**BWS**

Loss of methylation at KCNQ1OT1 TSS DMR (~50%)

Paternal UPD 11 (~20%)

Unknown cause (~15%)

Chromosomal aberrations within 11p (~1%) Gain of methylation at H19 TSS DMR (~5%) Mutation in CDKN1C (~10%)

**Figure 1. (Epi)genetic aberrations in BWS.**