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1) mean = 10.968 trimmed mean = 9.667 with K(\alpha)=45-45(.1)-(45(.1)+1)-1=.1 This suggests that there are some extreme values in the top 10 percent that are pulling the mean up 2) Weibull(\gamma,\alpha)=\frac{\gamma}{\alpha}(\frac{x}{\alpha})^{\gamma-1}e^{-(x/\alpha)^{\gamma}} parameter estimates: \gamma=1.132, \alpha=11.498
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shape scale
1.1326673 11.4982087
(0.1317455) (1.6007013)
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3) Assuming Weibull(\gamma=1.1326673,\alpha=11.4982087) P(x > 30) = 0.0517
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4)

[1] "Distribution Free Mean: 10.968 Distribution Free Sigma: 9.837"

5)

[1] "Distribution Free Median: 7.97 Distribution Free IQR: 15.22"

6)

[1] "Small Mean: 6.886 Small SD: 5.46"

7)

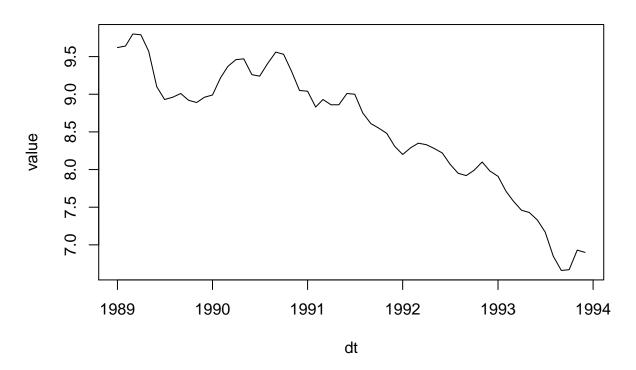
[1] "Small Mean: 5 Small MAD: 5.234"

- 8) Since both distributions are right skewed, median and MAD do the best job at describing the center and spread of the distributions.
- 9) Large litters on average have higher brain_wt/body_wt based on the centeroid measurements like median and mean, but large litters also have a much larger variation in brain_wt/body_wt where as the small litters brain_wt/body_wt mass is more concentrated.

11.

1)

Daily Yields of Moody AAA bonds



2) You can conclude that there is a strong positive correlation between adjacent values

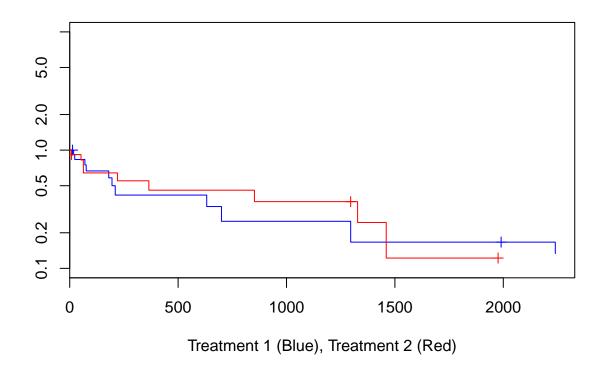
Autocorrelations of series 'ts\$value', by lag

3) No, it is easy so see that the mean has been declining over the course of time.

III.

1)

Call: survfit(formula = Surv(G1\$Time, G1\$Status) ~ 1, conf.type = "log-log") records n.max n.start events *rmean *se(rmean) 13 13 657 229 13 11 0.95LCL 0.95UCL median 202 23 1296 * restricted mean with upper limit = 2240 Call: survfit(formula = Surv(G1\$Time, G1\$Status) ~ 1, conf.type = "log-log") time n.risk n.event survival std.err lower 95% CI upper 95% CI 18 1 0.917 0.0798 0.5390 0.988 12 23 11 1 0.833 0.1076 0.4817 0.956 70 10 0.750 0.1250 1 0.4084 0.912 76 9 1 0.667 0.1361 0.3370 0.860 180 0.583 0.1423 8 0.2701 0.801 1 195 7 0.500 0.1443 0.2085 0.736 1 210 6 1 0.417 0.1423 0.1525 0.665 632 5 0.333 0.1361 0.1027 0.588 1 700 4 1 0.250 0.1250 0.0601 0.505 1296 3 0.167 0.1076 0.0265 0.413 1 0.000 2240 1 1 NaNNA NACall: survfit(formula = Surv(G2\$Time, G2\$Status) ~ 1, conf.type = "log-log") records n.max n.start events *rmean *se(rmean) 12 9 731 216 12 12 0.95LCL 0.95UCL median 1460 * restricted mean with upper limit = 1976 Call: survfit(formula = Surv(G2\$Time, G2\$Status) ~ 1, conf.type = "log-log") time n.risk n.event survival std.err lower 95% CI upper 95% CI 8 12 1 0.917 0.0798 0.53898 0.988 52 10 0.825 0.1128 0.953 1 0.46095 9 63 2 0.642 0.1441 0.30225 0.848 220 7 0.550 0.1499 1 0.23210 0.783 365 6 0.458 0.1503 0.16890 0.710 1 852 5 0.367 0.1456 0.630 1 0.11318 1328 3 0.244 0.1392 0.04456 0.528 1 1460 2 1 0.122 0.1110 0.00744 0.406



- 2) Treatment 1: Mean = 657, Median = 202; Treatment 2: Mean = 731, Median = 365
- 3) Treatment 1 seems to be more effective than treatment 2.

IV.

- 1) Type I censoring, the termination point of the experiment is fixed at 500 psi.
- 2) Left censored, we have complete data because all of the puppies already knew how to swim.
- 3) Uncensored, all puppies were able to learn how to swim so all data could be rec
- 4) Random censoring, all puppies in this group failed to learn to swim during the time of the study
- V. In this case SIQR = IQR/2

$$r = \tilde{\mu} - Q(.25)$$

$$= Q(.75) - \tilde{\mu}$$

$$SIQR = \frac{r + \tilde{\mu} - \tilde{\mu} - r}{2}$$

$$= r$$

$$\begin{split} P\Big[Q(.25) &\leq y \leq Q(.75)\Big] = .5 \\ P\Big[\tilde{\mu} - r \leq y \leq \tilde{\mu} + r\Big] = .5 \\ P\Big[-r \leq (y - \tilde{\mu} \leq r)\Big] = .5 \\ P\Big[|(y - \tilde{\mu}| \leq r)\Big] = .5 \\ r = SIQR = |y - \tilde{\mu}| \end{split}$$