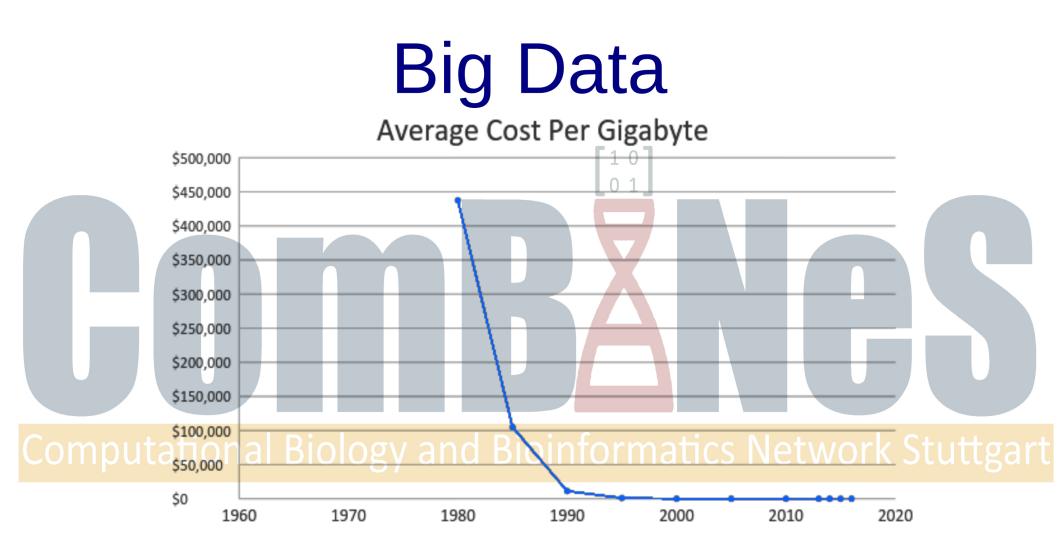
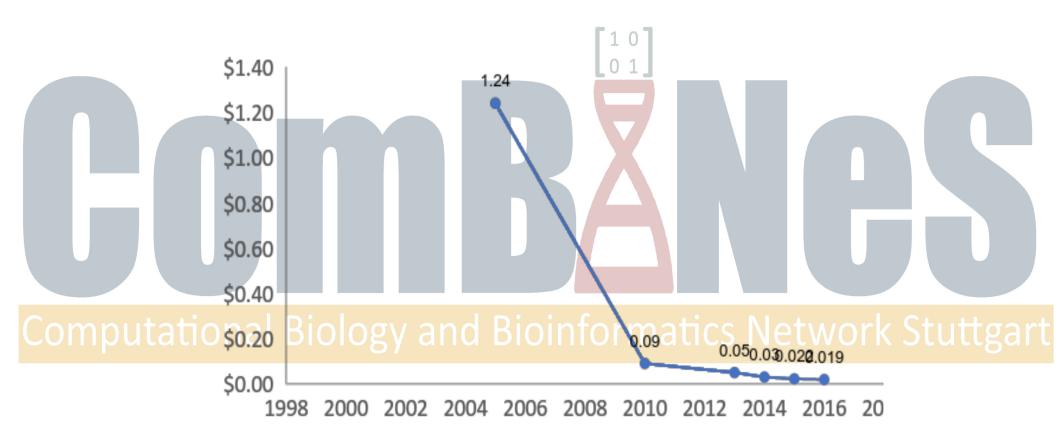
## ComBINeS

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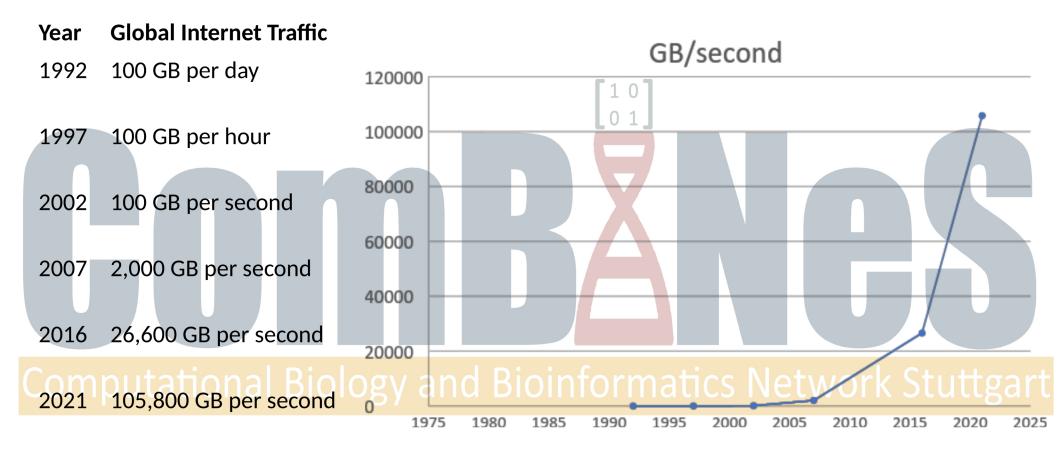
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#### Big Data in Biology & Medicine @ 06.06.2019 Dr. Nandor Poka





Data source: http://www.statisticbrain.com/average-cost-of-hard-drive-storage/



#### But what is Big Data?

- Like old data, just a lot more... too much to deal with traditional methods
- System logs of various kinds
- Financial transactions
- Online purchases

Etc.

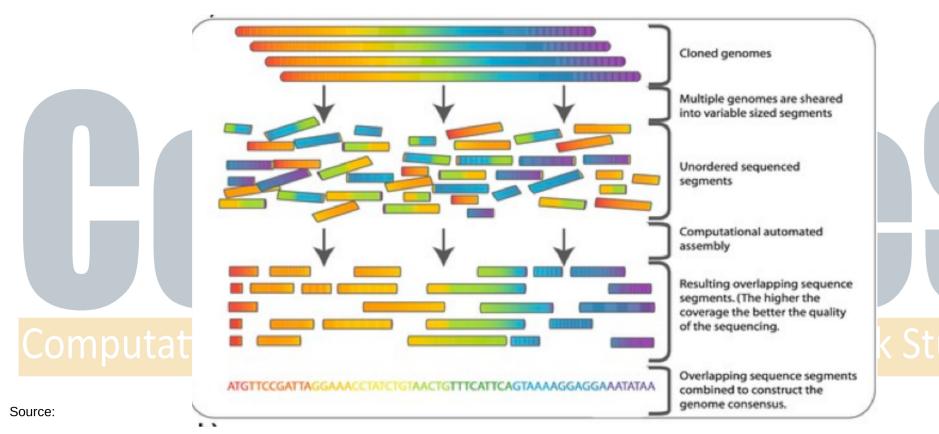
#### "New age" Big Data

- Social media likes, posts, contacts1 0
- User location data navigation searches, phone location data
- Community knowledge recommendations, product ratings
- Personal (non-professional) health data fitness trackers

# What sources of data we have in Biology and Medicine?

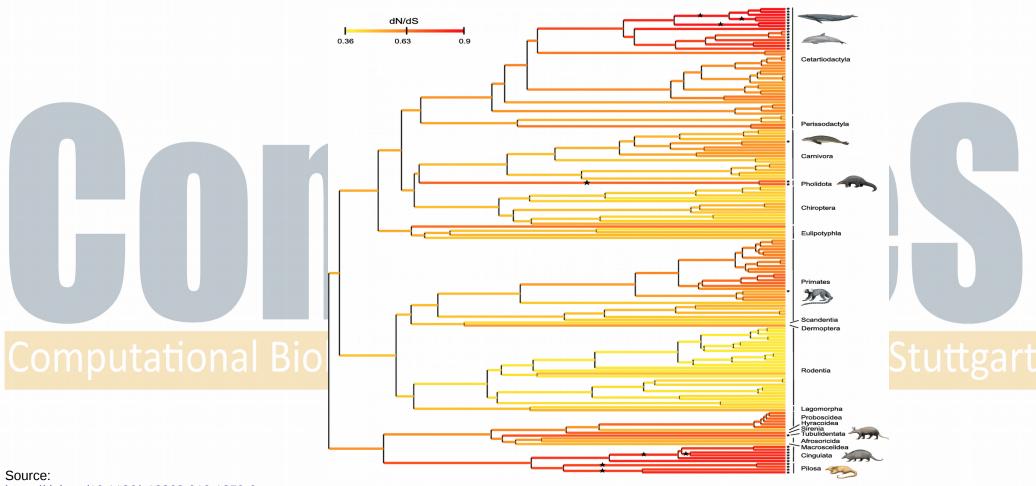
- Sequencing data
- Genomics / proteomics (and other "omics") data
- Gene / protein interaction networks
- Medical images
- Electronic personal medical data
- Wearable fitness data

#### Sequencing

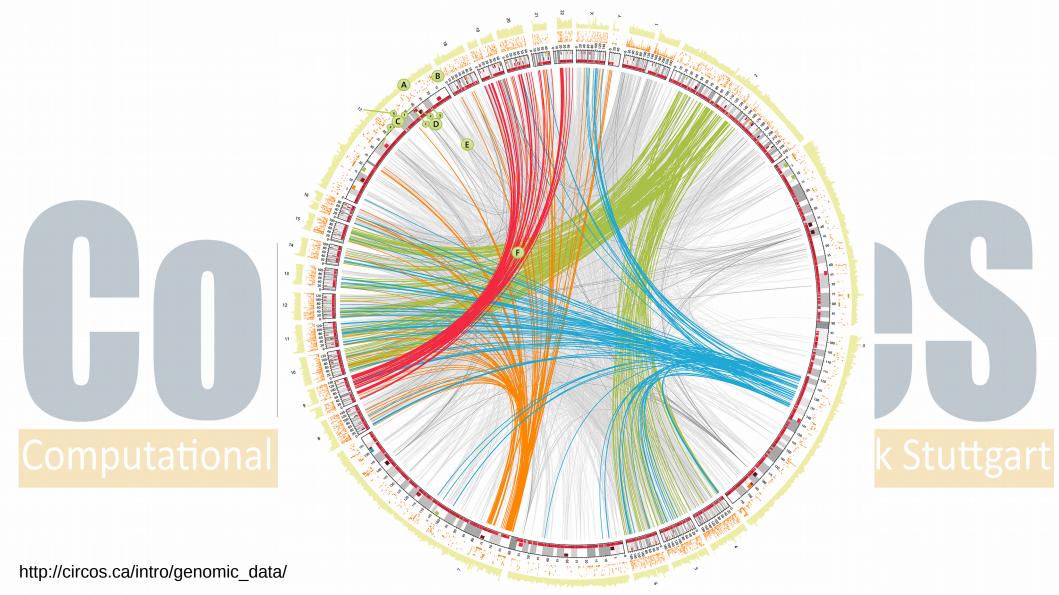


Computational Biology Methods and Their Application to the Comparative Genomics of Endocellular Symbiotic Bacteria of Insects. Biol. Procedures Online (2009). Commins, J., Toft, C., Fares, M. A.

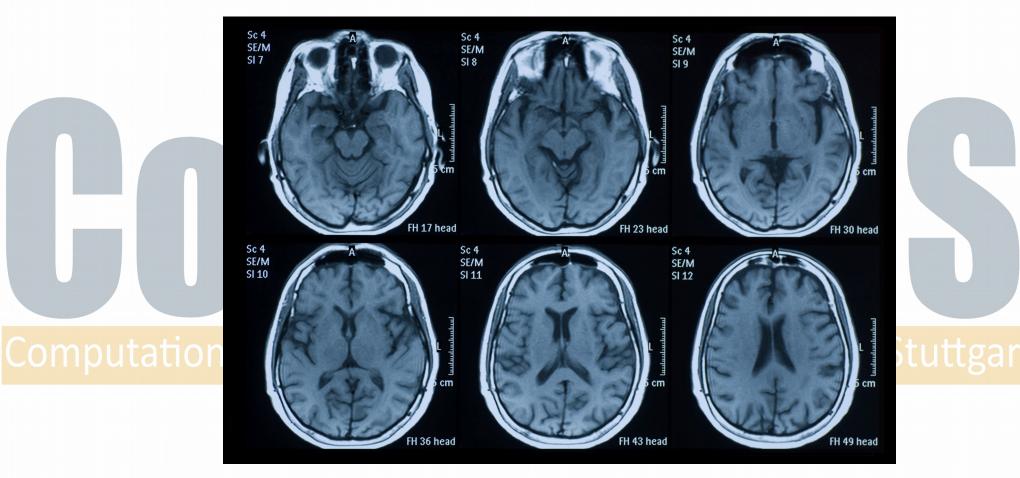
#### Genomics and other "omics"



https://doi.org/10.1186/s12862-019-1359-6



#### Medical imaging



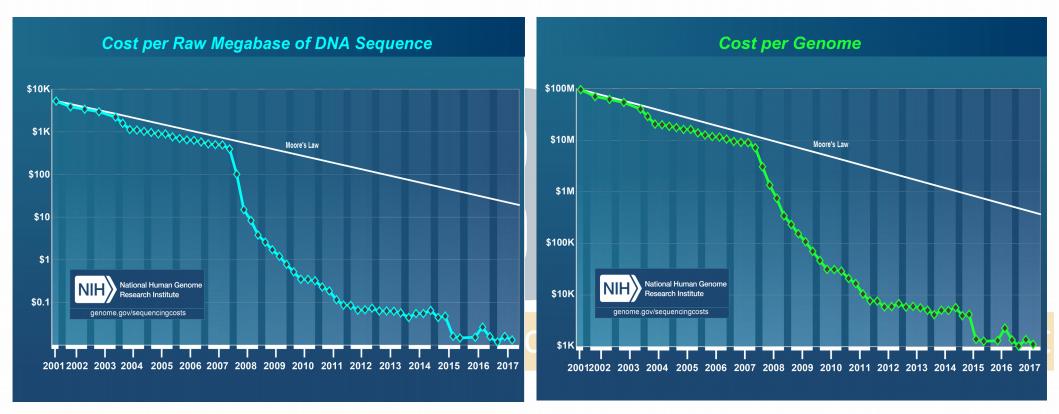
http://time.com/2860630/mri-scans-can-detect-early-onset-of-parkinsons-study-finds/

### What type of data?

- Sequences  $\rightarrow$  plain text, often compressed
- "Omics" data → heterogenous, but typically includes sequence data (often with added metadata / markup), tables / databases, graphs, 3D molecule models
- Medical images → high-res often multi-layered images / videos
- Medical records  $\rightarrow$  can be anything

• Personal fitness data  $\rightarrow$  varies, but somewhat unified  $\rightarrow$  eg. Google Fit data format / types informatics Network Stuttgart

#### How is this Big Data?



https://www.genome.gov/about-genomics/fact-sheets/DNA-Sequencing-Costs-Data

- The human diploid genome is approx 6.6Gb (GigaBase) → in plain ASCII it's 6.6GB (GigaByte)
- Typical sequencing is done with 100 base long reads and 40X coverage  $\rightarrow \sim 2.64*10^{9}$  reads  $\rightarrow 264$ GB raw sequence
- ~ 2x more with quality data  $\rightarrow$  0,5TB+
- Some organisms have genome size of 100-200Gb+ genome sizes

Computational Biology and Bioinformatics Network Stuttgart @SEQ\_ID GATTTGGGGTTCAAAGCAGTATCGATCA + !''\*(((((\*\*\*+))%%%++)(%%%%).1

#### For assembly

- de Bruijn graph with 66 Mill. Vertices
- Min. same amount of directed edges

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AAA

AAG

GAA

AGA

AGG

GAG

GGG

#### "Omics" data

- Sample from an gene expression experiment data
- 47191 rows by 405 colums =  $\sim$ 19,1M data points

	ID_REF	1NL.AVG_Sig nal	1NL.Detection Pval	1TU.AVG_Sig nal	1TU.Detection Pval	2NL.AVG_Sig nal	
	ILMN_17623 37	61.1	0.24156	59.8	0.31558	64.3	
	ILMN_20552 71	63.3	0.16364	68.8	0.07662	67.4	
	ILMN_17360 07	54.7	0.51818	61.6	0.24545	55.5	tuttgart

Dataset published: Wozniak MB, Le Calvez-Kelm F, Abedi-Ardekani B, Byrnes G et al. Integrative genome-wide gene expression profiling of clear cell renal cell carcinoma in Czech Republic and in the United States. PLoS One 2013;8(3):e57886. PMID: 23526956 Direct download: https://www.ncbi.nlm.nih.gov/geo/query/acc.cgi?acc=GSE40435 Dataset further used in this paper to make devise interaction network https://doi.org/10.1016/j.gdata.2017.10.006

#### Medical data

- MRI scans  $\rightarrow$  ~3Mpx per image  $\rightarrow$  ~1.7MB as lossless PNG
- 20 images per patient → 24MB / Scan
- In Germany, in 2015, 136.2 MRI scans / 1.000 resident
- (82.175M / 1000) \* 136.2 = ~11.2M Scans  $\rightarrow$  ~1,14PB (Petabyte) of data.

#### What can be achieved bio / medicine side?

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- "full stack" organism analysis
- Rapid pathogen detection in plants / animals as well
- Personalized medicine Cas9 / CRISPR → "on-site" gene repairing
- Rapid drug development

#### Challenges

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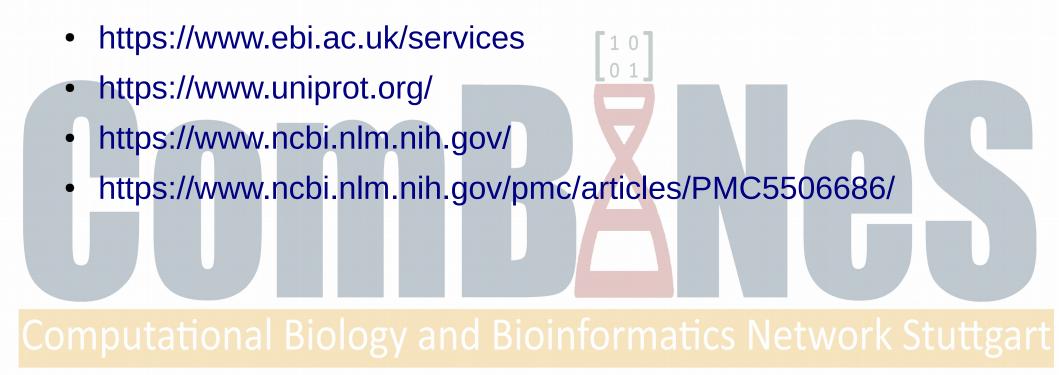
- Data privacy
- Cost manual labour, ingredient and equipment cost
- Quantity vs. quality data is abundant, but heterogeneous
- Visualization
- Speed(up)
- Free / commercial software

#### What can be done on IT side?

- Better, faster software / algorithms cluster / parallel computing
- AI for applied cases detecting micro fractures on X-rays, detecting tumors on MRI
- Better equipments, data acquirement
- Applied Data analysis / predictions:
  - Live predictions epilepsy, heart attacks

 Early onset detection of neurodegenerative diseases (eg. Compuparkinsons) 1010gy and Bioinformatics Network Stuttgart

#### Get your share of the data!



#### Questions?

