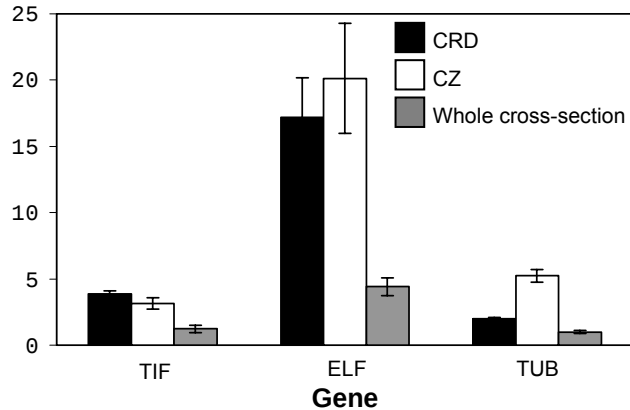


A**Relative transcript abundance****B****Relative transcript abundance**