**Figure 1:** The timeline for Mozambique was generated using Timeline storyteller and accessed using PowerBI software.

* For this timeline four categories were considered: pandemic milestones, reported case numbers, reported deaths, variant milestones. The data on reported case numbers, pandemic milestones and reported COVID-19 related deaths were accessed using Our World in Data (OWID), and variant milestones (illustrating the first reports of new variants within Mozambique) were accessed using GISAID (date of sampling).
* The following headings were included and saved as a comma delimited CSV file

|  |  |  |  |
| --- | --- | --- | --- |
| category | start\_date | end\_date | content\_text |

* CSV document is converted to .json format
* Data was imported into PowerBi
* Timeline Storyteller was added as a Visual
* Results were plotted using a spiral layout

**Supplementary figure S1** illustrates the workflow and overview of samples collected in Mozambique, submitted to sequencing facilities, submitted to GISAID, assemblies accessed through GISAID and used for downstream analyses. This overview was illustrated using Microsoft PowerPoint.

* The top level indicates the number of nasopharyngeal or oropharyngeal swabs collected in Mozambique,
* the second the sequencing facility that experimentally determined whole-genome sequences,
* the third indicates the number of high-quality genomes submitted to GISAID,
* the fourth level (bottom) illustrates the total number of Mozambican SARS-CoV-2 genome sequences used for analyses.