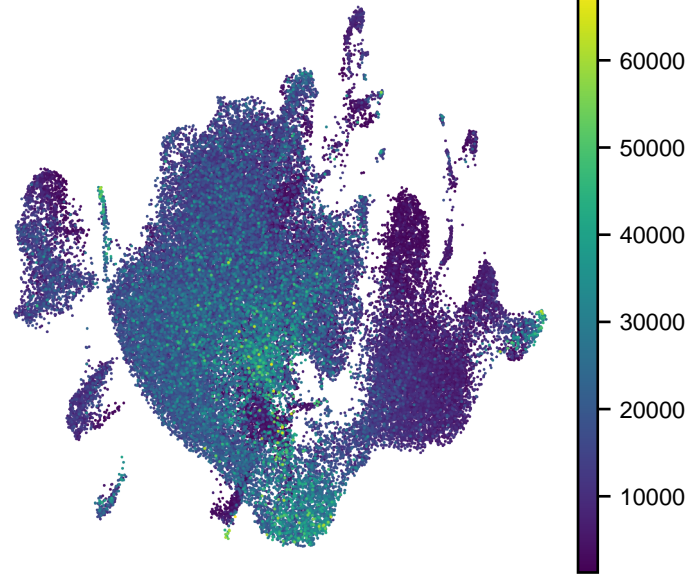


batch

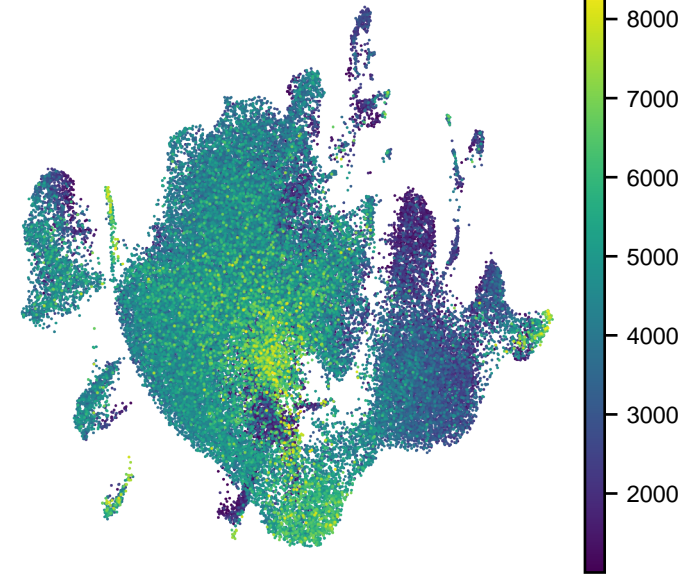


- Aire\_WT-1
- Aire\_WT-2
- Fezf2\_WT-1
- Fezf2\_WT-2
- Aire\_KO-1
- Aire\_KO-2
- Fezf2\_KO-1
- Fezf2\_KO-2

total\_counts



n\_genes\_by\_counts



pct\_counts\_mt

