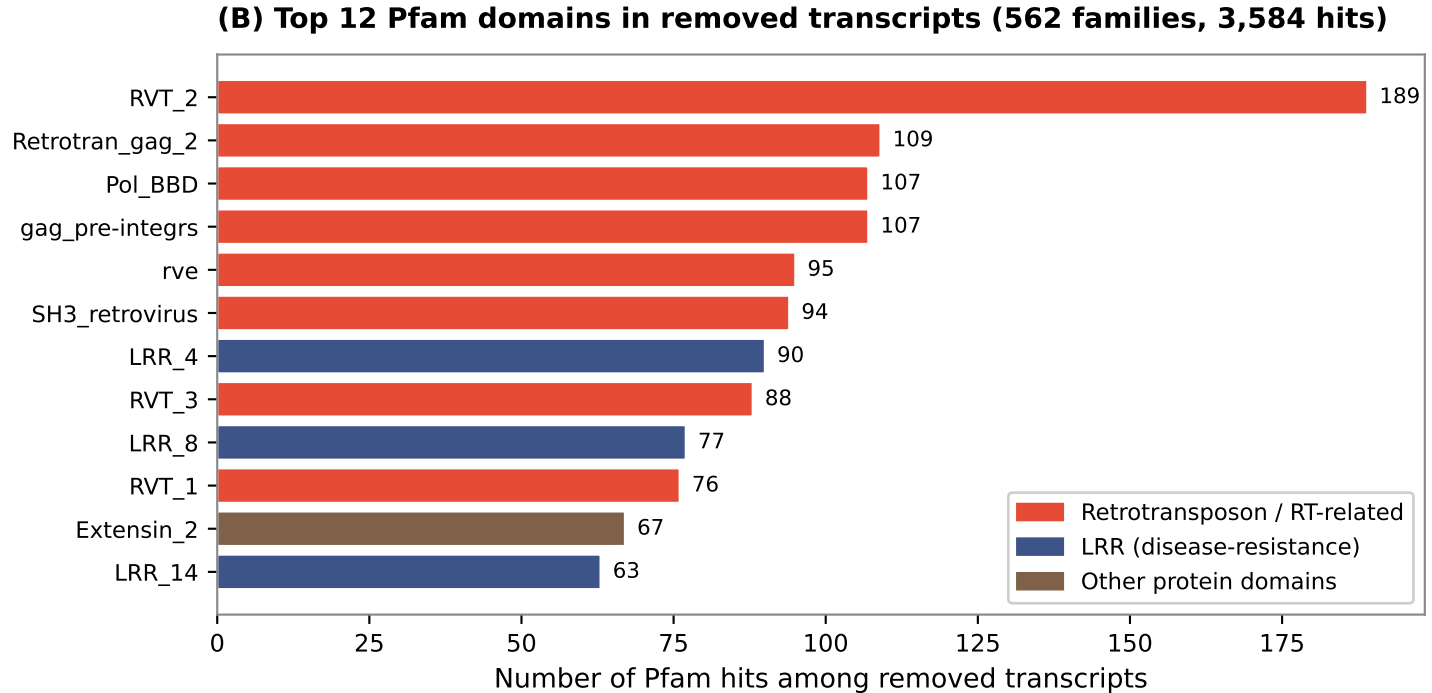
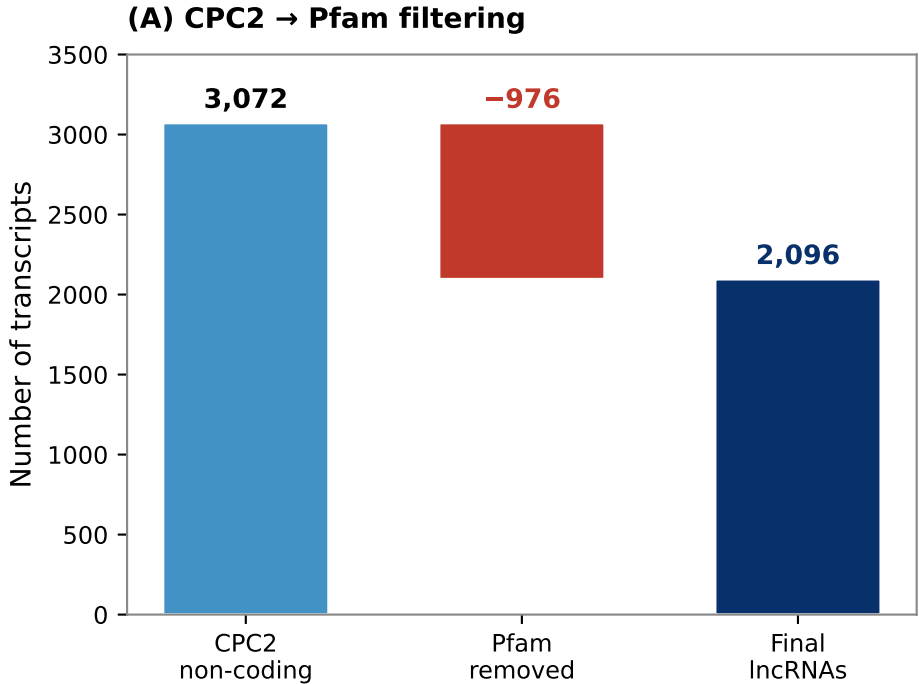


Figure S3

Coding-potential (CPC2) and protein-domain (Pfam) filtering of lncRNA candidates



CPC2 classified 3,072 candidates as non-coding; a further 976 transcripts carrying recognizable Pfam protein domains (predominantly retrotransposon-derived and LRR domains) were removed, yielding 2,096 high-confidence lncRNAs.