



Supplement Figure 1: PCR confirmation of selected fusion candidates from breast tumors. We selected for validation two redundant transcripts (ARL3_MTF2 and IGLL_LOC966610), a transcript that is expressed as two isoforms (GEMIN7_SCL39A14 isoforms 1 and 2), and five non-redundant, private fusion transcripts. The tumors in which these were validated are identified above each lane.