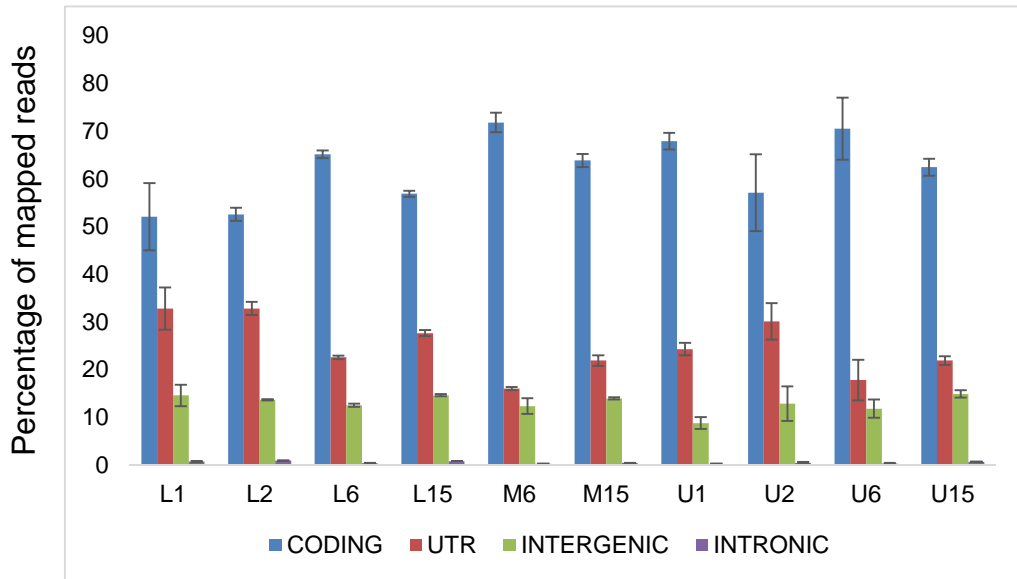


A



B

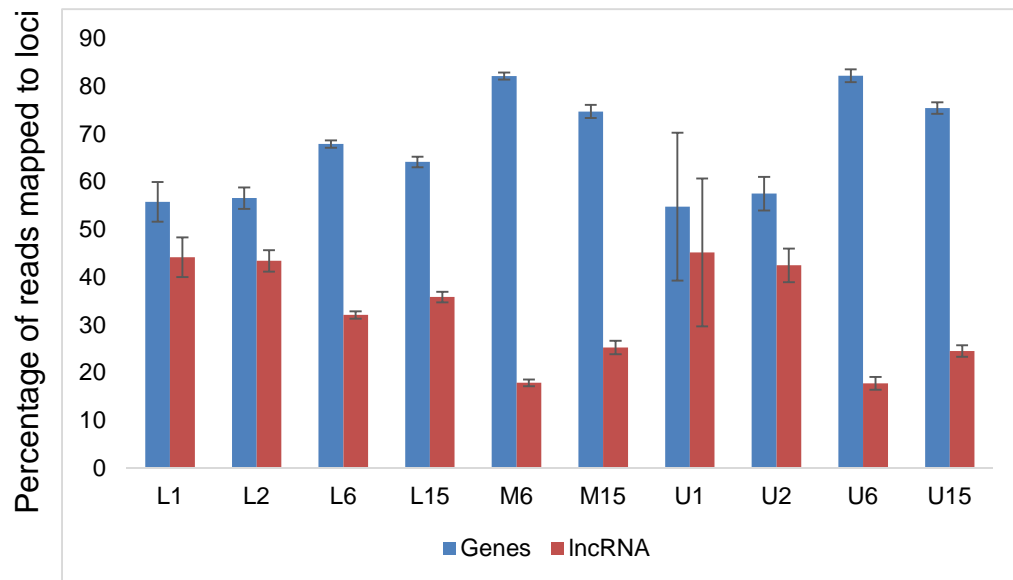


Figure S1

Read mapping. For each sample, the percentages of reads, mapping to different genomic features (A) and the percentages of locus-mapped reads, mapping to genes and to lncRNA (B), were calculated and plotted as bar charts.

Table S1: List of the primers

RDN25-1-int-for	GAGGAAAAGAAACCAACCG
RDN25-1-int2-rev	AGGCACTTTACAAAGAACCG
RPB2-int-for	GAAGACGATGAATCACTAGG
RPB2-int-rev	TCAGCATCAATGTATTCCAC
COG1-int-for	AGTTACTTCCACTACTCCG
COG1-int-rev	TGTCTTCATAGAGTCCACC
TAF6-int-for	ATTAGAGGTGCTATTGTCAC
TAF6-int-rev	CACTAAAGGTTTGACTTCAG
RAD52-int-for	AGAACAACAACAATACCCG
RAD52-int-rev	TACTGTTAGCCTCAGCAG
NCE102-int-for	CATCGGTTTAATCAGTTCG
NCE102-int2-rev	CCAATGGTTCAATGAAGTTG
VBA5-int-for	AGGAGATAGAAGGAGCATG
VBA5-int-rev	ACTGTAGCCTGTAACGAG
YDL218w-int-for	GAATGTACTTCTGTGCCG
YDL218w-int-rev	TACATGCTCTTCTATGACTG
MIG2-int-for	GATACACCAACCAATACCG
MIG2-int-rev	ATTGGCATAGGAGAAGTTG
CYC1-int-for	ATGTCTACAATGCCACACC
CYC1-int-rev	CTTCAACCCACCAAAGGC
CYC7-int-for	GTGTCAGCAGTGCATAC
CYC7-int-rev	CTTCTTCAACCCGGCAAAC
HSP30-int-for	GGTGGACTGGTGTTC AAG
HSP30-int-rev	CACGATTTGAATTAACAGCG
CTS1-int-for	ACTCAGCAGTCGTTGATG
CTS1-int-rev	TCCAACAAGTCACCAACAG
THI5-int-for	TACGGTATGAAGCCAGAAG
THI5-int-rev	AACGGTACAGAAGCAACAG
VHT1-int-for	CTGGACTTGGTTCACATTTG
VHT1-int-rev	CAGAAAGTCCTGAAGTCACG
INO1-int-for	CCCCTGTTCAAGACTACG
INO1-int-rev	TTAACGCCTTCCTTAGTTTG
ZRT1-int-for	TGGTGGAACGAGTACAATG
ZRT1-int-rev	CAATCGCACCATAAGCAG
ANS1-int-for	AGTAAGCAACTCCTCAGAC
ANS1-int-rev	CAACCACCCACATCATATAC
CWP1-int-for	CGGCTCGGATTTACAATAC
CWP1-int-rev	CATCCTCATTGACAACAGC
STL1-int-for	CTTGGTAGGAACACTAGACG
STL1-int-rev	GTATATGGCAGCGTTACAAC
PDR15-int2-for	ACTCACTCGGTTGTGTATG
PDR15-int-rev	CTGGTCTCCCAAGAACAAC