Figure S1

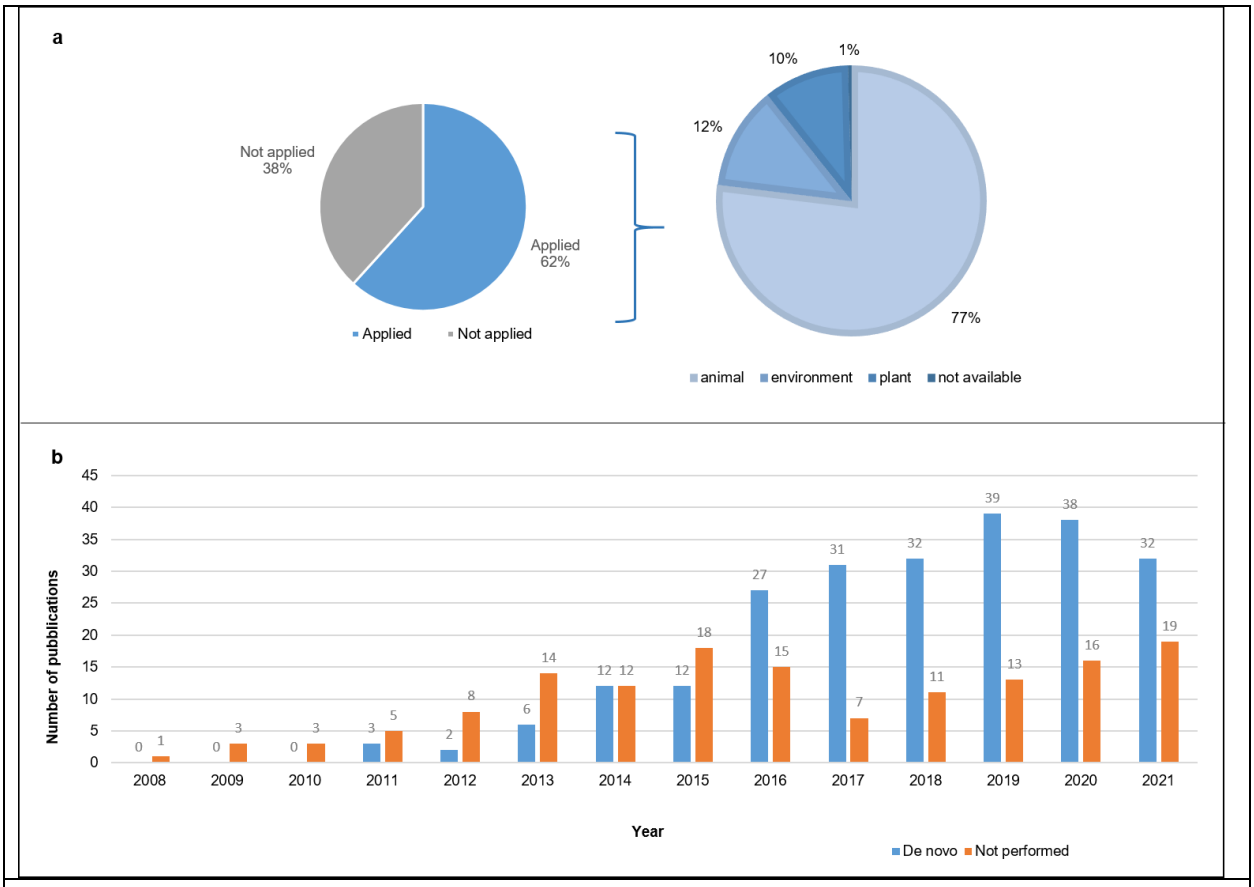

Figure S1*: Use of the " de novo Assembly". (a)* The graph shows the percentage of publications employing (62%) or not employing (38%) de novo Assembly for viral genomes identification. Among the 62% articles performing de novo Assembly, percentages for each sample type are also indicated: 77% animal, 12% environment, 10% plant. (**b**) use of de novo Assembly through the years**.** The graph shows the number of publications in which de novo Assembly is used (blue bars), compared with publications in which it is not used (orange bars), from year 2008. x-axis: year of publication, y-axis: number of publications.