Family Expressiveness Questionnaire

Fast Track Project Technical Report
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FAST Track Technical Report Reference

Greenberg, M., Mason, C., & Lengua, L. & Conduct Problems Prevention Research Group (1995). <u>Family expressiveness questionnaire: Technical report</u>. Unpublished Manuscript, University of Washington, Seattle, WA.

I. Source

This 13-item measure is a revision of the original 40-item measure developed by Halberstadt (1986). This parent report measure assesses the frequency and nature of family communication of both positive and negative emotional states.

II. Scale Derivation

A principal components analysis (with varimax rotation) of the "Family Expressiveness Questionnaire" based on Year One, Cohort One Normative Sample data was performed. By default, all components with eigenvalues greater than 1 were extracted. The resulting 2 factor solution accounted for 52% of the variance:

Mean

Scale		Fact	Fact.		Scale	Inter-Item
Name	Items	One	Two	Item Description	Alpha	Correlatio n
Negative Expression P1gfeneg	P1G1FE11 P1G1FE5 P1G1FE6 P1G1FE13 P1G1FE8 P1G1FE7 P1G1FE12 P1G1FE2	.77 .75 .70 .70 .67 .52 .52	.41 .45	Disappointed/empathy for another member Mentions another's behavior bothers them Talk to someone when their feelings hurt Discusses reasons for anger Explains point of view Cheer up another family member Happy Tell someone something is bothering them	.82	.44
Positive	P1G1FE4		.77	Someone shows love	.82	.39
Expression	P1G1FE9		.71	Hug or snuggle		
P1gfepos	P1G1FE10		.71	Praise another family member		
	P1G1FE1		.69	Tell someone they look nice		
	P1G1FE3		.58	Someone says "I am sorry"		
NOTE N				tee 1		

NOTE: Values less than 0.4 have been omitted.

Factor 1 corresponds to a "Negative Expression", while factor 2 appears to correspond to "Positive Expression". Factor based scales were calculated by: 1) rescaling all items to a possible 0-10 range, 2) summing the values of items on each factor, 3) dividing by the number of items present or responded to by each subject, and 4) multiplying by the number of items on each scale (6 items for negative expression and 7 for positive expression). While the loadings for items #7 and #12 were greater on Factor One than on Factor Two, both were included on the "Positive Expression" scale because 1) they had adequate loadings on Factor Two, and 2) they were conceptually consistent with the positive expression factor. Subscale scores were considered missing if greater than 25% of the items on a scale were missing.

III. Missing Data

One subject (Normative sample) was missing data on the negative and positive expression subscales. For the Normative sample, N = 386; for the High-risk sample, N = 310.

IV. Descriptives

The mean, standard deviation, skewness and kurtosis of each of the subscales for the Normative and High-Risk samples are presented in the following table. None of the subscale distributions deviated markedly formed a normal distribution.

Normative Sample

Subscale	Mean	S.D.	Skewness	Kurtosis
Positive	51.71	11.57	-0.58	0.50
Expression				
Negative	37.54	10.96	0.11	-0.48
Expression				

High-Risk Sample

Subscale	Mean	S.D.	Skewness	Kurtosis
Positive	49.26	11.59	-0.41	0.32
Expression				
Negative	36.55	11.45	0.28	-0.53
Expression				

IV. Intercorrelations

The correlations between the Positive Expression and Negative Expression subscales for the Normative and High-risk samples were 0.64 and 0.61 respectively.

VI. Site Differences

Mean site differences were explored using PROC GLM in SAS. There were no significant site, intervention, or site x intervention differences.

		F	p value	
Normative Sa	mple			
Positive Expression		2.47	0.06	
Negative Expression		2.29	0.08	
	Based on Cohort 1, Yea General Line			
Level of SITE N DURH 100 NASH 100 PENN 98 WASH 88	P1GFEPOS- Mean 53.7333333 49.4500000 51.2925170 52.4242424	SD 12.2202019 12.1878649 9.6518153 11.7523104	P1GFENEG Mean 39.4200000 35.533333 37.0748299 38.2196970	SD 11.6617710 10.7749137 10.1677263 10.9561866
		F	p value	
Normative Sa	mple			
Positive Expression		2.47	0.06	
Negative Expression		2.29	0.08	
	Based on Cohort 1, Yea General Line			
Level of SITE N DURH 78 NASH 81 PENN 81 WASH 71	P1GFEPOS- Mean 49.6153846 48.3470508 48.0833333 51.2206573	SD 12. 0242752 13.3581873 10.5111956 9.9560851	P1GFENEG Mean 36.7606838 36.5432099 34.9000000 38.1690141	SD 12.2761512 11.8725225 9.5943607 11.9361892