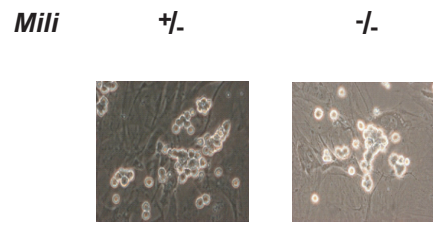
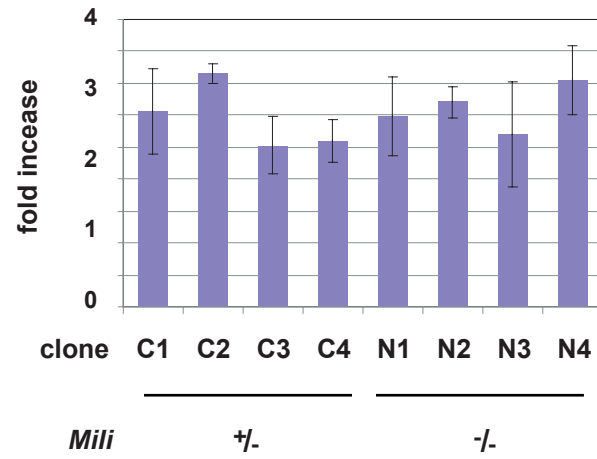
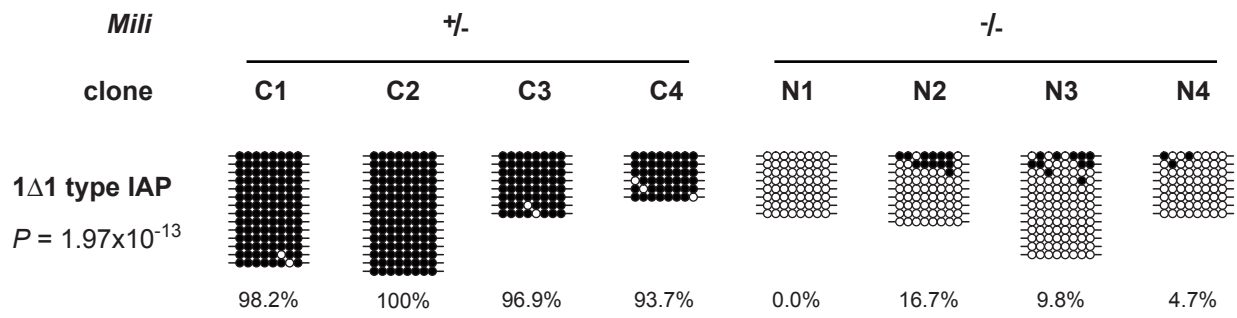
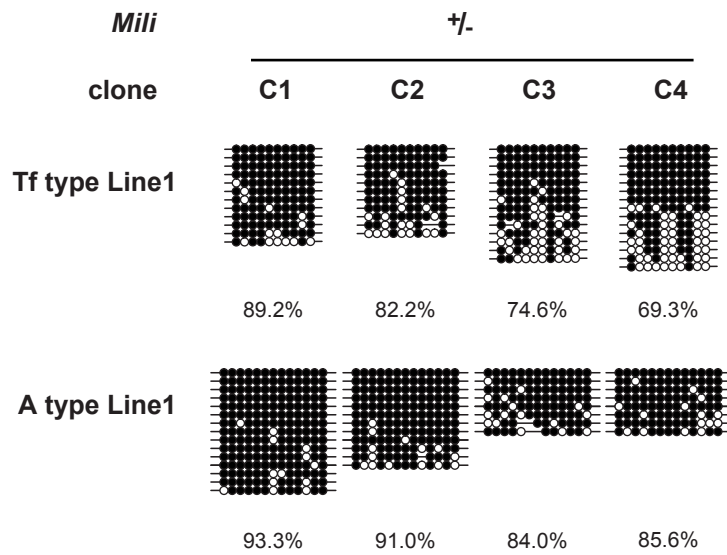


A**B** $P = 0.866$

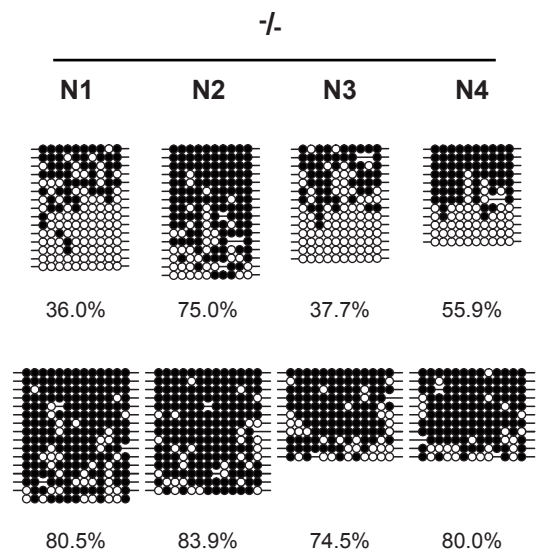
A

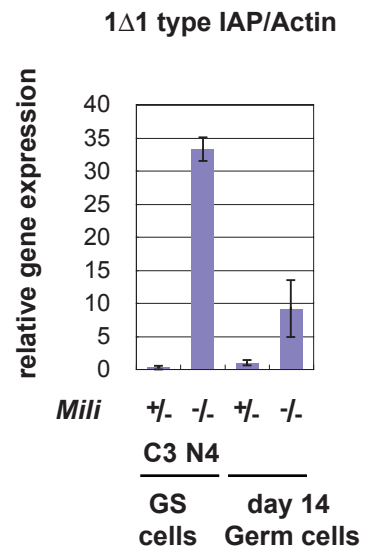


B



C

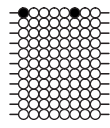






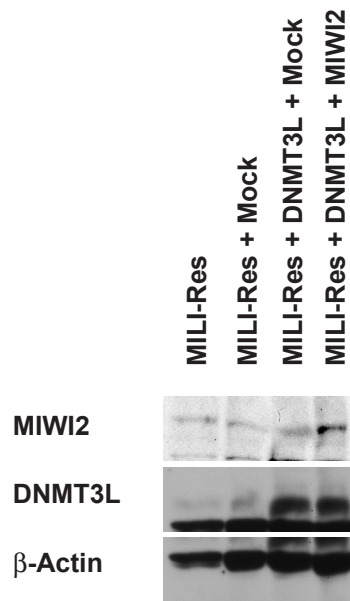
A

MILI-Res

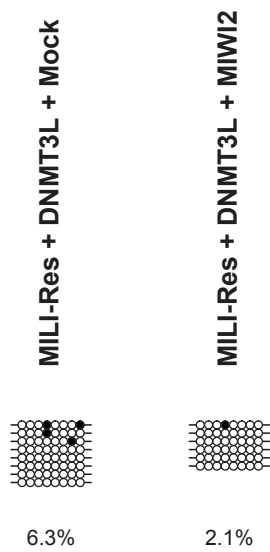


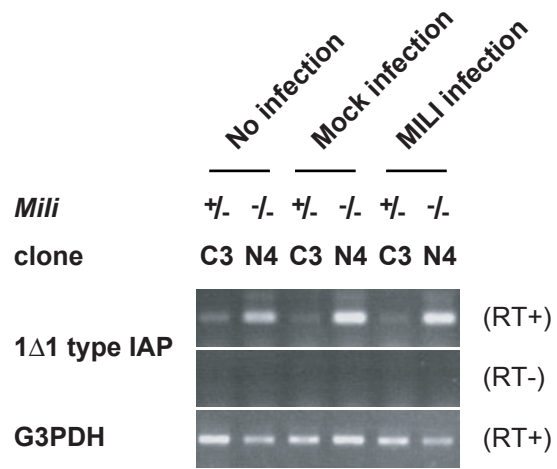
2.3%

B

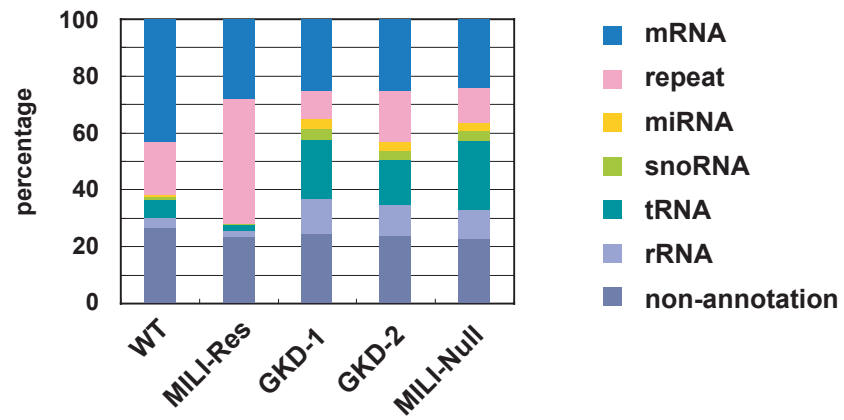


C



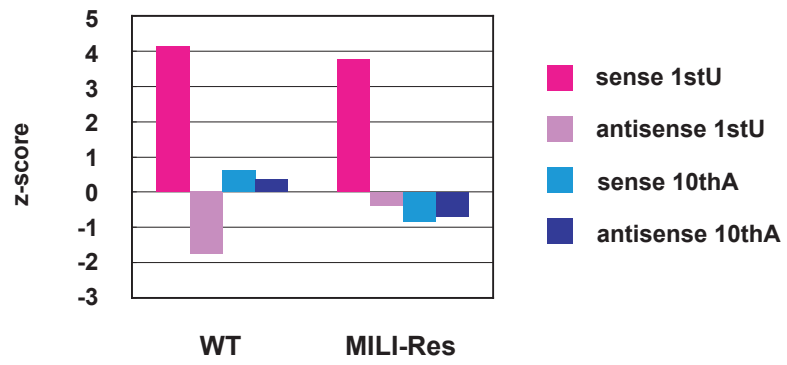


A

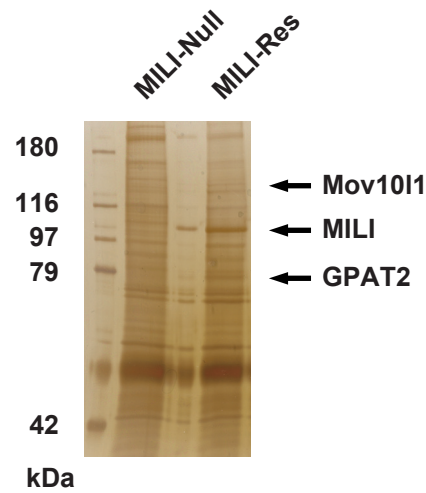


B

	WT	MILI-Res	GKD-1	GKD-2	MILI-Null
rRNA	130718	101238	160288	161183	141474
tRNA	240847	70886	270172	239171	326565
snoRNA	31311	19730	54533	50241	50492
miRNA	34413	14944	41999	45153	39613
repeat	691496	1893630	132039	265980	162519
mRNA	1611978	1211251	328322	378618	331647
non-annotation	993925	1016097	321599	358978	312048
total reads	3734688	4327776	1308952	1499324	1364358



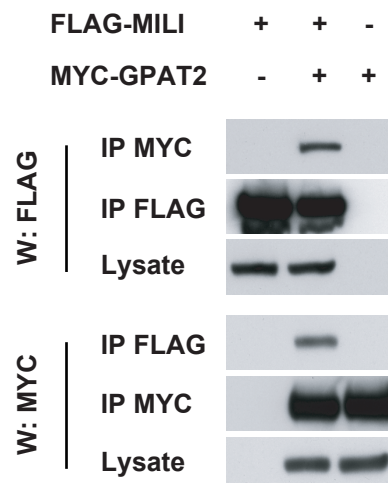
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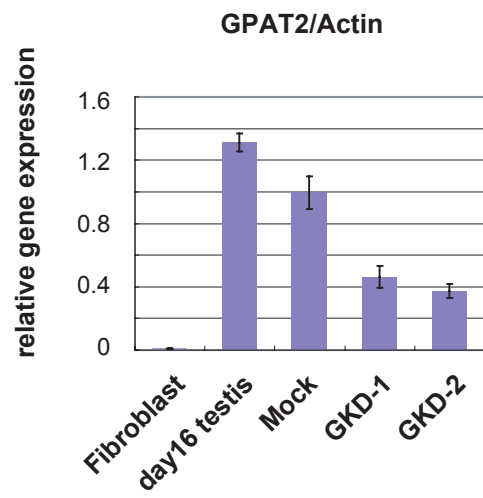
B

Band 1					
Accession	Entry	Description	mW (Da)	Peptides	Coverage (%)
IPI:IP100118099.2	IPI:IP100118099.2	Mov1011 Uncharacterized protein	138643	14	10.9766
Band 2					
Accession	Entry	Description	mW (Da)	Peptides	Coverage (%)
IPI:IP100461359.2	IPI:IP100461359.2	Piwil2 Piwi like protein 2	109417	42	38.414
Band 3					
Accession	Entry	Description	mW (Da)	Peptides	Coverage (%)
IPI:IP100889266.3	IPI:IP100889266.3	Gpat2 Isoform 2 of Glycerol 3 phosphate acyltransferase 2 mitochondr	88658	13	20.7026

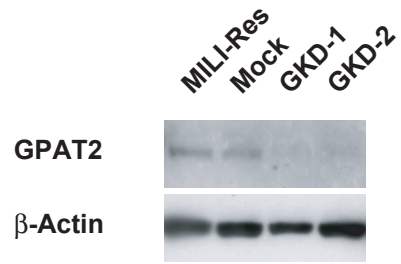
C



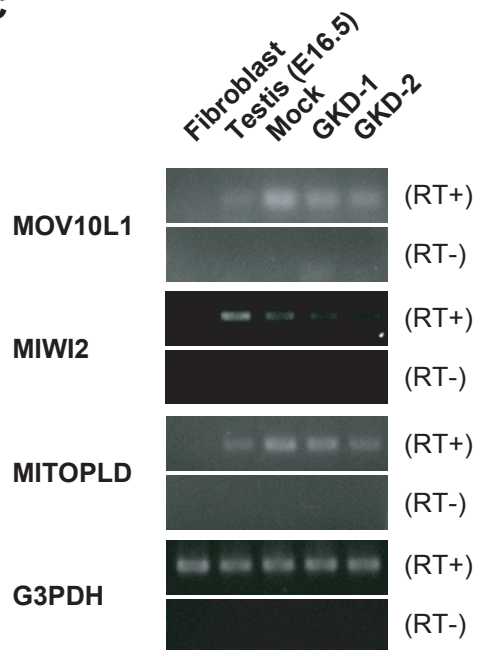
A



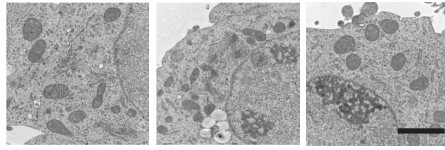
B



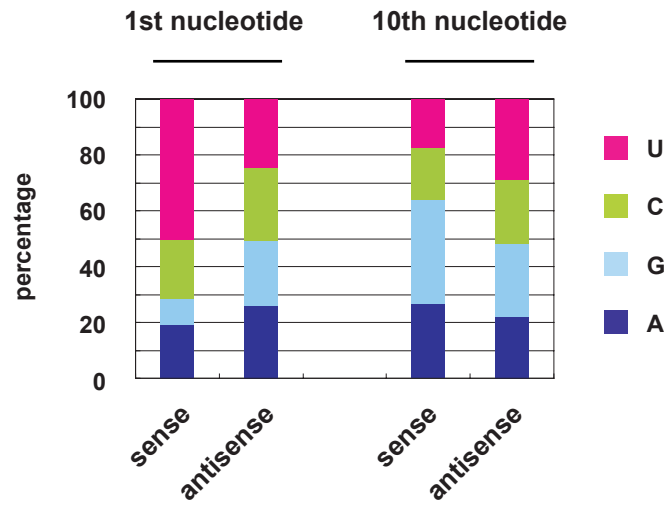
C



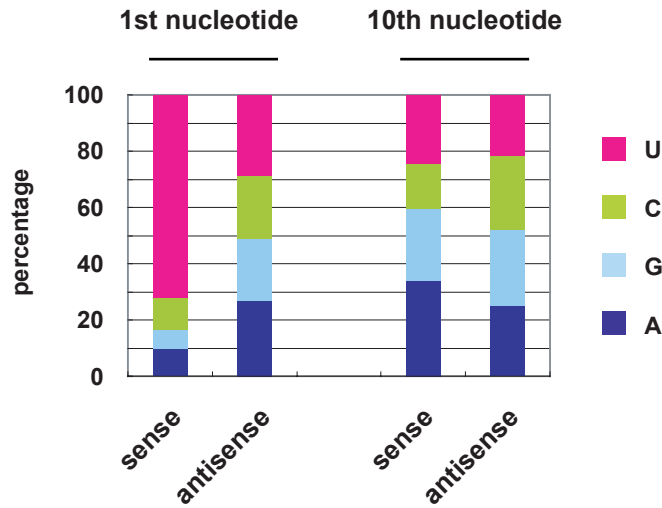
GKD-1 GKD-2 Mock



A



B



C

