Figure legend (right side):

Predictor

- ABC (A=scATAC, C=power law) (K562)
- Cicero (co–accessibility) (K562)
- FigR (p–value) (K562)
- In element (ABC (A=scATAC)) & distance to TSS (K562)
- Kendall correlation (access. at E x expr. at G) (K562)
- scE2G (Multiome) (K562)
- scE2G (scATAC) (K562)
- SCENIC+ (importance) (K562)
- SCENT (FDR) (K562)
- SnapATAC (p–value) (K562)
- STARE (gABC, power law) (K562)
- ArchR (high–confidence) (K562)
- DIRECTNET (high–confidence) (K562)

Axis labels:

Y-axis: Enrichment (eQTLs versus common variants)

X-axis: Recall (variants overlapping prediction linked to eGene)
618 variants