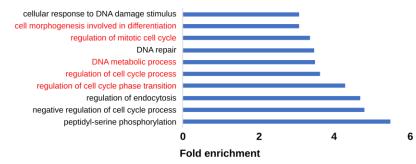


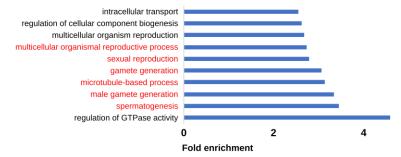
b

GO term analysis of MassSpec identified Round_WT circRNA



С

GO term analysis of MassSpec identified Elongating_WT circRNA



d

GO term analysis of MassSpec identified Sperm_WT circRNA

microtubule-based process				
male gamete generation	-			
spermatogenesis	-			1
macromolecule catabolic process	-			1
microtubule cytoskeleton organization	-			
protein modification by small protein conjugation	-			
modification-dependent protein catabolic process	-			
ubiquitin-dependent protein catabolic process				
modification-dependent macromolecule catabolic process				
cellular macromolecule catabolic process				
	0	1	2	3
Fold enrichment				

Fig.S13. GO term enrichment analyses of circRNA genes with circRNA junction peptides detected in LC-MS assays in pachytene spermatocytes (a), round (b) and elongating (c) spermatids, as well as spermatozoa (d). GO terms highly relevant to spermatogenesis are highlighted in red.