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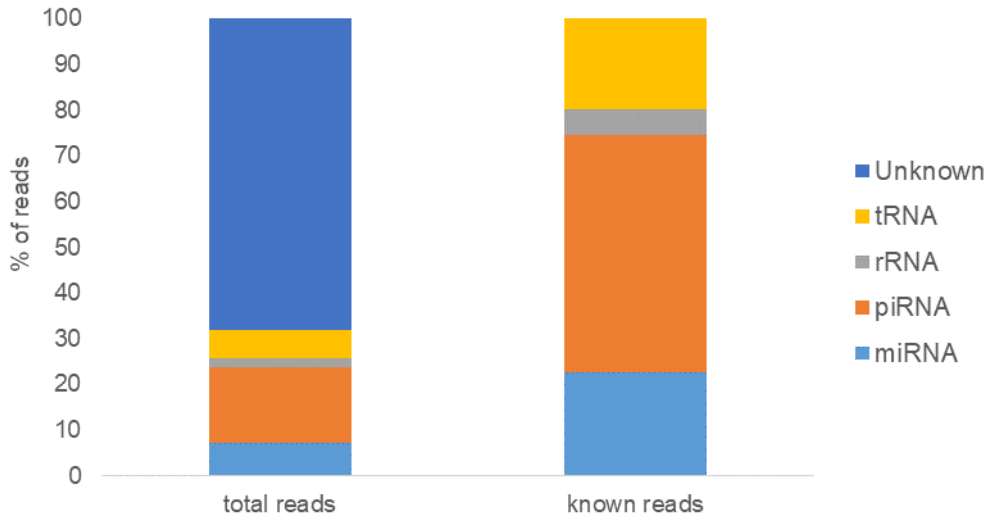


Figure S2. Percentage of mappable reads after sequencing of small RNAs from rat serum EVs. The first column represents the general profile of reads. The second column represents only the known reads mapped against *Rattus norvegicus* databases.