



Supplementary figure 3. Distribution of promoter powers measured as base 2 logarithm of transcription start site coverage in 5'-end enriched RNA sequencing (Mazin et al, NAR 2014). The promoters taken for the analysis are full consensus promoters as the promoter of *hsd* operon (TRTGNTAWAATN₆R). Arrow indicates measured power of HsdC promoter, which is significantly lower than it would be expected. Raw data is shown on Supplementary table 2.