Functional and non-functional enhancers: Binding site densities by transcription factor

- **BCD sites per 1kb**
  - Entire genome (non-coding)
- **CAD sites per 1kb**
- **HB sites per 1kb**
- **KR sites per 1kb**
- **KNI sites per 1kb**
- **BCD sites per 1kb**
  - Aligned BCD sites per 1kb
  - Aligned + Preserved BCD sites per 1kb
- **CAD sites per 1kb**
  - Aligned CAD sites per 1kb
  - Aligned + Preserved CAD sites per 1kb
- **HB sites per 1kb**
  - Aligned HB sites per 1kb
  - Aligned + Preserved HB sites per 1kb
- **KR sites per 1kb**
  - Aligned KR sites per 1kb
  - Aligned + Preserved KR sites per 1kb
- **KNI sites per 1kb**
  - Aligned KNI sites per 1kb
  - Aligned + Preserved KNI sites per 1kb

**Legend:**
- Green: True CRMs
- Red: Non-functional pCRMs
- Blue: Ambiguous expression pCRMs

Additional File 01  Berman, Pfeiffer et al. Genome Biol. 2004