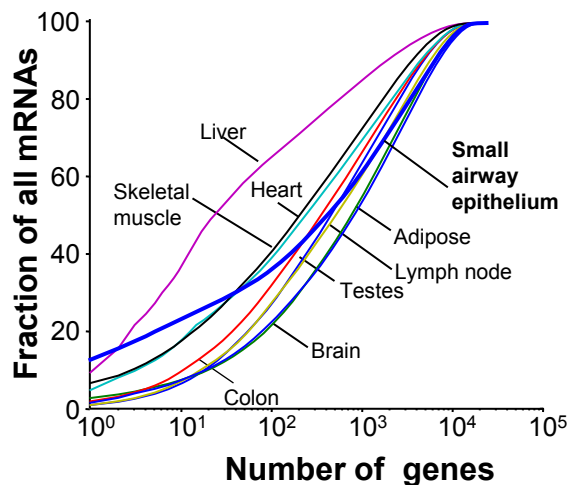
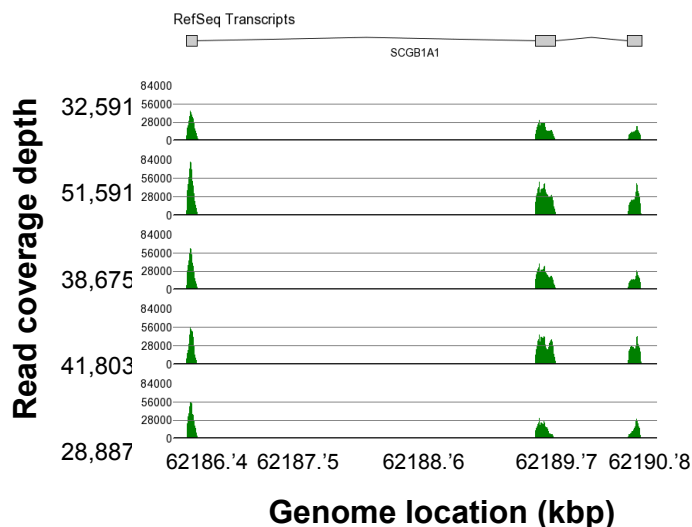


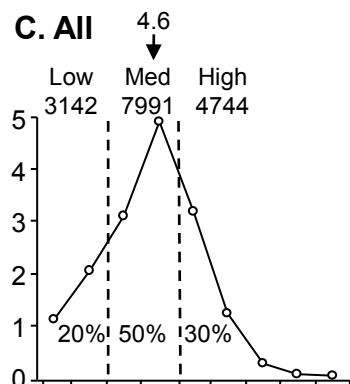
A. Transcript abundance



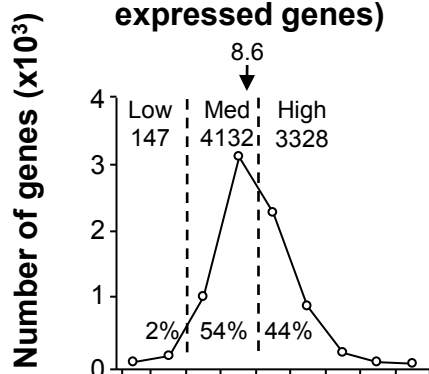
B. SCGB1A1



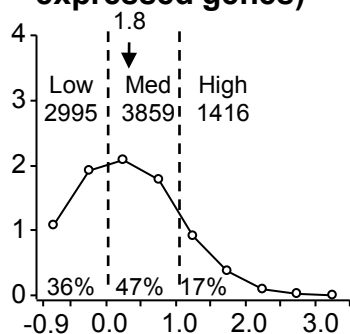
C. All



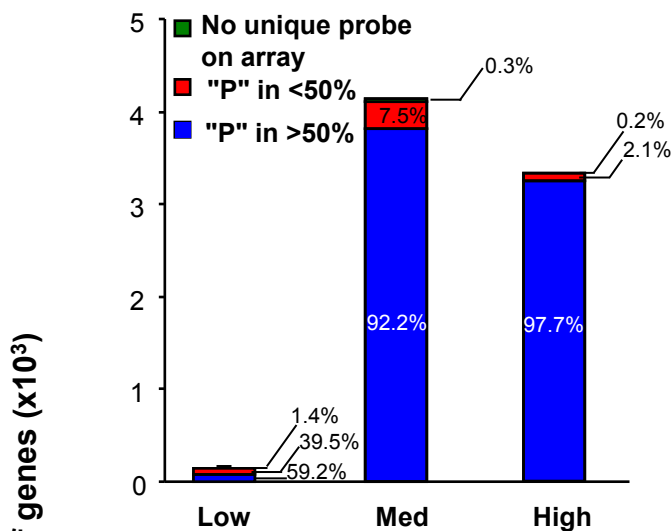
D. Ubiquitous (48% of all expressed genes)



E. SAE enriched (52% of all expressed genes)



F. Ubiquitous



G. SAE-enriched

