Supplementary Figure 4: Bisulfite sequencing of the HES5 promoter in 39 matched Tumour-Normal pairs. Each pair of plots shows benign (above) and tumour (below) for a single patient. Methylation percentage (y-axis, fixed 0-50% scale) is shown for each of 60 CpGs assayed (x-axis). Red box highlights the single sample pair without increased methylation in tumour vs benign.