

Table 1: Numbers of probes in the original data (Original) and after preprocessing (Processed) having the characteristics shown under the Measure column.

Measure		prefrontal	cerebellum	temporal	pons
<i>Total</i>					
	Number of probes	20000	20000	20000	20000
<i>known Chr. names</i>					
	Original	17945	17953	17957	17973
	Processed	19921	19922	19922	19928
<i>known txStart^a</i>					
	Original	17945	17953	17957	17973
	Processed	19917	19917	19919	19923
<i>LOCxx^b</i>					
	Original	452	461	461	462
	Processed	147	156	145	144
<i>Gene symbol: LOCxx and unknown Chromosome^c</i>					
	Original	89	88	92	89

^a transcription start position

^b gene symbols beginning with *LOC*; example: *LOC730412*

^c After preprocessing there were no probes with gene symbols *LOCxx* and with unknown chromosome names.