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# EORTC QLQ-C30 descriptive analysis with the `qlqc30` command

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**Abstract.** Health-related quality of life is often an endpoint in oncology clinical trials. The European Organization for Research and Treatment of Cancer (EORTC) developed the cancer-specific quality of life questionnaire (QLQ-C30), which includes five functions, nine symptoms, and a global health status. These questionnaires are completed by the patients themselves throughout the process of care. The recommended approaches for processing EORTC QLQ-C30 data are usually descriptive and graphic.

Our aim was to develop a user-written command that provided an automatic descriptive analysis of EORTC QLQ-C30 data, consisting of profile plots per visit and longitudinal plots per functional and symptom scale.

**Keywords:** dm0084, `qlqc30`, health-related quality of life, QLQ-C30, functional scale, symptom scale, profile plot, longitudinal plot

## 1 Introduction

The health-related quality of life (HRQoL) is a multidimensional, subjective, and dynamic concept incorporating at least three domains: physical, psychological, and social functioning. It overlaps the definition of health given by the World Health Organization

in 1948. This concept refers to a patient's perception of his or her treatment and illness, although indirect consequences such as unemployment or financial difficulties are sometimes considered.

HRQoL falls within the scope of "patient-reported outcomes" (Gotay et al. 2008), that is, of measures reported by the subjects themselves. According to the American Society of Clinical Oncology and to the Food and Drug Administration (Beitz, Gnecco, and Justice 1996), when an experimental intervention has no significant effect on overall survival, HRQoL should be considered a second primary endpoint.

The European Organization for Research and Treatment of Cancer (EORTC) has developed the cancer-specific quality of life questionnaire (QLQ-C30) (Aaronson et al. 1993), which consists of five functional scales (physical, role, cognitive, emotional, and social); nine symptom scales (fatigue, pain, nausea and vomiting, dyspnea, loss of appetite, insomnia, constipation, diarrhea, and financial difficulties); and a global health status/quality of life (GHS/QoL). Based on 30 questions in total, the scores range from 0 to 100. This questionnaire was approved and psychometrically validated (as well as each of its translations) for use in oncology clinical trials. Such standardization allows the comparison of results. Generally, the questionnaires are collected at different times predefined in the study protocol according to the EORTC recommendations (Fayers et al. 2001). The EORTC QLQ-C30 is a common basis often supplemented by tumor-specific modules, such as the QLQ-BR23, QLQ-OES18 or QLQ-OG25, and QLQ-PAN26 modules adapted to breast, esophageal, and pancreatic cancers, respectively. These uni- or multi-item scores allow one to indirectly evaluate the patients' quality of life in each dimension.

The EORTC proposed a descriptive and transversal approach for processing QLQ-C30 data. Using Stata, we developed a QLQ-C30-specific command, `qlqc30`, that implements the scoring procedures according to the algorithms recommended by the EORTC (Fayers et al. 2001) and includes descriptive analyses at each measurement time as well as transversal and longitudinal graphics. In section 2, we introduce the scoring procedure recommended by the EORTC. In section 3, we describe the command and present the obtained syntax, options, and outputs. In section 4, we provide an example with results based on data from a clinical trial. In section 5, we conclude.

## 2 Method

The scoring methodology is mainly based on the observation of scores that are supposed to be close to the "real" score. Therefore, the observed raw score is a direct measurement of the patients' HRQoL and is estimated by the average of the items composing the scale. The standardized score is expressed on a 0-to-100 scale after a linear transformation. Table 1 summarizes the items to be considered in the score calculation for the GHS, the functional, and the symptom scales (QLQ-C30 version 3). Only the five items of the physical functioning (`q1`–`q5`) were modified in the third version of the questionnaire: they switched from dichotomous items (no or yes) in QLQ-C30 version 2 to ordinal items (not at all, a little, quite a bit, or very much) in QLQ-C30 version 3. The command allows both versions of the questionnaire to be considered.

Table 1. GHS, functional, and symptom scales

	Scale	Number of items	Number of levels	Item numbers	Score	
Functional scales						
Physical functioning	PF2	5	4	q1-q5	$X = \left\{ 1 - \frac{\left( \frac{1}{n} \sum_{k=1}^n I_k \right) - 1}{r} \right\} \times 100$	
Role functioning	RF2	2	4	q6, q7		
Emotional functioning	EF	4	4	q21-q24		
Cognitive functioning	CF	2	4	q20, q25		
Social functioning	SF	2	4	q26, q27		
Symptom scales						
Fatigue	FA	3	4	q10, q12, q18	$X = \frac{\left( \frac{1}{n} \sum_{k=1}^n I_k \right) - 1}{r} \times 100$	
Nausea and vomiting	NV	2	4	q14, q15		
Pain	PA	2	4	q9, q19		
Dyspnea	DY	1	4	q8		
Insomnia	SL	1	4	q11		
Appetite loss	AP	1	4	q13		
Constipation	CO	1	4	q16		
Diarrhea	DI	1	4	q17		
Financial difficulties	FI	1	4	q28		
GHS						
		high response/score = high GHS				
GHS/QoL	QL2	2	7	q29, q30		

The EORTC recommends calculating algorithms as follows. We defined  $n$  ( $n = 1, \dots, 5$ ) as the number of items contributing to the scale being considered,  $(I_1, \dots, I_n)$  as the responses to items, and  $r$  (item range) as the difference between the possible maximum and the minimum response to individual items (for example, in the case of an item whose responses range from 1 to 4,  $r = 4 - 1 = 3$ ). The score is then equal to  $X = [ \{ (1/n \sum_{k=1}^n I_k) - 1 \} / r ] \times 100$  for symptom and GHS/QoL scales and  $X = (1 - [ \{ (1/n \sum_{k=1}^n I_k) - 1 \} / r ]) \times 100$  for functional scales. For functional scales and GHS/QoL, a high score reflects a high level of functional capacity (physical, role, etc.) or GHS with a good QoL. Conversely, for symptom scales, a high score is the expression of strong symptoms (fatigue, nausea and vomiting, etc.) associated with a poor QoL. Moreover, the scores can be calculated even in the event of missing data. Indeed, if at least half the items are known, the score is valid and becomes the standardized mean of the nonmissing items.

### 3 Program description

A flowchart describing `qlqc30` is presented in figure 1.

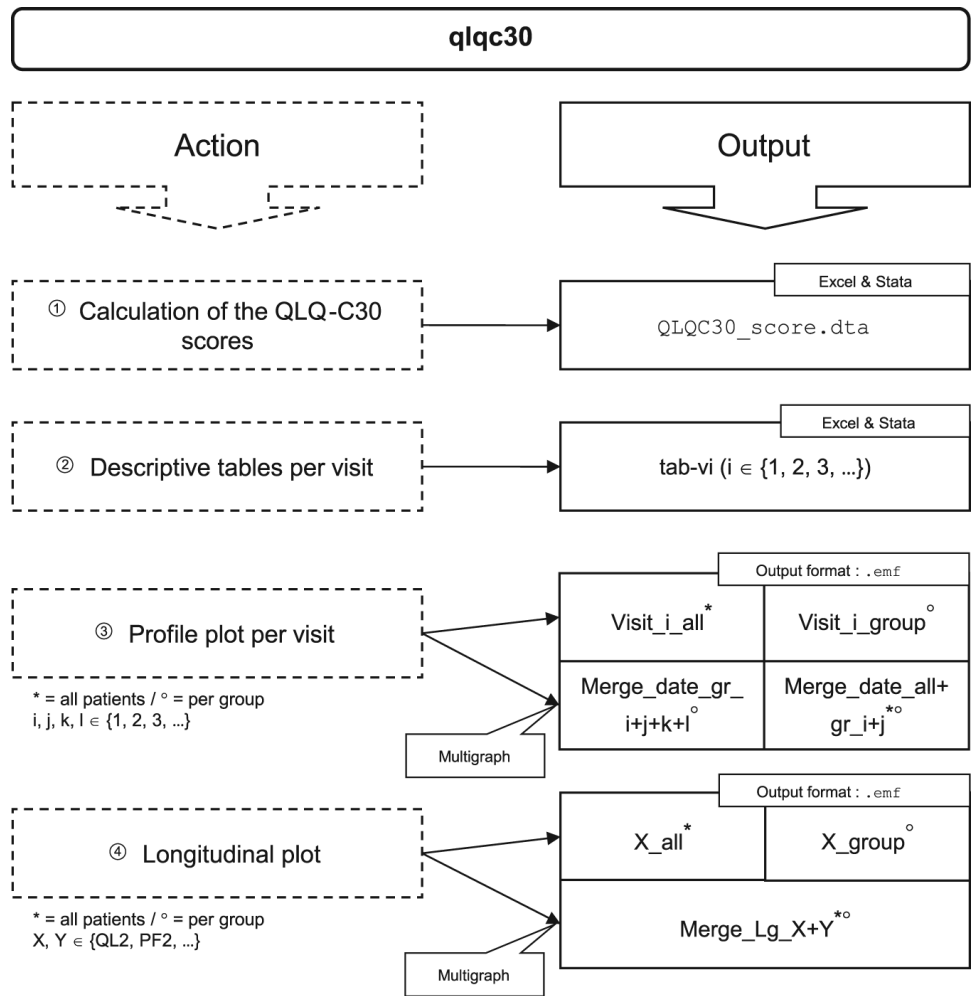


Figure 1. Flowchart describing the `qlqc30` command

The `qlqc30` command performs the following:

- It computes the different QLQ-C30 dimensions and creates a Stata dataset and an Excel file containing the calculated dimensions (named `QLQC30_score.dta` and `QLQC30_score.xls`, respectively).



- It computes the descriptive analysis of all scales for each visit and generates a Stata dataset and an Excel file containing the descriptive results for each visit.
- It displays profile plots (Fayers and Machin 2007) that are a form of transversal presentation particularly useful in HRQoL analyses (all scales are simultaneously presented at one specific visit).
- It displays longitudinal plots that represent the evolution over time of each dimension (the mean score is plotted for each visit with its 95% confidence interval if the number of patients is greater than or equal to 10).

The graphical representations may be performed for all patients or for a specified group, for example, according to the treatment arm.

### 3.1 Syntax

```
qlqc30, filename(string) version(integer) grp(integer) [path(string)
table(yes) graph(yes)]
```

`filename()` can be personalized, but the file structure should keep the same format. Each line contains an observation per pair patient-visit, whereas the columns contain the following variables:

Variable	Description
<code>npat</code>	patient number
<code>q1-q30</code>	QLQ-C30 items in numerical format: <code>q1</code> to <code>q5</code> variable values must range from 1 to 2 for the QLQ-C30 version 2 and from 1 to 4 for the QLQ-C30 version 3; <code>q6</code> to <code>q28</code> variable values must range from 1 to 4; and <code>q29</code> and <code>q30</code> variable values must range from 1 to 7
<code>arm</code>	coded 1 or 2; coded 2 only if the study comprises two arms and a comparison of those is sought; otherwise, the variable can be present and is always coded 1
<code>visit</code>	number of time units from the beginning of the study. For example, the value of this variable is equal to 0 at baseline, to 3 at three months, etc.

## 3.2 Options

`filename(string)` specifies the filename to use. `filename()` is required.

`version(integer)` specifies the version of QLQ-C30 used, either 2 or 3. `version()` is required.

`grp(integer)` specifies 1 for a single-arm study (one group) and 2 for a double-arm study (two groups). `grp()` is required.

`path(string)` specifies a direct file path.

`table(yes)` specifies that the descriptive tables in the specific subdirectory **Table** be obtained.

`graph(yes)` specifies that the profile and longitudinal plots in the specific subdirectory **Graph** be obtained.

The command does not support missing values in the following variables: **npat**, **arm**, and **visit**. Errors are due to missing data in the previous variables.

## 3.3 Displayed outputs

The `qlqc30` command displays results such as tables and graphs (only if desired) that are automatically saved in the working subdirectories named **Table** and **Graph**, respectively.

Output datasets (and Excel files), named `tab-vi.dta` (and `tab-vi.xls`), with  $i$  being the number of the described visit, contain the descriptive tables. Each scale is summarized by its mean, standard deviation (SD), median, and range for all patients or per treatment arm. The dimensions between the groups are compared using the Wilcoxon rank-sum test and the Student's  $t$  test.

Output profiles and longitudinal graphs are saved under Windows Enhanced Metafile format. Several output multigraphs are also saved; three kinds of combinations are proposed:

- a combination of global and per-group profile graphs for each visit:  
(`Merge_date_all+gr_i+j.emf`,  $i, j \in \{1, 2, 3, \dots\}$ );
- a combination of profile graphs for several consecutive visits:  
(`Merge_date_gr_i+j+k+l.emf`,  $i, j, k, l \in \{1, 2, 3, \dots\}$ ); and
- a combination of global and per-group longitudinal graphs for each dimension:  
(`Merge_Lg_X+Y.emf`,  $X, Y \in \{QL2, PF2, \dots\}$ ).

## 4 Example

We used the `qlqc30` command to analyze HRQoL data from the CO-HO-RT (concomitant hormono-radio therapy) clinical trial. This trial assessed the acute and late radiation-induced skin toxicities in 150 patients with breast cancer (Azria et al. 2010). After breast-conserving surgery, women were randomly assigned to receive concurrent radiotherapy and letrozole (arm 1,  $n = 74$ ) or sequential radiotherapy and letrozole (arm 2,  $n = 75$ ). QLQ-C30 evaluations were performed before the beginning of the treatment (baseline) and every 3 months over a 24-month period. Baseline compliance with the completion of QLQ-C30 was high: 73 (99%) and 70 (93%) questionnaires were completed in the concurrent and sequential groups, respectively. Thereafter, the compliance for the questionnaire completion decreased over time to reach 81% (60/74) in the concurrent group and 80% (60/75) in the sequential group at 24 months.

Table 2 provides a descriptive analysis of all domains at baseline, globally or per treatment arm. HRQoL at baseline did not differ significantly between the two groups. The results were similar for all the other visits (data not shown).

Table 2. Descriptive analysis of all scales at baseline

		Concurrent group (Arm 1)				
		N1	Mean1	SD1	Median1	Range1
GHS		73	69.63	(23.67)	75.00	[0,100]
	Missing	0				
Physical functioning		73	83.29	(19.72)	100.00	[20,100]
	Missing	0				
Role functioning		73	82.65	(26.71)	100.00	[0,100]
	Missing	0				
Emotional functioning		73	73.21	(23.42)	75.00	[0,100]
	Missing	0				
Cognitive functioning		73	77.85	(24.07)	83.33	[0,100]
	Missing	0				
Social functioning		73	89.73	(20.72)	100.00	[0,100]
	Missing	0				
Fatigue		73	27.32	(27.12)	22.22	[0,100]
	Missing	0				
Nausea and vomiting		73	4.79	(12.87)	0.00	[0,83.33]
	Missing	0				
Pain		73	21.00	(26.94)	16.67	[0,100]
	Missing	0				
Dyspnea		72	13.89	(24.86)	0.00	[0,100]
	Missing	1				
Insomnia		72	36.11	(33.45)	33.33	[0,100]
	Missing	1				
Appetite loss		71	9.39	(19.67)	0.00	[0,100]
	Missing	2				
Constipation		72	18.52	(29.01)	0.00	[0,100]
	Missing	1				
Diarrhea		72	9.26	(17.89)	0.00	[0,66.67]
	Missing	1				
Financial difficulties		73	6.85	(20.00)	0.00	[0,100]
	Missing	0				

*Continued on next page*

		Sequential group (Arm 2)				
		N2	Mean2	SD2	Median2	Range2
GHS		69	70.77	(23.60)	75.00	[0,100]
	Missing	1				
Physical functioning		70	84.50	(17.45)	90.00	[40,100]
	Missing	0				
Role functioning		70	84.29	(23.21)	100.00	[0,100]
	Missing	0				
Emotional functioning		69	73.35	(22.99)	83.33	[8.33,100]
	Missing	1				
Cognitive functioning		69	76.09	(22.59)	83.33	[0,100]
	Missing	1				
Social functioning		69	92.75	(15.25)	100.00	[33.33,100]
	Missing	1				
Fatigue		69	27.54	(22.84)	33.33	[0,100]
	Missing	1				
Nausea and vomiting		70	5.95	(12.70)	0.00	[0,66.67]
	Missing	0				
Pain		70	18.81	(25.68)	0.00	[0,100]
	Missing	0				
Dyspnea		70	12.86	(20.69)	0.00	[0,100]
	Missing	0				
Insomnia		70	32.86	(30.29)	33.33	[0,100]
	Missing	0				
Appetite loss		70	7.14	(16.92)	0.00	[0,66.67]
	Missing	0				
Constipation		69	15.94	(26.57)	0.00	[0,100]
	Missing	1				
Diarrhea		67	6.97	(18.84)	0.00	[0,100]
	Missing	3				
Financial difficulties		69	7.73	(22.25)	0.00	[0,100]
	Missing	1				

*Continued on next page*

		<i>N</i>	Mean	Total SD	Median	Range	Student's <i>t</i> test <i>p</i> -value	Wilcoxon rank-sum test <i>p</i> -value
GHS		142	70.19	(23.56)	75.00	[0,100]	0.775	0.835
	Missing	1						
Physical functioning		143	83.88	(18.59)	100.00	[20,100]	0.698	0.886
	Missing	0						
Role functioning		143	83.45	(24.98)	100.00	[0,100]	0.697	0.838
	Missing	0						
Emotional functioning		142	73.28	(23.13)	77.78	[0,100]	0.972	0.985
	Missing	1						
Cognitive functioning		142	77.00	(23.30)	83.33	[0,100]	0.653	0.485
	Missing	1						
Social functioning		142	91.20	(18.27)	100.00	[0,100]	0.325	0.673
	Missing	1						
Fatigue		142	27.43	(25.04)	22.22	[0,100]	0.959	0.469
	Missing	1						
Nausea and vomiting		143	5.36	(12.76)	0.00	[0,83.33]	0.589	0.526
	Missing	0						
Pain		143	19.93	(26.26)	16.67	[0,100]	0.619	0.601
	Missing	0						
Dyspnea		142	13.38	(22.83)	0.00	[0,100]	0.789	0.856
	Missing	1						
Insomnia		142	34.51	(31.86)	33.33	[0,100]	0.545	0.655
	Missing	1						
Appetite loss		141	8.27	(18.33)	0.00	[0,100]	0.469	0.593
	Missing	2						
Constipation		141	17.26	(27.78)	0.00	[0,100]	0.584	0.749
	Missing	2						
Diarrhea		139	8.15	(18.32)	0.00	[0,100]	0.463	0.405
	Missing	4						
Financial difficulties		142	7.28	(21.05)	0.00	[0,100]	0.804	0.977
	Missing	1						

Profile plots (figure 2) display all dimensions simultaneously, globally or divided according to the treatment variable. At baseline, the pattern was similar between the two groups.

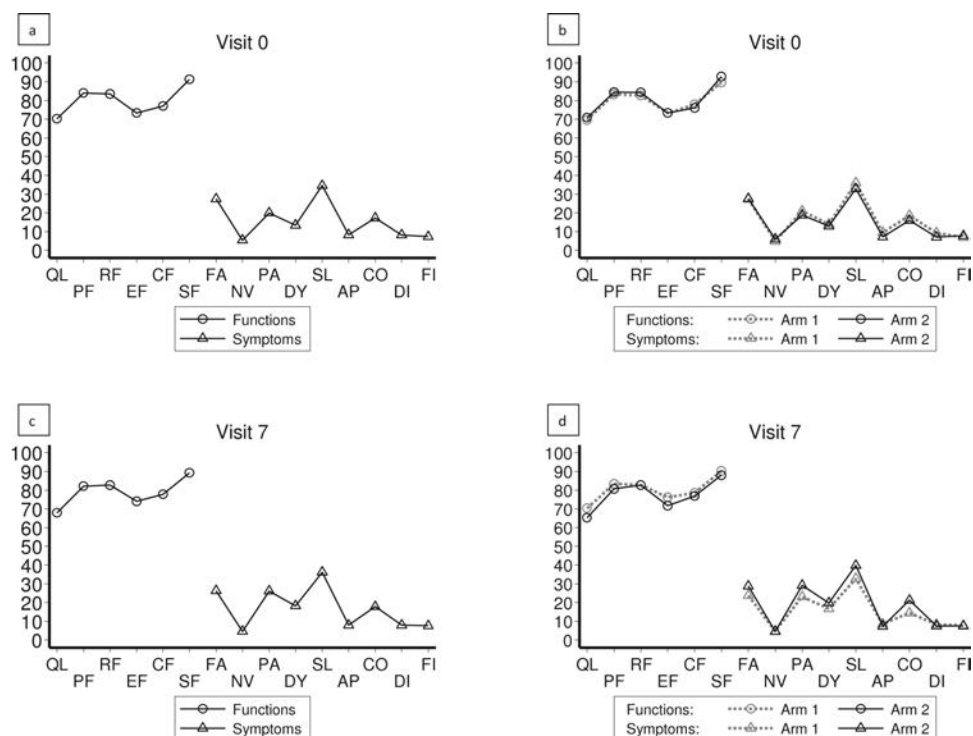


Figure 2. Profile plots at baseline (visit 0), globally (2a) and per group (2b), and at the 24th-month visit (visit 7), globally (2c) and per group (2d)

Longitudinal plots are also represented globally or per treatment arm for GHS/QoL (figure 3) and for some functional or symptom domains (figures 4 and 5). No difference could be observed between the treatment arms for any of the functional or symptom scales. Functional scores were high, and the less marked symptoms were nausea and vomiting, appetite loss, and diarrhea.

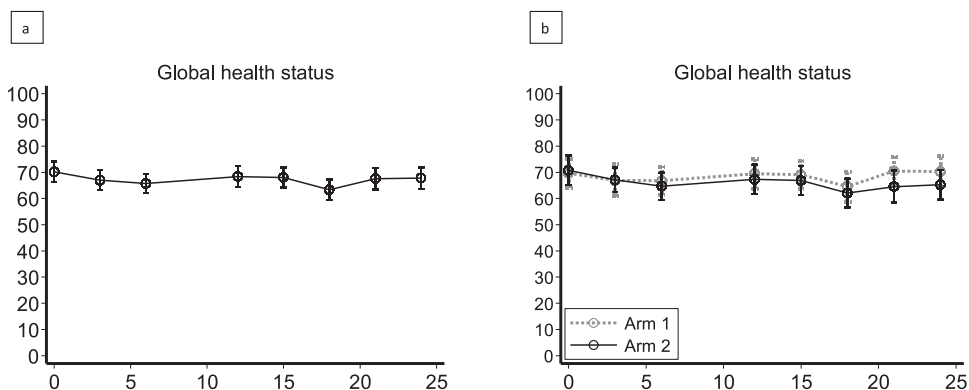


Figure 3. Change in mean EORTC QLQ-C30 GHS/QoL score over time, globally (3a) and per group (3b). For GHS/QoL, a high score indicates a good quality of life.

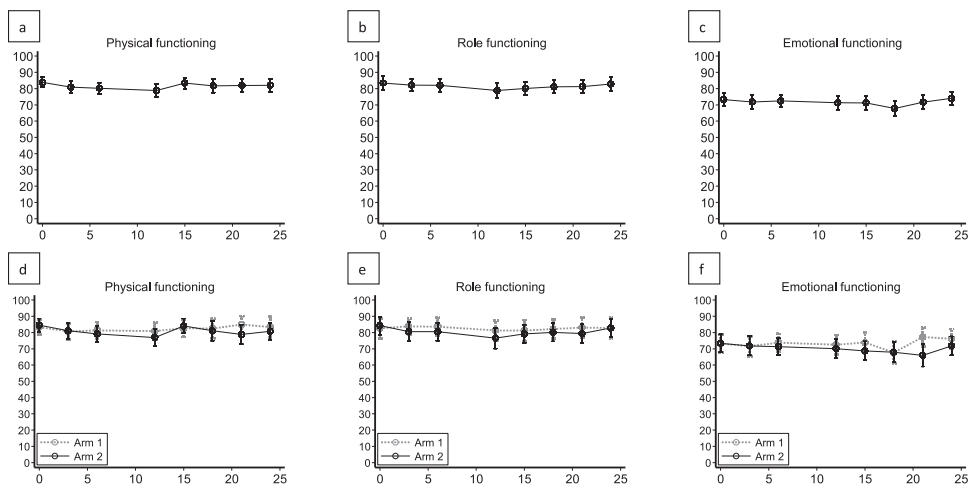
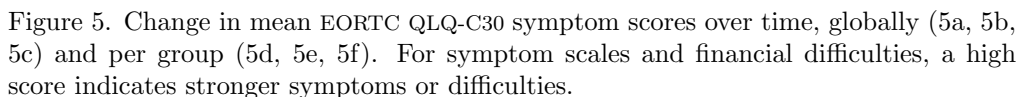


Figure 4. Change in mean EORTC QLQ-C30 functional scores over time, globally (4a, 4b, 4c) and per group (4d, 4e, 4f). For functional scales, a high score indicates a better function.





EORTC QLQ-C30 is completed by the patients themselves throughout the process of care. These uni- or multi-item scores indirectly reflect the patient's QoL. For the processing of such data, the EORTC recommends approaches that are usually descriptive and graphic. We developed a command providing an automatic descriptive analysis, profile plots per visit, and longitudinal plots per domain.

## 6 Acknowledgments

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## 7 References

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