



Using The Right Tool For the Right Tree in the Forest

Gen Pan, Roche



Section 1: Right Tree in the Forest

Why Forest Plot

- ▶ Quick overview instead of potentially producing many more pages with summary tables
- ▶ Useful catalyst for timely and informed decision making on potential further analysis

Two Kinds of the Forest Plot

- ▶ Meta-Analysis for Displaying Treatment Effects
- ▶ Subgroup Analysis in Randomized Controlled Trials

Questions Being Addressed

#Meta-Analysis

- ▶ In a Meta-Analysis, results comes from different clinical trials
- ▶ To combine the results of separate analyses

Questions Being Addressed

#Subgroup Analysis

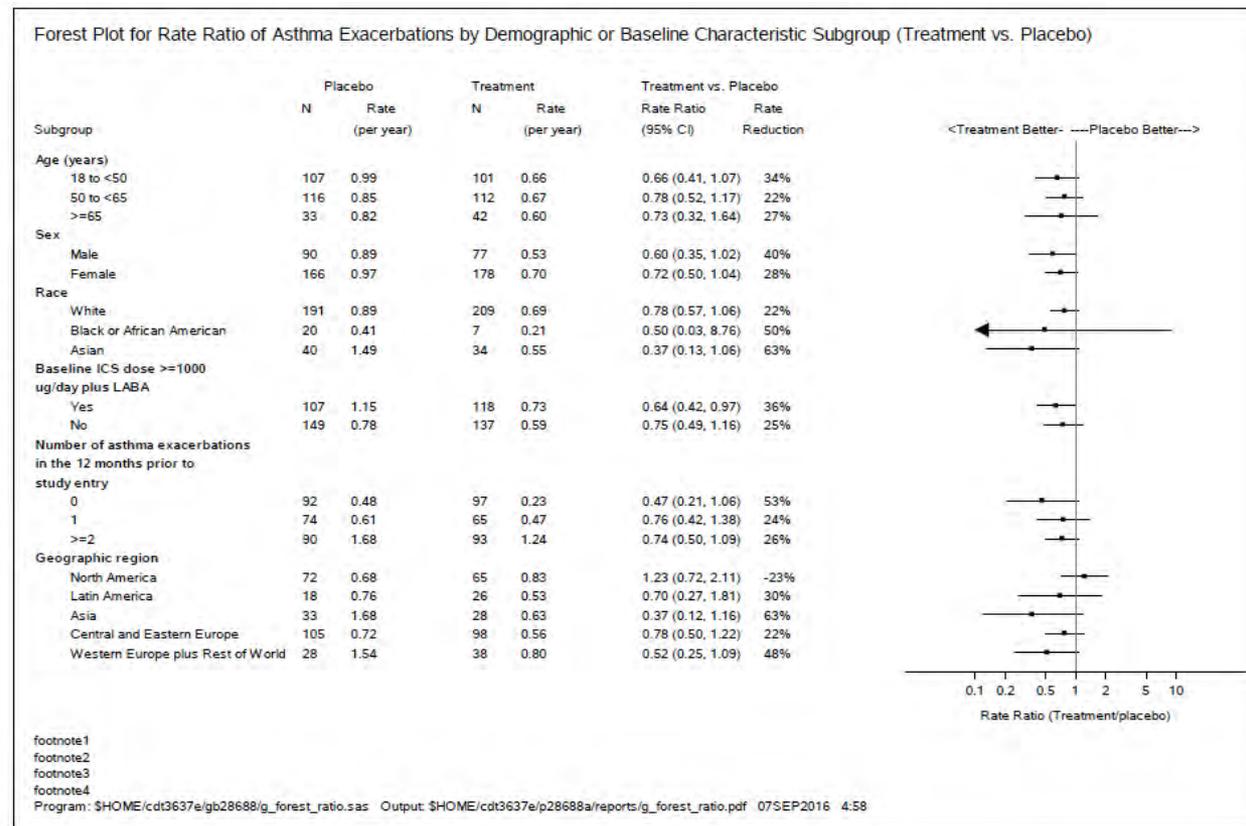
- ▶ In a Subgroup Analysis, patients are typically grouped according to common baseline characteristics
- ▶ To assess the homogeneity of a result across different patient subgroups (e.g. to demonstrate the result applies equally to all patient subgroups)
- ▶ To identify subgroups of patients in which the treatment effect differs from that of the main analysis



Section 2: Using the Right Tool

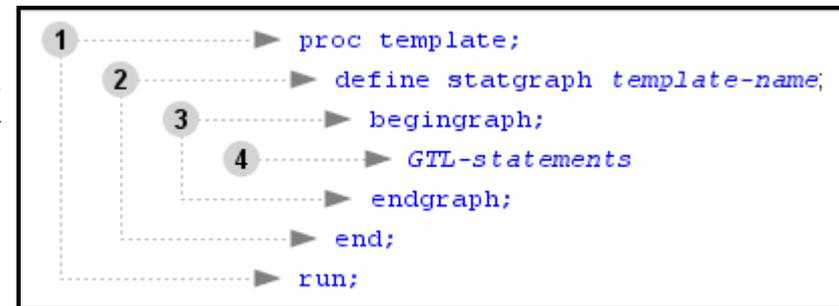
Case Study

- ▶ To create a custom Forest Plot to display Table Columns, Statistics, and Line Chart with arrow



Using the SAS 9.4 Graph Template Language (GTL)

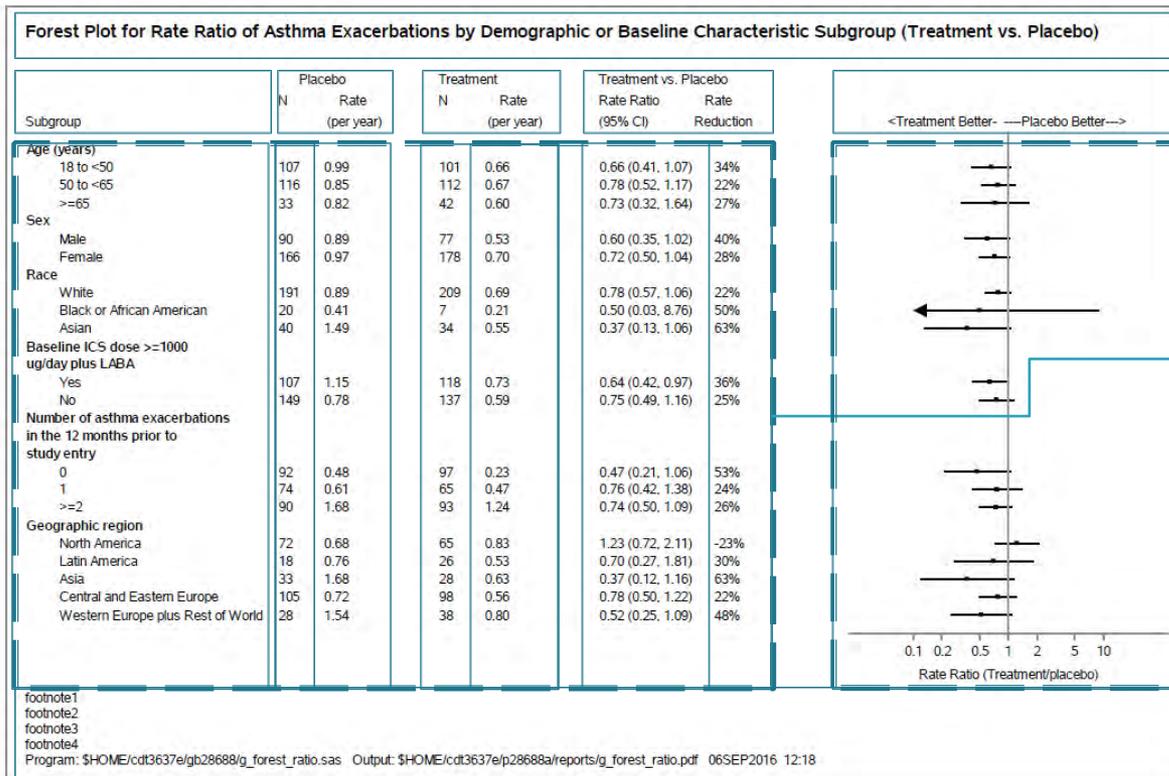
- ▶ Define the structure of the graph using the GTL syntax in a **DEFINE STATGRAPH** block in a **TEMPLATE** procedure statement
- ▶ Run the **TEMPLATE** procedure
- ▶ Run an **SGRENDER** procedure statement



```
□ proc sgrender data = data-source template = template-name;  
  run;
```

Structured Building-Block Approach

- ▶ To create a forest plot as below



Input Dataset

► What you see is what you get

	GROUP	N_P	Rate_P	N_125	Rate_125	Ratio_CI_125	Reduction_12	Ratio_N_125	Low_N_125	High_N_125	ZERO	INDENT	GRUPELVE
1	Age (years)										0	0	1
2	18 to <50	107	0.99	101	0.66	0.66 (0.41, 1.07)	34%	0.66	0.41	1.07	0	2	1
3	50 to <65	116	0.85	112	0.67	0.78 (0.52, 1.17)	22%	0.78	0.52	1.17	0	2	1
4	>=65	33	0.62	42	0.60	0.73 (0.32, 1.64)	27%	0.73	0.32	1.64	0	2	1
5											0	0	2
6	Sex										0	0	2
7	Male	90	0.89	77	0.53	0.60 (0.35, 1.02)	40%	0.6	0.35	1.02	0	2	2
8	Female	166	0.97	178	0.70	0.72 (0.50, 1.04)	28%	0.72	0.5	1.04	0	2	2
9											0	0	3
10	Race										0	0	3
11	White	191	0.89	209	0.69	0.78 (0.57, 1.06)	22%	0.78	0.57	1.06	0	2	3
12	Black or African American	20	0.41	7	0.21	0.50 (0.03, 8.76)	50%	0.5	0.03	8.76	0	2	3
13	Asian	40	1.49	34	0.55	0.37 (0.13, 1.06)	63%	0.37	0.13	1.06	0	2	3
14											0	0	4
15	Baseline ICS dose >=1000										0	0	4
16	ug/day plus LABA										0	0	4
17	Yes	107	1.15	118	0.73	0.64 (0.42, 0.97)	36%	0.64	0.42	0.97	0	2	4
18	No	149	0.78	137	0.59	0.75 (0.49, 1.16)	25%	0.75	0.49	1.16	0	2	4
19											0	0	5
20	Number of asthma exacerbations										0	0	5
21	in the 12 months prior to										0	0	5
22	study entry										0	0	5
23	0	92	0.48	97	0.23	0.47 (0.21, 1.06)	53%	0.47	0.21	1.06	0	2	5
24	1	74	0.61	65	0.47	0.76 (0.42, 1.38)	24%	0.76	0.42	1.38	0	2	5
25	>=2	90	1.68	93	1.24	0.74 (0.50, 1.09)	26%	0.74	0.5	1.09	0	2	5
26											0	0	6
27	Geographic region										0	0	6
28	North America	72	0.68	65	0.83	1.23 (0.72, 2.11)	-23%	1.23	0.72	2.11	0	2	6
29	Latin America	18	0.76	26	0.53	0.70 (0.27, 1.81)	30%	0.7	0.27	1.81	0	2	6
30	Asia	33	1.68	28	0.63	0.37 (0.12, 1.16)	63%	0.37	0.12	1.16	0	2	6
31	Central and Eastern Europe	105	0.72	98	0.56	0.78 (0.50, 1.22)	22%	0.78	0.5	1.22	0	2	6
32	Western Europe plus Rest of World	28	1.54	38	0.80	0.52 (0.25, 1.09)	48%	0.52	0.25	1.09	0	2	6

Some of the Code

- ▶ Add Error Bars
 - By using **ScatterPlot** Statement with **xErrorLower** and **xErrorUpper**
- ▶ Replace Error bars with caps for outlier value
 - Pre-processing for the input dataset
 - By using **HighLowPlot** Statement with **HighCap** and **LowCap**

```
layout overlay /      xaxisopts =( type=log label='Rate Ratio (Treatment/placebo)' labelattrs = (size = 7)
                        tickvalueattrs=(size = 7) offsetmin = 0.2 offsetmax = 0.2
                        logopts=(tickintervalstyle = linear minorticks=true tickvaluepriority=true
                        viewmin = 0.1 viewmax = 10 tickvaluelist = (0.1 0.2 0.5 1 2 5 10) )
                        )
yaxisopts = (reverse=true display=none) walldisplay=none;
highlowplot y=GROUP high=HIGH_N_125 low=LOW_N_125 /
            outlineattrs = (pattern=solid)
            fillattrs = (color = black)
            lineattrs = (color = black)
            type=line
            highcap = CAP_H lowcap = CAP_L;

scatterplot y=group x=Ratio_N_125 /
            markerattrs=(color=black symbol=squarefilled) ;
referenceline x=1;
endlayout;
```

Conclusions

- ▶ Time–Cost
 - Do not need to know anything about templates to create statistical graphics
 - With just a little knowledge of the graph and style template languages
- ▶ Users' Experience
 - Easy to use
 - Easy to modify based on stakeholders' comments
- ▶ Traceability
 - What you see is what you get

References:

- ▶ How to interpret the sample forest plot
- ▶ What is an effect size?

Center For Evidence-Based Intervention
Department of Social Policy and Intervention
University of OXFORD

- ▶ **Forest plots and the interpretation of subgroups**
Jack Cuzicka, Cancer Research UK Centre for Epidemiology,
Mathematics and Statistics, Wolfson Institute of Preventive Medicine,
Charterhouse Square, London EC1M 6BQ, UK, The Lancet, Volume
365, Issue 9467, 9–15 April 2005, Pages 1308