Supplementary Figure 1: Volcano plots showing the relationship between fold change of testicular versus spermatozoal transcript detection and the level of significance as represented by raw-processed $p$-value. Results of probe hybridizations, which are base-2 log-transformed normalized ratios of testis/spermatozoa intensities, are plotted against their respective probabilities. Thus, left panel presents transcripts more abundant in spermatozoa whereas right panel are highly represented clones in testis. A) is hybridization result of hot-TRIzol from the bottom 90% sample when “generous” option of the “Channel Tolerance” parameter was applied which engage that a gene value was not discarded because the intensity was too low for one or the other channel. B) represents the same sample with the stringent analysis (both channels passed). Significant microspots are presented as red dots in the Volcano graph indicating a $P$ value $< 0.001$, which corresponds to “$-\log_{10}(P \text{ value}) > 3$”.