Legend to Supplementary Fig. 2:

Sensitivity and 1 – specificity by ROC curves for *SIM2-s*, *SIM2-l* and *SIM2-total* mRNA relative expression values. Optimal cut-off points for classifying samples as malignant were 2.0, 8.0 and 3.2 for *SIM2-s*, *SIM2-l* and *SIM2-total*. The sensitivity and specificity for these cut-off points were 0.95 and 0.62 for *SIM2-s*, 0.86 and 0.72 for *SIM2-l*, and 0.89 and 0.69 for *SIM2-total*, respectively.