

sample	library	# reads	# nucleotides	N50	Max length	Mean length	Median length	Mean Q	Median Q
AQ0090	9867AE	2,131,636	1277968393	717	16571	599.52	483	6.29	6.42
AQ0091	9928AE	2,175,133	1281280919	697	20932	589.06	479	6.22	6.36
AQ0092	0318AF	637,670	437368503	843	20013	685.89	537	6.33	6.48
AQ0093	0827AF	2,353,202	1476728326	754	26246	627.54	499	6.18	6.32

Table S1. Statistics dRNA sequencing of *S. pombe*. Approximately 600 ngr polyA+ RNA in 10ul volume per run were employed. The samples were run in an ONT gridion X4 at the Centro Nacional de Análisis Genómicos (CNAG, Barcelona). Q: quality score.