**290524 - OH**

* Will proceed with Warfarin dataset
* Figure out the way to execute NONMEM from R
* Sent Erwin an email about calculating RSE for CC, and for T-distributed pooled parameter from MI
* Investigating the variability of the estimate, so that MI really shows . Cause what I have now is only from one seed, and what we observe can be due to randomness (cf. counter-intuitive result from 30% missingness)
  + Try 30 different seeds, compare the relative bias – report the mean and standard deviation of different techniques observed. Hopefully we can see that MI is actually the best.
  + Also we want to observe increasing relative bias with CC with increasing % of missingness