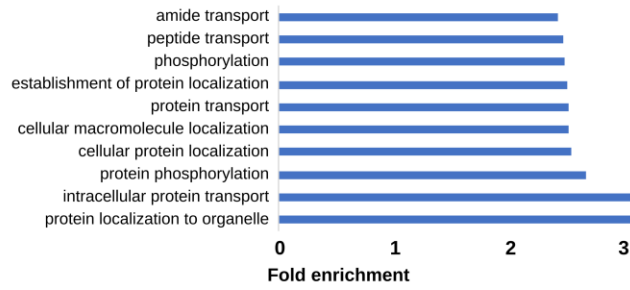


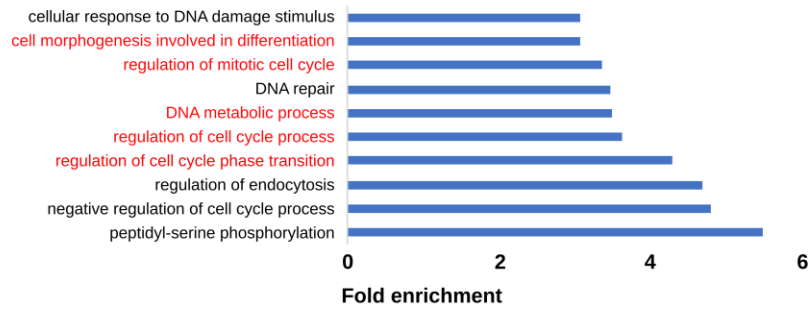
**a**

**GO term analysis of MassSpec identified Pachytene\_WT circRNA**



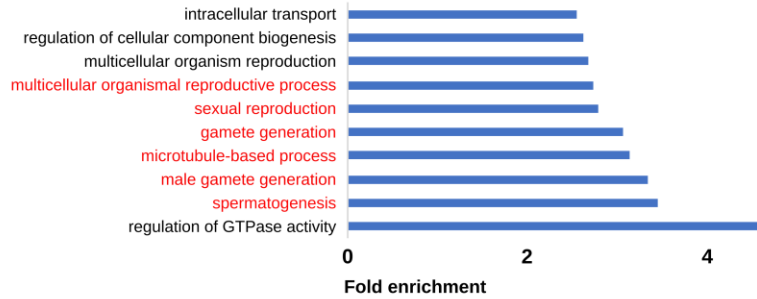
**b**

**GO term analysis of MassSpec identified Round\_WT circRNA**



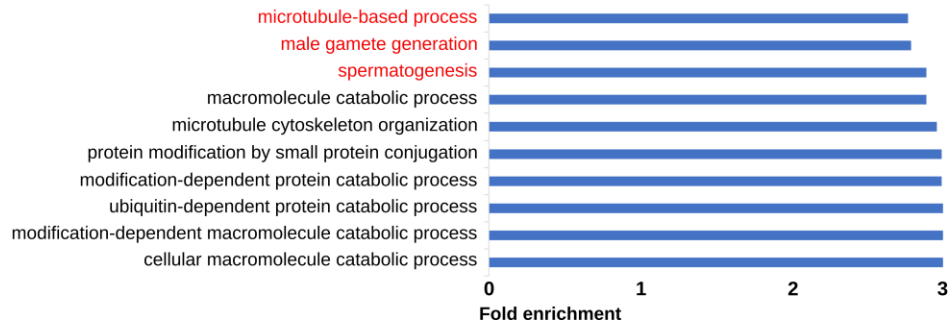
**c**

**GO term analysis of MassSpec identified Elongating\_WT circRNA**



**d**

**GO term analysis of MassSpec identified Sperm\_WT circRNA**



**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.**