

Figure S3. ML-trees of known and environmental RT sequences. (A) ML-tree of known RT sequences with 124 RT-like sequences identified in the metagenomic data.

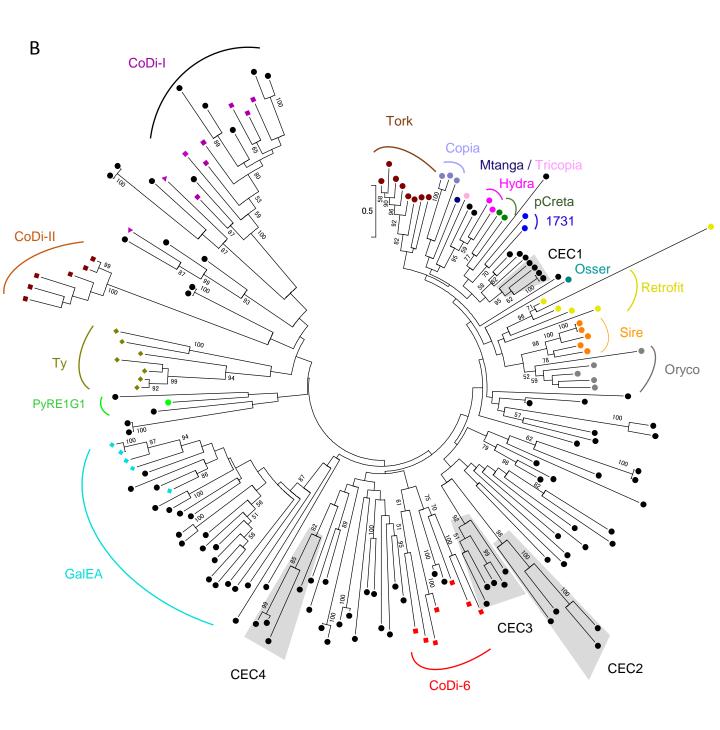


Figure S3. ML-trees of known and environmental RT sequences. (B) ML-tree of known RT sequences with 100 Copia RT-like sequences identified in the metagenomic data.

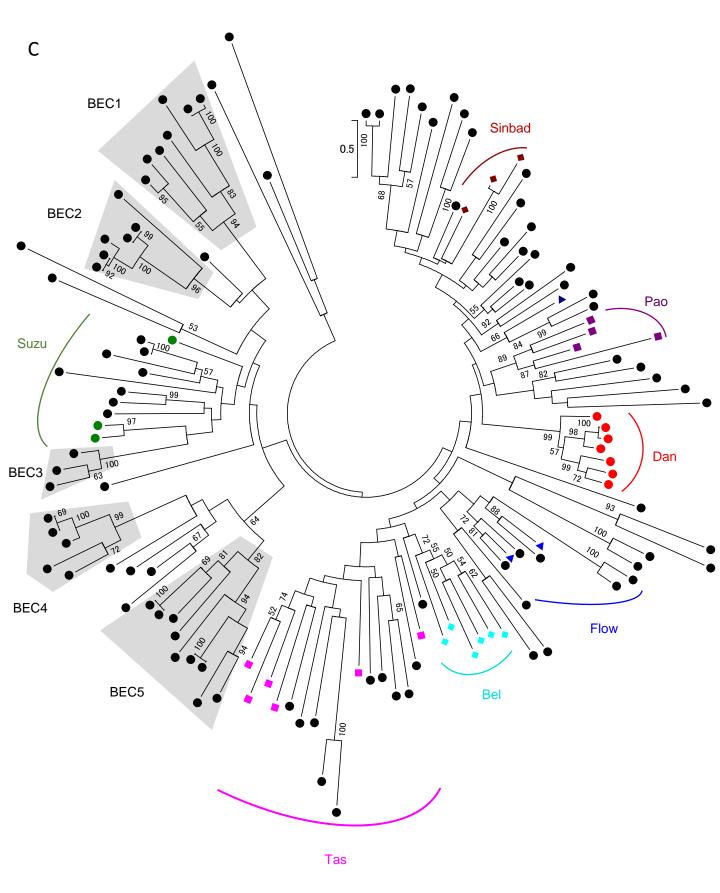


Figure S3. ML-trees of known and environmental RT sequences. (C) ML-tree of known RT sequences with 100 longest BEL RT-like sequences identified in the metagenomic data.