

Description of Additional Supplementary Files

File Name: Supplementary Data 1

Description: ChIP-chip data set for *S.coelicolor* BldC. Each row represents an enriched probe (probeID) with the mid-position (midpos) of each probe on the *S. coelicolor* genome recorded. Enrichment ratios for the wild type, relative to the *bldC* mutant are expressed as the log-fold change (logFC). Probes are listed in order of significance (adjusted p value - adj.P.Val). For each probe, the nearest gene to the left and right is identified (left/rightLocusTag), its distance to the midpos of the probe (left/rightDistance), whether the gene is on the forward (1) or reverse (-1) strand (inStrand) and the predicted function (left/rightProduct) based on annotation in strepdb (<http://strepdb.streptomyces.org.uk>). If the midpos of a probe falls within a gene, it's gene identifier (inLocusTag), distance to the probe (inDistance), whether the gene is on the forward (1) or reverse (-1) strand (inStrand) and predicted function (inProduct) is also listed.