

Additional file 1: Gene expression profiling analysis.

Material and Methods. 21 MGF and 12 MGF receptor genes (Tables 1 and 2) could not be studied owing to a lack of probe set (one gene: Wnt3A) or to a non-informative probe set. We considered a probe set to be non informative when it was characterized by an absent call in more than 95% of the samples studied here and more than 95% of a panel of 72 human normal tissues samples available from Hogenesch's group in a public database [1] and human embryonic stem cell samples [2] analysed with Affymetrix HG-U133A. The same global scaling and normalization was used for these and our data. The call (present or absent) is determined by Affymetrix GCOS software and indicates whether a gene is reliably expressed [3].

References.

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