

Considerable effort has been and is being expended in the development of screening assays, particularly as a response to the need for the evaluation of large numbers of samples in HTS and the expectation that many new targets will be identified in the wake of genome sequencing projects. High-throughput screening involves the robotic handling of very large numbers of candidate samples, the registering of appropriate signals from the assay system, and data management and interpretation. However, the advent of HTS, whereby lead discoveries may be identified in a matter of days from libraries of 103–105 compounds, may be limited by the provision of sufficient quantities of the assay components. The development of surrogate hosts, such as enzyme inhibition, receptor binding, and cell function assays, provides possible means of alleviating such bottlenecks.

A large number of drugs manufactured through recombinant methods are coming off patents, creating markets for biogeneric medicines.

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