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Computational approaches applied to TB have predominantly implemented standard commercially available cheminformatic methods as will be described in the following section. These methods have been generally used by specialists focused on a single target or series of compounds and rarely in combination with other computational tools.

*Computational databases, pathway and
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The amount of publicly available large-scale biomedical and pharmaceutical data is growing exponentially, and computational drug repositioning approaches using data mining, machine learning, and network analysis become ever more critical when it comes to systematic drug repositioning due to the ability to overcome classical statistical approaches limitations and unreliable conclusions.

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