

Table 8. Random genes

<u>Gene</u> <u>identification</u>	<u>30 min, %</u>	<u>45 min, %</u>	<u>120 min, %</u>
152963	-31.45	-29.29	-34.06
153085	6.76	-13.5	-3.39
155554	-5.96	4.65	56.97
156681	-1.98	-22.83	8.41
157184	-9.64	-7.2	14.74
157482	0.39	4.88	9.75
158830	-13.7	-0.96	-1.94
159026	36.24	-2.73	21.38
159393	-31.94	-30.87	35.62
159414	-5.94	-19.13	9.89
160388	11.17	-11.16	3.45
160518	-4	16	-14.7
162564	-8.42	-19.22	0.6
164658	-8.9	-17	29.08
165045	0.15	8.92	12.03
167309	4.27	-37.77	-23.03
167940	34.43	-18.57	21.57
170062	-4.16	-8.05	-8.13
170932	15.57	7.53	2.49
171282	-4.24	-10.2	0.56
171962	23.77	-11.69	16.86

Data set used to validate microarray detection of induction of transcripts and the strategy of flagellar gene identification based on induction during flagellar regeneration. Gene list was generated by a random number generator, without any information concerning gene identity. Entries sorted by gene identification no.