Co-financing model code documentation

Itamar Megiddo

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Contents

Setup

Loading packages.

```
library(tidyverse)
```

```
## -- Attaching packages -----------------------------------------------------------------------
```

```
## v ggplot2 3.2.0     v purrr   0.3.2
## v tibble  2.1.3     v dplyr   0.8.3
## v tidyr   0.8.3     v stringr 1.4.0
## v readr   1.3.1     v forcats 0.4.0
```

```
## -- Conflicts -------------------------------------------------------------------------------------
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()    masks stats::lag()
```

Load data.

```
gdp_data <- read_csv("data/data_gdp_per_capita.csv")
```

```
## Parsed with column specification:
## cols(
##   Country = col_character(),
##   `2001` = col_double(),
##   `2002` = col_double(),
##   `2003` = col_double(),
##   `2004` = col_number(),
##   `2005` = col_number(),
##   `2006` = col_number(),
##   `2007` = col_number(),
##   `2008` = col_number(),
##   `2009` = col_number(),
##   `2010` = col_number(),
##   `2011` = col_number(),
##   `2012` = col_number(),
##   `2013` = col_number(),
##   `2014` = col_number(),
##   `2015` = col_number(),
##   `2016` = col_number(),
##   `2017` = col_number(),
##   `2018` = col_number()
## )
```

```
cea_data <- read_csv("data/data_cea.csv")
```

```
## Warning: Missing column names filled in: 'X15' [15], 'X16' [16],
## 'X17' [17], 'X18' [18], 'X19' [19], 'X20' [20], 'X21' [21], 'X22' [22],
## 'X23' [23], 'X24' [24]
```

```
## Parsed with column specification:
## cols(
##   .default = col_logical(),
##   Country = col_character(),
##   Vaccine = col_character(),
##   `Indicator (e.g. deaths averted)` = col_character(),
##   `Total cost for the period` = col_number(),
##   `Costs per year` = col_number(),
##   `Total benefit for the period` = col_number(),
##   `Benefits per year` = col_number(),
##   `Cost-effectiveness` = col_number(),
##   `Start year` = col_double(),
##   `End year` = col_double(),
##   `Year of dollar rate used` = col_double(),
##   Perspective = col_character(),
##   Assumptions = col_character(),
##   `Author(s)` = col_character()
## )

## See spec(...) for full column specifications.
```

```
fund_data <- read_csv("data/data_vaccine_funding.csv")
```

```
## Warning: Missing column names filled in: 'X9' [9], 'X10' [10], 'X11' [11],
## 'X12' [12], 'X13' [13], 'X14' [14], 'X15' [15]

## Parsed with column specification:
## cols(
##   Country = col_character(),
##   Vaccine = col_character(),
##   Year = col_double(),
##   Total = col_double(),
##   `Domestic contribution ($)` = col_double(),
##   `Gavi contribution ($)` = col_double(),
##   `Domestic (%)` = col_double(),
##   `Gavi (%)` = col_double(),
##   X9 = col_logical(),
##   X10 = col_logical(),
##   X11 = col_logical(),
##   X12 = col_logical(),
##   X13 = col_logical(),
##   X14 = col_logical(),
##   X15 = col_logical()
## )
```

```
gdp_deflator <- read_csv("data/gdp_deflator.csv")
```

```
## Parsed with column specification:
## cols(
##   .default = col_double(),
##   `Country Name` = col_character(),
##   `Country Code` = col_character(),
```

```
##   `Indicator Name` = col_character(),
##   `Indicator Code` = col_character(),
##   `2019` = col_logical()
## )
## See spec(...) for full column specifications.
```

```
ex_rate <- read_csv("data/ex_rate.csv")
```

```
## Parsed with column specification:
## cols(
##   .default = col_double(),
##   `Country Name` = col_character(),
##   `Country Code` = col_character(),
##   `Indicator Name` = col_character(),
##   `Indicator Code` = col_character(),
##   `2019` = col_logical()
## )
## See spec(...) for full column specifications.
```

```
population_data <- read_csv("data/population_data.csv")
```

```
## Parsed with column specification:
## cols(
##   `Country Name` = col_character(),
##   `2012` = col_double(),
##   `2013` = col_double(),
##   `2014` = col_double(),
##   `2015` = col_double(),
##   `2016` = col_double(),
##   `2017` = col_double(),
##   `2018` = col_double()
## )
```

Set some global values.

```
countries_model <- c("Ghana", "India", "Kenya",
                     "Nigeria", "Angola",   "Senegal")
```

Capitlise first letter function (to be used later).

```
firstup <- function(x) {
  substr(x, 1, 1) <- toupper(substr(x, 1, 1))
  x
}
```

Round up nicely for plotting (to be used later).

```
roundUp <- function(x, nice_val=c(1, 2, 3, 4, 5, 6, 7,  8, 9, 10)) {
    if(length(x) != 1) stop("'x' can only take vectors of length 1")
    10^floor(log10(x)) * nice_val[[which(x <= 10^floor(log10(x)) * nice_val)[[1]]]]
}
```

Clean data

Population data

```r
# Change to long format data
population_data <-
  population_data %>%
  rename(country = `Country Name`) %>%
  gather(key = year, value = population, `2012`:`2018`) %>%
  mutate(year = as.numeric(year)) %>%
  filter(country %in% countries_model)
```

Gross domestic product deflator and exchange rate

Clean and combine Gross domestic product (GDP) delfator and exchange rate data.

```r
# Clean and remove unnecessary columns
gdp_deflator <- gdp_deflator %>% rename(country = `Country Name`) %>%
  select(country, `2000`:`2018`) %>%
  filter(country %in% countries_model) %>%
  gather(key = year, value = deflator, `2000`:`2018`)

# Add 2018 year to get multiplier of def_2018/def_year
gdp_deflator <- left_join(gdp_deflator,
          gdp_deflator %>% filter(year == 2018) %>% select(-year) %>%
            rename(deflator2018 = deflator),
          by = "country") %>%
  mutate(def_multiplier = deflator2018/deflator) %>%
  select(country, year, def_multiplier)

# Clean exchange rate tibble
ex_rate <- ex_rate %>% rename(country = `Country Name`) %>%
  select(country, `2000`:`2018`) %>%
  filter(country %in% countries_model) %>%
  gather(key = year, value = ex, `2000`:`2018`)

# Add exchange rate multiplier to get ex_2018/ex_year
#   multiplying by both multipliers will give
#   to local_currency = reported$$ * ex
#   to 2018_local-currency = local_currency * def_2018/def
#   to 2018_$ = 2018_local_currency / ex_2018
#   i.e. (reported$$ * ex * def_2018/def) / ex_2018 =
#   reported$$ * def_2018/def * ex/ex_2018
ex_rate <- left_join(ex_rate,
          ex_rate %>% filter(year == 2018) %>% select(-year) %>%
            rename(ex2018 = ex),
          by = "country") %>%
  mutate(ex_multiplier = ex/ex2018) %>%
  select(country, year, ex_multiplier)

# Create currentcy exchange deflating and exchanging
```

```r
currency_conv <- left_join(gdp_deflator, ex_rate, by = c("country", "year")) %>%
  mutate(currency_mult = def_multiplier * ex_multiplier) %>%
  mutate(dollar_year = as.numeric(year)) %>%
  select(country, dollar_year, currency_mult)
```

Cost-effectiveness data and GDP per capita

Clean the cost-effectivness analysis (CEA) data.

```r
cea_data <- cea_data %>%
  # Rename to more data friendly
  rename(country = Country, vaccine = Vaccine,
         indicator = `Indicator (e.g. deaths averted)`,
         cost_for_horizon = `Total cost for the period`,
         cost = `Costs per year`,
         benefit_for_horizon = `Total benefit for the period`,
         benefit = `Benefits per year`, cea = `Cost-effectiveness`,
         s_year = `Start year`, e_year = `End year`,
         dollar_year = `Year of dollar rate used`,
         perspective = Perspective, assumptions = Assumptions,
         authors = `Author(s)`) %>%
  select(country:authors) %>%
  # Consistent nameing of indicators
  mutate(indicator = firstup(indicator)) %>%
  mutate(indicator = gsub("DALY ", "DALYs ", indicator)) %>%
  mutate(indicator = gsub("Death ", "Deaths ", indicator)) %>%
  mutate(indicator = gsub("Case ", "Cases ", indicator)) %>%
  mutate(indicator = gsub("YLL", "YLLs averted", indicator)) %>%
  # Some of the vaccine names differ between fund_data and cea_data
  mutate(vaccine = ifelse(grepl("DTP-Hep", vaccine, ignore.case = TRUE),
                          "DTP-hep B",
                          vaccine)) %>%
  mutate(vaccine = ifelse(grepl("Yellow Fever", vaccine),
                          "Yellow Fever",
                          vaccine)) %>%
  mutate(vaccine = ifelse(grepl("Measles", vaccine),
                          "Measles",
                          vaccine)) %>%
  mutate(vaccine = ifelse(grepl("Pneumococcal", vaccine),
                          "PCV",
                          vaccine)) %>%
  mutate(vaccine = ifelse(grepl("Rota", vaccine),
                          "Rotavirus",
                          vaccine)) %>%
  mutate(vaccine = ifelse(grepl("BCG", vaccine),
                          "BCG",
                          vaccine)) %>%
  # Select only countries in the analysis
  filter(country %in% countries_model)
```

Perform some checks on CEA data.

```r
# Check diff between calculated and CEA reported in papers
#   Small differences are likely due to rounding errors
cea_data %>% select(cost, benefit, cea) %>%
  mutate(cea2 = round(cost/benefit,2)) %>%
  mutate(diff = cea - cea2) %>%
  arrange(desc(abs(diff))) %>%
  filter(abs(diff) > 1e-09)
```

```
## # A tibble: 19 x 5
##         cost benefit    cea   cea2     diff
##        <dbl>   <dbl>  <dbl>  <dbl>    <dbl>
##  1   3689149     303 12196. 12175. 20.1
##  2   1409658     303  4660.  4652.  7.69
##  3   6409006     634 10102. 10109. -6.37
##  4  11783120     888 13274. 13269.  4.48
##  5  18747118    1303 14392. 14388.  4.01
##  6   3721375     634  5866.  5870. -3.7
##  7   1505569     311  4844.  4841.  3.11
##  8   7390364     888  8325.  8322.  2.81
##  9   1083447     311  3486.  3484.  2.24
## 10    379910     311  1222.  1222.  0.780
## 11 176198173   10601 16621. 16621.  0.470
## 12 176198173   10601 16621. 16621.  0.470
## 13  34223066    9193  3723.  3723. -0.160
## 14   7766946    3053  2544.  2544.  0.160
## 15   7766946    3053  2544.  2544.  0.160
## 16  28770592    9193  3129.  3130. -0.140
## 17  18747118    9228  2031.  2032. -0.1000
## 18   3200000   12570   255.   255.  0.01
## 19   3721375   33380   111.   111. -0.01000
```

Transform GDP per capita data to long format, convert to $2018, and add cost & benefit columns.

```r
# Max cost +, to create a sequence for plotting
max_x <- max(cea_data$cost) + 50000000
seq_x <- seq(0, max_x, by = 100000)

# Convert GDP data to 2018 $
gdp_data <- gdp_data %>%
  gather(key = "year", value = "gdp_pc", -Country) %>%
  rename(country = Country) %>%
  mutate(year = as.numeric(year)) %>%
  # Convert currency
  left_join(currency_conv %>% rename(year = dollar_year),
            by = c("country", "year")) %>%
  mutate(gdp_pc = gdp_pc * currency_mult) %>%
  select(-currency_mult)

# Add cost and benefit columns to gdp_data to
#   draw the CEA threshold
gdp_data <- gdp_data %>%
  # Create rows for each analysis
  slice(rep(row_number(), length(seq_x))) %>%
```

```r
  # Add a cost column, for plotting cost x-axis
  mutate(cost = unlist(sapply(seq_x, function(x) rep(x, nrow(gdp_data)), simplify = FALSE))) %>%
  # Add a number of benefit lines for plotting
  mutate(benefit_line = cost/gdp_pc,
         benefit_half_line = cost/gdp_pc/0.5,
         benefit_3_line = cost/gdp_pc/3,
         benefit_4_5_line = cost/gdp_pc/4.5)

# Convert to long data tibble
gdp_data <- gdp_data %>%
  gather(key = threshold, value = cea_line, benefit_line:benefit_4_5_line) %>%
  # Write clear headings for different thresholds
  mutate(threshold = ifelse(threshold == "benefit_line",
                            "1 x GDP\nper capita",
                            threshold)) %>%
  mutate(threshold = ifelse(threshold == "benefit_half_line",
                            "0.5 x GDP\nper capita",
                            threshold)) %>%
  mutate(threshold = ifelse(threshold == "benefit_3_line",
                            "3 x GDP\nper capita",
                            threshold)) %>%
  mutate(threshold = ifelse(threshold == "benefit_4_5_line",
                            "4.5 x GDP\nper capita",
                            threshold))
```

Convert the CEA data into 2018 US Dollars using WB deflator.

```r
# Combine cea data and currency conversion data
cea_data <- left_join(cea_data, currency_conv, by = c("country", "dollar_year")) %>%
  # Convert all costs
  mutate(cost_for_horizon = cost_for_horizon * currency_mult,
         cost = cost * currency_mult,
         cea = cea * currency_mult,
         benefit = ifelse(
           grepl("Total healthcare cost averted|Out-of-pocket expenditure",
                 indicator),
           benefit * currency_mult,
           benefit))
```

Co-financing data

Clean co-financing data.

```r
fund_data <-
  fund_data %>%
  # Rename columns so they are easier to work with.
  rename(country = Country, vaccine = Vaccine, year = Year,
         total = Total,
         domestic = `Domestic contribution ($)`,
         gavi = `Gavi contribution ($)`,
         domestic_p = `Domestic (%)`, gavi_p = `Gavi (%)`) %>%
  # Remove some columns that are unnecesary
```

```r
  select(country:gavi_p) %>%
  # Remove rows with countries that are not part of the analysis (there are some
  #   unnecessary rows in the data that explain the data)
  filter(country %in% countries_model) %>%
  # Convert numbers to numeric
  mutate(year = as.numeric(year),
         total = as.numeric(gsub("\\,", "", total)),
         domestic = as.numeric(gsub("\\,", "", domestic)),
         gavi = as.numeric(gsub("\\,", "", gavi)),
         domestic_p = as.numeric(domestic_p)/100,
         gavi_p = as.numeric(gavi_p)/100) %>%
  # Convert currency using currency conversion table
  left_join(currency_conv %>% rename(year = dollar_year),
            by = c("country", "year")) %>%
  mutate(total = total * currency_mult,
         domestic = domestic * currency_mult,
         gavi = gavi * currency_mult) %>%
  select(-currency_mult) %>%
  # Some of the vaccine names differ between fund_data and cea_data;
  #   fix.
  mutate(vaccine = ifelse(grepl("DTP-Hep", vaccine),
                          "DTP-hep B",
                          vaccine)) %>%
  mutate(vaccine = ifelse(grepl("Yellow Fever", vaccine),
                          "Yellow Fever",
                          vaccine)) %>%
  mutate(vaccine = ifelse(grepl("Measles", vaccine),
                          "Measles",
                          vaccine)) %>%
  mutate(vaccine = ifelse(grepl("Pneumococcal", vaccine),
                          "PCV",
                          vaccine)) %>%
  mutate(vaccine = ifelse(grepl("Rota", vaccine),
                          "Rotavirus",
                          vaccine))
```

Combine CEA and funding data into a tibble

```r
# Check what vaccines in CEA data are not in co-financing data
temp <- (cea_data %>% distinct(vaccine) %>% unlist) %in%
(fund_data %>% distinct(vaccine) %>% unlist)
(cea_data %>% distinct(vaccine) %>% unlist)[!temp]
```

```
## vaccine3
##    "BCG"
```

```r
# Check what vaccines in co-financing data are not in CEA data
temp <- (fund_data %>% distinct(vaccine) %>% unlist) %in%
(cea_data %>% distinct(vaccine) %>% unlist)
(fund_data %>% distinct(vaccine) %>% unlist)[!temp]
```

```
##                 vaccine6                vaccine7                vaccine8
##                   "MenA" "MR Follow up campaign"          "MenA Routine"
##                 vaccine9
##                    "IPV"
```

```
# Combine CEA and co-financing data by country and vaccine.
#   Include all of the vaccines we have in the CEA analysis data,
#   but not ones we have only in the co-financing data.
data <- full_join(cea_data,
                  fund_data %>% rename(year_gavi = year),
                  by = c("country", "vaccine"))

# Add BCG vaccine, not supporte by GAVI for plotting
BCG_data <- data %>%
  # Get BCG data; at this point only includes Levin et al 2007
  filter(vaccine == "BCG") %>%
  # Copy rows years to create years 2006-2018
  slice(rep(row_number(), length(2006:2018))) %>%
  arrange(indicator) %>%
  mutate(year_gavi = rep(2006:2018, 3)) %>%
  # Remove 2015 since do not have data
  filter(year_gavi != 2015)

# Add the BCG data
data <- data %>%
  # Remove the NA BCG data row
  filter(vaccine != "BCG") %>%
  # Add rest of BCG data
  rbind(BCG_data)
```

Clean combined data.

```
# Remove non base-case
data <- data %>%
  # Include only DALY or YLL
  filter(grepl("DALY|YLL", indicator)) %>%
  # Filter years do not have any GAVI data or only for
  #   few countries and based on future projections
  #   First add a year for BCG, so plot despite no GAVI
  #   funding. Use year of paper.
  mutate(year_gavi = ifelse(is.na(year_gavi) &
                              authors == "Levin et al. 2007" &
                              vaccine == "BCG",
                            2007,
                            year_gavi)) %>%
  filter(year_gavi < 2019) %>%
  # Only include base-case of Abbot et al. 2012
  #  and use earleist year of GAVI funding (since intervention is 2003-08).
  filter(!(authors == "Abbott et al. 2012" &
             (assumptions != "$5 per dose"))) %>%# | year_gavi != 2012))) %>%
  # Do not include the CEA with GAVI Subsidy accounted for
  #   and focus on health system perspective.
  filter(!(authors == "Nonvignon et al. 2018" &
             (grepl("Ghana only", assumptions) |
```

```
            !grepl("Health system", perspective)))) %>%
  # Use relevant GAVI data year for Krishnamoorthy et al. 2019
  #   based on paper intervention years
  # filter(!(authors == "Krishnamoorthy et al. 2019" &
  #            year_gavi != 2018)) %>%
  # Use the analysis with the 90% intervention coverage and year 2018
  #   for GAVI funding based on paper intervention years
  filter(!(authors == "Megiddo et al. 2018" &
            (grepl("similar to DPT", assumptions)))) %>%# |
  #              year_gavi != 2018))) %>%
  # Use most recent year of GAVI funding for Ojal et al. 2019
  #   based on paper intervention years
  #filter(!(authors == "Ojal et al. 2019" & year_gavi < 2016)) %>%
  # Do not include the CEA with GAVI subsidy accounted for
  filter(!(authors == "Debellut et al., 2019" &
            (assumptions == "Vaccine programme costs with Gavi subsidy"))) %>%# |
               # (year_gavi < 2017 & country == "Ghana") |
               # (year_gavi < 2018 & country == "India") |
               # (year_gavi < 2015 & country == "Angola")))) %>%
  # Remove rows with no co-financing data
  filter(!(is.na(total) & (!is.na(gavi) | !is.na(domestic)))) %>%
  # Add a column with vaccine and paper
  mutate(vaccine_cea_paper = paste0(vaccine, "\n", authors)) %>%
  # Remove columns that do not need for figures
  select(-cost_for_horizon, -benefit_for_horizon, -currency_mult)
```

Figures

All Ghana figure

```
# Add plotting themes.
library(ggthemes)

# Extract Ghana data
data_ghana <- data %>%
  rename(year = year_gavi) %>%
  filter(country == "Ghana" &
           grepl("DALY|YLL", indicator)) %>%
  select(vaccine_cea_paper, indicator, cost, benefit, cea, year, total,
         domestic, gavi, domestic_p, gavi_p, s_year, e_year, vaccine, perspective) %>%
  # Data for 2018 only on BCG which is not supported
  filter(year < 2018)

# Max axes
max_ghana_x <- data_ghana %>%
  summarise(max(cost)) %>% unlist %>%
  roundUp + 10000000
max_ghana_y <- data_ghana %>%
  summarise(max(benefit)) %>% unlist %>%
  roundUp
```

```r
# Select relevant GDP data (depending on which years have data
#   for Ghana)
gdp_data_ghana <- left_join(data_ghana %>% select(year),
                            gdp_data %>% filter(country == "Ghana"),
                            by = "year") %>%
  # Data for 2018 only on BCG which is not supported
  filter(year < 2018)

# Plot legend lables
legend_lables <- data_ghana %>%
  distinct(vaccine_cea_paper) %>%
  unlist %>% sort



# Plot Ghana
p1 <- ggplot() +
  # Facet by year and threshold
  facet_grid(year~threshold, scales = "free") +
  # Threshold line
  geom_line(data = gdp_data_ghana %>%
              filter(country == "Ghana") %>%
              filter(cost < max_ghana_x),
            mapping = aes(x = cost/1000000, y = cea_line/1000), linetype = 2) +
  # Costs and effectivness point
  geom_point(data = data_ghana,
             mapping = aes(x = cost/1000000, y = benefit/1000,
                           color = vaccine_cea_paper,
                           shape = vaccine_cea_paper),
             size = 2.5) +
  # Arrow based on proportion of GAVI support relative to deomestic
  geom_segment(data = data_ghana,
               aes(x = cost/1000000, xend = (cost - gavi)/1000000,#(cost * domestic_p)/1000000,
                   y = benefit/1000, yend = benefit/1000),
               arrow = arrow(length = unit(0.1, "cm"))) +
  # x-axis name and $ formatting
  scale_x_continuous(name = "Cost (in millions USD 2018)",
                     labels = scales::dollar) +
  # y-axis name and formatting
  scale_y_continuous(name ="DALYs averted (in thousdands)",
                     labels = scales::comma) +
  # Remove color/shape legend name
  # Display color legend in two rows and remove shape legend
  # Set colors and shapes
  guides(color = guide_legend(nrow = 2, byrow = TRUE)) +
  scale_color_manual(name = "",
                     labels = legend_lables,
                     values = gdocs_pal()(7)) +

  scale_shape_manual(name = "",
                     labels = legend_lables,
                     values = c(15, 16, 18, 17, 17, 17, 7)) +
  # Change theme of plot
```

```r
  theme_fivethirtyeight() +
  theme(legend.position = "bottom",
        axis.title = element_text(),
        plot.caption = element_text(hjust = 0))

# Save figure
ggsave("figures/ghana_all.png", p1, dpi = 300,
       width = 8*2.5, height = 10*2.5, units = "cm")
```

```
## Warning: Removed 44 rows containing missing values (geom_segment).
```

```r
# Create and save figure with appendix caption
p1 +
  # Add a caption
  labs(caption = paste(
    "Figure A1. GAVI contribution to vaccines in Ghana.",
    "Columns differ by cost-effectiveness thresholds, represented by the dashed black line;",
    "Rows differ by the year of GAVI funding (years missing are due to lack of data);",
    "Points represent costs and benefits are extracted from cost-effectiveness (CE) studies;",
    "Arrows represent financing by GAVI (CE studies typically include costs not covered by GAVI",
    "such as distribution costs).",
    #"Arrows represent the proportion of costs funded by GAVI (CE studies and total funding do not matc
    sep = "\n"))
```

```
## Warning: Removed 44 rows containing missing values (geom_segment).
```

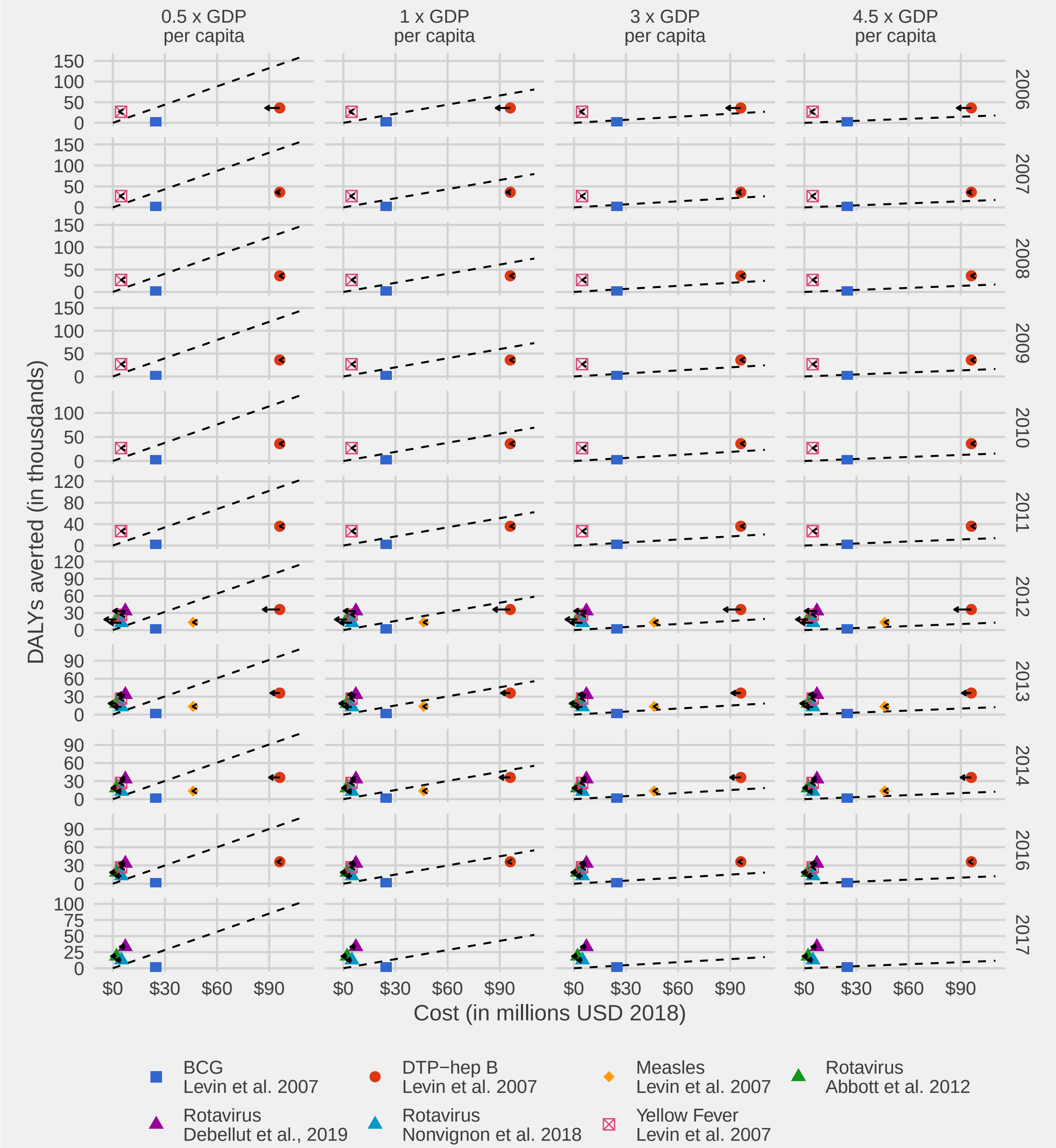


Figure A1. GAVI contribution to vaccines in Ghana.
Columns differ by cost−effectiveness thresholds, represented by the dashed black line;
Rows differ by the year of GAVI funding (years missing are due to lack of data);
Points represent costs and benefits are extracted from cost−effectiveness (CE) studies;
Arrows represent financing by GAVI (CE studies typically include costs not covered by GAVI
such as distribution costs).

```
ggsave("figures/A1_ghana_all_appendix_caption.png", dpi = 300,
       width = 8*2.5, height = 10*2.5, units = "cm")
```

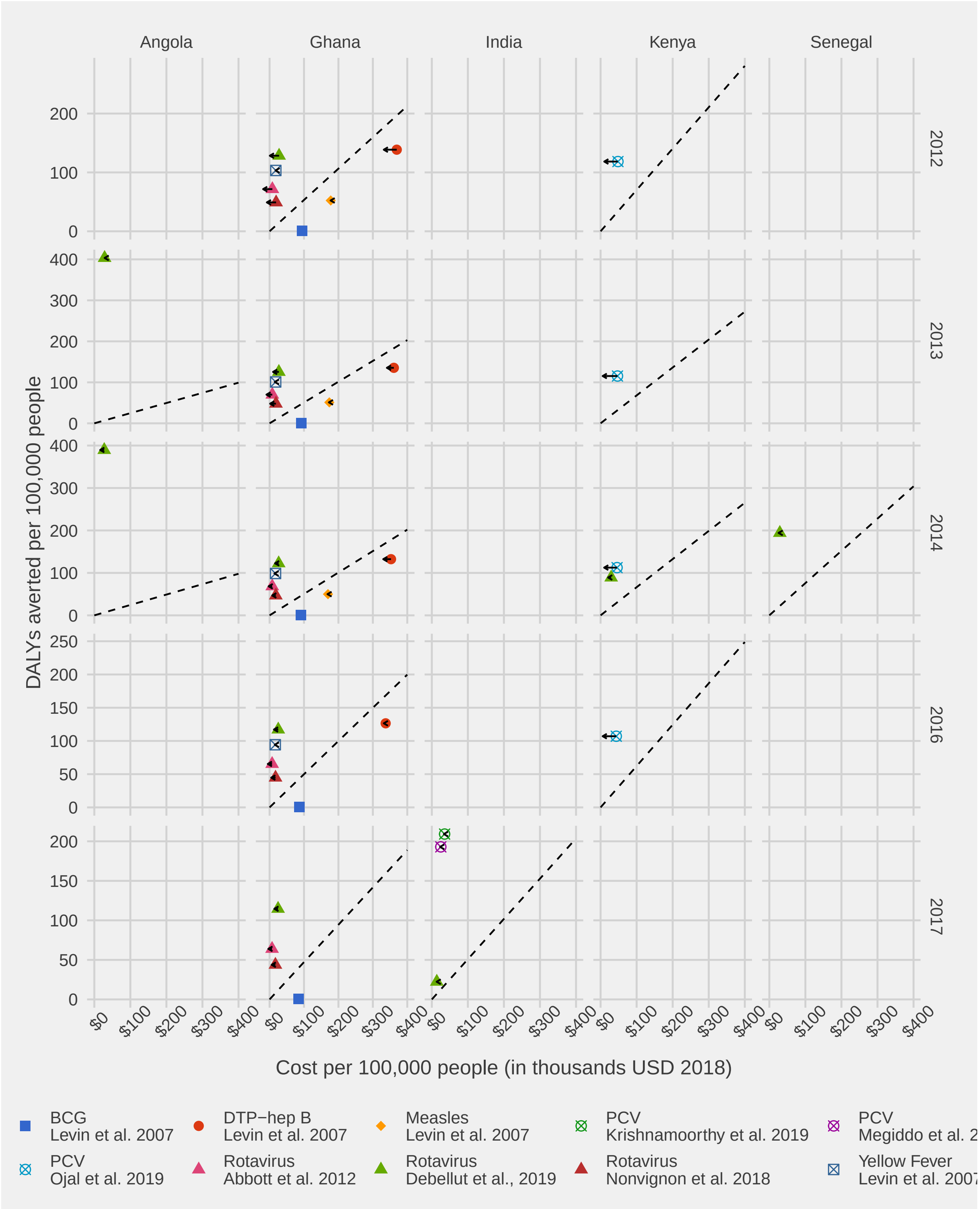
```
## Warning: Removed 44 rows containing missing values (geom_segment).
```

All cross country figure 1 x GDP

```r
# Get years that include non Ghana countries
years_include <- data %>%
  filter(country != "Ghana") %>% distinct(year_gavi) %>% unlist

# All country data
data_all_countries <-
  data %>%
  rename(year = year_gavi) %>%
  filter(grepl("DALY|YLL", indicator)) %>%
  filter(year %in% years_include) %>%
  select(country, vaccine_cea_paper, indicator, cost, benefit, cea, year, total,
         domestic, gavi, domestic_p, gavi_p, s_year, e_year, vaccine, perspective) %>%
  # Data for 2018 only on BCG which is not supported
  #   Also remove 2015 for which there is very limited data
  filter(year < 2018 & year != 2015) %>%
  # Add population data
  left_join(population_data, by = c("country", "year")) %>%
  # Convert values to per 100,000 population
  mutate(cost = cost/population*100000,
         benefit = benefit/population*100000,
         gavi = gavi/population*100000,
         domestic = domestic/population*100000)

# Max axes
max_all_countries_x <- data_all_countries %>%
  summarise(max(cost)) %>% unlist %>%
  roundUp + 10
max_all_countries_y <- data_all_countries %>%
  summarise(max(benefit)) %>% unlist %>%
  roundUp

# GDP data selct
gdp_data_all_countries <-
  left_join(data_all_countries %>% select(year, country),
            gdp_data,
            by = c("year", "country")) %>%
  filter(year %in% years_include) %>%
  # Data for 2018 only on BCG which is not supported
  #   Also remove 2015 for which there is very limited data
  filter(year < 2018 & year != 2015)

# Plot legend lables
legend_lables <- data_all_countries %>%
  distinct(vaccine_cea_paper) %>%
  unlist %>% sort

# Plot all countries using 1 X GDP per capita threshold
p2 <- ggplot() +
  # Facet by country and year
  facet_grid(year~country, scales = "free") +
  # Threshold line
```

```
  geom_line(data = gdp_data_all_countries %>%
              filter(threshold == "1 x GDP\nper capita") %>%
              filter(cost < max_all_countries_x),
            mapping = aes(x = cost/1000, y = cea_line), linetype = 2) +
  # Costs and effectivness point
  geom_point(data = data_all_countries,
             mapping = aes(x = cost/1000, y = benefit,
                           color = vaccine_cea_paper,
                           shape = vaccine_cea_paper),
             size = 2.5) +
  # Arrow based on proportion of GAVI support relative to deomestic
  geom_segment(data = data_all_countries,
               aes(x = cost/1000, xend = (cost - gavi)/1000,#(cost/1000 * domestic_p),
                   y = benefit, yend = benefit),
               arrow = arrow(length = unit(0.1, "cm"))) +
  # x-axis name and $ formatting
  scale_x_continuous(name = "Cost per 100,000 people (in thousands USD 2018)",
                     labels = scales::dollar) +
  # y-axis name and formatting
  scale_y_continuous(name ="DALYs averted per 100,000 people",
                     labels = scales::comma) +
  # Remove color/shape legend name
  # Display color legend in two rows and remove shape legend
  # Set colors and shapes
  guides(color = guide_legend(nrow = 2, byrow = TRUE)) +
  scale_color_manual(name = "",
                     labels = legend_lables,
                     values = gdocs_pal()(10)) +

  scale_shape_manual(name = "",
                     labels = legend_lables,
                     values = c(15, 16, 18,  13, 13, 13, 17, 17, 17, 7)) +
  # Display color legend in 4 columns
  # Change theme of plot
  theme_fivethirtyeight() +
  theme(legend.position = "bottom",
        axis.title = element_text(),
        axis.text.x = element_text(angle = 45),
        plot.caption = element_text(hjust = 0))
p2
```

```
## Warning: Removed 5 rows containing missing values (geom_segment).
```




```r
# Save figure
ggsave("figures/all_country_compare_1GDP.png", p2, dpi = 300,
       width = 8*2.5, height = 10*2.5, units = "cm")
```

```
## Warning: Removed 5 rows containing missing values (geom_segment).
```

```r
p2 + labs(
  caption = paste(
    "Figure A2. GAVI contribution to vaccines in Ghana, Angola, India, Kenya, and Senegal.",
    "The dashed black line is the 1 x GDP cost-effectiveness threshold;",
    "Points represent costs and benefits are extracted from cost-effectiveness (CE) studies;",
    "Arrows represent financing by GAVI (CE studies typically include costs not covered by GAVI",
    "such as distribution costs).",
    #"Arrows represent the proportion of costs funded by GAVI (CE studies and total funding do not matc
    sep = "\n"))
```

```
## Warning: Removed 5 rows containing missing values (geom_segment).
```

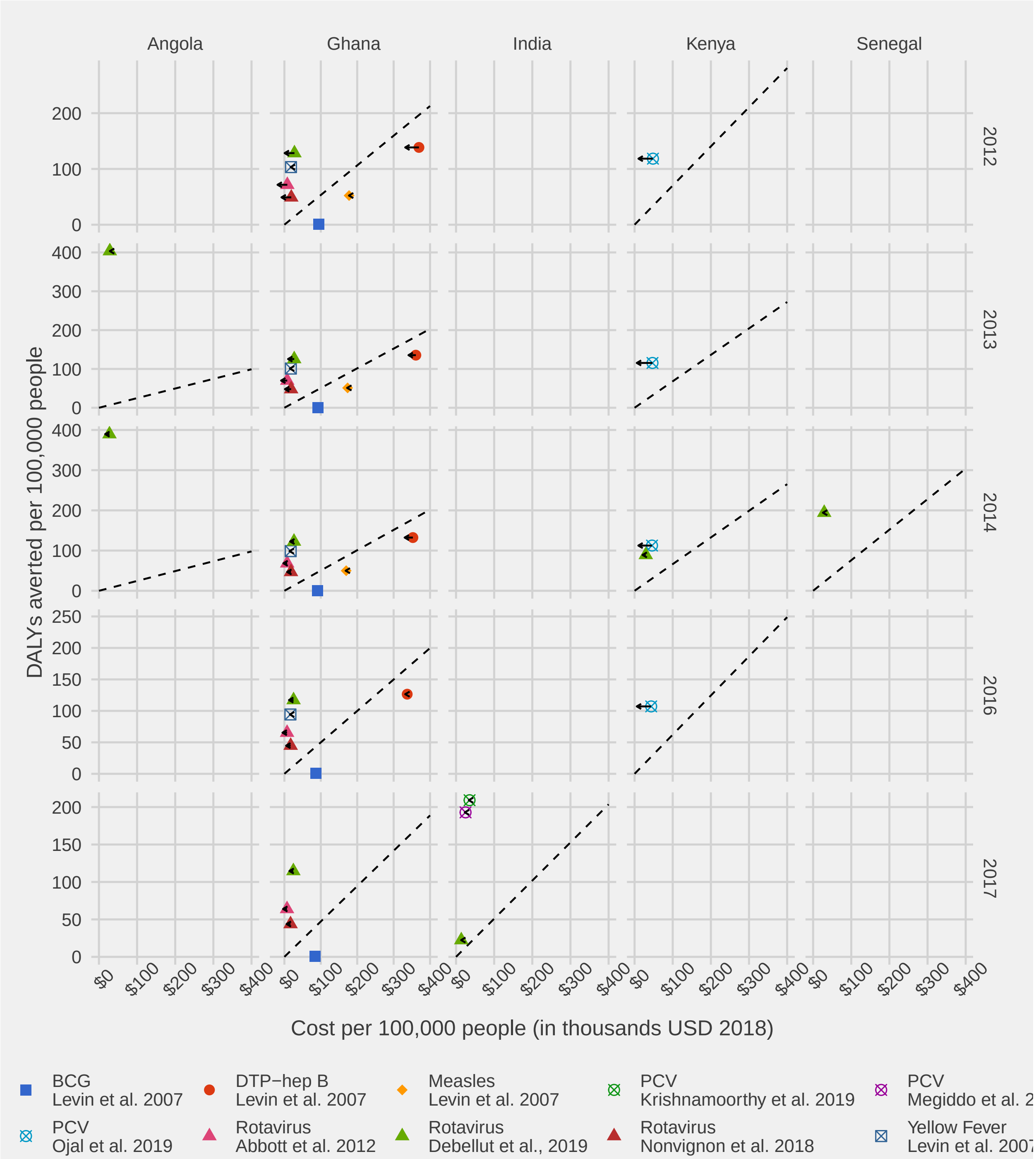


Figure A2. GAVI contribution to vaccines in Ghana, Angola, India, Kenya, and Senegal.
The dashed black line is the 1 x GDP cost−effectiveness threshold;
Points represent costs and benefits are extracted from cost−effectiveness (CE) studies;
Arrows represent financing by GAVI (CE studies typically include costs not covered by GAVI
such as distribution costs).

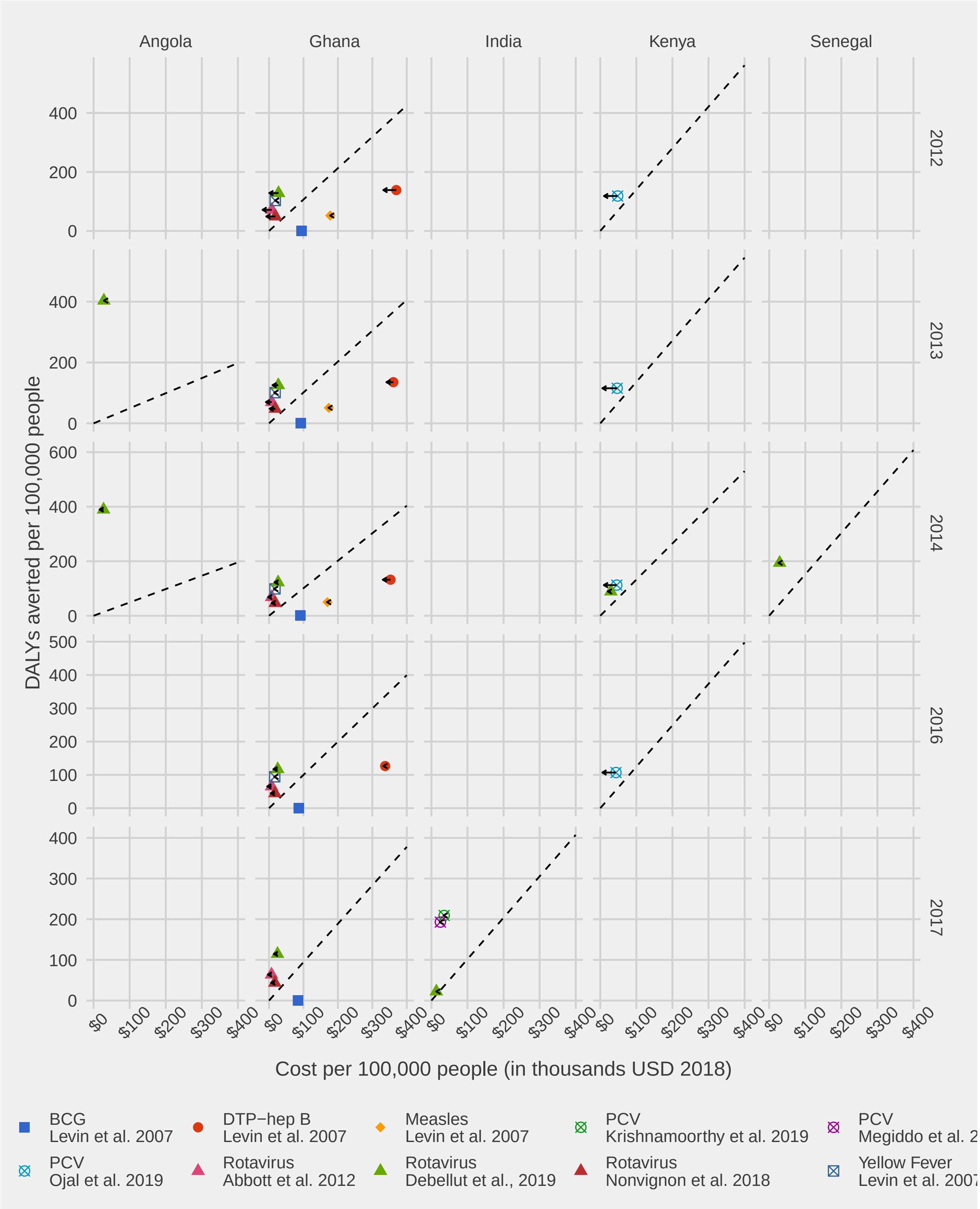
```r
# Create and save figure with appendix caption
ggsave("figures/A2_all_country_compare_1GDP.png", dpi = 300,
       width = 8*2.5, height = 10*2.5, units = "cm")
```

```
## Warning: Removed 5 rows containing missing values (geom_segment).
```

All cross country figure 0.5 x GDP

```r
# Plot all countries using 1 X GDP per capita threshold
p3 <- ggplot() +
  # Facet by country and year
  facet_grid(year~country, scales = "free") +
  # Threshold line
  geom_line(data = gdp_data_all_countries %>%
              filter(threshold == "0.5 x GDP\nper capita") %>%
              filter(cost < max_all_countries_x),
            mapping = aes(x = cost/1000, y = cea_line), linetype = 2) +
  # Costs and effectivness point
  geom_point(data = data_all_countries,
             mapping = aes(x = cost/1000, y = benefit,
                           color = vaccine_cea_paper,
                           shape = vaccine_cea_paper),
             size = 2.5) +
  # Arrow based on proportion of GAVI support relative to deomestic
  geom_segment(data = data_all_countries,
               aes(x = cost/1000, xend = (cost - gavi)/1000,#(cost/1000 * domestic_p),
                   y = benefit, yend = benefit),
               arrow = arrow(length = unit(0.1, "cm"))) +
  # x-axis name and $ formatting
  scale_x_continuous(name = "Cost per 100,000 people (in thousands USD 2018)",
                     labels = scales::dollar) +
  # y-axis name and formatting
  scale_y_continuous(name ="DALYs averted per 100,000 people",
                     labels = scales::comma) +
  # Remove color/shape legend name
  # Display color legend in four rows and remove shape legend
  # Set colors and shapes
  guides(color = guide_legend(nrow = 2, byrow = TRUE)) +
  scale_color_manual(name = "",
                     labels = legend_lables,
                     values = gdocs_pal()(10)) +

  scale_shape_manual(name = "",
                     labels = legend_lables,
                     values = c(15, 16, 18,  13, 13, 13, 17, 17, 17, 7)) +
  # Display color legend in 4 columns
  # Change theme of plot
  theme_fivethirtyeight() +
  theme(legend.position = "bottom",
        axis.title = element_text(),
        axis.text.x = element_text(angle = 45),
        plot.caption = element_text(hjust = 0))
p3
```

```
## Warning: Removed 5 rows containing missing values (geom_segment).
```




```r
# Save figure
ggsave("figures/all_country_compare_05GDP.png", p2, dpi = 300,
       width = 8*2.5, height = 10*2.5, units = "cm")
```

```
## Warning: Removed 5 rows containing missing values (geom_segment).
```

```r
p2 + labs(
  caption = paste(
    "Figure A2. GAVI contribution to vaccines in Ghana, Angola, India, Kenya, and Senegal.",
    "The dashed black line is the 0.5 x GDP cost-effectiveness threshold;",
    "Points represent costs and benefits are extracted from cost-effectiveness (CE) studies;",
    "Arrows represent financing by GAVI (CE studies typically include costs not covered by GAVI",
    "such as distribution costs).",
    #"Arrows represent the proportion of costs funded by GAVI (CE studies and total funding do not matc
    sep = "\n"))
```

```
## Warning: Removed 5 rows containing missing values (geom_segment).
```

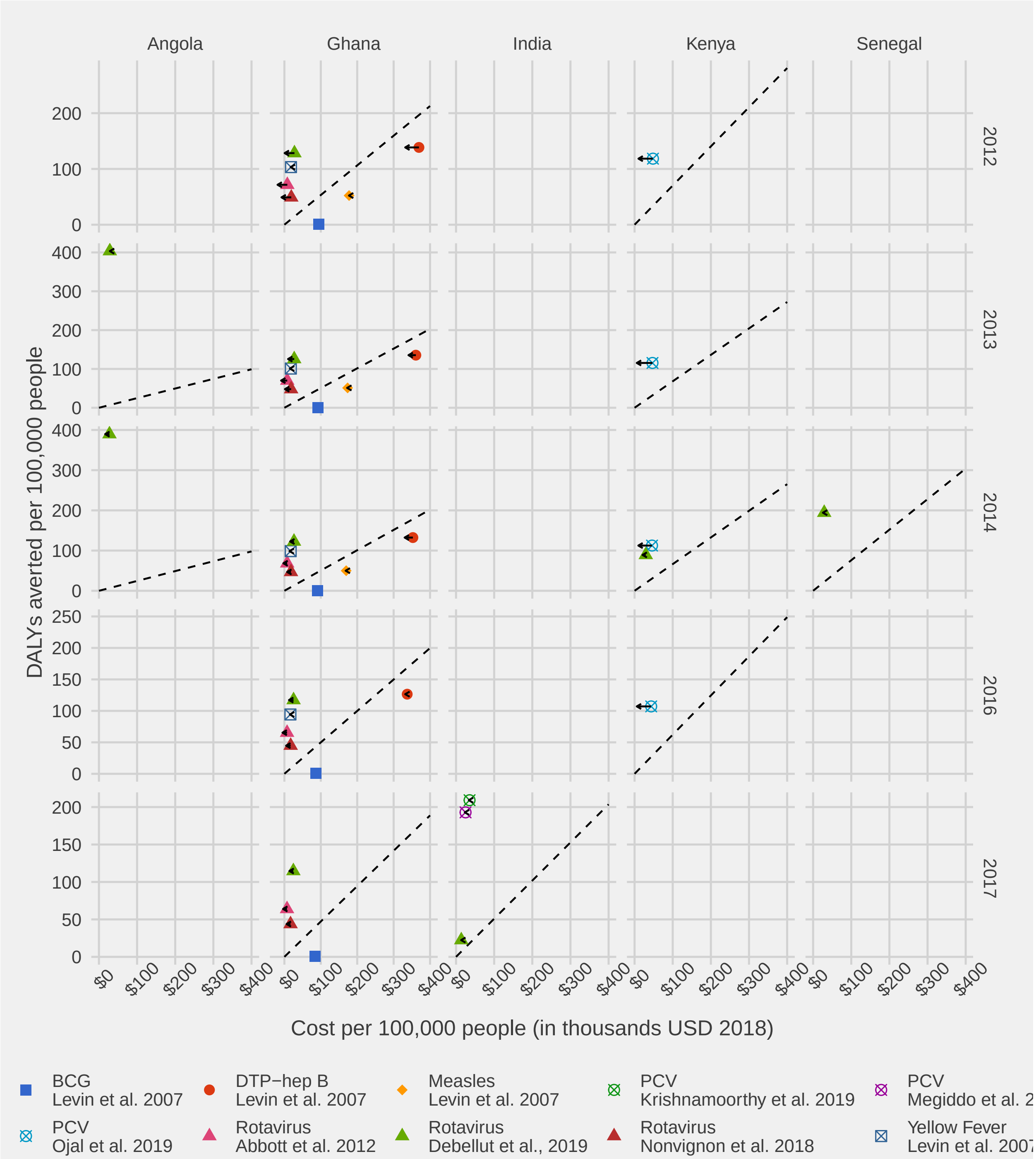


Figure A2. GAVI contribution to vaccines in Ghana, Angola, India, Kenya, and Senegal.
The dashed black line is the 0.5 x GDP cost−effectiveness threshold;
Points represent costs and benefits are extracted from cost−effectiveness (CE) studies;
Arrows represent financing by GAVI (CE studies typically include costs not covered by GAVI
such as distribution costs).

```
# Create and save figure with appendix caption
ggsave("figures/A2_all_country_compare_05GDP.png", dpi = 300,
       width = 8*2.5, height = 10*2.5, units = "cm")
```

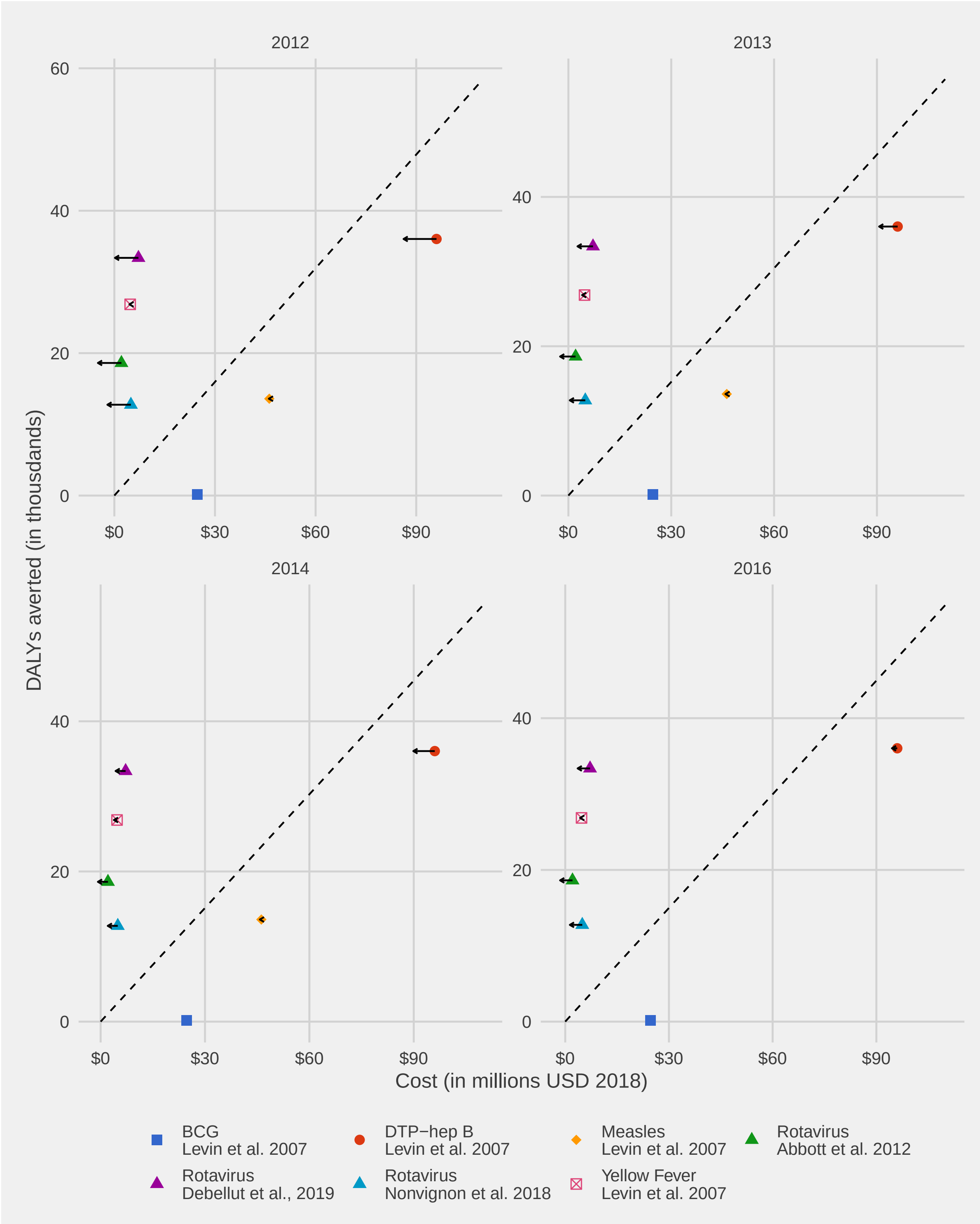
```
## Warning: Removed 5 rows containing missing values (geom_segment).
```

Ghana 1 x GDP 2012-2016 (most data for these years)

```
# Plot legend lables
legend_lables <- data_ghana %>%
  distinct(vaccine_cea_paper) %>%
  unlist %>% sort

# Plot Ghana
p4 <- ggplot() +
  # Facet by year and threshold
  facet_wrap(year~., scales = "free", ncol = 2) +
  # Threshold line
  geom_line(data = gdp_data_ghana %>%
              filter(country == "Ghana") %>%
              filter(cost < max_ghana_x) %>%
              filter(threshold == "1 x GDP\nper capita") %>%
              filter(year %in% 2012:2016),
            mapping = aes(x = cost/1000000, y = cea_line/1000), linetype = 2) +
  # Costs and effectivness point
  geom_point(data = data_ghana %>%
               filter(year %in% 2012:2016),
             mapping = aes(x = cost/1000000, y = benefit/1000,
                           color = vaccine_cea_paper,
                           shape = vaccine_cea_paper),
             size = 2.5) +
  # Arrow based on proportion of GAVI support relative to deomestic
  geom_segment(data = data_ghana %>%
                 filter(year %in% 2012:2016),
               aes(x = cost/1000000, xend = (cost - gavi)/1000000,#(cost * domestic_p)/1000000,
                   y = benefit/1000, yend = benefit/1000),
               arrow = arrow(length = unit(0.1, "cm"))) +
  # x-axis name and $ formatting
  scale_x_continuous(name = "Cost (in millions USD 2018)",
                     labels = scales::dollar) +
  # y-axis name and formatting
  scale_y_continuous(name ="DALYs averted (in thousdands)",
                     labels = scales::comma) +
  # Remove color/shape legend name
  # Display color legend in two rows and remove shape legend
  # Set colors and shapes
  guides(color = guide_legend(nrow = 2, byrow = TRUE)) +
  scale_color_manual(name = "",
                     labels = legend_lables,
                     values = gdocs_pal()(7)) +
  scale_shape_manual(name = "",
                     labels = legend_lables,
                     values = c(15, 16, 18, 17, 17, 17, 7)) +
  # Change theme of plot
  theme_fivethirtyeight() +
  theme(legend.position = "bottom",
```

```
        axis.title = element_text(),
        plot.caption = element_text(hjust = 0))
p4
```

```
## Warning: Removed 4 rows containing missing values (geom_segment).
```




```
# Save figure
ggsave("figures/ghana_x1gdp.png", p4, dpi = 300,
       width = 8*2.5, height = 10*2.5, units = "cm")
```

```
## Warning: Removed 4 rows containing missing values (geom_segment).
```

```r
# Create and save figure with appendix caption
p4 +
  # Add a caption
  labs(caption = paste(
    "Figure 1. GAVI contribution to vaccines in Ghana.",
    "The dashed black line is the 0.5 x GDP-based cost-effectiveness threshold;",
    "Each plot represents a differe year of GAVI financing;",
    "Points represents costs and benefits extracted from cost-effectiveness (CE) studies;",
    "Arrows represent financing by GAVI (CE studies typically include costs not covered by GAVI",
    "such as distribution costs).",
    #"Arrows represent the proportion of costs funded by GAVI (CE studies and total funding do not matc
    sep = "\n"))
```

```
## Warning: Removed 4 rows containing missing values (geom_segment).
```

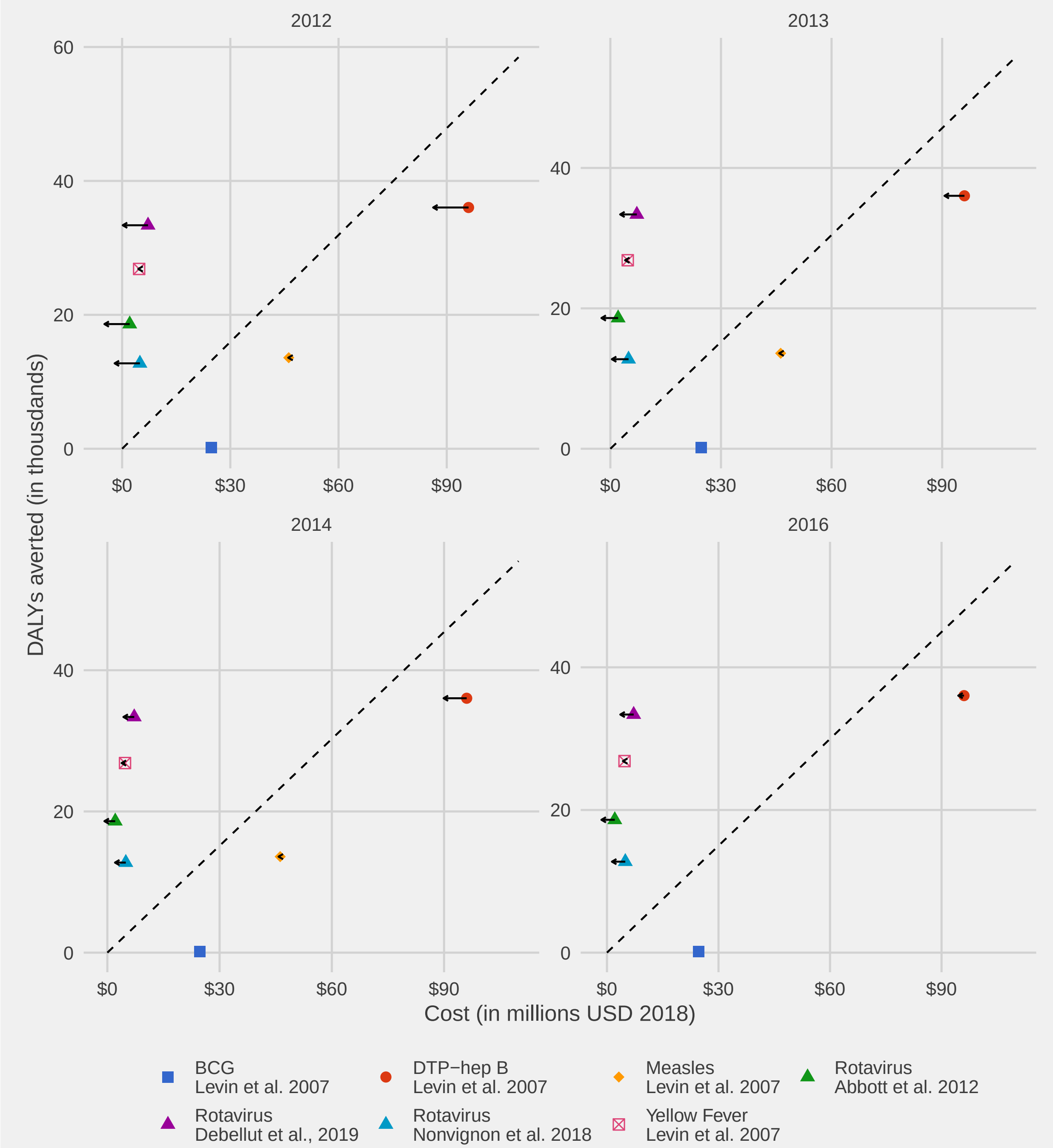


Figure 1. GAVI contribution to vaccines in Ghana.
The dashed black line is the 0.5 x GDP−based cost−effectiveness threshold;
Each plot represents a differe year of GAVI financing;
Points represents costs and benefits extracted from cost−effectiveness (CE) studies;
Arrows represent financing by GAVI (CE studies typically include costs not covered by GAVI
such as distribution costs).

```
ggsave("figures/A3_all_ghana_x1gdp_appendix_caption.png", dpi = 300,
       width = 8*2.5, height = 10*2.5, units = "cm")
```

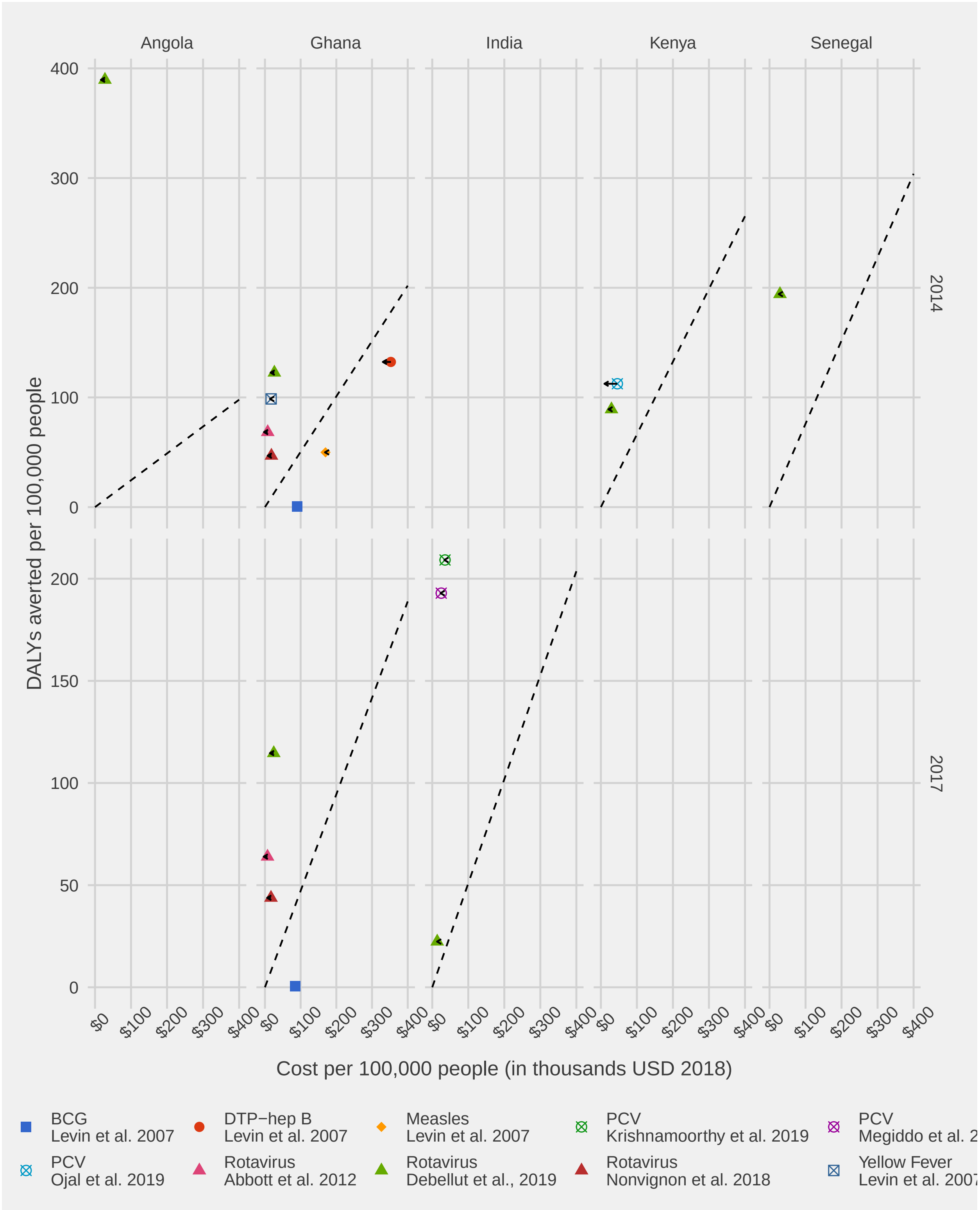
```
## Warning: Removed 4 rows containing missing values (geom_segment).
```

Country comparison 1 x GDP per capita

```
# Plot legend lables
legend_lables <- data_all_countries %>%
  distinct(vaccine_cea_paper) %>%
  unlist %>% sort

# Plot all countries using 1 X GDP per capita threshold
p2 <- ggplot() +
  # Facet by country and year
  facet_grid(year~country, scales = "free") +
  # Threshold line
  geom_line(data = gdp_data_all_countries %>%
              filter(threshold == "1 x GDP\nper capita") %>%
              filter(cost < max_all_countries_x) %>%
              filter(year %in% c(2014, 2017)),
            mapping = aes(x = cost/1000, y = cea_line), linetype = 2) +
  # Costs and effectivness point
  geom_point(data = data_all_countries %>%
              filter(year %in% c(2014, 2017)),
             mapping = aes(x = cost/1000, y = benefit,
                           color = vaccine_cea_paper,
                           shape = vaccine_cea_paper),
             size = 2.5) +
  # Arrow based on proportion of GAVI support relative to deomestic
  geom_segment(data = data_all_countries %>%
              filter(year %in% c(2014, 2017)),
               aes(x = cost/1000, xend = (cost - gavi)/1000,#(cost/1000 * domestic_p),
                   y = benefit, yend = benefit),
               arrow = arrow(length = unit(0.1, "cm"))) +
  # x-axis name and $ formatting
  scale_x_continuous(name = "Cost per 100,000 people (in thousands USD 2018)",
                     labels = scales::dollar) +
  # y-axis name and formatting
  scale_y_continuous(name ="DALYs averted per 100,000 people",
                     labels = scales::comma) +
  # Remove color/shape legend name
  # Display color legend in four rows and remove shape legend
  # Set colors and shapes
  guides(color = guide_legend(nrow = 2, byrow = TRUE)) +
  scale_color_manual(name = "",
                     labels = legend_lables,
                     values = gdocs_pal()(10)) +

  scale_shape_manual(name = "",
                     labels = legend_lables,
                     values = c(15, 16, 18,  13, 13, 13, 17, 17, 17, 7)) +
  # Display color legend in 4 columns
  # Change theme of plot
  theme_fivethirtyeight() +
  theme(legend.position = "bottom",
        axis.title = element_text(),
        axis.text.x = element_text(angle = 45),
```

```
        plot.caption = element_text(hjust = 0))
p2
```

```
## Warning: Removed 2 rows containing missing values (geom_segment).
```




```
# Save figure
ggsave("figures/country_compare_1GDP.png", p2, dpi = 300,
       width = 8*2.5, height = 10*2.5, units = "cm")
```

```
## Warning: Removed 2 rows containing missing values (geom_segment).
```

```r
p2 + labs(
  caption = paste(
    "Figure A2. GAVI contribution to vaccines in Ghana, Angola, India, Kenya, and Senegal.",
    "The dashed black line is the 1 x GDP cost-effectiveness threshold;",
    "Points represent costs and benefits are extracted from cost-effectiveness (CE) studies;",
    "Arrows represent financing by GAVI (CE studies typically include costs not covered by GAVI",
    "such as distribution costs).",
    #"Arrows represent the proportion of costs funded by GAVI (CE studies and total funding do not matc
    sep = "\n"))
```

```
## Warning: Removed 2 rows containing missing values (geom_segment).
```

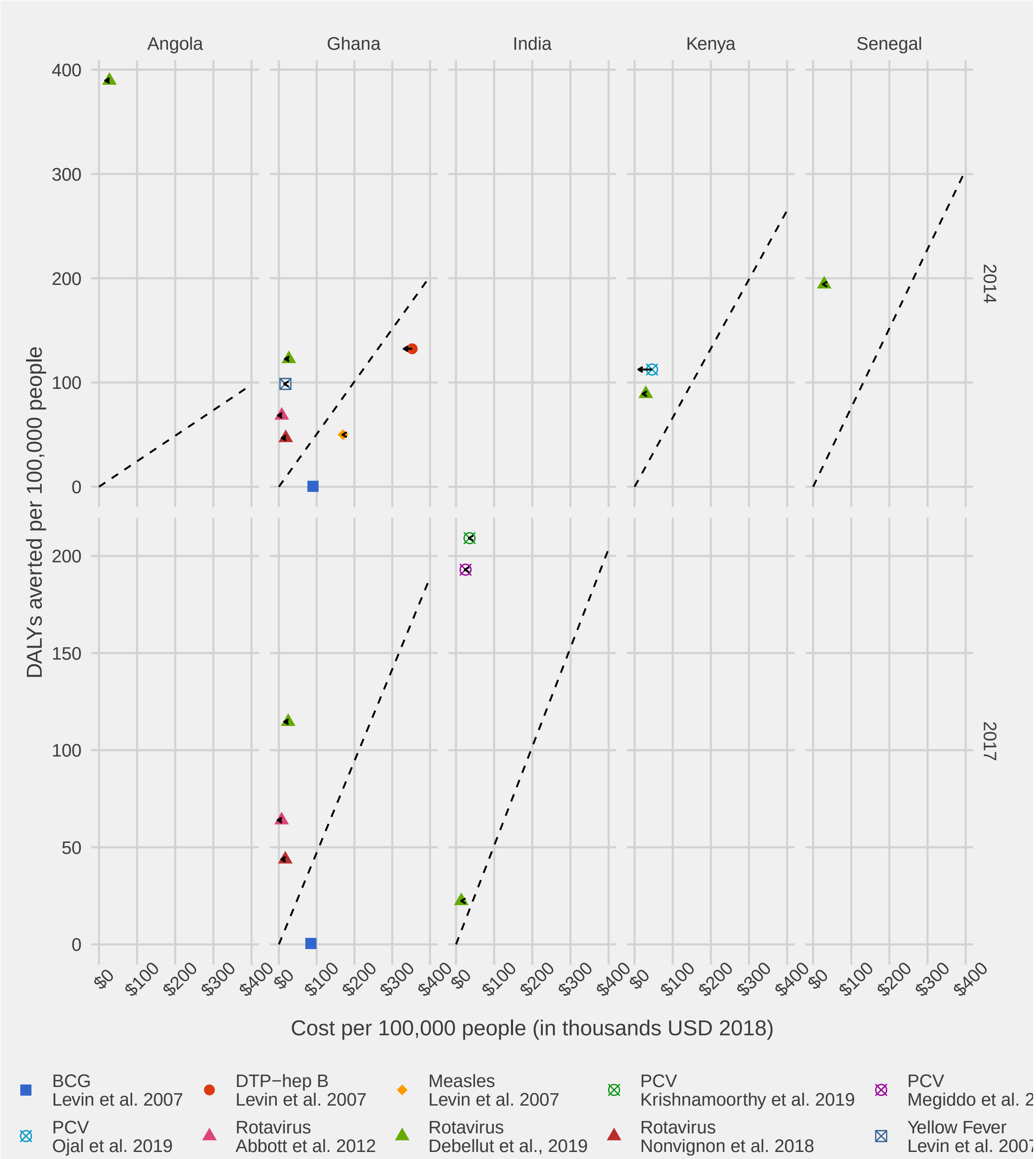


Figure A2. GAVI contribution to vaccines in Ghana, Angola, India, Kenya, and Senegal.
The dashed black line is the 1 x GDP cost−effectiveness threshold;
Points represent costs and benefits are extracted from cost−effectiveness (CE) studies;
Arrows represent financing by GAVI (CE studies typically include costs not covered by GAVI
such as distribution costs).

```
# Create and save figure with appendix caption
ggsave("figures/A4_country_compare_1GDP.png", dpi = 300,
       width = 8*2.5, height = 10*2.5, units = "cm")
```

```
## Warning: Removed 2 rows containing missing values (geom_segment).
```