



- MARCS: Strongly recruited to H3K4me3
- MARCS: Strongly excluded from H3K4me3
- Histones
- Other
- (border) ChIP-MS no significant difference me3 vs me1
- (border) ChIP-MS significant difference me3 vs me1
- (size) Estimate imputed or based on 1 datapoint
- ◀ (shape) Out of bounds