

**Table 3.** Comparison of model fits<sup>a</sup> (linear vs. logarithmic) for entire sample and by form of treatment (available at <http://www.psyres.de/transop>)

Time segment	Sample N (obs./s.s <sup>b</sup> )	Fit	df	AIC	BIC	logLik
Before and in treatment	PAP	Linear	6	401.11	424.78	-194.55
	382/108	Logarithmic	6	391.16	414.84	-189.58
	PD	Linear	6	1052.23	1081.5	-520.11
	972/308	Logarithmic	6	1037.13	1066.41	-512.56
	Total	Linear	6	1449.99	1481.25	-718.99
	1,354/416	Logarithmic	6	1429.98	1461.24	-708.99
In treatment	PAP	Linear	6	339.51	362.05	-163.76
	317/102	Logarithmic	6	326.94	349.48	-157.47
	PD	Linear	6	864.72	892.59	-426.36
	773/282	Logarithmic	6	846.3	874.18	-417.15
	Total	Linear	6	1200.19	1230.13	-594.09
	1,090/384	Logarithmic	6	1173.07	1203.01	-580.54

<sup>a</sup> Originating from the log-restricted likelihood (logLik), the Akaike information criterion (AIC) and the Bayesian information criterion (BIC) in addition consider number of model parameters and observations. Comparison of GOF indices between models calculated with different samples is not meaningful.

<sup>b</sup> Observations of subjects.