

Supplementary Table 1 Native RNA nanopore sequencing statistics

Organism	sample	"Click" type	pass reads	N50	adapted
<i>S. cerevisiae</i>	Untreated 1	None	1,391,578	531	0
<i>S. cerevisiae</i>	Untreated 2	None	1,755,395	1,092	0
<i>S. cerevisiae</i>	Untreated 3	None	3,222,192	1,034	0
<i>S. cerevisiae</i>	Untreated 4	None	861,210	947	0
<i>S. cerevisiae</i>	Untreated Pooled	None	7,230,375	957	0
<i>S. cerevisiae</i>	Treated 1	Cu ²⁺	2,576,122	798	0.2382
<i>S. cerevisiae</i>	Treated 2	Cu ²⁺	3,933,777	676	0.0889
<i>S. cerevisiae</i>	Treated 3	Cu ²⁺	1,848,281	722	0.1451
<i>S. cerevisiae</i>	Treated 4	Cu ²⁺	1,382,164	575	0.0696
<i>S. cerevisiae</i>	Treated 5	Cu ²⁺	600,225	435	0.0618
<i>S. cerevisiae</i>	Treated Pooled	Cu²⁺	10,340,569	692	0.134
<i>S. cerevisiae</i>	Treated 1	Cu-Free	1,128,595	737	0.3347
<i>S. cerevisiae</i>	Treated 2	Cu-Free	3,799,042	755	0.4138
<i>S. cerevisiae</i>	Treated 3	Cu-Free	1,275,112	715	0.3407
<i>S. cerevisiae</i>	Treated Pooled	Cu-Free	6,202,749	744	0.3844
GM12878	Untreated 1	None	1,332,182	1,572	0
GM12879	Untreated 2	None	2,497,808	1,691	0
GM12880	Untreated Pooled	None	3,829,990	1,615	0
GM12881	Treated 1	Cu-Free	640,221	1,334	0.1106
GM12882	Treated 2	Cu-Free	919,701	1,131	0.1021
GM12883	Treated 3	Cu-Free	797,451	1,212	0.1596
GM12884	Treated 4	Cu-Free	1,706,866	1,189	0.1653
GM12885	Treated Pooled	Cu-Free	4,064,239	1,207	0.1413