The comprehensive expression analysis of circular RNAs in gastric cancer and its association with field cancerization

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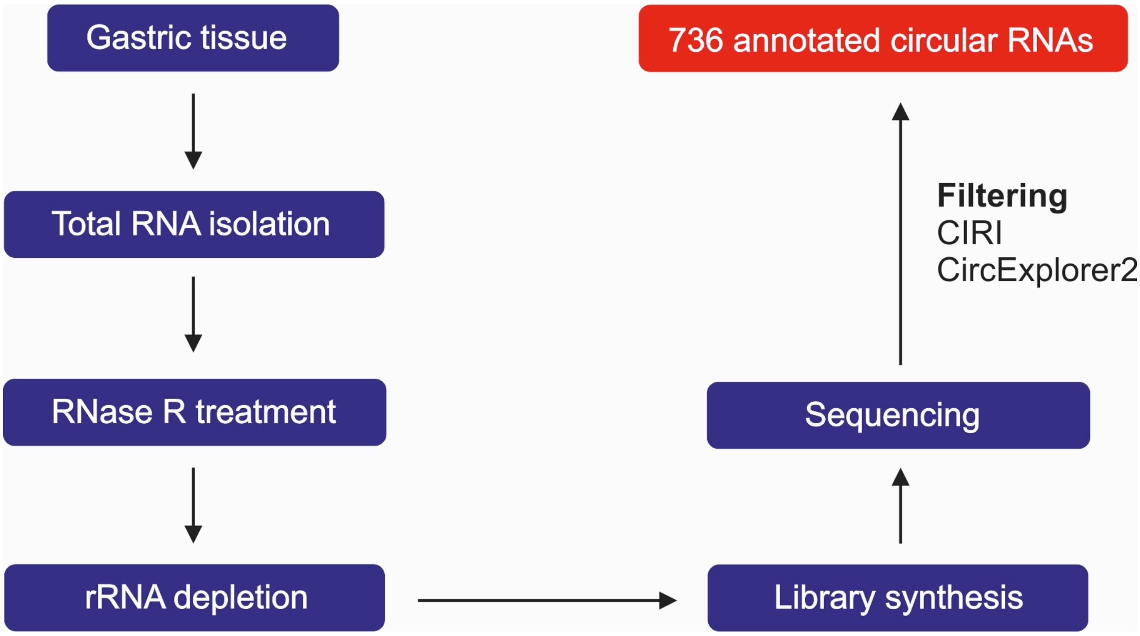
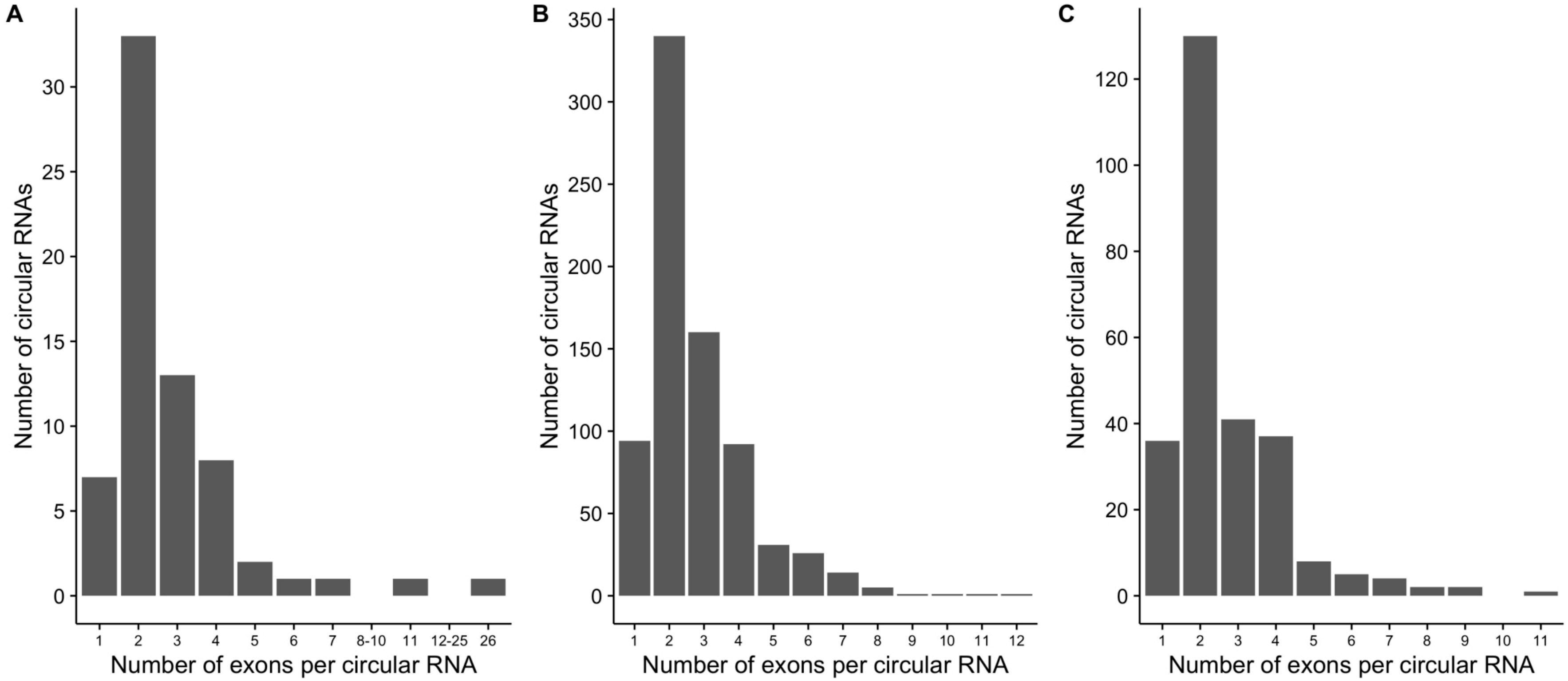
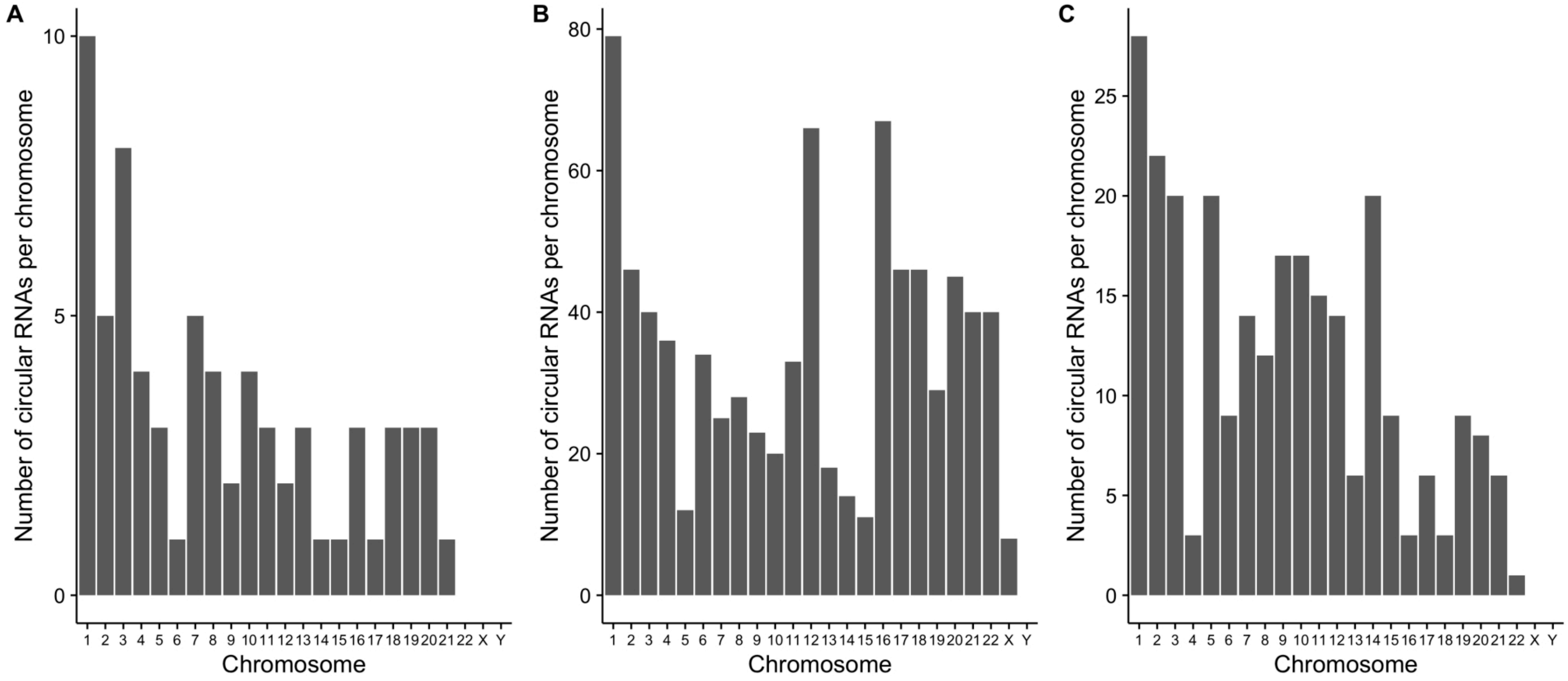
Supplementary information



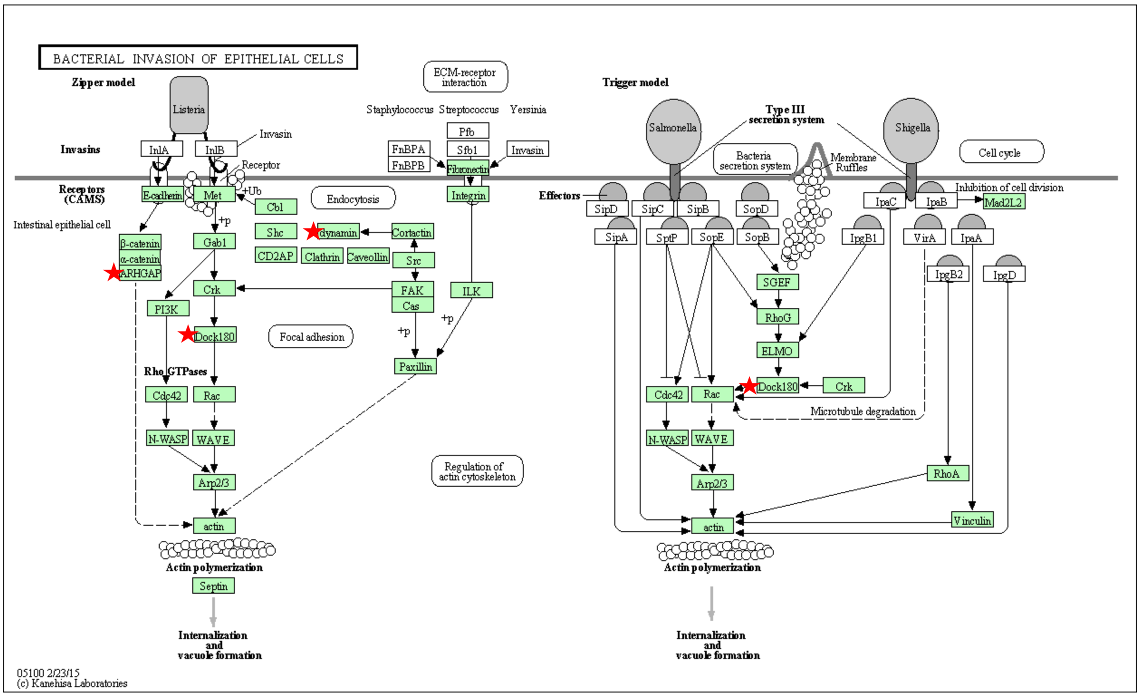
Supplementary Fig. 1 | Circular RNAs detection workflow.



Supplementary Fig. 2 | Number of exons per annotated circular RNAs in gastric tissue. (A) Gastric tissue without gastric cancer. (B) Matched tumor-adjacent gastric tissue. (C) Gastric cancer.



Supplementary Fig. 3 | Number of circular RNAs per human chromosome. (A) Gastric tissue without gastric cancer. (B) Matched tumor-adjacent gastric tissue. (C) Gastric cancer.



Supplementary Fig. 4 | Scheme of the KEGG pathway of bacterial invasion of epithelial cells[30]**.** This pathway was enriched in gastric tissue without gastric cancer and in matched adjacent tumor gastric tissue.