

POSITIVE AFFECT SCORING MANUAL

A brief guide to scoring the PROMIS® Positive Affect instruments:

ADULT	PEDIATRIC	EARLY CHILDHOOD PARENT-REPORT	PROXY
PROMIS Item Bank v1.0 Positive Affect PROMIS Short Form v1.0 Positive Affect 15a	PROMIS Pediatric Item Bank v1.0 Positive Affect PROMIS Pediatric Short Form v1.0 Positive Affect 4a PROMIS Pediatric Short Form v1.0 Positive Affect 8a	PROMIS Early Childhood Parent- Report Bank v1.0 – Positive Affect PROMIS Early Childhood Parent- Report Short Form v1.0 – Positive Affect 4a PROMIS Early Childhood Parent- Report Short Form v1.0 – Positive Affect 8a	PROMIS Proxy Item Bank v1.0 Positive Affect PROMIS Proxy Short Form v1.0 Positive Affect 4a PROMIS Proxy Short Form v1.0 Positive Affect 8a

SCORING THE INSTRUMENT

Short Forms: PROMIS instruments are scored using item-level calibrations. This means that the most accurate way to score a PROMIS instrument is to use the HealthMeasures Scoring Service (https://www.assessmentcenter.net/ac_scoring-service) or a data collection tool that automatically calculates scores (e.g., REDCap auto-score). This method of scoring uses responses to each item for each participant. We refer to this as “response pattern scoring.” Because response pattern scoring is more accurate than the use of raw score/scale score look up tables included in this manual, it is preferred. Response pattern scoring is especially useful when there is missing data (i.e., a respondent skipped an item), different groups of participants responded to different items, or you have created a new questionnaire using a subset of questions from a PROMIS item bank.

To use the scoring tables in this manual, calculate a summed score. Each question usually has five response options ranging in value from one to five. To find the total raw score for a short form with all questions answered, sum the values of the response to each question. For example, for the v1.0 pediatric 4-item form, the lowest possible raw score is 4; the highest possible raw score is 20 (see all short form scoring tables in the Appendix). **All questions must be answered in order to produce a valid score using the scoring tables.** If a participant has skipped a question, use the HealthMeasures Scoring Service (https://www.assessmentcenter.net/ac_scoring-service) to generate a final score.

With the total raw score for a measure, locate the applicable score conversion table in the Appendix and use this table to translate the total raw score into a T-score for each participant. The T-score rescales the raw score into a standardized score with a mean of 50 and a standard deviation (SD) of 10. Therefore a person with a T-score of 40 is one SD below the mean.

For the PROMIS Pediatric 4a short form v1.0, a raw score of 10 converts to a T-score of 35.4 with a standard error (SE) of 2.5 (see scoring table for the 4a v1.0 short form in the Appendix). Thus, the 95% confidence interval around the observed score ranges from 30.5 to 40.3 (T-score $\pm$ (1.96*SE) or 35.4 $\pm$ (1.96*2.5)).

CAT: A minimum number of items (4 for adult and 5 for peds and parent proxy CATs) must be answered in order to receive a score for Positive Affect CAT. The response to the first item will guide the system’s choice of the

next item for the participant. The participant's response to the second item will dictate the selection of the following question, and so on. As additional items are administered, the potential for error is reduced and confidence in the respondent's score increases. CAT will continue until either the standard error drops below a specified level (on the T-score metric 3.0 for adult and 4.0 for peds and parent proxy CATs), or the participant has answered the maximum number of questions (12), whichever occurs first.

SCORES

For most PROMIS instruments, a score of 50 is the average for the United States general population with a standard deviation of 10 because calibration testing was performed on a large sample of the general population. You can read more about the calibration and centering samples on HealthMeasures.net (<http://www.healthmeasures.net/score-and-interpret/interpret-scores/promis>). The T-score is provided with an error term (Standard Error or SE). The Standard Error is a statistical measure of variance and represents the "margin of error" for the T-score.

Important: *A higher PROMIS T-score represents more of the concept being measured. For positively-worded concepts like Positive Affect, a T-score of 60 is one SD better than average. By comparison, a Positive Affect T-score of 40 is one SD worse than average.*

Standard Error (SE): A PROMIS score includes a T-score and a standard error (SE). The standard error is a measure of the variability for a given T-score across hypothetical repeated measurements. The standard error can be used to construct confidence intervals around a T-score. A 95% confidence interval is common. A 95% confidence interval means there is a 95% probability that the true T-score is within this range. The formula for a 95% confidence interval is $(T\text{-score} \pm (1.96 * SE))$. For example, if $T=52$ and $SE=2$, the lower boundary of the confidence interval is $(52 - (1.96 * 2)) = 48$ and the upper boundary is $(52 + (1.96 * 2)) = 56$.

FREQUENTLY ASKED QUESTIONS (FAQs)

Q: I am interested in learning more. Where can I do that?

Review the HealthMeasures website at www.healthmeasures.net.

Q: How do I handle multiple responses when administering a short form on paper?

Guidelines on how to deal with multiple responses have been established. Resolution depends on the responses noted by the research participant.

- If two or more responses are marked by the respondent, and they are next to one another, then a data entry specialist will be responsible for randomly selecting one of them to be entered and will write down on the form which answer was selected. Note: To randomly select one of two responses, the data entry specialist will flip a coin (heads - higher number will be entered; tails – lower number will be entered). To randomly select one of three (or more) responses, a table of random numbers should be used with a statistician's assistance.
- If two or more responses are marked, and they are NOT all next to one another, the response will be considered missing.



Q: What is the minimum change on a PROMIS instrument that represents a clinically meaningful difference? To learn more about research on the meaning of a change in scores, we suggest conducting a literature review to identify the most current information. The HealthMeasures website (<http://www.healthmeasures.net/score-and-interpret/interpret-scores/promis>) has additional information on interpreting scores.

APPENDIX-SCORING TABLES – ADULT

Positive Affect 15a – Adult v1.0					
Short Form Conversion Table					
Raw Summed Score	T-score	SE*	Raw Summed Score	T-score	SE*
15	14.4	2.7	46	40.9	2.1
16	15.7	3.0	47	41.6	2.1
17	17.3	3.0	48	42.2	2.1
18	18.7	3.0	49	42.9	2.1
19	20.1	2.9	50	43.5	2.1
20	21.3	2.7	51	44.2	2.1
21	22.5	2.6	52	44.8	2.1
22	23.5	2.5	53	45.5	2.1
23	24.5	2.4	54	46.2	2.1
24	25.4	2.3	55	46.9	2.1
25	26.3	2.3	56	47.5	2.1
26	27.1	2.2	57	48.2	2.1
27	27.9	2.2	58	48.9	2.1
28	28.7	2.2	59	49.6	2.1
29	29.4	2.1	60	50.3	2.1
30	30.2	2.1	61	51.1	2.1
31	30.9	2.1	62	51.8	2.1
32	31.6	2.1	63	52.6	2.1
33	32.3	2.1	64	53.4	2.2
34	33.0	2.1	65	54.2	2.2
35	33.7	2.1	66	55.0	2.3
36	34.4	2.1	67	55.9	2.3
37	35.0	2.1	68	56.9	2.4
38	35.7	2.1	69	58.0	2.6
39	36.4	2.1	70	59.2	2.8
40	37.0	2.1	71	60.5	3.0
41	37.7	2.1	72	62.1	3.3
42	38.3	2.1	73	63.9	3.7
43	39.0	2.1	74	66.3	4.2
44	39.6	2.1	75	69.9	5.1
45	40.3	2.1			

*SE = Standard Error on T-score

Positive Affect 4a – Early Childhood Parent-Report v1.0		
<i>Short Form Conversion Table</i>		
Raw Summed Score	T-score	SE*
4	14.5	2.6
5	16.6	2.6
6	18.7	2.5
7	20.5	2.3
8	22.1	2.3
9	23.7	2.4
10	25.6	2.6
11	27.5	2.6
12	29.6	2.6
13	31.8	2.6
14	34.4	2.7
15	37.7	3.3
16	42.5	3.6
17	46.6	3.1
18	50.0	2.7
19	53.3	3.1
20	60.5	5.8

*Standard Error on the T-score metric

Positive Affect 8a – Early Childhood Parent-Report v1.0		
<i>Short Form Conversion Table</i>		
Raw Summed Score	T-score	SE*
8	12.5	1.9
9	13.6	2.2
10	14.7	2.3
11	15.9	2.3
12	17.0	2.2
13	18.1	2.1
14	19.2	2.0
15	20.2	2.0
16	21.1	1.9
17	22.0	2.0
18	23.0	2.0
19	24.0	2.0
20	25.0	2.1
21	26.2	2.2
22	27.4	2.3
23	28.6	2.3
24	29.8	2.3
25	31.0	2.3
26	32.4	2.2
27	33.8	2.3
28	35.3	2.5
29	37.1	2.8
30	39.0	3.0
31	41.2	3.1
32	43.3	3.0
33	45.5	2.8
34	47.7	2.5
35	49.6	2.3
36	51.3	2.3
37	53.1	2.5
38	55.4	3.1
39	58.5	4.0
40	63.6	5.6

*Standard Error on the T-score metric

Positive Affect 4a – Parent Proxy v1.0		
<i>Short Form Conversion Table</i>		
Raw Summed Score	T-score	SE*
4	16.8	2.9
5	19.9	2.5
6	22.3	2.4
7	24.5	2.5
8	26.5	2.5
9	28.5	2.5
10	30.8	2.6
11	33.3	2.6
12	36.1	2.6
13	38.7	2.5
14	41.3	2.5
15	44.3	2.8
16	48.3	3.1
17	52.0	2.8
18	54.9	2.5
19	57.8	2.8
20	63.8	5.1

*Standard Error on the T-score metric

Positive Affect 8a – Parent Proxy v1.0		
<i>Short Form Conversion Table</i>		
Raw Summed Score	T-score	SE*
8	13.4	2.2
9	14.7	2.4
10	16.4	2.4
11	18.0	2.3
12	19.6	2.2
13	21.0	2.1
14	22.3	2.1
15	23.6	2.1
16	24.9	2.1
17	26.1	2.1
18	27.3	2.1
19	28.6	2.1
20	29.9	2.1
21	31.3	2.2
22	32.8	2.3
23	34.3	2.3
24	35.8	2.3
25	37.3	2.3
26	38.8	2.2
27	40.3	2.2
28	41.8	2.2
29	43.5	2.4
30	45.3	2.5
31	47.2	2.6
32	49.1	2.6
33	51.1	2.5
34	53.0	2.3
35	54.7	2.2
36	56.3	2.2
37	58.0	2.4
38	60.0	2.7
39	62.7	3.5
40	67.3	5.0

*Standard Error on the T-score metric

APPENDIX-SCORING TABLES – PEDIATRIC

Positive Affect 4a – Pediatric v1.0		
<i>Short Form Conversion Table</i>		
Raw Summed Score	T-score	SE*
4	22.0	3.6
5	25.7	2.7
6	28.0	2.5
7	30.0	2.5
8	31.8	2.4
9	33.6	2.5
10	35.4	2.5
11	37.4	2.5
12	39.5	2.5
13	41.6	2.5
14	43.8	2.6
15	46.2	2.6
16	48.7	2.7
17	51.2	2.6
18	53.8	2.6
19	56.8	3.0
20	63.0	5.3

*Standard Error on the T-score
metric

Positive Affect 8a – Pediatric v1.0		
<i>Short Form Conversion Table</i>		
Raw Summed Score	T-score	SE*
8	19.1	3.4
9	21.9	2.9
10	23.8	2.6
11	25.3	2.3
12	26.6	2.2
13	27.8	2.1
14	28.9	2.1
15	30.0	2.1
16	31.0	2.1
17	32.0	2.1
18	33.0	2.1
19	34.0	2.1
20	35.1	2.1
21	36.2	2.1
22	37.3	2.2
23	38.4	2.2
24	39.5	2.2
25	40.7	2.2
26	41.9	2.2
27	43.1	2.2
28	44.3	2.2
29	45.6	2.2
30	46.9	2.3
31	48.3	2.3
32	49.6	2.3
33	50.9	2.2
34	52.3	2.2
35	53.7	2.2
36	55.2	2.3
37	56.8	2.5
38	58.7	2.8
39	61.3	3.4
40	66.2	5.1

*Standard Error on the T-score metric