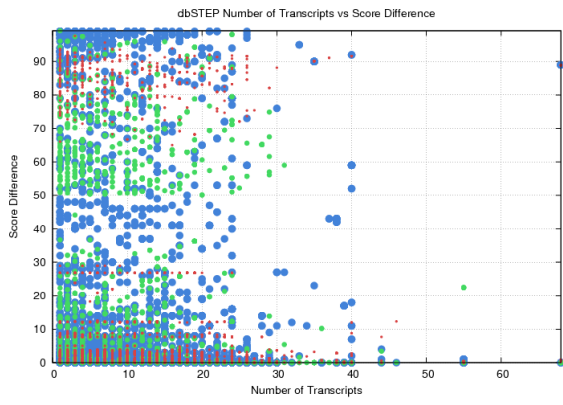
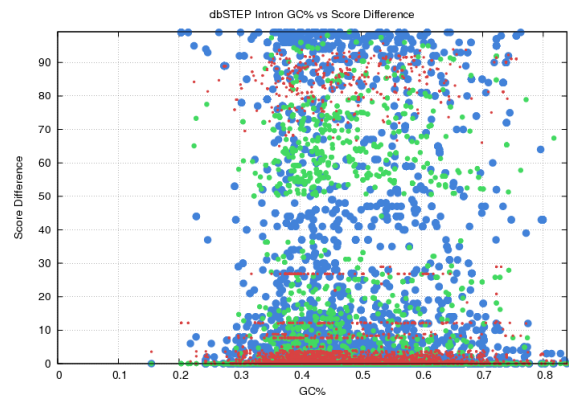


**Figure S1:** **A**, **B** and **C** show plots of allele difference scores from *in silico* prediction tools against 3 different attributes which may relate to STEPs: Number of transcripts (**A**), GC-richness (**B**) and intron length (**C**). Blue points are NNSplice results, green points are NetUTR results and red points are HSF results.

**A**



**B**



**C**

