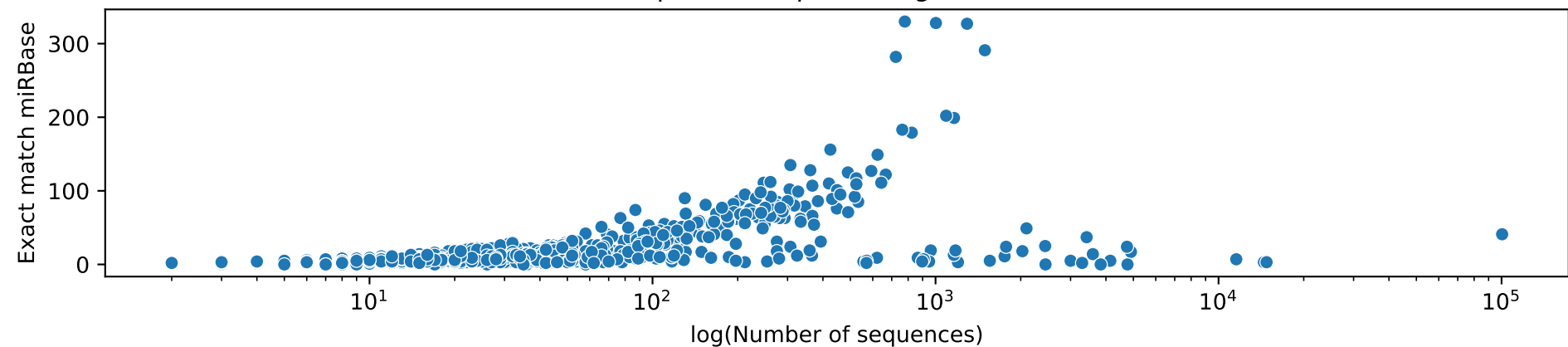


Rfam-miRBase correspondence percentage (FULL) n = 1146 (263573)



Rfam-miRBase correspondence percentage (SEED) n = 1146 (9852)

