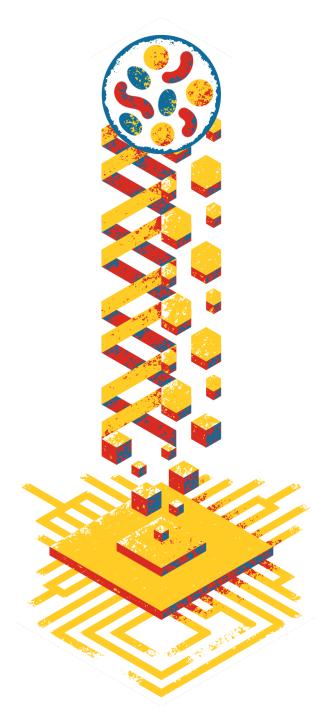
## Load genome from file .... -> select the <MAG>.contigs.fa -file

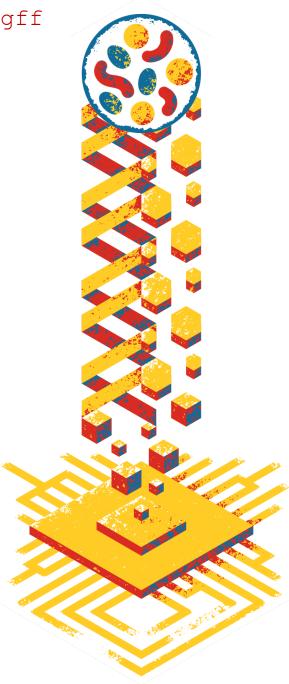
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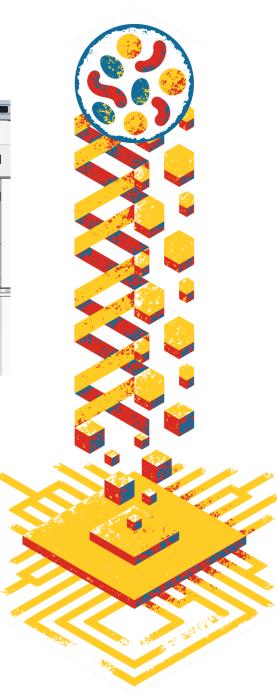


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	HJNANFJJ_00934												

gene representation (arrows are transcription direction), hover over it to get further annotation



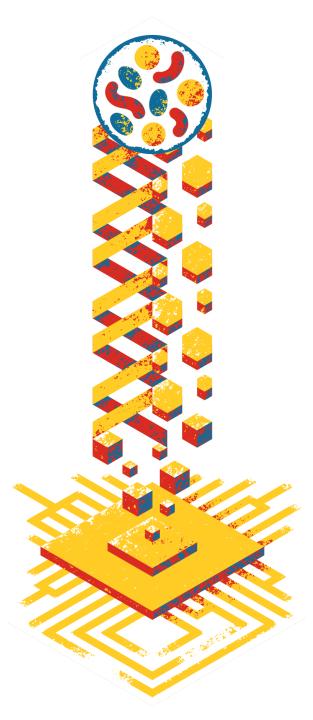


## Load from file .... -> select <MAG>.mg.bam

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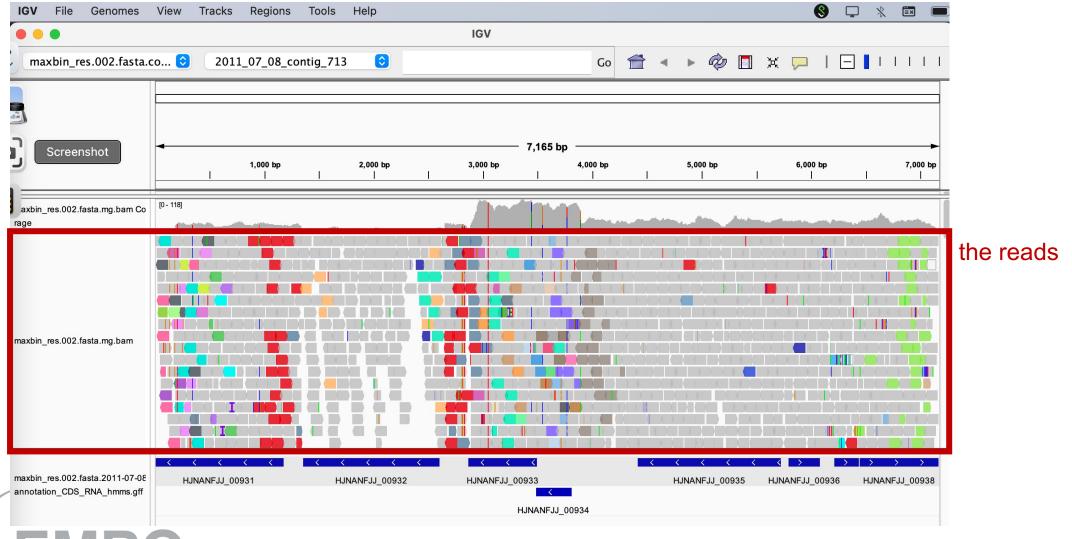
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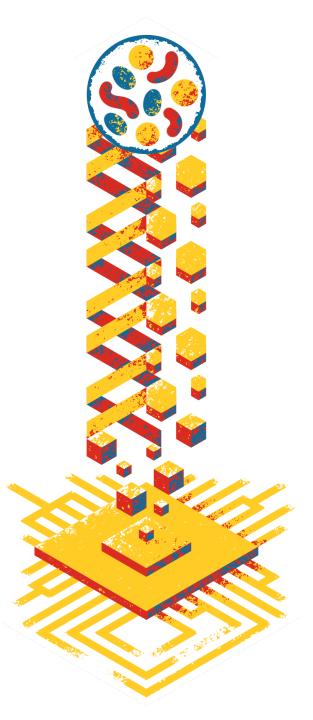
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### Load from file .... -> select <MAG>.mt.bam



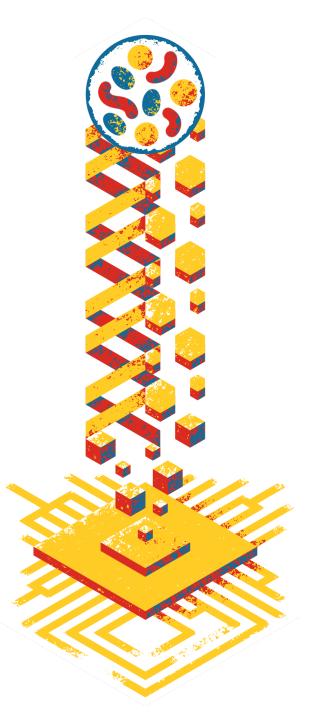




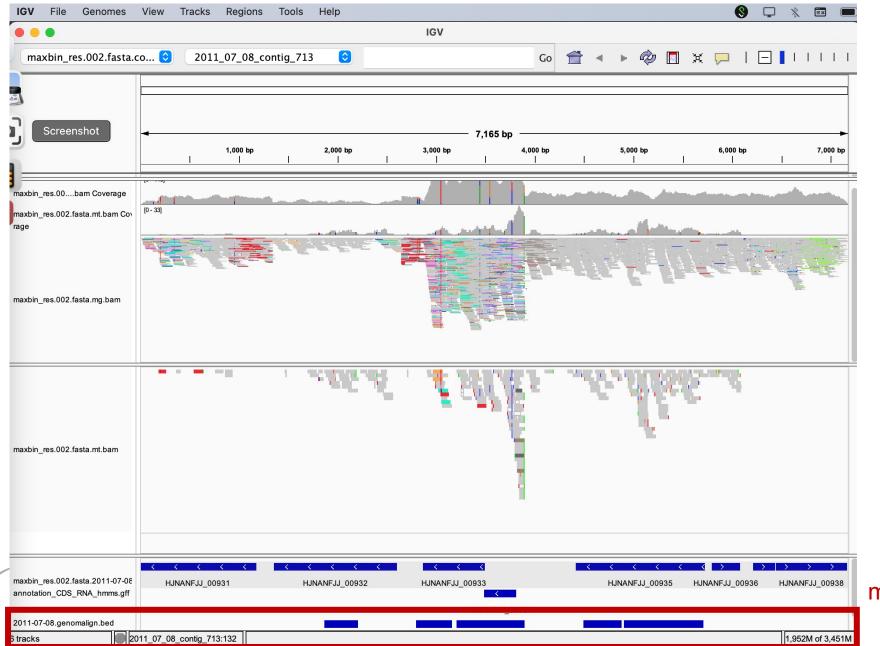


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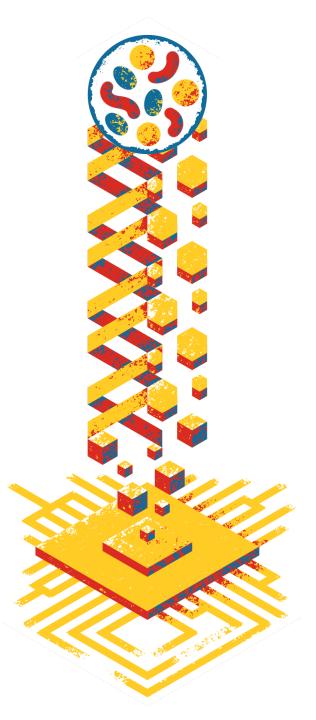
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metaT-assembly contigs

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