

Supplementary Figure 1. CNVs identified in 13 DNA samples replicated between independent SNP 6.0 arrays. Frequency analysis for CNVs was performed using Nexus Copy Number™ 5.1 Discovery Edition (BioDiscovery Inc.). Green shows the frequency of amplification and red shows the frequency of deletion across 13 DNA samples. Batch 1 and 2 show data for 13 DNA samples, replicated by SNP 6.0 array.

Supplementary Figure 2. Examples of common and sample specific amplification in tumor and plasma of patients on follow-up. (a) patient 13, showing amplification at 7q11.23 in tumor only, (b) patient 19, (c) patient 25 and (d) patient 55 showing common amplification at 7q11.23 in tumor and paired plasma and other sample specific amplification peaks, (e) patient 48, common amplification at 7q11.23 in tumor and the first plasma sample, a peak in the first plasma only at 7q35 and no amplification in the second plasma (f) patient 47 showing common amplification at 10q11. in tumor and paired plasma (2 peaks) and other sample specific amplification peaks, Top to bottom: L, normal leucocyte DNA, P1, first plasma DNA sample, P2, second plasma DNA sample, T, FFPE tumor DNA.