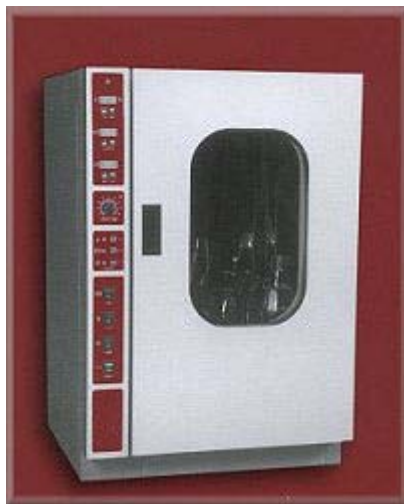


Vertiga-IM Shaker



Manufacturer: [Thomson Instrument Company](#)

Model: Vertiga 1M Shaker(#381150)

Primary Use: Small Scale Protein Expression in insect or Mammalian Cells

Description: Vertiga 1M has a platform that holds 12 plates and a top shelf. The orbit can be adjusted for doing Well Plates or Flasks, so whether one is screening or expressing, both needs are addressed. This unit has a cooling capability to 5°C, and can maintain between 5°C-60°C with 0.1 degree temperature fluctuation. Exterior dimensions are 26 in. width x 27 in. depth x 42in and a weight of 250 lbs.

The Vertiga IM shaker was developed as a collaboration project between JCIMPT and Thomson Instruments Company and is a modification of the Vertiga I Shaker which was developed for small scale expression in bacterial cells and which was adopted by the Joint Center for Structural Genomics (JCSG) for small scale expression studies. The Vertiga-IM is a second generation small-scale protein expression system [See References below] that enables screening of insect or mammalian (e.g. CHO and HEK293) cells in suspension. Three parameters were optimized using the Vertiga-I as a starting point: orbital path, throw distance, and temperature control. The throw distance for the circular orbit was optimized to one-half inch for use with the Thomson 24-well plates. However, the throw is adjustable to accommodate different vessel formats. Finally, the temperature control was upgraded from the peltier system used on the Vertiga-I to a compression refrigeration system, which is capable of maintaining a temperature from 5–60 °C (± 0.1 °C).

References

1. R. Page, K. Moy, E.C. Sims, J. Velasquez, B. McManus, C. Grittini, T.L. Clayton, R.C. Stevens, Scalable high-throughput micro-expression device for recombinant proteins, *BioTechniques* 37 (2004) 364–370.
2. W. Peti, R. Page, K. Moy, M. O’Neil-Johnson, I.A. Wilson, R.C. Stevens, K. Wu” thrich, Towards miniaturization of a structural genomics pipeline using micro-expression and microcoil NMR, *J. Struct. Funct. Genomics* 6 (2005) 259–267.
3. Hanson MA, Brooun A, Baker KA, Jaakola VP, Roth C, Chien EY, Alexandrov A, Velasquez J, Davis L, Griffith M, Moy K, Ganser-Pornillos BK, Hua Y, Kuhn P, Ellis S, Yeager M, Stevens RC., “Profiling of membrane protein variants in a baculovirus system by coupling cell-surface detection with small-scale parallel expression.”, *Protein Expr Purif.* (2007) ;**56**(1):85-92.