

Additional file 2. General statistics for the sequencing of triplicate floral sRNA libraries originated from apomictic and sexual *Paspalum notatum* genotypes.

Sample Name	Total number of reads	% duplicate reads	Sequence length	% GC
Apo1_S1	2,996,675	74.3%	50	54%
Apo2_S2	1,981,542	70.8%	50	55%
Apo3_S3	1,558,547	69.5%	50	54%
Sex1_S4	1,986,075	71.3%	50	55%
Sex2_S5	2,032,618	72.5%	50	55%
Sex3_S6	2,248,170	72.5%	50	55%