



Additional file 6: A comparison of breakpoints by 1Mb array painting and solexa sequencing data.

Hybridisation of flow-sorted chromosome peak G (Howarth et al 2008) (blue diamonds) to a 1Mb BAC array is shown compared to copy number data from solexa sequencing (red diamonds). Copy number data is shown for the region spanning 27-40 Mb on chromosome 1 (Hg19). The pattern observed is strikingly similar and allows breakpoints to be refined to around 70kb resolution.