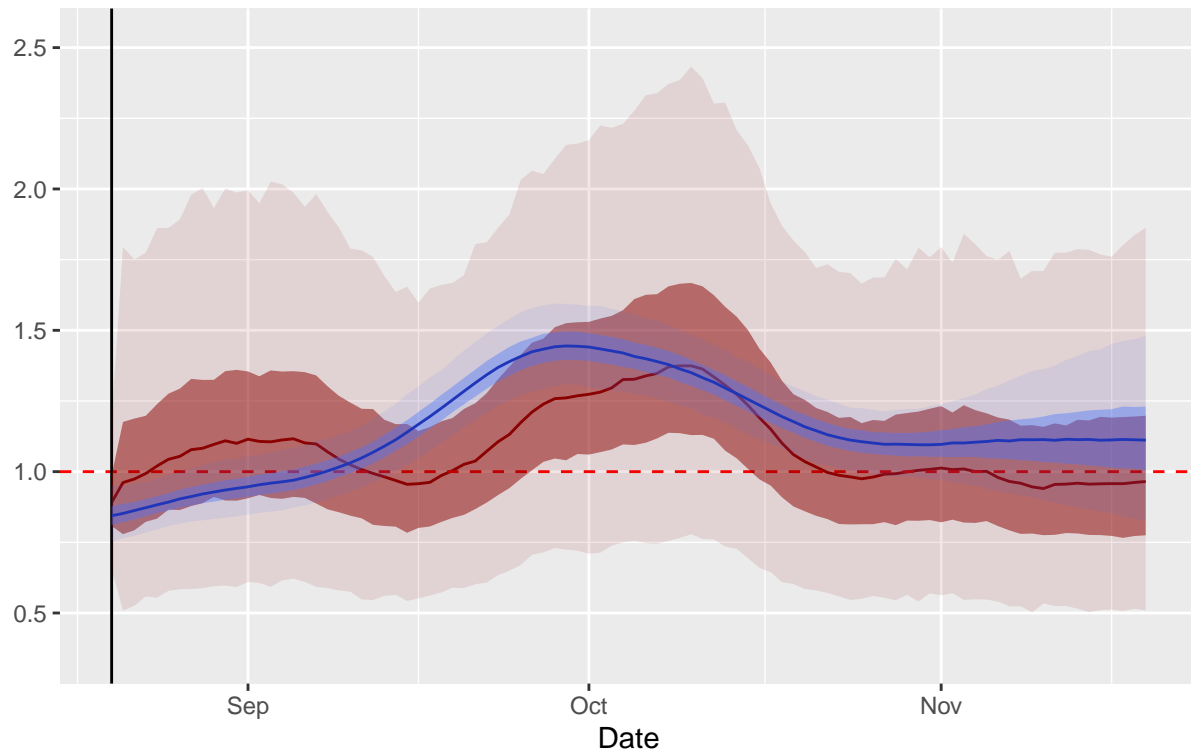
hierarchical model n.1 (red) and single region model (blue)



```
p2 + ylim(c(ymin, ymax))
```

Rt Friuli Venezia Giulia

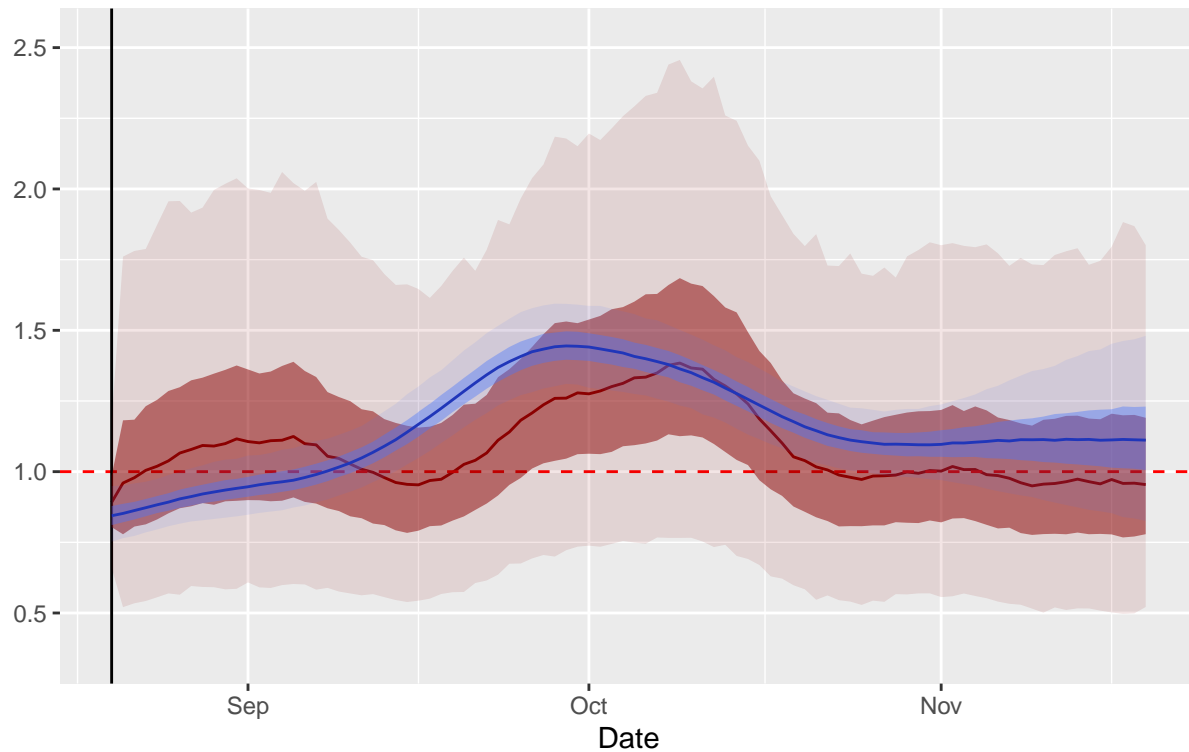
hierarchical model n.2 (red) and single region model (blue)



```
p3 + ylim(c(ymin, ymax))
```

Rt Friuli Venezia Giulia

hierarchical model n.3 (red) and single region model (blue)



Abruzzo

```
p1 <- plot_rt(hier_data, fit_hier, regions, 'Abruzzo')
p1 <- plot_overlay(p1, 'Abruzzo') + labs(subtitle = 'hierarchical model n.1 (red) and single region model')

p2 <- plot_rt(hier_data, fit_C05, regions, 'Lazio')
p2 <- plot_overlay(p2, 'Abruzzo') + labs(subtitle = 'hierarchical model n.2 (red) and single region model')

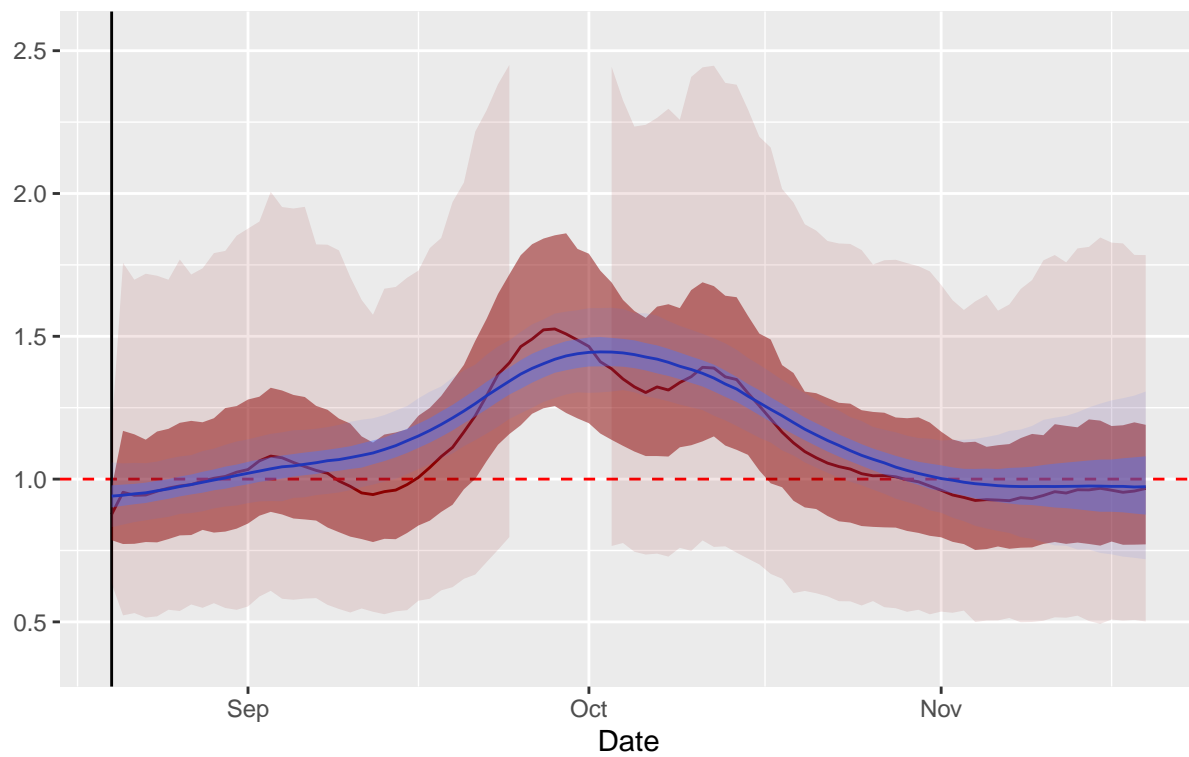
p3 <- plot_rt(hier_data, fit_C1, regions, 'Lazio')
p3 <- plot_overlay(p3, 'Abruzzo') + labs(subtitle = 'hierarchical model n.3 (red) and single region model')

ymin <- min(get_plot_limits(p1)$ymin, get_plot_limits(p2)$ymin, get_plot_limits(p3)$ymin)
ymax <- min(get_plot_limits(p1)$ymax, get_plot_limits(p2)$ymax, get_plot_limits(p3)$ymax)

p1 + ylim(c(ymin, ymax))
```

Rt Abruzzo

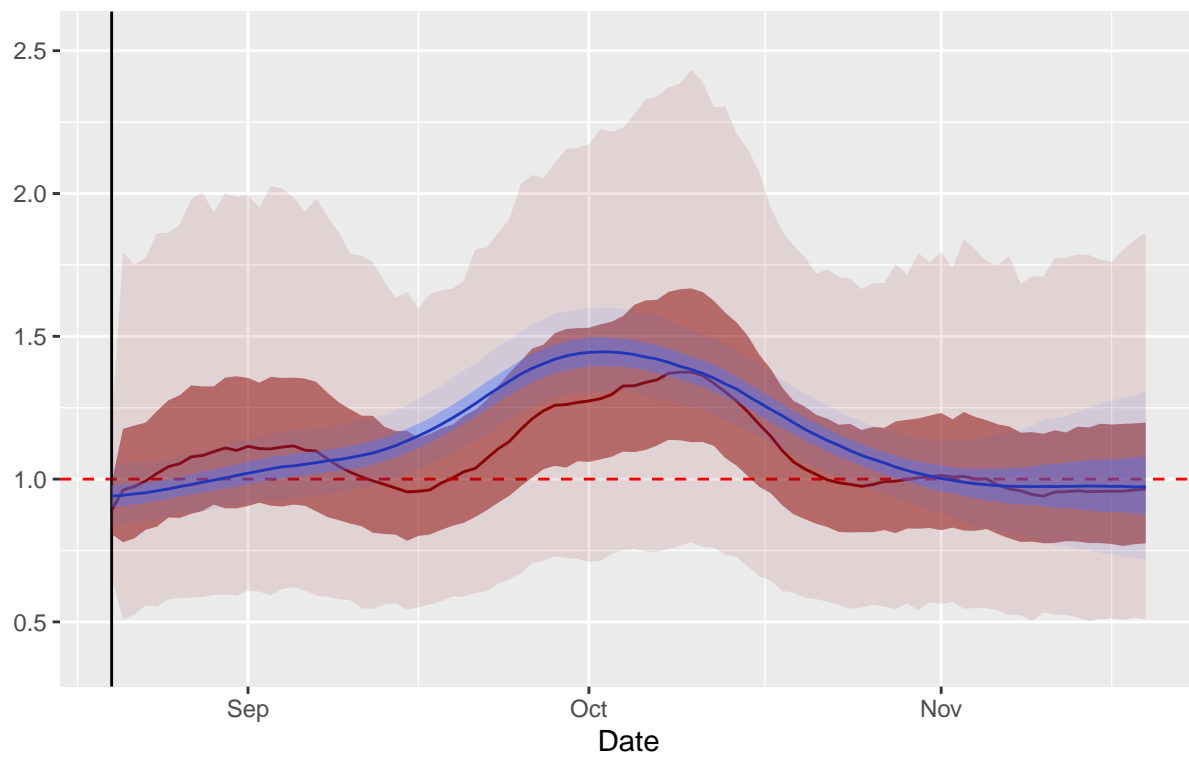
hierarchical model n.1 (red) and single region model (blue)



```
p2 + ylim(c(ymin, ymax))
```

Rt Abruzzo

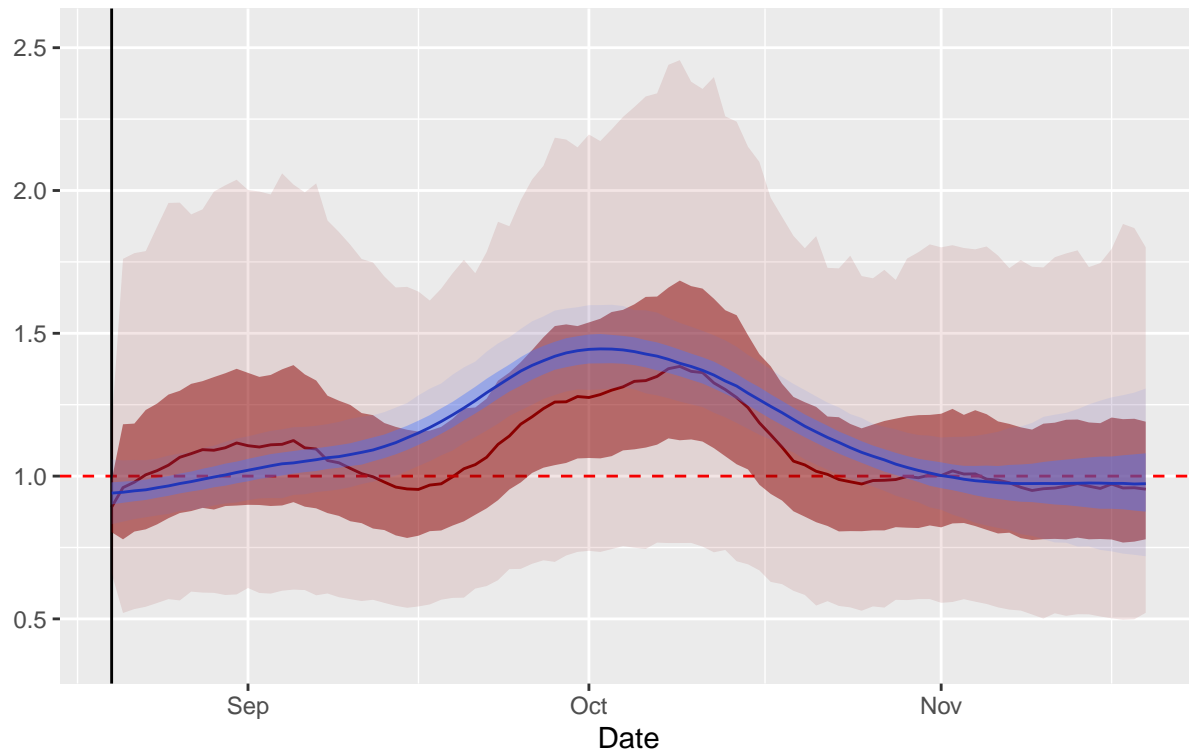
hierarchical model n.2 (red) and single region model (blue)



```
p3 + ylim(c(ymin, ymax))
```

Rt Abruzzo

hierarchical model n.3 (red) and single region model (blue)



Model comparison

```
log_lik_1 <- extract_log_lik(fit_hier)
loo_1 <- loo(log_lik_1)
waic_1 <- waic(log_lik_1)
```

```
log_lik_2 <- extract_log_lik(fit_C1)
loo_2 <- loo(log_lik_2)
waic_2 <- waic(log_lik_2)
```

```
log_lik_3 <- extract_log_lik(fit_C05)
loo_3 <- loo(log_lik_3)
waic_3 <- waic(log_lik_3)
```

```
loo_compare(loo_1, loo_2, loo_3)
```

```
##      elpd_diff se_diff
## model2  0.0      0.0
## model3 -0.2      0.2
## model1 -0.6      0.4
```

looic

```
loo_1$estimates[3,1]
```

```
## [1] 9243.193
```



```

loo_2$estimates[3,1]

## [1] 9242.059
loo_3$estimates[3,1]

## [1] 9242.554

waic
loo_compare(waic_1, waic_2, waic_3)

##          elpd_diff se_diff
## model2  0.0        0.0
## model3 -0.2        0.3
## model1 -0.5        0.3
waic_1$estimates[3,1]

## [1] 9241.65
waic_2$estimates[3,1]

## [1] 9240.689
waic_3$estimates[3,1]

## [1] 9241.102

```