Figure 3. Sperm production rates for different *C. elegans* male genetic backgrounds. (A) Box-plots for spermatid counts for each strain at time zero (left) and two hours later (right), indicating median and inter-quartile range (points beyond whiskers are candidate outlier values). (B) Estimates of sperm production rate for each strain, based on the linear regression of sperm counts between the two time periods. Error bars in (B) indicate 1 S.E. for the slope of the regression line estimates of sperm production rate.