Supplementary File 1. ChIP-seq statistics. Related to Figure 2.

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| --- | --- | --- | --- | --- |
| Target | Stage (hpf) | Replicates | Replicate overlap | FE≥10 peaks |
| Prep | 3.5 | 2 | 95.6% | 13,328 |
| Prep | 12 | 2 | 98.8% | 24,199 |

Prep ChIP-seq experiments were performed on 3.5hpf and 12hpf zebrafish embryos. Biological replicates (n=2) for each condition show a high degree of reproducibility as shown by the percent of peaks found in both replicates (replicate overlap). Peaks with Fold Enrichment (FE)≥10 were used for subsequent analysis.