

# Using JMP® and R integration to Assess Inter-rater Reliability in Diagnosing Penetrating Abdominal Injuries from MDCT Radiological Imaging

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## ABSTRACT

Fleiss' kappa (in JMP's Attribute Gauge platform) using ordinal rating scales helped assess inter-rater agreement between independent radiologists who diagnosed patients with penetrating abdominal injuries.

One drawback of Fleiss' kappa is that it does not estimate inter-rater reliability well enough since it is limited to disagreement distributions between raters being treated equally. Typically, disagreements are not all alike and have different magnitudes with multiple raters.

When compared to Fleiss' kappa, Krippendorff's alpha better differentiates between rater disagreements for various sample sizes; and estimates judgments, with or without missing data, across multiple measurement scales (binary, nominal, ordinal, interval, and ratio) for multiple raters.

Currently, Krippendorff's alpha is not available in JMP, but is available in the R open-source statistical programming language. JMP connects to R via JSL to execute R commands and exchange data.

This presentation will demonstrate how JMP and R integration allows users to take advantage of the powerful capabilities in both tools. Combining JMP and R helps users gain more insight and get better analytic results. Results helped radiologists discern which imaging signs detected injuries from signs that needed improved detection training.

## INTRODUCTION

This paper describes how JMP and R integration (via the JMP Scripting language) was used to prospectively assess the degree of agreement independent radiologists had in detecting penetrating GI tract- Bowel injuries (a.k.a. PBI). GI tract-Bowel injuries have led to serious morbidity and mortalities in trauma patients with sustaining penetrating torso injuries. Triple contrast (oral, rectal, and intravenous) multi-detector computed tomography (MDCT) has been used in the past decade as the primary means of evaluating penetrating GI tract- Bowel injuries (PBI) resulting from gunshot or stab wounds. Contrast media increases visibility of internal abdominal structures in CT imaging. Few studies have reported with high accuracy that triple contrast CT predicts the need for surgical treatment of penetrating abdominal injuries.

## MATERIALS AND METHODS

The Institutional Review Board (IRB) of the University of Maryland Medical Center (UMMC) approved the prospective observational study used in this example with a waiver of informed consent. Sixteen signs have been cited in the medical literature as key signs indicating penetrating GI Tract-Bowel injuries (PBIs). Such signs include: peritoneal violation; bone and bullet fragments extending to the GI tract; visible wound track hemorrhaging; gas and inflammation near the colon, intestines, and other abdominal organs; leakage of contrast media, etc.

Working with radiologists, a 5-point ordinal scale was developed to spot signs indicating PBI. This was the first rating scale (part of six-sigma's measurement phase) developed for MDCT PBI imaging.

Each sign used a 5-point confidence scale (1-definitely absent, 2-possibly present but unlikely, 3-equivocal, 4-likely present, 5-definitely present).

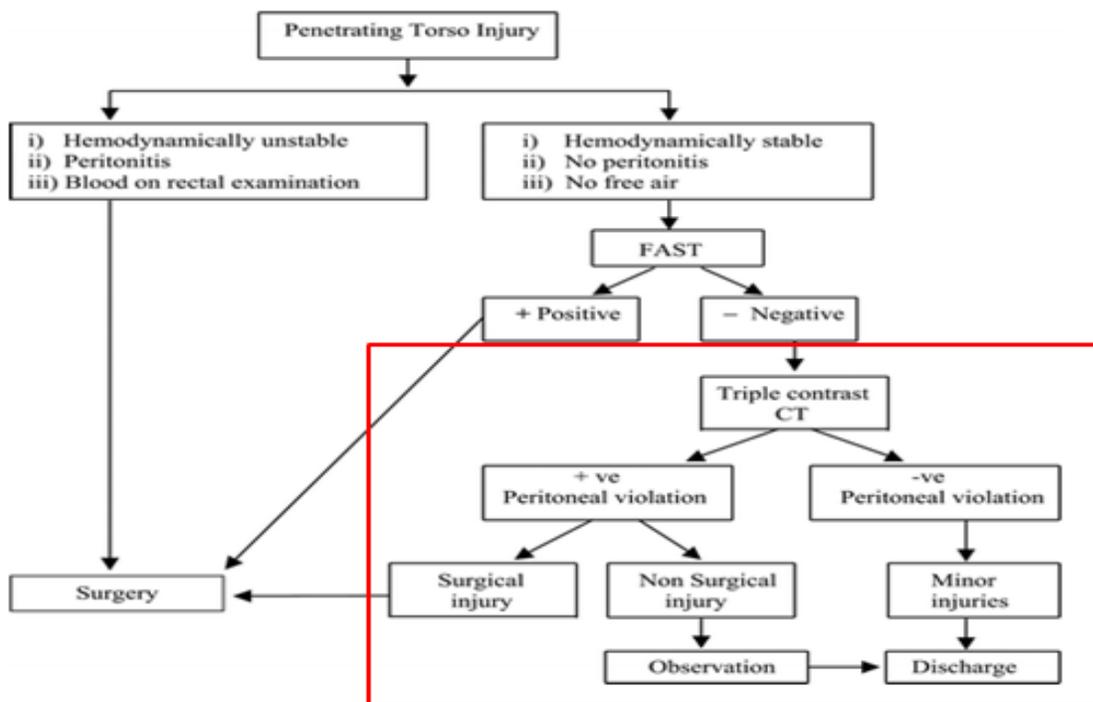
The attending trauma surgeons and radiologists (reference standard) determined the presence or absence of GI tract injury intra-operatively (1=BI injury, 0 = No BI injury) after patient admission.

All radiologists scored a binary variable indicating whether the patient required an operative management specifically for GI tract injury or not.

Figure 1 [1, p. 783] depicts the **Focused Assessment with Sonography for Trauma (FAST)** algorithm. FAST is the Rapid Sonographic, Ultrasound, MDCT Examination used by radiologists, surgeons, emergency physicians, and

certain paramedics to screen for blood around the heart or abdominal organs after trauma. The FAST protocol was applied to detect the key signs indicating penetrating GI Tract-Bowel injuries (PBIs).

**FIGURE 1: TRIPLE-CONTRAST IMAGING ALGORITHM (FAST) FOR STABLE PATIENTS WITH PENETRATING TRAUMA**  
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The FAST protocol in Figure 1 directs patient movement after they were admitted and imaged. No Peritonitis means that patients had no inflamed [peritoneum](#) (i.e., membrane lining the abdominal wall and internal tissues) nor any presence of free air or gas in the abdominal (peritoneal) cavity. The right side of the FAST path (inside the red rectangle) determines the patient treatment course options (surgery, observation, discharge) based on CT findings.

Table 1 lists the direct and indirect signs after the FAST protocol was used. A 17<sup>th</sup> overall CT diagnosis of GI injury (CToverall) was computed using the same 5-point confidence scale based on assessed combinations of the 16 signs.

**TABLE 1: 16 SIGNS CITED IN THE MEDICAL LITERATURE AS INDICATIVE OF PRESENCE OF GI INJURY.**

JMP Column	Sign	Peritoneum – Membrane (thin tissues) that lines the abdomen and inner abdominal organs
Q1	Peritoneal violation	Indirect sign of injury to the peritoneum
Q2	Retroperitoneal violation	Indirect sign of injury to retro-peritoneal colon
Q3a	Free gas at wound track	Indirect sign of free <u>intraperitoneal</u> retroperitoneal gas adjacent to the GI injury site
Q3b	Free gas away from wound track	Indirect sign of free <u>intraperitoneal</u> /retroperitoneal gas remote to the GI injury site
Q4	Intramural gas	Direct sign of intramural air
Q5	Leakage of luminal content	Direct sign bleeding into GI lumen (inside space of intestinal tubular structure)
Q6	Leakage of oral/rectal contrast	Direct Sign of leakage of enteric contrast material
Q7	Discontinuity of bowel wall	Direct sign of GI
Q8	Bowel wall thickening	Direct sign of peritoneal thickening or enhancement
Q9	Free fluid	Indirect sign of free <u>intraperitoneal</u> fluid
Q10	Mesenteric hematoma	Indirect sign of blood leakage into intestinal organs
Q11	Active mesenteric bleeding	Indirect sign of extensive hemorrhaging into intestinal organs
Q12	Signs of peritonitis	Indirect sign of peritoneum inflammation
Q13	Signs of solid organ injury	Indirect sign of injuries to other abdominal organs
Q14	Intraluminal bleeding	Direct sign of bleeding into GI lumen
Q15	Wound track extending up to bowel	Direct sign of visible penetrating wound track hemorrhage, air, and/or ballistic fragments that extended up to the GI wall



JMP's Attribute Gauge Platform [2] allowed comparisons: between raters; within each other; of raters vs. a standard; and across the scales. Kappa [3, 4] measured the degree of agreement by different raters beyond chance agreement over the rating scale. Krippendorff's alpha [5-14] is a coefficient of reliability that measures the extent of agreement among different raters. It is distinct among other similar measures of agreement (Cohen and Fleiss' kappas, Scott's pi, Intraclass Correlation Coefficient, Kendall's Coefficient of Concordance, etc.) in that it applies to any number of observers, and generalizes across scales of measurement (binary, nominal, ordinal, interval, ratio) with or without missing data.

Krippendorff alpha provided a more generalized measure of inter-rater reliability than kappa. Kappa treats disagreement distributions equally among different raters. Krippendorff alpha takes the degree of disagreement among different raters into account more generally for many types of measurement scales. For example, suppose three ratings were assigned to cases, like (+1), neutral (0), and dislike (-1). The degree of difference between two raters of like (+1) vs. neutral (0) or dislike (-1) vs. neutral (0) would be taken into account the same way as paired ratings of like (+1) vs. dislike (-1) using kappa.

Krippendorff alpha would treat the like (+1) vs. dislike (-1) more severely than like (+1) vs. neutral (0) or dislike (-1) vs. neutral (0) classification by two or more raters. This is partly due to the fact that kappa treats such classifications strictly as nominal. On the other hand, Krippendorff alpha discerns the measurement classifications as ordinal, especially when multiple raters have different magnitudes of disagreements or there are incomplete data. Different formulas are used to compute alphas, depending upon the nominal, ordinal, interval, ratio (even polar and circular) metrics for the data.

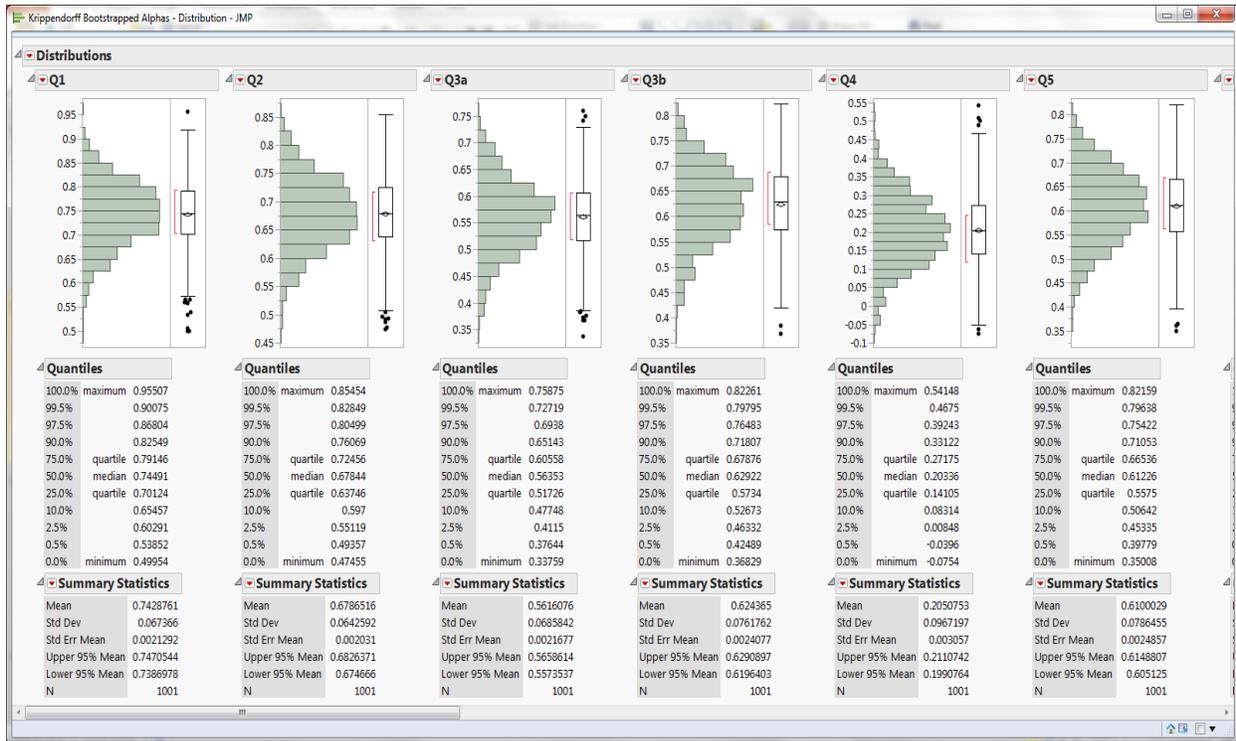
Since JMP did not compute this statistic, the kripp.alpha function in the lpSolve and irr packages of R [11] was used to obtain the metric using JMP and R integration (see Appendix 1). The bootalpha function [12] produced 95% confidence intervals of krippendorff alphas from 1000 bootstrapped resamples for each sign (Table 3).

**TABLE 3: SELECTED SIGNS OF 1001 KRIPPENDORFF'S ALPHAS (FIRST ROW IS ARE THE KRIPPENDORFF ALPHA VALUES FOR EACH SIGN OF THE ORIGINAL, SOURCE DATA)**

	Q1	Q2	Q3a	Q3b	Q4	Q5	Q6	Q7	Q8
1	0.7503407205	0.6863852424	0.5696314041	0.6282646652	0.2139495116	0.6155022449	0.7821915078	0.4424166103	0.477530414
2	0.7210360746	0.7859200824	0.5046050838	0.6056230326	0.1947785763	0.7059581525	0.8079280009	0.4604220466	0.389775604
3	0.5720361086	0.6482426264	0.514438717	0.6588734576	0.1879729034	0.5646124566	0.7947567094	0.1381087392	0.538654539
4	0.7506966294	0.6023449735	0.6158203202	0.6521212777	0.2025975456	0.5054917158	0.5698358853	0.3346991954	0.438355545
5	0.7715459011	0.6272030508	0.6195849086	0.6861634784	0.1673904465	0.5048999755	0.8252894444	0.5993458945	0.528086890
6	0.6280803851	0.5600545567	0.6716101968	0.6711610727	0.3420326486	0.539202459	0.5018901591	0.4802878074	0.72450041
7	0.7732603879	0.6761144729	0.5274906005	0.7066767284	0.1077347	0.7335028986	0.7750818821	0.3739184774	0.419341013
8	0.7338822678	0.6876724506	0.5193174203	0.6429510506	0.1451848429	0.5748235679	0.8108623581	0.4449492196	0.364786389
9	0.6769203931	0.6227643765	0.7497746122	0.6337941785	0.2846360342	0.457483872	0.8227957925	0.3241417287	0.361501689
10	0.7235056939	0.768018476	0.6630945567	0.7272999747	0.2305584332	0.5848968593	0.7870634607	0.3672754197	0.429156673
11	0.7605945339	0.6635798092	0.5727918574	0.6622085244	0.2106267704	0.6614732251	0.8006302235	0.4187850251	0.307332009
12	0.7708749225	0.682339275	0.5350938842	0.5976630835	0.1321558467	0.5854656389	0.7604468865	0.3719499532	0.589711358
13	0.6372039997	0.6426098698	0.5718887318	0.6707687996	0.1458339579	0.5999384779	0.7190363121	0.4021871538	0.545018387
14	0.7640478029	0.6141551123	0.5221268047	0.7026533601	0.1616578682	0.7105800187	0.9514525139	0.5125407724	0.40452465
15	0.7600176898	0.6494131183	0.6084140732	0.7979545441	0.180615877	0.6792807285	0.8240211237	0.4751102005	0.606648207
16	0.5807408431	0.7718705223	0.6321302269	0.5778995766	0.2933670118	0.6802872078	0.7666736916	0.3665353209	0.487058037
17	0.7805272207	0.6730492708	0.6605950713	0.5793462048	0.3540457742	0.6750115674	0.7197741931	0.4351805292	0.497322298
18	0.6507940291	0.7230874432	0.5974847323	0.444380781	0.2207898559	0.6709899714	0.6418147685	0.5783670043	0.409557003
19	0.7419538518	0.7715424138	0.53844472799	0.6425875489	0.2165720164	0.6810537925	0.7496650449	0.5140910758	0.375117370
20	0.7616445982	0.7040156087	0.4802663329	0.7130578605	0.2429206431	0.7159862525	0.7384735901	0.5340342135	0.508478506
21	0.6912089883	0.5453255004	0.4940759801	0.600797481	0.124466469	0.6323482586	0.7156257938	0.5262515436	0.453879355

Figure 3 shows the bootstrapped resampling distribution of Krippendorff alphas for selected signs.

**FIGURE 3: DISTRIBUTION OF 1001 BOOTSTRAPPED KRIPPENDORFF'S ALPHAS FOR SELECTED SIGNS**



The remaining script in Appendix 1 vertically concatenated the 95% Krippendorff alpha confidence interval matrices of all signs and formed a final data table. Table 4 lists the resulting Data Table of the observed Krippendorff alpha values and the 95% confidence intervals for each sign from the original data in Table 2.

**TABLE 4: KRIPPENDORFF'S ALPHA DATA TABLE WITH 95% CONFIDENCE INTERVALS COMPUTED FROM 1000 BOOTSTRAPPED RESAMPLES**

Sign	Alpha	Standard Error of the median	Lower 95% Alpha	Upper 95% Alpha
1 Q1	0.7503407205	0.0673660191	0.605123872	0.8678345655
2 Q2	0.6863852424	0.0642591695	0.5515136157	0.8042550042
3 Q3a	0.5696314041	0.068584244	0.4117809287	0.693727753
4 Q3b	0.6282646652	0.0761761889	0.4640509678	0.7631710414
5 Q4	0.2139495116	0.0967197495	0.0133032772	0.3909735248
6 Q5	0.6155022449	0.0786455012	0.4539879004	0.7533003295
7 Q6	0.7821915078	0.1089618458	0.5092853062	0.9356726452
8 Q7	0.4424166103	0.0939232452	0.237861742	0.6028726855
9 Q8	0.4775304141	0.0769103174	0.3276420937	0.6268635406
10 Q9	0.6790435774	0.0637603594	0.5384322913	0.7895230257
11 Q10	0.4134978718	0.0818086061	0.2397866828	0.5604381094
12 Q11	0.4477476322	0.1623261924	0.0834909273	0.7251071094
13 Q12	-0.019908968	0.0650933494	-0.038277248	-0.002173913
14 Q13	0.8223807001	0.047156654	0.720483304	0.8992787812
15 Q14	0.1285898652	0.1073809241	-0.088778552	0.3285515847
16 Q15	0.5571417835	0.0715141086	0.4083534376	0.6852612036
17 Ctoverall	0.792743252	0.0391764202	0.6985028447	0.8509944042

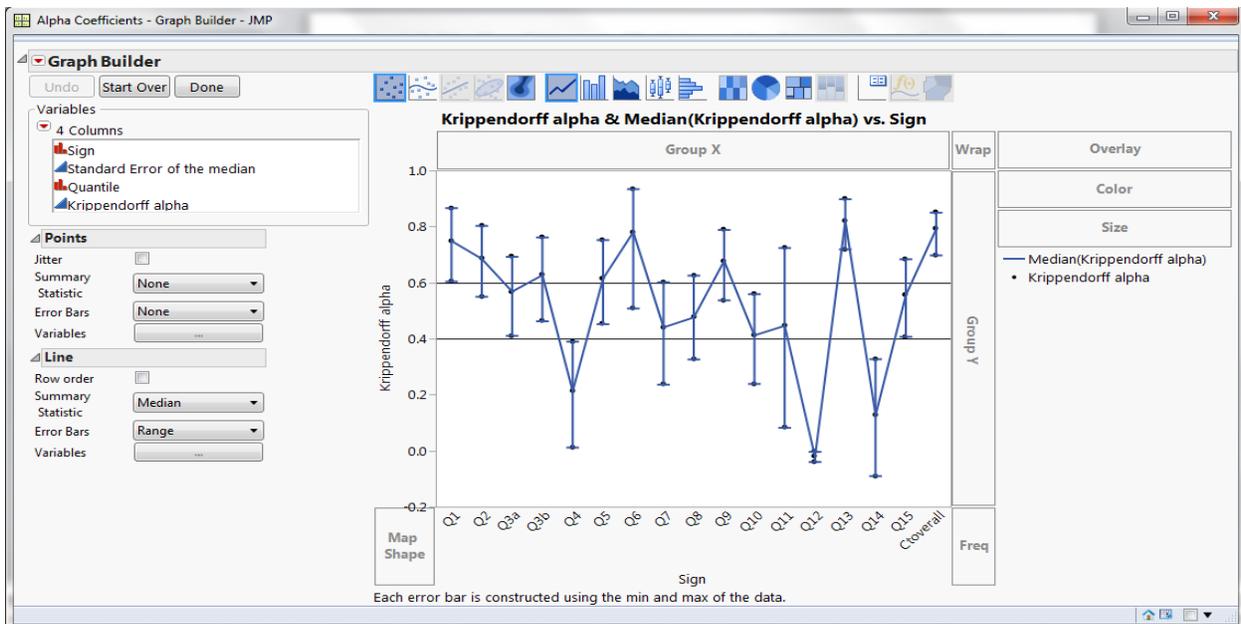
Table 5 shows the stacked columns of Table 4 that was used to prepare Figure 4.

**TABLE 5: STACKED COLUMN OF KRIPPENDORFF'S ALPHAS (COEFFICIENT AND 95% CONFIDENCE INTERVAL VALUES)**

Sign	Standard Error of the median	Quantile	Krippendorff alpha
1 Q1	0.0673660191	Alpha	0.7503407205
2 Q1	0.0673660191	Lower 95% Alpha	0.605123872
3 Q1	0.0673660191	Upper 95% Alpha	0.8678345655
4 Q2	0.0642591695	Alpha	0.6863852424
5 Q2	0.0642591695	Lower 95% Alpha	0.5515136157
6 Q2	0.0642591695	Upper 95% Alpha	0.8042550042
7 Q3a	0.068584244	Alpha	0.5696314041
8 Q3a	0.068584244	Lower 95% Alpha	0.4117809287
9 Q3a	0.068584244	Upper 95% Alpha	0.693727753
10 Q3b	0.0761761889	Alpha	0.6282646652
11 Q3b	0.0761761889	Lower 95% Alpha	0.4640509678
12 Q3b	0.0761761889	Upper 95% Alpha	0.7631710414
13 Q4	0.0967197495	Alpha	0.2139495116
14 Q4	0.0967197495	Lower 95% Alpha	0.0133032772
15 Q4	0.0967197495	Upper 95% Alpha	0.3909735248
16 Q5	0.0786455012	Alpha	0.6155022449
17 Q5	0.0786455012	Lower 95% Alpha	0.4539879004
18 Q5	0.0786455012	Upper 95% Alpha	0.7533003295
19 Q6	0.1089618458	Alpha	0.7821915078
20 Q6	0.1089618458	Lower 95% Alpha	0.5092853062
21 Q6	0.1089618458	Upper 95% Alpha	0.9356726452

Figure 4 is a Graph-Builder plot of the Krippendorff alpha 95% Confidence Intervals. Alpha values above 0.8 (Q13 and Coverall) indicated good reliability in detection between raters. Three Signs (Q4, Q12, and Q14) with confidence intervals below 0.4 had poor reliability. These signs indicated the need for more medical training so that radiologists can improve their ability to detect them as indicators of penetrating bowel injuries.

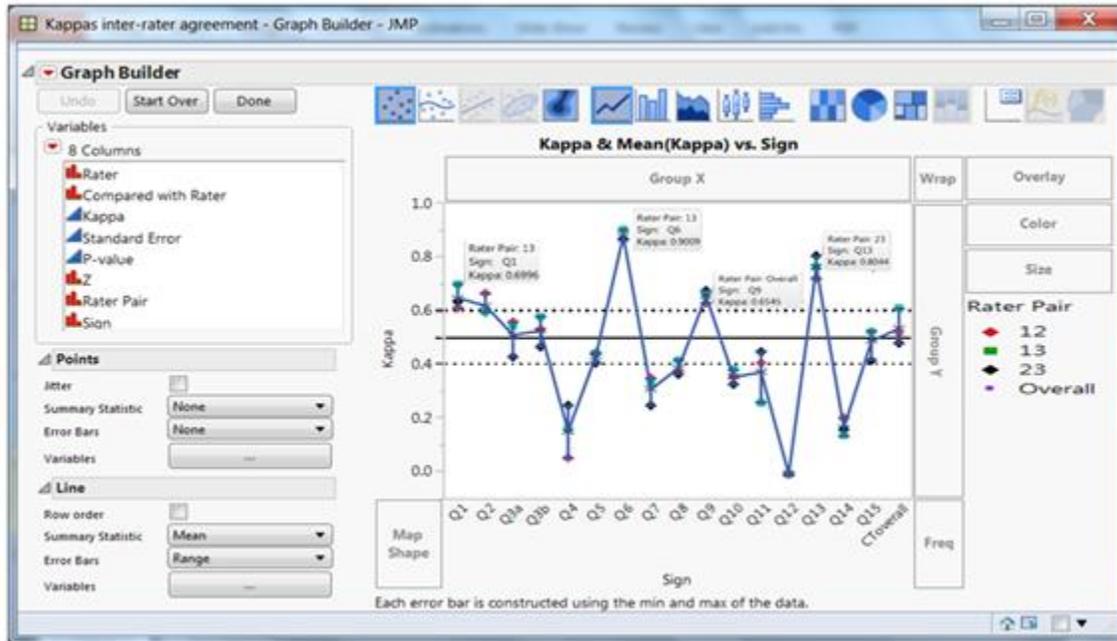
**FIGURE 4: GRAPH-BUILDER PLOT OF KRIPPENDORFF'S ALPHA AND CONFIDENCE INTERVALS FOR EACH SIGN**



The Kappa values above 0.6 (Q1, Q2, Q6, Q9, and Q13) in Figure 5 indicate good agreement between raters. Five Signs (Q3a, Q3b, Q5, Q15, and Coverall) had moderate kappa values between 0.4-0.6. Four signs (Q7, Q8, Q10,

and Q11) had fair agreement with kappa values between 0.2-0.4. Signs Q4, Q12, and Q14 had poor agreement. Like the Krippendorff alpha, the fair-to-poor kappa values (below 0.2) suggested the need for increased medical training in detecting these signs in order to improve the measurement system capability of the rating scales.

FIGURE 5: GRAPH-BUILDER PLOT OF PAIRED RATERS AND OVERALL KAPPA VALUES FOR EACH SIGN



Next, I joined the Fleiss overall kappa statistics with the Krippendorff alpha data table as shown in Figure 6 to form Table 6 and Figure 7.

FIGURE 6: OPENED FLEISS KAPPA AND KRIPPENDORFF ALPHA DATA TABLES WITH SCRIPT TO JOIN THEM

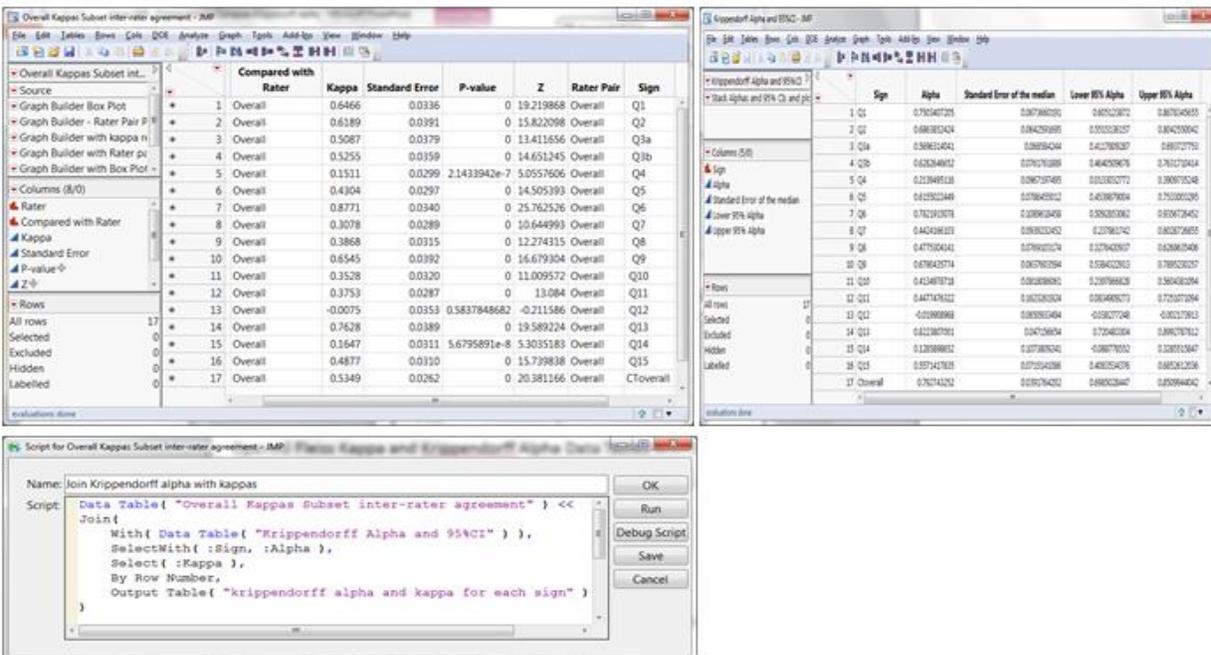
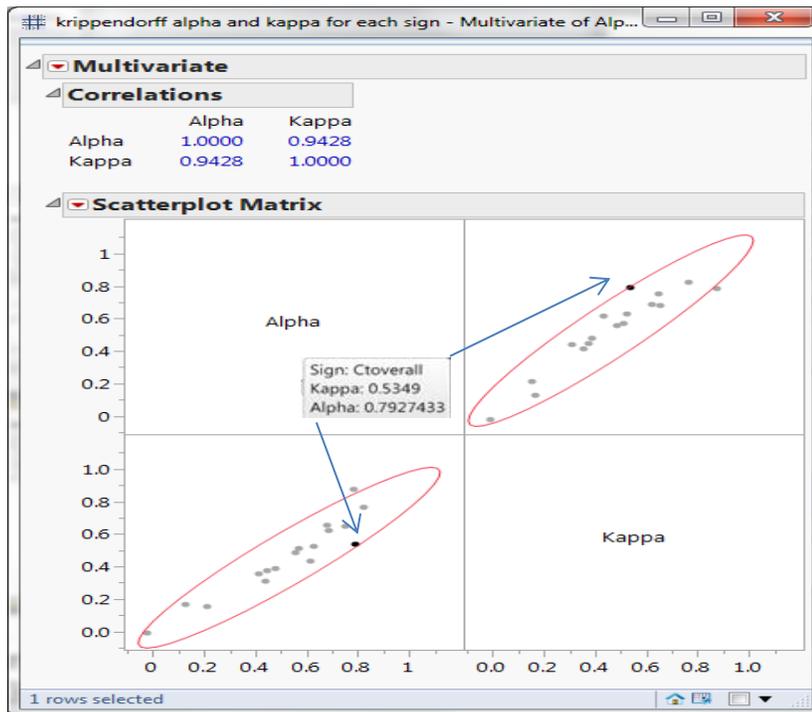


TABLE 6: KRIPPENDORFF ALPHA AND OVERALL KAPPA DATA TABLE

	Sign	Alpha	Kappa
1	Q1	0.7503407205	0.6466
2	Q2	0.6863852424	0.6189
3	Q3a	0.5696314041	0.5087
4	Q3b	0.6282646652	0.5255
5	Q4	0.2139495116	0.1511
6	Q5	0.6155022449	0.4304
7	Q6	0.7821915078	0.8771
8	Q7	0.4424166103	0.3078
9	Q8	0.4775304141	0.3868
10	Q9	0.6790435774	0.6545
11	Q10	0.4134978718	0.3528
12	Q11	0.4477476322	0.3753
13	Q12	-0.019908968	-0.0075
14	Q13	0.8223807001	0.7628
15	Q14	0.1285898652	0.1647
16	Q15	0.5571417835	0.4877
17	Ctoverall	0.792743252	0.5349

The Scatterplot Matrix density ellipses of Figure 7 showed the strong positive correlation (0.9428) between the overall Fleiss' kappas and Krippendorff alphas across all signs.

FIGURE 7: MULTIVARIATE CORRELATIONS BETWEEN THE KRIPPENDORFF ALPHA AND OVERALL FLEISS' KAPPA STATISTICS



## CONCLUSION

This presentation showed how JMP's JSL with R integration computed a Krippendorff's alpha inter-rater reliability statistic that complements Fleiss' kappa in JMP. Although Krippendorff alpha is not as widely used and is more computationally complex than Fleiss' kappa, it has gained more acceptance by researchers in content analysis and measurement reproducibility studies as a robust, flexible, inter-coder reliability metric. Combining JMP's dynamic interactivity with R's unique functionalities and packages gives users the ability to develop custom applications that can be implemented in JMP.

## REFERENCES

- [1]. Shanmuganathan, K., Mirvis, S.E., Chiu, W.C., Killeen, K.L., Hogan, G.J.F., Scalea, T.M. (2004), "Penetrating Torso Trauma: Triple-Contrast Helical CT in Peritoneal Violation and Organ Injury—A Prospective Study in 200 Patients," *Radiology*, 231, 775-784.
- [2]. SAS Institute Inc. (2014). "Attribute Gauge Charts," *JMP® 11 Quality and Process Methods*. Cary, NC: SAS Institute Inc., pp. 193-205.
- [3]. De Mast, J and van Wieringen, W. (2004). "Measurement System Analysis for Bounded Ordinal Data," *Quality and Reliability Engineering International*, 20, 383-395.
- [4]. Fleiss, J.L., Levin, B., and Paik, M.C. (2003), *Statistical Methods for Rates and Proportions (3<sup>rd</sup> ed.)*. New York: Wiley.
- [5]. Wikipedia, "Krippendorff's Alpha," [http://en.wikipedia.org/wiki/Krippendorff's\\_alpha](http://en.wikipedia.org/wiki/Krippendorff's_alpha).
- [6]. Hayes, A. F., & Krippendorff, K. (2007). "Answering the Call for a Standard Reliability Measure for Coding Data," *Communication Methods and Measures*, 1, 77-89.
- [7]. Kang, N. et al. (1993), "A SAS® Macro for Calculating Inter-coder Agreement in Content Analysis," *Journal of Advertising*, 23, 17-28.
- [8]. Krippendorff, K. (2013). *Content Analysis: An Introduction to its Methodology (3rd ed.)*. Thousand Oaks, CA: Sage.
- [9]. Krippendorff, K. (2007), "Computing Krippendorff's Alpha-Reliability," <http://www.asc.upenn.edu/usr/krippendorff/mwebreliability5.pdf>.
- [10]. Gwet, K. (2011), "On the Krippendorff's Alpha Coefficient," [http://www.agreestat.com/research\\_papers/onkrippendorffalpha\\_old.pdf](http://www.agreestat.com/research_papers/onkrippendorffalpha_old.pdf).
- [11]. Gamer, M. et al., (2012) "Various Coefficients of Interrater Reliability and Agreement," <http://cran.r-project.org/web/packages/irr/irr.pdf>.
- [12]. Gruszczynski, M. (2013). "R Convenience functions to perform bootstrap resampling on inter-coder reliability data for Krippendorff's statistics," <https://github.com/MikeGruz/kripp.boot/blob/master/R/kripp.boot.R>.
- [13]. Hayes, A.F. (2011), "KALPHA SAS Macro," <http://www.afhayes.com/public/kalpha.sas>.
- [14]. Artstein, R. and Poesio, M. (2008). "Inter-Coder Agreement for Computational Linguistics," *Computational Linguistics*, 34, 555-596.

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## APPENDIX 1: JSL SCRIPT TO COMPUTE KRIPPENDORFF'S ALPHA USING JMP AND R INTEGRATION

```
/******  
This JMP and R Integration example computes Krippendorff's alpha  
reliability coefficient that measures agreement among  
different raters or measuring instruments drawing distinctions  
among unstructured phenomena or assign computable values to  
items.  
  
Alpha emerged from content analysis whenever two or more  
methods of generating data are applied to the same subjects.  
Alpha answers the question of how much the resulting data  
can be trusted to represent something real.  
  
The R implementation uses the lpSolve and irr library packages  
with the kripp.alpha function with ordinal measures.  
  
Visit http://en.wikipedia.org/wiki/Krippendorff's\_alpha  
for more information.  
  
Author: Melvin Alexander  
Version: 1, 12Feb14 - Initial coding  
Updates:  
  
*****/  
  
R Init() ;  
//Open Data Table with Response variable (BI) and Signs (Explanatory variables)  
//pbi 11 for Mel July_02_2013 contains all data  
dt = Open("G:\Nitima\Surgical Subset of pbi 11 for Mel July_02_2013.jmp",  
Select Columns("No", "Readernumber", "BI", "Q1","Q2", "Q3a", "Q3b", "Q4",  
"Q5", "Q6", "Q7", "Q8", "Q9", "Q10", "Q11", "Q12", "Q13", "Q14", "Q15", "Ctoverall" )  
) << New Column( "Numeric_BI",  
Nominal,  
Format( "Best", 12 ),  
Formula( :BI == "Y" ));  
  
// Form column vectors of signs (Q1-Ctoverall)  
col=Column("Q1");  
q1mtx = col<<GetAsMatrix;  
dt=Current Data Table() ;  
q2mtx=Column("Q2")<<GetAsMatrix;  
dt=Current Data Table();  
  
< Code Removed for other column vectors >  
Dt=Current Data Table();  
qoallmtx=Column("Ctoverall")<<GetAsMatrix;  
  
//Send JMP vectors to R variables and compute Krippendorff alphas  
R Send(q1mtx);  
R Submit( "\[  
# load lpSolve and irr packages  
library (lpSolve)  
library (irr)
```

```

Q1<-matrix(q1mtx,nrow=3)
Q1alpha <- kripp.alpha(Q1,"ordinal")
# Mel Alexander's adaptation of Mike Gruscynski's kripp.boot.R code
# Reference: https://github.com/MikeGruz/kripp.boot/blob/master/R/kripp.boot.R
# bootalpha generates bootstrap resamples of krippendorff alphas several times
(default iter=100)

bootalpha <- function(x, raters='rows', probs=c(0.025,0.975), iter=100,
method=c('nominal','ordinal','interval','ratio'))
{
  alphas <- numeric(iter)

  for( i in 1:iter)
  {
    alphas[i] <- kripp.alpha(x[,sample(ncol(x),
size=ncol(x),
replace=TRUE)],
method=method)$value
  }

  kripp.ci <- quantile(alphas, probs=probs, na.rm=TRUE)
  boot.stats <- list(mean.alpha=mean(alphas, na.rm=TRUE),
upper=kripp.ci[2],
lower=kripp.ci[1],
alphas=alphas,
raters=nrow(x),
iter=iter,
probs=probs,
sealpha = sqrt(var(alphas)/length(alphas)),
size=ncol(x))
  class(boot.stats) <- 'bootalpha'
  return(boot.stats)
}

Q1boot <- bootalpha(Q1, iter=1000, method='ordinal')
#Q1boot
# combine the original alpha (Q1alpha$value) with the bootstrapped alphas
(Q1boot$alphas)
Q1allalphas <- matrix(c(Q1alpha$value, Q1boot$alphas))
#Q1allalphas

#hist(allalphas)
# Get the standard error of the median
Q1.median <- sapply(Q1allalphas,median)
Q1semed <- sqrt(var(Q1.median))

# join the original alpha (Q1alpha$value) with the standard error of the median
(Q1semed),
# 95% ci alpha values (Q1boot$lower, Q1boot$upper)
Q195ci <- matrix(c(Q1alpha$value, Q1semed, Q1boot$lower, Q1boot$upper), nrow=1)
Q195ci
]\" );

R Send(q2mtx);
R Submit( "[
#library (irr)
Q2<-matrix(q2mtx,nrow=3)
Q2alpha <- kripp.alpha(Q2,"ordinal")
Q2boot <- bootalpha(Q2, iter=1000, method='ordinal')
#Q2boot
# combine the original alpha (Q2alpha$value) with the bootstrapped alphas
(Q2boot$alphas)
Q2allalphas <- matrix(c(Q2alpha$value, Q2boot$alphas))
#Q2allalphas

```

```

#hist(Q2allalphas)
# Get the standard error of the median
Q2.median <- sapply(Q2allalphas,median)
Q2semed <- sqrt(var(Q2.median))
# join the original alpha (Q2alpha$value) with the standard error of the median
(Q2semed)
# 95% ci alpha values (Q2boot$lower, Q2boot$upper)
Q295ci <- matrix(c(Q2alpha$value, Q2semed, Q2boot$lower, Q2boot$upper), nrow=1)
Q295ci
]\" );

```

< Code removed for other R Send and Submit commands >

```

/*Collect R krippendorff bootstrapped alphas as JMP variables*/
Q1allalpha = R Get (Q1allalphas);
Q2allalpha = R Get (Q2allalphas);
Q3aallalpha = R Get (Q3aallalphas);
Q3ballalpha = R Get (Q3ballalphas);
Q4allalpha = R Get (Q4allalphas);
Q5allalpha = R Get (Q5allalphas);
Q6allalpha = R Get (Q6allalphas);
Q7allalpha = R Get (Q7allalphas);
Q8allalpha = R Get (Q8allalphas);
Q9allalpha = R Get (Q9allalphas);
Q10allalpha = R Get (Q10allalphas);
Q11allalpha = R Get (Q11allalphas);
Q12allalpha = R Get (Q12allalphas);
Q13allalpha = R Get (Q13allalphas);
Q14allalpha = R Get (Q14allalphas);
Q15allalpha = R Get (Q15allalphas);
Qoallalpha = R Get (Qoallalphas);
//Collect R krippendorff alphas as JMP variables
Q1alpha95 = R Get (Q195ci) ;
Q2alpha95 = R Get (Q295ci) ;
Q3aalpha95 = R Get (Q3a95ci) ;
Q3balph95 = R Get (Q3b95ci) ;
Q4alpha95 = R Get (Q495ci) ;
Q5alpha95 = R Get (Q595ci) ;
Q6alpha95 = R Get (Q695ci) ;
Q7alpha95 = R Get (Q795ci) ;
Q8alpha95 = R Get (Q895ci) ;
Q9alpha95 = R Get (Q995ci) ;
Q10alpha95 = R Get (Q1095ci) ;
Q11alpha95 = R Get (Q1195ci) ;
Q12alpha95 = R Get (Q1295ci) ;
Q13alpha95 = R Get (Q1395ci) ;
Q14alpha95 = R Get (Q1495ci) ;
Q15alpha95 = R Get (Q1595ci) ;
Qoallalpha95 = R Get (Qoall95ci) ;

/*Send krippendorff bootstrapped alphas and cis from R to JMP matrices and data tables
*/
allalpha=Concat(Q1allalpha,Q2allalpha,Q3aallalpha,Q3ballalpha,
Q4allalpha,Q5allalpha,Q6allalpha,Q7allalpha,Q8allalpha,
Q9allalpha,Q10allalpha, Q11allalpha,Q12allalpha,Q13allalpha,
Q14allalpha,Q15allalpha,Qoallalpha );
// Vertically Concatenate the alphas and 95%ci alpha matrices
alpha95ci = VConcat(Q1alpha95, Q2alpha95,Q3aalpha95, Q3balph95, Q4alpha95, Q5alpha95,
Q6alpha95, Q7alpha95, Q8alpha95, Q9alpha95, Q10alpha95, Q11alpha95,
Q12alpha95, Q13alpha95, Q14alpha95, Q15alpha95, Qoallalpha95 );

dtallalpha = As Table(allalpha)<< Set Name("Krippendorff Bootstrapped Alphas");

```

```

Data Table ("Krippendorff Bootstrapped Alphas");
col = Col1 << Set Name("Q1");
col = Col2 << Set Name("Q2");
col = Col3 << Set Name("Q3a");
col = Col4 << Set Name("Q3b");
col = Col5 << Set Name("Q4");
col = Col6 << Set Name("Q5");
col = Col7 << Set Name("Q6");
col = Col8 << Set Name("Q7");
col = Col9 << Set Name("Q8");
col = Col10 << Set Name("Q9");
col = Col11 << Set Name("Q10");
col = Col12 << Set Name("Q11");
col = Col13 << Set Name("Q12");
col = Col14 << Set Name("Q13");
col = Col15 << Set Name("Q14");
col = Col16 << Set Name("Q15");
col = Col17 << Set Name("Ctoverall");

dt95cialpha = As Table(alpha95ci)<< Set Name("Krippendorff Alpha and 95%CI");
col = Col1 << Set Name("Alpha");
col = Col2 << Set Name("Standard Error of the median");
col = Col3 << Set Name("Lower 95% Alpha ");
col = Col4 << Set Name("Upper 95% Alpha ");

// Add Sign column to Data Table and move it to first
dt95cialpha = Data Table ("Krippendorff Alpha and 95%CI") << New Column("Sign",
Character, Values({"Q1", "Q2", "Q3a", "Q3b", "Q4",
"Q5", "Q6", "Q7", "Q8", "Q9", "Q10", "Q11", "Q12", "Q13", "Q14", "Q15", "Ctoverall"})) ;

dt95cialpha = Data Table("Krippendorff Alpha and 95%CI")
<< Get Selected Columns ("Sign")
<< Move Selected Columns({"Sign"}, To First);

// Open data table of kappas
dtkappa = Open("G:\Nitima\Kappas inter-rater agreement.jmp");
R Term( );

```