

# xCELLigence System

## 实时、无标记、高通量细胞分析系统

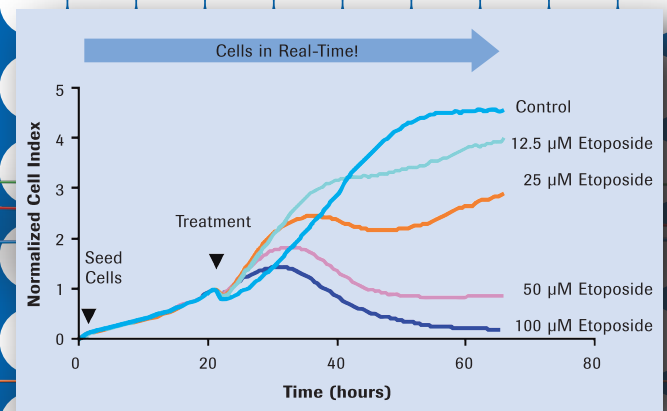
集合了分子生物学、细胞生物学与微电子方法学，突破了传统终点法细胞分析技术的瓶颈，为细胞学基础研究和药物研发提供了新型的、高通量筛选和检测平台。

### 性能

- 轻松检测从数分钟到数周细胞效应
- 高精确性和高重复性，提供更大的动态检测范围
- 灵活性高、易操作，软件友好直观

### 应用领域

- 细胞增殖
- Compound介导的细胞毒作用
- G蛋白偶联受体活性

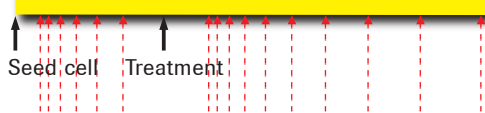


**Figure 1: Real-time monitoring of cytotoxicity**  
 Etoposide is a topoisomerase II inhibitor. At high concentrations it causes DNA damage and cell death, while at lower concentrations it leads to S-Phase and/or G2-M phase arrest.

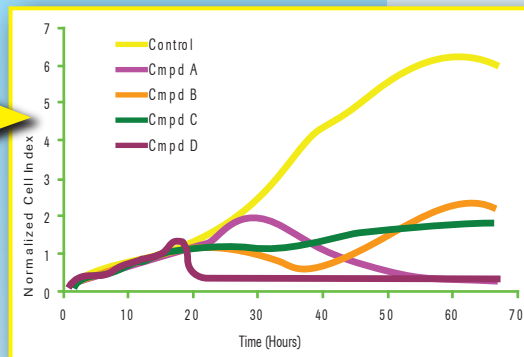
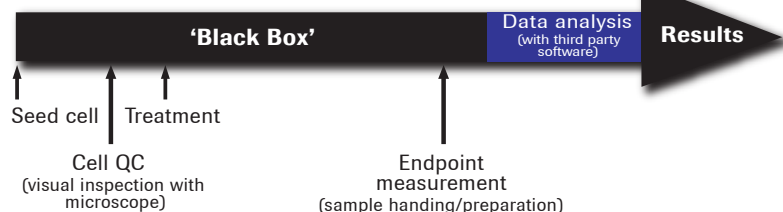


## xCELLigence workflow (seamless)

Automated real-time continuous measurements & data analysis → Results



## Endpoint assay workflow (step-wise)



▲ 化合物效应曲线图谱

◀ xCELLigence创新检测技术完全消除了传统终点法检测过程可能存在的“黑箱”问题

The xCELLigence system provides dynamic, real-time cell-based assays without the use of labels. The assays provide valuable data that traditional end-point analysis can't, leading to faster assay development and improved attrition rates. The assays offer several throughput options, from 16 to 576 (6x96) wells run simultaneously.

**SCIENCE VOL 320 20 JUNE 2008**

In summary, the RT-CES method based on electrical impedance measurement can provide real-time assessment of the cytotoxic activity induced by particulate matter without interference from the insoluble particles. The fully automated measurement without any labeling materials and reagents is potentially useful for large scale screening of particle-induced cytotoxicity. This provides a unique approach for biomonitoring of air quality as demonstrated by the SRM testing results.

**Analyst, 2008, 133, 643-648 | 643**

By dynamically measuring the response of the cells to chemotherapy using the RT-CES system, we identified that 6 to 8 h following etoposide treatment was the critical time to determine the cell death or cell survival.

**Mol Cancer Ther 2007; 6 (12). December 2007**

## 参考文献:

1. Identification of ALK as a major familial neuroblastoma predisposition gene. Mossé, YP, Laudenslager M, Longo L, Cole KA, Wood A, Attiyeh EF, Laquaglia MJ, Sennett R, Lynch JE, Perri P, Laurens G, Speleman F, Kim C, Hou C, Hakonarson H, Torkamani A, Schork NJ, Brodeur GM, Tonini GP, Rappaport E, Devoto M, Maris JM. Nature advance online publication 2008, August 24; doi: 10.1038/nature07261. (实时细胞分析系统用于siRNA研究)
2. A functional screen identifies miR-34a as a candidate neuroblastoma tumor suppressor gene. Cole KA, Attiyeh EF, Mosse YP, La Guaglia MJ, Diskin SJ, Brodeur GM, Mar JM. Mol Cancer Res 2008 May; 6(5): 735-42. (实时细胞分析系统用于micro RNA筛选)
3. Compound cytotoxicity profiling using quantitative high-throughput screening. Xia M, Huang R, Witt KL, Southall N, Fostel J, Cho MH, Jadhav A, Smith CS, Inglesse J, Portier CJ, Tice RR, Austin CP. Environmental Health Perspectives 2008 March; 116(3): 284-291. (实时细胞分析系统用于化合物高通量筛选)