SAVEMORE analysis

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1. Data overview

- Exposure: anakinra vs. placebo.
- $\bullet\,$ Survival outcome: 28-day and 90-day survival.
- $\bullet\,$ Mediator: IL6 on days 1, 4 and 7.

2. Descriptives

2.1. Table 1

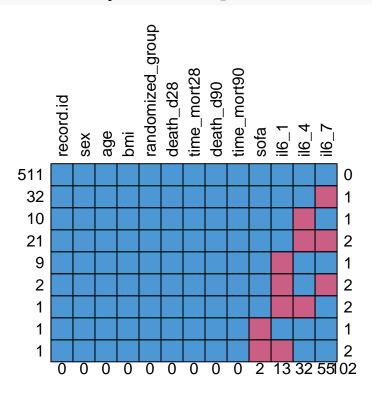
##	:	Stratified by	randomized_group
##		Placebo	Anakinra
##	n	187	401
##	death_d28 = 1 (%)	13 (7.0)	13 (3.2)
##	$death_d90 = 1 (\%)$	19 (10.2)	22 (5.5)
##	il6_1 (mean (SD))	1.26 (0.57)	1.20 (0.54)
##	il6_4 (mean (SD))	1.05 (0.59)	0.91 (0.55)
##	il6_7 (mean (SD))	0.99 (0.63)	0.80 (0.57)
##	sex = male (%)	106 (56.7)	235 (58.6)
##	age (mean (SD))	61.53 (11.33)	62.09 (12.43)

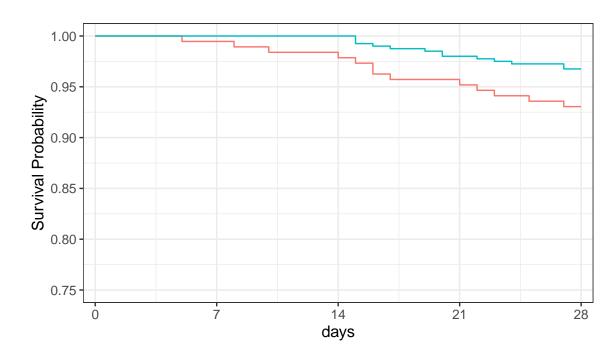
```
## bmi (mean (SD)) 29.73 (5.49) 29.40 (5.53)
## sofa (mean (SD)) 2.46 (1.15) 2.38 (1.11)
```

2.2. Missingness Of 593 patients, 5 have no IL6 biomarker measures. They are not included in any of the analyses.

For the 588 subjects who do have IL6 measures, this is the pattern of missingness:

missing_savemore <- mice::md.pattern(savemore_wide, rotate.names = TRUE, plot = TRUE)</pre>





Placebo -

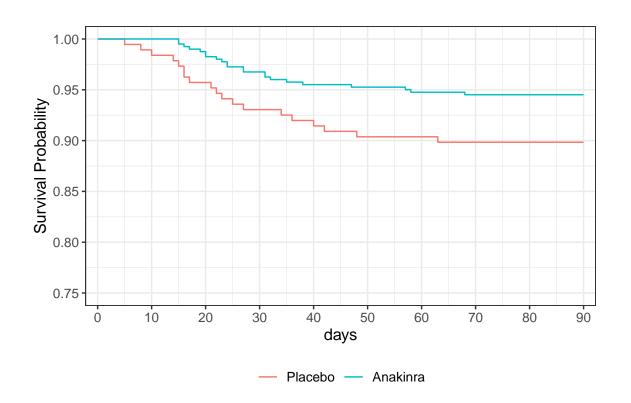
Anakinra

2.3. 28-day survival

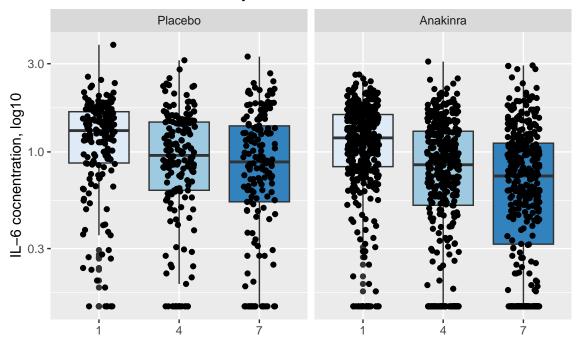
2.4. 90-day survival

```
class(savemore_surv$death_d90) <- "integer"

survfit2(Surv(time_mort90, death_d90) ~ randomized_group, data = savemore_surv) %>%
    ggsurvfit() +
    scale_ggsurvfit(x_scales= list(breaks = c(0, 10, 20, 30, 40, 50, 60, 70, 80, 90)))+
    ylim(c(.75, 1))+
    xlab("days")
```



IL-6 concentration on days 1, 4 and 7



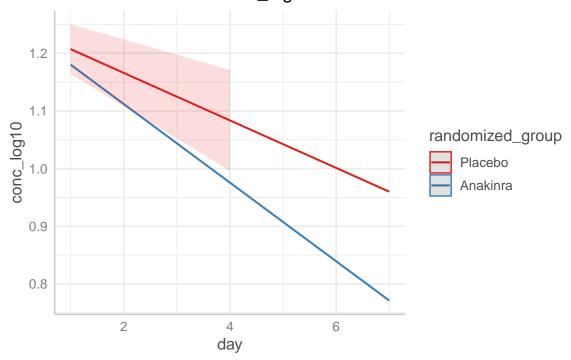
2.5. IL-6 over time

3. Models

3.1. Linear-mixed model for IL-6 over time

```
## Linear mixed-effects model fit by REML
     Data: savemore_long
##
##
          AIC
                   BIC
                          logLik
     2663.727 2701.633 -1324.863
##
##
## Random effects:
## Formula: ~day | record.id
## Structure: General positive-definite, Log-Cholesky parametrization
##
              StdDev
                          Corr
## (Intercept) 0.34478060 (Intr)
## day
              0.05371349 -0.28
## Residual
              0.42544483
##
## Fixed effects: conc_log10 ~ day:randomized_group + day
                                     Value
                                             Std.Error
                                                         DF t-value p-value
                                 1.2484091 0.024396881 1074 51.17085
## (Intercept)
                                                                       0e+00
## day
                                -0.0412061 0.007195113 1074 -5.72695
                                                                       0e+00
## day:randomized_groupAnakinra -0.0269798 0.007669486 1074 -3.51780
                                                                       5e-04
## Correlation:
##
                                (Intr) day
## day
                                -0.467
## day:randomized_groupAnakinra 0.004 -0.736
##
## Standardized Within-Group Residuals:
           Min
                        Q1
                                   Med
                                                QЗ
## -2.54974446 -0.58093507 -0.06755401 0.53782487 4.71086242
##
## Number of Observations: 1664
## Number of Groups: 588
## Approximate 95% confidence intervals
##
## Fixed effects:
##
                                      lower
                                                   est.
                                                               upper
## (Intercept)
                                 1.20053819 1.24840915 1.29628010
                                -0.05532416 -0.04120608 -0.02708801
## day
## day:randomized_groupAnakinra -0.04202863 -0.02697975 -0.01193088
library(sjPlot)
library(sjmisc)
theme_set(theme_sjplot())
plot_model(lmefit.savemore, type = "int", terms = c("randomized_group", "day"))
```

Predicted values of conc_log10



```
# save the interaction estimate
a_res <- get_int(lmefit.savemore, "randomized_groupAnakinra")
saveRDS(a_res, "SAVEMORE_beta_est.rds")</pre>
```

3.2. Cox proportional hazards models For 28-day and 90-day survival.

```
# Fit cox proportional hazard model
coxfit.savemore28 <- coxph(Surv(time_mort28, death_d28) ~ randomized_group, data = savemore_surv, x = T.
summary(coxfit.savemore28)</pre>
```

3.2.1. 28-day survival

```
## Call:
## coxph(formula = Surv(time_mort28, death_d28) ~ randomized_group,
##
      data = savemore_surv, x = TRUE)
##
    n= 588, number of events= 26
##
##
##
                              coef exp(coef) se(coef)
                                                          z Pr(>|z|)
## randomized_groupAnakinra -0.7883
                                      0.4546 0.3922 -2.01
##
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
##
##
                           exp(coef) exp(-coef) lower .95 upper .95
                                            2.2
                                                   0.2108
                                                             0.9807
## randomized_groupAnakinra
                              0.4546
## Concordance= 0.595 (se = 0.049)
```

```
= 4.04 on 1 df, p=0.04
## Wald test
## Score (logrank) test = 4.25 on 1 df,
                                        p=0.04
confint(coxfit.savemore28) %>% exp() %>% round(3)
                          2.5 % 97.5 %
## randomized_groupAnakinra 0.211 0.981
# Fit cox proportional hazard model
coxfit.savemore90 <- coxph(Surv(time_mort90, death_d90) ~ randomized_group, data = savemore_surv, x = T
summary(coxfit.savemore90)
3.2.2. 90-day survival
## Call:
## coxph(formula = Surv(time_mort90, death_d90) ~ randomized_group,
      data = savemore_surv, x = TRUE)
##
##
    n= 588, number of events= 41
##
##
##
                             coef exp(coef) se(coef)
                                                        z Pr(>|z|)
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
##
##
                          exp(coef) exp(-coef) lower .95 upper .95
## randomized_groupAnakinra
                             0.5225
                                        1.914
                                                 0.2828
## Concordance= 0.577 (se = 0.039)
## Likelihood ratio test= 4.16 on 1 df, p=0.04
                    = 4.3 on 1 df,
                                       p=0.04
## Score (logrank) test = 4.45 on 1 df,
                                        p=0.03
confint(coxfit.savemore90) %>% exp() %>% round(3)
                          2.5 % 97.5 %
## randomized_groupAnakinra 0.283 0.965
3.3. Joint models Using 28- and 90-day survival as endpoints.
set.seed(15)
# Fit joint model for 28 day survival
jointfit.savemore_28 <- JMbayes2::jm(coxfit.savemore28, lmefit.savemore, time_var = "day", n_chains = "
# save results
saveRDS(jointfit.savemore_28, "jointfit.savemore_28.rds")
# Fit joint model for 90 day survival
jointfit.savemore_90 <- JMbayes2::jm(coxfit.savemore90, lmefit.savemore, time_var = "day", n_chains = "
saveRDS(jointfit.savemore_90, "jointfit.savemore_90.rds")
```

p=0.05

Likelihood ratio test= 3.94 on 1 df,

3.3.1. 28-day endpoint

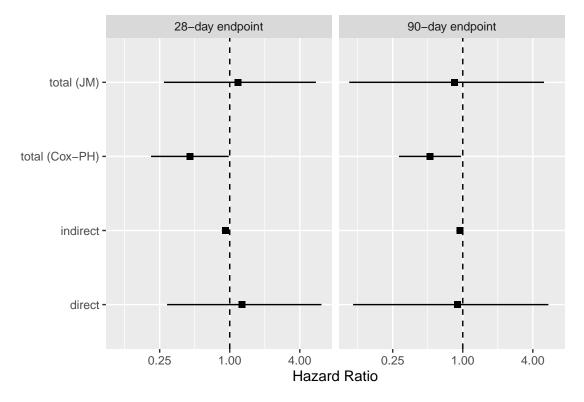
```
##
## Call:
## JMbayes2::jm(Surv_object = coxfit.savemore28, Mixed_objects = lmefit.savemore,
       time_var = "day", n_chains = 2, n_iter = 50000L, n_burnin = 5000L,
##
       n thin = 5)
##
##
## Data Descriptives:
## Number of Groups: 592
                                Number of events: 26 (4.4%)
## Number of Observations:
     conc_log10: 1661
##
##
##
                            WAIC
                                       LPML
                    DIC
## marginal
               2907.000 2971.415 -1550.877
## conditional 1460.744 2731.342 -21749.354
## Random-effects covariance matrix:
##
          StdDev
##
                   Corr
## (Intr) 0.3143 (Intr)
## day
         0.0415 0.0869
## Survival Outcome:
                              Mean StDev
                                             2.5% 97.5%
                                                              Р
## randomized_groupAnakinra 0.2423 0.7734 -1.2396 1.8143 0.7589 1.0001
                            3.0819 0.9965 1.4228 5.3825 0.0000 1.0314
## value(conc_log10)
## Longitudinal Outcome: conc_log10 (family = gaussian, link = identity)
                  Mean StDev
                                 2.5%
                                        97.5%
                                                   Ρ
## (Intercept) 1.1850 0.0212 1.1427 1.2265 0.0000 1.0038
## day
               -0.0409 0.0077 -0.0560 -0.0258 0.0000 1.0017
               -0.0260 0.0085 -0.0431 -0.0094 0.0016 1.0001
## d:_A
               0.4344 0.0109 0.4133 0.4563 0.0000 1.0011
## sigma
##
## MCMC summary:
## chains: 2
## iterations per chain: 50000
## burn-in per chain: 5000
## thinning: 5
## time: 12.1 min
3.3.2. 90-day endpoint
##
## JMbayes2::jm(Surv_object = coxfit.savemore90, Mixed_objects = lmefit.savemore,
       time_var = "day", n_chains = 2, n_iter = 50000L, n_burnin = 5000L,
##
##
       n_{thin} = 5
## Data Descriptives:
## Number of Groups: 592
                                Number of events: 41 (6.9%)
## Number of Observations:
##
     conc_log10: 1661
##
```

```
##
                       DIC
                                    WAIC
                                                  LPML
                  3311.347 6.021164e+03 -4.206220e+03
## marginal
## conditional 3447233.293 6.678743e+16 -6.634874e+10
## Random-effects covariance matrix:
##
          StdDev
##
                   Corr
## (Intr) 0.3224 (Intr)
##
  day
          0.0667 -0.1341
##
## Survival Outcome:
                               Mean StDev
                                               2.5% 97.5%
##
                                                                     Rhat
## randomized_groupAnakinra -0.1051 0.9359 -2.1706 1.6944 0.9277 1.0621
                             1.9649 0.5242 0.9024 2.7771 0.0000 1.3804
## value(conc_log10)
##
## Longitudinal Outcome: conc_log10 (family = gaussian, link = identity)
##
                  Mean StDev
                                 2.5%
                                         97.5%
                                                    Ρ
## (Intercept) 1.1629 0.0295 1.1044 1.2164 0.0000 1.3484
               -0.0308 0.0136 -0.0535 -0.0030 0.0254 1.4917
## day
## d: A
               -0.0290 0.0105 -0.0506 -0.0094 0.0030 1.0976
## sigma
                0.4282 0.0113 0.4063 0.4508 0.0000 1.0032
##
## MCMC summary:
## chains: 2
## iterations per chain: 50000
## burn-in per chain: 5000
## thinning: 5
## time: 5 min
4. Results
4.1. Indirect, direct, and total effects Of anakinra through IL6 on the hazard of death.
res28 <- get_effects(jointfit.savemore_28, coxfit.savemore28,</pre>
                     "randomized_groupAnakinra") %>% cbind(endpoint =c("28-day endpoint"))
res28
##
             effect
                                  CI_lower
                                               CI_upper
                                                                endpoint
                             est
## 1
             direct 0.24229995 -1.2395534 1.81427247 28-day endpoint
## 2
           indirect -0.08003438 -0.1567002 -0.02451719 28-day endpoint
## 3 total (Cox-PH) -0.78825892 -1.5570289 -0.01948896 28-day endpoint
         total (JM) 0.16226557 -1.3027106 1.70492160 28-day endpoint
res90 <- get_effects(jointfit.savemore_90, coxfit.savemore90,</pre>
                     "randomized_groupAnakinra") %>% cbind(endpoint =c("90-day endpoint"))
res90
##
                            est CI_lower
                                              CI_upper
                                                               endpoint
             effect
## 1
             direct -0.10505573 -2.170580
                                           1.69444173 90-day endpoint
           indirect -0.05689723 -0.126946 -0.01364629 90-day endpoint
## 3 total (Cox-PH) -0.64909520 -1.262941 -0.03524972 90-day endpoint
```

total (JM) -0.16195296 -2.246248 1.60819108 90-day endpoint

```
res <- rbind(res28, res90)
saveRDS(res, "savemore_res.rds")

res %>%
    ggplot(aes(y = effect))+
    theme_grey()+
    geom_point(aes(x=exp(est)), shape=15, size=2) +
    geom_linerange(aes(xmin=exp(CI_lower), xmax=exp(CI_upper))) +
    geom_vline(xintercept = 1, linetype="dashed") +
    labs(x="Hazard Ratio", y= "")+
    scale_x_continuous(trans = "log2")+
    facet_grid(~endpoint)
```



4.2. Association parameter Hazard ratio estimate and 95% CI for the association parameter α for a one unit increase (at any time point) of IL-6 and the hazard of death over 28 days.

```
# save association estimates
alpha_28 <- get_alpha(jointfit.savemore_28, "28-day endpoint")
alpha_90 <- get_alpha(jointfit.savemore_90, "90-day endpoint")
alpha_est <- rbind(alpha_28, alpha_90)
saveRDS(alpha_est, "SAVEMORE_alpha_est.rds")</pre>
```

4.4. Conclusions

- From lime and JM we can conclude that there is an effect of anakinra over time on IL-6.
- From joint model we conclude 1) that there is no direct effect of anakinra on survival when controlling

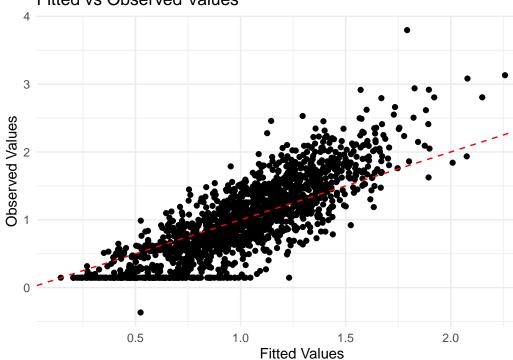
for IL-6, 2) there is an association between IL-6 and survival, and (3) some of the effect of anakinra is mediated by IL-6.

5. Model checks

```
# get fitted values
fitted_values<- fitted(lmefit.savemore)
savemore_long <- savemore_long %>% drop_na(conc_log10)

# plot observed vs fitted values
ggplot(data = savemore_long, aes(x = fitted_values, y = conc_log10)) +
    geom_point() +
    geom_abline(slope = 1, intercept = 0, linetype = "dashed", color = "red") + # Line of perfect fit
    labs(x = "Fitted Values", y = "Observed Values") +
    ggtitle("Fitted vs Observed Values") +
    theme_minimal()
```

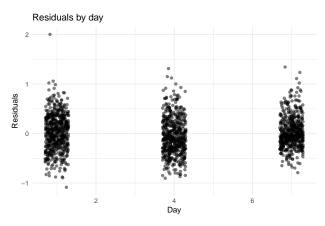
Fitted vs Observed Values



5.1. Longitudinal submodel

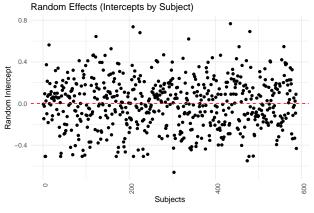
```
# get residuals
residuals_values <- resid(lmefit.savemore)

# plot residuals vs time
ggplot(savemore_long, aes(x = day, y = residuals_values)) +
  geom_jitter(width = 0.3, alpha = 0.5) +
  labs(x = "Day", y = "Residuals") +
  ggtitle("Residuals by day") +
  theme_minimal()</pre>
```

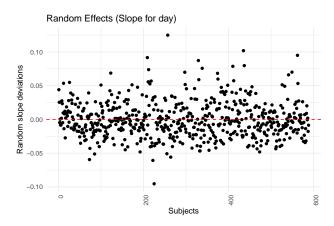


```
# get random effects
random_effects <- ranef(lmefit.savemore)

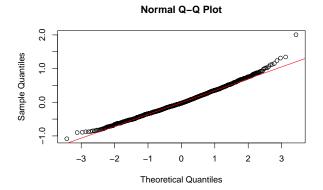
#plot random effects
ggplot(random_effects, aes(x = c(1:nrow(savemore_surv)), y = `(Intercept)`)) +
    geom_point() +
    geom_hline(yintercept = 0, linetype = "dashed", color = "red") +
    labs(x = "Subjects", y = "Random Intercept") +
    ggtitle("Random Effects (Intercepts by Subject)") +
    theme_minimal() +
    theme(axis.text.x = element_text(angle = 90, hjust = 1))</pre>
```

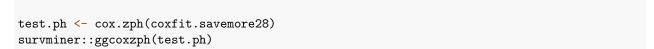


```
#plot random effects
ggplot(random_effects, aes(x = c(1:nrow(savemore_surv)), y = `day`)) +
  geom_point() +
  geom_hline(yintercept = 0, linetype = "dashed", color = "red") +
  labs(x = "Subjects", y = "Random slope deviations") +
  ggtitle("Random Effects (Slope for day)") +
  theme_minimal() +
  theme(axis.text.x = element_text(angle = 90, hjust = 1))
```

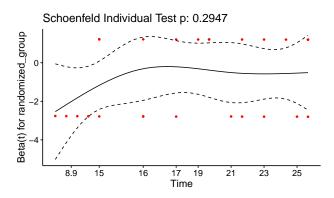


qq plot for residuals
qqnorm(resid(lmefit.savemore))
qqline(resid(lmefit.savemore), col = "red")



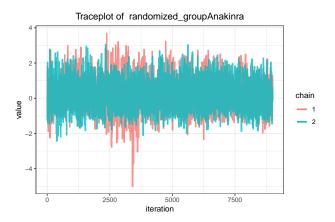


Global Schoenfeld Test p: 0.2947



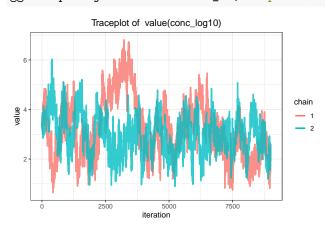
5.2 Survival submodel

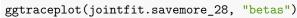
```
ggtraceplot(jointfit.savemore_28, "gammas")
```

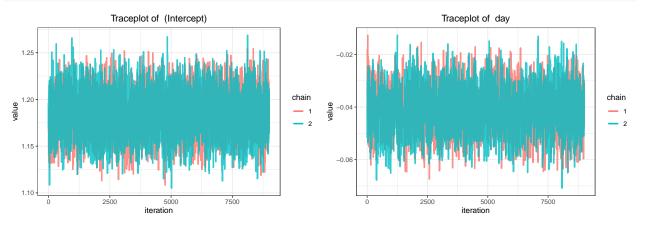


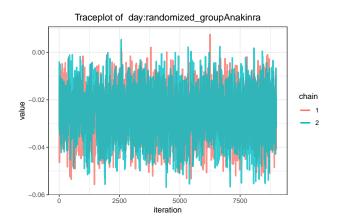
5.1. 28-day endpoint

ggtraceplot(jointfit.savemore_28, "alphas")

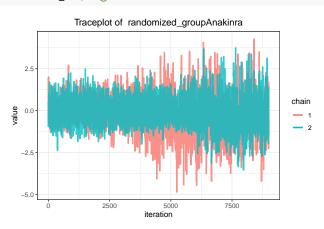






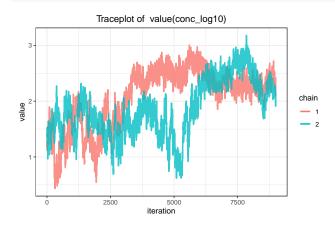


ggtraceplot(jointfit.savemore_90, "gammas")



5.2. 90-day endpoint

ggtraceplot(jointfit.savemore_90, "alphas")



ggtraceplot(jointfit.savemore_90, "betas")

