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SORT CASES  BY HasValidConcl
SPLIT FILE SEPARATE BY HasValidConcl
DATASET ACTIVATE DataSet1.
CLUSTER  Acc204
/METHOD WARD
/MEASURE=SEUCLID
/ID=syllogism
/PRINT SCHEDULE
/PLOT DENDROGRAM VICICLE.

```

## Cluster

### HasValidConcl = N

**Case Processing Summary<sup>a,b,c</sup>**

Cases					
Valid		Missing		Total	
N	Percent	N	Percent	N	Percent
37	100.0	0	.0	37	100.0

a. HasValidConcl = N

b. Squared Euclidean Distance used

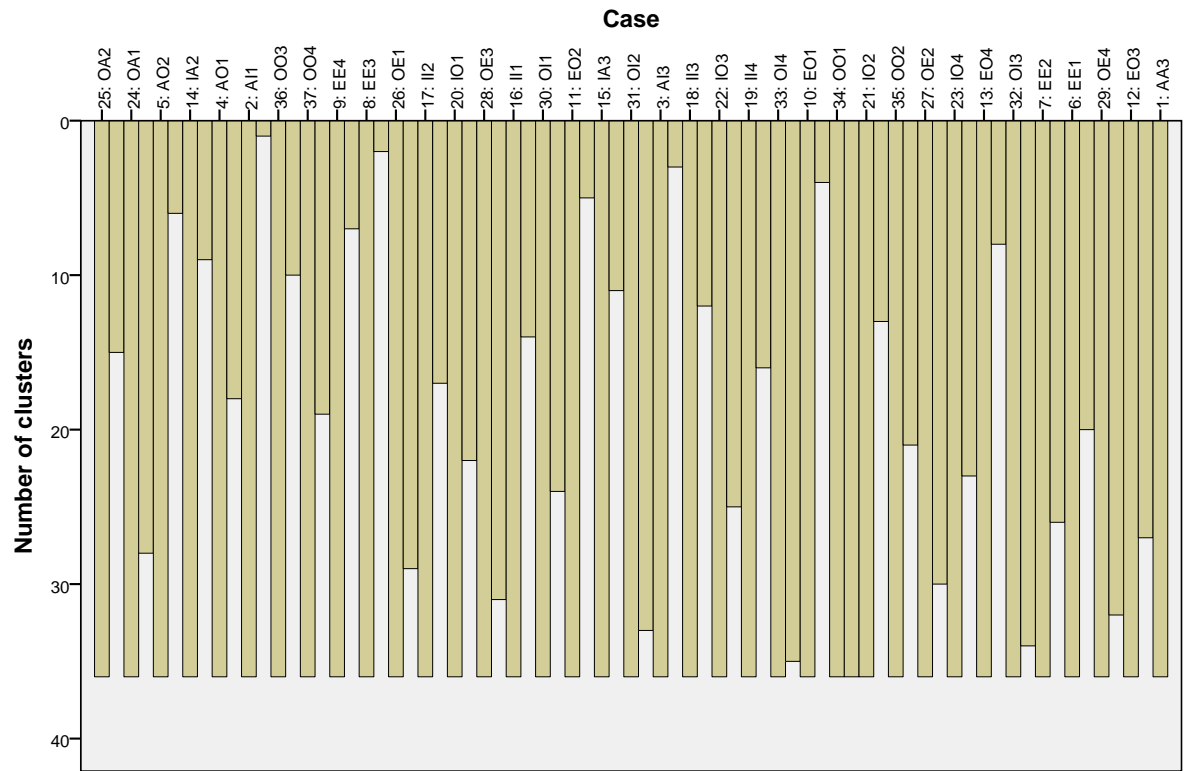
c. Ward Linkage

## Ward Linkage

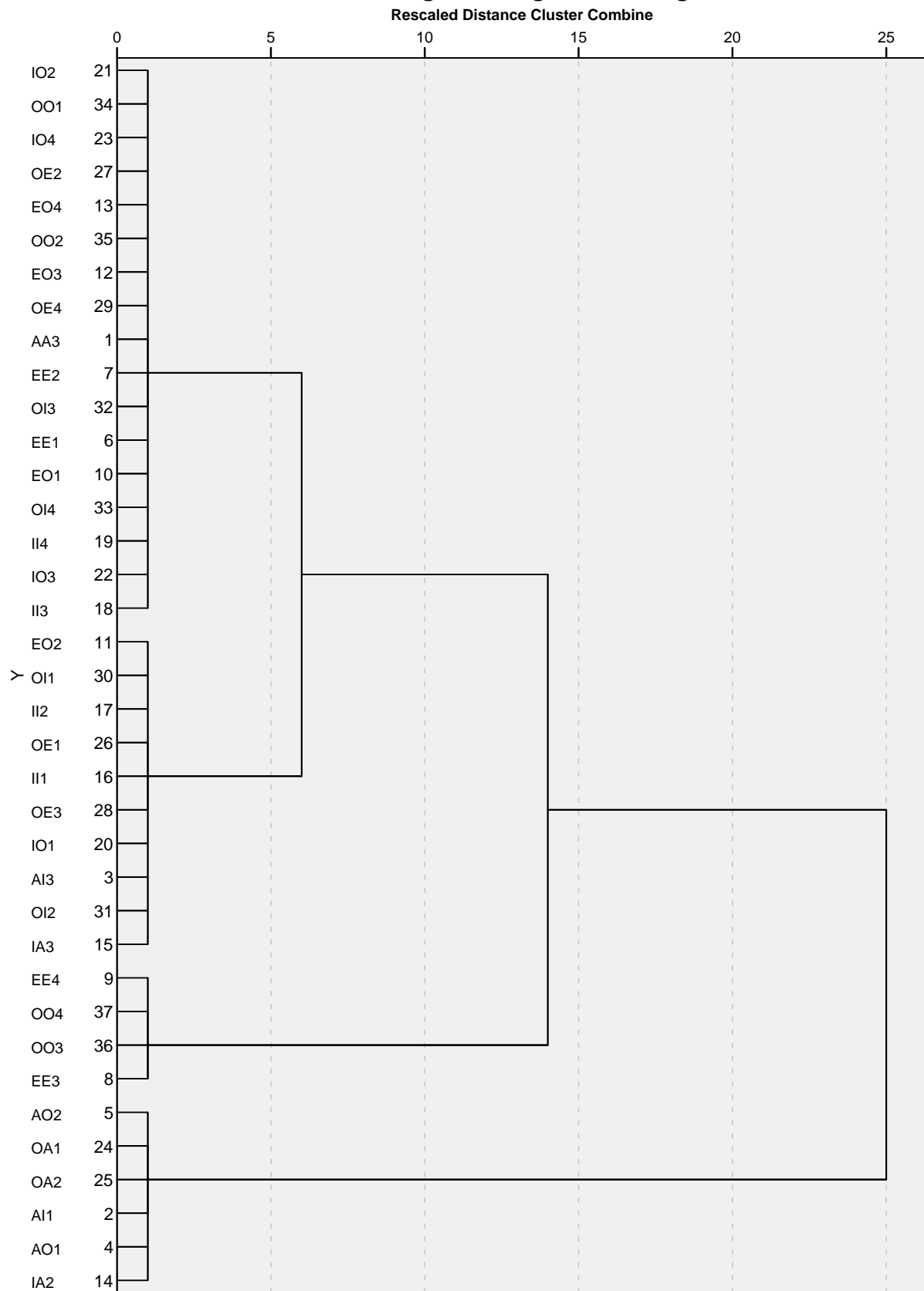
**Agglomeration Schedule<sup>a</sup>**

Stage	Cluster Combined		Coefficients	Stage Cluster First Appears		Next Stage
	Cluster 1	Cluster 2		Cluster 1	Cluster 2	
1	21	34	.000	0	0	24
2	10	33	.000	0	0	21
3	7	32	.000	0	0	11
4	3	31	.000	0	0	26
5	12	29	.000	0	0	10
6	16	28	.000	0	0	15
7	23	27	.000	0	0	14
8	17	26	.000	0	0	20
9	5	24	.000	0	0	22
10	1	12	.000	0	5	17
11	6	7	.000	0	3	17
12	19	22	.000	0	0	21
13	11	30	.000	0	0	23
14	13	23	.000	0	7	16
15	16	20	.000	6	0	20
16	13	35	.000	14	0	24
17	1	6	.000	10	11	29
18	9	37	.000	0	0	27
19	2	4	.000	0	0	28
20	16	17	.000	15	8	23
21	10	19	.000	2	12	25
22	5	25	.001	9	0	31
23	11	16	.001	13	20	32
24	13	21	.002	16	1	29
25	10	18	.002	21	0	33
26	3	15	.003	4	0	32
27	9	36	.003	18	0	30
28	2	14	.004	19	0	31
29	1	13	.006	17	24	33
30	8	9	.007	0	27	35
31	2	5	.011	28	22	36
32	3	11	.015	26	23	34
33	1	10	.021	29	25	34
34	1	3	.067	33	32	35
35	1	8	.172	34	30	36
36	1	2	.364	35	31	0

a. HasValidConcl = N



## Dendrogram using Ward Linkage



HasValidConcl = Y

### Case Processing Summary<sup>a,b,c</sup>

Cases					
Valid		Missing		Total	
N	Percent	N	Percent	N	Percent
27	100.0	0	.0	27	100.0

a. HasValidConcl = Y

b. Squared Euclidean Distance used

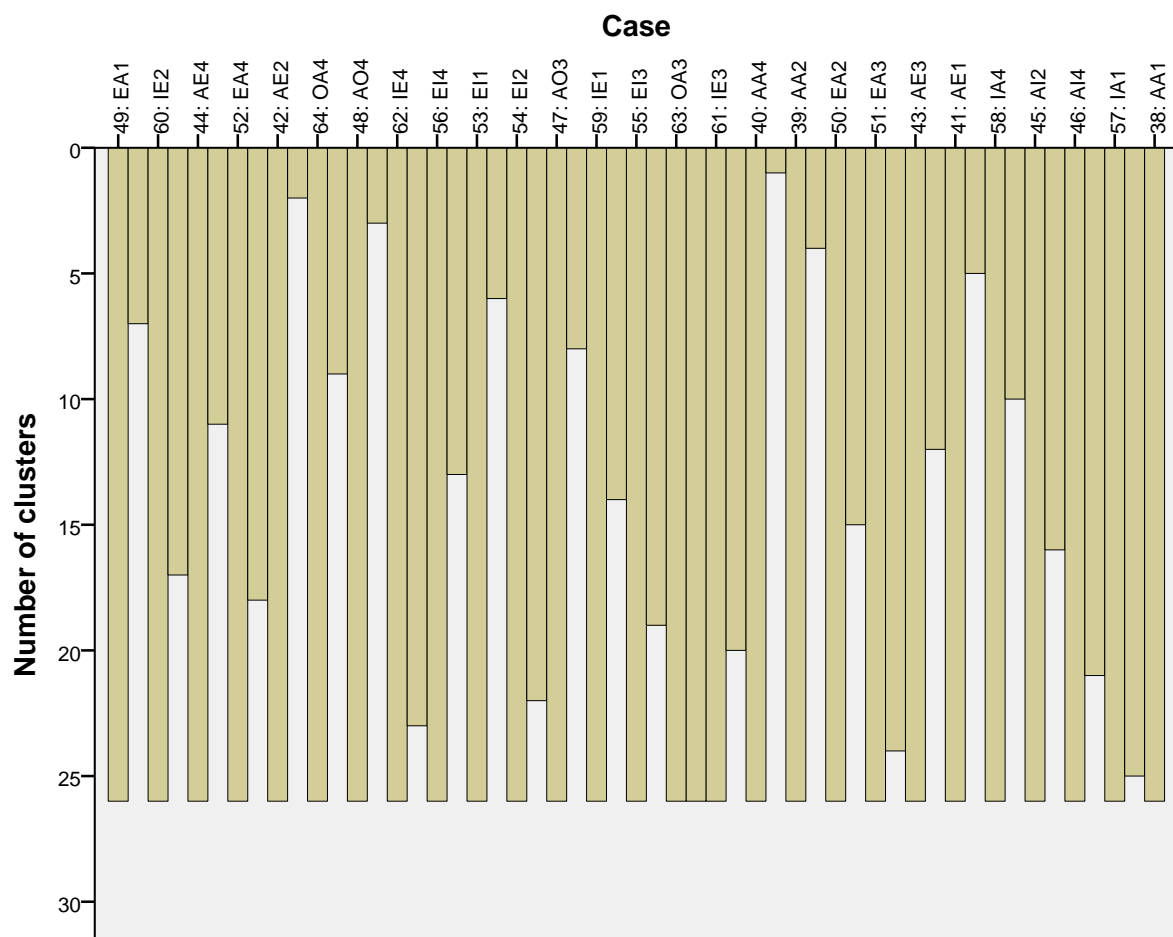
c. Ward Linkage

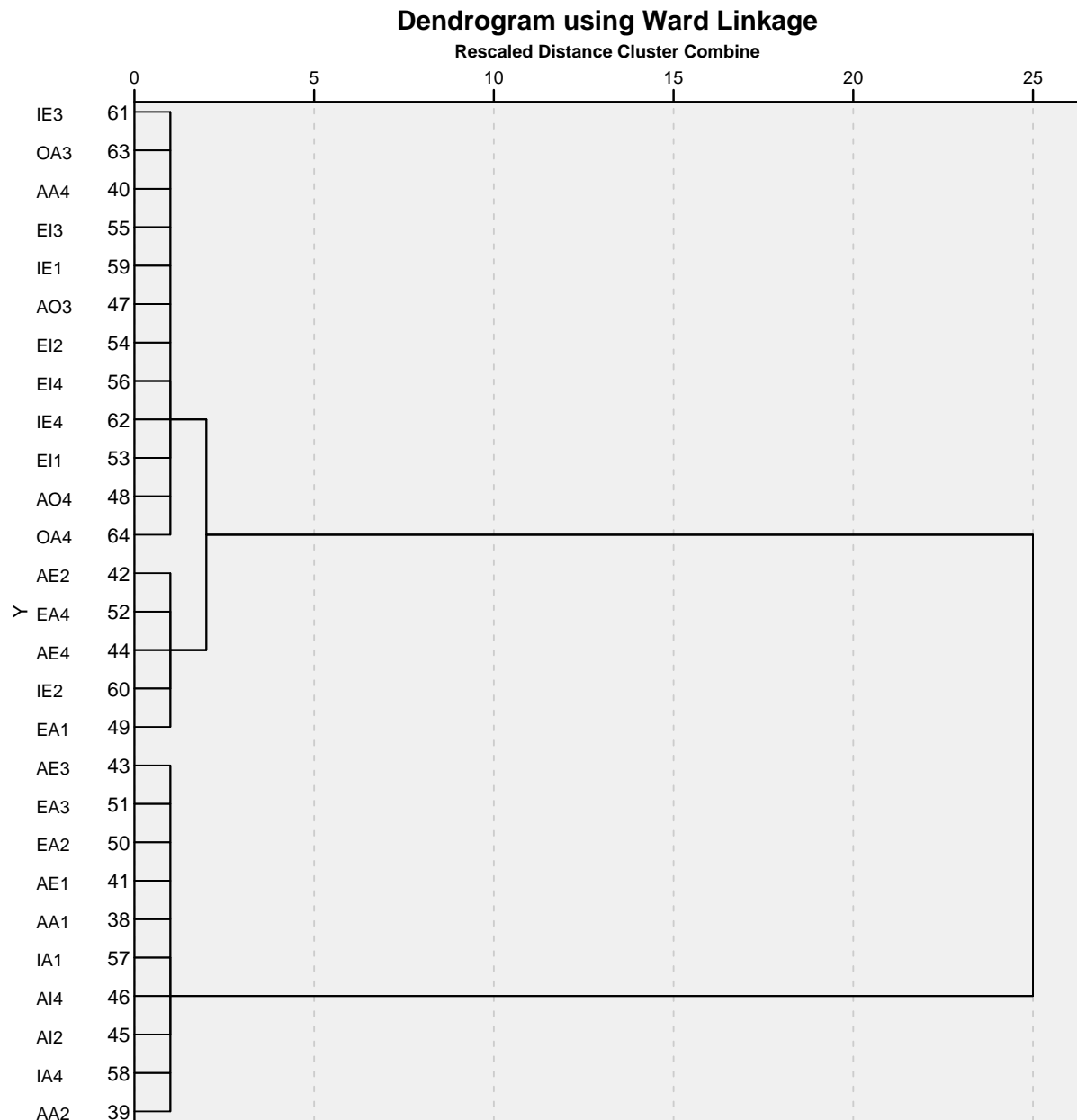
## Ward Linkage

### Agglomeration Schedule<sup>a</sup>

Stage	Cluster Combined		Coefficients	Stage Cluster First Appears		Next Stage
	Cluster 1	Cluster 2		Cluster 1	Cluster 2	
1	61	63	.000	0	0	7
2	38	57	.000	0	0	6
3	43	51	.000	0	0	12
4	56	62	.000	0	0	14
5	47	54	.000	0	0	19
6	38	46	.000	2	0	11
7	40	61	.000	0	1	8
8	40	55	.000	7	0	13
9	42	52	.000	0	0	16
10	44	60	.000	0	0	16
11	38	45	.001	6	0	17
12	43	50	.001	3	0	15
13	40	59	.001	8	0	19
14	53	56	.001	0	4	21
15	41	43	.002	0	12	22
16	42	44	.003	9	10	20
17	38	58	.004	11	0	22
18	48	64	.007	0	0	24
19	40	47	.009	13	5	21
20	42	49	.014	16	0	25
21	40	53	.020	19	14	24
22	38	41	.030	17	15	23
23	38	39	.049	22	0	26
24	40	48	.101	21	18	25
25	40	42	.205	24	20	26
26	38	40	1.768	23	25	0

a. HasValidConcl = Y





```

QUICK CLUSTER Acc204
/MISSING=LISTWISE
/CRITERIA=CLUSTER(3) MXITER(10) CONVERGE(0)
/METHOD=KMEANS(NOUPDATE)
/SAVE CLUSTER
/PRINT ID(syllogism) INITIAL ANOVA CLUSTER DISTAN.

```

**K-Means, 3 clusters**

**HasValidConcl = N**

### Initial Cluster Centers<sup>a</sup>

	Cluster		
	1	2	3
Acc204	.4804	.0539	.2647

a. HasValidConcl = N

### Iteration History<sup>a,b</sup>

Iteration	Change in Cluster Centers		
	1	2	3
1	.037	.043	.006
2	.000	.010	.004
3	.000	.000	.000

a. HasValidConcl = N

b. Convergence achieved due to no or small change in cluster centers. The maximum absolute coordinate change for any center is .000. The current iteration is 3. The minimum distance between initial centers is ....

### Cluster Membership<sup>a</sup>

Case Number	syllogism	Cluster	Distance
1	AA3	3	.025
2	AI1	2	.029
3	AI3	3	.078
4	AO1	2	.019
5	AO2	2	.020
6	EE1	3	.029
7	EE2	3	.029
8	EE3	1	.037
9	EE4	1	.002
10	EO1	3	.049
11	EO2	3	.029
12	EO3	3	.025
13	EO4	3	.005
14	IA2	2	.053
15	IA3	2	.060
16	II1	3	.049
17	II2	3	.054
18	II3	3	.078
19	II4	3	.054
20	IO1	3	.044



**Cluster Membership<sup>a</sup>**

Case Number	syllogism	Cluster	Distance
21	IO2	3	.010
22	IO3	3	.059
23	IO4	3	.010
24	OA1	2	.020
25	OA2	2	.001
26	OE1	3	.054
27	OE2	3	.010
28	OE3	3	.049
29	OE4	3	.025
30	OI1	3	.034
31	OI2	3	.078
32	OI3	3	.029
33	OI4	3	.049
34	OO1	3	.010
35	OO2	3	.015
36	OO3	1	.032
37	OO4	1	.007

a. HasValidConcl = N

**Final Cluster Centers<sup>a</sup>**

	Cluster		
	1	2	3
Acc204	.4436	.1071	.2745

a. HasValidConcl = N

**Distances between Final Cluster Centers<sup>a</sup>**

Cluster	1	2	3
1		.336	.169
2	.336		.167
3	.169	.167	

a. HasValidConcl = N

**ANOVA<sup>a</sup>**

	Cluster		Error		F	Sig.
	Mean Square	df	Mean Square	df		
Acc204	.152	2	.002	34	85.994	.000

The F tests should be used only for descriptive purposes because the clusters have been chosen to maximize the differences among cases in different clusters. The observed significance levels are not corrected for this and thus cannot be interpreted as tests of the hypothesis that the cluster means are equal.

a. HasValidConcl = N

**Number of Cases in each Cluster<sup>a</sup>**

Cluster	1	4.000
	2	7.000
	3	26.000
Valid		37.000
Missing		.000

a. HasValidConcl = N

**HasValidConcl = Y****Initial Cluster Centers<sup>a</sup>**

	Cluster		
	1	2	3
Acc204	.7990	.3922	.0490

a. HasValidConcl = Y

**Iteration History<sup>a,b</sup>**

Iteration	Change in Cluster Centers		
	1	2	3
1	.072	.090	.097
2	.000	.000	.000

a. HasValidConcl = Y

b. Convergence achieved due to no or small change in cluster centers. The maximum absolute coordinate change for any center is .000. The current iteration is 2. The minimum distance between initial centers is ....

**Cluster Membership<sup>a</sup>**

Case Number	syllogism	Cluster	Distance
38	AA1	1	.001
39	AA2	1	.129
40	AA4	2	.047
41	AE1	1	.072
42	AE2	3	.044
43	AE3	1	.052
44	AE4	3	.001
45	AI2	1	.021
46	AI4	1	.006
47	AO3	2	.008
48	AO4	2	.159
49	EA1	3	.097
50	EA2	1	.033
51	EA3	1	.052
52	EA4	3	.034
53	EI1	3	.050
54	EI2	2	.003
55	EI3	2	.057
56	EI4	3	.069
57	IA1	1	.001
58	IA4	1	.050
59	IE1	2	.032
60	IE2	3	.019
61	IE3	2	.052
62	IE4	3	.074
63	OA3	2	.052
64	OA4	2	.090

a. HasValidConcl = Y

**Final Cluster Centers<sup>a</sup>**

	Cluster		
	1	2	3
Acc204	.7270	.3017	.1464

a. HasValidConcl = Y

**Distances between Final Cluster Centers<sup>a</sup>**

Cluster	1	2	3
1		.425	.581
2	.425		.155
3	.581	.155	

a. HasValidConcl = Y

**ANOVA<sup>a</sup>**

	Cluster		Error		F	Sig.
	Mean Square	df	Mean Square	df		
Acc204	.833	2	.004	24	195.191	.000

The F tests should be used only for descriptive purposes because the clusters have been chosen to maximize the differences among cases in different clusters. The observed significance levels are not corrected for this and thus cannot be interpreted as tests of the hypothesis that the cluster means are equal.

a. HasValidConcl = Y

**Number of Cases in each Cluster<sup>a</sup>**

Cluster	1	10.000
	2	9.000
	3	8.000
Valid		27.000
Missing		.000

a. HasValidConcl = Y