































## **CIM** potential Examples Healthcare: DNA sequencing • we assume we have 200 GB of DNA data to be compared to A healthy reference of 3GB for 50% coverage\*\* [\*\*E. A. Worthey, Current Protocols in Human Genetics, 2001] • Mathematic: 10<sup>6</sup> parallel additions Assumptions • Conventional architecture · FinFET 22nm multi-core implementation, with scalable number of clusters, each with 32 ALU (e.g comparator) • 64 clusters; each cluster share a 8KB L1 cache CIM architecture • Memristor 5nm crossbar implementation • The crossbar size equals to total cache size of CMOS computer [Source: S. Hamdioui, et.al, DATE 2015] © Said Hamdioui, Computer Engineering, TUDelft 17

CIM potential					
<ul> <li>Metrics         <ul> <li>Energy-delay/operation</li> <li>Computing efficiency : number of operations per required energy</li> <li>Performance area : number of operations per required area</li> </ul> </li> <li>Results</li> </ul>					
	Metric	Archit.	DNA sequencing	10 <sup>6</sup> additions	
	Energy –Delay/ operations	Conv.	2.02e-03	1.5043e-18 9.25702-21	> x100
	Computing Efficiency	Conv.	4.11e01 3.70e04	6.5226e+9 3.9063e+12	> x100
	Performance Area	Conv.	5.73e06	5.1118e+09	> x100
CIM8.28e094.9164e+12Key drives:Reduced memory bottleneck, non-volatile technology, massive parallelismHamdioui, et.al, DATE 2015					
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