

Visualization 2: Matching Results

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Preparation

```
## Clean Up Space
rm(list=ls())

## Set Working Directory (Automatically) ##
require(rstudioapi); require(rprojroot)
if (rstudioapi::isAvailable() == TRUE) {
  setwd(dirname(rstudioapi::getActiveDocumentContext()$path));
}
projdir <- find_root(has_file("thisishome.txt"))
cat(paste("Working Directory Set to:\n",projdir))

## Working Directory Set to:
## /home/gentok/GoogleDrive/Projects/Fan-Gento-Lab/ForeignerJapan

setwd(projdir)

## Import Matched Data

d <- readRDS(paste0(projdir, "/data/sifcct_zip_latest_v5.rds"))
dy <- readRDS(paste0(projdir, "/data/sifcct_unmatched_v5.rds"))
dym1 <- readRDS(paste0(projdir, "/data/sifcct_matched_1_all_v5.rds"))
dym2 <- readRDS(paste0(projdir, "/data/sifcct_matched_2_all_v5.rds"))
dym3 <- readRDS(paste0(projdir, "/data/sifcct_matched_3_all_v5.rds"))
dym4 <- readRDS(paste0(projdir, "/data/sifcct_matched_4_all_v5.rds"))
dym5 <- readRDS(paste0(projdir, "/data/sifcct_matched_5_all_v5.rds"))

## Fix Pair ID
updatepairid <- function(dym1) {
  dym1$pair_id[which(dym1$femalebefore==1)] <- paste0("fb_",dym1$pair_id[which(dym1$femalebefore==1)])
  dym1$pair_id[which(dym1$femaleafter==1)] <- paste0("fa_",dym1$pair_id[which(dym1$femaleafter==1)])
  dym1$pair_id[which(dym1$malebefore==1)] <- paste0("mb_",dym1$pair_id[which(dym1$malebefore==1)])
  dym1$pair_id[which(dym1$maleafter==1)] <- paste0("ma_",dym1$pair_id[which(dym1$maleafter==1)])
  return(dym1)
}
dym1 <- updatepairid(dym1)
dym2 <- updatepairid(dym2)
dym3 <- updatepairid(dym3)
dym4 <- updatepairid(dym4)
dym5 <- updatepairid(dym5)
```

```
## Packages
library(lmtest) # For Statistical Test
library(sandwich) # Cluster Robust Standard Error
library(ggplot2) # Plotting
library(grid) # Plotting
library(gridExtra) # Plotting
library(sf) # Plotting Map
library(ggimage) # Plotting Map
library(jpndistrict) # Plotting Japanese Map
library(magrittr) # Data Management/Plotting
library(purrr) # Data Management
library(pbapply) # Apply with Progress Bar
require(ebal) # Matching Balance
require(Matching) # Matching Balance
```

Plotting Individual-Level Predictors Balance

```
## Import Data
bal_dy_unmatched <- readRDS(paste0(projdir, "/data/sifcct_unmatched_balance_v5.rds")) # Young, No Distanc
bal_dy_matched_1 <- readRDS(paste0(projdir, "/data/sifcct_matched_1_balance_v5.rds")) # Young, No Distanc
bal_dy_matched_2 <- readRDS(paste0(projdir, "/data/sifcct_matched_2_balance_v5.rds")) # Young, Distanc
bal_dy_matched_3 <- readRDS(paste0(projdir, "/data/sifcct_matched_3_balance_v5.rds")) # Young, Distanc
bal_dy_matched_4 <- readRDS(paste0(projdir, "/data/sifcct_matched_4_balance_v5.rds")) # Young, Distanc
bal_dy_matched_5 <- readRDS(paste0(projdir, "/data/sifcct_matched_5_balance_v5.rds")) # Young, Distanc

## Raw data balance

## Matching Function
source(paste0(projdir, "/src/findmatch.R"))

fmbal = formula(edu2 ~ female + age + bornyr + lvlen + lvpr +
  c10_sreg_fper + I(sqrt(c10_sreg_fper)) + c10_sreg_foreignN + c10_sreg_pop +
  c10_sreg_edu_ugsP + c10_sreg_edu_ugs + c10_sreg_edu_graduated +
  c10_mun_fper + I(sqrt(c10_mun_fper)) + c10_mun_foreignN + c10_mun_pop +
  c10_mun_edu_ugsP + c10_mun_edu_ugs + c10_mun_edu_graduated +
  zip_did + didper + wave + after + panel)
vnbal = c("Gender", "Age", "Born Year", "Living Length", "Living Proportion",
  "Foreigner Percentage (zip)", "Foreigner Percentage sqrt. (zip)",
  "Foreigner Population (zip)", "Population (zip)",
  "University Percentage (zip)",
  "University Population (zip)", "Graduated Population (zip)",
  "Foreigner Percentage (mun.)", "Foreigner Percentage sqrt. (mun.)",
  "Foreigner Population (mun.)", "Population (mun.)",
  "University Percentage (mun.)",
  "University Population (mun.)", "Graduated Population (mun.)",
  "DID Residence", "DID Proportion", "Wave", "Aug. 2012 or After", "Panel")

### Female
balf_dy_unmatched <- findbalance(dy[dy$female==1,], fmbal, vnbal)
round(balf_dy_unmatched, 3)[, 1:7]
```

```
## mean.Tr mean.Co sdiff sdiff.pooled var.ratio T pval KS pval
```

## Gender	1.000	1.000	0.000	0.000	NaN	1.000	NaN
## Age	35.614	44.297	-68.600	-67.072	0.916	0.000	0.000
## Born Year	1976.308	1967.666	68.299	66.794	0.916	0.000	0.000
## Livng Length	31.581	41.358	-70.775	-69.669	0.940	0.000	0.000
## Living Proportion	0.870	0.924	-32.565	-36.994	1.819	0.000	0.000
## Foreigner Percentage (zip)	1.487	1.332	4.831	5.651	2.167	0.130	0.000
## Foreigner Percentage sqrt. (zip)	1.038	0.955	12.966	12.880	0.974	0.000	0.000
## Foreigner Population (zip)	127.311	92.861	12.784	14.230	1.629	0.000	0.000
## Population (zip)	6898.757	5559.082	18.482	19.867	1.368	0.000	0.000
## University Percentage (zip)	21.319	17.098	46.515	48.944	1.240	0.000	0.000
## University Population (zip)	1320.148	858.543	27.479	31.602	1.952	0.000	0.000
## Graduated Population (zip)	5489.361	4428.532	18.265	19.646	1.373	0.000	0.000
## Foreigner Percentage (mun.)	1.447	1.279	13.905	12.678	0.711	0.000	0.000
## Foreigner Percentage sqrt. (mun.)	1.124	1.028	22.181	21.143	0.832	0.000	0.000
## Foreigner Population (mun.)	3644.498	2976.128	15.311	15.792	1.136	0.000	0.000
## Population (mun.)	238367.915	217214.812	11.825	12.246	1.156	0.001	0.000
## University Percentage (mun.)	19.910	16.898	44.492	45.851	1.132	0.000	0.000
## University Population (mun.)	40337.562	32009.598	22.331	24.394	1.479	0.000	0.000
## Graduated Population (mun.)	189577.702	172755.826	11.767	12.236	1.177	0.001	0.000
## DID Residence	0.738	0.677	13.967	13.530	0.884	0.000	NaN
## DID Proportion	0.781	0.704	28.269	26.907	0.828	0.000	0.000
## Wave	11.649	12.030	-6.024	-6.133	1.076	0.093	0.030
## Aug. 2012 or After	0.545	0.576	-6.286	-6.310	1.016	0.083	NaN
## Panel	0.047	0.052	-2.203	-2.153	0.914	0.552	NaN

Male

```

balm_dy_unmatched <- findbalance(dy[dy$male==1,], fmbal, vnbal)
round(balm_dy_unmatched,3)[,1:7]

```

##	mean.Tr	mean.Co	sdiff	sdiff.pooled	var.ratio	T	pval	KS	pval
## Gender	0.000	0.000	0.000	0.000	NaN	1.000	NaN		
## Age	44.016	45.479	-10.370	-10.580	1.085	0.000	0.000		
## Born Year	1967.901	1966.461	10.195	10.396	1.083	0.001	0.000		
## Livng Length	40.305	42.622	-15.328	-15.653	1.089	0.000	0.000		
## Living Proportion	0.902	0.928	-19.177	-20.225	1.253	0.000	0.000		
## Foreigner Percentage (zip)	1.357	1.126	12.843	14.015	1.472	0.000	0.000		
## Foreigner Percentage sqrt. (zip)	1.006	0.898	18.348	18.704	1.082	0.000	0.000		
## Foreigner Population (zip)	106.053	71.883	15.496	17.307	1.657	0.000	0.000		
## Population (zip)	6393.051	4935.819	21.400	23.130	1.405	0.000	0.000		
## University Percentage (zip)	19.695	15.798	45.201	47.934	1.285	0.000	0.000		
## University Population (zip)	1161.691	724.864	28.244	32.963	2.135	0.000	0.000		
## Graduated Population (zip)	5101.522	3942.693	21.242	22.959	1.405	0.000	0.000		
## Foreigner Percentage (mun.)	1.405	1.174	17.590	19.557	1.619	0.000	0.000		
## Foreigner Percentage sqrt. (mun.)	1.103	1.002	23.025	23.656	1.118	0.000	0.000		
## Foreigner Population (mun.)	3482.812	2456.075	24.170	27.236	1.739	0.000	0.000		
## Population (mun.)	234059.670	190426.378	23.929	25.812	1.391	0.000	0.000		
## University Percentage (mun.)	18.965	16.032	43.664	45.625	1.202	0.000	0.000		
## University Population (mun.)	38297.775	27569.190	29.116	32.893	1.763	0.000	0.000		
## Graduated Population (mun.)	186083.053	151516.953	23.831	25.785	1.412	0.000	0.000		
## DID Residence	0.713	0.591	26.963	25.820	0.847	0.000	NaN		
## DID Proportion	0.747	0.635	38.766	36.383	0.787	0.000	0.000		
## Wave	11.670	11.755	-1.389	-1.404	1.045	0.639	0.150		
## Aug. 2012 or After	0.557	0.562	-0.937	-0.938	1.002	0.755	NaN		
## Panel	0.050	0.052	-1.169	-1.155	0.954	0.701	NaN		

```

## Matched Sample Proportions
matchprdt <- data.frame(
  labs = c("Unmatched",
            "Matched without Distance Adjustment",
            "Matched with Lambda = 350km",
            "Matched with Lambda = 200km",
            "Matched with Lambda = 100km",
            "Matched with Lambda = 50km"),
  notF = c(table(dy[dy$female==1,]$edu2)[1],
            table(dym1[dym1$female==1,]$edu2)[1],
            table(dym5[dym5$female==1,]$edu2)[1],
            table(dym4[dym4$female==1,]$edu2)[1],
            table(dym3[dym3$female==1,]$edu2)[1],
            table(dym2[dym2$female==1,]$edu2)[1]),
  treatedF = c(table(dy[dy$female==1,]$edu2)[2],
               table(dym1[dym1$female==1,]$edu2)[2],
               table(dym5[dym5$female==1,]$edu2)[2],
               table(dym4[dym4$female==1,]$edu2)[2],
               table(dym3[dym3$female==1,]$edu2)[2],
               table(dym2[dym2$female==1,]$edu2)[2]),
  notM = c(table(dy[dy$female==0,]$edu2)[1],
            table(dym1[dym1$female==0,]$edu2)[1],
            table(dym5[dym5$female==0,]$edu2)[1],
            table(dym4[dym4$female==0,]$edu2)[1],
            table(dym3[dym3$female==0,]$edu2)[1],
            table(dym2[dym2$female==0,]$edu2)[1]),
  treatedM = c(table(dy[dy$female==0,]$edu2)[2],
               table(dym1[dym1$female==0,]$edu2)[2],
               table(dym5[dym5$female==0,]$edu2)[2],
               table(dym4[dym4$female==0,]$edu2)[2],
               table(dym3[dym3$female==0,]$edu2)[2],
               table(dym2[dym2$female==0,]$edu2)[2]))
matchprdt$prF <- round((matchprdt$notF/table(dy[dy$female==1,]$edu2)[2])*100,1)
matchprdt$prM <- round((matchprdt$notM/table(dy[dy$female==0,]$edu2)[1])*100,1)
matchprdt <- matchprdt[,c("labs", "notF", "treatedF", "prF", "notM", "treatedM", "prM")]

matchprdt <- as.matrix(matchprdt)
colnames(matchprdt) <- c("", "No Univ.", "Univ.", "% Matched",
                        "No Univ.", "Univ.", "% Matched")

### Table of Data Sizes
require(knitr)
require(kableExtra)
tmp <- add_header_above(kable(matchprdt, "latex", booktabs = TRUE, linesep = ""),
                        c(" ", "Female"=3, "Male"=3))
cat(tmp)

##
## \begin{tabular}{l1l1l1l1l1}
## \toprule
## \multicolumn{1}{c}{ } & \multicolumn{3}{c}{Female} & \multicolumn{3}{c}{Male} \\
## \cmidrule{1{3pt}r{3pt}}{2-4} \cmidrule{1{3pt}r{3pt}}{5-7}
## & No Univ. & Univ. & \% Matched & No Univ. & Univ. & \% Matched \\
## \midrule

```

```
## Unmatched & 1778 & 1317 & 135.0 & 1778 & 2954 & 100.0\\
## Matched without Distance Adjustment & 856 & 856 & 65.0 & 1451 & 1451 & 81.6\\
## Matched with Lambda = 350km & 785 & 785 & 59.6 & 1355 & 1355 & 76.2\\
## Matched with Lambda = 200km & 692 & 692 & 52.5 & 1201 & 1201 & 67.5\\
## Matched with Lambda = 100km & 530 & 530 & 40.2 & 934 & 934 & 52.5\\
## Matched with Lambda = 50km & 406 & 406 & 30.8 & 655 & 655 & 36.8\\
## \bottomrule
## \end{tabular}
```

```
writeLines(tmp, paste0(projdir, "/out/matchedsizes.tex"))
```

```
### Balance Data
```

```
baldt <- as.data.frame(rbind(balf_dy_unmatched,
  bal_dy_matched_1$f,
  bal_dy_matched_5$f,
  bal_dy_matched_4$f,
  bal_dy_matched_3$f,
  bal_dy_matched_2$f,
  balm_dy_unmatched,
  bal_dy_matched_1$m,
  bal_dy_matched_5$m,
  bal_dy_matched_4$m,
  bal_dy_matched_3$m,
  bal_dy_matched_2$m))
baldt <- data.frame(stat = c(baldt$sdiff, baldt$`T pval`))
baldt$data <- rep(c("Unmatched",
  "Matched without Distance Adjustment",
  "Matched with Lambda = 350km",
  "Matched with Lambda = 200km",
  "Matched with Lambda = 100km",
  "Matched with Lambda = 50km"), each=nrow(balf_dy_unmatched))
baldt$data <- factor(baldt$data, levels=unique(baldt$data))
baldt$vn <- factor(rownames(balf_dy_unmatched),
  levels=rev(rownames(balf_dy_unmatched)))
baldt$stat_cat <- rep(c("Standardized Difference in Means",
  "p-value: Difference of Means Test"),
  each = nrow(balf_dy_unmatched)*12)
baldt$stat_cat <- factor(baldt$stat_cat, levels=unique(baldt$stat_cat))
baldt$gender <- rep(c("Female", "Male"), each=nrow(balf_dy_unmatched)*6)
baldt$gender <- factor(baldt$gender, levels=unique(baldt$gender))
```

```
require(ggplot2)
```

```
p <- ggplot(baldt, aes(x=vn, y=stat)) +
  geom_hline(aes(yintercept=0), size=0.25, linetype=1) +
  geom_point(aes(alpha=gender, shape=data), color="black",
    position=position_dodge(width=-0.5), size=2) +
  facet_grid( ~ stat_cat, scales="free_x", switch="x") +
  scale_shape_discrete(name="Matching\nStatus",
    labels = c("Unmatched",
      "Matched without Distance Adjustment",
      bquote("Matched with"~lambda~"= 350km"),
      bquote("Matched with"~lambda~"= 200km"),
      bquote("Matched with"~lambda~"= 100km"),
```

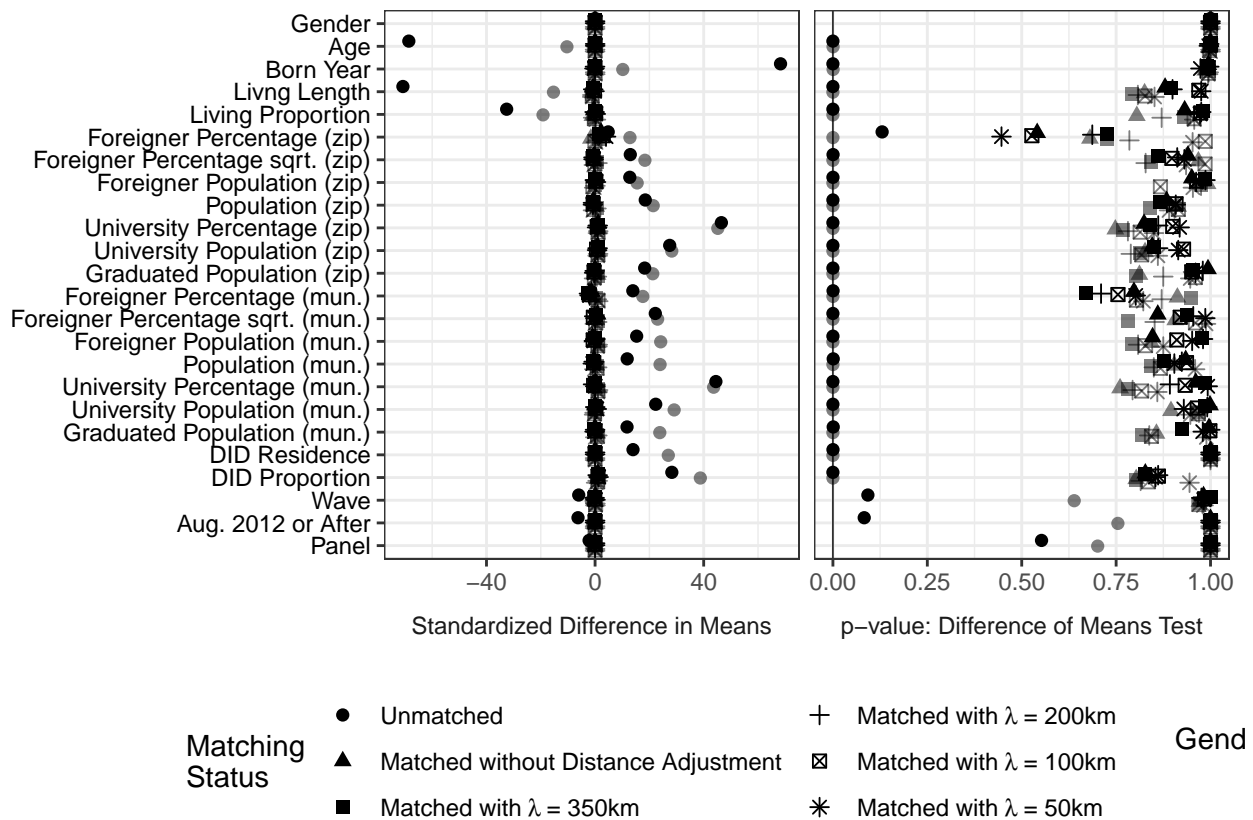
```

      bquote("Matched with ~lambda~" = 50km")) +
scale_alpha_manual(name="Gender", values=c("Female"=1,"Male"=0.5)) +
coord_flip() +
ylab(NULL) + xlab(NULL) +
guides(alpha = guide_legend(nrow = 2),
       shape = guide_legend(nrow = 3)) +
theme_bw() +
theme(legend.position="bottom",
      axis.text.y = element_text(color="black"),
      strip.background.x = element_blank(),
      strip.text.y = element_text(angle=0,size=10),
      strip.placement = "outside")
p

```

```
## Warning: position_dodge requires non-overlapping x intervals
```

```
## Warning: position_dodge requires non-overlapping x intervals
```



```
ggsave(paste0(projdir,"/out/matchbalanceplot_sifcct_v5.pdf"),p,width=8,height=5)
```

Plotting Geographic Distance Balance

Prepare Japanese Map Data

```

## Referenced from https://uribo.hatenablog.com/entry/2017/12/08/144549
## All w/o Tokyo and Okinawa
alljp_no1347 <- do.call("rbind", pblapply(seq(1,47)[-c(13,47)],

```

```

                                function(k) jpn_pref(pref_code = k, district=FALSE))

## Tokyo
tokyo13 <- jpn_pref(pref_code = 13, district = FALSE) %>%
  st_simplify(dTolerance = 0.01)

## Warning in st_simplify.sfc(st_geometry(x), preserveTopology, dTolerance): st_simplify does not corre
## longitude/latitude data, dTolerance needs to be in decimal degrees

## (Excluding Southern Islands) # Deprecated
# tokyo13 <- jpn_pref(pref_code = 13, district = TRUE) %>%
#   st_simplify(dTolerance = 0.01) %>%
#   mutate(city_code = as.numeric(city_code)) %>%
#   filter(city_code != 13421) %>% st_union() %>%
#   as.data.frame() %>% mutate(jis_code = "13",
#                               prefecture = "東京都") %>% magrittr::set_names(c("geometry",
#                                           "jis_code", "prefecture"))
## Okinawa
okinawa47 <- jpn_pref(pref_code = 47, district = FALSE)
okinawa47 <- okinawa47 %>% st_set_crs(value = 4326)

```

Prepare Respondents Data

```

# Sample N Respondents
N = 200
set.seed(3451)
dymap1 <- dym1[which(dym1$pair_id%in%sample(dym1$pair_id,N)),]
set.seed(5412)
dymap2 <- dym2[which(dym2$pair_id%in%sample(dym2$pair_id,N)),]
set.seed(5241)
dymap3 <- dym3[which(dym3$pair_id%in%sample(dym3$pair_id,N)),]
set.seed(5441)
dymap4 <- dym4[which(dym4$pair_id%in%sample(dym4$pair_id,N)),]
set.seed(5141)
dymap5 <- dym5[which(dym5$pair_id%in%sample(dym5$pair_id,N)),]

# Move Okinawa location to left-upper corner (not done for now)
# okinawa47$geometry <- okinawa47$geometry %>% magrittr::add(c(5.6, 17.5))
# dymap1$zip_lon[which(dymap1$zip_pref=="沖縄県")] <- dymap1$zip_lon[which(dymap1$zip_pref=="沖縄県")]
# dymap1$zip_lat[which(dymap1$zip_pref=="沖縄県")] <- dymap1$zip_lat[which(dymap1$zip_pref=="沖縄県")]
# dymap2$zip_lon[which(dymap2$zip_pref=="沖縄県")] <- dymap2$zip_lon[which(dymap2$zip_pref=="沖縄県")]
# dymap2$zip_lat[which(dymap2$zip_pref=="沖縄県")] <- dymap2$zip_lat[which(dymap2$zip_pref=="沖縄県")]
# dmmap1$zip_lon[which(dmmap1$zip_pref=="沖縄県")] <- dmmap1$zip_lon[which(dmmap1$zip_pref=="沖縄県")]
# dmmap1$zip_lat[which(dmmap1$zip_pref=="沖縄県")] <- dmmap1$zip_lat[which(dmmap1$zip_pref=="沖縄県")]
# dmmap2$zip_lon[which(dmmap2$zip_pref=="沖縄県")] <- dmmap2$zip_lon[which(dmmap2$zip_pref=="沖縄県")]
# dmmap2$zip_lat[which(dmmap2$zip_pref=="沖縄県")] <- dmmap2$zip_lat[which(dmmap2$zip_pref=="沖縄県")]
# demap1$zip_lon[which(demap1$zip_pref=="沖縄県")] <- demap1$zip_lon[which(demap1$zip_pref=="沖縄県")]
# demap1$zip_lat[which(demap1$zip_pref=="沖縄県")] <- demap1$zip_lat[which(demap1$zip_pref=="沖縄県")]
# demap2$zip_lon[which(demap2$zip_pref=="沖縄県")] <- demap2$zip_lon[which(demap2$zip_pref=="沖縄県")]
# demap2$zip_lat[which(demap2$zip_pref=="沖縄県")] <- demap2$zip_lat[which(demap2$zip_pref=="沖縄県")]

```

Plot


```

p1 <- ggplot() +
  geom_sf(data=alljp_no1347 %>% st_simplify(dTolerance = 0.01), size=0.3) +
  geom_sf(data = tokyo13, inherit.aes = TRUE, size=0.3) +
  geom_sf(data = okinawa47 %>% st_simplify(dTolerance = 0.01), inherit.aes = TRUE, size=0.3) +
  # geom_segment(aes(x = round(st_bbox(alljp_no1347)[1], 0), xend = 132.5, y = 40, yend = 40)) +
  # geom_segment(aes(x = 132.5, xend = 138, y = 40, yend = 42)) +
  # geom_segment(aes(x = 138, xend = 138, y = 42, yend = round(st_bbox(alljp_no1347)[4],0))) +
  geom_point(data = dymap1, aes(x=zip_lon,y=zip_lat, color=as.factor(1-female)), alpha=0.5, size=0.4) +
  geom_path(data = dymap1, aes(x=zip_lon,y=zip_lat, group=pair_id, color=as.factor(1-female)), alpha=0.5) +
  scale_color_manual(name="Gender", values=c("darkred","navyblue")) +
  coord_sf(xlim=c(124.5,148.5)) +
  #coord_sf(xlim=c(128,148.5),ylim=c(27,46)) +
  labs(x=paste0(table(dym1[dym1$female==1,]$treated)[1],"/",table(dy[dy$female==1,]$edu2)[2]," Female and",
    table(dym1[dym1$female==0,]$treated)[1],"/",table(dy[dy$female==0,]$edu2)[1],
    " Male Matched Pairs Found"),
    y=NULL,title="No Distance Adjustment") + theme_light() +
  theme(plot.title=element_text(hjust=0.5),
    panel.background = element_rect(color="black",fill="white"),
    axis.ticks = element_blank(),
    axis.text = element_blank(),
    line = element_blank(),
    axis.title.x = element_text(size=10),
    legend.position = "none")

```

```

## Warning in st_simplify.sfc(st_geometry(x), preserveTopology, dTolerance): st_simplify does not correct
## longitude/latitude data, dTolerance needs to be in decimal degrees

```

```

## Warning in st_simplify.sfc(st_geometry(x), preserveTopology, dTolerance): st_simplify does not correct
## longitude/latitude data, dTolerance needs to be in decimal degrees

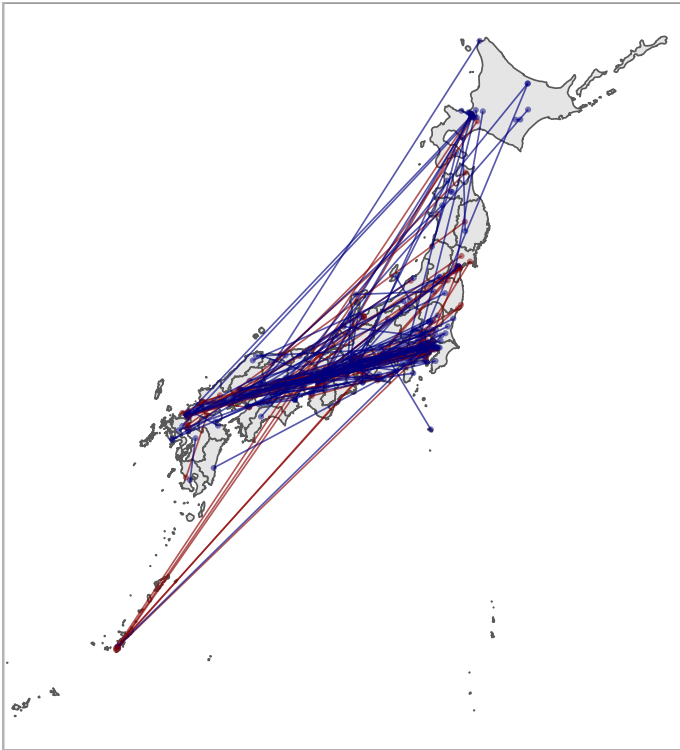
```

```

p1

```


No Distance Adjustment



856/1317 Female and 1451/1778 Male Matched Pairs Found

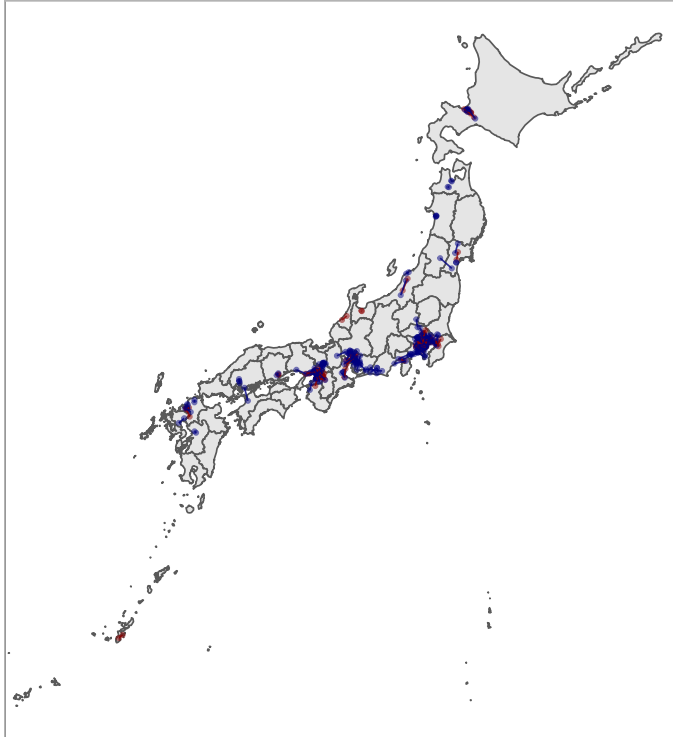
```
p2 <- ggplot() +
  geom_sf(data=alljp_no1347 %>% st_simplify(dTolerance = 0.01), size=0.3) +
  geom_sf(data = tokyo13, inherit.aes = TRUE, size=0.3) +
  geom_sf(data = okinawa47 %>% st_simplify(dTolerance = 0.01), inherit.aes = TRUE, size=0.3) +
  # geom_segment(aes(x = round(st_bbox(alljp_no1347)[1], 0), xend = 132.5, y = 40, yend = 40)) +
  # geom_segment(aes(x = 132.5, xend = 138, y = 40, yend = 42)) +
  # geom_segment(aes(x = 138, xend = 138, y = 42, yend = round(st_bbox(alljp_no1347)[4], 0))) +
  geom_point(data = dymap2, aes(x=zip_lon, y=zip_lat, color=as.factor(1-female)), alpha=0.5, size=0.4) +
  geom_path(data = dymap2, aes(x=zip_lon, y=zip_lat, group=pair_id, color=as.factor(1-female)), alpha=0.5) +
  scale_color_manual(name="Gender", values=c("darkred", "navyblue")) +
  coord_sf(xlim=c(124.5, 148.5)) +
  # coord_sf(xlim=c(128, 148.5), ylim=c(27, 46)) +
  labs(x=paste0(table(dym2[dym2$female==1,]$treated)[1], "/", table(dy[dy$female==1,]$edu2)[2], " Female and ",
    table(dym2[dym2$female==0,]$treated)[1], "/", table(dy[dy$female==0,]$edu2)[1],
    " Male Matched Pairs Found"),
    y=NULL, title=bquote("Distance Adjusted (~lambda~" = 50km))) + theme_light() +
  theme(plot.title=element_text(hjust=0.5),
    panel.background = element_rect(color="black", fill="white"),
    axis.ticks = element_blank(),
    axis.text = element_blank(),
    line = element_blank(),
    axis.title.x = element_text(size=10),
    legend.position = "none")
```

```
## Warning in st_simplify.sfc(st_geometry(x), preserveTopology, dTolerance): st_simplify does not correct
## longitude/latitude data, dTolerance needs to be in decimal degrees
```

```
## Warning in st_simplify.sfc(st_geometry(x), preserveTopology, dTolerance): st_simplify does not corre
## longitude/latitude data, dTolerance needs to be in decimal degrees
```

p2

Distance Adjusted ($\lambda = 50\text{km}$)



406/1317 Female and 655/1778 Male Matched Pairs Found

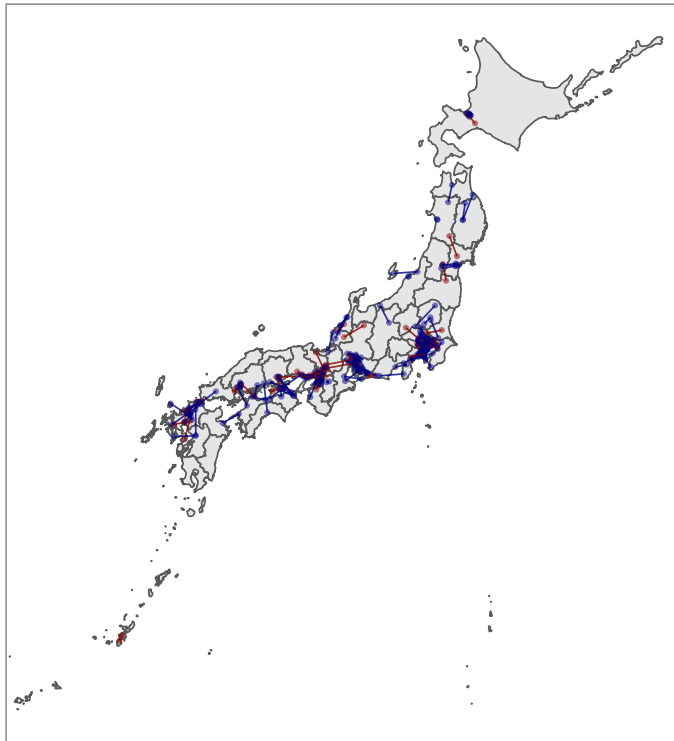
```
p3 <- ggplot() +
  geom_sf(data=alljp_no1347 %>% st_simplify(dTolerance = 0.01), size=0.3) +
  geom_sf(data = tokyo13, inherit.aes = TRUE, size=0.3) +
  geom_sf(data = okinawa47 %>% st_simplify(dTolerance = 0.01), inherit.aes = TRUE, size=0.3) +
  # geom_segment(aes(x = round(st_bbox(alljp_no1347)[1], 0), xend = 132.5, y = 40, yend = 40)) +
  # geom_segment(aes(x = 132.5, xend = 138, y = 40, yend = 42)) +
  # geom_segment(aes(x = 138, xend = 138, y = 42, yend = round(st_bbox(alljp_no1347)[4], 0))) +
  geom_point(data = dymap3, aes(x=zip_lon, y=zip_lat, color=as.factor(1-female)), alpha=0.5, size=0.4) +
  geom_path(data = dymap3, aes(x=zip_lon, y=zip_lat, group=pair_id, color=as.factor(1-female)), alpha=0.5) +
  scale_color_manual(name="Gender", values=c("darkred", "navyblue")) +
  coord_sf(xlim=c(124.5, 148.5)) +
  #coord_sf(xlim=c(128, 148.5), ylim=c(27, 46)) +
  labs(x=paste0(table(dym3[dym3$female==1,]$treated)[1], "/", table(dy[dy$female==1,]$edu2)[2], " Female and",
    table(dym3[dym3$female==0,]$treated)[1], "/", table(dy[dy$female==0,]$edu2)[1],
    " Male Matched Pairs Found"),
    y=NULL, title=bquote("Distance Adjusted (~lambda~ = 100km))) + theme_light() +
  theme(plot.title=element_text(hjust=0.5),
    panel.background = element_rect(color="black", fill="white"),
    axis.ticks = element_blank(),
    axis.text = element_blank(),
    line = element_blank(),
    axis.title.x = element_text(size=10),
    legend.position = "none")
```

```
## Warning in st_simplify.sfc(st_geometry(x), preserveTopology, dTolerance): st_simplify does not corre
## longitude/latitude data, dTolerance needs to be in decimal degrees
```

```
## Warning in st_simplify.sfc(st_geometry(x), preserveTopology, dTolerance): st_simplify does not corre
## longitude/latitude data, dTolerance needs to be in decimal degrees
```

p3

Distance Adjusted ($\lambda = 100\text{km}$)



530/1317 Female and 934/1778 Male Matched Pairs Found

```
p4 <- ggplot() +
  geom_sf(data=alljp_no1347 %>% st_simplify(dTolerance = 0.01), size=0.3) +
  geom_sf(data = tokyo13, inherit.aes = TRUE, size=0.3) +
  geom_sf(data = okinawa47 %>% st_simplify(dTolerance = 0.01), inherit.aes = TRUE, size=0.3) +
  # geom_segment(aes(x = round(st_bbox(alljp_no1347)[1], 0), xend = 132.5, y = 40, yend = 40)) +
  # geom_segment(aes(x = 132.5, xend = 138, y = 40, yend = 42)) +
  # geom_segment(aes(x = 138, xend = 138, y = 42, yend = round(st_bbox(alljp_no1347)[4],0))) +
  geom_point(data = dymap4, aes(x=zip_lon,y=zip_lat, color=as.factor(1-female)), alpha=0.5, size=0.4) +
  geom_path(data = dymap4, aes(x=zip_lon,y=zip_lat, group=pair_id, color=as.factor(1-female)), alpha=0.5) +
  scale_color_manual(name="Gender", values=c("darkred","navyblue")) +
  coord_sf(xlim=c(124.5,148.5)) +
  #coord_sf(xlim=c(128,148.5),ylim=c(27,46)) +
  labs(x=paste0(table(dym4[dym4$female==1,$treated])[1],"/",table(dy[dy$female==1,$edu2])[2], " Female and
      table(dym4[dym4$female==0,$treated])[1],"/",table(dy[dy$female==0,$edu2])[1],
      " Male Matched Pairs Found"),
      y=NULL,title=bquote("Distance Adjusted (~lambda~ = 200km)")) + theme_light() +
  theme(plot.title=element_text(hjust=0.5),
        panel.background = element_rect(color="black",fill="white"),
```

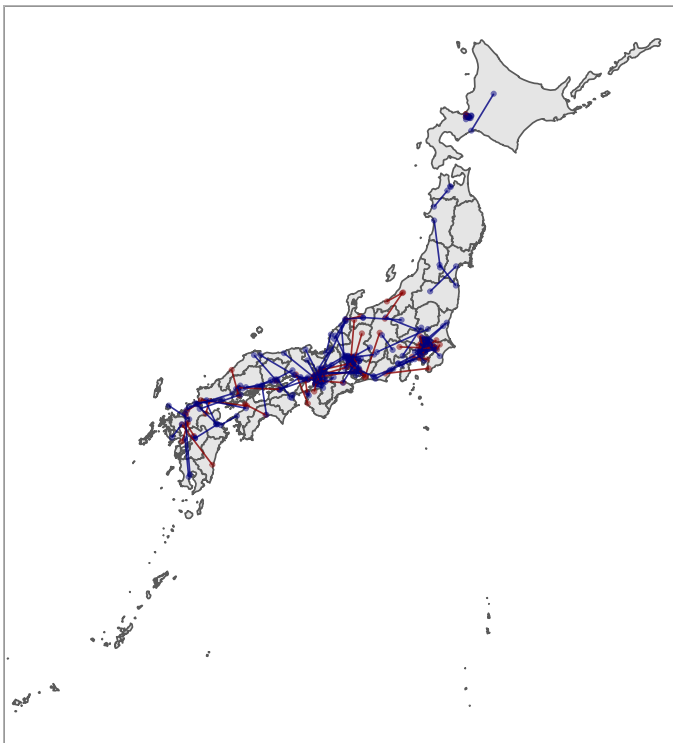
```
axis.ticks = element_blank(),
axis.text = element_blank(),
line = element_blank(),
axis.title.x = element_text(size=10),
legend.position = "none")
```

```
## Warning in st_simplify.sfc(st_geometry(x), preserveTopology, dTolerance): st_simplify does not correct
## longitude/latitude data, dTolerance needs to be in decimal degrees
```

```
## Warning in st_simplify.sfc(st_geometry(x), preserveTopology, dTolerance): st_simplify does not correct
## longitude/latitude data, dTolerance needs to be in decimal degrees
```

p4

Distance Adjusted ($\lambda = 200\text{km}$)



692/1317 Female and 1201/1778 Male Matched Pairs Found

```
p5 <- ggplot() +
  geom_sf(data=alljp_no1347 %>% st_simplify(dTolerance = 0.01), size=0.3) +
  geom_sf(data = tokyo13, inherit.aes = TRUE, size=0.3) +
  geom_sf(data = okinawa47 %>% st_simplify(dTolerance = 0.01), inherit.aes = TRUE, size=0.3) +
  # geom_segment(aes(x = round(st_bbox(alljp_no1347)[1], 0), xend = 132.5, y = 40, yend = 40)) +
  # geom_segment(aes(x = 132.5, xend = 138, y = 40, yend = 42)) +
  # geom_segment(aes(x = 138, xend = 138, y = 42, yend = round(st_bbox(alljp_no1347)[4], 0))) +
  geom_point(data = dymap5, aes(x=zip_lon, y=zip_lat, color=as.factor(1-female)), alpha=0.5, size=0.4) +
  geom_path(data = dymap5, aes(x=zip_lon, y=zip_lat, group=pair_id, color=as.factor(1-female)), alpha=0.5) +
  scale_color_manual(name="Gender", values=c("darkred", "navyblue")) +
  coord_sf(xlim=c(124.5, 148.5)) +
  #coord_sf(xlim=c(128, 148.5), ylim=c(27, 46)) +
  labs(x=paste0(table(dym5[dym5$female==1,]$treated)[1], "/", table(dy[dy$female==1,]$edu2)[2], " Female and Male Matched Pairs Found"))
```

```

        table(dym5[dym5$female==0,]$treated)[1],"/",table(dy[dy$female==0,]$edu2)[1],
        " Male Matched Pairs Found"),
    y=NULL,title=bquote("Distance Adjusted ( $\lambda$  = 350km)")) + theme_light() +
    theme(plot.title=element_text(hjust=0.5),
    panel.background = element_rect(color="black",fill="white"),
    axis.ticks = element_blank(),
    axis.text = element_blank(),
    line = element_blank(),
    axis.title.x = element_text(size=10),
    legend.position = "none")

```

```

## Warning in st_simplify.sfc(st_geometry(x), preserveTopology, dTolerance): st_simplify does not correct
## longitude/latitude data, dTolerance needs to be in decimal degrees

```

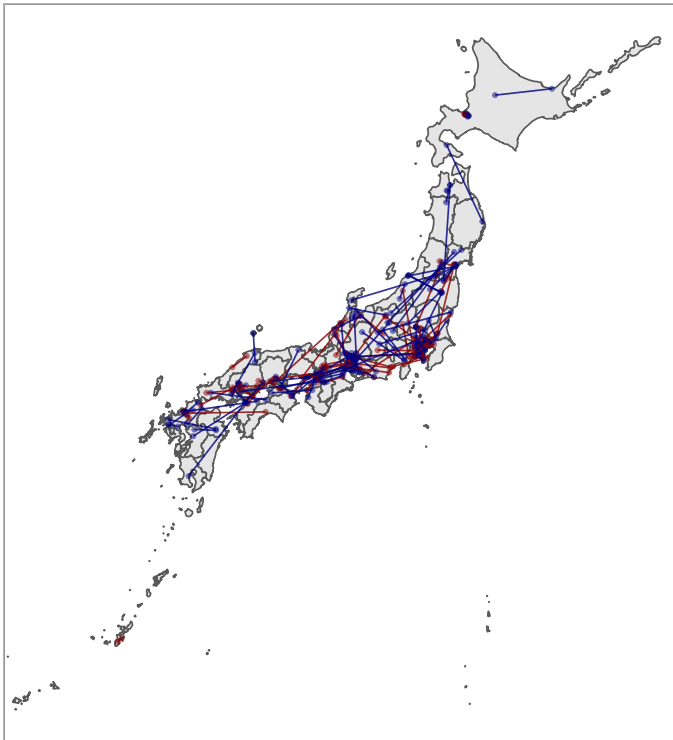
```

## Warning in st_simplify.sfc(st_geometry(x), preserveTopology, dTolerance): st_simplify does not correct
## longitude/latitude data, dTolerance needs to be in decimal degrees

```

p5

Distance Adjusted (λ = 350km)



785/1317 Female and 1355/1778 Male Matched Pairs Found

Export Map Plots

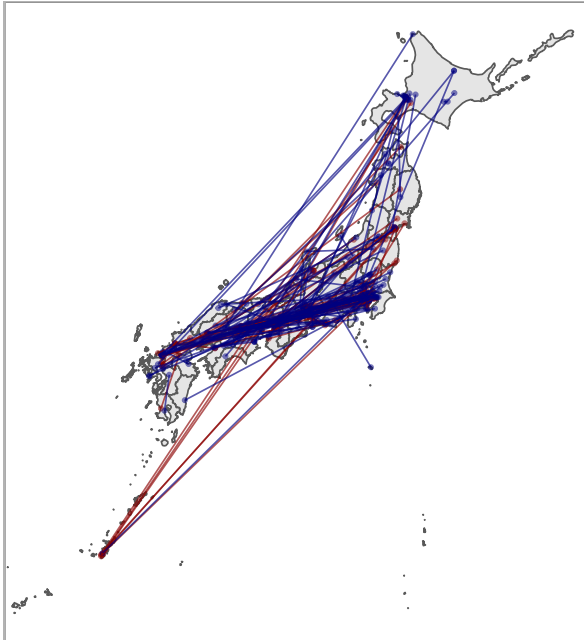
```

foottxt <- paste0("Dots represent randomly sampled ", N, " matched respondent pairs",
  " and lines connect two matched pairs ",
  "on the map (red = female, \nblue = male). The left panel shows the matching outcome
  \"without geographic distance adjustment and ",
  "the right panel shows \nthe outcome of matching with geographic distance adjustment.

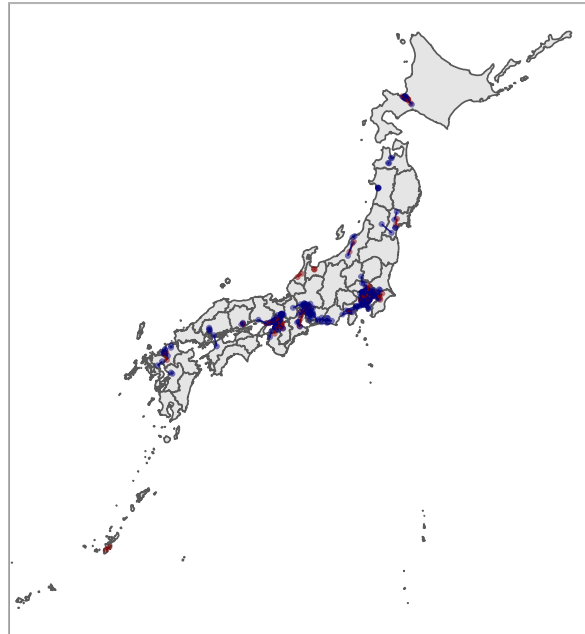
```

```
p <- arrangeGrob(p1,p2, nrow=1,
  bottom=textGrob(foottxt, vjust=0.5,just="left",x=unit(0.05,"npc"),
    gp=gpar(fontsize=9)))
grid.arrange(p)
```

No Distance Adjustment



Distance Adjusted ($\lambda = 50\text{km}$)



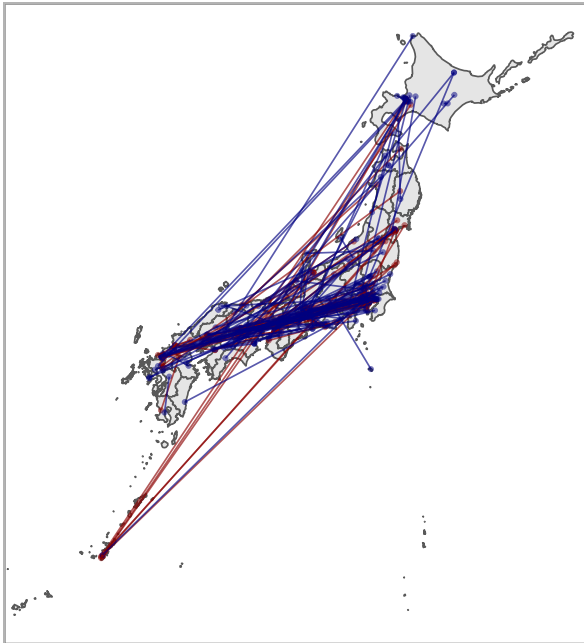
/1317 Female and 1451/1778 Male Matched Pairs 46/1317 Female and 655/1778 Male Matched Pairs For

Dots represent randomly sampled 200 matched respondent pairs and lines connect two matched pairs on the map (blue = male). The left panel shows the matching outcome without geographic distance adjustment and the right panel shows the outcome of matching with geographic distance adjustment.

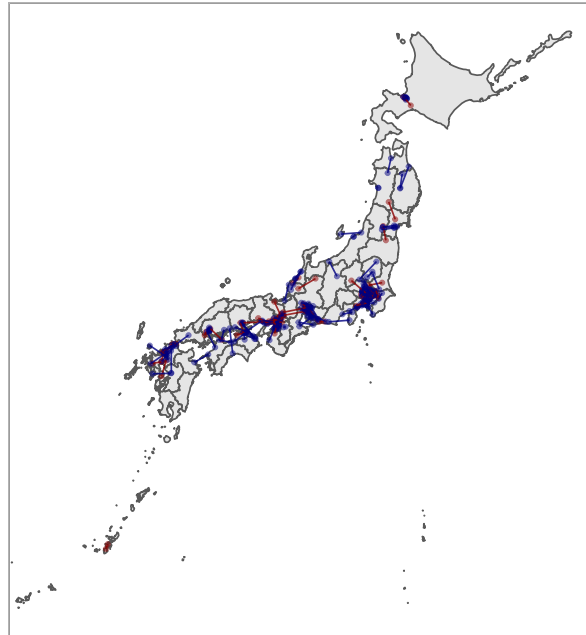
```
ggsave(paste0(projdir,"/out/geomatchplot_150_sifcct_v5.pdf"),p,width=8,height=5)
```

```
p <- arrangeGrob(p1,p3, nrow=1,
  bottom=textGrob(foottxt,
    vjust=0.5,just="left",x=unit(0.05,"npc"),
    gp=gpar(fontsize=9)))
grid.arrange(p)
```

No Distance Adjustment



Distance Adjusted ($\lambda = 100\text{km}$)



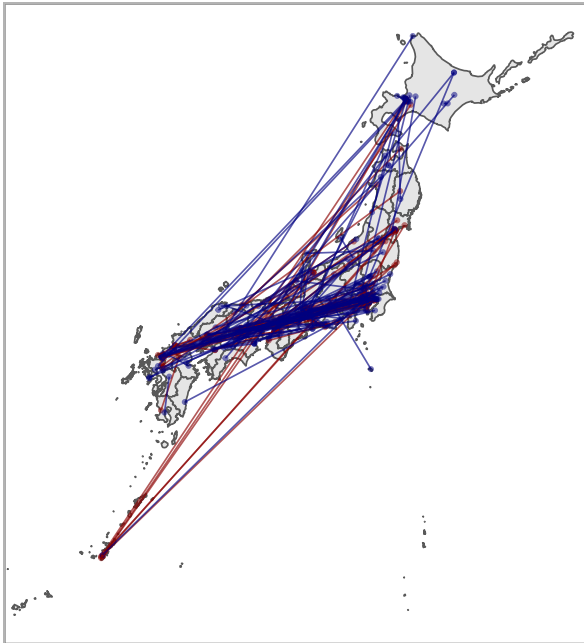
50/1317 Female and 1451/1778 Male Matched Pairs 50/1317 Female and 934/1778 Male Matched Pairs For

Dots represent randomly sampled 200 matched respondent pairs and lines connect two matched pairs on the map (red = female, blue = male). The left panel shows the matching outcome without geographic distance adjustment and the right panel shows the outcome of matching with geographic distance adjustment.

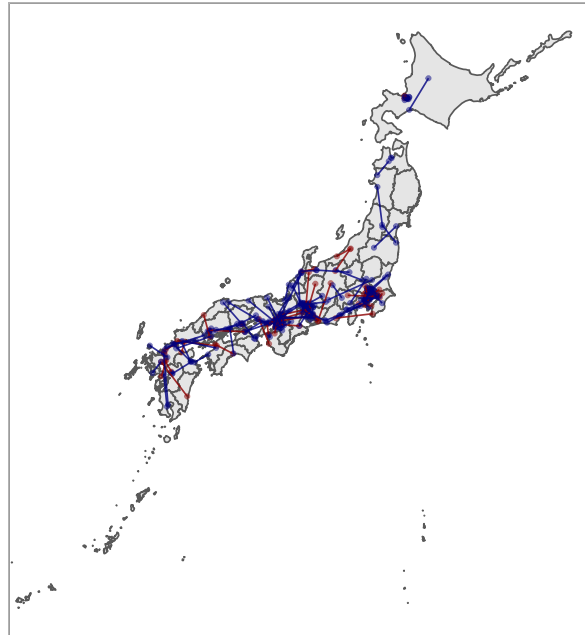
```
ggsave(paste0(projdir, "/out/geomatchplot_l100_sifcct_v5.pdf"), p, width=8, height=5)
```

```
p <- arrangeGrob(p1, p4, nrow=1,
  bottom=textGrob(foottxt, vjust=0.5, just="left", x=unit(0.05, "npc"),
    gp=gpar(fontsize=9)))
grid.arrange(p)
```


No Distance Adjustment



Distance Adjusted ($\lambda = 200\text{km}$)



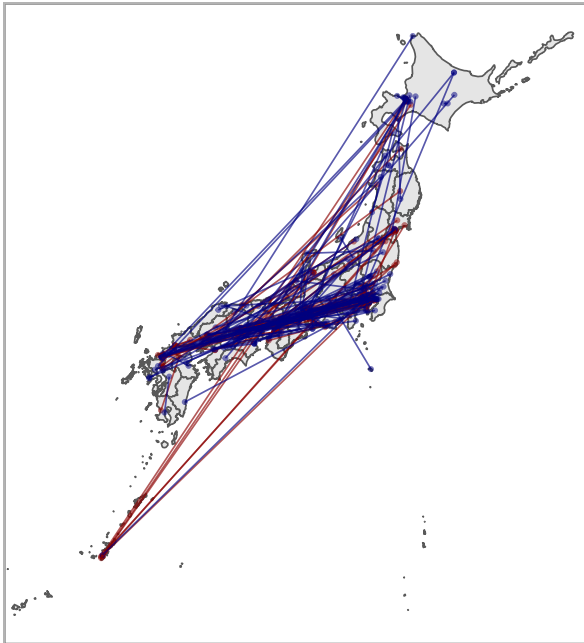
1317 Female and 1451/1778 Male Matched Pairs 1317 Female and 1201/1778 Male Matched Pairs

Dots represent randomly sampled 200 matched respondent pairs and lines connect two matched pairs on the map (blue = male). The left panel shows the matching outcome without geographic distance adjustment and the right panel shows the outcome of matching with geographic distance adjustment.

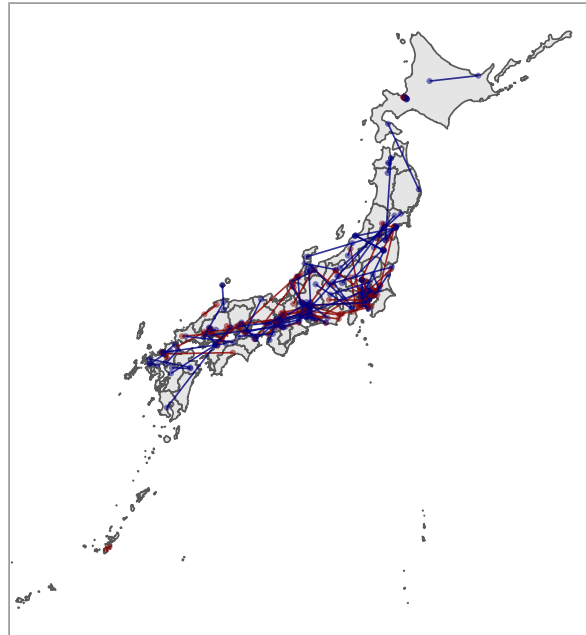
```
ggsave(paste0(projdir, "/out/geomatchplot_l200_sifcct_v5.pdf"), p, width=8, height=5)
```

```
p <- arrangeGrob(p1, p5, nrow=1,
  bottom=textGrob(foottxt, vjust=0.5, just="left", x=unit(0.05, "npc"),
    gp=gpar(fontsize=9)))
grid.arrange(p)
```

No Distance Adjustment



Distance Adjusted ($\lambda = 350\text{km}$)



1317 Female and 1451/1778 Male Matched Pairs 1317 Female and 1355/1778 Male Matched Pairs

Dots represent randomly sampled 200 matched respondent pairs and lines connect two matched pairs on the map (blue = male). The left panel shows the matching outcome without geographic distance adjustment and the right panel shows the outcome of matching with geographic distance adjustment.

```
ggsave(paste0(projdir, "/out/geomatchplot_l350_sifcct_v5.pdf"), p, width=8, height=5)
```