

8 March 2012

# Southeast Asphalt User-Producer Group Interlaboratory Study to Determine the Precision of AASHTO TP70 – the Multiple- Stress Creep-Recovery (MSCR) Test

Prepared for the Southeast Asphalt User-Producer Group (SEAUPG)

R. Michael Anderson, P.E.  
Asphalt Institute

in cooperation with  
the Federal Highway Administration

through  
DTFH61-08-H-00030  
and  
DTFH61-11-H-00033



## **Introduction**

In 2011, the Southeast Asphalt User-Producer Group (SEAUPG) initiated an interlaboratory study (ILS) for participating labs to evaluate the repeatability and reproducibility of AASHTO TP70, *Multiple Stress Creep Recovery (MSCR) Test of Asphalt Binder Using a Dynamic Shear Rheometer (DSR)*. This ILS was seen as a follow-up to the ILS conducted by the Asphalt Binder Expert Task Group in 2008-09 that established the initial precision statement and the ILS conducted by the Northeast Asphalt User Producer Group (NEAUPG) in 2010.

Twenty-three (23) laboratories, principally located in the SEAUPG, agreed to participate in the study. Included were Users (13 labs), Producers (8 labs), and Industry/Academia (2 labs). These labs are shown in Table 1. More detailed contact information can be found in the Appendix.

Table 1: Participating Labs

<b>Users</b>	<b>Producers</b>	<b>Industry/Academia</b>
Alabama DOT	BETA R&D	Asphalt Institute
Federal Highway Administration (FHWA) Office of Pavement Technology (3 labs/DSRs)	Ergon Asphalt & Emulsions (5 Labs)	National Center for Asphalt Technology
Florida DOT	Marathon	
Georgia DOT	Paragon Technical Services	
Louisiana Transportation Research Center		
Mississippi DOT		
North Carolina DOT		
Oklahoma DOT		
South Carolina DOT		
Texas DOT		
Virginia Center for Transportation Innovation and Research		

Each laboratory was asked to perform a temperature verification on the DSR that would be used before beginning the testing. Technicians were also requested to be familiar with the MSCR test before participating. Other than reviewing the AASHTO TP70 test procedure, technicians were referred to the MSCR seminar (<http://www.ct.gov/dot/cwp/view.asp?a=1617&q=448116>) hosted by the Connecticut DOT Division of Research and conducted by the Asphalt Institute and FHWA in 2009.

Each participating laboratory received samples of three asphalt binders (labeled as A, B, and C): PG 64-22, PG 76-22, and a lab blend of the two (most similar to a PG 70-22). For each binder, there were three 1-oz tins and one 6-oz tin of unaged material supplied. Labs were asked to do the following:

- Before conducting any testing, heat the 6-oz tin of each unaged asphalt binder and pour two RTFO bottles. Conduct the RTFO aging in accordance with AASHTO T240. After the RTFO aging is complete, scrape the two bottles for each asphalt binder into a single container and stir. At this point, it is recommended that the RTFO-aged binder be split into three separate 1-oz containers, if possible, so that reheating can be minimized. Label each tin with the Sample Identification and “RTFO”.
- Heat the 1-oz tins of unaged (“ORIG”) asphalt binder only once when preparing to conduct testing.
- Perform the MSCR test (AASHTO TP70) at 64°C on each asphalt binder sample as identified on the supplied data form. Perform the testing in the order indicated on the data form. A total of 18 tests will be performed (3 binders x 2 aged conditions x 3 replicates).
- Report all results and either e-mail or scan/fax the data form to AI when complete.

Each participating lab also received a randomized test plan (data sheet) indicating the experimental testing sequence. Note that three different data sheets were used for participating labs (as shown in the Appendix). Asphalt binder samples were tested using the MSCR test at 0.1 kPa and 3.2 kPa shear stress at 64°C for all three asphalt binders.

It was stressed to the participating lab that it was important to follow the randomized testing sequence indicated on the experimental test plan. The testing was randomized to minimize the chance of a systematic error affecting just one set of test results. Labs were also informed that it was preferable that the testing technician and DSR should be the same for all samples, but at a minimum, must be the same for each asphalt binder sample with the same letter designation (i.e., A, B, or C). This last requirement allows for a proper estimate of single operator precision.

### **ILS Procedure**

The ILS procedure followed the guidelines in ASTM E691, *Standard Practice for Conducting an Interlaboratory Study to Determine the Precision of a Test Method*. The purpose of an ILS is to develop information needed for a precision statement. However, the practice cautions the reader that:

“The precision statistics obtained by an ILS such as described in this practice must not be treated as exact mathematical quantities which are applicable to all circumstances and uses...The repeatability limit and reproducibility limit should be considered as general guides, and the associated probability of 95% as only a rough indicator of what can be expected.”

### **Results**

Data was analyzed separately for each asphalt binder sample and test result –  $J_{nr}$  @ 0.1 kPa shear stress (Jnr-0.1), Recovery @ 0.1 kPa shear stress (Rec-0.1),  $J_{nr}$  @ 3.2 kPa shear stress (Jnr-3.2), Recovery @ 3.2 kPa shear stress (Rec-3.2), and Stress Sensitivity (Jnr-Diff). An example of the data is shown in Table 2. The Appendix contains all thirty data tables.

Table 2: Sample AR (PG 64-22 Binder, RTFO-aged) Data for Jnr-3.2  
 Binder AR  
**PG 64-22**

**Jnr (3.2 kPa)**

Lab No.	Lab Name	Results, x			Average	s	d	h	k
		1	2	3					
1		2.292	2.279	2.268	2.27967	0.01201	0.12040	0.73	0.12
2		2.290	2.318	2.377	2.32833	0.04441	0.16907	1.03	0.45
3		2.057	2.070	2.042	2.05633	0.01401	-0.10293	-0.63	0.14
4		2.266	2.228	2.263	2.25233	0.02113	0.09307	0.57	0.21
5		1.762	1.999	1.859	1.87307	0.11886	-0.28620	-1.74	1.21
6		2.361	1.916	1.886	2.05403	0.26610	-0.10523	-0.64	2.70
7		1.854	2.129	2.210	2.06433	0.18660	-0.09493	-0.58	1.89
8		2.291	2.149	2.200	2.21333	0.07193	0.05407	0.33	0.73
9		1.915	1.957	1.892	1.92153	0.03319	-0.23773	-1.45	0.34
10		2.491	2.260	2.276	2.34233	0.12900	0.18307	1.11	1.31
11		2.065	2.078	2.236	2.12633	0.09495	-0.03293	-0.20	0.96
12		2.053	2.046	1.996	2.03160	0.03125	-0.12767	-0.78	0.32
13		2.465	2.454	2.447	2.45533	0.00907	0.29607	1.80	0.09
14		2.311	2.345	2.309	2.32153	0.01991	0.16227	0.99	0.20
15		2.037	2.015	2.080	2.04380	0.03303	-0.11547	-0.70	0.34
16		2.152	2.156	2.128	2.14533	0.01499	-0.01393	-0.08	0.15
17		2.143	2.242	2.279	2.22133	0.07044	0.06207	0.38	0.71
18		2.244	2.248	2.243	2.24520	0.00263	0.08593	0.52	0.03
19		2.221	2.315	2.424	2.32000	0.10174	0.16073	0.98	1.03
20		2.521	2.182	2.156	2.28643	0.20383	0.12717	0.77	2.07
21		2.264	2.295	2.236	2.26497	0.02957	0.10570	0.64	0.30
22		1.897	1.935	1.977	1.93633	0.04002	-0.22293	-1.36	0.41
23		1.938	1.821		1.87960	0.08259	-0.27967	-1.70	0.84
24									
25									
26									
27									
28									
Average of Cell Averages		X-bar			2.159				
Standard Deviation of Cell Averages		S <sub>x-bar</sub>			0.164				
Repeatability Standard Deviation		S <sub>r</sub>			0.09855	0.183			
Reproducibility Standard Deviation		S <sub>R</sub>			0.183				

*Shaded cells were removed after outlier analysis*

A quality review of the data was initiated to ensure that there were no obvious errors that would require removal of some data points from the participating labs. Although there were some reporting errors, these were corrected before conducting the analysis.

The values for Repeatability Standard Deviation ( $s_r$ ) and Reproducibility Standard Deviation ( $s_R$ ) are used to determine the  $1s$ ,  $1s\%$ ,  $d2s$ , and  $d2s\%$  values for repeatability (within-lab) and reproducibility (between-lab), respectively.

The data can also be represented graphically as shown in Figure 1 and Figure 2 (with bands shown for data within one, two, and three standard deviations).

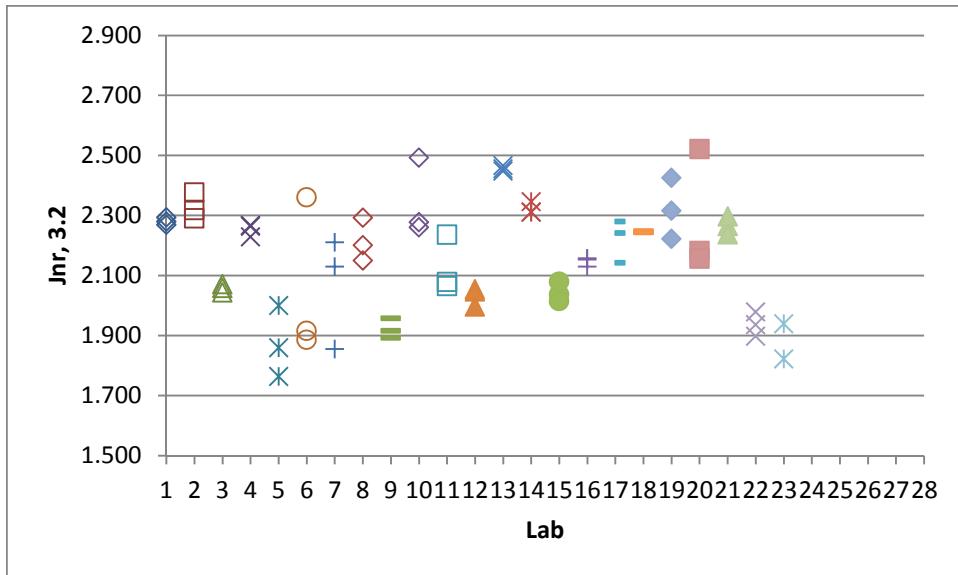


Figure 1: Binder AR (PG 64-22, RTFO-aged) Data for Jnr-3.2

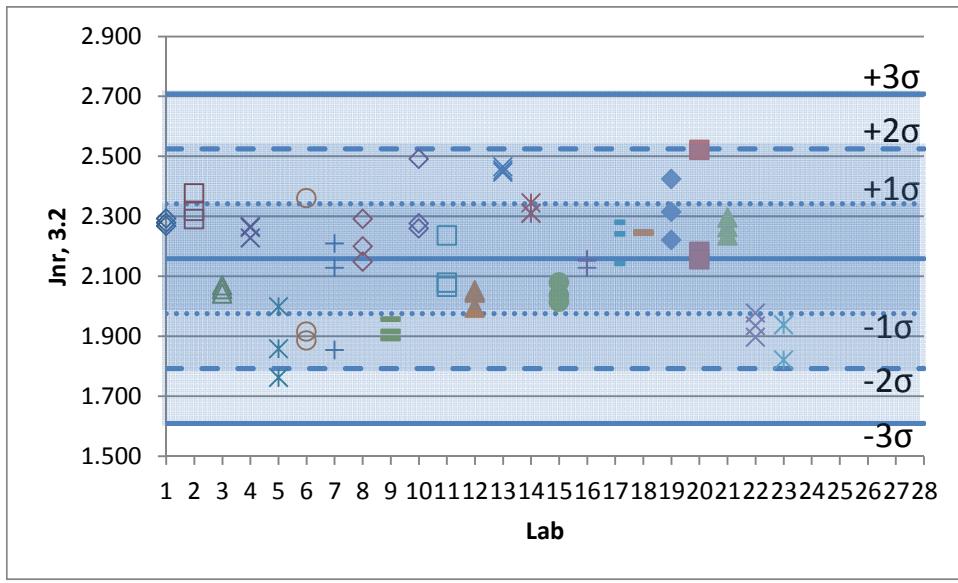


Figure 2: Binder AR (PG 64-22, RTFO-aged) Data for Jnr-3.2 with Standard Deviation Bands

The ILS procedure recommends that outliers should be identified through a consistency analysis. The consistency analysis in ASTM E691 considers two consistency statistics. The “h” consistency statistic is “...an indicator of how one laboratory’s cell average, for a particular material, compares with the average of the other laboratories.” The “k” consistency statistic is “...an indicator of how one laboratory’s within-laboratory variability, under repeatability conditions, on a particular material, compares with all of the laboratories combined. Values of ‘k’ larger than 1 indicate greater within-laboratory variability than the average for all laboratories.” Critical values are chosen at a significance level of 0.5%.

Tables of the “h” and “k” consistency statistics were created for each of the five test parameters: Jnr-0.1, Rec-0.1, Jnr-3.2, Rec-3.2, and Jnr-Diff. These tables are provided in the Appendix with an example table provided below.

Table 3: Consistency Statistic “h” for Rec-3.2

**h**

**Critical Value = 2.59**

**Recovery (3.2 kPa)**

Lab No.	Lab Name	Material					
		Binder AO	Binder BO	Binder CO	Binder AR	Binder BR	Binder CR
1		0.56	-0.21	-0.75	0.48	0.25	0.10
2		0.57	-0.20	0.21	0.46	0.01	0.02
3		0.75	0.28	0.37	0.69	0.36	0.44
4		0.66	-0.03	0.37	0.52	0.17	0.30
5		0.30	-0.40	-0.20	0.51	-0.32	1.32
6		-1.32	1.40	1.49	-1.04	1.51	0.71
7		0.73	-0.67	-0.31	0.64	-1.06	-0.80
8		0.73	<b>-3.82</b>	-1.89	0.50	<b>-3.77</b>	-1.32
9		0.44	0.31	0.08	0.42	0.04	0.06
10		0.61	-0.11	-0.43	0.48	0.33	-0.78
11		-1.31	-0.47	0.00	-1.18	0.62	-0.64
12		0.13	0.55	1.04	0.27	0.39	0.19
13		0.57	0.10	-0.34	0.39	0.12	-0.33
14		0.06	1.27	0.40	0.14	0.38	0.17
15		0.03	0.67	0.61	0.19	0.70	1.29
16		0.12	0.40	0.49	0.26	0.10	1.14
17		0.02	0.14	-0.31	0.15	0.15	0.98
18		0.16	0.62	0.28	0.17	0.38	0.14
19		<b>-3.30</b>	-0.28	<b>-2.24</b>	<b>-3.45</b>	-0.46	<b>-2.50</b>
20		-1.83	0.07	-1.46	-1.99	-0.07	-1.80
21		0.33	-0.49	-0.37	0.14	-1.19	-0.63
22		0.62	-0.15	0.71	0.70	0.45	0.74
23		0.36	1.02	<b>2.24</b>	0.53	0.88	1.22
24		0.00	0.00	0.00	0.00	0.00	0.00
25		0.00	0.00	0.00	0.00	0.00	0.00
26		0.00	0.00	0.00	0.00	0.00	0.00
27		0.00	0.00	0.00	0.00	0.00	0.00
28		0.00	0.00	0.00	0.00	0.00	0.00

exceeds h  
exceeds h-0.5

In Table 3, purple-shaded cells highlight values that exceed the critical value of “h”, indicating inconsistent or questionable data. Yellow-shaded cells are warning indicators that the value is approaching the critical value of “h”.

The data in the consistency tables can be used with graphs to identify if inconsistent data should be eliminated when determining the precision estimates. For example, Lab 8 has “h” values

exceeding the critical value for both Binder BO and Binder BR. Lab 19 has “h” values exceeding the critical value for Binders AO and AR, and values approaching the critical value for Binders CO and CR. An examination of the plot (shown in Figure 3, with Labs 8 and 19 highlighted by dashed lines), verifies that the Rec-3.2 values for Labs 8 and 19 are lower than the average lab values. This can also be seen graphically in the following six figures, where Lab 19 data (Figures 4,5,8, and 9) and Lab 8 data (Figures 6 and 7) are circled.

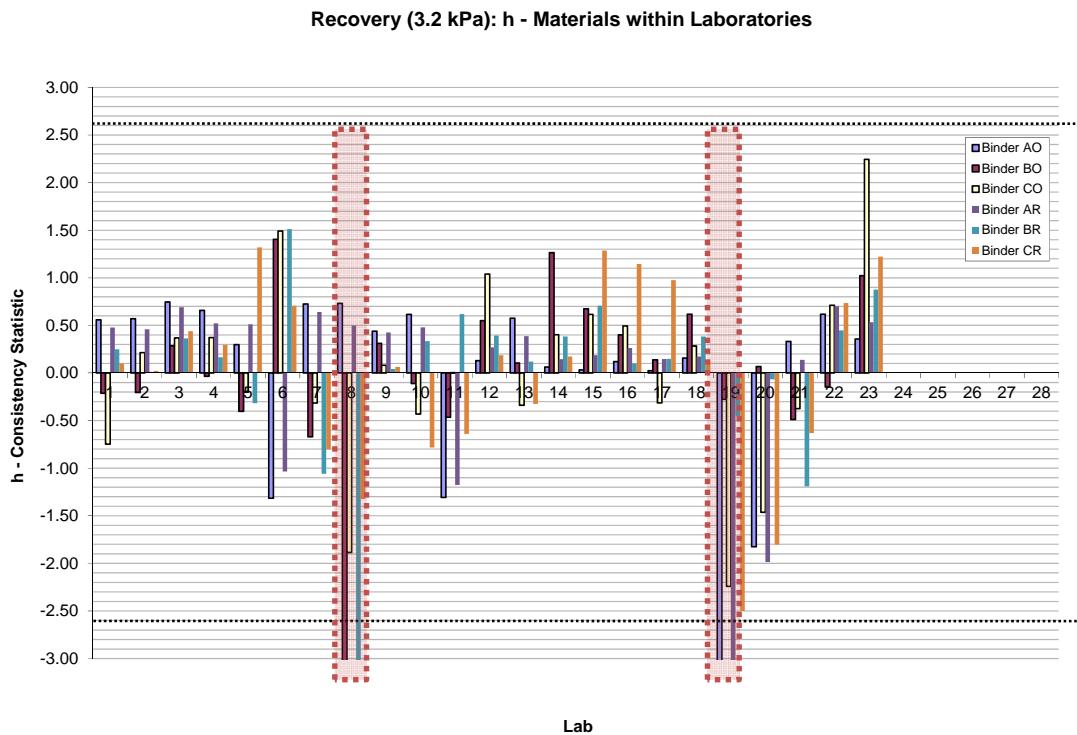


Figure 3: Consistency Statistic “h” for Jnr-3.2 – All Labs and Materials

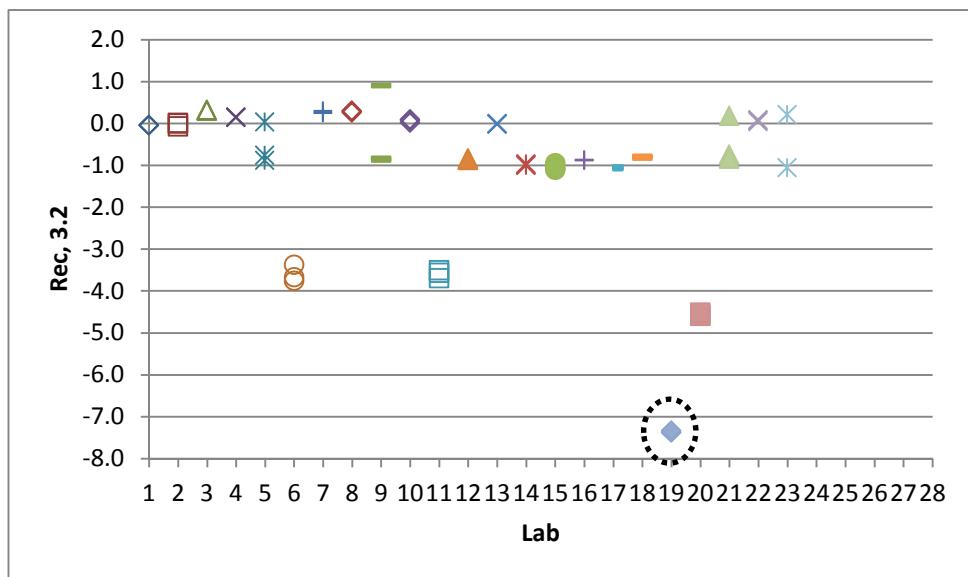


Figure 4: Rec-3.2 Data for Binder AO

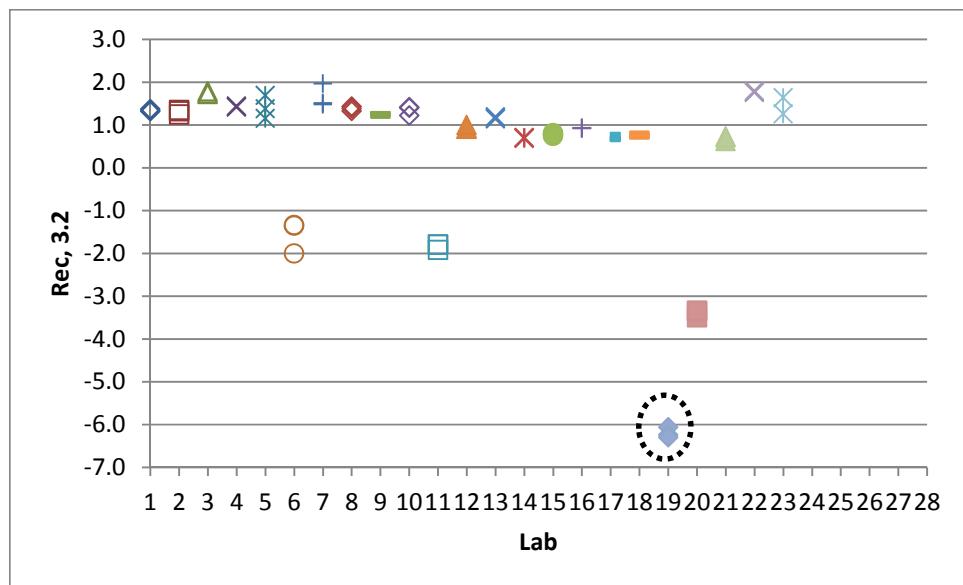


Figure 5: Rec-3.2 Data for Binder AR

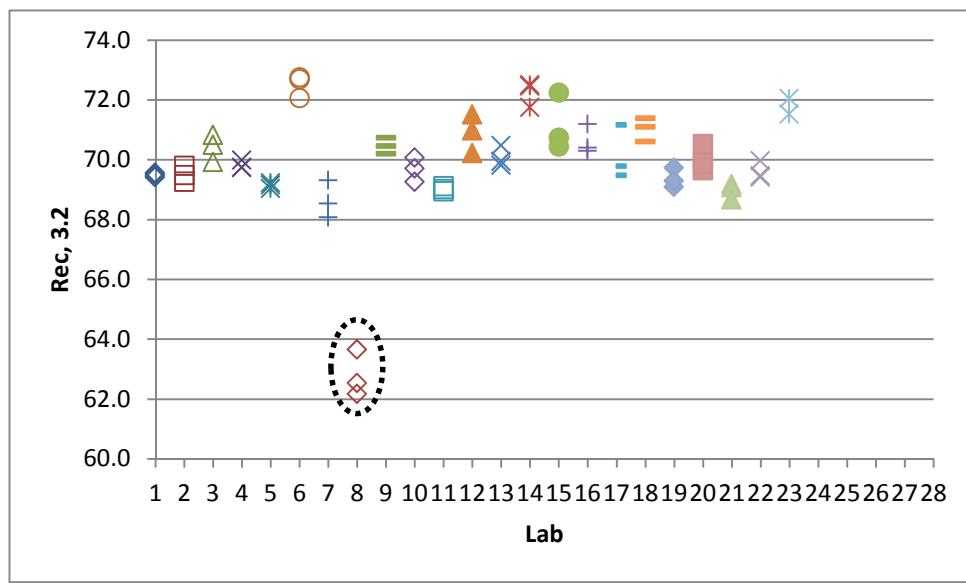


Figure 6: Rec-3.2 Data for Binder BO

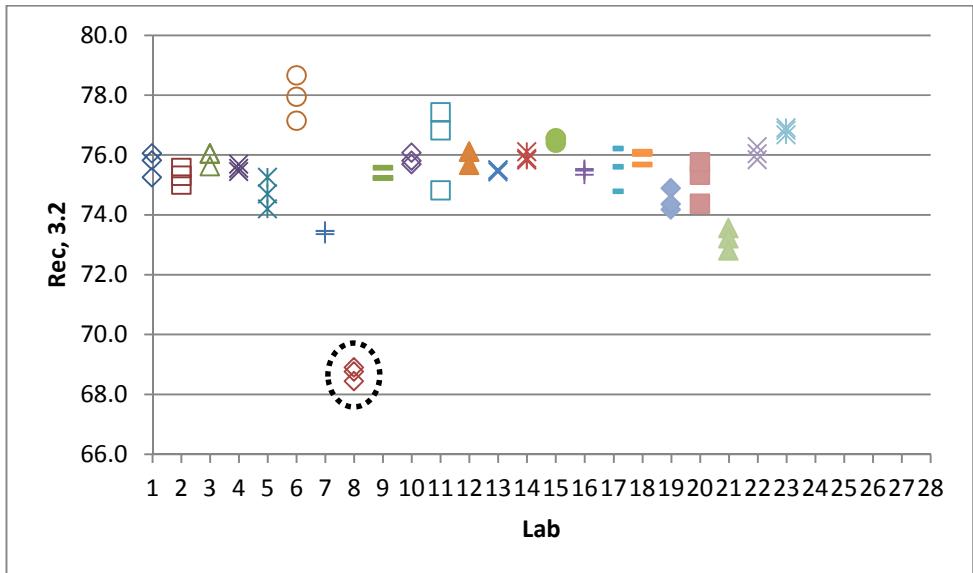


Figure 7: Rec-3.2 Data for Binder BR

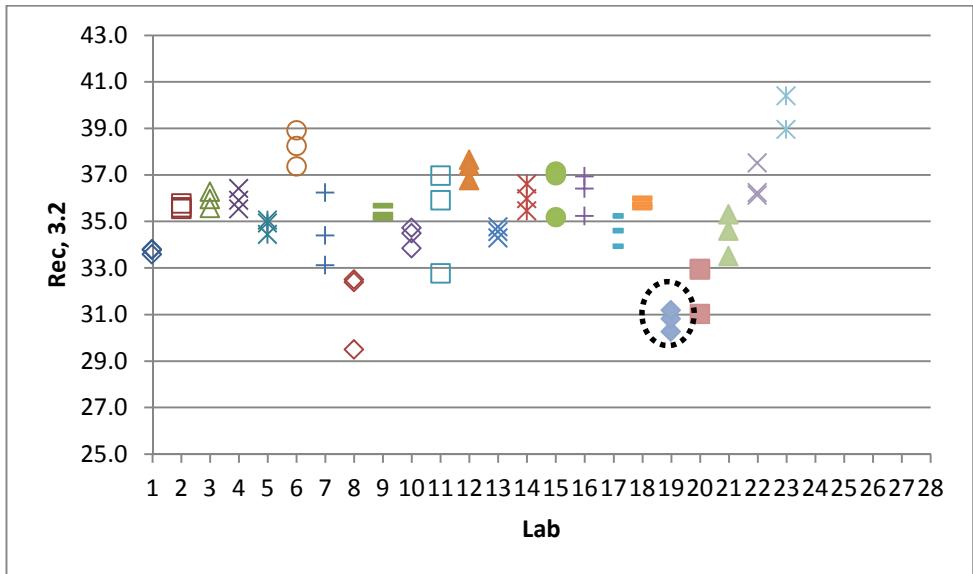


Figure 8: Rec-3.2 Data for Binder CO

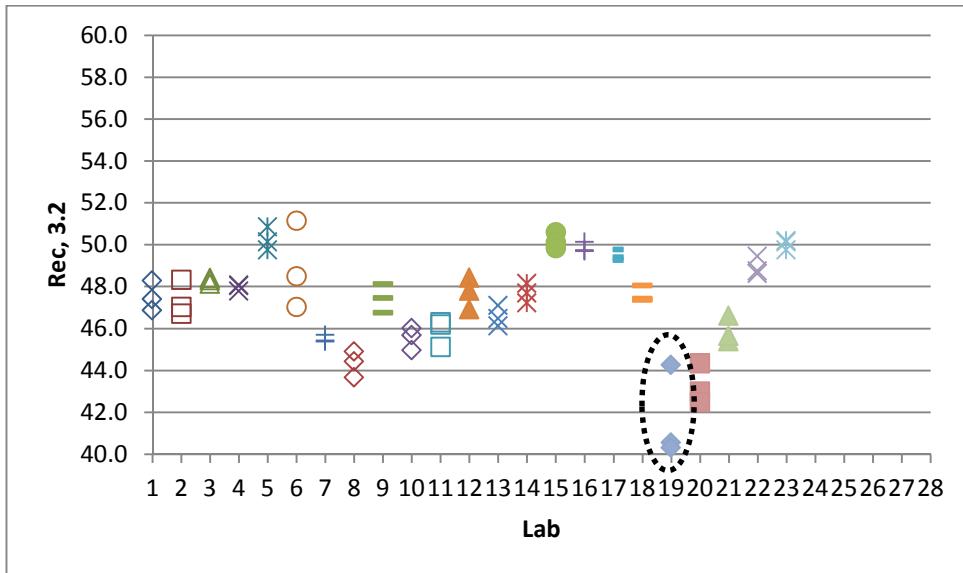


Figure 9: Rec-3.2 Data for Binder CR

As can be seen in Figures 4-7, the circled data is clearly different than the rest of the data – indicating an outlier. Based on the consistency analysis, the Rec-3.2 data from Lab 19 for Binders AO and AR (Figures 4 and 5) and from Lab 8 for Binders BO and BR (Figures 6 and 7) was eliminated from the repeatability and reproducibility calculation.

In Figures 8 and 9, while the circled data looks a little low, it is not clearly different than the rest of the data. As such, the Rec-3.2 data from Lab 19 for Binders CO and CR was left in the repeatability and reproducibility calculation.

This same analysis was conducted for each of the “flagged” data in the consistency tables. It should be noted that ASTM E691 generally does not advocate excluding data except in special instances and recommends against excluding more than 5% of the ILS data to prevent presenting precision that cannot be met in routine use. Thus, to meet this requirement, no more than 1 cell should be eliminated per binder for each MSCR parameter.

Based on the consistency analysis, the data in Table 4 was excluded from the calculation of repeatability and reproducibility.

Table 4: Data Excluded from Repeatability and Reproducibility Calculations

	Jnr-0.1	Jnr-3.2	Rec-0.1	Rec-3.2	Jnr-Diff
<b>Binder AO</b>	none <sup>1</sup>	none <sup>1</sup>	Lab 19	Lab 19	none
<b>Binder AR</b>	none	none	Lab 19	Lab 19	none
<b>Binder BO</b>	none	none	Lab 8	Lab 8	Lab 6
<b>Binder BR</b>	none	none	Lab 8	Lab 8	none
<b>Binder CO</b>	none	none	none	none	none
<b>Binder CR</b>	none	none	none	none	none

<sup>1</sup> although data was not eliminated from any one lab, two suspect data points were eliminated based on analysis of the “k” statistic.



Table 9: Repeatability and Reproducibility Calculations for Jnr-Diff

ID	Binder	X-bar	$s_{X\bar{}}^{} \bar{}$	$s_r$	$s_R$	r	R	Repeatability		Reproducibility	
								1s%	d2s%	1s%	d2s%
AO	PG 64-22	0.06517	0.01337	0.01931	0.02067	0.05407	0.05789	29.6%	83.0%	31.7%	88.8%
BO	PG 76-22	0.11513	0.02814	0.01316	0.03012	0.03684	0.08433	11.4%	32.0%	26.2%	73.2%
CO	PG 70-22	0.36367	0.04617	0.03197	0.05304	0.08952	0.14850	8.8%	24.6%	14.6%	40.8%
AR	PG 64-22	0.07742	0.01646	0.01864	0.02242	0.05220	0.06278	24.1%	67.4%	29.0%	81.1%
BR	PG 76-22	0.07326	0.02389	0.02335	0.03057	0.06537	0.08559	31.9%	89.2%	41.7%	116.8%
CR	PG 70-22	0.21118	0.02754	0.01614	0.03053	0.04519	0.08548	7.6%	21.4%	14.5%	40.5%

In the tables above:

- “X-bar” represents the average value from the participating labs for each binder sample
- “ $s_r$ ” represents the within-lab (single operator) standard deviation, or 1s. Dividing this value by the average value generates the Repeatability coefficient of variation, or 1s%. Multiplying  $s_r$  by 2.8 generates the acceptable range of two test results under repeatability conditions, or d2s. Dividing this product by the average value results in the d2s%.
- “ $s_R$ ” represents the between-lab standard deviation, or 1s. Dividing this value by the average value generates the Reproducibility coefficient of variation, or 1s%. Multiplying  $s_R$  by 2.8 generates the acceptable range of two test results under reproducibility conditions, or d2s. Dividing this product by the average value results in the d2s%.

The original MSCR ILS conducted through the Asphalt Binder Expert Task Group (ETG) resulted in the development of a tiered precision statement for  $J_{nr}$  and a regular precision statement for Recovery. The “tiers” for  $J_{nr}$  were established as follows:

- $J_{nr}$  greater than  $1.00 \text{ kPa}^{-1}$ ;
- $J_{nr}$  greater than  $0.25 \text{ kPa}^{-1}$  and less than or equal to  $1.00 \text{ kPa}^{-1}$ ; and
- $J_{nr}$  greater than  $0.10 \text{ kPa}^{-1}$  and less than or equal to  $0.25 \text{ kPa}^{-1}$

The NEAUPG ILS did not support a tiered precision statement for  $J_{nr}$ . A review of the data in Tables 5 and 6 indicates that this ILS also does not support the tiered precision statement developed by the Asphalt Binder ETG. Repeatability and Reproducibility values appear very similar for  $J_{nr}$  for all asphalt binders. As such, average values were calculated for  $J_{nr}$  repeatability and reproducibility.

Similarly, the data in Tables 7 and 8 were used to calculate average values for Recovery repeatability and reproducibility. Note that the Recovery values for Binders AO and AR were not used in calculating the average repeatability and reproducibility. The actual Recovery value is low for both binders (expected since they are unmodified asphalt binders), leading to unreasonably high 1s% and d2s% values. The repeatability and reproducibility values for Recovery were determined by only using the data from Binders BO, BR, CO, and CR.

In some cases for Binders AO and AR negative Recovery values were reported by the participating lab. Negative Recovery values are usually only seen with unmodified asphalt binders. To address the issue, guidance from the Binder ETG suggests that negative Recovery values should be simply reported as 0%.

Jnr-Diff has not been determined in previous ILS programs so the results of this ILS cannot be compared to other data. A review of the data in Table 9 suggests that the within-lab standard

deviation ( $s_r$ ) and between-lab standard deviation ( $s_R$ ) are relatively unaffected by the average test value. As such, it appears that coefficient of variation is not the best way of representing this data. For Jnr-Diff, the precision data is shown in Table 10.

Table 10: Precision Estimates for Jnr-Diff from SEAUPG ILS

<b>Condition</b>	<b>Acceptable Range of Two Test Results (d2s)</b>
Single-Operator Precision Jnr-Diff	0.057
Multilaboratory Precision Jnr-Diff	0.085

Table 11 provides a comparison of the SEAUPG ILS with the 2010 NEAUPG ILS and the ILS previously performed by the Asphalt Binder ETG.

Table 11: Comparison of SEAUPG ILS with NEAUPG ILS and ETG ILS

<i>Condition</i>	<i>Acceptable Range of Two Test Results (d2s%)</i>		
	<i>SEAUPG ILS</i>	<i>NEAUPG ILS</i>	<i>ETG ILS</i>
<i>Single-Operator Precision:</i>			
Recovery <sub>0.1kPa</sub> (%)	3.2%	9.7%	6.7%
Recovery <sub>3.2kPa</sub> (%)	3.9%	7.7%	8.5%
<i>Multilaboratory Precision:</i>			
Recovery <sub>0.1kPa</sub> (%)	6.8%	14.0%	15.0%
Recovery <sub>3.2kPa</sub> (%)	9.8%	18.7%	18.1%
<i>Single-Operator Precision:</i>			
$J_{nr}@0.1kPa$ (kPa <sup>-1</sup> )			
>1.00	13.3%	14.5%	12.8%
0.26 - 1.00	13.3%	14.5%	15.2%
0.10 - 0.25	13.3%	14.5%	38.3%
<0.10	n/a	n/a	n/a
$J_{nr}@3.2kPa$ (kPa <sup>-1</sup> )			
>1.00	13.0%	14.5%	16.0%
0.26 - 1.00	13.0%	14.5%	15.3%
0.10 - 0.25	13.0%	n/a	26.6%
<0.10	n/a	n/a	n/a
<i>Multilaboratory Precision:</i>			
$J_{nr}@0.1kPa$ (kPa <sup>-1</sup> )			
>1.00	26.1%	25.6%	25.6%
0.26 - 1.00	26.1%	25.6%	35.6%
0.10 - 0.25	26.1%	25.6%	46.8%
<0.10	n/a	n/a	n/a
$J_{nr}@3.2kPa$ (kPa <sup>-1</sup> )			
>1.00	28.0%	33.7%	22.0%
0.26 - 1.00	28.0%	33.7%	39.0%
0.10 - 0.25	28.0%	n/a	42.6%
<0.10	n/a	n/a	n/a

The data in Table 11 indicates that the SEAUPG ILS has an equal or lower value (indicating better repeatability and/or reproducibility) in nearly all cases than was generated from the NEAUPG ILS and ETG ILS.

## **Summary**

1. After removing potential outliers, the data indicates SEAUPG ILS has an equal or lower d<sub>2s</sub>% value, thus indicating better repeatability and/or reproducibility than generated from the NEAUPG ILS and ETG ILS.
2. The analysis of the SEAUPG ILS data did not indicate that a tiered precision statement for J<sub>nr</sub> was warranted. Values for d<sub>2s</sub>% were approximately equal, regardless of the measured J<sub>nr</sub> value. This finding matched the results from the NEAUPG ILS, which contradicted the ETG ILS.
3. Analysis of Jnr-Diff data indicated that the standard deviation was unaffected by the test value. As such, unlike the other parameters precision estimates are reported as d<sub>2s</sub>, not d<sub>2s</sub>% (coefficient of variation).

## **Acknowledgments**

The information presented in this report would not have been possible without the willingness of the SEAUPG labs to provide data. Thanks to all the participating labs. Thanks in particular to Don Siler (Marathon), Andy Menapace (Paragon Technical Services), Rick Holmgreen (ConocoPhillips), Danny Gierhart (Asphalt Institute), and Bob Horan (Asphalt Institute) for coordinating sample supply and assisting with the development of the ILS program.

The analysis was conducted as part of the Asphalt Institute's Cooperative Agreements with the Federal Highway Administration through projects DTFH61-08-H-00030 and DTFH61-11-H-00033. Thanks to John Bukowski (FHWA, AOTR for DTFH61-08-H-00030) and Michael Arasteh (FHWA, AOTR for DTFH61-11-H-00033) for working with us on this project.

Finally, the support of the Member Companies of the Asphalt Institute is gratefully acknowledged. Without their interest and support, none of the work presented could have been accomplished.

## **Disclaimer**

The analyses, conclusions, and recommendations expressed in this report are those of the author and do not necessarily reflect the views of the Federal Highway Administration, the Asphalt Institute or its individual Member Companies.

# APPENDIX

## **SEAUPG MSCR Round Robin Participating Labs**

**October 2011**

North Carolina DOT  
Materials and Test Unit  
1801 Blue Ridge Road  
Raleigh, NC 27607  
Attn: Brian Hunter  
[bhunter@ncdot.gov](mailto:bhunter@ncdot.gov)

Oklahoma DOT  
Materials Division - Bituminous Engineer  
200 NE 21st Street  
Oklahoma City, OK 73105-3204  
Attn: Kenneth Ray Hobson, P.E.  
[khobson@odot.org](mailto:khobson@odot.org)

Florida DOT  
State Materials Office  
5007 NE 39 Ave  
Gainesville, Florida 32609  
Attn: Tanya Nash  
[tanya.nash@dot.state.fl.us](mailto:tanya.nash@dot.state.fl.us)

Virginia Center for Transportation Innovation and Research  
530 Edgemont Road  
Charlottesville, VA 22903  
Attn: Donnie Dodds  
[Donald.Dodds@VDOT.Virginia.gov](mailto:Donald.Dodds@VDOT.Virginia.gov)

Texas DOT - Construction Division  
Asphalt Lab  
9500 N. Lake Creek Parkway  
Austin, TX 78717  
Attn: Jerry Peterson  
[jerry.peterson@txdot.gov](mailto:jerry.peterson@txdot.gov)

Georgia DOT  
Office of Materials & Research  
15 Kennedy Drive  
Forest Park, GA 30297  
Attention: Freddie Sexton (Bit. Control)  
[pwu@dot.ga.gov](mailto:pwu@dot.ga.gov)

Alabama DOT  
Materials and Tests  
3704 Fairground Road  
Montgomery, AL 36177-9965  
ATTN: Douglas Greenwood  
[greenwoodd@dot.state.al.us](mailto:greenwoodd@dot.state.al.us)

Louisiana Transportation Research Center  
4101 Gourrier Avenue  
Baton Rouge, LA 70808-4443  
Attn: Md Sharear Kabir  
[mdsharear.kabir@la.gov](mailto:mdsharear.kabir@la.gov)

South Carolina DOT  
Office of Materials & Research  
1406 Shop Road  
Columbia, SC 29201  
Attn: Cliff Selkinghaus  
[selkinghcb@scdot.org](mailto:selkinghcb@scdot.org)

Mississippi DOT  
412 East Woodrow Wilson  
Jackson, MS 39216  
Attn: Chemical Lab  
[bstrum@mdot.state.ms.us](mailto:bstrum@mdot.state.ms.us)

Paragon Technical Services  
390 Carrier Blvd  
Richland, MS 39218  
Attn: Andy Menapace  
[andy.menapace@ptsilab.com](mailto:andy.menapace@ptsilab.com)

Marathon Petroleum Company  
Asphalt Certification Lab  
11631 U.S. Route 23  
Catlettsburg, KY 41129  
Attn: Paul Hartwig  
[pfhartwig@marathonpetroleum.com](mailto:pfhartwig@marathonpetroleum.com)

National Center for Asphalt Technology  
277 Technology Pkwy  
Auburn, AL 36830  
Attn: Pamela Turner  
[turnepa@auburn.edu](mailto:turnepa@auburn.edu)

BETA R&D  
11310 Reichold Rd  
Gulfport, MS 39503  
Attn: Codrin Daranga  
[cdaranga@be-inc.com](mailto:cdaranga@be-inc.com)

Matthew Corrigan  
Federal Highway Administration  
HIPT-10, Room E73-465  
1200 New Jersey Ave. SE  
Washington, DC 20590  
[matthew.corrigan@dot.gov](mailto:matthew.corrigan@dot.gov)

Ergon Asphalt and Emulsions  
300 Industrial Road  
Halstead, KS 67056  
Attn: Tim Weatherman

Lion Oil Company  
2501 Port Place  
Muskogee, Oklahoma 74403  
Attn: Tim Weatherman

Ergon Asphalt and Emulsions  
3901 West Ponderosa Way  
Las Vegas, NV 89118  
Attn: Joel Shealy

Ergon Asphalt and Emulsions  
95 West 1100 North  
North Salt Lake, UT 84054  
Attn: Joel Shealy

Ergon Asphalt and Emulsions  
1114 Visco Drive  
Nashville, TN 37210  
Attn: Shelly Cowley

Asphalt Institute  
2696 Research Park Dr.  
Lexington, KY 40511-8480  
Attn: Gary Irvine  
[girvine@asphaltinstitute.org](mailto:girvine@asphaltinstitute.org)



SEAUPG Interlaboratory Study for Determining the Precision of AASHTO TP70, the Multiple Stress Creep Recovery (MSCR) Test

October 2011

To: Participating Labs

From: Mike Anderson, Asphalt Institute  
859.288.4984 office  
859.422.1301 FAX  
[manderson@asphalteinstitute.org](mailto:manderson@asphalteinstitute.org)

Thanks for agreeing to participate in the SEAUPG Interlaboratory Study (ILS) on the MSCR test. Enclosed are asphalt binder samples, a data reporting form, and appropriate MSDS. You should be receiving three different asphalt binders (labeled as A, B, and C) representing PG 64-22 and PG 76-22 binders, as well as a lab blend of the two binders. For each binder, there will be three 1-oz tins and one 6-oz tin of unaged material. Please do the following:

- The 6-oz tin includes enough unaged asphalt binder to perform RTFO-aging using two bottles. **Before conducting any testing**, heat the 6-oz tin for each asphalt binder and pour two RTFO bottles. Conduct the RTFO aging in accordance with AASHTO T240. After the RTFO aging is complete, scrape the two bottles for each asphalt binder into a single container and stir. At this point, we would recommend that the RTFO-aged binder be split into three separate 1-oz containers, if possible. Label each tin with the Sample Identification and "RTFO".
- The 1-oz tins are replicates of unaged ("ORIG") asphalt binder. They should only be heated once.
- Perform the MSCR test (AASHTO TP70) at 64°C on each asphalt binder sample as identified on the data form. **Please perform the testing in the order indicated on the data form.** There are a total of 18 tests to be performed (3 binders x 2 aged conditions x 3 replicates).
- Report all results and either e-mail or scan/fax the data form to me when complete.

If you have any questions, please contact me. Thanks for your participation!

**SEAUPG - Interlaboratory Study for Determining the Precision of AASHTO TP70, the Multiple Stress Creep Recovery (MSCR) Test**

**Data Sheet - 1**

**October 2011**

**Lab**

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Please return this form to: [manderson@asphaltinstitute.org](mailto:manderson@asphaltinstitute.org)

***Please follow the testing order indicated in the table below (i.e., top to bottom).***

Sample	Cond.	Test Temp, C	0.1 kPa Shear Stress		3.2 kPa Shear Stress		Jnr Diff*	Date	Tech	DSR Make &Model
			J <sub>nr</sub> , kPa <sup>-1</sup>	%Recovery	J <sub>nr</sub> , kPa <sup>-1</sup>	%Recovery				
A ORIG		64								
A ORIG		64								
A RTFO		64								
B RTFO		64								
C RTFO		64								
B ORIG		64								
A ORIG		64								
C ORIG		64								
B RTFO		64								
C ORIG		64								
B ORIG		64								
C ORIG		64								
B ORIG		64								
B RTFO		64								
A RTFO		64								
A RTFO		64								
C RTFO		64								
C RTFO		64								

\* calculated as (J<sub>n</sub>r,3.2kPa - J<sub>n</sub>r,0.1kPa)/(J<sub>n</sub>r,0.1kPa)

**SEAUPG - Interlaboratory Study for Determining the Precision of AASHTO TP70, the Multiple Stress Creep Recovery (MSCR) Test****Data Sheet - 2****October 2011****Lab**Please return this form to: [manderson@asphaltinstitute.org](mailto:manderson@asphaltinstitute.org)***Please follow the testing order indicated in the table below (i.e., top to bottom).***

Sample	Cond.	Test Temp, C	0.1 kPa Shear Stress		3.2 kPa Shear Stress		Jnr Diff*	Date	Tech	DSR Make &Model
			J <sub>nr</sub> , kPa <sup>-1</sup>	%Recovery	J <sub>nr</sub> , kPa <sup>-1</sup>	%Recovery				
C RTFO		64								
C ORIG		64								
B RTFO		64								
A ORIG		64								
B ORIG		64								
B RTFO		64								
C ORIG		64								
B RTFO		64								
C ORIG		64								
C RTFO		64								
A ORIG		64								
C RTFO		64								
A RTFO		64								
A RTFO		64								
B ORIG		64								
A ORIG		64								
A RTFO		64								
B ORIG		64								

\* calculated as (Jnr,3.2kPa - Jnr,0.1kPa)/(Jnr,0.1kPa)

**SEAUPG - Interlaboratory Study for Determining the Precision of AASHTO TP70, the Multiple Stress Creep Recovery (MSCR) Test**

**Data Sheet - 3**

**October 2011**

**Lab**

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Please return this form to: [manderson@asphaltinstitute.org](mailto:manderson@asphaltinstitute.org)

***Please follow the testing order indicated in the table below (i.e., top to bottom).***

Sample	Cond.	Test Temp, C	0.1 kPa Shear Stress		3.2 kPa Shear Stress		Jnr Diff*	Date	Tech	DSR Make &Model
			J <sub>nr</sub> , kPa <sup>-1</sup>	%Recovery	J <sub>nr</sub> , kPa <sup>-1</sup>	%Recovery				
B RTFO		64								
C RTFO		64								
B ORIG		64								
A ORIG		64								
C ORIG		64								
A RTFO		64								
C RTFO		64								
B RTFO		64								
A ORIG		64								
C ORIG		64								
B ORIG		64								
B ORIG		64								
A RTFO		64								
C ORIG		64								
A RTFO		64								
A ORIG		64								
B RTFO		64								
C RTFO		64								

\* calculated as (J<sub>n</sub>r,3.2kPa - J<sub>n</sub>r,0.1kPa)/(J<sub>n</sub>r,0.1kPa)

Binder AO

**PG 64-22**

**Jnr (0.1 kPa)**

Lab No.	Lab Name	Results, x			Average	s	d	h	k
		1	2	3					
1		4.924	4.848	4.851	4.87433	0.04304	0.46350	1.79	0.18
2		4.675	4.513	4.503	4.56367	0.09655	0.15283	0.59	0.41
3		4.491	4.458	4.549	4.49933	0.04607	0.08850	0.34	0.20
4		4.316	4.327	4.338	4.32700	0.01100	-0.08383	-0.32	0.05
5		2.768	4.453	4.212	4.33270	0.17041	-0.07813	-0.30	0.72
6		4.664	4.487	3.952	4.36747	0.37087	-0.04337	-0.17	1.57
7		4.520	4.670	4.770	4.65333	0.12583	0.24250	0.93	0.53
8		4.152	4.291	4.172	4.20500	0.07515	-0.20583	-0.79	0.32
9		4.317	4.255	4.423	4.33163	0.08520	-0.07920	-0.31	0.36
10		5.269	4.870	4.781	4.97333	0.25989	0.56250	2.17	1.10
11		4.169	4.615	4.068	4.28370	0.29094	-0.12713	-0.49	1.24
12		4.194	4.323	4.283	4.26687	0.06636	-0.14397	-0.55	0.28
13		4.419	5.042	4.660	4.70700	0.31415	0.29617	1.14	1.33
14		4.726	4.671	4.772	4.72320	0.05082	0.31237	1.20	0.22
15		4.124	4.501	4.324	4.31640	0.18817	-0.09443	-0.36	0.80
16		4.244	4.250	4.267	4.25373	0.01151	-0.15710	-0.61	0.05
17		4.836	4.765	4.727	4.77600	0.05543	0.36517	1.41	0.24
18		4.198	4.138	4.126	4.15410	0.03879	-0.25673	-0.99	0.16
19		4.131	4.281	4.059	4.15710	0.11300	-0.25373	-0.98	0.48
20		4.096	4.096	3.884	4.02490	0.12246	-0.38593	-1.49	0.52
21		4.182	4.403	2.389	4.29240	0.15684	-0.11843	-0.46	0.67
22		4.236	4.112	4.414	4.25400	0.15180	-0.15683	-0.60	0.64
23		3.523	4.701		4.11200	0.83297	-0.29883	-1.15	3.54
24									
25									
26									
27									
28									

Average of Cell Averages

X-bar

4.411

Standard Deviation of Cell Averages

s<sub>X-bar</sub>

0.259

Repeatability Standard Deviation

s<sub>r</sub>

0.23554

0.323

Reproducibility Standard Deviation

s<sub>R</sub>

0.323

*Shaded cells were removed after outlier analysis*

Binder AO

**PG 64-22**

**Jnr (3.2 kPa)**

Lab No.	Lab Name	Results, x								
		1	2	3	Average	s	d	h	k	
1		5.212	5.116	5.159	5.16233	0.04809	0.47020	1.76	0.21	
2		4.935	4.854	4.848	4.87900	0.04859	0.18687	0.70	0.21	
3		4.661	4.668	4.760	4.69633	0.05525	0.00420	0.02	0.24	
4		4.769	4.739	4.713	4.74033	0.02802	0.04820	0.18	0.12	
5		2.866	4.744	4.469	4.60630	0.19445	-0.08583	-0.32	0.83	
6		4.953	4.785	4.222	4.65317	0.38269	-0.03897	-0.15	1.64	
7		4.630	4.900	4.920	4.81667	0.16197	0.12453	0.47	0.69	
8		4.737	4.550	4.400	4.56233	0.16884	-0.12980	-0.49	0.72	
9		4.570	4.544	4.707	4.60690	0.08746	-0.08523	-0.32	0.37	
10		5.706	5.254	5.167	5.37567	0.28937	0.68353	2.56	1.24	
11		4.515	4.779	4.398	4.56417	0.19541	-0.12797	-0.48	0.84	
12		4.490	4.572	4.541	4.53417	0.04133	-0.15797	-0.59	0.18	
13		4.981	5.047	4.916	4.98133	0.06550	0.28920	1.08	0.28	
14		4.998	4.960	5.057	5.00513	0.04895	0.31300	1.17	0.21	
15		4.418	4.779	4.576	4.59060	0.18107	-0.10153	-0.38	0.77	
16		4.565	4.542	4.557	4.55450	0.01172	-0.13763	-0.52	0.05	
17		5.067	4.971	4.965	5.00100	0.05689	0.30887	1.16	0.24	
18		4.529	4.419	4.463	4.47037	0.05490	-0.22177	-0.83	0.23	
19		4.329	4.524	4.297	4.38303	0.12268	-0.30910	-1.16	0.53	
20		4.424	4.424	4.116	4.32130	0.17823	-0.37083	-1.39	0.76	
21		4.455	4.689	2.544	4.57210	0.16518	-0.12003	-0.45	0.71	
22		4.546	4.112	4.762	4.47333	0.33104	-0.21880	-0.82	1.42	
23		3.800	4.938		4.36900	0.80469	-0.32313	-1.21	3.44	
24										
25										
26										
27										
28										

Average of Cell Averages

X-bar                          4.692

Standard Deviation of Cell Averages

s<sub>X-bar</sub>                          0.267

Repeatability Standard Deviation

s<sub>r</sub>                                  0.23367                          0.328

Reproducibility Standard Deviation

s<sub>R</sub>                                  0.328

*Shaded cells were removed after outlier analysis*

Binder AO

**PG 64-22**

**Recovery (0.1 kPa)**

Lab No.	Lab Name	Results, x			Average	s	d	h	k
		1	2	3					
1		0.8	0.7	1.0	0.838	0.177	0.394	0.33	0.13
2		0.5	1.1	1.7	1.113	0.600	0.669	0.56	0.46
3		1.1	1.2	1.2	1.169	0.064	0.725	0.61	0.05
4		2.3	1.5	0.8	1.538	0.733	1.094	0.92	0.56
5		1.3	0.4	0.4	0.703	0.508	0.259	0.22	0.39
6		-2.2	-1.8	-1.4	-1.808	0.380	-2.252	-1.90	0.29
7		0.9	2.1	1.1	1.367	0.615	0.923	0.78	0.47
8		2.4	2.7	2.5	2.530	0.160	2.086	1.76	0.12
9		0.4	0.7	0.1	0.387	0.298	-0.057	-0.05	0.23
10		1.3	0.5	1.1	0.981	0.430	0.537	0.45	0.33
11		-0.4	-4.1	-0.9	-1.824	1.966	-2.268	-1.91	1.50
12		0.7	-0.1	0.0	0.198	0.454	-0.246	-0.21	0.35
13		6.2	-4.5	0.5	0.732	5.360	0.288	0.24	4.09
14		0.1	0.2	0.1	0.141	0.083	-0.303	-0.26	0.06
15		0.5	0.2	0.1	0.268	0.241	-0.176	-0.15	0.18
16		0.3	0.3	0.1	0.238	0.115	-0.206	-0.17	0.09
17		0.0	0.0	0.1	0.011	0.040	-0.433	-0.36	0.03
18		1.2	0.8	1.4	1.126	0.312	0.682	0.57	0.24
19		-6.5	-6.2	-6.2					
20		-2.3	-2.3	-3.5	-2.692	0.681	-3.137	-2.64	0.52
21		0.7	1.1	0.9	0.915	0.186	0.471	0.40	0.14
22		1.0	0.2	1.8	1.029	0.790	0.585	0.49	0.60
23		1.7	-0.1		0.810	1.299	0.365	0.31	0.99
24									
25									
26									
27									
28									

Average of Cell Averages

X-bar                    0.444

Standard Deviation of Cell Averages

s<sub>X-bar</sub>                1.187

Repeatability Standard Deviation

s<sub>r</sub>                    1.31083      1.598

Reproducibility Standard Deviation

s<sub>R</sub>                    1.598

*Shaded cells were removed after outlier analysis*









# Binder AR

**PG 64-22**

## Recovery (0.1 kPa)

Lab No.	Lab Name	Results, x			Average	s	d	h	k
		1	2	3					
1		3.9	3.8	3.8	3.827	0.051	0.390	0.26	0.03
2		3.7	5.6	2.9	4.066	1.360	0.629	0.42	0.91
3		4.3	4.6	4.8	4.560	0.220	1.123	0.74	0.15
4		3.8	4.1	5.1	4.326	0.665	0.890	0.59	0.45
5		6.0	4.3	4.9	5.049	0.879	1.612	1.07	0.59
6		1.4	1.7	2.2	1.786	0.397	-1.651	-1.09	0.27
7		5.2	3.6	3.5	4.093	0.953	0.657	0.43	0.64
8		5.1	5.1	4.7	4.993	0.237	1.557	1.03	0.16
9		4.4	4.3	4.7	4.474	0.205	1.038	0.69	0.14
10		3.4	4.3	4.0	3.885	0.471	0.448	0.30	0.32
11		2.1	4.2	3.8	3.386	1.135	-0.051	-0.03	0.76
12		3.2	3.3	4.1	3.543	0.527	0.106	0.07	0.35
13		8.0	1.3	-2.0	2.442	5.089	-0.995	-0.66	3.42
14		3.8	3.2	3.0	3.312	0.432	-0.125	-0.08	0.29
15		3.4	3.5	3.6	3.508	0.088	0.071	0.05	0.06
16		3.6	3.5	3.4	3.456	0.103	0.020	0.01	0.07
17		3.2	3.1	3.1	3.122	0.078	-0.314	-0.21	0.05
18		4.1	3.6	3.5	3.702	0.312	0.265	0.18	0.21
19		-3.4	-3.5	-3.8					
20		-1.8	-2.0	-1.2	-1.675	0.398	-5.111	-3.38	0.27
21		2.6	2.1	3.4	2.675	0.661	-0.762	-0.50	0.44
22		4.4	5.2	6.9	5.528	1.297	2.091	1.38	0.87
23		-1.1	4.2		1.548	3.776	-1.888	-1.25	2.54
24									
25									
26									
27									
28									

Average of Cell Averages

X-bar                    3.437

Standard Deviation of Cell Averages

s<sub>X-bar</sub>                1.510

Repeatability Standard Deviation

s<sub>r</sub>                    1.48950      1.939

Reproducibility Standard Deviation

s<sub>R</sub>                    1.939

*Shaded cells were removed after outlier analysis*

## Binder AR

**PG 64-22****Recovery (3.2 kPa)**

Lab No.	Lab Name	Results, x			Average	s	d	h	k
		1	2	3					
1		1.4	1.3	1.3	1.33567	0.02730	0.61476	0.47	0.20
2		1.3	1.3	1.2	1.30100	0.05311	0.58009	0.45	0.39
3		1.7	1.7	1.8	1.74667	0.03453	1.02576	0.79	0.25
4		1.4	1.4	1.4	1.42467	0.01419	0.70376	0.54	0.10
5		1.7	1.2	1.4	1.40263	0.26793	0.68173	0.53	1.98
6		-2.0	-1.3	-1.4	-1.56900	0.37705	-2.28991	-1.77	2.78
7		2.0	1.5	1.5	1.65000	0.26889	0.92909	0.72	1.98
8		1.4	1.4	1.3	1.37667	0.05508	0.65576	0.51	0.41
9		1.3	1.2	1.2	1.23617	0.02746	0.51526	0.40	0.20
10		1.2	1.4	1.4	1.34333	0.10637	0.62243	0.48	0.79
11		-1.8	-1.8	-1.9	-1.83847	0.07356	-2.55937	-1.97	0.54
12		0.9	0.9	1.0	0.93493	0.05053	0.21403	0.16	0.37
13		1.1	1.2	1.2	1.16433	0.02654	0.44343	0.34	0.20
14		0.7	0.7	0.7	0.69507	0.01240	-0.02584	-0.02	0.09
15		0.8	0.8	0.7	0.77707	0.03095	0.05616	0.04	0.23
16		0.9	0.9	0.9	0.92330	0.00433	0.20239	0.16	0.03
17		0.8	0.7	0.7	0.70267	0.05714	-0.01824	-0.01	0.42
18		0.8	0.7	0.7	0.75343	0.03928	0.03253	0.03	0.29
19		-6.1	-6.2	-6.3					
20		-3.5	-3.4	-3.3	-3.39703	0.08560	-4.11794	-3.17	0.63
21		0.7	0.6	0.7	0.68440	0.05794	-0.03651	-0.03	0.43
22		1.8	1.8	1.8	1.76900	0.01970	1.04809	0.81	0.15
23		1.6	1.3		1.44345	0.26806	0.72254	0.56	1.98
24									
25									
26									
27									
28									

Average of Cell Averages

X-bar                    0.721

Standard Deviation of Cell Averages

S<sub>X-bar</sub>                1.297

Repeatability Standard Deviation

S<sub>r</sub>                    0.13549            1.302

Reproducibility Standard Deviation

S<sub>R</sub>                    1.302*Shaded cells were removed after outlier analysis*

# Binder AR

**PG 64-22**

**Jnr, Diff**

Lab No.	Lab Name	Results, x			Average	s	d	h	k
		1	2	3					
1		0.081	0.079	0.080	0.08003	0.00063	0.00261	0.16	0.03
2		0.082	0.111	0.078	0.09039	0.01808	0.01297	0.79	0.97
3		0.067	0.067	0.072	0.06830	0.00288	-0.00911	-0.55	0.15
4		0.104	0.106	0.120	0.11000	0.00872	0.03258	1.98	0.47
5		0.099	0.083	0.086	0.08915	0.00838	0.01174	0.71	0.45
6		0.086	0.074	0.082	0.08070	0.00611	0.00328	0.20	0.33
7		0.055	0.049	0.043	0.04893	0.00606	-0.02848	-1.73	0.32
8		0.078	0.061	0.067	0.06880	0.00838	-0.00862	-0.52	0.45
9		0.079	0.081	0.085	0.08197	0.00306	0.00456	0.28	0.16
10		0.083	0.086	0.095	0.08815	0.00593	0.01073	0.65	0.32
11		0.085	0.120	0.112	0.10591	0.01848	0.02850	1.73	0.99
12		0.072	0.073	0.085	0.07688	0.00696	-0.00053	-0.03	0.37
13		0.144	0.050	0.013	0.06890	0.06741	-0.00852	-0.52	3.62
14		0.080	0.077	0.069	0.07534	0.00555	-0.00207	-0.13	0.30
15		0.071	0.072	0.073	0.07233	0.00113	-0.00509	-0.31	0.06
16		0.080	0.075	0.076	0.07728	0.00263	-0.00013	-0.01	0.14
17		0.066	0.068	0.070	0.06800	0.00201	-0.00942	-0.57	0.11
18		0.088	0.079	0.077	0.08109	0.00606	0.00368	0.22	0.33
19		0.070	0.072	0.067	0.06960	0.00298	-0.00782	-0.47	0.16
20		0.068	0.065	0.072	0.06818	0.00355	-0.00924	-0.56	0.19
21		0.061	0.058	0.070	0.06280	0.00627	-0.01462	-0.89	0.34
22		0.093	0.100	0.125	0.10607	0.01710	0.02866	1.74	0.92
23		0.010	0.073		0.04175	0.04423	-0.03567	-2.17	2.37
24									
25									
26									
27									
28									

Average of Cell Averages

X-bar                    0.077

Standard Deviation of Cell Averages

s<sub>X-bar</sub>                0.016

Repeatability Standard Deviation

s<sub>r</sub>                    0.01864                0.022

Reproducibility Standard Deviation

s<sub>R</sub>                    0.022

*Shaded cells were removed after outlier analysis*

Binder BO

**PG 76-22**

**Jnr (0.1 kPa)**

Lab No.	Lab Name	Results, x			Average	s	d	h	k
		1	2	3					
1		0.306	0.312	0.306	0.30817	0.00315	0.03049	1.13	0.25
2		0.272	0.290	0.287	0.28327	0.00959	0.00559	0.21	0.77
3		0.251	0.269	0.283	0.26777	0.01604	-0.00991	-0.37	1.29
4		0.283	0.281	0.271	0.27810	0.00645	0.00042	0.02	0.52
5		0.291	0.279	0.279	0.28317	0.00696	0.00549	0.20	0.56
6		0.257	0.245	0.244	0.24853	0.00707	-0.02915	-1.08	0.57
7		0.287	0.324	0.300	0.30367	0.01877	0.02599	0.97	1.51
8		0.335	0.356	0.350	0.34700	0.01082	0.06932	2.58	0.87
9		0.306	0.270	0.273	0.28287	0.01972	0.00519	0.19	1.59
10		0.306	0.301	0.285	0.29737	0.01103	0.01969	0.73	0.89
11		0.298	0.292	0.292	0.29390	0.00312	0.01622	0.60	0.25
12		0.237	0.268	0.247	0.25037	0.01566	-0.02731	-1.02	1.26
13		0.296	0.296	0.290	0.29403	0.00335	0.01635	0.61	0.27
14		0.273	0.261	0.258	0.26410	0.00767	-0.01358	-0.51	0.62
15		0.215	0.258	0.268	0.24707	0.02845	-0.03061	-1.14	2.29
16		0.234	0.270	0.283	0.26240	0.02539	-0.01528	-0.57	2.05
17		0.301	0.302	0.288	0.29703	0.00751	0.01935	0.72	0.61
18		0.267	0.274	0.266	0.26910	0.00401	-0.00858	-0.32	0.32
19		0.275	0.270	0.280	0.27493	0.00495	-0.00275	-0.10	0.40
20		0.264	0.272	0.274	0.26993	0.00503	-0.00775	-0.29	0.41
21		0.275	0.284	0.282	0.28050	0.00493	0.00282	0.10	0.40
22		0.273	0.275	0.279	0.27567	0.00295	-0.00201	-0.07	0.24
23		0.200		0.216	0.20770	0.01117	-0.06998	-2.60	0.90
24									
25									
26									
27									
28									

Average of Cell Averages

X-bar                    0.278

Standard Deviation of Cell Averages

s<sub>X-bar</sub>                0.027

Repeatability Standard Deviation

s<sub>r</sub>                    0.01240            0.029

Reproducibility Standard Deviation

s<sub>R</sub>                    0.029

*Shaded cells were removed after outlier analysis*

# Binder BO

**PG 76-22**

**Jnr (3.2 kPa)**

Lab No.	Lab Name	Results, x			Average	s	d	h	k
		1	2	3					
1		0.349	0.356	0.350	0.35173	0.00399	0.04277	1.19	0.26
2		0.316	0.338	0.327	0.32673	0.01090	0.01777	0.50	0.70
3		0.279	0.301	0.317	0.29907	0.01909	-0.00990	-0.28	1.23
4		0.325	0.324	0.309	0.31943	0.00889	0.01047	0.29	0.57
5		0.331	0.320	0.316	0.32237	0.00800	0.01340	0.37	0.51
6		0.261	0.238	0.236	0.24473	0.01369	-0.06423	-1.79	0.88
7		0.320	0.359	0.331	0.33667	0.02011	0.02770	0.77	1.29
8		0.374	0.406	0.397	0.39233	0.01650	0.08337	2.33	1.06
9		0.352	0.306	0.311	0.32287	0.02515	0.01390	0.39	1.62
10		0.356	0.348	0.332	0.34510	0.01254	0.03613	1.01	0.81
11		0.315	0.315	0.316	0.31523	0.00093	0.00627	0.17	0.06
12		0.261	0.296	0.274	0.27697	0.01796	-0.03200	-0.89	1.15
13		0.340	0.331	0.315	0.32870	0.01297	0.01973	0.55	0.83
14		0.295	0.279	0.275	0.28283	0.01032	-0.02613	-0.73	0.66
15		0.236	0.289	0.301	0.27560	0.03455	-0.03337	-0.93	2.22
16		0.260	0.302	0.316	0.29253	0.02911	-0.01643	-0.46	1.87
17		0.335	0.339	0.316	0.32987	0.01244	0.02090	0.58	0.80
18		0.302	0.309	0.299	0.30320	0.00537	-0.00577	-0.16	0.34
19		0.292	0.289	0.299	0.29340	0.00492	-0.01557	-0.43	0.32
20		0.283	0.294	0.295	0.29093	0.00654	-0.01803	-0.50	0.42
21		0.313	0.323	0.322	0.31940	0.00563	0.01043	0.29	0.36
22		0.308	0.321	0.317	0.31520	0.00626	0.00623	0.17	0.40
23		0.209		0.234	0.22135	0.01789	-0.08762	-2.44	1.15
24									
25									
26									
27									
28									

Average of Cell Averages

X-bar                  0.309

Standard Deviation of Cell Averages

s<sub>X-bar</sub>            0.036

Repeatability Standard Deviation

s<sub>r</sub>                0.01557      0.038

Reproducibility Standard Deviation

s<sub>R</sub>                0.038

*Shaded cells were removed after outlier analysis*

# Binder BO

**PG 76-22**

## Recovery (0.1 kPa)

Lab No.	Lab Name	Results, x			Average	s	d	h	k
		1	2	3					
1		72.4	72.4	72.5	72.437	0.055	-0.077	-0.10	0.13
2		73.1	72.7	72.2	72.650	0.405	0.137	0.17	0.94
3		73.4	73.2	72.8	73.107	0.330	0.593	0.76	0.77
4		72.4	72.4	72.7	72.453	0.170	-0.060	-0.08	0.40
5		72.2	72.2	72.1	72.169	0.088	-0.344	-0.44	0.20
6		72.1	71.9	71.7	71.911	0.213	-0.602	-0.77	0.49
7		72.0	71.2	71.8	71.663	0.417	-0.850	-1.08	0.97
8		66.3	65.7	65.9					
9		73.7	72.9	73.6	73.391	0.387	0.877	1.12	0.90
10		72.4	72.8	73.1	72.763	0.330	0.250	0.32	0.77
11		70.4	70.8	71.0	70.725	0.292	-1.788	-2.28	0.68
12		73.6	73.2	72.6	73.124	0.524	0.611	0.78	1.22
13		73.1	72.3	72.2	72.507	0.517	-0.007	-0.01	1.20
14		73.3	73.7	73.7	73.594	0.249	1.081	1.38	0.58
15		74.3	73.3	73.1	73.575	0.657	1.062	1.35	1.53
16		73.5	72.8	72.8	73.042	0.402	0.528	0.67	0.94
17		72.3	72.2	73.1	72.534	0.518	0.021	0.03	1.21
18		73.9	73.3	73.6	73.579	0.277	1.065	1.36	0.64
19		71.3	70.9	71.0	71.074	0.164	-1.439	-1.83	0.38
20		72.0	71.4	71.5	71.606	0.309	-0.907	-1.16	0.72
21		72.3	72.2	71.9	72.144	0.216	-0.370	-0.47	0.50
22		72.5	72.3	72.2	72.320	0.114	-0.193	-0.25	0.26
23		72.1		73.8	72.925	1.209	0.412	0.52	2.81
24									
25									
26									
27									
28									

Average of Cell Averages

X-bar                    72.513

Standard Deviation of Cell Averages

s<sub>X-bar</sub>                0.785

Repeatability Standard Deviation

s<sub>r</sub>                    0.42972            0.860

Reproducibility Standard Deviation

s<sub>R</sub>                    0.860

*Shaded cells were removed after outlier analysis*

# Binder BO

**PG 76-22**

## Recovery (3.2 kPa)

Lab No.	Lab Name	Results, x			Average	s	d	h	k
		1	2	3					
1		69.6	69.4	69.5	69.49333	0.07024	-0.71686	-0.68	0.15
2		69.8	69.3	69.5	69.51000	0.26627	-0.70020	-0.67	0.58
3		70.8	70.5	69.9	70.41667	0.46069	0.20647	0.20	1.01
4		69.7	69.7	70.0	69.82333	0.14434	-0.38686	-0.37	0.32
5		69.2	69.0	69.2	69.14040	0.09748	-1.06980	-1.02	0.21
6		72.1	72.7	72.7	72.49543	0.38433	2.28524	2.17	0.84
7		69.3	68.1	68.5	68.64333	0.62148	-1.56686	-1.49	1.36
8		63.7	62.2	62.5					
9		70.5	70.2	70.7	70.46520	0.26682	0.25500	0.24	0.58
10		69.3	69.7	70.1	69.68000	0.40583	-0.53020	-0.50	0.89
11		69.1	69.0	68.9	69.02307	0.08548	-1.18713	-1.13	0.19
12		71.5	71.0	70.2	70.90833	0.65152	0.69814	0.66	1.43
13		69.9	69.8	70.5	70.08000	0.35157	-0.13020	-0.12	0.77
14		71.7	72.5	72.5	72.23723	0.42810	2.02704	1.93	0.94
15		72.2	70.7	70.4	71.13810	0.96660	0.92790	0.88	2.12
16		71.2	70.4	70.3	70.63097	0.49144	0.42077	0.40	1.08
17		69.8	69.5	71.2	70.14270	0.89821	-0.06750	-0.06	1.97
18		71.4	70.6	71.1	71.03350	0.39367	0.82330	0.78	0.86
19		69.7	69.1	69.3	69.37063	0.32566	-0.83956	-0.80	0.71
20		70.5	69.6	69.9	70.01333	0.45107	-0.19686	-0.19	0.99
21		69.2	69.1	68.7	68.97807	0.27142	-1.23213	-1.17	0.59
22		70.0	69.4	69.5	69.61667	0.30665	-0.59353	-0.56	0.67
23		71.5		72.0	71.78400	0.37335	1.57380	1.50	0.82
24									
25									
26									
27									
28									

Average of Cell Averages

X-bar                    70.210

Standard Deviation of Cell Averages

s<sub>X-bar</sub>                1.051

Repeatability Standard Deviation

s<sub>r</sub>                    0.45632            1.115

Reproducibility Standard Deviation

s<sub>R</sub>                    1.115

*Shaded cells were removed after outlier analysis*

# Binder BO

**PG 76-22**

**Jnr, Diff**

Lab No.	Lab Name	Results, x			Average	s	d	h	k
		1	2	3					
1		0.139	0.143	0.143	0.14127	0.00240	0.02613	0.93	0.18
2		0.160	0.163	0.138	0.15343	0.01393	0.03830	1.36	1.06
3		0.113	0.117	0.121	0.11673	0.00420	0.00160	0.06	0.32
4		0.151	0.152	0.142	0.14863	0.00544	0.03350	1.19	0.41
5		0.137	0.147	0.132	0.13853	0.00754	0.02340	0.83	0.57
6		0.015	0.028	0.035					
7		0.130	0.110	0.103	0.11423	0.01401	-0.00090	-0.03	1.07
8		0.105	0.125	0.117	0.11573	0.00981	0.00060	0.02	0.75
9		0.151	0.133	0.139	0.14105	0.00941	0.02592	0.92	0.72
10		0.140	0.134	0.140	0.13837	0.00352	0.02323	0.83	0.27
11		0.058	0.077	0.083	0.07264	0.01309	-0.04249	-1.51	0.99
12		0.100	0.107	0.111	0.10601	0.00523	-0.00912	-0.32	0.40
13		0.152	0.117	0.084	0.11762	0.03368	0.00248	0.09	2.56
14		0.080	0.067	0.066	0.07092	0.00794	-0.04422	-1.57	0.60
15		0.101	0.120	0.123	0.11455	0.01223	-0.00058	-0.02	0.93
16		0.111	0.115	0.117	0.11463	0.00320	-0.00050	-0.02	0.24
17		0.114	0.122	0.095	0.11027	0.01405	-0.00487	-0.17	1.07
18		0.129	0.129	0.121	0.12669	0.00468	0.01156	0.41	0.36
19		0.061	0.072	0.068	0.06715	0.00557	-0.04799	-1.71	0.42
20		0.073	0.082	0.078	0.07755	0.00468	-0.03758	-1.34	0.36
21		0.138	0.136	0.142	0.13863	0.00278	0.02350	0.84	0.21
22		0.128	0.167	0.135	0.14333	0.02103	0.02820	1.00	1.60
23		0.045		0.085	0.06494	0.02885	-0.05019	-1.78	2.19
24									
25									
26									
27									
28									

Average of Cell Averages

X-bar                    0.115

Standard Deviation of Cell Averages

s<sub>X-bar</sub>                0.028

Repeatability Standard Deviation

s<sub>r</sub>                    0.01316            0.030

Reproducibility Standard Deviation

s<sub>R</sub>                    0.030

*Shaded cells were removed after outlier analysis*

Binder BR

**PG 76-22**

**Jnr (0.1 kPa)**

Lab No.	Lab Name	Results, x					h	k
		1	2	3	Average	s		
1		0.119	0.133	0.122	0.12437	0.00726	-0.00470	-0.36
2		0.124	0.128	0.134	0.12863	0.00502	-0.00043	-0.03
3		0.124	0.124	0.129	0.12570	0.00286	-0.00337	-0.26
4		0.122	0.125	0.122	0.12317	0.00159	-0.00590	-0.45
5		0.136	0.149	0.131	0.13873	0.00916	0.00967	0.74
6		0.145	0.127	0.115	0.12893	0.01525	-0.00013	-0.01
7		0.141	0.142	0.143	0.14200	0.00100	0.01293	1.00
8		0.166	0.174	0.173	0.17100	0.00436	0.04193	3.23
9		0.121	0.124	0.115	0.11983	0.00480	-0.00923	-0.71
10		0.136	0.135	0.129	0.13343	0.00395	0.00437	0.34
11		0.155	0.136	0.112	0.13437	0.02131	0.00530	0.41
12		0.124	0.122	0.127	0.12450	0.00256	-0.00457	-0.35
13		0.137	0.134	0.129	0.13333	0.00381	0.00427	0.33
14		0.137	0.138	0.133	0.13577	0.00268	0.00670	0.52
15		0.105	0.105	0.104	0.10463	0.00047	-0.02443	-1.88
16		0.121	0.125	0.126	0.12377	0.00294	-0.00530	-0.41
17		0.128	0.127	0.126	0.12723	0.00110	-0.00183	-0.14
18		0.128	0.130	0.129	0.12917	0.00110	0.00010	0.01
19		0.119	0.129	0.123	0.12387	0.00497	-0.00520	-0.40
20		0.124	0.131	0.129	0.12797	0.00357	-0.00110	-0.08
21		0.140	0.144	0.146	0.14313	0.00303	0.01407	1.08
22		0.113	0.109	0.111	0.11093	0.00191	-0.01813	-1.40
23		0.114	0.115	0.113	0.11403	0.00097	-0.01503	-1.16
24								0.15
25								
26								
27								
28								

Average of Cell Averages

X-bar                    0.129

Standard Deviation of Cell Averages

S<sub>X-bar</sub>                0.013

Repeatability Standard Deviation

S<sub>r</sub>                    0.00662            0.014

Reproducibility Standard Deviation

S<sub>R</sub>                    0.014

*Shaded cells were removed after outlier analysis*

# Binder BR

**PG 76-22**

**Jnr (3.2 kPa)**

Lab No.	Lab Name	Results, x			Average	s	d	h	k
		1	2	3					
1		0.128	0.144	0.131	0.13417	0.00832	-0.00286	-0.19	1.53
2		0.132	0.137	0.143	0.13733	0.00565	0.00031	0.02	1.04
3		0.132	0.131	0.137	0.13333	0.00335	-0.00369	-0.25	0.62
4		0.134	0.138	0.135	0.13543	0.00199	-0.00159	-0.11	0.37
5		0.145	0.162	0.140	0.14923	0.01142	0.01221	0.82	2.10
6		0.125	0.113	0.100	0.11297	0.01260	-0.02406	-1.61	2.32
7		0.148	0.146	0.154	0.14933	0.00416	0.01231	0.83	0.77
8		0.177	0.187	0.184	0.18267	0.00513	0.04564	3.06	0.95
9		0.133	0.132	0.134	0.13260	0.00100	-0.00442	-0.30	0.18
10		0.150	0.146	0.144	0.14660	0.00302	0.00958	0.64	0.56
11		0.148	0.134	0.130	0.13763	0.00938	0.00061	0.04	1.73
12		0.132	0.130	0.136	0.13240	0.00321	-0.00462	-0.31	0.59
13		0.151	0.147	0.140	0.14617	0.00550	0.00914	0.61	1.01
14		0.143	0.145	0.139	0.14223	0.00283	0.00521	0.35	0.52
15		0.111	0.112	0.110	0.11100	0.00078	-0.02602	-1.75	0.14
16		0.129	0.133	0.136	0.13257	0.00349	-0.00446	-0.30	0.64
17		0.138	0.142	0.134	0.13787	0.00431	0.00084	0.06	0.79
18		0.138	0.140	0.138	0.13843	0.00095	0.00141	0.09	0.17
19		0.130	0.132	0.127	0.12970	0.00271	-0.00732	-0.49	0.50
20		0.131	0.138	0.130	0.13273	0.00423	-0.00429	-0.29	0.78
21		0.151	0.157	0.159	0.15567	0.00405	0.01864	1.25	0.75
22		0.125	0.120	0.124	0.12270	0.00267	-0.01432	-0.96	0.49
23		0.119	0.118	0.120	0.11873	0.00121	-0.01829	-1.23	0.22
24									
25									
26									
27									
28									

Average of Cell Averages

X-bar                    0.137

Standard Deviation of Cell Averages

s<sub>X-bar</sub>                0.015

Repeatability Standard Deviation

s<sub>r</sub>                    0.00543            0.016

Reproducibility Standard Deviation

s<sub>R</sub>                    0.016

*Shaded cells were removed after outlier analysis*

# Binder BR

**PG 76-22**

## Recovery (0.1 kPa)

Lab No.	Lab Name	Results, x			Average	s	d	h	k
		1	2	3					
1		77.4	76.7	77.2	77.093	0.322	0.363	0.43	0.39
2		76.6	76.5	76.2	76.403	0.206	-0.327	-0.39	0.25
3		77.4	77.3	77.0	77.223	0.195	0.493	0.59	0.24
4		76.9	76.9	77.0	76.927	0.083	0.196	0.23	0.10
5		76.1	75.9	76.5	76.169	0.303	-0.562	-0.67	0.37
6		74.2	75.7	76.1	75.346	1.013	-1.385	-1.65	1.23
7		75.4	75.1	75.5	75.330	0.192	-1.401	-1.67	0.23
8		70.5	70.7	70.4					
9		77.0	76.4	78.7	77.375	1.191	0.644	0.77	1.44
10		77.4	77.2	77.5	77.377	0.167	0.646	0.77	0.20
11		74.0	76.8	80.6	77.123	3.351	0.393	0.47	4.06
12		77.2	77.3	77.0	77.142	0.145	0.412	0.49	0.18
13		77.5	77.2	77.1	77.257	0.175	0.526	0.63	0.21
14		76.8	76.7	77.0	76.815	0.143	0.084	0.10	0.17
15		77.7	77.7	77.7	77.685	0.034	0.954	1.14	0.04
16		76.9	76.8	76.8	76.825	0.028	0.095	0.11	0.03
17		77.0	76.9	77.3	77.074	0.242	0.343	0.41	0.29
18		77.4	77.3	77.0	77.237	0.201	0.506	0.60	0.24
19		75.9	75.0	75.3	75.418	0.469	-1.312	-1.56	0.57
20		76.4	75.4	75.6	75.838	0.517	-0.892	-1.06	0.63
21		75.3	75.1	74.7	75.032	0.301	-1.699	-2.03	0.36
22		77.6	78.0	78.1	77.883	0.225	1.153	1.37	0.27
23		77.6	77.1	77.8	77.503	0.393	0.773	0.92	0.48
24									
25									
26									
27									
28									

Average of Cell Averages

X-bar

76.731

Standard Deviation of Cell Averages

s<sub>X-bar</sub>

0.839

Repeatability Standard Deviation

s<sub>r</sub>

0.82517      1.076

Reproducibility Standard Deviation

s<sub>R</sub>

1.076

*Shaded cells were removed after outlier analysis*

# Binder BR

**PG 76-22**

## Recovery (3.2 kPa)

Lab No.	Lab Name	Results, x			Average	s	d	h	k
		1	2	3					
1		76.1	75.3	75.8	75.70667	0.41187	0.13784	0.14	0.90
2		75.6	75.3	75.0	75.29000	0.27514	-0.27883	-0.27	0.60
3		76.0	76.1	75.6	75.90333	0.24542	0.33451	0.33	0.53
4		75.6	75.4	75.7	75.56000	0.12530	-0.00883	-0.01	0.27
5		74.7	74.2	75.3	74.71527	0.52715	-0.85356	-0.84	1.15
6		77.1	77.9	78.7	77.90983	0.75886	2.34101	2.30	1.65
7		73.5	73.5	73.4	73.42333	0.06351	-2.14549	-2.10	0.14
8		68.8	68.9	68.4					
9		75.2	75.2	75.6	75.34570	0.19503	-0.22313	-0.22	0.42
10		75.8	76.1	75.7	75.85333	0.19553	0.28451	0.28	0.43
11		74.8	76.8	77.4	76.34917	1.36693	0.78034	0.77	2.98
12		76.1	76.1	75.7	75.95790	0.25582	0.38907	0.38	0.56
13		75.4	75.5	75.5	75.47667	0.04041	-0.09216	-0.09	0.09
14		75.9	75.8	76.1	75.93860	0.14580	0.36977	0.36	0.32
15		76.5	76.4	76.5	76.49980	0.08126	0.93097	0.91	0.18
16		75.5	75.5	75.3	75.44533	0.09615	-0.12349	-0.12	0.21
17		75.6	74.8	76.2	75.53087	0.71625	-0.03796	-0.04	1.56
18		76.0	76.1	75.7	75.93650	0.22842	0.36767	0.36	0.50
19		74.9	74.2	74.4	74.47273	0.37144	-1.09609	-1.07	0.81
20		75.3	74.4	75.8	75.15287	0.71005	-0.41596	-0.41	1.55
21		73.6	73.2	72.8	73.19227	0.37722	-2.37656	-2.33	0.82
22		75.8	76.3	76.0	76.05333	0.22030	0.48451	0.47	0.48
23		76.9	76.8	76.7	76.80067	0.12017	1.23184	1.21	0.26
24									
25									
26									
27									
28									

Average of Cell Averages

X-bar                    75.569

Standard Deviation of Cell Averages

s<sub>X-bar</sub>                1.020

Repeatability Standard Deviation

s<sub>r</sub>                    0.45928                    1.087

Reproducibility Standard Deviation

s<sub>R</sub>                    1.087

*Shaded cells were removed after outlier analysis*

## Binder BR

PG 76-22

## Jnr, Diff

Lab No.	Lab Name	Results, x			Average	s	d	h	k
		1	2	3					
1		0.076	0.083	0.078	0.07887	0.00380	0.00561	0.23	0.16
2		0.063	0.072	0.068	0.06754	0.00459	-0.00572	-0.24	0.20
3		0.061	0.058	0.063	0.06073	0.00251	-0.01253	-0.52	0.11
4		0.098	0.102	0.100	0.09984	0.00220	0.02658	1.11	0.09
5		0.070	0.088	0.068	0.07557	0.01124	0.00231	0.10	0.48
6		0.136	0.107	0.127	0.12333	0.01455	0.05007	2.10	0.62
7		0.050	0.030	0.078	0.05247	0.02425	-0.02079	-0.87	1.04
8		0.063	0.067	0.061	0.06373	0.00281	-0.00953	-0.40	0.12
9		0.094	0.063	0.166	0.10776	0.05320	0.03450	1.44	2.28
10		0.092	0.074	0.103	0.08985	0.01462	0.01659	0.69	0.63
11		0.042	0.013	0.161	0.07216	0.07867	-0.00110	-0.05	3.37
12		0.060	0.061	0.068	0.06319	0.00430	-0.01007	-0.42	0.18
13		0.107	0.094	0.087	0.09583	0.01028	0.02257	0.94	0.44
14		0.047	0.048	0.047	0.04743	0.00042	-0.02583	-1.08	0.02
15		0.061	0.064	0.058	0.06084	0.00315	-0.01242	-0.52	0.13
16		0.069	0.069	0.075	0.07106	0.00365	-0.00220	-0.09	0.16
17		0.072	0.118	0.060	0.08317	0.03029	0.00991	0.41	1.30
18		0.073	0.070	0.072	0.07176	0.00180	-0.00150	-0.06	0.08
19		0.032	0.025	0.029	0.02842	0.00351	-0.04484	-1.88	0.15
20		0.054	0.053	0.004	0.03721	0.02840	-0.03605	-1.51	1.22
21		0.081	0.091	0.090	0.08730	0.00582	0.01404	0.59	0.25
22		0.107	0.099	0.112	0.10576	0.00621	0.03250	1.36	0.27
23		0.042	0.022	0.060	0.04117	0.01906	-0.03209	-1.34	0.82
24									
25									
26									
27									
28									

Average of Cell Averages

X-bar                    0.073

Standard Deviation of Cell Averages

S<sub>X-bar</sub>                0.024

Repeatability Standard Deviation

S<sub>r</sub>                    0.02335      0.031

Reproducibility Standard Deviation

S<sub>R</sub>                    0.031*Shaded cells were removed after outlier analysis*

Binder CO

**PG 70-22**

**Jnr (0.1 kPa)**

Lab No.	Lab Name	Results, x								
		1	2	3	Average	s	d	h	k	
1		0.860	0.881	0.899	0.87977	0.01957	0.09917	1.49	0.55	
2		0.806	0.784	0.802	0.79727	0.01132	0.01667	0.25	0.32	
3		0.819	0.808	0.782	0.80290	0.01922	0.02230	0.33	0.54	
4		0.783	0.739	0.760	0.76060	0.02182	-0.02000	-0.30	0.61	
5		0.824	0.730	0.762	0.77173	0.04778	-0.00886	-0.13	1.33	
6		0.663	0.600	0.626	0.62943	0.03165	-0.15116	-2.27	0.88	
7		0.831	0.810	0.879	0.84000	0.03537	0.05940	0.89	0.99	
8		0.905	0.879	0.874	0.88600	0.01664	0.10540	1.58	0.46	
9		0.814	0.814	0.661	0.76303	0.08802	-0.01756	-0.26	2.46	
10		0.856	0.820	0.819	0.83147	0.02108	0.05087	0.76	0.59	
11		0.828	0.872	0.740	0.81343	0.06731	0.03284	0.49	1.88	
12		0.738	0.755	0.700	0.73087	0.02825	-0.04973	-0.75	0.79	
13		0.818	0.869	0.825	0.83713	0.02746	0.05654	0.85	0.77	
14		0.801	0.799	0.766	0.78853	0.01978	0.00794	0.12	0.55	
15		0.744	0.686	0.780	0.73683	0.04711	-0.04376	-0.66	1.31	
16		0.793	0.731	0.746	0.75667	0.03218	-0.02393	-0.36	0.90	
17		0.803	0.857	0.845	0.83510	0.02803	0.05450	0.82	0.78	
18		0.781	0.771	0.770	0.77390	0.00629	-0.00670	-0.10	0.18	
19		0.843	0.777	0.780	0.80003	0.03742	0.01944	0.29	1.04	
20		0.776	0.831	0.763	0.79037	0.03613	0.00977	0.15	1.01	
21		0.784	0.746	0.816	0.78203	0.03512	0.00144	0.02	0.98	
22		0.723	0.770	0.746	0.74617	0.02345	-0.03443	-0.52	0.65	
23		0.592	0.609		0.60045	0.01195	-0.18015	-2.70	0.33	
24										
25										
26										
27										
28										

Average of Cell Averages

X-bar                    0.781

Standard Deviation of Cell Averages

s<sub>X-bar</sub>                0.067

Repeatability Standard Deviation

s<sub>r</sub>                    0.03584            0.073

Reproducibility Standard Deviation

s<sub>R</sub>                    0.073

*Shaded cells were removed after outlier analysis*

## Binder CO

PG 70-22

Jnr (3.2 kPa)

Lab No.	Lab Name	Results, x					h	k
		1	2	3	Average	s		
1		1.201	1.234	1.258	1.23100	0.02862	0.15865	1.52
2		1.094	1.137	1.122	1.11767	0.02183	0.04531	0.44
3		1.121	1.116	1.063	1.10000	0.03214	0.02765	0.27
4		1.107	1.050	1.070	1.07567	0.02892	0.00331	0.03
5		1.142	0.998	1.037	1.05907	0.07469	-0.01329	-0.13
6		0.862	0.779	0.806	0.81593	0.04245	-0.25642	-2.46
7		1.163	1.087	1.185	1.14500	0.05142	0.07265	0.70
8		1.177	1.181	1.130	1.16267	0.02836	0.09031	0.87
9		1.130	1.126	1.042	1.09920	0.05000	0.02685	0.26
10		1.246	1.175	1.163	1.19467	0.04486	0.12231	1.18
11		1.125	1.195	0.983	1.10087	0.10820	0.02851	0.27
12		1.053	1.047	0.955	1.01820	0.05518	-0.05415	-0.52
13		1.117	1.175	1.146	1.14600	0.02900	0.07365	0.71
14		1.107	1.106	1.052	1.08803	0.03164	0.01568	0.15
15		1.059	0.924	1.075	1.01900	0.08301	-0.05335	-0.51
16		1.079	0.991	1.010	1.02683	0.04621	-0.04552	-0.44
17		1.104	1.189	1.175	1.15613	0.04559	0.08378	0.81
18		1.095	1.061	1.086	1.08050	0.01759	0.00815	0.08
19		1.160	1.051	1.057	1.08940	0.06138	0.01705	0.16
20		1.046	1.132	1.034	1.07047	0.05325	-0.00189	-0.02
21		1.078	1.009	1.121	1.06923	0.05686	-0.00312	-0.03
22		0.986	1.056	1.044	1.02867	0.03743	-0.04369	-0.42
23		0.734	0.806		0.76995	0.05028	-0.30240	-2.91
24								0.98
25								
26								
27								
28								

Average of Cell Averages

X-bar                    1.072

Standard Deviation of Cell Averages

S<sub>X-bar</sub>                0.104

Repeatability Standard Deviation

S<sub>r</sub>                    0.05119                0.112

Reproducibility Standard Deviation

S<sub>R</sub>                    0.112*Shaded cells were removed after outlier analysis*

Binder CO

**PG 70-22**

**Recovery (0.1 kPa)**

Lab No.	Lab Name	Results, x			Average	s	d	h	k
		1	2	3					
1		49.1	49.0	49.2	49.097	0.091	-0.482	-0.29	0.11
2		49.0	52.0	50.3	50.433	1.492	0.855	0.51	1.82
3		50.3	51.0	50.4	50.557	0.421	0.978	0.59	0.51
4		50.3	51.2	50.4	50.623	0.522	1.045	0.63	0.64
5		50.0	49.3	48.9	49.442	0.554	-0.137	-0.08	0.68
6		49.7	50.8	50.0	50.179	0.584	0.600	0.36	0.71
7		51.4	48.9	48.1	49.490	1.729	-0.089	-0.05	2.11
8		45.0	44.9	44.9	44.947	0.070	-4.632	-2.78	0.09
9		50.3	50.1	52.2	50.857	1.163	1.278	0.77	1.42
10		50.3	50.5	49.5	50.127	0.524	0.548	0.33	0.64
11		47.6	47.2	48.6	47.806	0.757	-1.773	-1.07	0.93
12		52.9	51.4	51.1	51.809	0.944	2.231	1.34	1.15
13		49.0	48.1	49.4	48.830	0.675	-0.749	-0.45	0.82
14		50.7	50.5	51.0	50.728	0.249	1.149	0.69	0.30
15		52.5	50.6	50.1	51.092	1.239	1.513	0.91	1.51
16		49.5	50.6	50.1	50.094	0.553	0.516	0.31	0.67
17		49.9	49.3	49.9	49.711	0.324	0.132	0.08	0.40
18		50.8	50.4	51.2	50.821	0.398	1.243	0.75	0.49
19		46.4	46.0	46.4	46.239	0.225	-3.340	-2.01	0.28
20		47.3	46.2	47.5	46.977	0.710	-2.602	-1.56	0.87
21		49.6	49.5	48.7	49.298	0.506	-0.281	-0.17	0.62
22		51.1	49.9	51.0	50.653	0.665	1.075	0.65	0.81
23		49.5	51.5		50.500	1.457	0.921	0.55	1.78
24									
25									
26									
27									
28									

Average of Cell Averages

X-bar                  49.579

Standard Deviation of Cell Averages

s<sub>X-bar</sub>              1.664

Repeatability Standard Deviation

s<sub>r</sub>                  0.81864                  1.793

Reproducibility Standard Deviation

s<sub>R</sub>                  1.793

*Shaded cells were removed after outlier analysis*

Binder CO

**PG 70-22**

**Recovery (3.2 kPa)**

Lab No.	Lab Name	Results, x			Average	s	d	h	k
		1	2	3					
1		33.8	33.6	33.8	33.72000	0.12166	-1.48919	-0.75	0.14
2		35.6	35.8	35.5	35.63333	0.12342	0.42414	0.21	0.14
3		35.6	36.3	36.0	35.94000	0.35043	0.73081	0.37	0.39
4		35.5	36.4	35.9	35.95000	0.44136	0.74081	0.37	0.49
5		34.9	35.0	34.4	34.80643	0.32772	-0.40276	-0.20	0.37
6		37.4	38.9	38.2	38.17013	0.78130	2.96094	1.49	0.87
7		36.2	34.4	33.1	34.58333	1.56896	-0.62586	-0.31	1.76
8		32.4	29.5	32.5	31.46000	1.70696	-3.74919	-1.89	1.91
9		35.3	35.1	35.7	35.36990	0.27368	0.16071	0.08	0.31
10		33.8	34.7	34.5	34.35000	0.45640	-0.85919	-0.43	0.51
11		35.9	37.0	32.8	35.21307	2.18167	0.00388	0.00	2.44
12		37.6	36.8	37.4	37.27383	0.45013	2.06464	1.04	0.50
13		34.8	34.6	34.3	34.53667	0.23587	-0.67252	-0.34	0.26
14		36.0	35.5	36.6	36.00970	0.57537	0.80051	0.40	0.64
15		37.1	37.0	35.2	36.43110	1.08516	1.22191	0.61	1.21
16		35.2	36.9	36.4	36.19080	0.87118	0.98161	0.49	0.97
17		35.2	33.9	34.6	34.58737	0.64850	-0.62182	-0.31	0.73
18		35.6	35.7	36.0	35.77127	0.19685	0.56208	0.28	0.22
19		30.3	30.8	31.2	30.75393	0.46610	-4.45526	-2.24	0.52
20		33.0	31.0	32.9	32.30110	1.11147	-2.90809	-1.46	1.24
21		34.6	35.3	33.5	34.46610	0.90880	-0.74309	-0.37	1.02
22		37.5	36.2	36.1	36.62333	0.76950	1.41414	0.71	0.86
23		40.4	39.0		39.67000	1.01823	4.46081	2.24	1.14
24									
25									
26									
27									
28									

Average of Cell Averages

X-bar    35.209

Standard Deviation of Cell Averages

s<sub>X-bar</sub>    1.987

Repeatability Standard Deviation

s<sub>r</sub>    0.89385                                    2.117

Reproducibility Standard Deviation

s<sub>R</sub>    2.117

*Shaded cells were removed after outlier analysis*

Binder CO

**PG 70-22**

**Jnr, Diff**

Lab No.	Lab Name	Results, x			Average	s	d	h	k
		1	2	3					
1		0.398	0.401	0.400	0.39937	0.00164	0.03570	0.77	0.05
2		0.358	0.449	0.399	0.40230	0.04552	0.03863	0.84	1.42
3		0.368	0.382	0.360	0.36970	0.01100	0.00603	0.13	0.34
4		0.414	0.421	0.409	0.41457	0.00576	0.05090	1.10	0.18
5		0.387	0.367	0.362	0.37180	0.01308	0.00813	0.18	0.41
6		0.301	0.299	0.288	0.29628	0.00709	-0.06739	-1.46	0.22
7		0.400	0.345	0.348	0.36430	0.03104	0.00063	0.01	0.97
8		0.230	0.256	0.227	0.23777	0.01589	-0.12590	-2.73	0.50
9		0.389	0.384	0.575	0.44891	0.10887	0.08524	1.85	3.41
10		0.313	0.302	0.296	0.30373	0.00854	-0.05993	-1.30	0.27
11		0.359	0.370	0.328	0.35213	0.02182	-0.01154	-0.25	0.68
12		0.428	0.386	0.364	0.39273	0.03233	0.02906	0.63	1.01
13		0.365	0.352	0.389	0.36887	0.01851	0.00520	0.11	0.58
14		0.382	0.384	0.373	0.37972	0.00561	0.01605	0.35	0.18
15		0.422	0.346	0.378	0.38205	0.03846	0.01838	0.40	1.20
16		0.361	0.355	0.354	0.35696	0.00350	-0.00671	-0.15	0.11
17		0.374	0.388	0.390	0.38427	0.00870	0.02060	0.45	0.27
18		0.402	0.376	0.411	0.39616	0.01820	0.03249	0.70	0.57
19		0.376	0.353	0.355	0.36133	0.01268	-0.00234	-0.05	0.40
20		0.348	0.361	0.354	0.35414	0.00662	-0.00953	-0.21	0.21
21		0.376	0.351	0.373	0.36683	0.01343	0.00317	0.07	0.42
22		0.364	0.372	0.400	0.37877	0.01881	0.01510	0.33	0.59
23		0.241	0.323		0.28170	0.05820	-0.08197	-1.78	1.82
24									
25									
26									
27									
28									

Average of Cell Averages

X-bar                    0.364

Standard Deviation of Cell Averages

s<sub>X-bar</sub>                0.046

Repeatability Standard Deviation

s<sub>r</sub>                    0.03197      0.053

Reproducibility Standard Deviation

s<sub>R</sub>                    0.053

*Shaded cells were removed after outlier analysis*

Binder CR

**PG 70-22**

**Jnr (0.1 kPa)**

Lab No.	Lab Name	Results, x			Average	s	d	h	k
		1	2	3					
1		0.338	0.343	0.351	0.34387	0.00657	-0.00505	-0.17	0.48
2		0.351	0.363	0.369	0.36087	0.00922	0.01195	0.39	0.67
3		0.344	0.345	0.338	0.34220	0.00383	-0.00672	-0.22	0.28
4		0.335	0.339	0.336	0.33640	0.00219	-0.01252	-0.41	0.16
5		0.315	0.297	0.285	0.29913	0.01521	-0.04978	-1.64	1.10
6		0.370	0.317	0.294	0.32677	0.03875	-0.02215	-0.73	2.81
7		0.371	0.369	0.384	0.37467	0.00814	0.02575	0.85	0.59
8		0.380	0.401	0.381	0.38733	0.01185	0.03842	1.27	0.86
9		0.342	0.323	0.343	0.33597	0.01150	-0.01295	-0.43	0.83
10		0.387	0.381	0.373	0.38027	0.00731	0.03135	1.04	0.53
11		0.381	0.374	0.387	0.38073	0.00637	0.03182	1.05	0.46
12		0.333	0.344	0.354	0.34387	0.01081	-0.00505	-0.17	0.78
13		0.389	0.363	0.373	0.37497	0.01286	0.02605	0.86	0.93
14		0.366	0.352	0.353	0.35700	0.00764	0.00808	0.27	0.55
15		0.282	0.287	0.293	0.28700	0.00577	-0.06192	-2.05	0.42
16		0.326	0.331	0.316	0.32450	0.00743	-0.02442	-0.81	0.54
17		0.332	0.319	0.335	0.32860	0.00865	-0.02032	-0.67	0.63
18		0.351	0.356	0.357	0.35443	0.00317	0.00552	0.18	0.23
19		0.362	0.420	0.431	0.40437	0.03726	0.05545	1.83	2.70
20		0.377	0.379	0.394	0.38313	0.00903	0.03422	1.13	0.65
21		0.359	0.370	0.372	0.36703	0.00690	0.01812	0.60	0.50
22		0.324	0.325	0.323	0.32403	0.00086	-0.02488	-0.82	0.06
23		0.309	0.313	0.302	0.30797	0.00572	-0.04095	-1.35	0.41
24									
25									
26									
27									
28									

Average of Cell Averages

X-bar                    0.349

Standard Deviation of Cell Averages

s<sub>X-bar</sub>                0.030

Repeatability Standard Deviation

s<sub>r</sub>                    0.01380            0.032

Reproducibility Standard Deviation

s<sub>R</sub>                    0.032

*Shaded cells were removed after outlier analysis*

Binder CR

**PG 70-22**

**Jnr (3.2 kPa)**

Lab No.	Lab Name	Results, x			Average	s	d	h	k
		1	2	3					
1		0.411	0.421	0.431	0.42100	0.01030	-0.00383	-0.09	0.54
2		0.446	0.459	0.451	0.45183	0.00639	0.02700	0.65	0.34
3		0.417	0.415	0.406	0.41280	0.00577	-0.01203	-0.29	0.30
4		0.416	0.421	0.418	0.41817	0.00225	-0.00666	-0.16	0.12
5		0.378	0.359	0.345	0.36020	0.01656	-0.06463	-1.56	0.87
6		0.436	0.364	0.332	0.37703	0.05323	-0.04780	-1.15	2.81
7		0.434	0.436	0.453	0.44100	0.01044	0.01617	0.39	0.55
8		0.499	0.471	0.443	0.47100	0.02800	0.04617	1.11	1.48
9		0.415	0.397	0.435	0.41570	0.01895	-0.00913	-0.22	1.00
10		0.494	0.482	0.474	0.48313	0.01032	0.05830	1.40	0.54
11		0.464	0.452	0.457	0.45740	0.00612	0.03257	0.78	0.32
12		0.404	0.420	0.436	0.41973	0.01585	-0.00510	-0.12	0.84
13		0.477	0.471	0.450	0.46613	0.01391	0.04130	0.99	0.73
14		0.450	0.431	0.433	0.43787	0.01081	0.01304	0.31	0.57
15		0.337	0.340	0.349	0.34197	0.00621	-0.08286	-1.99	0.33
16		0.393	0.398	0.381	0.39053	0.00850	-0.03430	-0.83	0.45
17		0.400	0.385	0.404	0.39627	0.01026	-0.02856	-0.69	0.54
18		0.437	0.438	0.438	0.43770	0.00062	0.01287	0.31	0.03
19		0.440	0.518	0.531	0.49607	0.04917	0.07124	1.71	2.59
20		0.455	0.466	0.485	0.46853	0.01482	0.04370	1.05	0.78
21		0.444	0.459	0.459	0.45410	0.00875	0.02927	0.70	0.46
22		0.386	0.395	0.379	0.38647	0.00778	-0.03836	-0.92	0.41
23		0.370	0.367	0.362	0.36647	0.00438	-0.05836	-1.40	0.23
24									
25									
26									
27									
28									

Average of Cell Averages

X-bar 0.425

Standard Deviation of Cell Averages

s<sub>X-bar</sub> 0.042

Repeatability Standard Deviation

s<sub>r</sub> 0.01895 0.044

Reproducibility Standard Deviation

s<sub>R</sub> 0.044

*Shaded cells were removed after outlier analysis*

Binder CR

**PG 70-22**

**Recovery (0.1 kPa)**

Lab No.	Lab Name	Results, x			Average	s	d	h	k
		1	2	3					
1		56.0	55.6	55.3	55.630	0.372	0.342	0.19	0.55
2		57.3	55.9	54.8	56.010	1.253	0.722	0.40	1.84
3		56.3	56.1	56.2	56.200	0.106	0.912	0.50	0.16
4		55.9	55.5	55.9	55.783	0.237	0.496	0.27	0.35
5		57.3	57.1	58.2	57.512	0.608	2.225	1.22	0.89
6		54.2	54.7	56.4	55.087	1.172	-0.201	-0.11	1.72
7		53.1	53.6	53.6	53.427	0.318	-1.861	-1.02	0.47
8		51.7	50.7	51.1	51.147	0.496	-4.141	-2.27	0.73
9		56.0	57.6	56.4	56.652	0.810	1.364	0.75	1.19
10		54.4	54.9	55.4	54.920	0.495	-0.368	-0.20	0.73
11		53.9	54.5	53.5	53.970	0.503	-1.317	-0.72	0.74
12		56.2	55.8	55.4	55.768	0.380	0.480	0.26	0.56
13		54.5	57.0	54.8	55.447	1.381	0.159	0.09	2.03
14		55.8	56.3	56.0	56.007	0.274	0.719	0.39	0.40
15		57.4	57.1	57.0	57.147	0.225	1.859	1.02	0.33
16		57.1	57.1	57.5	57.208	0.228	1.921	1.05	0.34
17		57.2	56.8	56.9	56.980	0.214	1.693	0.93	0.31
18		56.7	56.0	55.9	56.179	0.425	0.892	0.49	0.62
19		53.0	50.4	50.2	51.180	1.567	-4.108	-2.25	2.30
20		52.4	51.9	51.7	52.010	0.344	-3.278	-1.80	0.51
21		55.2	54.5	54.7	54.778	0.384	-0.510	-0.28	0.56
22		55.3	56.2	55.5	55.637	0.484	0.349	0.19	0.71
23		56.9	56.5	57.4	56.937	0.475	1.649	0.90	0.70
24									
25									
26									
27									
28									

Average of Cell Averages

X-bar

55.288

Standard Deviation of Cell Averages

s<sub>X-bar</sub>

1.826

Repeatability Standard Deviation

s<sub>r</sub>

0.68068      1.909

Reproducibility Standard Deviation

s<sub>R</sub>

1.909

*Shaded cells were removed after outlier analysis*

Binder CR

PG 70-22

**Recovery (3.2 kPa)**

Lab No.	Lab Name	Results, x							
		1	2	3	Average	s	d	h	k
1		48.3	47.4	46.9	47.52333	0.71675	0.22459	0.10	0.87
2		48.3	46.7	47.0	47.35000	0.85422	0.05125	0.02	1.04
3		48.3	48.1	48.4	48.28000	0.14107	0.98125	0.44	0.17
4		48.1	47.8	48.0	47.96333	0.15948	0.66459	0.30	0.19
5		50.1	49.7	50.9	50.24920	0.56284	2.95045	1.32	0.69
6		47.0	48.5	51.1	48.87920	2.08772	1.58045	0.71	2.54
7		45.4	45.7	45.4	45.50333	0.17214	-1.79541	-0.80	0.21
8		44.9	43.7	44.4	44.33667	0.62525	-2.96208	-1.32	0.76
9		48.1	47.5	46.8	47.44137	0.68303	0.14262	0.06	0.83
10		45.0	45.7	46.0	45.55000	0.53694	-1.74875	-0.78	0.65
11		45.1	46.3	46.2	45.86363	0.65148	-1.43511	-0.64	0.79
12		48.4	47.8	46.9	47.71747	0.75712	0.41872	0.19	0.92
13		46.1	46.5	47.1	46.56667	0.49217	-0.73208	-0.33	0.60
14		47.2	48.1	47.7	47.68563	0.46288	0.38689	0.17	0.56
15		50.6	50.1	49.8	50.17567	0.38244	2.87692	1.29	0.47
16		49.7	49.7	50.1	49.85920	0.24149	2.56045	1.14	0.29
17		49.8	49.4	49.3	49.48280	0.26203	2.18405	0.98	0.32
18		48.0	47.4	47.4	47.60833	0.38064	0.30959	0.14	0.46
19		44.3	40.6	40.3	41.70960	2.21848	-5.58915	-2.50	2.70
20		44.3	43.0	42.5	43.26430	0.96403	-4.03445	-1.80	1.17
21		46.6	45.4	45.7	45.88477	0.63985	-1.41398	-0.63	0.78
22		48.6	48.8	49.5	48.94333	0.44287	1.64459	0.74	0.54
23		49.8	50.1	50.2	50.03333	0.23861	2.73459	1.22	0.29
24									
25									
26									
27									
28									

Average of Cell Averages

X-bar                  47.299

Standard Deviation of Cell Averages

s<sub>X-bar</sub>              2.237

Repeatability Standard Deviation

s<sub>r</sub>                  0.82159          2.335

Reproducibility Standard Deviation

s<sub>R</sub>                  2.335

*Shaded cells were removed after outlier analysis*

## Binder CR

PG 70-22

Jnr, Diff

Lab No.	Lab Name	Results, x			Average	s	d	h	k
		1	2	3					
1		0.217	0.226	0.230	0.22417	0.00683	0.01299	0.47	0.42
2		0.271	0.263	0.223	0.25247	0.02563	0.04129	1.50	1.59
3		0.222	0.205	0.202	0.20963	0.01085	-0.00155	-0.06	0.67
4		0.244	0.241	0.244	0.24293	0.00163	0.03175	1.15	0.10
5		0.197	0.207	0.208	0.20434	0.00604	-0.00684	-0.25	0.37
6		0.179	0.148	0.128	0.15183	0.02552	-0.05935	-2.16	1.58
7		0.169	0.181	0.181	0.17683	0.00652	-0.03435	-1.25	0.40
8		0.154	0.149	0.140	0.14773	0.00712	-0.06345	-2.30	0.44
9		0.215	0.230	0.267	0.23722	0.02672	0.02604	0.95	1.66
10		0.216	0.210	0.213	0.21290	0.00310	0.00172	0.06	0.19
11		0.216	0.207	0.181	0.20156	0.01820	-0.00962	-0.35	1.13
12		0.214	0.219	0.229	0.22045	0.00791	0.00927	0.34	0.49
13		0.227	0.297	0.208	0.24383	0.04734	0.03265	1.19	2.93
14		0.231	0.224	0.225	0.22650	0.00394	0.01532	0.56	0.24
15		0.197	0.187	0.191	0.19156	0.00490	-0.01962	-0.71	0.30
16		0.204	0.202	0.205	0.20351	0.00148	-0.00767	-0.28	0.09
17		0.204	0.207	0.207	0.20600	0.00139	-0.00518	-0.19	0.09
18		0.246	0.230	0.229	0.23498	0.00933	0.02381	0.86	0.58
19		0.215	0.232	0.231	0.22624	0.00931	0.01506	0.55	0.58
20		0.208	0.229	0.231	0.22268	0.01282	0.01150	0.42	0.79
21		0.236	0.240	0.236	0.23717	0.00215	0.02599	0.94	0.13
22		0.190	0.215	0.173	0.19250	0.02076	-0.01868	-0.68	1.29
23		0.198	0.173	0.198	0.19008	0.01445	-0.02110	-0.77	0.90
24									
25									
26									
27									
28									

Average of Cell Averages

X-bar                    0.211

Standard Deviation of Cell Averages

S<sub>X-bar</sub>                0.028

Repeatability Standard Deviation

S<sub>r</sub>                    0.01614      0.031

Reproducibility Standard Deviation

S<sub>R</sub>                    0.031*Shaded cells were removed after outlier analysis*

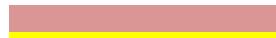
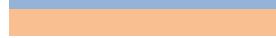
**h**

**Critical Value = 2.59**

2.59      2.59      2.59      2.59      2.59      2.59

**Jnr (0.1 kPa)**

Lab No.	Lab Name	Material					
		Binder AO	Binder BO	Binder CO	Binder AR	Binder BR	Binder CR
1	0	1.79	1.13	1.49	0.70	-0.36	-0.17
2	0	0.59	0.21	0.25	0.86	-0.03	0.39
3	0	0.34	-0.37	0.33	-0.51	-0.26	-0.22
4	0	-0.32	0.02	-0.30	0.17	-0.45	-0.41
5	0	-0.30	0.20	-0.13	-1.84	0.74	-1.64
6	0	-0.17	-1.08	-2.27	-0.67	-0.01	-0.73
7	0	0.93	0.97	0.89	-0.23	1.00	0.85
8	0	-0.79	2.58	1.58	0.37	3.23	1.27
9	0	-0.31	0.19	-0.26	-1.48	-0.71	-0.43
10	0	2.17	0.73	0.76	0.86	0.34	1.04
11	0	-0.49	0.60	0.49	-0.53	0.41	1.05
12	0	-0.55	-1.02	-0.75	-0.76	-0.35	-0.17
13	0	1.14	0.61	0.85	1.94	0.33	0.86
14	0	1.20	-0.51	0.12	1.01	0.52	0.27
15	0	-0.36	-1.14	-0.66	-0.64	-1.88	-2.05
16	0	-0.61	-0.57	-0.36	-0.08	-0.41	-0.81
17	0	1.41	0.72	0.82	0.49	-0.14	-0.67
18	0	-0.99	-0.32	-0.10	0.48	0.01	0.18
19	0	-0.98	-0.10	0.29	1.08	-0.40	1.83
20	0	-1.49	-0.29	0.15	0.89	-0.08	1.13
21	0	-0.46	0.10	0.02	0.83	1.08	0.60
22	0	-0.60	-0.07	-0.52	-1.65	-1.40	-0.82
23	0	-1.15	-2.60	-2.70	-1.28	-1.16	-1.35
24	0	0.00	0.00	0.00	0.00	0.00	0.00
25	0	0.00	0.00	0.00	0.00	0.00	0.00
26	0	0.00	0.00	0.00	0.00	0.00	0.00
27	0	0.00	0.00	0.00	0.00	0.00	0.00
28	0	0.00	0.00	0.00	0.00	0.00	0.00

 exceeds h before removal of outliers  
 exceeds h-0.5 before removal of outliers  
 exceeds h after removal of outliers  
 exceeds h-0.5 after removal of outliers

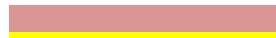
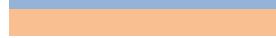
**h**

**Critical Value = 2.59**

2.59      2.59      2.59      2.59      2.59      2.59

**Jnr (3.2 kPa)**

Lab No.	Lab Name	Material					
		Binder AO	Binder BO	Binder CO	Binder AR	Binder BR	Binder CR
1	0	1.76	1.19	1.52	0.73	-0.19	-0.09
2	0	0.70	0.50	0.44	1.03	0.02	0.65
3	0	0.02	-0.28	0.27	-0.63	-0.25	-0.29
4	0	0.18	0.29	0.03	0.57	-0.11	-0.16
5	0	-0.32	0.37	-0.13	-1.74	0.82	-1.56
6	0	-0.15	-1.79	-2.46	-0.64	-1.61	-1.15
7	0	0.47	0.77	0.70	-0.58	0.83	0.39
8	0	-0.49	2.33	0.87	0.33	3.06	1.11
9	0	-0.32	0.39	0.26	-1.45	-0.30	-0.22
10	0	2.56	1.01	1.18	1.11	0.64	1.40
11	0	-0.48	0.17	0.27	-0.20	0.04	0.78
12	0	-0.59	-0.89	-0.52	-0.78	-0.31	-0.12
13	0	1.08	0.55	0.71	1.80	0.61	0.99
14	0	1.17	-0.73	0.15	0.99	0.35	0.31
15	0	-0.38	-0.93	-0.51	-0.70	-1.75	-1.99
16	0	-0.52	-0.46	-0.44	-0.08	-0.30	-0.83
17	0	1.16	0.58	0.81	0.38	0.06	-0.69
18	0	-0.83	-0.16	0.08	0.52	0.09	0.31
19	0	-1.16	-0.43	0.16	0.98	-0.49	1.71
20	0	-1.39	-0.50	-0.02	0.77	-0.29	1.05
21	0	-0.45	0.29	-0.03	0.64	1.25	0.70
22	0	-0.82	0.17	-0.42	-1.36	-0.96	-0.92
23	0	-1.21	-2.44	-2.91	-1.70	-1.23	-1.40
24	0	0.00	0.00	0.00	0.00	0.00	0.00
25	0	0.00	0.00	0.00	0.00	0.00	0.00
26	0	0.00	0.00	0.00	0.00	0.00	0.00
27	0	0.00	0.00	0.00	0.00	0.00	0.00
28	0	0.00	0.00	0.00	0.00	0.00	0.00

 exceeds h before removal of outliers  
 exceeds h-0.5 before removal of outliers  
 exceeds h after removal of outliers  
 exceeds h-0.5 after removal of outliers

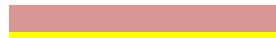
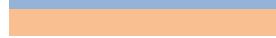
**h**

**Critical Value = 2.59**

2.59      2.59      2.59      2.59      2.59      2.59

**Recovery (0.1 kPa)**

Lab No.	Lab Name	Material					
		Binder AO	Binder BO	Binder CO	Binder AR	Binder BR	Binder CR
1	0	0.33	-0.10	-0.29	0.26	0.43	0.19
2	0	0.56	0.17	0.51	0.42	-0.39	0.40
3	0	0.61	0.76	0.59	0.74	0.59	0.50
4	0	0.92	-0.08	0.63	0.59	0.23	0.27
5	0	0.22	-0.44	-0.08	1.07	-0.67	1.22
6	0	-1.90	-0.77	0.36	-1.09	-1.65	-0.11
7	0	0.78	-1.08	-0.05	0.43	-1.67	-1.02
8	0	1.76	0.00	-2.78	1.03	0.00	-2.27
9	0	-0.05	1.12	0.77	0.69	0.77	0.75
10	0	0.45	0.32	0.33	0.30	0.77	-0.20
11	0	-1.91	-2.28	-1.07	-0.03	0.47	-0.72
12	0	-0.21	0.78	1.34	0.07	0.49	0.26
13	0	0.24	-0.01	-0.45	-0.66	0.63	0.09
14	0	-0.26	1.38	0.69	-0.08	0.10	0.39
15	0	-0.15	1.35	0.91	0.05	1.14	1.02
16	0	-0.17	0.67	0.31	0.01	0.11	1.05
17	0	-0.36	0.03	0.08	-0.21	0.41	0.93
18	0	0.57	1.36	0.75	0.18	0.60	0.49
19	0	0.00	-1.83	-2.01	0.00	-1.56	-2.25
20	0	-2.64	-1.16	-1.56	-3.38	-1.06	-1.80
21	0	0.40	-0.47	-0.17	-0.50	-2.03	-0.28
22	0	0.49	-0.25	0.65	1.38	1.37	0.19
23	0	0.31	0.52	0.55	-1.25	0.92	0.90
24	0	0.00	0.00	0.00	0.00	0.00	0.00
25	0	0.00	0.00	0.00	0.00	0.00	0.00
26	0	0.00	0.00	0.00	0.00	0.00	0.00
27	0	0.00	0.00	0.00	0.00	0.00	0.00
28	0	0.00	0.00	0.00	0.00	0.00	0.00

 exceeds h before removal of outliers  
 exceeds h-0.5 before removal of outliers  
 exceeds h after removal of outliers  
 exceeds h-0.5 after removal of outliers

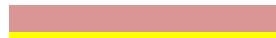
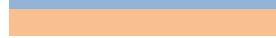
**h**

**Critical Value = 2.59**

2.59      2.59      2.59      2.59      2.59      2.59

**Recovery (3.2 kPa)**

Lab No.	Lab Name	Material					
		Binder AO	Binder BO	Binder CO	Binder AR	Binder BR	Binder CR
1	0	0.57	-0.68	-0.75	0.47	0.14	0.10
2	0	0.59	-0.67	0.21	0.45	-0.27	0.02
3	0	0.84	0.20	0.37	0.79	0.33	0.44
4	0	0.71	-0.37	0.37	0.54	-0.01	0.30
5	0	0.21	-1.02	-0.20	0.53	-0.84	1.32
6	0	-2.06	2.17	1.49	-1.77	2.30	0.71
7	0	0.81	-1.49	-0.31	0.72	-2.10	-0.80
8	0	0.82	0.00	-1.89	0.51	0.00	-1.32
9	0	0.41	0.24	0.08	0.40	-0.22	0.06
10	0	0.65	-0.50	-0.43	0.48	0.28	-0.78
11	0	-2.05	-1.13	0.00	-1.97	0.77	-0.64
12	0	-0.03	0.66	1.04	0.16	0.38	0.19
13	0	0.60	-0.12	-0.34	0.34	-0.09	-0.33
14	0	-0.12	1.93	0.40	-0.02	0.36	0.17
15	0	-0.16	0.88	0.61	0.04	0.91	1.29
16	0	-0.04	0.40	0.49	0.16	-0.12	1.14
17	0	-0.18	-0.06	-0.31	-0.01	-0.04	0.98
18	0	0.01	0.78	0.28	0.03	0.36	0.14
19	0	0.00	-0.80	-2.24	0.00	-1.07	-2.50
20	0	-2.77	-0.19	-1.46	-3.17	-0.41	-1.80
21	0	0.26	-1.17	-0.37	-0.03	-2.33	-0.63
22	0	0.66	-0.56	0.71	0.81	0.47	0.74
23	0	0.29	1.50	2.24	0.56	1.21	1.22
24	0	0.00	0.00	0.00	0.00	0.00	0.00
25	0	0.00	0.00	0.00	0.00	0.00	0.00
26	0	0.00	0.00	0.00	0.00	0.00	0.00
27	0	0.00	0.00	0.00	0.00	0.00	0.00
28	0	0.00	0.00	0.00	0.00	0.00	0.00

 exceeds h before removal of outliers  
 exceeds h-0.5 before removal of outliers  
 exceeds h after removal of outliers  
 exceeds h-0.5 after removal of outliers

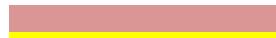
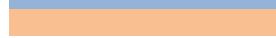
**h**

**Critical Value = 2.59**

2.59      2.59      2.59      2.59      2.59      2.59

**Jnr, Diff**

Lab No.	Lab Name	Material					
		Binder AO	Binder BO	Binder CO	Binder AR	Binder BR	Binder CR
1	0	-0.46	0.93	0.77	0.16	0.23	0.47
2	0	0.31	1.36	0.84	0.79	-0.24	1.50
3	0	-1.48	0.06	0.13	-0.55	-0.52	-0.06
4	0	2.26	1.19	1.10	1.98	1.11	1.15
5	0	-0.85	0.83	0.18	0.71	0.10	-0.25
6	0	0.03	0.00	-1.46	0.20	2.10	-2.16
7	0	-2.23	-0.03	0.01	-1.73	-0.87	-1.25
8	0	0.91	0.02	-2.73	-0.52	-0.40	-2.30
9	0	-0.12	0.92	1.85	0.28	1.44	0.95
10	0	0.71	0.83	-1.30	0.65	0.69	0.06
11	0	0.11	-1.51	-0.25	1.73	-0.05	-0.35
12	0	-0.18	-0.32	0.63	-0.03	-0.42	0.34
13	0	-0.32	0.09	0.11	-0.52	0.94	1.19
14	0	-0.41	-1.57	0.35	-0.13	-1.08	0.56
15	0	-0.11	-0.02	0.40	-0.31	-0.52	-0.71
16	0	0.41	-0.02	-0.15	-0.01	-0.09	-0.28
17	0	-1.13	-0.17	0.45	-0.57	0.41	-0.19
18	0	0.82	0.41	0.70	0.22	-0.06	0.86
19	0	-0.81	-1.71	-0.05	-0.47	-1.88	0.55
20	0	0.62	-1.34	-0.21	-0.56	-1.51	0.42
21	0	0.00	0.84	0.07	-0.89	0.59	0.94
22	0	1.96	1.00	0.33	1.74	1.36	-0.68
23	0	-0.05	-1.78	-1.78	-2.17	-1.34	-0.77
24	0	0.00	0.00	0.00	0.00	0.00	0.00
25	0	0.00	0.00	0.00	0.00	0.00	0.00
26	0	0.00	0.00	0.00	0.00	0.00	0.00
27	0	0.00	0.00	0.00	0.00	0.00	0.00
28	0	0.00	0.00	0.00	0.00	0.00	0.00

 exceeds h before removal of outliers  
 exceeds h-0.5 before removal of outliers  
 exceeds h after removal of outliers  
 exceeds h-0.5 after removal of outliers

k

**Critical Value = 2.22**

2.22      2.22      2.22      2.22      2.22      2.22

**Jnr (0.1 kPa)**

Lab No.	Lab Name	Material					
		Binder AO	Binder BO	Binder CO	Binder AR	Binder BR	Binder CR
1	0	0.18	0.25	0.55	0.10	1.10	0.48
2	0	0.41	0.77	0.32	0.62	0.76	0.67
3	0	0.20	1.29	0.54	0.18	0.43	0.28
4	0	0.05	0.52	0.61	0.21	0.24	0.16
5	0	0.72	0.56	1.33	1.20	1.38	1.10
6	0	1.57	0.57	0.88	2.35	2.30	2.81
7	0	0.53	1.51	0.99	1.85	0.15	0.59
8	0	0.32	0.87	0.46	0.48	0.66	0.86
9	0	0.36	1.59	2.46	0.33	0.72	0.83
10	0	1.10	0.89	0.59	1.26	0.60	0.53
11	0	1.24	0.25	1.88	0.79	3.22	0.46
12	0	0.28	1.26	0.79	0.41	0.39	0.78
13	0	1.33	0.27	0.77	1.32	0.58	0.93
14	0	0.22	0.62	0.55	0.19	0.40	0.55
15	0	0.80	2.29	1.31	0.29	0.07	0.42
16	0	0.05	2.05	0.90	0.14	0.44	0.54
17	0	0.24	0.61	0.78	0.62	0.17	0.63
18	0	0.16	0.32	0.18	0.12	0.17	0.23
19	0	0.48	0.40	1.04	0.98	0.75	2.70
20	0	0.52	0.41	1.01	1.90	0.54	0.65
21	0	0.67	0.40	0.98	0.40	0.46	0.50
22	0	0.64	0.24	0.65	0.13	0.29	0.06
23	0	3.54	0.90	0.33	1.55	0.15	0.41
24	0	0.00	0.00	0.00	0.00	0.00	0.00
25	0	0.00	0.00	0.00	0.00	0.00	0.00
26	0	0.00	0.00	0.00	0.00	0.00	0.00
27	0	0.00	0.00	0.00	0.00	0.00	0.00
28	0	0.00	0.00	0.00	0.00	0.00	0.00

exceeds k before removal of outliers

exceeds k-0.5 before removal of outliers

exceeds k after removal of outliers

exceeds k-0.5 after removal of outliers

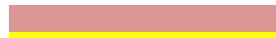
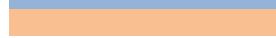
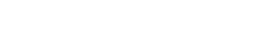
k

**Critical Value = 2.22**

2.22      2.22      2.22      2.22      2.22      2.22

**Jnr (3.2 kPa)**

Lab No.	Lab Name	Material					
		Binder AO	Binder BO	Binder CO	Binder AR	Binder BR	Binder CR
1	0	0.21	0.26	0.56	0.12	1.53	0.54
2	0	0.21	0.70	0.43	0.45	1.04	0.34
3	0	0.24	1.23	0.63	0.14	0.62	0.30
4	0	0.12	0.57	0.56	0.21	0.37	0.12
5	0	0.83	0.51	1.46	1.21	2.10	0.87
6	0	1.64	0.88	0.83	2.70	2.32	2.81
7	0	0.69	1.29	1.00	1.89	0.77	0.55
8	0	0.72	1.06	0.55	0.73	0.95	1.48
9	0	0.37	1.62	0.98	0.34	0.18	1.00
10	0	1.24	0.81	0.88	1.31	0.56	0.54
11	0	0.84	0.06	2.11	0.96	1.73	0.32
12	0	0.18	1.15	1.08	0.32	0.59	0.84
13	0	0.28	0.83	0.57	0.09	1.01	0.73
14	0	0.21	0.66	0.62	0.20	0.52	0.57
15	0	0.77	2.22	1.62	0.34	0.14	0.33
16	0	0.05	1.87	0.90	0.15	0.64	0.45
17	0	0.24	0.80	0.89	0.71	0.79	0.54
18	0	0.23	0.34	0.34	0.03	0.17	0.03
19	0	0.53	0.32	1.20	1.03	0.50	2.59
20	0	0.76	0.42	1.04	2.07	0.78	0.78
21	0	0.71	0.36	1.11	0.30	0.75	0.46
22	0	1.42	0.40	0.73	0.41	0.49	0.41
23	0	3.44	1.15	0.98	0.84	0.22	0.23
24	0	0.00	0.00	0.00	0.00	0.00	0.00
25	0	0.00	0.00	0.00	0.00	0.00	0.00
26	0	0.00	0.00	0.00	0.00	0.00	0.00
27	0	0.00	0.00	0.00	0.00	0.00	0.00
28	0	0.00	0.00	0.00	0.00	0.00	0.00

 exceeds k before removal of outliers  
 exceeds k-0.5 before removal of outliers  
 exceeds k after removal of outliers  
 exceeds k-0.5 after removal of outliers

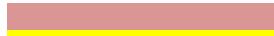
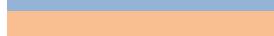
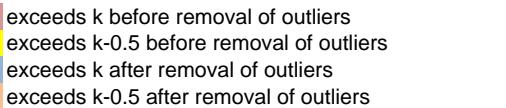
k

**Critical Value = 2.22**

2.22      2.22      2.22      2.22      2.22      2.22

**Recovery (0.1 kPa)**

Lab No.	Lab Name	Material					
		Binder AO	Binder BO	Binder CO	Binder AR	Binder BR	Binder CR
1	0	0.13	0.13	0.11	0.03	0.39	0.55
2	0	0.46	0.94	1.82	0.91	0.25	1.84
3	0	0.05	0.77	0.51	0.15	0.24	0.16
4	0	0.56	0.40	0.64	0.45	0.10	0.35
5	0	0.39	0.20	0.68	0.59	0.37	0.89
6	0	0.29	0.49	0.71	0.27	1.23	1.72
7	0	0.47	0.97	2.11	0.64	0.23	0.47
8	0	0.12	0.00	0.09	0.16	0.00	0.73
9	0	0.23	0.90	1.42	0.14	1.44	1.19
10	0	0.33	0.77	0.64	0.32	0.20	0.73
11	0	1.50	0.68	0.93	0.76	4.06	0.74
12	0	0.35	1.22	1.15	0.35	0.18	0.56
13	0	4.09	1.20	0.82	3.42	0.21	2.03
14	0	0.06	0.58	0.30	0.29	0.17	0.40
15	0	0.18	1.53	1.51	0.06	0.04	0.33
16	0	0.09	0.94	0.67	0.07	0.03	0.34
17	0	0.03	1.21	0.40	0.05	0.29	0.31
18	0	0.24	0.64	0.49	0.21	0.24	0.62
19	0	0.00	0.38	0.28	0.00	0.57	2.30
20	0	0.52	0.72	0.87	0.27	0.63	0.51
21	0	0.14	0.50	0.62	0.44	0.36	0.56
22	0	0.60	0.26	0.81	0.87	0.27	0.71
23	0	0.99	2.81	1.78	2.54	0.48	0.70
24	0	0.00	0.00	0.00	0.00	0.00	0.00
25	0	0.00	0.00	0.00	0.00	0.00	0.00
26	0	0.00	0.00	0.00	0.00	0.00	0.00
27	0	0.00	0.00	0.00	0.00	0.00	0.00
28	0	0.00	0.00	0.00	0.00	0.00	0.00

 exceeds k before removal of outliers  
 exceeds k-0.5 before removal of outliers  
 exceeds k after removal of outliers  
 exceeds k-0.5 after removal of outliers

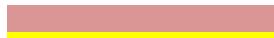
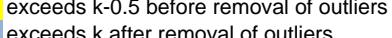
k

**Critical Value = 2.22**

2.22      2.22      2.22      2.22      2.22      2.22

**Recovery (3.2 kPa)**

Lab No.	Lab Name	Material					
		Binder AO	Binder BO	Binder CO	Binder AR	Binder BR	Binder CR
1	0	0.01	0.15	0.14	0.20	0.90	0.87
2	0	0.14	0.58	0.14	0.39	0.60	1.04
3	0	0.03	1.01	0.39	0.25	0.53	0.17
4	0	0.03	0.32	0.49	0.10	0.27	0.19
5	0	1.49	0.21	0.37	1.98	1.15	0.69
6	0	0.60	0.84	0.87	2.78	1.65	2.54
7	0	0.08	1.36	1.76	1.98	0.14	0.21
8	0	0.06	0.00	1.91	0.41	0.00	0.76
9	0	3.04	0.58	0.31	0.20	0.42	0.83
10	0	0.12	0.89	0.51	0.79	0.43	0.65
11	0	0.29	0.19	2.44	0.54	2.98	0.79
12	0	0.06	1.43	0.50	0.37	0.56	0.92
13	0	0.03	0.77	0.26	0.20	0.09	0.60
14	0	0.05	0.94	0.64	0.09	0.32	0.56
15	0	0.24	2.12	1.21	0.23	0.18	0.47
16	0	0.02	1.08	0.97	0.03	0.21	0.29
17	0	0.05	1.97	0.73	0.42	1.56	0.32
18	0	0.06	0.86	0.22	0.29	0.50	0.46
19	0	0.00	0.71	0.52	0.00	0.81	2.70
20	0	0.12	0.99	1.24	0.63	1.55	1.17
21	0	1.69	0.59	1.02	0.43	0.82	0.78
22	0	0.07	0.67	0.86	0.15	0.48	0.54
23	0	2.67	0.82	1.14	1.98	0.26	0.29
24	0	0.00	0.00	0.00	0.00	0.00	0.00
25	0	0.00	0.00	0.00	0.00	0.00	0.00
26	0	0.00	0.00	0.00	0.00	0.00	0.00
27	0	0.00	0.00	0.00	0.00	0.00	0.00
28	0	0.00	0.00	0.00	0.00	0.00	0.00

 exceeds k before removal of outliers  
 exceeds k-0.5 before removal of outliers  
 exceeds k after removal of outliers  
 exceeds k-0.5 after removal of outliers

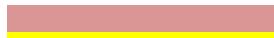
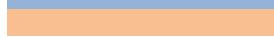
k

**Critical Value = 2.22**

2.22      2.22      2.22      2.22      2.22      2.22

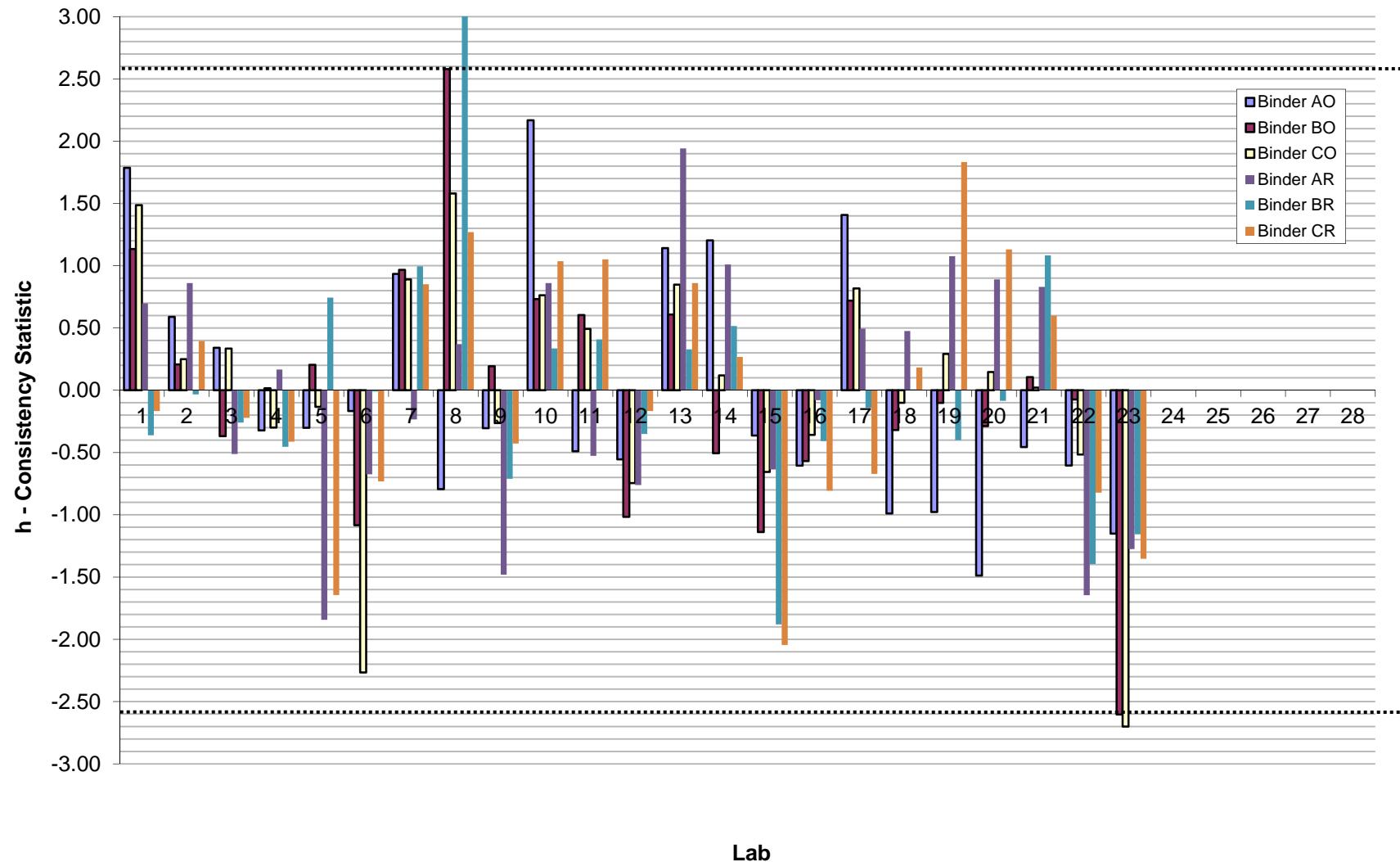
**Jnr, Diff**

Lab No.	Lab Name	Material					
		Binder AO	Binder BO	Binder CO	Binder AR	Binder BR	Binder CR
1	0	0.22	0.18	0.05	0.03	0.16	0.42
2	0	0.61	1.06	1.42	0.97	0.20	1.59
3	0	0.36	0.32	0.34	0.15	0.11	0.67
4	0	0.48	0.41	0.18	0.47	0.09	0.10
5	0	0.84	0.57	0.41	0.45	0.48	0.37
6	0	0.17	0.00	0.22	0.33	0.62	1.58
7	0	0.71	1.07	0.97	0.32	1.04	0.40
8	0	2.07	0.75	0.50	0.45	0.12	0.44
9	0	0.24	0.72	3.41	0.16	2.28	1.66
10	0	0.09	0.27	0.27	0.32	0.63	0.19
11	0	1.39	0.99	0.68	0.99	3.37	1.13
12	0	0.36	0.40	1.01	0.37	0.18	0.49
13	0	3.28	2.56	0.58	3.62	0.44	2.93
14	0	0.12	0.60	0.18	0.30	0.02	0.24
15	0	0.35	0.93	1.20	0.06	0.13	0.30
16	0	0.21	0.24	0.11	0.14	0.16	0.09
17	0	0.22	1.07	0.27	0.11	1.30	0.09
18	0	0.37	0.36	0.57	0.33	0.08	0.58
19	0	0.30	0.42	0.40	0.16	0.15	0.58
20	0	0.61	0.36	0.21	0.19	1.22	0.79
21	0	0.02	0.21	0.42	0.34	0.25	0.13
22	0	1.39	1.60	0.59	0.92	0.27	1.29
23	0	1.03	2.19	1.82	2.37	0.82	0.90
24	0	0.00	0.00	0.00	0.00	0.00	0.00
25	0	0.00	0.00	0.00	0.00	0.00	0.00
26	0	0.00	0.00	0.00	0.00	0.00	0.00
27	0	0.00	0.00	0.00	0.00	0.00	0.00
28	0	0.00	0.00	0.00	0.00	0.00	0.00

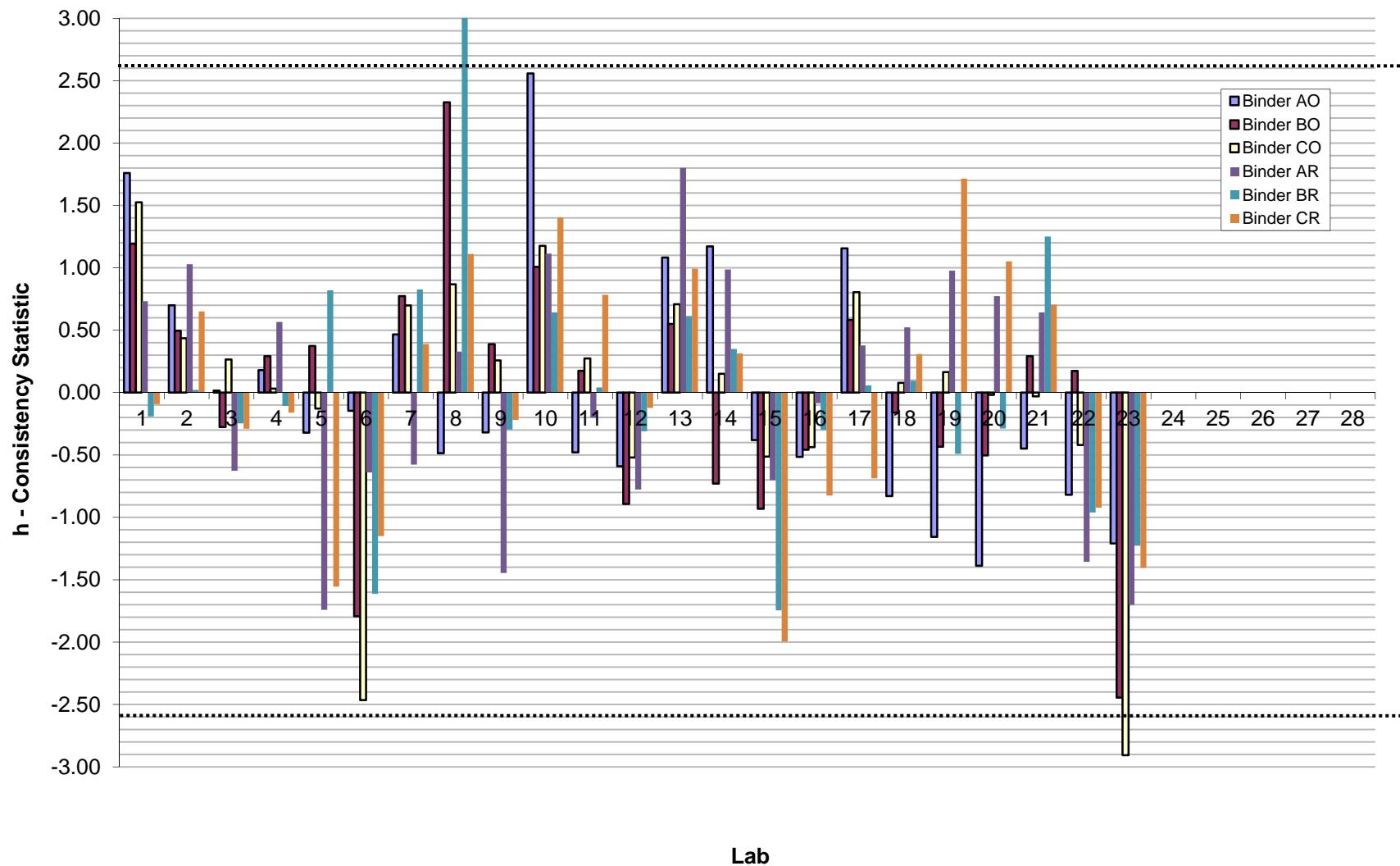
 exceeds k before removal of outliers  
 exceeds k-0.5 before removal of outliers  
 exceeds k after removal of outliers  
 exceeds k-0.5 after removal of outliers



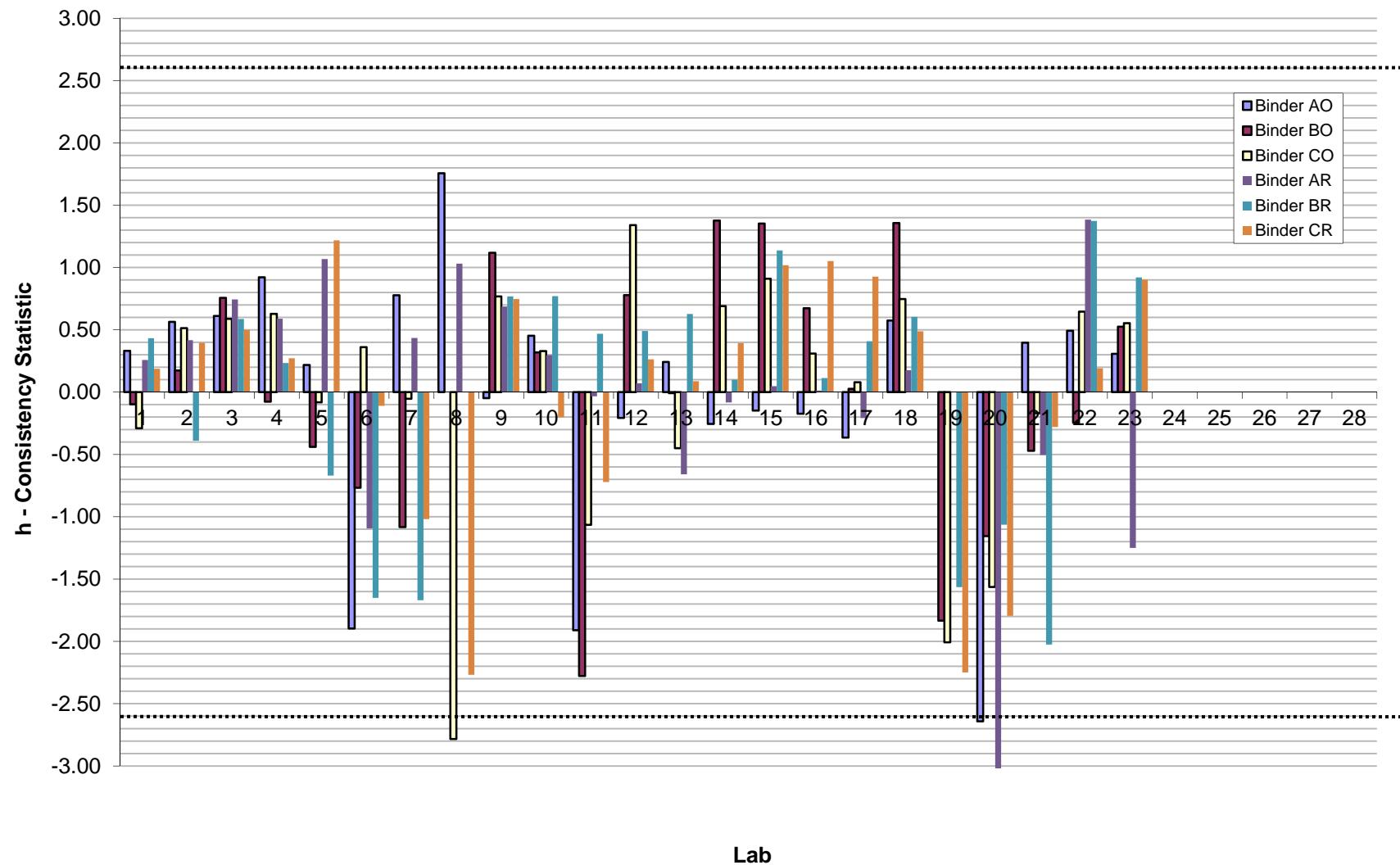
### Jnr (0.1 kPa): h - Materials within Laboratories



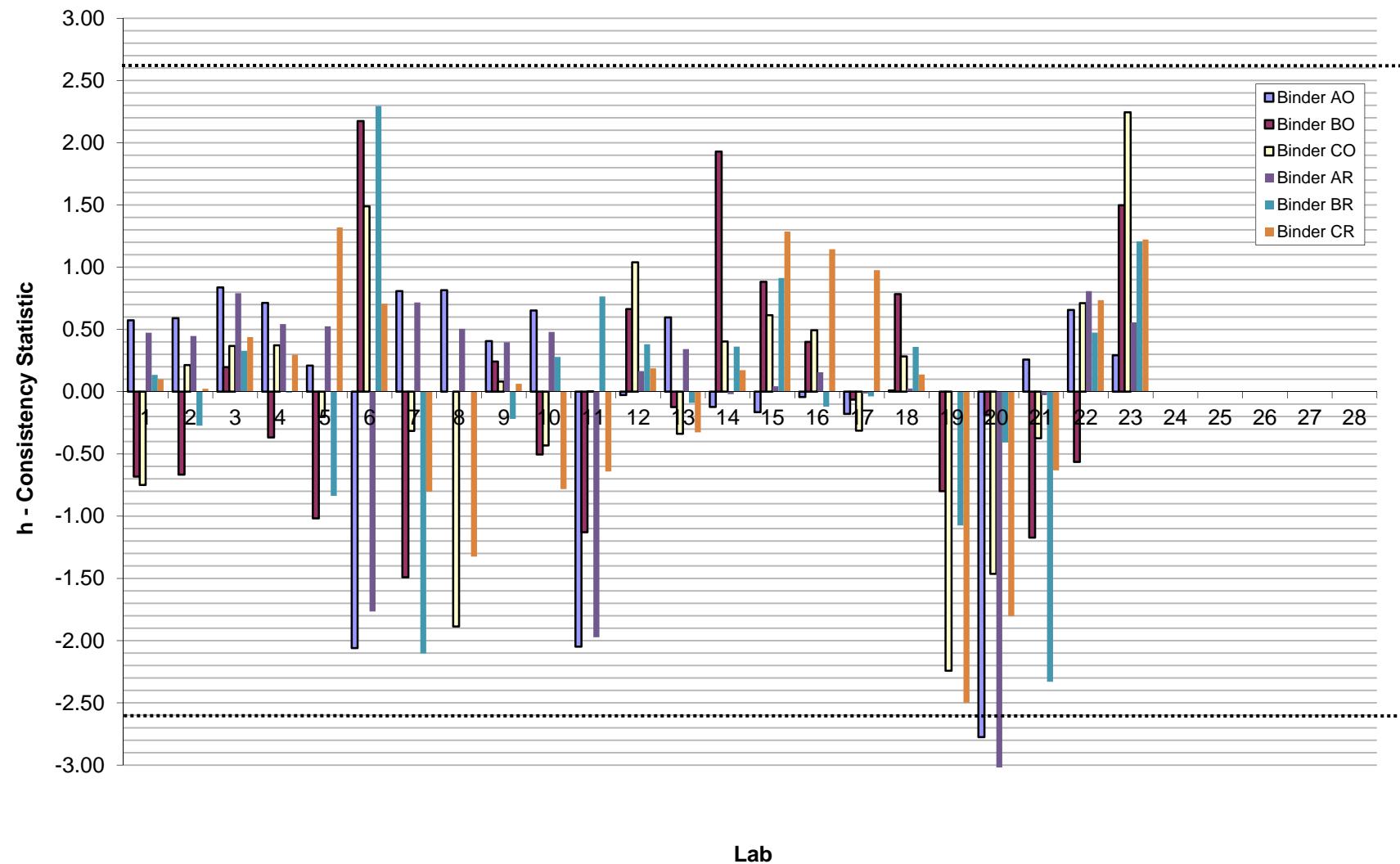
### Jnr (3.2 kPa): h - Materials within Laboratories



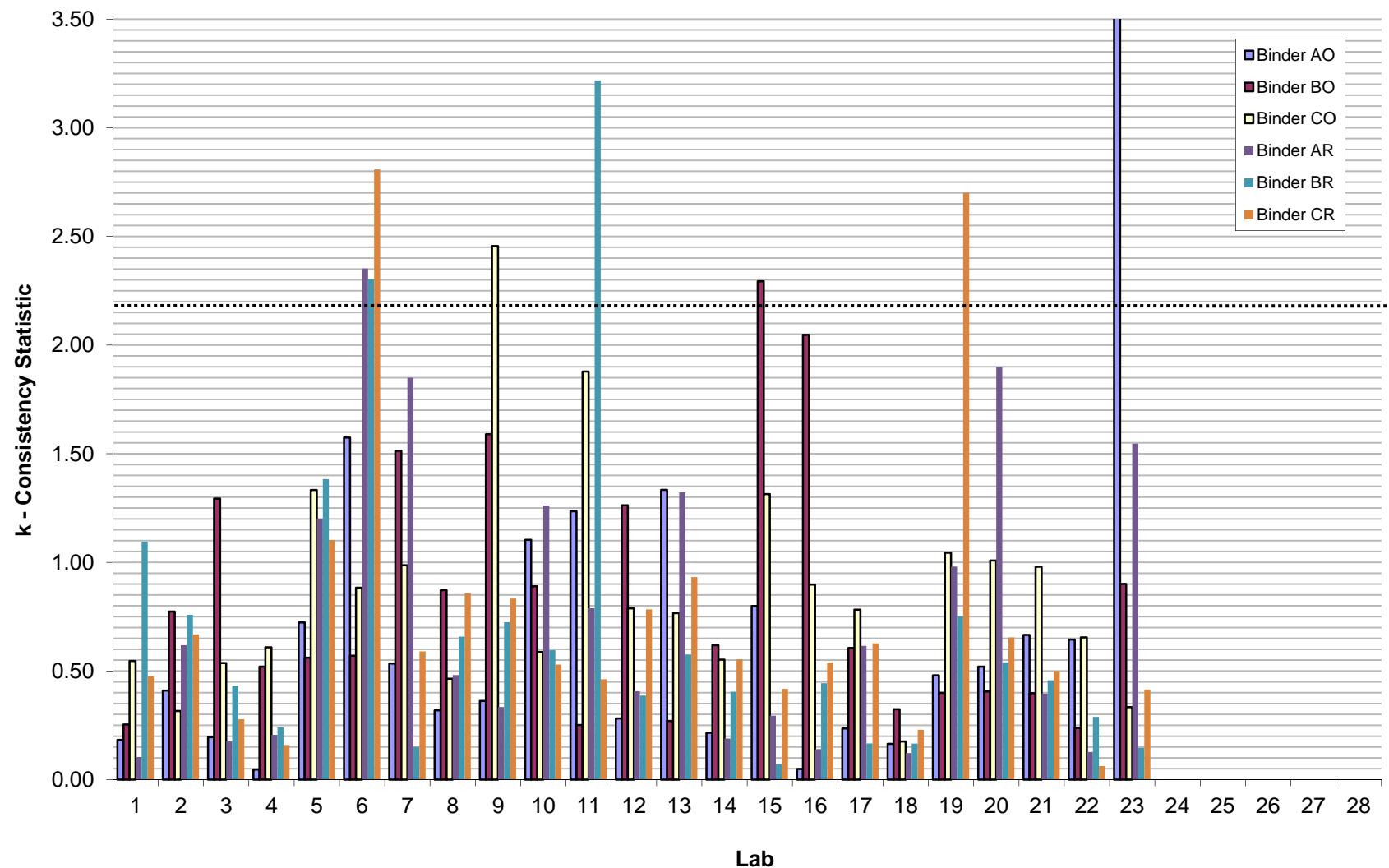
### Recovery (0.1 kPa): h - Materials within Laboratories



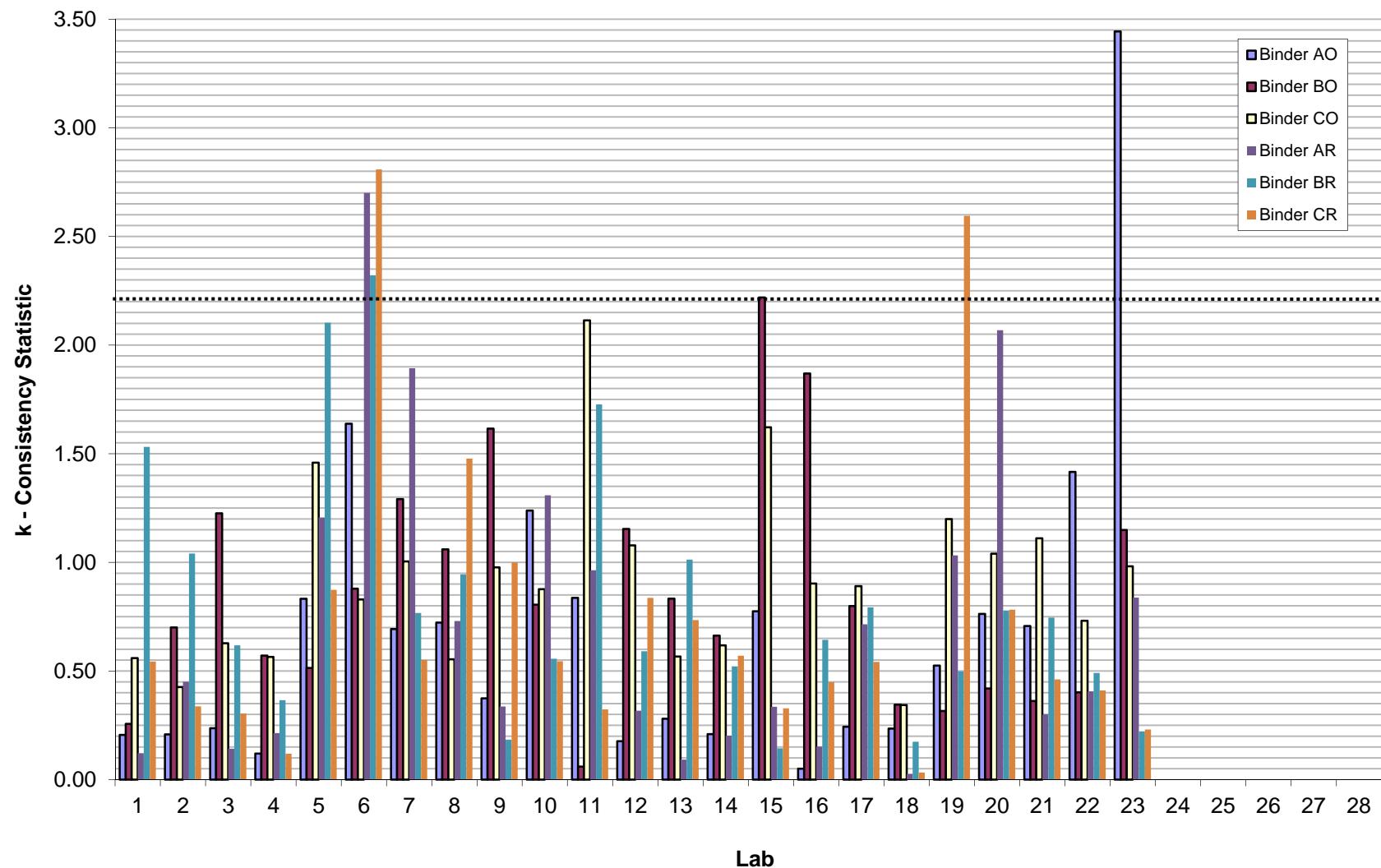
### Recovery (3.2 kPa): h - Materials within Laboratories



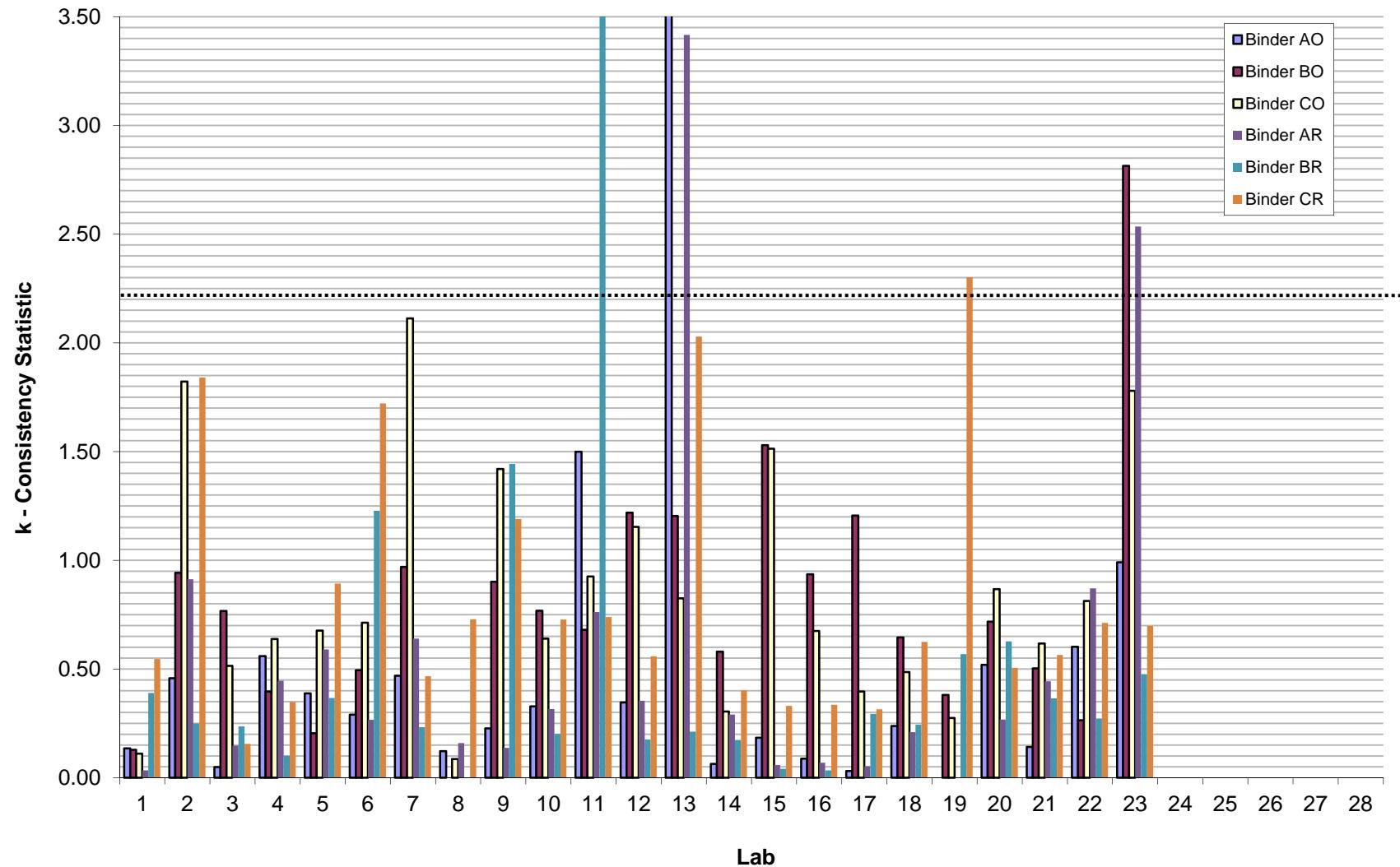
### Jnr (0.1 kPa): k - Materials within Laboratories



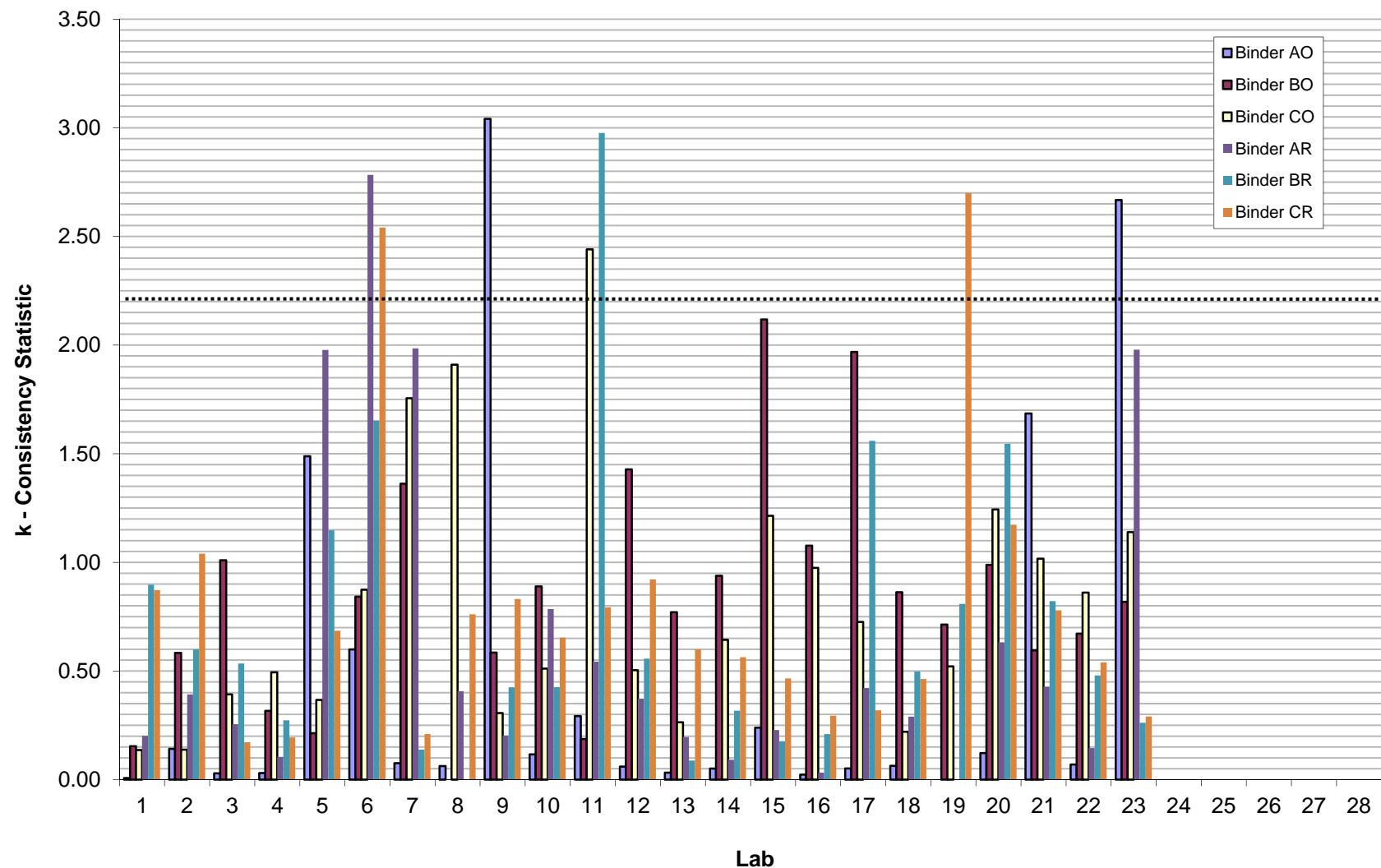
### Jnr (3.2 kPa): k - Materials within Laboratories



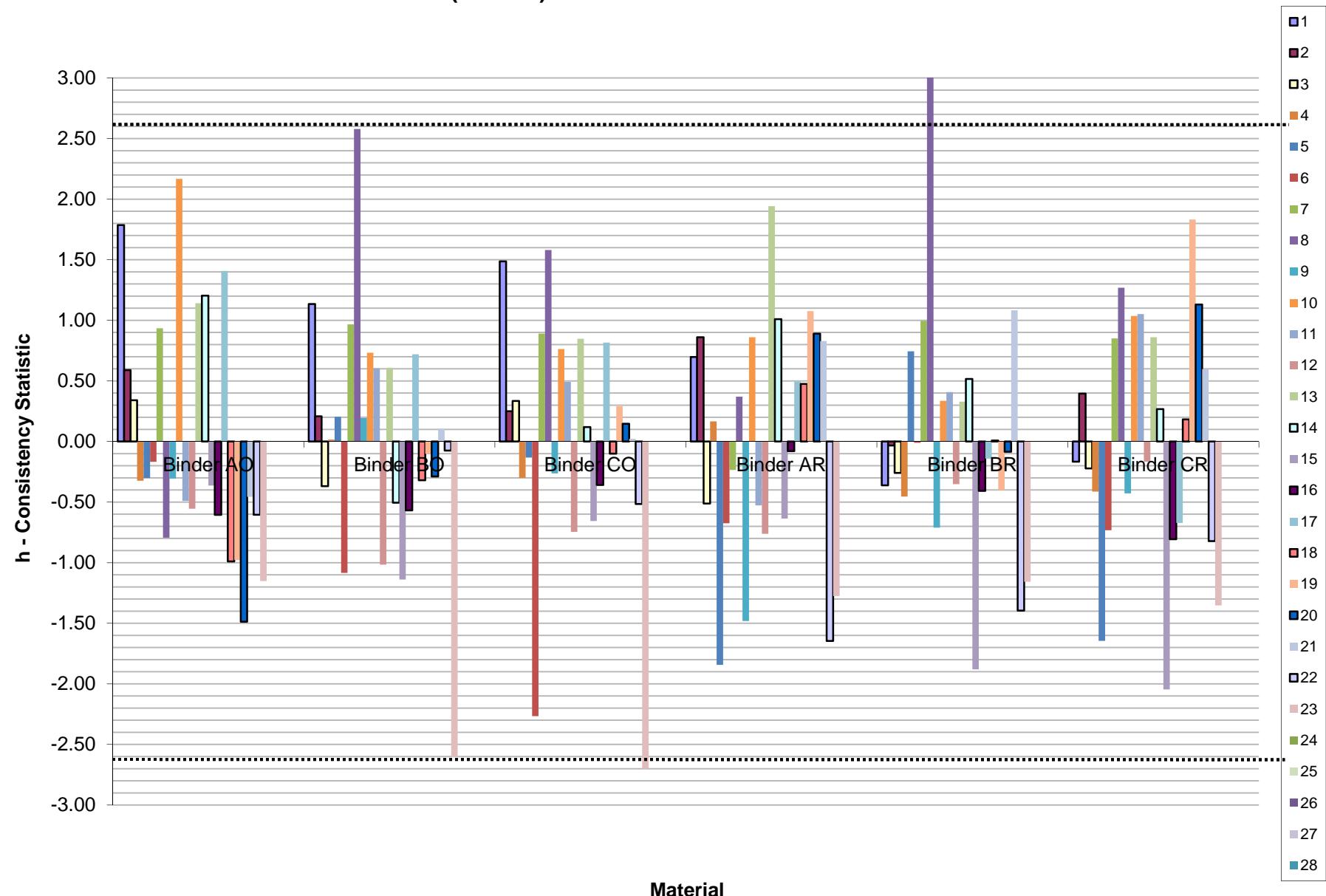
### Recovery (0.1 kPa): k - Materials within Laboratories



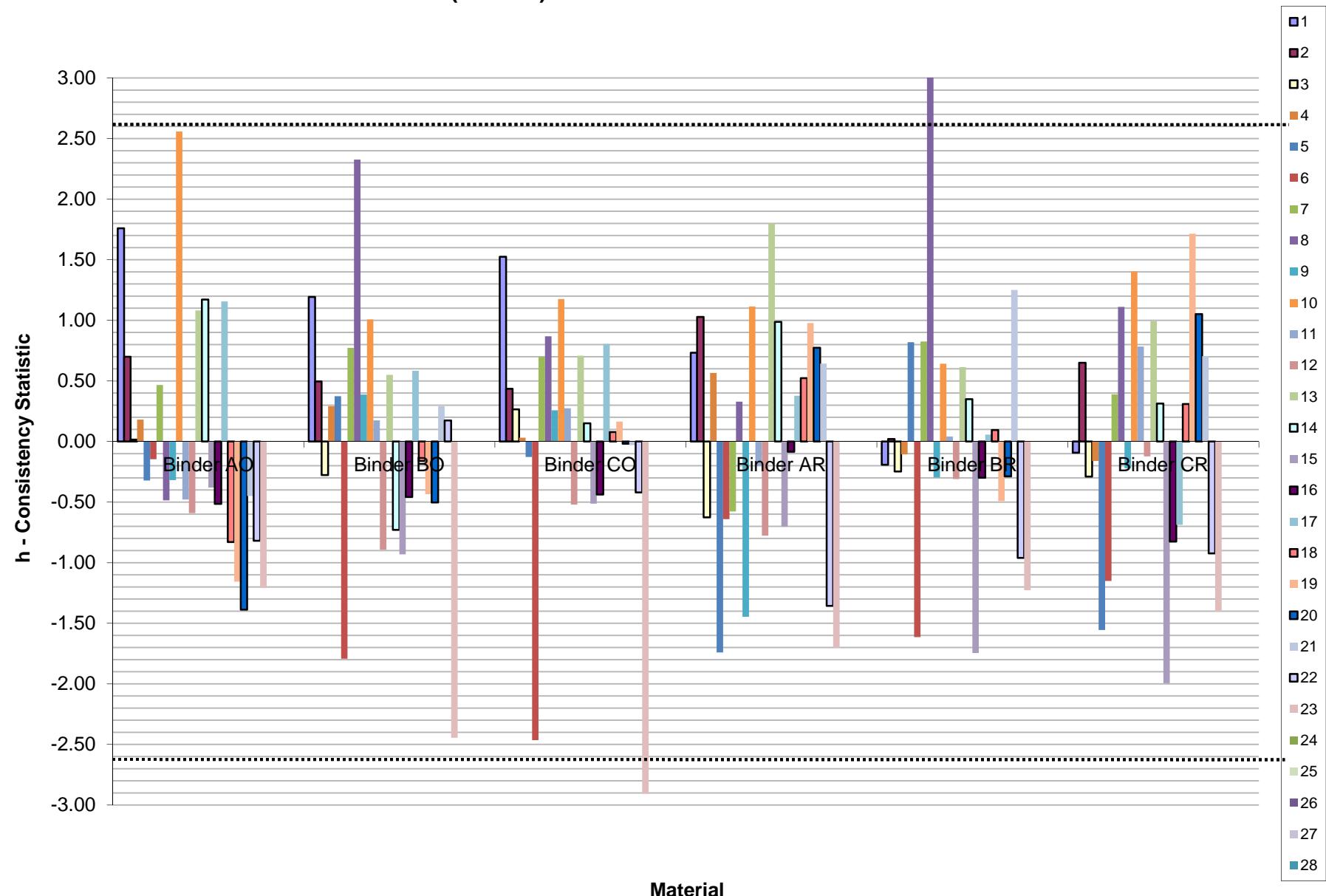
### Recovery (3.2 kPa): k - Materials within Laboratories



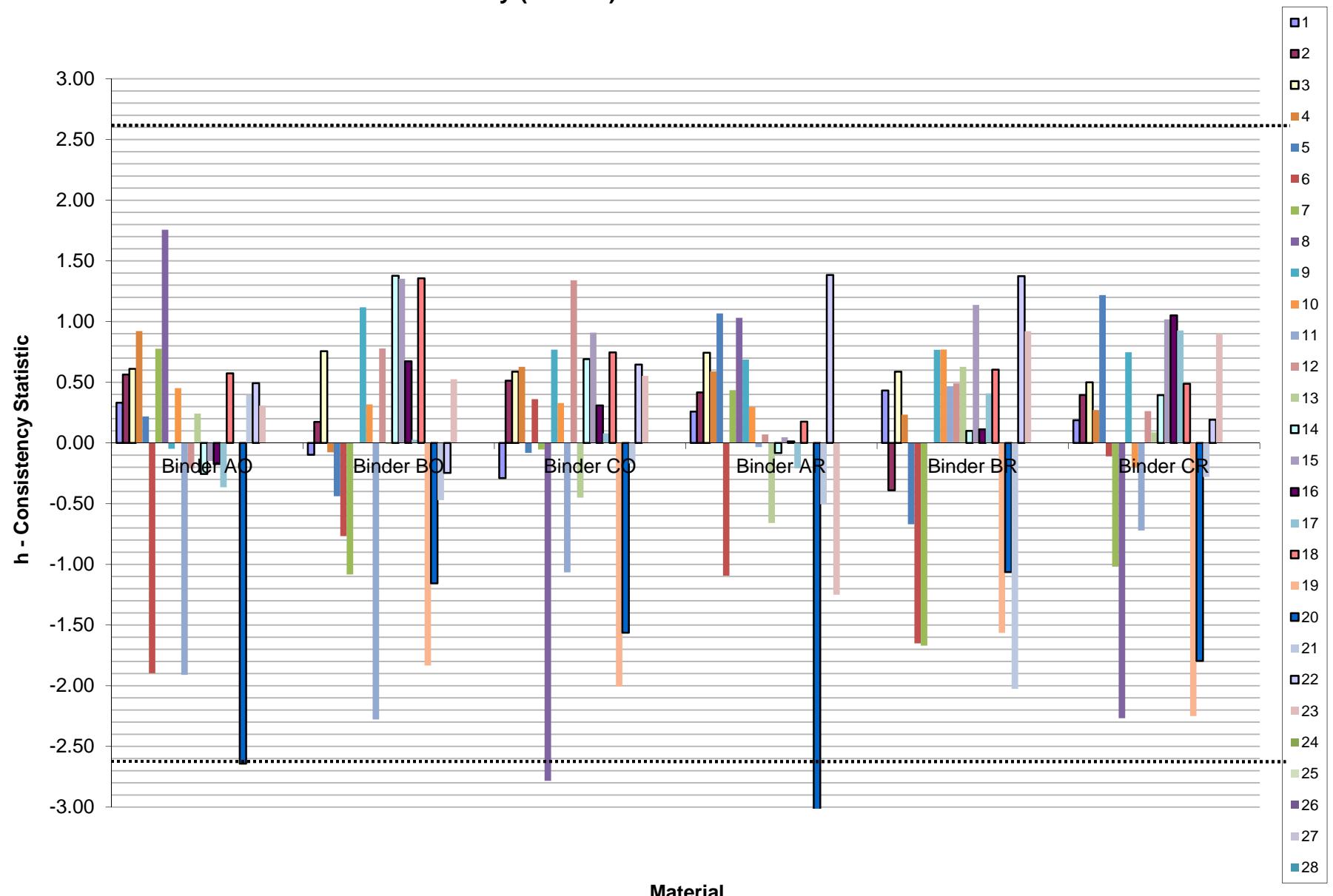
Jnr (0.1 kPa): h - Laboratories within Materials



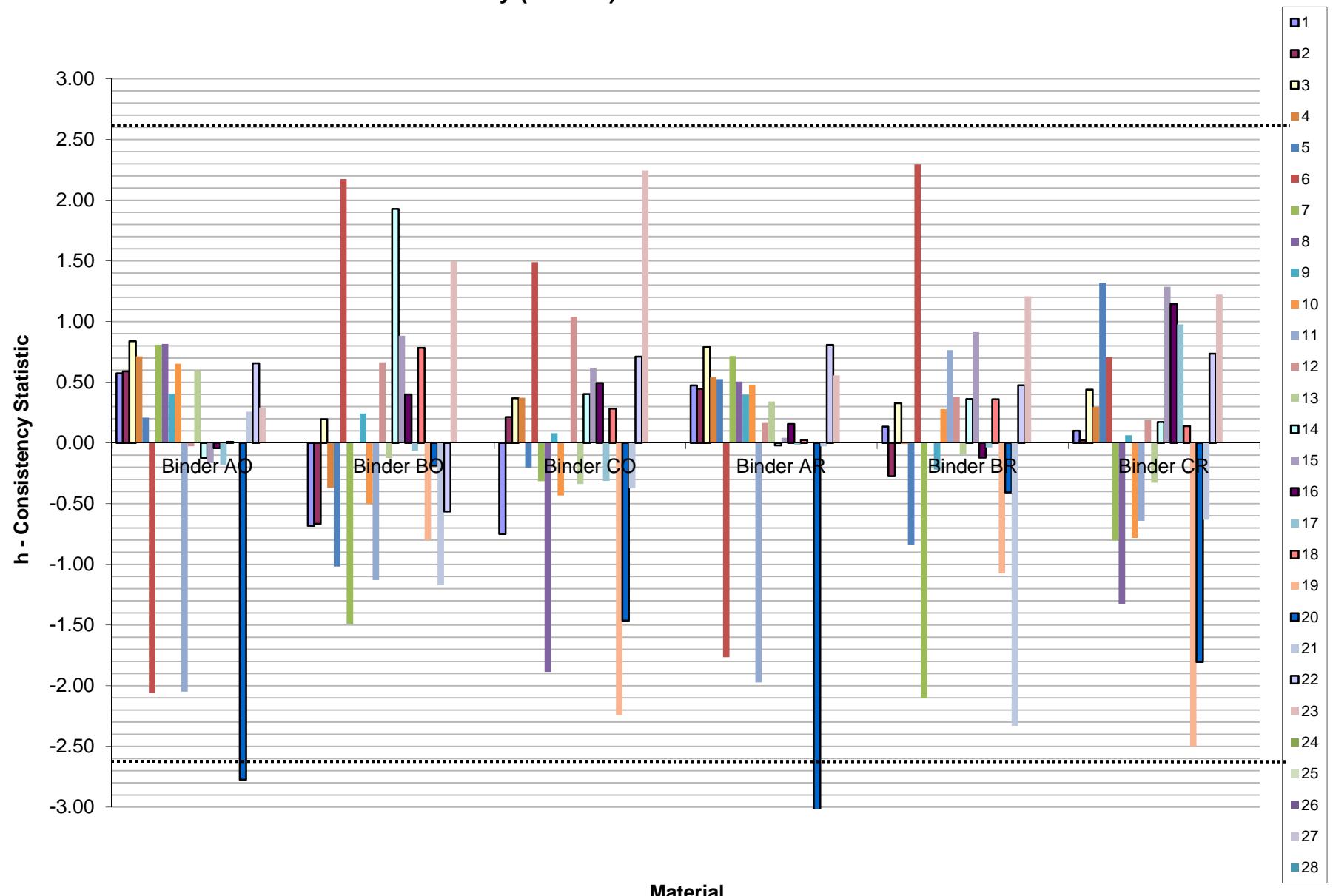
### **Jnr (3.2 kPa): h - Laboratories within Materials**



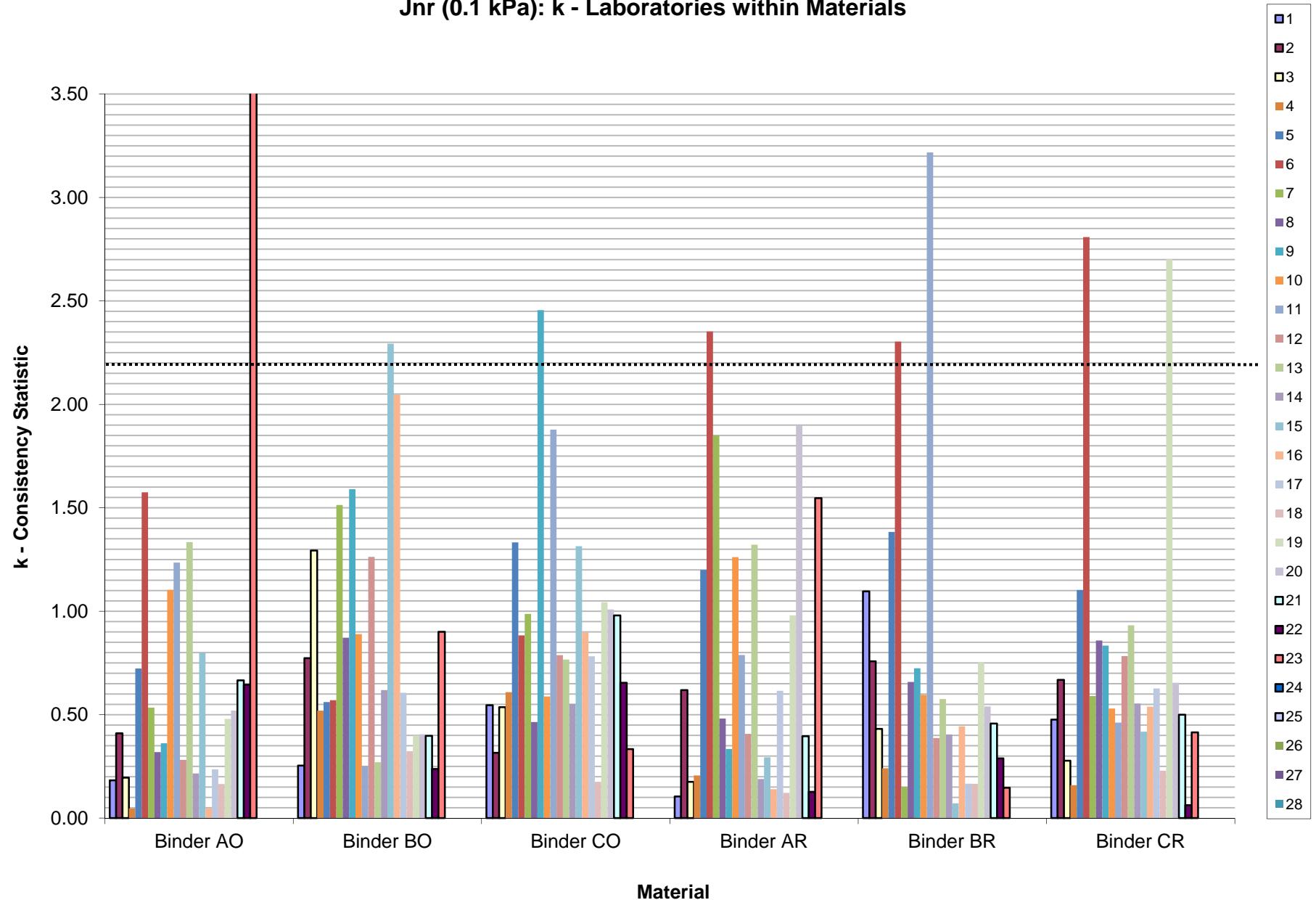
### Recovery (0.1 kPa): h - Laboratories within Materials



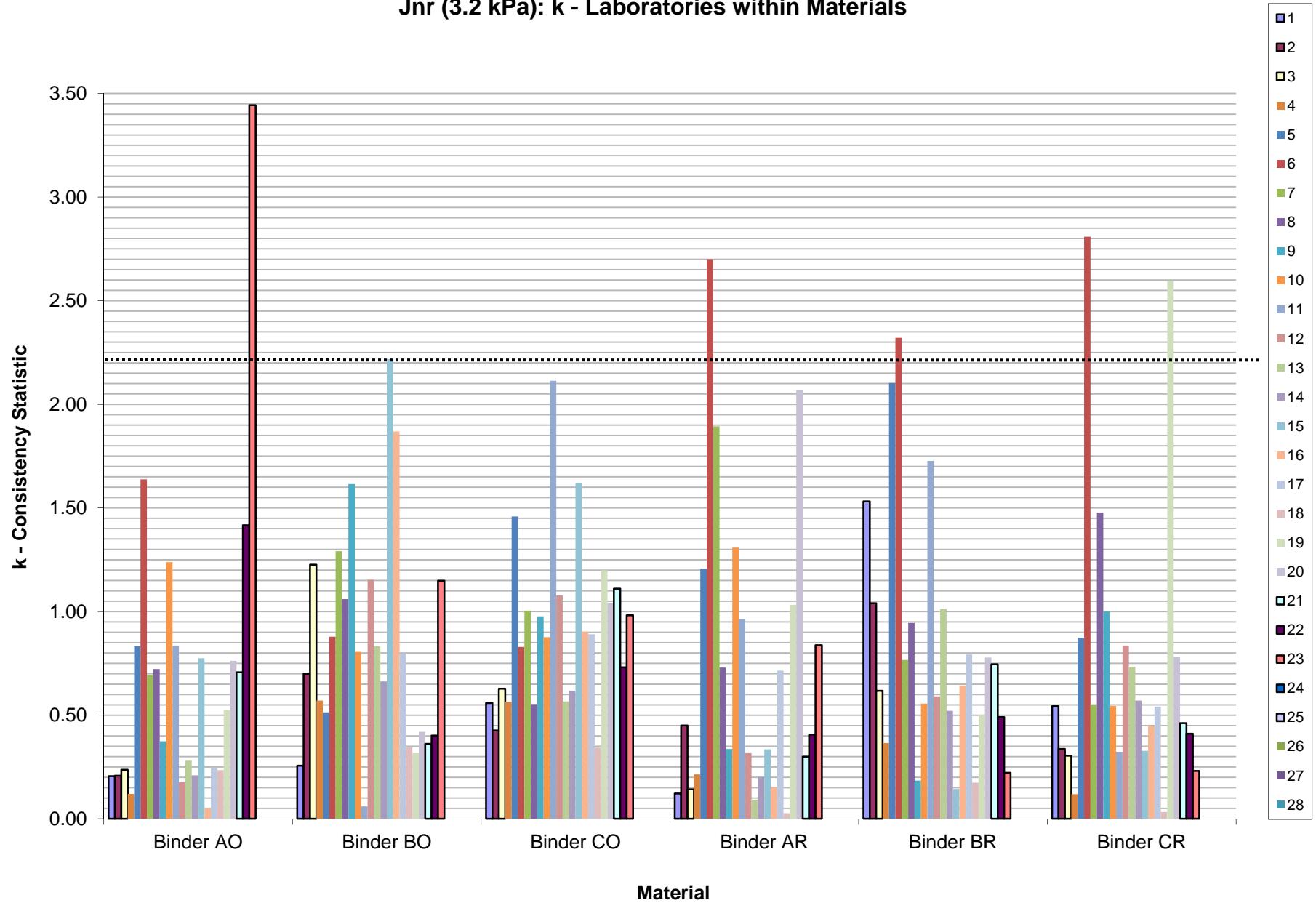
### Recovery (3.2 kPa): h - Laboratories within Materials



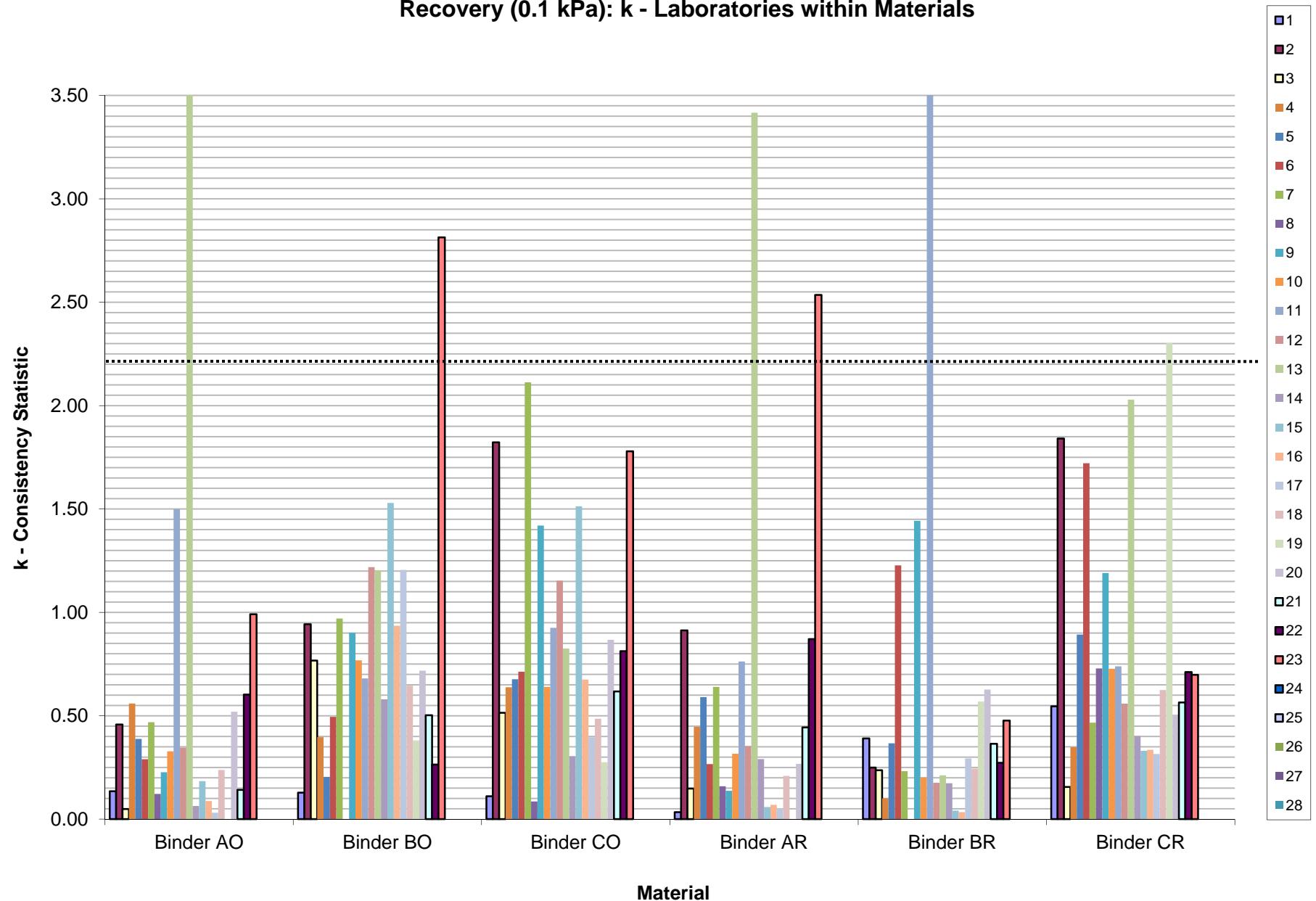
## Jnr (0.1 kPa): k - Laboratories within Materials



### Jnr (3.2 kPa): k - Laboratories within Materials



## **Recovery (0.1 kPa): k - Laboratories within Materials**



### Recovery (3.2 kPa): k - Laboratories within Materials

