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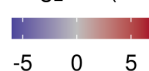
### More accessible regions in adult spermatogonia

TF	Enriched motif	% of target sequences
JUND		24.35%
FOS		23.98%
ATF3		25.94%
RFX2		30.71%
USF1		23.00%
POU3F1		9.01%

### Less accessible regions in adult spermatogonia

TF	Enriched motif	% of target sequences
EHF		37.63%
FOXO3		36.71%
FOXA1		27.63%
FOXL2		35.53%
GABPA		45.13%
ETV4		38.55%

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Log<sub>2</sub> FC (RNA-seq)-Log<sub>10</sub> adjusted *P* (Motif enrichment)