

Fine-scale differentiation between *B. anthracis* and *B. cereus* group signatures in metagenome shotgun data

Packages and Functions

```
library(dplyr)

##
## Attaching package: 'dplyr'

## The following objects are masked from 'package:stats':
##
##     filter, lag

## The following objects are masked from 'package:base':
##
##     intersect, setdiff, setequal, union

library(ggplot2)
options(scipen=999)
library(grid)
library(gridExtra)

##
## Attaching package: 'gridExtra'

## The following object is masked from 'package:dplyr':
##
##     combine

PATH <- '/home/rpettit/ba-paper'
FIGURE <- '/home/rpettit/ba-paper/results/figures'

read_samples <- function(gzip_file, distance) {
  s = merge(
    read.table(gzfile(gzip_file), header=TRUE),
    data.frame(sample=distance$accession, distance=distance$distance),
    by=c('sample')
  )

  s$bacillus = ifelse(
    s$is_bcg == 'True', ifelse(s$is_ba=='True', 'ba', 'bcg'), 'nonbcg'
  )

  return(s)
}

format_plot <- function(ggplot_object, xlab="Mean k-mer Coverage", ylab="Count", title=FALSE) {
  plot_title = theme(legend.title = element_blank())
  if (title != FALSE) {
    plot_title <- labs(color=title)
  }
  p <- ggplot_object +
```

```

    xlab(xlab) +
    ylab(ylab) +
    theme_bw() +
    theme(axis.text=element_text(size=12),
          axis.title=element_text(size=14,face="bold")) +
    plot_title
  return(p)
}

plot_hamming <- function(df) {
  p <- ggplot(df, aes(hamming_distance, hit)) +
    xlab("Hamming Distance") +
    ylab("Count") +
    geom_bar(stat='identity') +
    scale_x_continuous(breaks = seq(
      min(df$hamming_distance), max(df$hamming_distance), by = 1
    )) +
    geom_text(aes(label=hit), vjust = -0.5) +
    theme_bw() +
    theme(text=element_text(size=20),
          title=element_text(size=14, face="bold"),
          legend.position="none")
  return (p)
}

lm_eqn <- function(m, zero_intercept=TRUE){
  # GET EQUATION AND R-SQUARED AS STRING
  # SOURCE: http://goo.gl/K4yh
  a = ifelse(zero_intercept, 0, coef(m)[1])
  b = ifelse(zero_intercept, coef(m)[1], coef(m)[2])
  eq <- substitute(italic(y) == a + b %.% italic(x)*", "~~italic(r)^2~~"=~r2,
    list(a = format(a, digits = 2), b = format(b, digits = 2),
         r2 = format(summary(m)$r.squared, digits = 3)))
  as.character(as.expression(eq));
}

#' write_plot
#'
#' A wrapper for to validate given vector is multiple ids and proper type. This
#' function should not be directly used by the user.
#'
#' @param plot_object A ggplot object
#' @param name Basename for the output PDF and PNG files
#' @param height The PDF height of the output (Default: 5)
#' @param width The PDF width of the object (Default: 12)
#'
#' @export
#' @return bool TRUE is multiple ids else FALSE.
write_plot <- function(plot_object, name, height = 5, width = 12) {
  pdf(paste0(FIGURE, '/', name, ".pdf"), width=width, height=height)
  print(plot_object)
  dev_null <- dev.off()
}

```

```

  png(paste0(FIGURE, '/', name, ".png"), width=width*100, height=height*100)
  print(plot_object)
  dev_null <- dev.off()
}

```

Lethal Factor Gray Zone

Inputs

```

gray_ba_samples = read.table(
  gzfile(paste0(PATH, "/results/limit-of-detection-lef31/lef-ba-summary.txt.gz")),
  header=TRUE, sep="\t"
)

gray_bcg_samples = read.table(
  gzfile(paste0(PATH, "/results/limit-of-detection-lef31/lef-bcg-summary.txt.gz")),
  header=TRUE, sep="\t"
)

gray_lef_samples = read.table(
  gzfile(paste0(PATH, "/results/limit-of-detection-lef31/lef-lef-summary.txt.gz")),
  header=TRUE, sep="\t"
)

gray_summary = read.table(
  paste0(PATH, "/results/limit-of-detection-lef31/lef-subsample.txt"),
  header=TRUE, sep="\t"
)

gray_summary <- merge(
  gray_summary,
  gray_ba_samples %>%
    group_by(sample, coverage) %>%
    summarise(ba_tp=mean(tp), ba_coverage=mean(kmer_cov_mean),
              ba_coverage_nonzero=mean(non_zero_kmer_cov_mean)),
    by=c('sample', 'coverage')
)

gray_summary <- merge(
  gray_summary,
  gray_bcg_samples %>%
    group_by(sample, coverage) %>%
    summarise(bcg_tp=mean(tp), bcg_coverage=mean(kmer_cov_mean),
              bcg_coverage_nonzero=mean(non_zero_kmer_cov_mean)),
    by=c('sample', 'coverage')
)

gray_summary <- merge(
  gray_summary,
  gray_lef_samples %>%

```

```

    group_by(sample, coverage) %>%
    summarise(lef_tp=mean(tp), lef_coverage=mean(kmer_cov_mean),
              lef_coverage_nonzero=mean(non_zero_kmer_cov_mean)),
    by=c('sample', 'coverage')
)

gray_area_summary <- function(df, group, status) {
  temp_df <- data.frame(table(round(df, digits = 5)))
  colnames(temp_df) <- c('coverage', 'count')
  temp_df$group <- group
  temp_df$status <- status
  temp_df$coverage <- as.numeric(as.character(temp_df$coverage))
  temp_df$cumsum <- cumsum(temp_df$count)
  temp_df$percent <- temp_df$cumsum / max(temp_df$cumsum) * 100
  return(temp_df)
}

grid_arrange_shared_lef31 <- function(..., ncol = length(list(...)), nrow = 1, position = c("bottom", "top"),
plots <- list(...)
position <- match.arg(position)
g <- ggplotGrob(plots[[1]] + theme(legend.position = position))$grobs
legend <- g[[which(sapply(g, function(x) x$name) == "guide-box")]]
lheight <- sum(legend$height)
lwidth <- sum(legend$width)
gl <- lapply(plots, function(x) x + theme(legend.position="none"))
gl[[1]] <-arrangeGrob(gl[[1]], top = textGrob(
  "A", x = unit(0, "npc"), y = unit(1, "npc"), just=c("left","top"),
  gp=gpar(col="black", fontsize=18, fontface="bold"))
)
gl[[2]] <-arrangeGrob(gl[[2]], top = textGrob(
  "B", x = unit(0, "npc"), y = unit(1, "npc"), just=c("left","top"),
  gp=gpar(col="black", fontsize=18, fontface="bold"))
)

gl <- c(gl, ncol = ncol, nrow = nrow)

combined <- switch(position,
  "bottom" = arrangeGrob(do.call(arrangeGrob, gl),
                         legend,
                         ncol = 1,
                         heights = unit.c(unit(1, "npc") - lheight, lheight)),
  "right" = arrangeGrob(do.call(arrangeGrob, gl),
                        legend,
                        ncol = 2,
                        widths = unit.c(unit(1, "npc") - lwidth, lwidth)))

grid.newpage()
grid.draw(combined)

# return gtable invisibly
invisible(combined)
}

```

```

head(gray_summary)

##      sample coverage tests successes status     ba_tp ba_coverage
## 1 ERR930296 0.0400    100       95 relaxed  5562.642 0.02347827
## 2 ERR930296 0.0700    100      100 strict   9555.610 0.04074087
## 3 ERR930297 0.0589    100       99 relaxed  8441.505 0.03589068
## 4 ERR930297 0.0900    100      100 strict  12692.440 0.05445297
## 5 ERR930298 0.0590    100       97 relaxed  8437.495 0.03590081
## 6 ERR930298 0.1000    100      100 strict  14226.180 0.06123230
##   ba_coverage_nonzero   bcg_tp bcg_coverage bcg_coverage_nonzero   lef_tp
## 1              1.010975 258.5684  0.02576120             1.013722 189.2316
## 2              1.021273 436.5200  0.04388294             1.023961 301.1000
## 3              1.018254 368.9789  0.03705247             1.023516 208.8737
## 4              1.027492 553.4500  0.05609447             1.031130 340.3700
## 5              1.019040 371.2577  0.03710551             1.017710 204.9175
## 6              1.030878 609.7900  0.06219680             1.038531 304.3400
##   lef_coverage lef_coverage_nonzero
## 1 0.07564708          1.048649
## 2 0.12483760          1.081128
## 3 0.08272228          1.030974
## 4 0.14339702          1.092726
## 5 0.08183211          1.032480
## 6 0.12476118          1.065953

```

Gray Area Group Sumamry

```

gray_area_group_summary <- rbind(
  gray_area_summary(gray_summary[gray_summary$status == 'strict',]$ba_coverage, 'ba', '100%'),
  gray_area_summary(gray_summary[gray_summary$status == 'relaxed',]$ba_coverage, 'ba', '95%'),
  gray_area_summary(gray_summary[gray_summary$status == 'strict',]$bcg_coverage, 'bcg', '100%'),
  gray_area_summary(gray_summary[gray_summary$status == 'relaxed',]$bcg_coverage, 'bcg', '95%'),
  gray_area_summary(gray_summary[gray_summary$status == 'strict',]$lef_coverage, 'lef', '100%'),
  gray_area_summary(gray_summary[gray_summary$status == 'relaxed',]$lef_coverage, 'lef', '95%')
)

```

group column - ba: Ba31 kmer coverage - bcg: BCerG kmer coverage - lef: lef31 kmer coverage

status column - **strict** lef kmer found in 100% of subsamples - **relaxed** lef kmer found in > 95% of subsamples

Ba31 k-mer Coverage & BCerG31 k-mer Coverage

```

temp_df <- gray_area_group_summary[gray_area_group_summary$group == 'ba',]
print(temp_df[temp_df$percent >= 0.95,])

```

```

##      coverage count group status cumsum    percent
## 2    0.01645    1    ba  100%      2  1.219512
## 3    0.01724    1    ba  100%      3  1.829268
## 4    0.01806    1    ba  100%      4  2.439024
## 5    0.01833    1    ba  100%      5  3.048780
## 6    0.01890    1    ba  100%      6  3.658537
## 7    0.01904    1    ba  100%      7  4.268293
## 8    0.01970    1    ba  100%      8  4.878049

```

## 9	0.01989	1	ba	100%	9	5.487805
## 10	0.02065	1	ba	100%	10	6.097561
## 11	0.02071	1	ba	100%	11	6.707317
## 12	0.02136	1	ba	100%	12	7.317073
## 13	0.02147	1	ba	100%	13	7.926829
## 14	0.02159	1	ba	100%	14	8.536585
## 15	0.02206	1	ba	100%	15	9.146341
## 16	0.02367	1	ba	100%	16	9.756098
## 17	0.02390	1	ba	100%	17	10.365854
## 18	0.02453	1	ba	100%	18	10.975610
## 19	0.02483	1	ba	100%	19	11.585366
## 20	0.02501	1	ba	100%	20	12.195122
## 21	0.02514	1	ba	100%	21	12.804878
## 22	0.02531	1	ba	100%	22	13.414634
## 23	0.02551	1	ba	100%	23	14.024390
## 24	0.02558	1	ba	100%	24	14.634146
## 25	0.02594	1	ba	100%	25	15.243902
## 26	0.02600	1	ba	100%	26	15.853659
## 27	0.02668	1	ba	100%	27	16.463415
## 28	0.02681	1	ba	100%	28	17.073171
## 29	0.02701	1	ba	100%	29	17.682927
## 30	0.02741	1	ba	100%	30	18.292683
## 31	0.02784	1	ba	100%	31	18.902439
## 32	0.02933	1	ba	100%	32	19.512195
## 33	0.02943	1	ba	100%	33	20.121951
## 34	0.02950	1	ba	100%	34	20.731707
## 35	0.02974	1	ba	100%	35	21.341463
## 36	0.02995	1	ba	100%	36	21.951220
## 37	0.03002	1	ba	100%	37	22.560976
## 38	0.03022	1	ba	100%	38	23.170732
## 39	0.03063	1	ba	100%	39	23.780488
## 40	0.03076	1	ba	100%	40	24.390244
## 41	0.03117	1	ba	100%	41	25.000000
## 42	0.03131	1	ba	100%	42	25.609756
## 43	0.03159	1	ba	100%	43	26.219512
## 44	0.03163	1	ba	100%	44	26.829268
## 45	0.03172	1	ba	100%	45	27.439024
## 46	0.03199	1	ba	100%	46	28.048780
## 47	0.03211	1	ba	100%	47	28.658537
## 48	0.03227	1	ba	100%	48	29.268293
## 49	0.03228	1	ba	100%	49	29.878049
## 50	0.03243	1	ba	100%	50	30.487805
## 51	0.03249	1	ba	100%	51	31.097561
## 52	0.03263	1	ba	100%	52	31.707317
## 53	0.03278	1	ba	100%	53	32.317073
## 54	0.03312	1	ba	100%	54	32.926829
## 55	0.03315	1	ba	100%	55	33.536585
## 56	0.03323	1	ba	100%	56	34.146341
## 57	0.03334	1	ba	100%	57	34.756098
## 58	0.03340	1	ba	100%	58	35.365854
## 59	0.03373	1	ba	100%	59	35.975610
## 60	0.03420	1	ba	100%	60	36.585366
## 61	0.03424	1	ba	100%	61	37.195122
## 62	0.03432	1	ba	100%	62	37.804878

## 63	0.03492	1	ba	100%	63	38.414634
## 64	0.03499	2	ba	100%	65	39.634146
## 65	0.03501	1	ba	100%	66	40.243902
## 66	0.03537	1	ba	100%	67	40.853659
## 67	0.03538	1	ba	100%	68	41.463415
## 68	0.03555	1	ba	100%	69	42.073171
## 69	0.03585	1	ba	100%	70	42.682927
## 70	0.03596	1	ba	100%	71	43.292683
## 71	0.03651	1	ba	100%	72	43.902439
## 72	0.03661	1	ba	100%	73	44.512195
## 73	0.03720	1	ba	100%	74	45.121951
## 74	0.03723	1	ba	100%	75	45.731707
## 75	0.03725	1	ba	100%	76	46.341463
## 76	0.03814	2	ba	100%	78	47.560976
## 77	0.03844	1	ba	100%	79	48.170732
## 78	0.03845	1	ba	100%	80	48.780488
## 79	0.03869	1	ba	100%	81	49.390244
## 80	0.03949	1	ba	100%	82	50.000000
## 81	0.03953	1	ba	100%	83	50.609756
## 82	0.03965	1	ba	100%	84	51.219512
## 83	0.03969	1	ba	100%	85	51.829268
## 84	0.03971	1	ba	100%	86	52.439024
## 85	0.03973	1	ba	100%	87	53.048780
## 86	0.03983	2	ba	100%	89	54.268293
## 87	0.04006	1	ba	100%	90	54.878049
## 88	0.04025	1	ba	100%	91	55.487805
## 89	0.04043	1	ba	100%	92	56.097561
## 90	0.04049	1	ba	100%	93	56.707317
## 91	0.04052	1	ba	100%	94	57.317073
## 92	0.04074	1	ba	100%	95	57.926829
## 93	0.04090	1	ba	100%	96	58.536585
## 94	0.04099	1	ba	100%	97	59.146341
## 95	0.04126	1	ba	100%	98	59.756098
## 96	0.04129	1	ba	100%	99	60.365854
## 97	0.04144	1	ba	100%	100	60.975610
## 98	0.04167	1	ba	100%	101	61.585366
## 99	0.04231	1	ba	100%	102	62.195122
## 100	0.04245	1	ba	100%	103	62.804878
## 101	0.04285	1	ba	100%	104	63.414634
## 102	0.04326	1	ba	100%	105	64.024390
## 103	0.04351	2	ba	100%	107	65.243902
## 104	0.04373	1	ba	100%	108	65.853659
## 105	0.04380	1	ba	100%	109	66.463415
## 106	0.04571	2	ba	100%	111	67.682927
## 107	0.04576	1	ba	100%	112	68.292683
## 108	0.04584	1	ba	100%	113	68.902439
## 109	0.04637	1	ba	100%	114	69.512195
## 110	0.04661	1	ba	100%	115	70.121951
## 111	0.04670	1	ba	100%	116	70.731707
## 112	0.04713	1	ba	100%	117	71.341463
## 113	0.04751	1	ba	100%	118	71.951220
## 114	0.04779	1	ba	100%	119	72.560976
## 115	0.04809	1	ba	100%	120	73.170732
## 116	0.04823	1	ba	100%	121	73.780488

## 117	0.04829	1	ba	100%	122	74.390244
## 118	0.04836	1	ba	100%	123	75.000000
## 119	0.04918	1	ba	100%	124	75.609756
## 120	0.04936	1	ba	100%	125	76.219512
## 121	0.04964	1	ba	100%	126	76.829268
## 122	0.04998	1	ba	100%	127	77.439024
## 123	0.05054	1	ba	100%	128	78.048780
## 124	0.05062	1	ba	100%	129	78.658537
## 125	0.05125	1	ba	100%	130	79.268293
## 126	0.05147	1	ba	100%	131	79.878049
## 127	0.05247	1	ba	100%	132	80.487805
## 128	0.05255	1	ba	100%	133	81.097561
## 129	0.05322	1	ba	100%	134	81.707317
## 130	0.05335	1	ba	100%	135	82.317073
## 131	0.05369	1	ba	100%	136	82.926829
## 132	0.05373	1	ba	100%	137	83.536585
## 133	0.05445	1	ba	100%	138	84.146341
## 134	0.05582	1	ba	100%	139	84.756098
## 135	0.05661	1	ba	100%	140	85.365854
## 136	0.05732	1	ba	100%	141	85.975610
## 137	0.05740	1	ba	100%	142	86.585366
## 138	0.05792	1	ba	100%	143	87.195122
## 139	0.05884	1	ba	100%	144	87.804878
## 140	0.06049	1	ba	100%	145	88.414634
## 141	0.06123	1	ba	100%	146	89.024390
## 142	0.06237	1	ba	100%	147	89.634146
## 143	0.07122	1	ba	100%	148	90.243902
## 144	0.07932	1	ba	100%	149	90.853659
## 145	0.08033	1	ba	100%	150	91.463415
## 146	0.08270	1	ba	100%	151	92.073171
## 147	0.08536	1	ba	100%	152	92.682927
## 148	0.09070	1	ba	100%	153	93.292683
## 149	0.09600	1	ba	100%	154	93.902439
## 150	0.10695	1	ba	100%	155	94.512195
## 151	0.11361	1	ba	100%	156	95.121951
## 152	0.11532	1	ba	100%	157	95.731707
## 153	0.11549	1	ba	100%	158	96.341463
## 154	0.11791	1	ba	100%	159	96.951220
## 155	0.12036	1	ba	100%	160	97.560976
## 156	0.12039	1	ba	100%	161	98.170732
## 157	0.12113	1	ba	100%	162	98.780488
## 158	0.12284	1	ba	100%	163	99.390244
## 159	0.13870	1	ba	100%	164	100.000000
## 161	0.00905	1	ba	95%	2	1.219512
## 162	0.00916	1	ba	95%	3	1.829268
## 163	0.00956	1	ba	95%	4	2.439024
## 164	0.01157	1	ba	95%	5	3.048780
## 165	0.01239	1	ba	95%	6	3.658537
## 166	0.01284	1	ba	95%	7	4.268293
## 167	0.01298	1	ba	95%	8	4.878049
## 168	0.01312	1	ba	95%	9	5.487805
## 169	0.01336	1	ba	95%	10	6.097561
## 170	0.01344	1	ba	95%	11	6.707317
## 171	0.01489	1	ba	95%	12	7.317073

##	172	0.01494	1	ba	95%	13	7.926829
##	173	0.01504	1	ba	95%	14	8.536585
##	174	0.01507	1	ba	95%	15	9.146341
##	175	0.01523	1	ba	95%	16	9.756098
##	176	0.01531	1	ba	95%	17	10.365854
##	177	0.01559	1	ba	95%	18	10.975610
##	178	0.01568	1	ba	95%	19	11.585366
##	179	0.01573	1	ba	95%	20	12.195122
##	180	0.01584	1	ba	95%	21	12.804878
##	181	0.01635	1	ba	95%	22	13.414634
##	182	0.01657	1	ba	95%	23	14.024390
##	183	0.01659	1	ba	95%	24	14.634146
##	184	0.01673	1	ba	95%	25	15.243902
##	185	0.01689	1	ba	95%	26	15.853659
##	186	0.01691	1	ba	95%	27	16.463415
##	187	0.01706	1	ba	95%	28	17.073171
##	188	0.01719	1	ba	95%	29	17.682927
##	189	0.01738	1	ba	95%	30	18.292683
##	190	0.01766	1	ba	95%	31	18.902439
##	191	0.01771	1	ba	95%	32	19.512195
##	192	0.01853	1	ba	95%	33	20.121951
##	193	0.01859	1	ba	95%	34	20.731707
##	194	0.01868	1	ba	95%	35	21.341463
##	195	0.01914	1	ba	95%	36	21.951220
##	196	0.01916	1	ba	95%	37	22.560976
##	197	0.01951	1	ba	95%	38	23.170732
##	198	0.01955	1	ba	95%	39	23.780488
##	199	0.01960	2	ba	95%	41	25.000000
##	200	0.01969	1	ba	95%	42	25.609756
##	201	0.01972	1	ba	95%	43	26.219512
##	202	0.01974	1	ba	95%	44	26.829268
##	203	0.01975	1	ba	95%	45	27.439024
##	204	0.01980	1	ba	95%	46	28.048780
##	205	0.01999	1	ba	95%	47	28.658537
##	206	0.02005	1	ba	95%	48	29.268293
##	207	0.02030	1	ba	95%	49	29.878049
##	208	0.02056	1	ba	95%	50	30.487805
##	209	0.02060	1	ba	95%	51	31.097561
##	210	0.02077	1	ba	95%	52	31.707317
##	211	0.02079	1	ba	95%	53	32.317073
##	212	0.02112	1	ba	95%	54	32.926829
##	213	0.02118	1	ba	95%	55	33.536585
##	214	0.02138	1	ba	95%	56	34.146341
##	215	0.02156	1	ba	95%	57	34.756098
##	216	0.02159	1	ba	95%	58	35.365854
##	217	0.02171	1	ba	95%	59	35.975610
##	218	0.02203	1	ba	95%	60	36.585366
##	219	0.02206	1	ba	95%	61	37.195122
##	220	0.02216	1	ba	95%	62	37.804878
##	221	0.02226	1	ba	95%	63	38.414634
##	222	0.02235	1	ba	95%	64	39.024390
##	223	0.02238	1	ba	95%	65	39.634146
##	224	0.02260	1	ba	95%	66	40.243902
##	225	0.02262	1	ba	95%	67	40.853659

## 226	0.02264	1	ba	95%	68	41.463415
## 227	0.02284	1	ba	95%	69	42.073171
## 228	0.02288	1	ba	95%	70	42.682927
## 229	0.02292	1	ba	95%	71	43.292683
## 230	0.02310	1	ba	95%	72	43.902439
## 231	0.02319	1	ba	95%	73	44.512195
## 232	0.02325	1	ba	95%	74	45.121951
## 233	0.02329	1	ba	95%	75	45.731707
## 234	0.02348	1	ba	95%	76	46.341463
## 235	0.02363	1	ba	95%	77	46.951220
## 236	0.02367	1	ba	95%	78	47.560976
## 237	0.02380	1	ba	95%	79	48.170732
## 238	0.02406	1	ba	95%	80	48.780488
## 239	0.02450	1	ba	95%	81	49.390244
## 240	0.02454	1	ba	95%	82	50.000000
## 241	0.02504	1	ba	95%	83	50.609756
## 242	0.02532	1	ba	95%	84	51.219512
## 243	0.02589	1	ba	95%	85	51.829268
## 244	0.02593	1	ba	95%	86	52.439024
## 245	0.02603	1	ba	95%	87	53.048780
## 246	0.02614	1	ba	95%	88	53.658537
## 247	0.02626	1	ba	95%	89	54.268293
## 248	0.02649	1	ba	95%	90	54.878049
## 249	0.02659	1	ba	95%	91	55.487805
## 250	0.02663	1	ba	95%	92	56.097561
## 251	0.02679	1	ba	95%	93	56.707317
## 252	0.02712	2	ba	95%	95	57.926829
## 253	0.02721	1	ba	95%	96	58.536585
## 254	0.02725	1	ba	95%	97	59.146341
## 255	0.02733	1	ba	95%	98	59.756098
## 256	0.02746	1	ba	95%	99	60.365854
## 257	0.02755	1	ba	95%	100	60.975610
## 258	0.02766	1	ba	95%	101	61.585366
## 259	0.02767	2	ba	95%	103	62.804878
## 260	0.02768	1	ba	95%	104	63.414634
## 261	0.02783	1	ba	95%	105	64.024390
## 262	0.02801	1	ba	95%	106	64.634146
## 263	0.02808	1	ba	95%	107	65.243902
## 264	0.02809	1	ba	95%	108	65.853659
## 265	0.02836	1	ba	95%	109	66.463415
## 266	0.02837	1	ba	95%	110	67.073171
## 267	0.02844	1	ba	95%	111	67.682927
## 268	0.02870	1	ba	95%	112	68.292683
## 269	0.02877	1	ba	95%	113	68.902439
## 270	0.02887	1	ba	95%	114	69.512195
## 271	0.02916	1	ba	95%	115	70.121951
## 272	0.02938	1	ba	95%	116	70.731707
## 273	0.02939	1	ba	95%	117	71.341463
## 274	0.02946	1	ba	95%	118	71.951220
## 275	0.03038	1	ba	95%	119	72.560976
## 276	0.03077	1	ba	95%	120	73.170732
## 277	0.03081	1	ba	95%	121	73.780488
## 278	0.03115	1	ba	95%	122	74.390244
## 279	0.03121	1	ba	95%	123	75.000000

```

## 280 0.03129    1   ba  95%    124 75.609756
## 281 0.03144    1   ba  95%    125 76.219512
## 282 0.03169    1   ba  95%    126 76.829268
## 283 0.03206    1   ba  95%    127 77.439024
## 284 0.03211    1   ba  95%    128 78.048780
## 285 0.03259    1   ba  95%    129 78.658537
## 286 0.03276    1   ba  95%    130 79.268293
## 287 0.03294    1   ba  95%    131 79.878049
## 288 0.03345    1   ba  95%    132 80.487805
## 289 0.03380    1   ba  95%    133 81.097561
## 290 0.03490    1   ba  95%    134 81.707317
## 291 0.03568    1   ba  95%    135 82.317073
## 292 0.03589    1   ba  95%    136 82.926829
## 293 0.03590    2   ba  95%    138 84.146341
## 294 0.03602    1   ba  95%    139 84.756098
## 295 0.03625    1   ba  95%    140 85.365854
## 296 0.03630    1   ba  95%    141 85.975610
## 297 0.03798    1   ba  95%    142 86.585366
## 298 0.03869    1   ba  95%    143 87.195122
## 299 0.04046    1   ba  95%    144 87.804878
## 300 0.04049    1   ba  95%    145 88.414634
## 301 0.04083    1   ba  95%    146 89.024390
## 302 0.04167    1   ba  95%    147 89.634146
## 303 0.04246    1   ba  95%    148 90.243902
## 304 0.04355    1   ba  95%    149 90.853659
## 305 0.04382    2   ba  95%    151 92.073171
## 306 0.04511    1   ba  95%    152 92.682927
## 307 0.04727    1   ba  95%    153 93.292683
## 308 0.04838    1   ba  95%    154 93.902439
## 309 0.05029    1   ba  95%    155 94.512195
## 310 0.05044    1   ba  95%    156 95.121951
## 311 0.05258    1   ba  95%    157 95.731707
## 312 0.05520    1   ba  95%    158 96.341463
## 313 0.05736    1   ba  95%    159 96.951220
## 314 0.06137    1   ba  95%    160 97.560976
## 315 0.06878    1   ba  95%    161 98.170732
## 316 0.07455    1   ba  95%    162 98.780488
## 317 0.08211    1   ba  95%    163 99.390244
## 318 0.10294    1   ba  95%    164 100.000000

max_95 = round(max(temp_df[temp_df$status == "95%",$coverage], digits=3)
max_100 = round(max(temp_df[temp_df$status == "100%",$coverage], digits=3)
g <- ggplot(temp_df, aes(coverage, percent, color=status)) + geom_point() +
  scale_x_continuous(breaks = seq(0.01, max(temp_df$coverage)+0.01, by = 0.03)) +
  geom_vline(xintercept = max_95, linetype="dashed") +
  annotate("text", x = max_95 - 0.01, y = 0.05,
           label = paste0(as.character(max_95), "x"), size=4) +
  geom_vline(xintercept = max(temp_df[temp_df$status == '100%',$coverage], linetype="dashed") +
  annotate("text", x = max_100 - 0.01, y = 0.05,
           label = paste0(as.character(max_100), "x"), size=4)

p1 <- format_plot(g, xlab="Ba31 Coverage", ylab="Cumulative Percentage", title="Threshold")
remove(temp_df)

```

```
temp_df <- gray_area_group_summary[gray_area_group_summary$group == 'bcg',]
print(temp_df[temp_df$percent >= 0.95,])
```

##	coverage	count	group	status	cumsum	percent
## 320	0.01665	1	bcg	100%	2	1.219512
## 321	0.01854	1	bcg	100%	3	1.829268
## 322	0.01910	1	bcg	100%	4	2.439024
## 323	0.02009	1	bcg	100%	5	3.048780
## 324	0.02041	1	bcg	100%	6	3.658537
## 325	0.02058	1	bcg	100%	7	4.268293
## 326	0.02116	1	bcg	100%	8	4.878049
## 327	0.02240	1	bcg	100%	9	5.487805
## 328	0.02298	1	bcg	100%	10	6.097561
## 329	0.02315	1	bcg	100%	11	6.707317
## 330	0.02317	1	bcg	100%	12	7.317073
## 331	0.02387	1	bcg	100%	13	7.926829
## 332	0.02399	1	bcg	100%	14	8.536585
## 333	0.02444	1	bcg	100%	15	9.146341
## 334	0.02449	1	bcg	100%	16	9.756098
## 335	0.02545	1	bcg	100%	17	10.365854
## 336	0.02569	1	bcg	100%	18	10.975610
## 337	0.02592	1	bcg	100%	19	11.585366
## 338	0.02645	1	bcg	100%	20	12.195122
## 339	0.02810	1	bcg	100%	21	12.804878
## 340	0.02861	1	bcg	100%	22	13.414634
## 341	0.02871	1	bcg	100%	23	14.024390
## 342	0.02907	1	bcg	100%	24	14.634146
## 343	0.03012	1	bcg	100%	25	15.243902
## 344	0.03058	1	bcg	100%	26	15.853659
## 345	0.03116	1	bcg	100%	27	16.463415
## 346	0.03118	1	bcg	100%	28	17.073171
## 347	0.03155	1	bcg	100%	29	17.682927
## 348	0.03244	1	bcg	100%	30	18.292683
## 349	0.03288	1	bcg	100%	31	18.902439
## 350	0.03291	1	bcg	100%	32	19.512195
## 351	0.03304	1	bcg	100%	33	20.121951
## 352	0.03326	1	bcg	100%	34	20.731707
## 353	0.03337	1	bcg	100%	35	21.341463
## 354	0.03339	1	bcg	100%	36	21.951220
## 355	0.03385	1	bcg	100%	37	22.560976
## 356	0.03394	1	bcg	100%	38	23.170732
## 357	0.03419	1	bcg	100%	39	23.780488
## 358	0.03425	2	bcg	100%	41	25.000000
## 359	0.03447	1	bcg	100%	42	25.609756
## 360	0.03494	1	bcg	100%	43	26.219512
## 361	0.03495	1	bcg	100%	44	26.829268
## 362	0.03521	1	bcg	100%	45	27.439024
## 363	0.03533	1	bcg	100%	46	28.048780
## 364	0.03570	1	bcg	100%	47	28.658537
## 365	0.03571	1	bcg	100%	48	29.268293
## 366	0.03574	1	bcg	100%	49	29.878049
## 367	0.03579	1	bcg	100%	50	30.487805
## 368	0.03588	1	bcg	100%	51	31.097561

## 369	0.03625	1	bcg	100%	52	31.707317
## 370	0.03654	1	bcg	100%	53	32.317073
## 371	0.03740	1	bcg	100%	54	32.926829
## 372	0.03783	1	bcg	100%	55	33.536585
## 373	0.03792	1	bcg	100%	56	34.146341
## 374	0.03796	1	bcg	100%	57	34.756098
## 375	0.03823	2	bcg	100%	59	35.975610
## 376	0.03829	1	bcg	100%	60	36.585366
## 377	0.03830	1	bcg	100%	61	37.195122
## 378	0.03861	1	bcg	100%	62	37.804878
## 379	0.03919	1	bcg	100%	63	38.414634
## 380	0.03926	1	bcg	100%	64	39.024390
## 381	0.03961	1	bcg	100%	65	39.634146
## 382	0.03982	1	bcg	100%	66	40.243902
## 383	0.04071	1	bcg	100%	67	40.853659
## 384	0.04084	1	bcg	100%	68	41.463415
## 385	0.04124	1	bcg	100%	69	42.073171
## 386	0.04145	1	bcg	100%	70	42.682927
## 387	0.04151	1	bcg	100%	71	43.292683
## 388	0.04185	1	bcg	100%	72	43.902439
## 389	0.04246	1	bcg	100%	73	44.512195
## 390	0.04264	1	bcg	100%	74	45.121951
## 391	0.04291	1	bcg	100%	75	45.731707
## 392	0.04295	1	bcg	100%	76	46.341463
## 393	0.04345	1	bcg	100%	77	46.951220
## 394	0.04388	1	bcg	100%	78	47.560976
## 395	0.04394	1	bcg	100%	79	48.170732
## 396	0.04403	1	bcg	100%	80	48.780488
## 397	0.04408	1	bcg	100%	81	49.390244
## 398	0.04463	1	bcg	100%	82	50.000000
## 399	0.04473	1	bcg	100%	83	50.609756
## 400	0.04482	2	bcg	100%	85	51.829268
## 401	0.04489	1	bcg	100%	86	52.439024
## 402	0.04494	1	bcg	100%	87	53.048780
## 403	0.04521	1	bcg	100%	88	53.658537
## 404	0.04545	1	bcg	100%	89	54.268293
## 405	0.04562	1	bcg	100%	90	54.878049
## 406	0.04607	1	bcg	100%	91	55.487805
## 407	0.04624	1	bcg	100%	92	56.097561
## 408	0.04688	1	bcg	100%	93	56.707317
## 409	0.04714	1	bcg	100%	94	57.317073
## 410	0.04758	1	bcg	100%	95	57.926829
## 411	0.04798	1	bcg	100%	96	58.536585
## 412	0.04829	1	bcg	100%	97	59.146341
## 413	0.04835	1	bcg	100%	98	59.756098
## 414	0.04852	1	bcg	100%	99	60.365854
## 415	0.04867	1	bcg	100%	100	60.975610
## 416	0.04871	1	bcg	100%	101	61.585366
## 417	0.04889	1	bcg	100%	102	62.195122
## 418	0.04892	1	bcg	100%	103	62.804878
## 419	0.04896	1	bcg	100%	104	63.414634
## 420	0.04996	2	bcg	100%	106	64.634146
## 421	0.05071	1	bcg	100%	107	65.243902
## 422	0.05077	1	bcg	100%	108	65.853659

## 423	0.05085	1	bcg	100%	109	66.463415
## 424	0.05116	1	bcg	100%	110	67.073171
## 425	0.05148	1	bcg	100%	111	67.682927
## 426	0.05172	1	bcg	100%	112	68.292683
## 427	0.05182	1	bcg	100%	113	68.902439
## 428	0.05208	1	bcg	100%	114	69.512195
## 429	0.05221	1	bcg	100%	115	70.121951
## 430	0.05226	1	bcg	100%	116	70.731707
## 431	0.05250	1	bcg	100%	117	71.341463
## 432	0.05331	1	bcg	100%	118	71.951220
## 433	0.05346	1	bcg	100%	119	72.560976
## 434	0.05405	2	bcg	100%	121	73.780488
## 435	0.05464	1	bcg	100%	122	74.390244
## 436	0.05518	1	bcg	100%	123	75.000000
## 437	0.05534	1	bcg	100%	124	75.609756
## 438	0.05588	1	bcg	100%	125	76.219512
## 439	0.05609	1	bcg	100%	126	76.829268
## 440	0.05710	1	bcg	100%	127	77.439024
## 441	0.05752	1	bcg	100%	128	78.048780
## 442	0.05760	1	bcg	100%	129	78.658537
## 443	0.05820	1	bcg	100%	130	79.268293
## 444	0.05829	1	bcg	100%	131	79.878049
## 445	0.05865	1	bcg	100%	132	80.487805
## 446	0.05881	1	bcg	100%	133	81.097561
## 447	0.05951	1	bcg	100%	134	81.707317
## 448	0.06074	1	bcg	100%	135	82.317073
## 449	0.06138	1	bcg	100%	136	82.926829
## 450	0.06220	1	bcg	100%	137	83.536585
## 451	0.06227	1	bcg	100%	138	84.146341
## 452	0.06239	1	bcg	100%	139	84.756098
## 453	0.06317	1	bcg	100%	140	85.365854
## 454	0.06347	1	bcg	100%	141	85.975610
## 455	0.06377	1	bcg	100%	142	86.585366
## 456	0.06577	1	bcg	100%	143	87.195122
## 457	0.06685	1	bcg	100%	144	87.804878
## 458	0.06724	1	bcg	100%	145	88.414634
## 459	0.06888	1	bcg	100%	146	89.024390
## 460	0.07258	1	bcg	100%	147	89.634146
## 461	0.07324	1	bcg	100%	148	90.243902
## 462	0.09215	1	bcg	100%	149	90.853659
## 463	0.09456	1	bcg	100%	150	91.463415
## 464	0.09727	1	bcg	100%	151	92.073171
## 465	0.09767	1	bcg	100%	152	92.682927
## 466	0.10626	1	bcg	100%	153	93.292683
## 467	0.11041	1	bcg	100%	154	93.902439
## 468	0.11820	1	bcg	100%	155	94.512195
## 469	0.13167	1	bcg	100%	156	95.121951
## 470	0.13231	1	bcg	100%	157	95.731707
## 471	0.13375	1	bcg	100%	158	96.341463
## 472	0.13451	1	bcg	100%	159	96.951220
## 473	0.13605	1	bcg	100%	160	97.560976
## 474	0.14008	1	bcg	100%	161	98.170732
## 475	0.14128	1	bcg	100%	162	98.780488
## 476	0.14351	1	bcg	100%	163	99.390244

## 477	0.16458	1	bcg	100%	164	100.000000
## 479	0.00827	1	bcg	95%	2	1.219512
## 480	0.00938	1	bcg	95%	3	1.829268
## 481	0.01024	1	bcg	95%	4	2.439024
## 482	0.01233	1	bcg	95%	5	3.048780
## 483	0.01276	1	bcg	95%	6	3.658537
## 484	0.01349	1	bcg	95%	7	4.268293
## 485	0.01352	1	bcg	95%	8	4.878049
## 486	0.01427	1	bcg	95%	9	5.487805
## 487	0.01455	1	bcg	95%	10	6.097561
## 488	0.01560	1	bcg	95%	11	6.707317
## 489	0.01579	1	bcg	95%	12	7.317073
## 490	0.01596	1	bcg	95%	13	7.926829
## 491	0.01670	1	bcg	95%	14	8.536585
## 492	0.01726	1	bcg	95%	15	9.146341
## 493	0.01745	1	bcg	95%	16	9.756098
## 494	0.01788	1	bcg	95%	17	10.365854
## 495	0.01860	1	bcg	95%	18	10.975610
## 496	0.01863	1	bcg	95%	19	11.585366
## 497	0.01865	1	bcg	95%	20	12.195122
## 498	0.01899	1	bcg	95%	21	12.804878
## 499	0.01920	1	bcg	95%	22	13.414634
## 500	0.01938	1	bcg	95%	23	14.024390
## 501	0.01949	1	bcg	95%	24	14.634146
## 502	0.01977	1	bcg	95%	25	15.243902
## 503	0.01984	1	bcg	95%	26	15.853659
## 504	0.01998	1	bcg	95%	27	16.463415
## 505	0.02007	1	bcg	95%	28	17.073171
## 506	0.02019	1	bcg	95%	29	17.682927
## 507	0.02020	1	bcg	95%	30	18.292683
## 508	0.02045	1	bcg	95%	31	18.902439
## 509	0.02047	1	bcg	95%	32	19.512195
## 510	0.02076	1	bcg	95%	33	20.121951
## 511	0.02085	1	bcg	95%	34	20.731707
## 512	0.02086	1	bcg	95%	35	21.341463
## 513	0.02095	1	bcg	95%	36	21.951220
## 514	0.02099	1	bcg	95%	37	22.560976
## 515	0.02147	1	bcg	95%	38	23.170732
## 516	0.02171	1	bcg	95%	39	23.780488
## 517	0.02214	1	bcg	95%	40	24.390244
## 518	0.02220	1	bcg	95%	41	25.000000
## 519	0.02240	1	bcg	95%	42	25.609756
## 520	0.02254	1	bcg	95%	43	26.219512
## 521	0.02261	1	bcg	95%	44	26.829268
## 522	0.02275	1	bcg	95%	45	27.439024
## 523	0.02279	1	bcg	95%	46	28.048780
## 524	0.02283	1	bcg	95%	47	28.658537
## 525	0.02289	1	bcg	95%	48	29.268293
## 526	0.02292	1	bcg	95%	49	29.878049
## 527	0.02321	1	bcg	95%	50	30.487805
## 528	0.02330	1	bcg	95%	51	31.097561
## 529	0.02338	1	bcg	95%	52	31.707317
## 530	0.02343	1	bcg	95%	53	32.317073
## 531	0.02354	1	bcg	95%	54	32.926829

## 532	0.02364	1	bcg	95%	55	33.536585
## 533	0.02373	1	bcg	95%	56	34.146341
## 534	0.02378	1	bcg	95%	57	34.756098
## 535	0.02385	1	bcg	95%	58	35.365854
## 536	0.02387	1	bcg	95%	59	35.975610
## 537	0.02417	1	bcg	95%	60	36.585366
## 538	0.02447	1	bcg	95%	61	37.195122
## 539	0.02456	1	bcg	95%	62	37.804878
## 540	0.02482	1	bcg	95%	63	38.414634
## 541	0.02486	1	bcg	95%	64	39.024390
## 542	0.02488	2	bcd	95%	66	40.243902
## 543	0.02498	1	bcd	95%	67	40.853659
## 544	0.02500	1	bcd	95%	68	41.463415
## 545	0.02503	1	bcd	95%	69	42.073171
## 546	0.02504	1	bcd	95%	70	42.682927
## 547	0.02514	1	bcd	95%	71	43.292683
## 548	0.02515	1	bcd	95%	72	43.902439
## 549	0.02560	1	bcd	95%	73	44.512195
## 550	0.02562	2	bcd	95%	75	45.731707
## 551	0.02573	1	bcd	95%	76	46.341463
## 552	0.02576	1	bcd	95%	77	46.951220
## 553	0.02603	1	bcd	95%	78	47.560976
## 554	0.02612	1	bcd	95%	79	48.170732
## 555	0.02619	1	bcd	95%	80	48.780488
## 556	0.02621	1	bcd	95%	81	49.390244
## 557	0.02638	1	bcd	95%	82	50.000000
## 558	0.02686	1	bcd	95%	83	50.609756
## 559	0.02794	1	bcd	95%	84	51.219512
## 560	0.02803	1	bcd	95%	85	51.829268
## 561	0.02869	1	bcd	95%	86	52.439024
## 562	0.02929	1	bcd	95%	87	53.048780
## 563	0.02940	1	bcd	95%	88	53.658537
## 564	0.02965	1	bcd	95%	89	54.268293
## 565	0.02992	1	bcd	95%	90	54.878049
## 566	0.02996	1	bcd	95%	91	55.487805
## 567	0.03001	1	bcd	95%	92	56.097561
## 568	0.03048	1	bcd	95%	93	56.707317
## 569	0.03053	1	bcd	95%	94	57.317073
## 570	0.03060	1	bcd	95%	95	57.926829
## 571	0.03086	1	bcd	95%	96	58.536585
## 572	0.03094	1	bcd	95%	97	59.146341
## 573	0.03117	1	bcd	95%	98	59.756098
## 574	0.03119	1	bcd	95%	99	60.365854
## 575	0.03123	1	bcd	95%	100	60.975610
## 576	0.03151	2	bcd	95%	102	62.195122
## 577	0.03156	1	bcd	95%	103	62.804878
## 578	0.03158	1	bcd	95%	104	63.414634
## 579	0.03160	1	bcd	95%	105	64.024390
## 580	0.03161	1	bcd	95%	106	64.634146
## 581	0.03193	1	bcd	95%	107	65.243902
## 582	0.03218	1	bcd	95%	108	65.853659
## 583	0.03231	1	bcd	95%	109	66.463415
## 584	0.03247	1	bcd	95%	110	67.073171
## 585	0.03263	1	bcd	95%	111	67.682927

```

## 586 0.03286    1 bcg 95%   112 68.292683
## 587 0.03295    1 bcg 95%   113 68.902439
## 588 0.03307    1 bcg 95%   114 69.512195
## 589 0.03430    1 bcg 95%   115 70.121951
## 590 0.03454    1 bcg 95%   116 70.731707
## 591 0.03455    1 bcg 95%   117 71.341463
## 592 0.03458    1 bcg 95%   118 71.951220
## 593 0.03533    1 bcg 95%   119 72.560976
## 594 0.03549    1 bcg 95%   120 73.170732
## 595 0.03669    1 bcg 95%   121 73.780488
## 596 0.03692    1 bcg 95%   122 74.390244
## 597 0.03705    1 bcg 95%   123 75.000000
## 598 0.03711    1 bcg 95%   124 75.609756
## 599 0.03717    1 bcg 95%   125 76.219512
## 600 0.03741    2 bcg 95%   127 77.439024
## 601 0.03813    1 bcg 95%   128 78.048780
## 602 0.03823    1 bcg 95%   129 78.658537
## 603 0.03827    1 bcg 95%   130 79.268293
## 604 0.03891    1 bcg 95%   131 79.878049
## 605 0.03894    1 bcg 95%   132 80.487805
## 606 0.03908    1 bcg 95%   133 81.097561
## 607 0.03911    1 bcg 95%   134 81.707317
## 608 0.03936    1 bcg 95%   135 82.317073
## 609 0.03988    1 bcg 95%   136 82.926829
## 610 0.04054    1 bcg 95%   137 83.536585
## 611 0.04058    2 bcg 95%   139 84.756098
## 612 0.04269    1 bcg 95%   140 85.365854
## 613 0.04398    1 bcg 95%   141 85.975610
## 614 0.04402    1 bcg 95%   142 86.585366
## 615 0.04403    1 bcg 95%   143 87.195122
## 616 0.04448    1 bcg 95%   144 87.804878
## 617 0.04581    1 bcg 95%   145 88.414634
## 618 0.04852    1 bcg 95%   146 89.024390
## 619 0.04962    1 bcg 95%   147 89.634146
## 620 0.04982    2 bcg 95%   149 90.853659
## 621 0.05057    1 bcg 95%   150 91.463415
## 622 0.05169    1 bcg 95%   151 92.073171
## 623 0.05209    1 bcg 95%   152 92.682927
## 624 0.05384    1 bcg 95%   153 93.292683
## 625 0.05544    1 bcg 95%   154 93.902439
## 626 0.05560    1 bcg 95%   155 94.512195
## 627 0.06060    1 bcg 95%   156 95.121951
## 628 0.06089    1 bcg 95%   157 95.731707
## 629 0.06674    1 bcg 95%   158 96.341463
## 630 0.06706    1 bcg 95%   159 96.951220
## 631 0.06936    1 bcg 95%   160 97.560976
## 632 0.08222    1 bcg 95%   161 98.170732
## 633 0.08519    1 bcg 95%   162 98.780488
## 634 0.09806    1 bcg 95%   163 99.390244
## 635 0.11180    1 bcg 95%   164 100.000000

max_95 = round(max(temp_df[temp_df$status == "95%"]$coverage), digits=3)
max_100 = round(max(temp_df[temp_df$status == "100%"]$coverage), digits=3)
g <- ggplot(temp_df, aes(coverage, percent, color=status)) + geom_point() +

```

```

scale_x_continuous(breaks = seq(0.01, max(temp_df$coverage)+0.01, by = 0.03)) +
geom_vline(xintercept = max_95, linetype="dashed") +
annotate("text", x = max_95 - 0.01, y = 0.05,
label = paste0(as.character(max_95), "x"), size=4) +
geom_vline(xintercept = max(temp_df[temp_df$status == '100%',])$coverage, linetype="dashed") +
annotate("text", x = max_100 - 0.01, y = 0.05,
label = paste0(as.character(max_100), "x"), size=4)

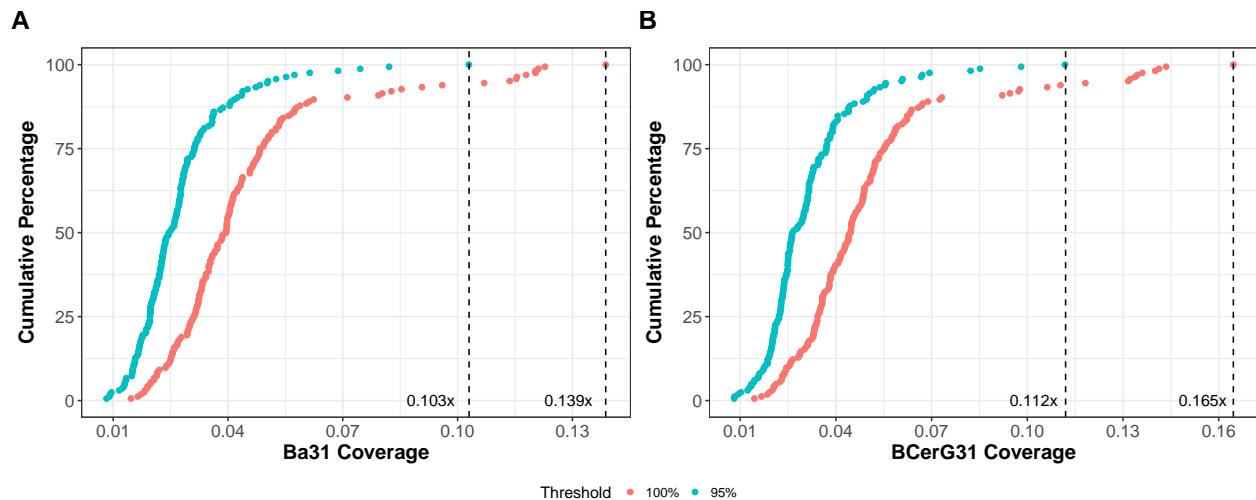
p2 <- format_plot(g, xlab="BCerG31 Coverage", ylab="Cumulative Percentage", title="Threshold")
remove(temp_df)

pdf(paste0(FIGURE, '/figure-02-lef31-lod.pdf'), height=6, width=12, onefile=FALSE)
grid_arrange_shared_lef31(p1, p2)
dev_null <- dev.off()

png(paste0(FIGURE, '/figure-02-lef31-lod.png'), height=600, width=1200)
grid_arrange_shared_lef31(p1, p2)
dev_null <- dev.off()

grid_arrange_shared_lef31(p1, p2)

```



lef31 k-mer Coverage

```

temp_df <- gray_area_group_summary[gray_area_group_summary$group == 'lef',]
print(temp_df[temp_df$percent >= 0.95,])

```

	coverage	count	group	status	cumsum	percent
##	0.07806	1	lef	100%	2	1.219512
##	0.07807	1	lef	100%	3	1.829268
##	0.07815	1	lef	100%	4	2.439024
##	0.07840	1	lef	100%	5	3.048780
##	0.07992	1	lef	100%	6	3.658537
##	0.08024	1	lef	100%	7	4.268293
##	0.08067	1	lef	100%	8	4.878049
##	0.08278	1	lef	100%	9	5.487805

## 645	0.08337	1	lef	100%	10	6.097561
## 646	0.08356	1	lef	100%	11	6.707317
## 647	0.08378	1	lef	100%	12	7.317073
## 648	0.08465	2	lef	100%	14	8.536585
## 649	0.08499	1	lef	100%	15	9.146341
## 650	0.08504	1	lef	100%	16	9.756098
## 651	0.08570	1	lef	100%	17	10.365854
## 652	0.08730	1	lef	100%	18	10.975610
## 653	0.08780	1	lef	100%	19	11.585366
## 654	0.08825	1	lef	100%	20	12.195122
## 655	0.08899	1	lef	100%	21	12.804878
## 656	0.08968	1	lef	100%	22	13.414634
## 657	0.09023	1	lef	100%	23	14.024390
## 658	0.09100	1	lef	100%	24	14.634146
## 659	0.09155	1	lef	100%	25	15.243902
## 660	0.09316	1	lef	100%	26	15.853659
## 661	0.09464	1	lef	100%	27	16.463415
## 662	0.09614	2	lef	100%	29	17.682927
## 663	0.09687	1	lef	100%	30	18.292683
## 664	0.09713	1	lef	100%	31	18.902439
## 665	0.09758	1	lef	100%	32	19.512195
## 666	0.09848	1	lef	100%	33	20.121951
## 667	0.09903	1	lef	100%	34	20.731707
## 668	0.10039	1	lef	100%	35	21.341463
## 669	0.10162	1	lef	100%	36	21.951220
## 670	0.10186	1	lef	100%	37	22.560976
## 671	0.10252	1	lef	100%	38	23.170732
## 672	0.10387	1	lef	100%	39	23.780488
## 673	0.10392	1	lef	100%	40	24.390244
## 674	0.10460	1	lef	100%	41	25.000000
## 675	0.10577	1	lef	100%	42	25.609756
## 676	0.10639	1	lef	100%	43	26.219512
## 677	0.10676	1	lef	100%	44	26.829268
## 678	0.10733	1	lef	100%	45	27.439024
## 679	0.10789	1	lef	100%	46	28.048780
## 680	0.10891	1	lef	100%	47	28.658537
## 681	0.10908	1	lef	100%	48	29.268293
## 682	0.10941	1	lef	100%	49	29.878049
## 683	0.11007	1	lef	100%	50	30.487805
## 684	0.11021	1	lef	100%	51	31.097561
## 685	0.11034	1	lef	100%	52	31.707317
## 686	0.11130	1	lef	100%	53	32.317073
## 687	0.11156	1	lef	100%	54	32.926829
## 688	0.11234	1	lef	100%	55	33.536585
## 689	0.11250	1	lef	100%	56	34.146341
## 690	0.11260	1	lef	100%	57	34.756098
## 691	0.11263	1	lef	100%	58	35.365854
## 692	0.11385	1	lef	100%	59	35.975610
## 693	0.11462	1	lef	100%	60	36.585366
## 694	0.11504	1	lef	100%	61	37.195122
## 695	0.11508	1	lef	100%	62	37.804878
## 696	0.11521	1	lef	100%	63	38.414634
## 697	0.11542	1	lef	100%	64	39.024390
## 698	0.11545	1	lef	100%	65	39.634146

## 699	0.11547	1	lef	100%	66	40.243902
## 700	0.11603	1	lef	100%	67	40.853659
## 701	0.11691	1	lef	100%	68	41.463415
## 702	0.11697	1	lef	100%	69	42.073171
## 703	0.11704	1	lef	100%	70	42.682927
## 704	0.11770	1	lef	100%	71	43.292683
## 705	0.11820	1	lef	100%	72	43.902439
## 706	0.11923	1	lef	100%	73	44.512195
## 707	0.11988	1	lef	100%	74	45.121951
## 708	0.12016	1	lef	100%	75	45.731707
## 709	0.12067	1	lef	100%	76	46.341463
## 710	0.12105	1	lef	100%	77	46.951220
## 711	0.12117	1	lef	100%	78	47.560976
## 712	0.12168	1	lef	100%	79	48.170732
## 713	0.12185	1	lef	100%	80	48.780488
## 714	0.12249	1	lef	100%	81	49.390244
## 715	0.12259	1	lef	100%	82	50.000000
## 716	0.12263	1	lef	100%	83	50.609756
## 717	0.12281	1	lef	100%	84	51.219512
## 718	0.12387	1	lef	100%	85	51.829268
## 719	0.12428	1	lef	100%	86	52.439024
## 720	0.12476	1	lef	100%	87	53.048780
## 721	0.12484	1	lef	100%	88	53.658537
## 722	0.12497	1	lef	100%	89	54.268293
## 723	0.12569	1	lef	100%	90	54.878049
## 724	0.12623	1	lef	100%	91	55.487805
## 725	0.12743	1	lef	100%	92	56.097561
## 726	0.12757	1	lef	100%	93	56.707317
## 727	0.12794	1	lef	100%	94	57.317073
## 728	0.12841	1	lef	100%	95	57.926829
## 729	0.12903	1	lef	100%	96	58.536585
## 730	0.12931	1	lef	100%	97	59.146341
## 731	0.12932	1	lef	100%	98	59.756098
## 732	0.12984	1	lef	100%	99	60.365854
## 733	0.13029	1	lef	100%	100	60.975610
## 734	0.13100	1	lef	100%	101	61.585366
## 735	0.13106	1	lef	100%	102	62.195122
## 736	0.13131	1	lef	100%	103	62.804878
## 737	0.13235	1	lef	100%	104	63.414634
## 738	0.13327	1	lef	100%	105	64.024390
## 739	0.13418	1	lef	100%	106	64.634146
## 740	0.13483	1	lef	100%	107	65.243902
## 741	0.13502	1	lef	100%	108	65.853659
## 742	0.13509	1	lef	100%	109	66.463415
## 743	0.13534	1	lef	100%	110	67.073171
## 744	0.13610	1	lef	100%	111	67.682927
## 745	0.13706	1	lef	100%	112	68.292683
## 746	0.13789	1	lef	100%	113	68.902439
## 747	0.13804	1	lef	100%	114	69.512195
## 748	0.13946	1	lef	100%	115	70.121951
## 749	0.14160	1	lef	100%	116	70.731707
## 750	0.14217	1	lef	100%	117	71.341463
## 751	0.14248	1	lef	100%	118	71.951220
## 752	0.14270	1	lef	100%	119	72.560976

## 753	0.14325	1	lef	100%	120	73.170732
## 754	0.14340	2	lef	100%	122	74.390244
## 755	0.14350	1	lef	100%	123	75.000000
## 756	0.14475	1	lef	100%	124	75.609756
## 757	0.14719	1	lef	100%	125	76.219512
## 758	0.14764	1	lef	100%	126	76.829268
## 759	0.14809	1	lef	100%	127	77.439024
## 760	0.14866	1	lef	100%	128	78.048780
## 761	0.15060	1	lef	100%	129	78.658537
## 762	0.15158	1	lef	100%	130	79.268293
## 763	0.15171	1	lef	100%	131	79.878049
## 764	0.15217	1	lef	100%	132	80.487805
## 765	0.15319	1	lef	100%	133	81.097561
## 766	0.15446	1	lef	100%	134	81.707317
## 767	0.15571	1	lef	100%	135	82.317073
## 768	0.15790	1	lef	100%	136	82.926829
## 769	0.15867	1	lef	100%	137	83.536585
## 770	0.16011	1	lef	100%	138	84.146341
## 771	0.16055	1	lef	100%	139	84.756098
## 772	0.16183	1	lef	100%	140	85.365854
## 773	0.16191	1	lef	100%	141	85.975610
## 774	0.16273	1	lef	100%	142	86.585366
## 775	0.16448	1	lef	100%	143	87.195122
## 776	0.16685	1	lef	100%	144	87.804878
## 777	0.17215	1	lef	100%	145	88.414634
## 778	0.17669	1	lef	100%	146	89.024390
## 779	0.18470	1	lef	100%	147	89.634146
## 780	0.19211	1	lef	100%	148	90.243902
## 781	0.19443	1	lef	100%	149	90.853659
## 782	0.19832	1	lef	100%	150	91.463415
## 783	0.20548	1	lef	100%	151	92.073171
## 784	0.20708	1	lef	100%	152	92.682927
## 785	0.20895	1	lef	100%	153	93.292683
## 786	0.21306	1	lef	100%	154	93.902439
## 787	0.21684	1	lef	100%	155	94.512195
## 788	0.21959	1	lef	100%	156	95.121951
## 789	0.22638	1	lef	100%	157	95.731707
## 790	0.23091	1	lef	100%	158	96.341463
## 791	0.23906	1	lef	100%	159	96.951220
## 792	0.24570	1	lef	100%	160	97.560976
## 793	0.25712	1	lef	100%	161	98.170732
## 794	0.27658	1	lef	100%	162	98.780488
## 795	0.29204	1	lef	100%	163	99.390244
## 796	0.30020	1	lef	100%	164	100.000000
## 798	0.06238	1	lef	95%	2	1.219512
## 799	0.06558	1	lef	95%	3	1.829268
## 800	0.06590	1	lef	95%	4	2.439024
## 801	0.06609	1	lef	95%	5	3.048780
## 802	0.06626	1	lef	95%	6	3.658537
## 803	0.06641	1	lef	95%	7	4.268293
## 804	0.06650	1	lef	95%	8	4.878049
## 805	0.06734	1	lef	95%	9	5.487805
## 806	0.06776	1	lef	95%	10	6.097561
## 807	0.06789	1	lef	95%	11	6.707317

## 808	0.06817	1	lef	95%	12	7.317073
## 809	0.06842	1	lef	95%	13	7.926829
## 810	0.06863	1	lef	95%	14	8.536585
## 811	0.06916	1	lef	95%	15	9.146341
## 812	0.06975	1	lef	95%	16	9.756098
## 813	0.06987	1	lef	95%	17	10.365854
## 814	0.07045	1	lef	95%	18	10.975610
## 815	0.07060	1	lef	95%	19	11.585366
## 816	0.07074	1	lef	95%	20	12.195122
## 817	0.07158	1	lef	95%	21	12.804878
## 818	0.07173	1	lef	95%	22	13.414634
## 819	0.07193	1	lef	95%	23	14.024390
## 820	0.07195	1	lef	95%	24	14.634146
## 821	0.07204	1	lef	95%	25	15.243902
## 822	0.07259	2	lef	95%	27	16.463415
## 823	0.07267	1	lef	95%	28	17.073171
## 824	0.07284	1	lef	95%	29	17.682927
## 825	0.07285	1	lef	95%	30	18.292683
## 826	0.07299	1	lef	95%	31	18.902439
## 827	0.07306	1	lef	95%	32	19.512195
## 828	0.07307	1	lef	95%	33	20.121951
## 829	0.07312	1	lef	95%	34	20.731707
## 830	0.07314	1	lef	95%	35	21.341463
## 831	0.07319	1	lef	95%	36	21.951220
## 832	0.07327	1	lef	95%	37	22.560976
## 833	0.07391	1	lef	95%	38	23.170732
## 834	0.07409	1	lef	95%	39	23.780488
## 835	0.07419	1	lef	95%	40	24.390244
## 836	0.07425	1	lef	95%	41	25.000000
## 837	0.07460	1	lef	95%	42	25.609756
## 838	0.07480	1	lef	95%	43	26.219512
## 839	0.07484	1	lef	95%	44	26.829268
## 840	0.07495	1	lef	95%	45	27.439024
## 841	0.07508	1	lef	95%	46	28.048780
## 842	0.07509	1	lef	95%	47	28.658537
## 843	0.07523	1	lef	95%	48	29.268293
## 844	0.07545	1	lef	95%	49	29.878049
## 845	0.07546	1	lef	95%	50	30.487805
## 846	0.07565	1	lef	95%	51	31.097561
## 847	0.07571	1	lef	95%	52	31.707317
## 848	0.07586	1	lef	95%	53	32.317073
## 849	0.07603	1	lef	95%	54	32.926829
## 850	0.07618	1	lef	95%	55	33.536585
## 851	0.07627	1	lef	95%	56	34.146341
## 852	0.07635	1	lef	95%	57	34.756098
## 853	0.07636	1	lef	95%	58	35.365854
## 854	0.07642	1	lef	95%	59	35.975610
## 855	0.07661	1	lef	95%	60	36.585366
## 856	0.07668	1	lef	95%	61	37.195122
## 857	0.07671	1	lef	95%	62	37.804878
## 858	0.07676	1	lef	95%	63	38.414634
## 859	0.07677	1	lef	95%	64	39.024390
## 860	0.07738	1	lef	95%	65	39.634146
## 861	0.07749	1	lef	95%	66	40.243902

## 862	0.07750	1	lef	95%	67	40.853659
## 863	0.07759	1	lef	95%	68	41.463415
## 864	0.07764	1	lef	95%	69	42.073171
## 865	0.07791	1	lef	95%	70	42.682927
## 866	0.07809	1	lef	95%	71	43.292683
## 867	0.07812	1	lef	95%	72	43.902439
## 868	0.07826	1	lef	95%	73	44.512195
## 869	0.07827	1	lef	95%	74	45.121951
## 870	0.07850	1	lef	95%	75	45.731707
## 871	0.07860	1	lef	95%	76	46.341463
## 872	0.07862	1	lef	95%	77	46.951220
## 873	0.07870	1	lef	95%	78	47.560976
## 874	0.07874	2	lef	95%	80	48.780488
## 875	0.07889	1	lef	95%	81	49.390244
## 876	0.07920	1	lef	95%	82	50.000000
## 877	0.07926	1	lef	95%	83	50.609756
## 878	0.07940	1	lef	95%	84	51.219512
## 879	0.07942	1	lef	95%	85	51.829268
## 880	0.07955	2	lef	95%	87	53.048780
## 881	0.07968	1	lef	95%	88	53.658537
## 882	0.07984	1	lef	95%	89	54.268293
## 883	0.07993	1	lef	95%	90	54.878049
## 884	0.07999	1	lef	95%	91	55.487805
## 885	0.08020	2	lef	95%	93	56.707317
## 886	0.08024	1	lef	95%	94	57.317073
## 887	0.08039	1	lef	95%	95	57.926829
## 888	0.08049	1	lef	95%	96	58.536585
## 889	0.08051	1	lef	95%	97	59.146341
## 890	0.08111	1	lef	95%	98	59.756098
## 891	0.08114	1	lef	95%	99	60.365854
## 892	0.08131	1	lef	95%	100	60.975610
## 893	0.08134	1	lef	95%	101	61.585366
## 894	0.08150	1	lef	95%	102	62.195122
## 895	0.08151	1	lef	95%	103	62.804878
## 896	0.08166	1	lef	95%	104	63.414634
## 897	0.08183	1	lef	95%	105	64.024390
## 898	0.08199	1	lef	95%	106	64.634146
## 899	0.08243	1	lef	95%	107	65.243902
## 900	0.08249	2	lef	95%	109	66.463415
## 901	0.08269	1	lef	95%	110	67.073171
## 902	0.08272	1	lef	95%	111	67.682927
## 903	0.08301	1	lef	95%	112	68.292683
## 904	0.08314	1	lef	95%	113	68.902439
## 905	0.08319	1	lef	95%	114	69.512195
## 906	0.08366	2	lef	95%	116	70.731707
## 907	0.08367	1	lef	95%	117	71.341463
## 908	0.08372	1	lef	95%	118	71.951220
## 909	0.08407	1	lef	95%	119	72.560976
## 910	0.08409	1	lef	95%	120	73.170732
## 911	0.08411	1	lef	95%	121	73.780488
## 912	0.08417	1	lef	95%	122	74.390244
## 913	0.08491	1	lef	95%	123	75.000000
## 914	0.08507	1	lef	95%	124	75.609756
## 915	0.08522	1	lef	95%	125	76.219512

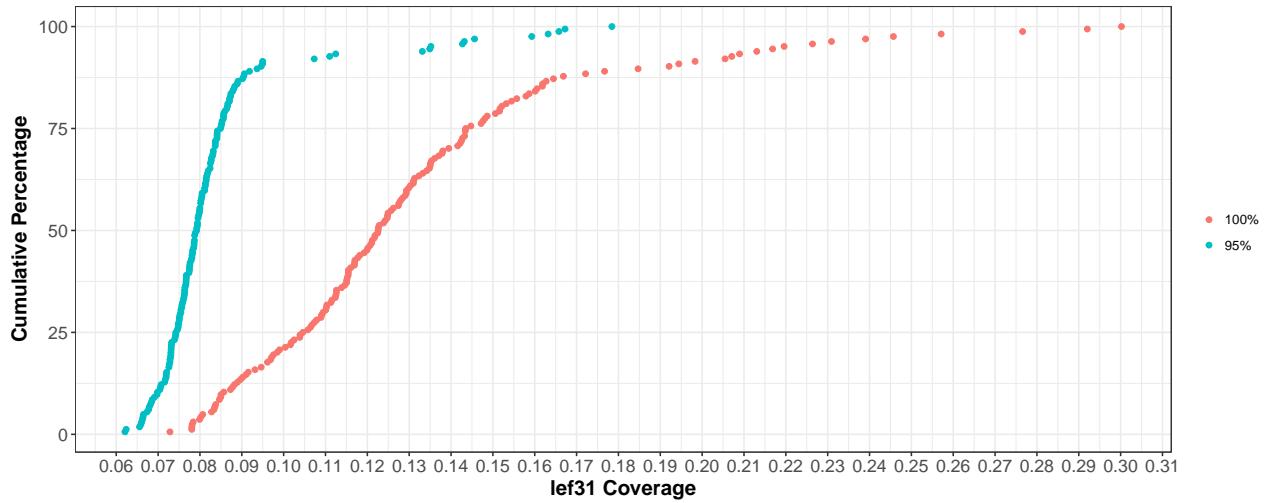
```

## 916 0.08530    1 lef 95%   126 76.829268
## 917 0.08561    1 lef 95%   127 77.439024
## 918 0.08564    1 lef 95%   128 78.048780
## 919 0.08570    1 lef 95%   129 78.658537
## 920 0.08598    1 lef 95%   130 79.268293
## 921 0.08642    1 lef 95%   131 79.878049
## 922 0.08658    1 lef 95%   132 80.487805
## 923 0.08669    1 lef 95%   133 81.097561
## 924 0.08706    1 lef 95%   134 81.707317
## 925 0.08707    1 lef 95%   135 82.317073
## 926 0.08734    1 lef 95%   136 82.926829
## 927 0.08740    1 lef 95%   137 83.536585
## 928 0.08784    1 lef 95%   138 84.146341
## 929 0.08810    1 lef 95%   139 84.756098
## 930 0.08834    1 lef 95%   140 85.365854
## 931 0.08893    1 lef 95%   141 85.975610
## 932 0.08910    1 lef 95%   142 86.585366
## 933 0.09019    1 lef 95%   143 87.195122
## 934 0.09047    1 lef 95%   144 87.804878
## 935 0.09063    1 lef 95%   145 88.414634
## 936 0.09184    1 lef 95%   146 89.024390
## 937 0.09365    1 lef 95%   147 89.634146
## 938 0.09464    1 lef 95%   148 90.243902
## 939 0.09497    1 lef 95%   149 90.853659
## 940 0.09500    1 lef 95%   150 91.463415
## 941 0.10731    1 lef 95%   151 92.073171
## 942 0.11100    1 lef 95%   152 92.682927
## 943 0.11244    1 lef 95%   153 93.292683
## 944 0.13315    1 lef 95%   154 93.902439
## 945 0.13493    1 lef 95%   155 94.512195
## 946 0.13519    1 lef 95%   156 95.121951
## 947 0.14273    1 lef 95%   157 95.731707
## 948 0.14318    1 lef 95%   158 96.341463
## 949 0.14562    1 lef 95%   159 96.951220
## 950 0.15930    1 lef 95%   160 97.560976
## 951 0.16321    1 lef 95%   161 98.170732
## 952 0.16578    1 lef 95%   162 98.780488
## 953 0.16725    1 lef 95%   163 99.390244
## 954 0.17844    1 lef 95%   164 100.000000

g <- ggplot(temp_df, aes(coverage, percent, color=status)) + geom_point() +
  scale_x_continuous(breaks = seq(0.01, max(temp_df$coverage)+0.01, by = 0.01))

p <- format_plot(g, xlab="lef31 Coverage", ylab="Cumulative Percentage")
print(p)

```



```
remove(temp_df)
```

Simulations

Inputs

```
ba_kmer_info = read.table(paste0(PATH, "/results/ba-kmer-info.txt"),
                           header=TRUE, sep="\t")

bcg_kmer_info = read.table(paste0(PATH, "/results/bcg-kmer-info.txt"),
                           header=TRUE, sep="\t")

simulation_ba = read_samples(
  paste0(PATH, "/results/simulations/simulation-ba-summary.txt.gz"),
  ba_kmer_info
)

simulation_bcg = read_samples(
  paste0(PATH, "/results/simulations/simulation-bcg-summary.txt.gz"),
  ba_kmer_info
)

simulation_lef = read_samples(
  paste0(PATH, "/results/simulations/simulation-lef-summary.txt.gz"),
  ba_kmer_info
)

simulation_ba = merge(
  simulation_ba,
  data.frame(
    sample=simulation_bcg$sample,
    simulated_coverage=simulation_bcg$simulated_coverage,
    bcg_cov_mean=simulation_bcg$kmer_cov_mean
  ),
  by=c('sample', 'simulated_coverage')
```

```
)
```

Coverages Simulated

A total of **341** coverages were simulated.

```
print(sort(unique(simulation_ba$simulated_coverage)))
```

```
## [1] 0.01 0.03 0.05 0.07 0.09 0.11 0.13 0.15 0.17 0.19 0.21  
## [12] 0.23 0.25 0.27 0.29 0.31 0.33 0.35 0.37 0.39 0.41 0.43  
## [23] 0.45 0.47 0.49 0.51 0.53 0.55 0.57 0.59 0.61 0.63 0.65  
## [34] 0.67 0.69 0.71 0.73 0.75 0.77 0.79 0.81 0.83 0.85 0.87  
## [45] 0.89 0.91 0.93 0.95 0.97 0.99 1.01 1.03 1.05 1.07 1.09  
## [56] 1.11 1.13 1.15 1.17 1.19 1.21 1.23 1.25 1.27 1.29 1.31  
## [67] 1.33 1.35 1.37 1.39 1.41 1.43 1.45 1.47 1.49 1.51 1.53  
## [78] 1.55 1.57 1.59 1.61 1.63 1.65 1.67 1.69 1.71 1.73 1.75  
## [89] 1.77 1.79 1.81 1.83 1.85 1.87 1.89 1.91 1.93 1.95 1.97  
## [100] 1.99 2.01 2.03 2.05 2.07 2.09 2.11 2.13 2.15 2.17 2.19  
## [111] 2.21 2.23 2.25 2.27 2.29 2.31 2.33 2.35 2.37 2.39 2.41  
## [122] 2.43 2.45 2.47 2.49 2.51 2.53 2.55 2.57 2.59 2.61 2.63  
## [133] 2.65 2.67 2.69 2.71 2.73 2.75 2.77 2.79 2.81 2.83 2.85  
## [144] 2.87 2.89 2.91 2.93 2.95 2.97 2.99 3.00 3.05 3.10 3.15  
## [155] 3.20 3.25 3.30 3.35 3.40 3.45 3.50 3.55 3.60 3.65 3.70  
## [166] 3.75 3.80 3.85 3.90 3.95 4.00 4.05 4.10 4.15 4.20 4.25  
## [177] 4.30 4.35 4.40 4.45 4.50 4.55 4.60 4.65 4.70 4.75 4.80  
## [188] 4.85 4.90 4.95 5.00 5.05 5.10 5.15 5.20 5.25 5.30 5.35  
## [199] 5.40 5.45 5.50 5.55 5.60 5.65 5.70 5.75 5.80 5.85 5.90  
## [210] 5.95 6.00 6.05 6.10 6.15 6.20 6.25 6.30 6.35 6.40 6.45  
## [221] 6.50 6.55 6.60 6.65 6.70 6.75 6.80 6.85 6.90 6.95 7.00  
## [232] 7.05 7.10 7.15 7.20 7.25 7.30 7.35 7.40 7.45 7.50 7.55  
## [243] 7.60 7.65 7.70 7.75 7.80 7.85 7.90 7.95 8.00 8.05 8.10  
## [254] 8.15 8.20 8.25 8.30 8.35 8.40 8.45 8.50 8.55 8.60 8.65  
## [265] 8.70 8.75 8.80 8.85 8.90 8.95 9.00 9.05 9.10 9.15 9.20  
## [276] 9.25 9.30 9.35 9.40 9.45 9.50 9.55 9.60 9.65 9.70 9.75  
## [287] 9.80 9.85 9.90 9.95 10.00 10.10 10.20 10.30 10.40 10.50 10.60  
## [298] 10.70 10.80 10.90 11.00 11.10 11.20 11.30 11.40 11.50 11.60 11.70  
## [309] 11.80 11.90 12.00 12.10 12.20 12.30 12.40 12.50 12.60 12.70 12.80  
## [320] 12.90 13.00 13.10 13.20 13.30 13.40 13.50 13.60 13.70 13.80 13.90  
## [331] 14.00 14.10 14.20 14.30 14.40 14.50 14.60 14.70 14.80 14.90 15.00
```

Samples

A total of **379** reference *Bacillus* genus genomes had reads simulated for each of the above coverages.

```
table(simulation_ba[simulation_ba$simulated_coverage == 0.01,]$bacillus)
```

```
##  
##      ba      bcg nonbcg  
##      48      95     236
```

Results

Lethal Factor

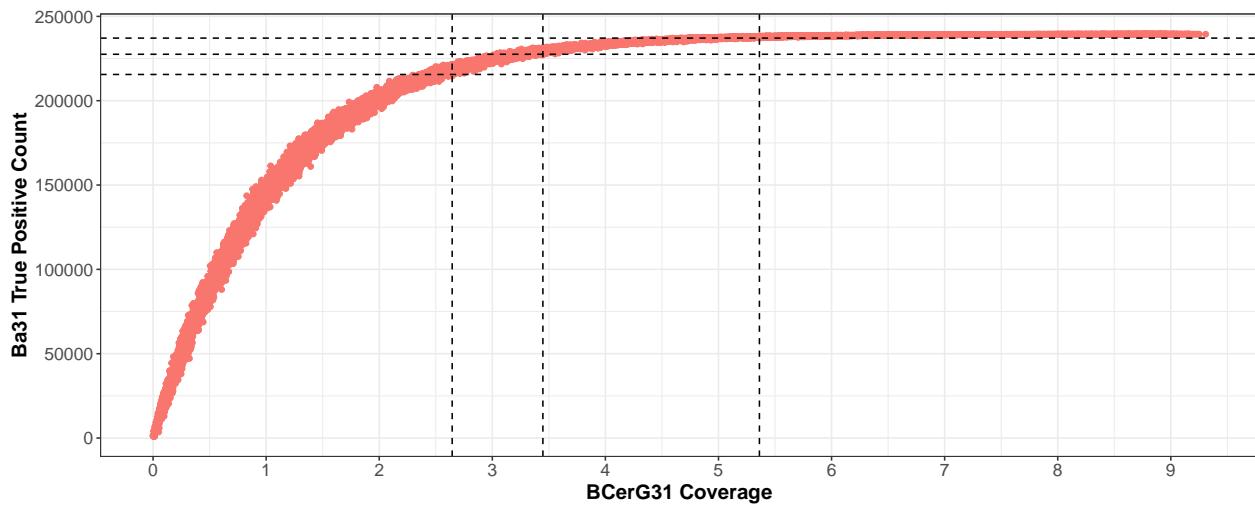
There were **0** simulations with hits to the lethal factor kmers.

Plots

Ba31 Sensitivity

```
temp_df <- simulation_ba[simulation_ba$is_ba == 'True',]
temp_total <- max(temp_df$total_kmers)
temp_90 <- floor(temp_total*0.90)
temp_95 <- floor(temp_total*0.95)
temp_99 <- floor(temp_total*0.99)
temp_999 <- floor(temp_total*0.999)
g <- ggplot(temp_df, aes(bcg_cov_mean, tp)) +
  geom_point(aes(color=is_ba)) +
  geom_hline(yintercept=temp_90, linetype="dashed") +
  geom_hline(yintercept=temp_95, linetype="dashed") +
  geom_hline(yintercept=temp_99, linetype="dashed") +
  geom_vline(xintercept=max(temp_df[temp_df$tp <= temp_90,]$bcg_cov_mean), linetype="dashed") +
  geom_vline(xintercept=max(temp_df[temp_df$tp <= temp_95,]$bcg_cov_mean), linetype="dashed") +
  geom_vline(xintercept=max(temp_df[temp_df$tp <= temp_99,]$bcg_cov_mean), linetype="dashed") +
  scale_x_continuous(breaks = seq(0, ceiling(max(temp_df$bcg_cov_mean)), by = 1))

p <- format_plot(g, xlab="BCerG31 Coverage", ylab="Ba31 True Positive Count") +
  theme(legend.position="none")
print(p)
```



```
print(
  data.frame(
    percent=c("90%", "95%", "99%", "99.9%"),
    total=c(temp_90, temp_95, temp_99, temp_999),
    min=c(
      min(temp_df[temp_df$tp >= temp_90,]$bcg_cov_mean),
      min(temp_df[temp_df$tp >= temp_95,]$bcg_cov_mean),
```

```

        min(temp_df[temp_df$tp >= temp_99,]$bcg_cov_mean),
        min(temp_df[temp_df$tp >= temp_999,]$bcg_cov_mean)
    ),
    max=c(
        max(temp_df[temp_df$tp <= temp_90,]$bcg_cov_mean),
        max(temp_df[temp_df$tp <= temp_95,]$bcg_cov_mean),
        max(temp_df[temp_df$tp <= temp_99,]$bcg_cov_mean),
        max(temp_df[temp_df$tp <= temp_999,]$bcg_cov_mean)
    )
)
)

##   percent total     min     max
## 1      90% 215552 2.3684 2.6451
## 2      95% 227527 3.0363 3.4473
## 3     99% 237107 4.6638 5.3620
## 4  99.9% 239263 6.8874 8.0657

```

Ba31 Coverage and BCerG Coverage

```

ba_bcgg_coverage_lm = lm(kmer_cov_mean ~ 0 + bcg_cov_mean, temp_df)
summary(ba_bcgg_coverage_lm)

##
## Call:
## lm(formula = kmer_cov_mean ~ 0 + bcg_cov_mean, data = temp_df)
##
## Residuals:
##       Min     1Q Median     3Q    Max
## -0.39812 -0.03832  0.00094  0.04225  0.41704
##
## Coefficients:
##             Estimate Std. Error t value            Pr(>|t|)
## bcg_cov_mean 0.9247825  0.0001525   6065 <0.0000000000000002 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.07775 on 16367 degrees of freedom
## Multiple R-squared:  0.9996, Adjusted R-squared:  0.9996
## F-statistic: 3.679e+07 on 1 and 16367 DF,  p-value: < 0.0000000000000022

cor.test(temp_df$bcg_cov_mean, temp_df$kmer_cov_mean, method = "pearson")

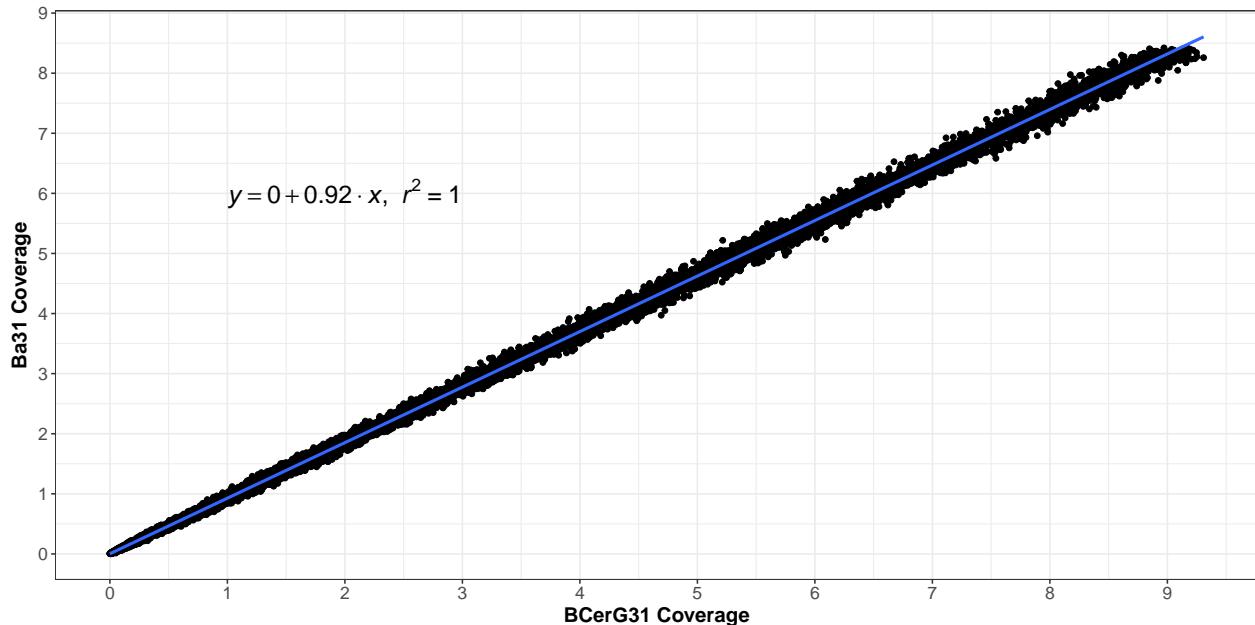
##
## Pearson's product-moment correlation
##
## data: temp_df$bcg_cov_mean and temp_df$kmer_cov_mean
## t = 3773.2, df = 16366, p-value < 0.0000000000000022
## alternative hypothesis: true correlation is not equal to 0
## 95 percent confidence interval:
##  0.9994079 0.9994431
## sample estimates:
##          cor
## 0.9994257

```

```

g <- ggplot(temp_df, aes(bcg_cov_mean, kmer_cov_mean)) +
  geom_point() +
  annotate("text", x = 2, y = 6, label = lm_eqn(ba_bcerg31_anthraxis_lm), parse = TRUE, size=6) +
  geom_smooth(method='lm') +
  scale_y_continuous(breaks = seq(0, ceiling(max(temp_df$kmer_cov_mean)), by = 1)) +
  scale_x_continuous(breaks = seq(0, ceiling(max(temp_df$bcg_cov_mean)), by = 1))
p <- format_plot(g, xlab="BCerG31 Coverage", ylab="Ba31 Coverage")
print(p)

```



```
write_plot(p, 'supplementary-figure-02-ba31-bcerg31-anthraxis', height=6, width=12)
```

Ba31 Coverage and Genome Coverage

```

ba_genome_coverages_lm = lm(kmer_cov_mean ~ 0 + simulated_coverage, temp_df)
summary(ba_genome_coverages_lm)

```

```

##
## Call:
## lm(formula = kmer_cov_mean ~ 0 + simulated_coverage, data = temp_df)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -0.131475 -0.010506  0.000081  0.010775  0.132931
##
## Coefficients:
##             Estimate Std. Error t value     Pr(>|t|)
## simulated_coverage 0.55641329 0.00002439 22814 <0.0000000000000002 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.02067 on 16367 degrees of freedom
## Multiple R-squared:      1, Adjusted R-squared:      1

```

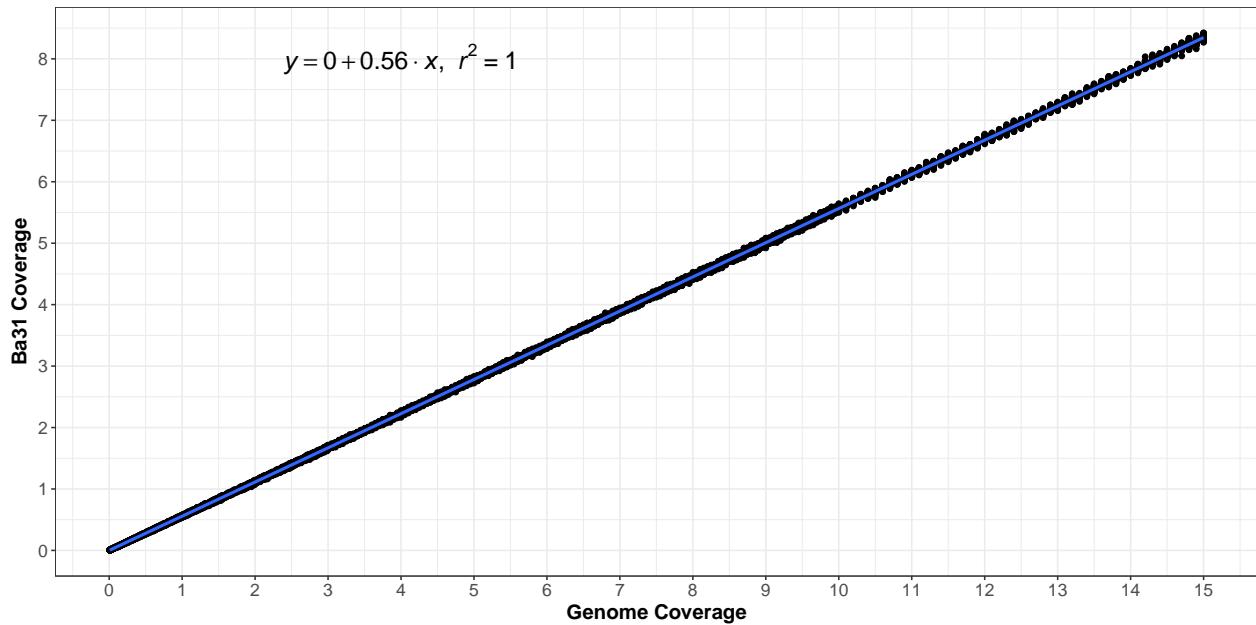
```

## F-statistic: 5.205e+08 on 1 and 16367 DF, p-value: < 0.00000000000000022
cor.test(temp_df$kmer_cov_mean, temp_df$simulated_coverage, method = "pearson")

##
## Pearson's product-moment correlation
##
## data: temp_df$kmer_cov_mean and temp_df$simulated_coverage
## t = 14192, df = 16366, p-value < 0.00000000000000022
## alternative hypothesis: true correlation is not equal to 0
## 95 percent confidence interval:
## 0.9999581 0.9999606
## sample estimates:
## cor
## 0.9999594

g <- ggplot(temp_df, aes(simulated_coverage, kmer_cov_mean)) +
  geom_point() +
  annotate("text", x = 4, y = 8, label = lm_eqn(ba_genome_coverages_lm), parse = TRUE, size=6) +
  geom_smooth(method='lm') +
  scale_y_continuous(breaks = seq(0, ceiling(max(temp_df$kmer_cov_mean)), by = 1)) +
  scale_x_continuous(breaks = seq(0, ceiling(max(temp_df$simulated_coverage)), by = 1))
ba_coverage_plot <- format_plot(g, xlab="Genome Coverage", ylab="Ba31 Coverage")
print(ba_coverage_plot)

```



```

remove(temp_df)
remove(temp_total)
remove(temp_90)
remove(temp_95)
remove(temp_99)
remove(temp_999)

```

Ba31 Coverage and BCerG Coverage (non-anthracis)

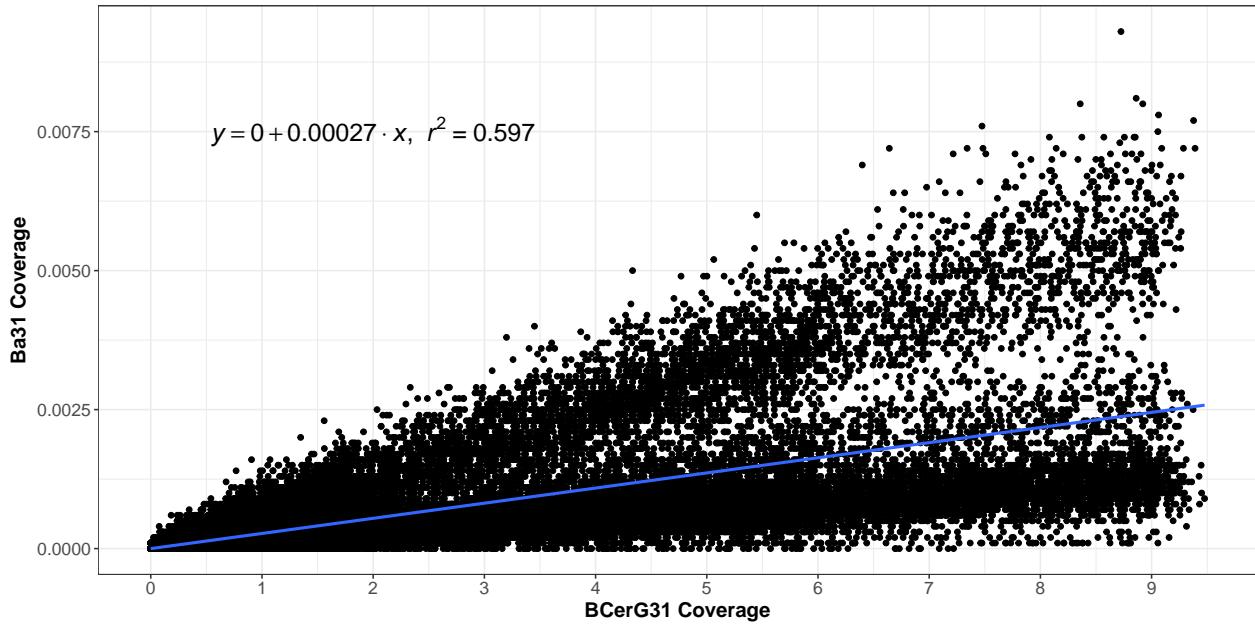
```
temp_df <- simulation_ba[simulation_ba$is_ba == 'False',]
ba_bcg_coverages_lm_non_anthracis = lm(kmer_cov_mean ~ 0 + bcg_cov_mean, temp_df)
summary(ba_bcg_coverages_lm_non_anthracis)

##
## Call:
## lm(formula = kmer_cov_mean ~ 0 + bcg_cov_mean, data = temp_df)
##
## Residuals:
##       Min        1Q    Median        3Q       Max
## -0.002365  0.000000  0.000000  0.000000  0.006925
##
## Coefficients:
##             Estimate Std. Error t value     Pr(>|t|)    
## bcg_cov_mean 0.0002722235 0.0000006653   409.2 <0.000000000000002 *** 
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.0004821 on 112870 degrees of freedom
## Multiple R-squared:  0.5973, Adjusted R-squared:  0.5973 
## F-statistic: 1.674e+05 on 1 and 112870 DF,  p-value: < 0.0000000000000022

cor.test(temp_df$bcg_cov_mean, temp_df$kmer_cov_mean, method = "pearson")

##
## Pearson's product-moment correlation
##
## data: temp_df$bcg_cov_mean and temp_df$kmer_cov_mean
## t = 371.5, df = 112870, p-value < 0.0000000000000022
## alternative hypothesis: true correlation is not equal to 0
## 95 percent confidence interval:
##  0.7390547 0.7443041
## sample estimates:
## cor
## 0.7416908

g <- ggplot(temp_df, aes(bcg_cov_mean, kmer_cov_mean)) +
  geom_point() +
  annotate("text", x = 2, y = 0.0075, label = lm_eqn(ba_bcg_coverages_lm_non_anthracis), parse = TRUE)
  geom_smooth(method='lm') +
  scale_x_continuous(breaks = seq(0, ceiling(max(temp_df$bcg_cov_mean)), by = 1))
p <- format_plot(g, xlab="BCerG31 Coverage", ylab="Ba31 Coverage")
print(p)
```



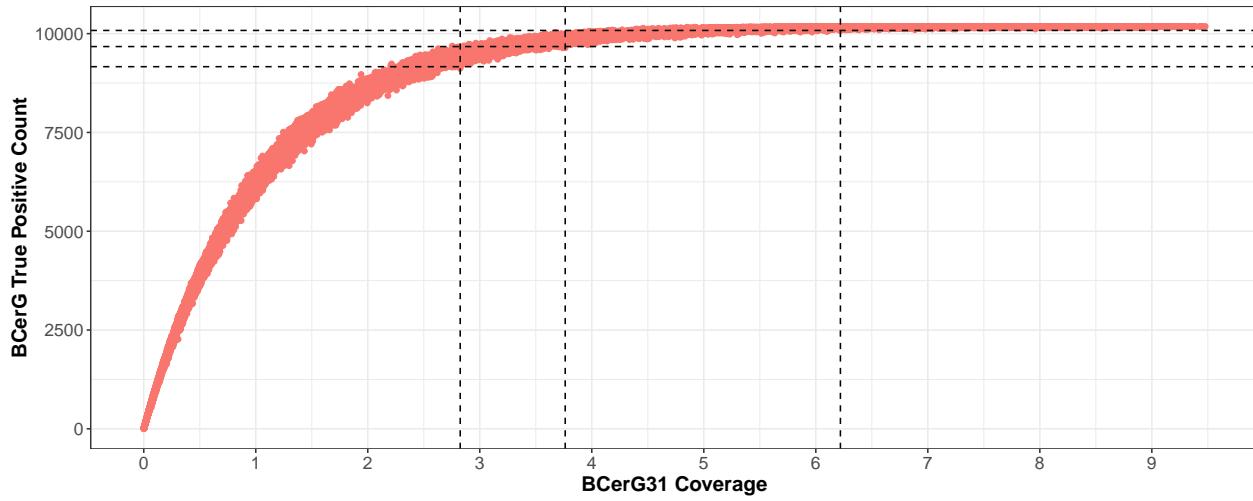
```
write_plot(p, 'supplementary-figure-03-ba31-bcerg31-nonanthracis', height=6, width=12)

remove(temp_df)
```

BCerG31 Sensitivity

```
temp_df <- simulation_bcg[simulation_bcg$is_bcg == 'True',]
temp_total <- max(temp_df$total_kmers)
temp_90 <- floor(temp_total*0.90)
temp_95 <- floor(temp_total*0.95)
temp_99 <- floor(temp_total*0.99)
temp_999 <- floor(temp_total*0.999)
g <- ggplot(temp_df, aes(kmer_cov_mean, tp)) +
  geom_point(aes(color=is_bcg)) +
  geom_hline(yintercept=temp_90, linetype="dashed") +
  geom_hline(yintercept=temp_95, linetype="dashed") +
  geom_hline(yintercept=temp_99, linetype="dashed") +
  geom_vline(xintercept=max(temp_df[temp_df$tp <= temp_90,]$kmer_cov_mean), linetype="dashed") +
  geom_vline(xintercept=max(temp_df[temp_df$tp <= temp_95,]$kmer_cov_mean), linetype="dashed") +
  geom_vline(xintercept=max(temp_df[temp_df$tp <= temp_99,]$kmer_cov_mean), linetype="dashed") +
  scale_x_continuous(breaks = seq(0, ceiling(max(temp_df$kmer_cov_mean)), by = 1))

p <- format_plot(g, xlab="BCerG31 Coverage", ylab="BCerG True Positive Count") +
  theme(legend.position="none")
print(p)
```



```

print(
  data.frame(
    percent=c("90%", "95%", "99%", "99.9%"),
    total=c(temp_90, temp_95, temp_99, temp_999),
    min=c(
      min(temp_df[temp_df$tp >= temp_90,]$kmer_cov_mean),
      min(temp_df[temp_df$tp >= temp_95,]$kmer_cov_mean),
      min(temp_df[temp_df$tp >= temp_99,]$kmer_cov_mean),
      min(temp_df[temp_df$tp >= temp_999,]$kmer_cov_mean)
    ),
    max=c(
      max(temp_df[temp_df$tp <= temp_90,]$kmer_cov_mean),
      max(temp_df[temp_df$tp <= temp_95,]$kmer_cov_mean),
      max(temp_df[temp_df$tp <= temp_99,]$kmer_cov_mean),
      max(temp_df[temp_df$tp <= temp_999,]$kmer_cov_mean)
    )
  )
)

##   percent total     min     max
## 1    90%  9164 2.2064 2.8252
## 2    95%  9673 2.7546 3.7627
## 3    99% 10081 4.1588 6.2195
## 4  99.9% 10172 5.3514 9.1855

```

BCerG31 coverage and Genome Coverage

```

bcg_genome_coverages_lm = lm(kmer_cov_mean ~ 0 + simulated_coverage, temp_df)
summary(bcg_genome_coverages_lm)

##
## Call:
## lm(formula = kmer_cov_mean ~ 0 + simulated_coverage, data = temp_df)
##
## Residuals:
##       Min     1Q Median     3Q    Max 
## -0.55991 -0.04287 -0.00093  0.04227  0.52956

```

```

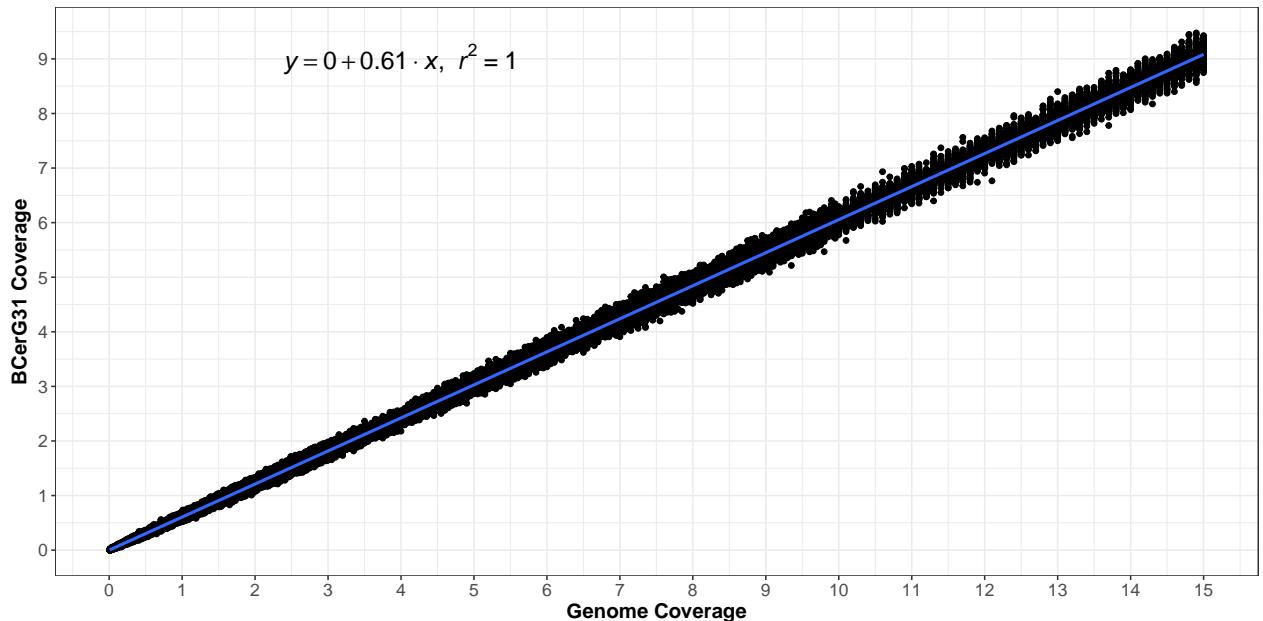
## 
## Coefficients:
##                               Estimate Std. Error t value      Pr(>|t|)    
## simulated_coverage  0.60547993  0.00005889   10282 <0.0000000000000002 *** 
## --- 
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1  
## 
## Residual standard error: 0.08616 on 48762 degrees of freedom
## Multiple R-squared:  0.9995, Adjusted R-squared:  0.9995 
## F-statistic: 1.057e+08 on 1 and 48762 DF,  p-value: < 0.0000000000000022 

cor.test(temp_df$kmer_cov_mean, temp_df$simulated_coverage, method = "pearson")

## 
## Pearson's product-moment correlation
## 
## data: temp_df$kmer_cov_mean and temp_df$simulated_coverage
## t = 6396.7, df = 48761, p-value < 0.0000000000000022
## alternative hypothesis: true correlation is not equal to 0
## 95 percent confidence interval:
##  0.9993940 0.9994152
## sample estimates:
##        cor
## 0.9994047

g <- ggplot(temp_df, aes(simulated_coverage, kmer_cov_mean)) +
  geom_point() +
  annotate("text", x = 4, y = 9, label = lm_eqn(bcg_genome_coverages_lm), parse = TRUE, size=6) +
  geom_smooth(method='lm') +
  scale_y_continuous(breaks = seq(0, ceiling(max(temp_df$kmer_cov_mean)), by = 1)) +
  scale_x_continuous(breaks = seq(0, ceiling(max(temp_df$simulated_coverage)), by = 1))
bcg_coverage_plot <- format_plot(g, xlab="Genome Coverage", ylab="BCerG31 Coverage")
print(bcg_coverage_plot)

```



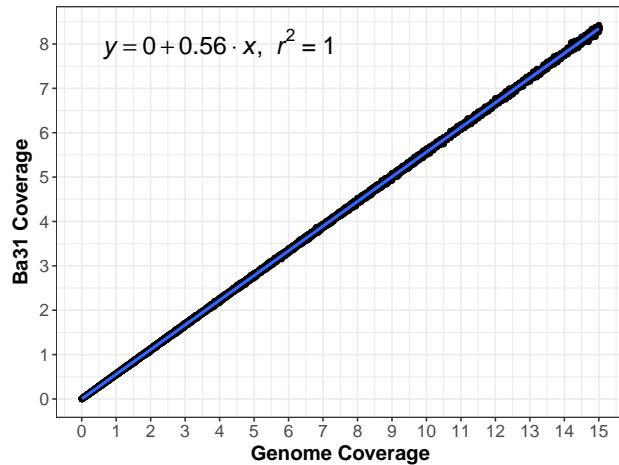
```

remove(temp_df)
remove(temp_total)
remove(temp_90)
remove(temp_95)
remove(temp_99)
remove(temp_999)

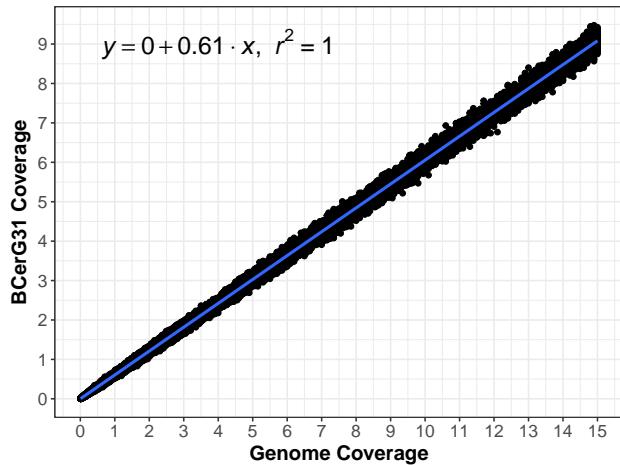
p1 <- arrangeGrob(ba_coverage_plot, top = textGrob(
  "A", x = unit(0, "npc"), y = unit(1, "npc"), just=c("left","top"),
  gp=gpar(col="black", fontsize=18, fontface="bold"))
)
p2 <- arrangeGrob(bc_gCoverage_plot, top = textGrob(
  "B", x = unit(0, "npc"), y = unit(1, "npc"), just=c("left","top"),
  gp=gpar(col="black", fontsize=18, fontface="bold"))
)
plots=list()
plots[[1]] = p1
plots[[2]] = p2
grid.arrange(grobs=plots, ncol=2)

```

A



B



```

pdf(paste0(FIGURE, '/supplementary-figure-01-31-genome-coverage.pdf'),
  height=6, width=12, onefile=FALSE)
grid.arrange(grobs=plots, ncol=2)
dev_null <- dev.off()

png(paste0(FIGURE, '/supplementary-figure-01-31-genome-coverage.png'),
  height=600, width=1200)
grid.arrange(grobs=plots, ncol=2)
dev_null <- dev.off()

```

non-anthraxis BCerG False Positive Model

Parse Results

```

bcg_model_set = read_samples(
  paste0(PATH, "/results/model-set/model-set-ba-summary.txt.gz"),
  ba_kmer_info
)

temp_df = read_samples(
  paste0(PATH, "/results/model-set/model-set-bcg-summary.txt.gz"),
  ba_kmer_info
)

bcg_model_set = merge(
  bcg_model_set,
  data.frame(
    sample=temp_df$sample,
    coverage=temp_df$coverage,
    replicate=temp_df$replicate,
    bcg_cov_mean=temp_df$kmer_cov_mean,
    bcg_non_zero_kmer_cov_mean=temp_df$non_zero_kmer_cov_mean
  ),
  by=c('sample', 'coverage', 'replicate')
)

bcg_model_set = merge(
  bcg_model_set,
  bcg_model_set %>%
    group_by(sample, coverage) %>%
    summarise(fp_mean=mean(fp), ba_mean=mean(kmer_cov_mean), bcg_mean=mean(bcg_cov_mean)),
  by=c('sample', 'coverage')
)
remove(temp_df)

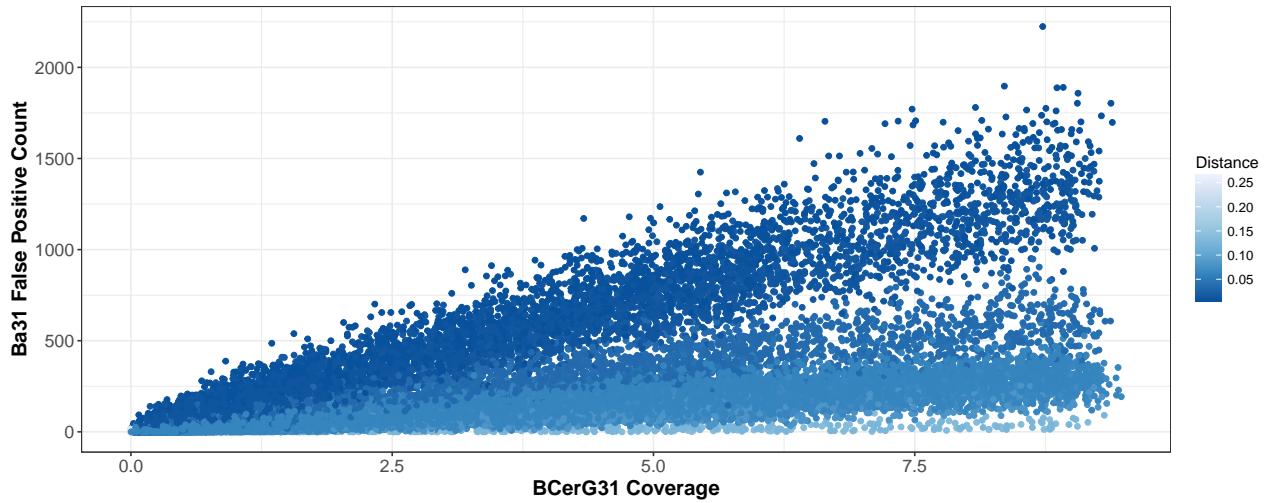
```

BA31 False Positive Count and Distance

```

p <- ggplot(simulation_ba[simulation_ba$is_ba == 'False',],
             aes(bcg_cov_mean, fp, color=distance)) +
  geom_point() +
  xlab("BCerG31 Coverage") +
  ylab("Ba31 False Positive Count") +
  theme_bw() +
  theme(axis.text=element_text(size=12),
        axis.title=element_text(size=14,face="bold")) +
  scale_colour_distiller(palette="Blues") +
  labs(color='Distance')
print(p)

```



```
write_plot(p, 'supplementary-figure-04-distance-fp', height=6, width=12)
```

non-*B. anthracis* BCG Model Set

```
print(as.character(sort(unique(bcg_model_set$sample))))  
  
## [1] "NC_011773"    "NZ_CP009335"  "NZ_CP009596"  "NZ_CP009605"  "NZ_CP009720"  
## [6] "NZ_CP018931"  "NZ_CP018933"  "NZ_CP018935"
```

Model BCerG kmer coverage and False Positive Match

```
bcg_model = lm(total_count ~ 0 + bcg_cov_mean, bcg_model_set)  
anova(bcg_model)  
  
## Analysis of Variance Table  
##  
## Response: total_count  
##                   Df     Sum Sq   Mean Sq F value  
## bcg_cov_mean      1 719180371096 719180371096 43457509  
## Residuals       311231  5150576486          16549  
##                         Pr(>F)  
## bcg_cov_mean < 0.0000000000000022 ***  
## Residuals  
## ---  
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1  
summary(bcg_model)  
  
##  
## Call:  
## lm(formula = total_count ~ 0 + bcg_cov_mean, data = bcg_model_set)  
##  
## Residuals:  
##      Min       1Q   Median       3Q      Max  
## -1463.46   -53.09   -0.07   51.61  1794.24  
##
```

```

## Coefficients:
##              Estimate Std. Error t value      Pr(>|t|)
## bcg_cov_mean 171.84251    0.02607   6592 <0.0000000000000002 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 128.6 on 311231 degrees of freedom
## Multiple R-squared:  0.9929, Adjusted R-squared:  0.9929
## F-statistic: 4.346e+07 on 1 and 311231 DF,  p-value: < 0.0000000000000022

```

Create a function to make predictions with 99% prediction intervals

```

predict_false_positive <- function(model, coverages) {
  temp_df <- cbind(
    data.frame(bcg_cov_mean=coverages),
    predict(
      model,
      newdata = data.frame(bcg_cov_mean=coverages),
      se=TRUE,
      interval='prediction',
      level=0.99
    )
  )
  return(temp_df)
}

```

Make a few predictions

```

predict_false_positive(bcg_model, seq(0, 10, 2))

##   bcg_cov_mean   fit.fit   fit.lwr   fit.upr   se.fit     df
## 1          0  0.0000 -331.3647  331.3647 0.00000000 311231
## 2          2 343.6850   12.3203  675.0498 0.05213486 311231
## 3          4 687.3701  356.0052 1018.7349 0.10426972 311231
## 4          6 1031.0551  699.6901 1362.4200 0.15640457 311231
## 5          8 1374.7401 1043.3750 1706.1052 0.20853943 311231
## 6         10 1718.4251 1387.0598 2049.7905 0.26067429 311231
##   residual.scale
## 1        128.6431
## 2        128.6431
## 3        128.6431
## 4        128.6431
## 5        128.6431
## 6        128.6431

```

Plot: BCerG kmer Coverage and False Positive Matches

```

plot_model_fit <- function(samples, model, coverage, coverages) {
  predicted_fit <- predict_false_positive(bcg_model, coverages)
  g <- ggplot(samples[samples$bcg_cov_mean <= coverage,], aes(bcg_cov_mean, fp))+
```

```

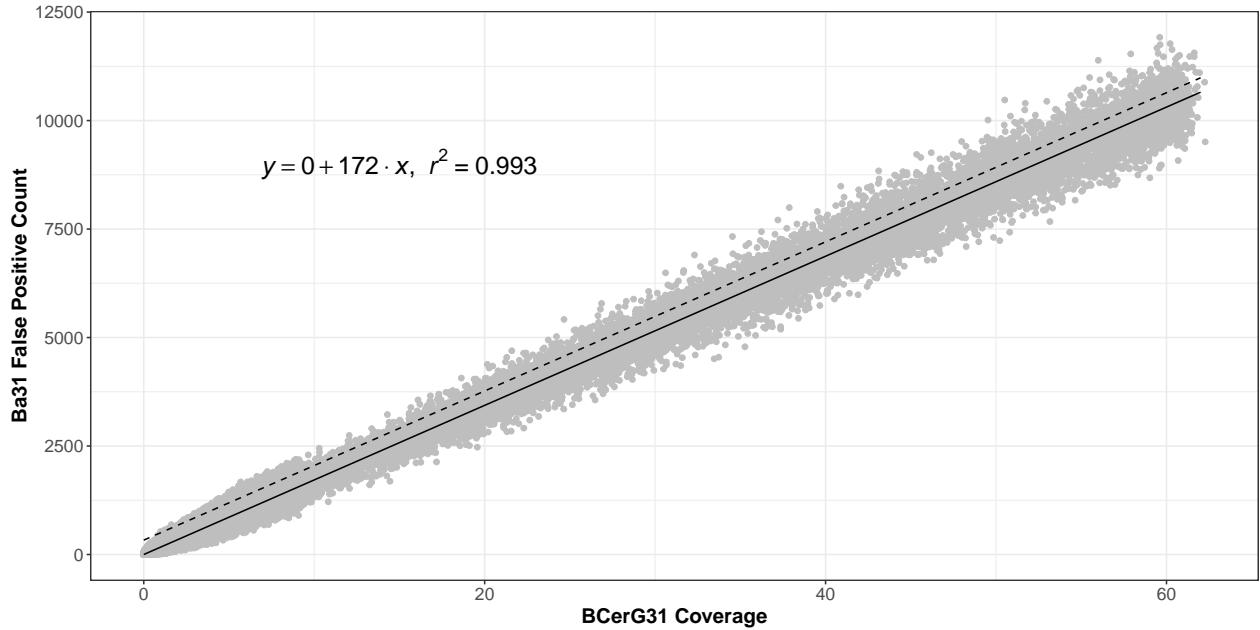
    geom_point(aes(color=is_ba)) +
    geom_line(data=predicted_fit, aes(x=bcg_cov_mean, y=fit.fit)) +
    geom_line(data=predicted_fit, aes(x=bcg_cov_mean, y=fit.upr), linetype="dashed")

  p <- format_plot(g, ylab="Ba31 False Positive Count", xlab="BCerG31 Coverage") +
    theme(legend.position="none")
  remove(predicted_fit)
  return(p)
}

predicted_fit <- predict_false_positive(
  bcg_model, seq(0, max(bcg_model_set$bcg_cov_mean), 1)
)
g <- ggplot(bcg_model_set, aes(bcg_cov_mean, total_count)) +
  geom_point(color="gray") +
  annotate("text", x = 15, y = 9000, label = lm_eqn(bcg_model), parse = TRUE, size=6) +
  geom_line(data=predicted_fit, aes(x=bcg_cov_mean, y=fit.fit)) +
  geom_line(data=predicted_fit, aes(x=bcg_cov_mean, y=fit.upr), linetype="dashed") +
  theme(legend.position="none")

p <- format_plot(g, ylab="Ba31 False Positive Count", xlab="BCerG31 Coverage")
remove(predicted_fit)
print(p)

```



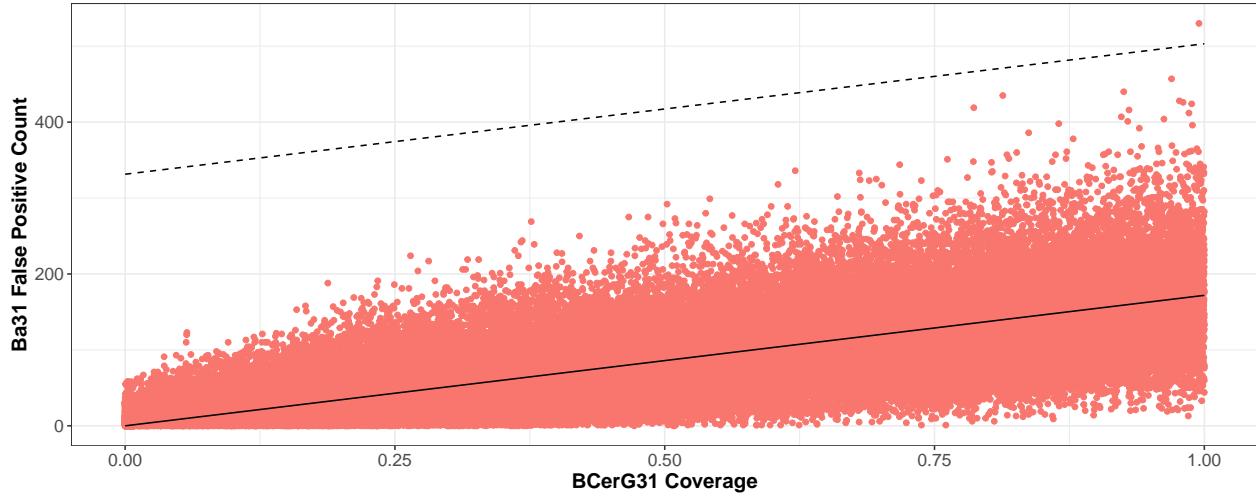
```
write_plot(p, 'figure-03-model', height=6, width=12)
```

Plot the fit between 0 and 1x BCerG kmer coverage

```

p <- plot_model_fit(bcg_model_set, bcg_model, 1, seq(0, 1, 0.1))
print(p)

```



Plot: BCerG False positives and the equivalent true positive BCerG coverage using counts from simulations

```

matches = max(bcg_model_set$fp)
g <- ggplot(bcg_model_set, aes(bcg_cov_mean, fp))+
  geom_point()
p1 <- format_plot(g, ylab="Ba31 False Positive Count", xlab="BCerG31 Coverage") +
  theme(legend.position="none")

p1 <- arrangeGrob(p1, top = textGrob(
  "A", x = unit(0, "npc"), y = unit(1, "npc"), just=c("left","top"),
  gp=gpar(col="black", fontsize=18, fontface="bold"))
)

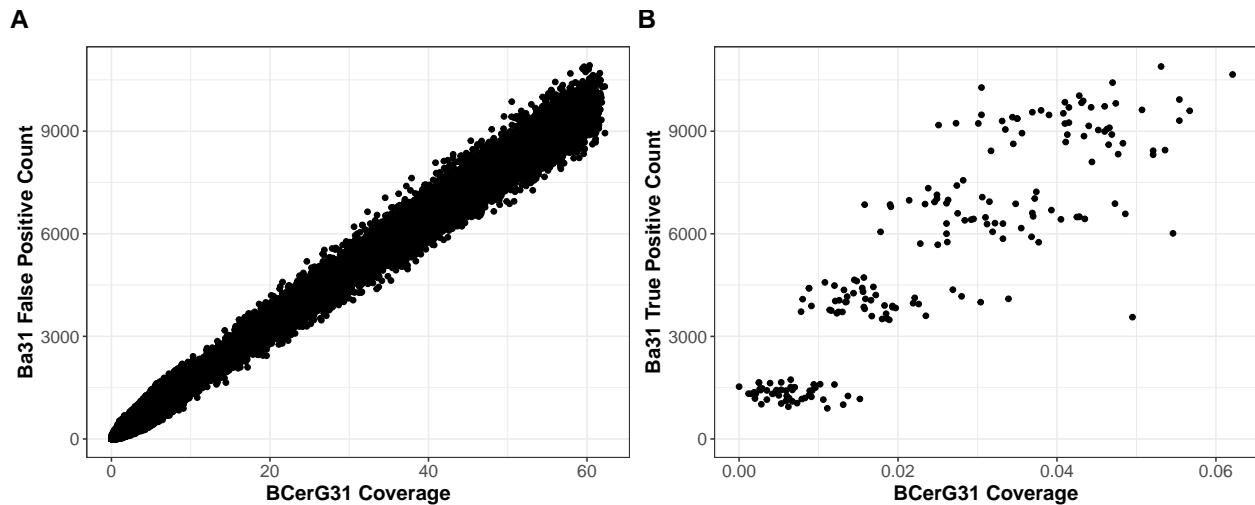
g <- ggplot(simulation_ba[simulation_ba$tp <= matches & simulation_ba$is_ba == 'True',],
            aes(bcg_cov_mean, tp)) +
  geom_point() +
  ylim(0, matches)

p2 <- format_plot(g, ylab="Ba31 True Positive Count", xlab="BCerG31 Coverage") +
  theme(legend.position="none")

p2 <- arrangeGrob(p2, top = textGrob(
  "B", x = unit(0, "npc"), y = unit(1, "npc"), just=c("left","top"),
  gp=gpar(col="black", fontsize=18, fontface="bold"))
)

grid.arrange(grobs=list(p1, p2), ncol=2)

```



B. anthracis Limit of Detection in *B. anthracis* and *B. cereus* Mixtures

Parse Results

```

kraken_report = read.table(
  paste0(PATH, "/results/limit-of-detection-ba31/kraken-report.txt"),
  header=TRUE, sep="\t"
)

subsample_ba_samples = read.table(
  gzfile(paste0(PATH, "/results/limit-of-detection-ba31/mixture-ba-summary.txt.gz")),
  header=TRUE, sep="\t"
)

subsample_bcg_samples = read.table(
  gzfile(paste0(PATH, "/results/limit-of-detection-ba31/mixture-bcg-summary.txt.gz")),
  header=TRUE, sep="\t"
)

subsample_ba_samples = merge(
  subsample_ba_samples,
  data.frame(
    replicate=subsample_bcg_samples$replicate,
    bcg_tp=subsample_bcg_samples$tp,
    bcg_coverage=subsample_bcg_samples$bcg_coverage,
    ba_coverage=subsample_bcg_samples$ba_coverage,
    bcg_cov_mean=subsample_bcg_samples$kmer_cov_mean
  ),
  by=c('replicate', 'ba_coverage', 'bcg_coverage')
)

subsample_summary <- subsample_ba_samples %>%
  group_by(.dots=c('bcg_coverage', 'ba_coverage')) %>%

```

```

    summarise(
      ba31_hits=mean(tp + fp),
      ba31_coverage=mean(kmer_cov_mean),
      bcerg31_coverage=mean(bcrg_cov_mean)
    )

subsample_summary$model_fit <- predict_false_positive(bcrg_model, subsample_summary$bcerg31_coverage)$fit
subsample_summary$model_99pi <- predict_false_positive(bcrg_model, subsample_summary$bcerg31_coverage)$fit.upr
subsample_summary$ba_detectable <- subsample_summary$ba31_hits > subsample_summary$model_99pi
subsample_summary$grey_zone <- subsample_summary$ba31_coverage < 0.1

subsample_summary <- merge(
  subsample_summary,
  kraken_report,
  by=c('bcg_coverage', 'ba_coverage')
)

exceeds_model <- function(mf, cov, count) {
  return(count > mf[mf$bcg_cov_mean == cov,]$fit.upr)
}

plot_ba_lod <- function(summary, bcg, ba) {
  temp_df <- summary[summary$bcg_coverage == bcg & summary$ba_coverage < ba,]

  predicted_fit <- predict_false_positive(
    bcg_model, unique(temp_df$bcerg31_coverage)
  )
  lod <- min(temp_df[
    apply(temp_df, 1, function(x) {
      exceeds_model(
        predicted_fit,
        x[["bcerg31_coverage"]],
        x[["ba31_hits"]]
      )
    })
  ]$ba_coverage)
  temp_df$detectable <- ifelse(temp_df$ba_coverage >= lod, TRUE, FALSE)
  temp_df$ba_coverage_text <- ifelse(temp_df$ba_coverage == lod, paste0(lod, "x"), "")
  g <- ggplot(temp_df, aes(bcerg31_coverage, ba31_hits)) +
    geom_point(aes(color=detectable)) +
    # geom_text(aes(label=ba_coverage), hjust=-0.2, vjust=0) +
    geom_text(aes(label=ba_coverage_text), hjust=-0.2, vjust=0) +
    geom_line(data=predicted_fit, aes(x=bcg_cov_mean, y=fit.fit)) +
    geom_line(data=predicted_fit, aes(x=bcg_cov_mean, y=fit.upr), linetype="dashed")

  p <- format_plot(g, xlab="BCerG31 Coverage", ylab="Ba31 Count", title="Ba31 Detectable")
  remove(predicted_fit)
  remove(lod)
  return(p)
}

write.table(
  subsample_summary,

```

```

paste0(PATH, "/results/limit-of-detection-ba31/ba31-lod-full-table.txt"),
sep="\t",
quote = FALSE,
row.names = FALSE
)

```

B. anthracis Subsampled Coverages

A total of **39** *B. anthracis* subsamples were used.

```

print(sort(unique(subsample_ba_samples$ba_coverage)))

## [1] 0.0000 0.0001 0.0002 0.0003 0.0004 0.0005 0.0006 0.0007 0.0008 0.0009
## [11] 0.0010 0.0020 0.0030 0.0040 0.0050 0.0060 0.0070 0.0080 0.0090 0.0100
## [21] 0.0200 0.0300 0.0400 0.0500 0.0600 0.0700 0.0800 0.0900 0.1000 0.1100
## [31] 0.1200 0.1300 0.1400 0.1500 0.1600 0.1700 0.1800 0.1900 0.2000

```

non-*B. anthracis* BCerG Subsample Coverages

A total of **8** non-*B. anthracis* BCerG subsamples were used.

```

print(sort(unique(subsample_ba_samples$bco_coverage)))

## [1] 0 1 5 10 25 50 75 100

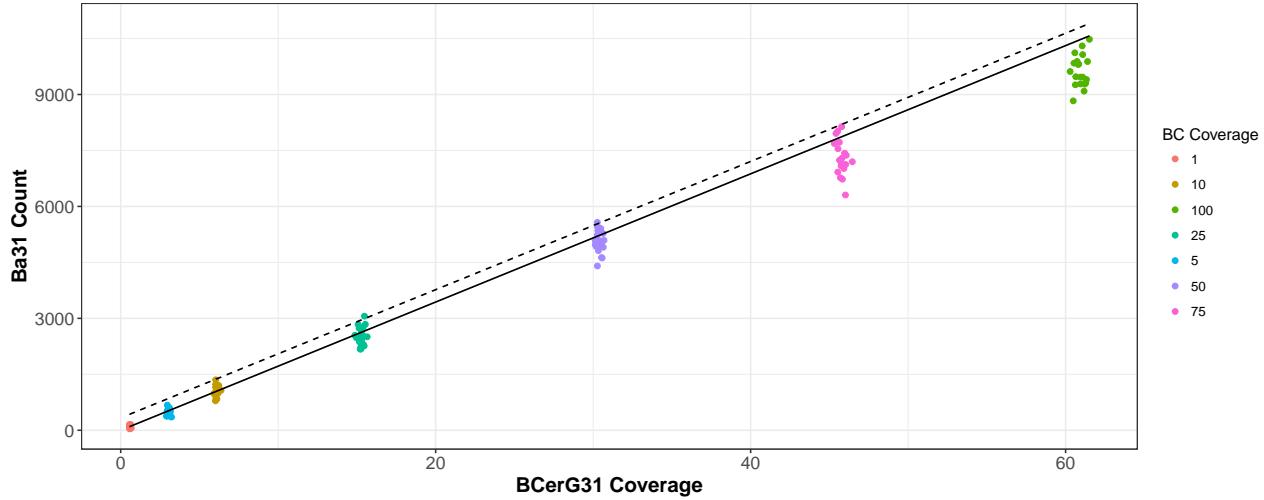
```

B. cereus Only

```

subsample_ba_samples$bco_coverage_text <- as.character(subsample_ba_samples$bco_coverage)
predicted_fit <- predict_false_positive(
  bcg_model, seq(min(subsample_ba_samples[subsample_ba_samples$ba_coverage == 0 ,]$bcg_cov_mean), max)
)
g <- ggplot(subsample_ba_samples[subsample_ba_samples$ba_coverage == 0 ,], aes(bcg_cov_mean, fp)) +
  geom_point(aes(color=bcg_coverage_text)) +
  geom_line(data=predicted_fit, aes(x=bcg_cov_mean, y=fit.fit)) +
  geom_line(data=predicted_fit, aes(x=bcg_cov_mean, y=fit.upr), linetype="dashed")
p <- format_plot(g, xlab="BCerG31 Coverage", ylab="Ba31 Count", title= "BC Coverage")
print(p)

```



```

p1 <- plot_ba_lod(subsample_summary, 0, 0.01)
p2 <- plot_ba_lod(subsample_summary, 1, 0.01)
p3 <- plot_ba_lod(subsample_summary, 5, 0.01)
p4 <- plot_ba_lod(subsample_summary, 10, 0.01)
p5 <- plot_ba_lod(subsample_summary, 25, 0.05)
p6 <- plot_ba_lod(subsample_summary, 50, 0.05)
p7 <- plot_ba_lod(subsample_summary, 75, 0.05)
p8 <- plot_ba_lod(subsample_summary, 100, 0.05)

gridArrangeSharedBa31 <- function(..., ncol = length(list(...)), nrow = 1, position = c("bottom", "right"))
  plots <- list(...)
  position <- match.arg(position)
  g <- ggplotGrob(plots[[1]] + theme(legend.position = position))$grobs
  legend <- g[[which(sapply(g, function(x) x$name) == "guide-box")]]
  lheight <- sum(legend$height)
  lwidth <- sum(legend$width)
  gl <- lapply(plots, function(x) x + theme(legend.position="none"))
  gl[[1]] <-arrangeGrob(gl[[1]], top = textGrob(
    "A) 0x", x = unit(0, "npc"), y = unit(1, "npc"), just=c("left","top"),
    gp=gpar(col="black", fontsize=18, fontface="bold"))
  )
  gl[[2]] <-arrangeGrob(gl[[2]], top = textGrob(
    "B) 1x", x = unit(0, "npc"), y = unit(1, "npc"), just=c("left","top"),
    gp=gpar(col="black", fontsize=18, fontface="bold"))
  )
  gl[[3]] <-arrangeGrob(gl[[3]], top = textGrob(
    "C) 5x", x = unit(0, "npc"), y = unit(1, "npc"), just=c("left","top"),
    gp=gpar(col="black", fontsize=18, fontface="bold"))
  )
  gl[[4]] <-arrangeGrob(gl[[4]], top = textGrob(
    "D) 10x", x = unit(0, "npc"), y = unit(1, "npc"), just=c("left","top"),
    gp=gpar(col="black", fontsize=18, fontface="bold"))
  )
  gl[[5]] <-arrangeGrob(gl[[5]], top = textGrob(
    "E) 25x", x = unit(0, "npc"), y = unit(1, "npc"), just=c("left","top"),
    gp=gpar(col="black", fontsize=18, fontface="bold"))
  )
  gl[[6]] <-arrangeGrob(gl[[6]], top = textGrob(
    "F) 50x", x = unit(0, "npc"), y = unit(1, "npc"), just=c("left","top"),
    gp=gpar(col="black", fontsize=18, fontface="bold"))
  )
  
```

```

    "F) 50x", x = unit(0, "npc"), y = unit(1, "npc"), just=c("left","top"),
    gp=gpar(col="black", fontsize=18, fontface="bold"))
)
g1[[7]] <- arrangeGrob(g1[[7]], top = textGrob(
    "G) 75x", x = unit(0, "npc"), y = unit(1, "npc"), just=c("left","top"),
    gp=gpar(col="black", fontsize=18, fontface="bold"))
)
g1[[8]] <- arrangeGrob(g1[[8]], top = textGrob(
    "H) 100x", x = unit(0, "npc"), y = unit(1, "npc"), just=c("left","top"),
    gp=gpar(col="black", fontsize=18, fontface="bold"))
)

g1 <- c(g1, ncol = ncol, nrow = nrow)

combined <- switch(position,
    "bottom" = arrangeGrob(do.call(arrangeGrob, g1),
        legend,
        ncol = 1,
        heights = unit.c(unit(1, "npc") - lheight, lheight)),
    "right" = arrangeGrob(do.call(arrangeGrob, g1),
        legend,
        ncol = 2,
        widths = unit.c(unit(1, "npc") - lwidth, lwidth)))
}

grid.newpage()
grid.draw(combined)

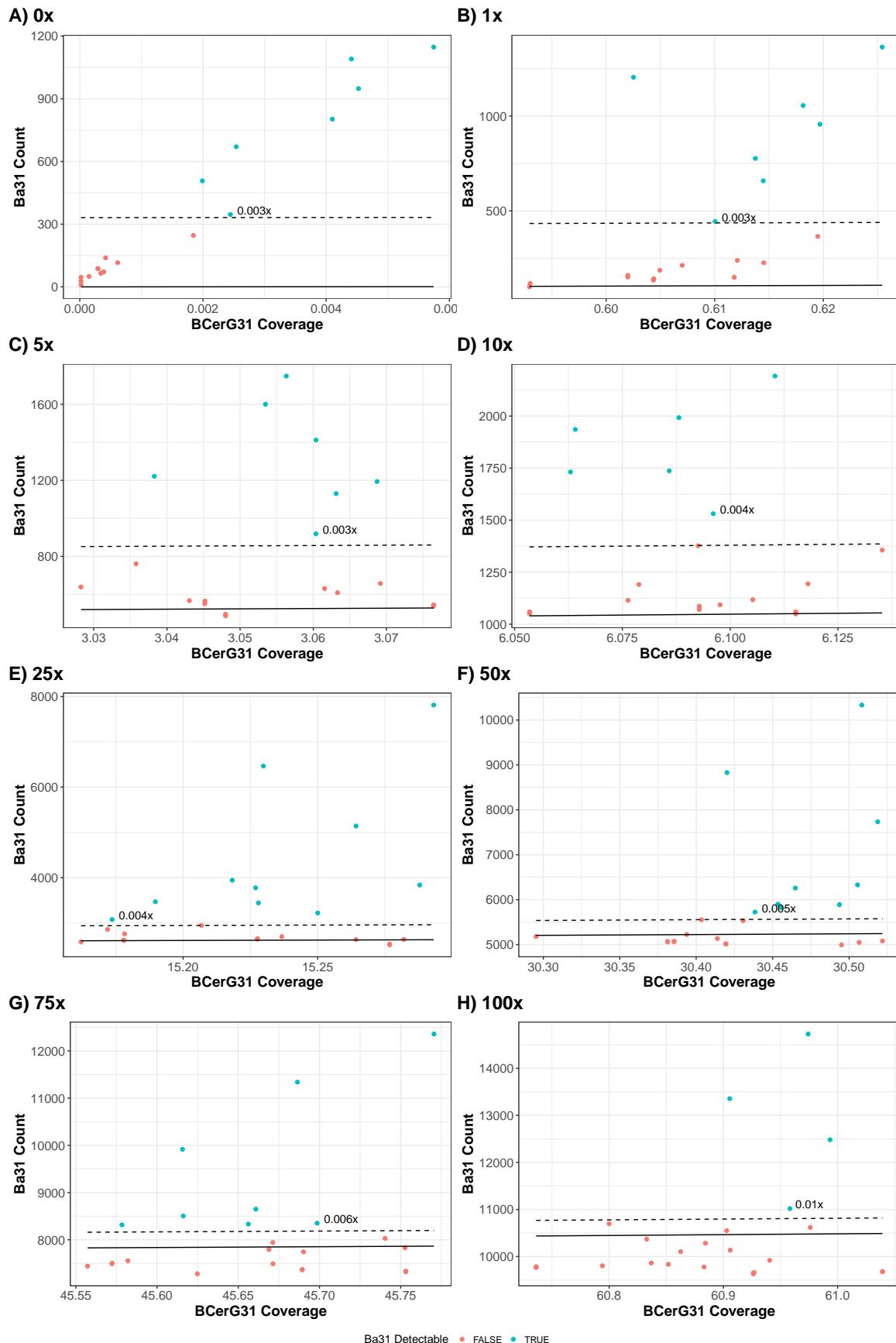
# return gtable invisibly
invisible(combined)
}

pdf(paste0(FIGURE, '/supplementary-figure-05-ba31-lod.pdf'), onefile=FALSE,
    height=12, width=8)
gridArrangeSharedBa31(p1, p2, p3, p4, p5, p6, p7, p8, ncol=2, nrow=4)
dev_null <- dev.off()

png(paste0(FIGURE, '/supplementary-figure-05-ba31-lod.png'), height=1200, width=800)
gridArrangeSharedBa31(p1, p2, p3, p4, p5, p6, p7, p8, ncol=2, nrow=4)
dev_null <- dev.off()

gridArrangeSharedBa31(p1, p2, p3, p4, p5, p6, p7, p8, ncol=2, nrow=4)

```



```

remove(p1)
remove(p2)
remove(p3)
remove(p4)
remove(p5)
remove(p6)
remove(p7)
remove(p8)

```

NYC Subway

Inputs

```

nyc_samples = read.table(
  gzipfile(paste0(PATH, "/results/nyc/nyc-ba-summary.txt.gz")),
  header=TRUE, sep="\t"
)

temp_df = read.table(
  gzipfile(paste0(PATH, "/results/nyc/nyc-bcg-summary.txt.gz")),
  header=TRUE, sep="\t"
)

nyc_samples = merge(
  nyc_samples,
  data.frame(
    run=temp_df$run,
    bcerg31_coverage=temp_df$kmer_cov_mean,
    bcerg31_hit=temp_df$hit
  ),
  by=c('run')
)

temp_df = read.table(
  gzipfile(paste0(PATH, "/results/nyc/nyc-lef-summary.txt.gz")),
  header=TRUE, sep="\t"
)

nyc_samples = merge(
  nyc_samples,
  data.frame(
    run=temp_df$run,
    lef31_hit=temp_df$hit
  ),
  by=c('run')
)

remove(temp_df)
names(nyc_samples)[names(nyc_samples) == 'hit'] <- 'ba31_hits'
names(nyc_samples)[names(nyc_samples) == 'kmer_cov_mean'] <- 'ba31_coverage'
colnames(nyc_samples)

```

```

## [1] "run"                      "is_bcg"
## [3] "is_ba"                     "has_lethal"
## [5] "group"                     "total_kmers"
## [7] "ba31_hits"                 "miss"
## [9] "kmer_cov_min"              "ba31_coverage"
## [11] "kmer_cov_median"           "kmer_cov_max"
## [13] "non_zero_kmer_cov_min"    "non_zero_kmer_cov_mean"
## [15] "non_zero_kmer_cov_median" "non_zero_kmer_cov_max"
## [17] "bcerg31_coverage"         "bcerg31_hit"
## [19] "lef31_hit"

```

Summary of NYC samples

```

nyc_summary <- nyc_samples[,c("run", "bcerg31_coverage", "ba31_hits", "lef31_hit")]
nyc_summary$model_fit <- predict_false_positive(bcg_model, nyc_summary$bcerg31_coverage)$fit.fit
nyc_summary$model_99pi <- predict_false_positive(bcg_model, nyc_summary$bcerg31_coverage)$fit.upr
nyc_summary$exceeds_99pi <- nyc_summary$ba31_hits > nyc_summary$model_99pi
nyc_summary$gray_zone <- nyc_summary$bcerg31_coverage < 0.1
head(nyc_summary)

##          run bcerg31_coverage ba31_hits lef31_hit model_fit model_99pi
## 1 GCSS-00      0.001964058     0        0  0.3375086   331.7022
## 2 GCSS-01      0.146224099     6        0 25.1275166   356.4922
## 3 GCSS-02      0.018854954     6        0  3.2400827   334.6048
## 4 GCSS-03      0.017381911     6        0  2.9869513   334.3516
## 5 GCSS-04      0.008838260     0        0  1.5187888   332.8835
## 6 GCSS-05      0.027103997     6        0  4.6576189   336.0223
## exceeds_99pi gray_zone
## 1      FALSE    TRUE
## 2      FALSE   FALSE
## 3      FALSE    TRUE
## 4      FALSE    TRUE
## 5      FALSE    TRUE
## 6      FALSE    TRUE

write.table(
  nyc_summary,
  paste0(PATH, "/results/nyc/nyc-summary.txt"),
  sep="\t",
  quote = FALSE,
  row.names = FALSE
)

```

NYC Samples which 0 Ba31 matches

```

nrow(nyc_summary[nyc_summary$ba31_hits == 0,])
## [1] 373
nrow(nyc_summary[nyc_summary$ba31_hits > 0 & nyc_summary$exceeds_99pi == FALSE,])
## [1] 1051

```

NYC Samples which exceed 99% Prediction interval

```
nrow(nyc_summary[nyc_summary$exceeds_99pi,])  
## [1] 34
```

In the Gray Zone (lef not detectable)

```
nyc_summary[nyc_summary$exceeds_99pi & nyc_summary$gray_zone,]  
  
##          run bcerg31_coverage ba31_hits lef31_hit model_fit model_99pi  
## 689 P00738      0.002160464     425        0 0.3712595   331.736  
## exceeds_99pi gray_zone  
## 689      TRUE      TRUE
```

Outside the Gray Zone (lef should have been detected)

```
nyc_summary[nyc_summary$exceeds_99pi & nyc_summary$gray_zone == FALSE,]  
  
##          run bcerg31_coverage ba31_hits lef31_hit model_fit model_99pi  
## 107 P00139      0.7789453    793        0 133.85592   465.2206  
## 163 P00200      7.9085731   2114        0 1359.02908  1690.3942  
## 167 P00204      2.0751252    993        0 356.59473   687.9595  
## 173 P00210      2.6157321   1884        0 449.49398   780.8587  
## 176 P00213      0.2041638    423        0 35.08402    366.4487  
## 180 P00217      0.2259648    1103       0 38.83037   370.1951  
## 204 P00241      0.4130413    484        0 70.97806   402.3428  
## 251 P00294      0.4412256    430        0 75.82131   407.1860  
## 277 P00320      1.7179613    715        0 295.21879   626.5835  
## 293 P00336      0.4416184    439        0 75.88881   407.2535  
## 301 P00344      2.1282530    847        0 365.72434   697.0891  
## 304 P00348      1.6077777   3380       0 276.28455   607.6493  
## 324 P00369      0.5699696    1951       0 97.94500   429.3097  
## 359 P00404      0.5555337    1299       0 95.46431   426.8290  
## 403 P00450      0.2016105    927        0 34.64526   366.0100  
## 451 P00498      1.6161249   1602       0 277.71897   609.0837  
## 457 P00504      10.7371109   2205       0 1845.09211  2176.4576  
## 486 P00535      0.2464892   1621       0 42.35733   373.7220  
## 578 P00627      0.6937052   1040       0 119.20804   450.5727  
## 733 P00783      0.2456054   1045       0 42.20545   373.5702  
## 741 P00791      0.4188353    951        0 71.97371   403.3384  
## 916 P00970      3.2344103   1206       0 555.80919   887.1740  
## 923 P00977      0.3976235    426        0 68.32862   399.6933  
## 927 P00981      1.3215163   20079      0 227.09267   558.4574  
## 940 P00994      0.1888442    375        0 32.45145   363.8162  
## 951 P01005      0.2707454    865        0 46.52556   377.8903  
## 953 P01007      1.8512226   1610       0 318.11875   649.4835  
## 955 P01009      0.2562113    489        0 44.02800   375.3927  
## 1010 P01071     2.4528135   893        0 421.49764   752.8624  
## 1055 P01116     0.1215752    731        0 20.89178   352.2565  
## 1261 P01337     0.5111460   1840       0 87.83662   419.2013  
## 1271 P01347     0.1404301    566        0 24.13187   355.4966
```

```

## 1304 P01380      0.4721595      981      0   81.13707   412.5018
##     exceeds_99pi gray_zone
## 107      TRUE    FALSE
## 163      TRUE    FALSE
## 167      TRUE    FALSE
## 173      TRUE    FALSE
## 176      TRUE    FALSE
## 180      TRUE    FALSE
## 204      TRUE    FALSE
## 251      TRUE    FALSE
## 277      TRUE    FALSE
## 293      TRUE    FALSE
## 301      TRUE    FALSE
## 304      TRUE    FALSE
## 324      TRUE    FALSE
## 359      TRUE    FALSE
## 403      TRUE    FALSE
## 451      TRUE    FALSE
## 457      TRUE    FALSE
## 486      TRUE    FALSE
## 578      TRUE    FALSE
## 733      TRUE    FALSE
## 741      TRUE    FALSE
## 916      TRUE    FALSE
## 923      TRUE    FALSE
## 927      TRUE    FALSE
## 940      TRUE    FALSE
## 951      TRUE    FALSE
## 953      TRUE    FALSE
## 955      TRUE    FALSE
## 1010     TRUE    FALSE
## 1055     TRUE    FALSE
## 1261     TRUE    FALSE
## 1271     TRUE    FALSE
## 1304     TRUE    FALSE

```

Session Info

```

sessionInfo()

## R version 3.4.3 (2017-11-30)
## Platform: x86_64-pc-linux-gnu (64-bit)
## Running under: Ubuntu 16.04.2 LTS
##
## Matrix products: default
## BLAS: /usr/lib/libblas/libblas.so.3.6.0
## LAPACK: /usr/lib/lapack/liblapack.so.3.6.0
##
## locale:
## [1] LC_CTYPE=en_US.UTF-8        LC_NUMERIC=C
## [3] LC_TIME=en_US.UTF-8         LC_COLLATE=en_US.UTF-8
## [5] LC_MONETARY=en_US.UTF-8     LC_MESSAGES=en_US.UTF-8

```

```

## [7] LC_PAPER=en_US.UTF-8      LC_NAME=C
## [9] LC_ADDRESS=C              LC_TELEPHONE=C
## [11] LC_MEASUREMENT=en_US.UTF-8 LC_IDENTIFICATION=C
##
## attached base packages:
## [1] grid      stats     graphics grDevices utils     datasets  methods
## [8] base
##
## other attached packages:
## [1] bindrcpp_0.2  gridExtra_2.3  ggplot2_2.2.1  dplyr_0.7.4
##
## loaded via a namespace (and not attached):
## [1] Rcpp_0.12.15    knitr_1.20      bindr_0.1.1
## [4] magrittr_1.5    munsell_0.4.3   colorspace_1.3-2
## [7] R6_2.2.2        rlang_0.1.6    plyr_1.8.4
## [10] stringr_1.2.0   tools_3.4.3    gtable_0.2.0
## [13] htmltools_0.3.6 lazyeval_0.2.1   yaml_2.1.18
## [16] assertthat_0.2.0 rprojroot_1.3-2  digest_0.6.15
## [19] tibble_1.4.2    RColorBrewer_1.1-2 glue_1.2.0
## [22] evaluate_0.10.1 rmarkdown_1.9    labeling_0.3
## [25] stringi_1.1.6   compiler_3.4.3  pillar_1.1.0
## [28] scales_0.5.0    backports_1.1.2  pkgconfig_2.0.1

```