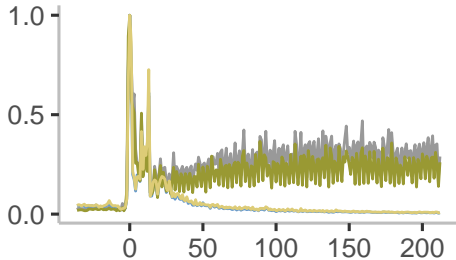


Normalized read count



P-site position w.r.t start codon

sample_name

- sgfluc_dms
- sgfluc_hht
- sggn1_dms
- sggn1_hht