

Additional data file 3. Distribution of BLAST bit-score for the second best hit of the 60-mer oligonucleotide probes from the microarray. Each probe sequence was used as query in a BLAST search against all human genome sequences for which there is any evidence of transcription. For each probe, the bit-score result for the second best hit was recorded. Bit-score 42.1 was used as a high-end cutoff for the custom-designed intronic probes selected for our arrays.

