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# Descargar Genexus X Evolution 2 Full Extra Quality 18

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the new version, x evolution 2 has been reengineered from the ground up to provide unprecedented sequencing flexibility, a dramatically enhanced workflow, new features and improvements in every area of the platform. the x evolution 2 platform has undergone extensive testing to ensure its reliability and robustness in clinical application.

the new system is based on a new generation of semiconductor, leading to more than a doubling in sequence length and a major reduction in raw sequence time. in addition, x evolution 2 reduces the time from raw data to finished sequences by over a factor of three. x evolution 2 also includes a new workflow that allows users to choose from among four different sequencing paths, and will be available as a kit, a cartridge, or the combination of both. we present the results of an evaluation of the fully automated sequencer genexus x evolution 2.

for this validation, we used samples from the first wave of the pandemic in the campania region of italy, providing the first data on the progress of the epidemic in a region with low levels of spread of the virus in the first half of 2020. results demonstrate that the genexus x evolution 2 system is capable of rapidly and accurately determining the sars-cov-2 genomics sequence, and that this method can be implemented in a clinical setting. the data also suggest that the system may be useful for further phylogenetic and epidemiological investigations. the results obtained, however, highlight the potentiality of the genexus system: (i) the users are involved only in the sample preparation and quantification, (ii) the automated process allows the users to focus on ngs raw data check and subsequent bioinformatic analysis, (iii) the technology allowed the complete genome sequencing of 78% of the samples (21/27) obtained from routine sars-cov-2 diagnosis process, with additional 3 samples showing nearly completed sequencing of the genome (>95%). moreover, the genexus system permits the sequencing of 32 multiplexed samples in less than 24h, representing a useful method for sars-cov-2 surveillance during a pandemic event. the genexus system could be useful to analyse in a few days, respect to the sanger sequencing, several viral strains from

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patients with an abnormal clinical presentation such as a very late viral clearance or to identify possible new variants in situations of rapid increase of contagions.

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