



The Elements of Bioinformatics

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1970 2014

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OPEN
SOURCE
TOOLS

Bn	So
Hm	Mq
Mu	Bo
Ss	Bw
Ex	Bf
Fa	Sm
Ac	Ja
	MI
	Po

KEY TO TOOL TYPE (stripes indicate new update)	
Aligners (pairwise)	Gene Prediction (mRNA)
Aligners (short read)	Gene Prediction (ncRNA)
Other Aligners	Sequence Tools
Multiple Sequence Aligners	Database/Warehouse
Assemblers Genomic (long read)	Workflows
Assemblers Genomic (short read)	Genome Browsers
Assemblers (mRNA)	ToolKits and APIs
Structure Modelling	Structure Visualisation

1970 2014

Py	Em	Bj	Bc	Bi	Ms
Bs	Eh	En	Br	La	Mve
Bm	Ta	Eb	Bp	Pl	Sa
In	Gx	Ap	Ig	Ut	Am
Ch	Kn	Gb	Nc	By	Pic
Pd	Up	Iv	Ag	At	As
Ct	Tg	Da	Jb	Mb	Pn
Ea	Pf	Sk	Sv	TP	
Dp	Nm	Gk	Ip	Cm	Cb
					Tm

TOOLS FREE FOR
ACADEMICS ONLY

Bl	Vm	Md	Rt	Av	Fg	Gn	Bh	Ga	Pm	Cr	DI	Pc	Td	Ts	Tgp
Cm	Sh	Ab	Mt	Mx	Mk	Mr	Pro	Pt	Sw	Pm	No	Uc	Mo	Ro	Sgp

COMMERCIAL
TOOLS

Zm	Ca	Gd	It	Sr	Gp	An	Pk	Dx
Oz	Ne	Gp	Mh	Mc	Pp	Gs	Gu	Cb

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Eagle integrates any of these tools into any bioinformatics analysis or pipeline. We combine your and all public data: on the cloud, on your internal system and, coming soon, through a simple web-interface. [Learn more.](#)

The initial selection focused on genomics but we are now opening it up to cover the whole of bioinformatics - proteomics, metabolomics, etc. Please [send us your suggestions](#) for additional tools - we need your input!

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