DEMO2002 Code Shortcuts

Remove "+" from Cohort column

```
fx$Cohort = gsub("[+]", "", fx$Cohort)
```

ggplot boilerplate/guidance

```
ggplot()+
 geom_line(LT_p[LT_p$Age==0,], mapping =aes(x=Year, y=ex, color="e(0) at age 0"))+
 scale_x_continuous(n.breaks = 10)+
 scale_y_continuous(breaks = seq(0,110,10),limits = c(0,100))+
 scale_color_manual(values = c("navy","red"))+
 guides(color = guide_legend(title = "Year"))+
 labs(x = "Age", y = "lx", title="This is a title")+
 theme_bw()
```

Life Table

```
LifeTableMx<-function(mx,sex){
# and mx would just be age-specific deaths/pop
### N is the number of ages that we have or length(mx)
N<-length(mx)
# person years lived by those dying in the interval
ax<-rep(0.5, N)
## for the first age ax depends on infant mortality and sex
  ax[1]<-ifelse(mx[1]<0.107,0.045+mx[1]*2.684,0.330)}
if(sex=="f"){
  ax[1]<-ifelse(mx[1]<0.107,0.053+2.800*mx[1],0.350)
## transform from death rates to probabilities of dying
qx < -mx / (1 + (1 - ax) * mx)
### with the last value of qx = 1
qx[N] <- 1
## Probability of surviving
## calculating the survival function
lx<-100000
for(y in 1:(N-1)){
  lx[y+1] < -lx[y] *px[y]
## the distribution of deaths
dx < -ax * lx
## the life tables person-years depends on lx, dx and ax
Lx \leftarrow (lx - dx) + ax * dx
## and Lx for the last age-group
Lx[N] \leftarrow ifelse(mx[N] > 0, mx[N], 0)
## person-years from a given age upwards
Tx<-c()
for(y in 1:N){
  Tx[y] \leftarrow sum(Lx[y:N])
## finally life expectancy
ex<- Tx / lx
# ALL<-data.frame(Age,mx,lx,dx,Lx,Tx,ex)
ALL<-data.frame(Age, mx, qx, ax, lx, dx, Lx, Tx, ex)
return(ALL)
```

Focus Points

- 1. Always connect demographic measures to their real-world implications
- 2. Include more policy-relevant interpretations
- 3. Consider historical and social context when analyzing trends
- 4. Strengthen technical explanations with concrete examples
- 5. Make better use of the word limit some answers could be more detailed while staying within 200 words