

Genome &



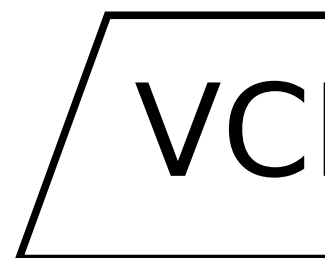
Bam

assembly



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Filter map
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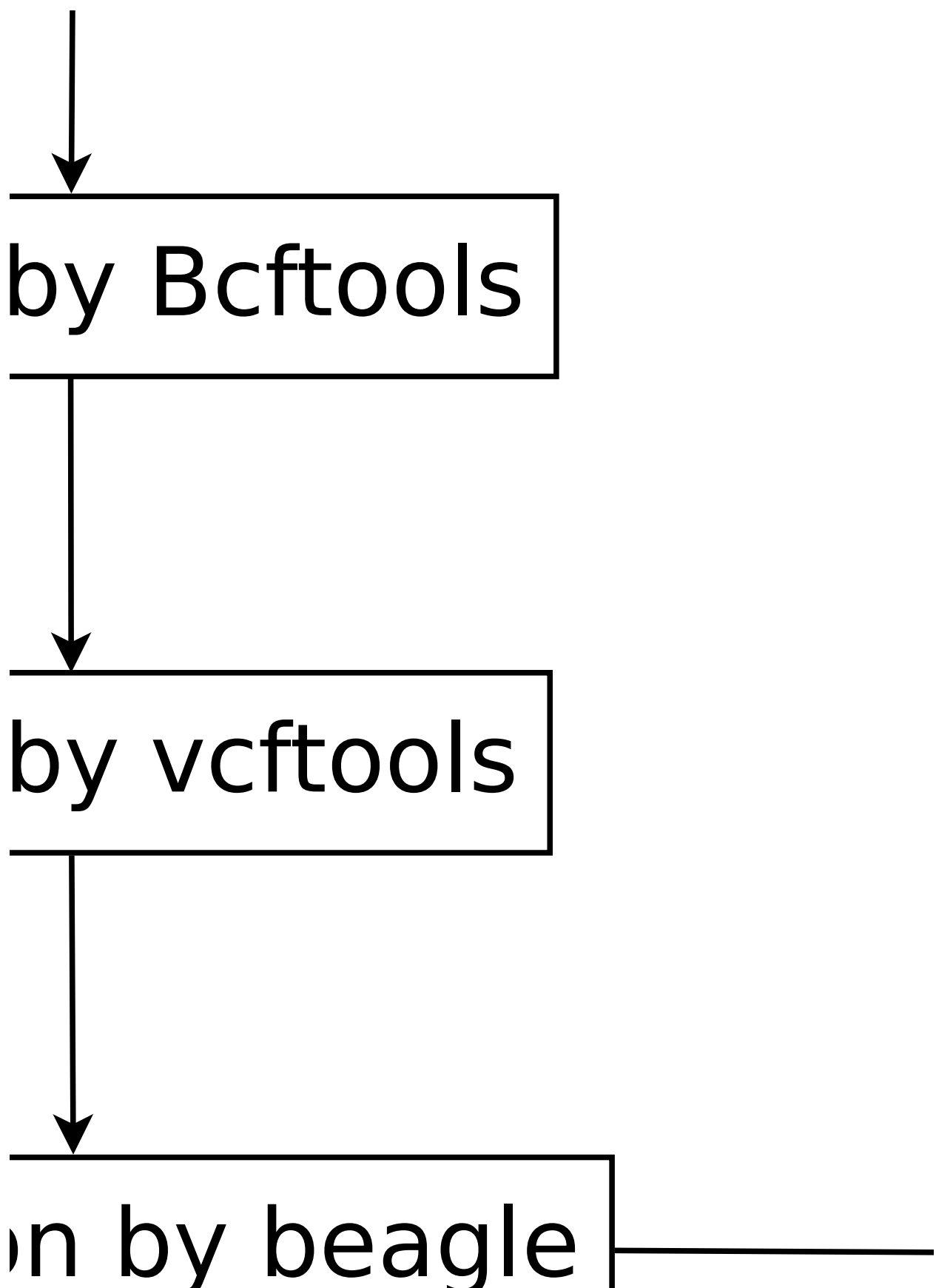
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Merged

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structure

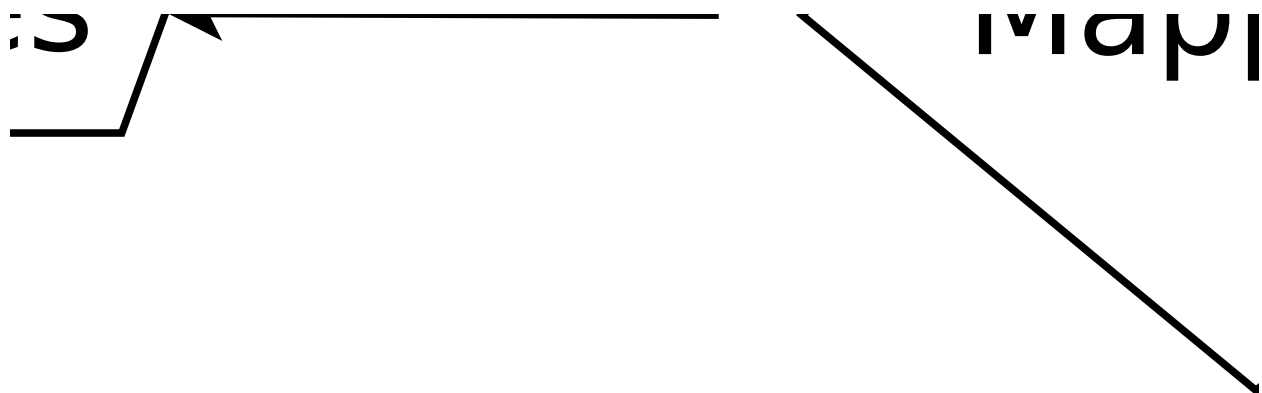
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Mapped BAM file



Extract sequences from
first genome (bedtools)

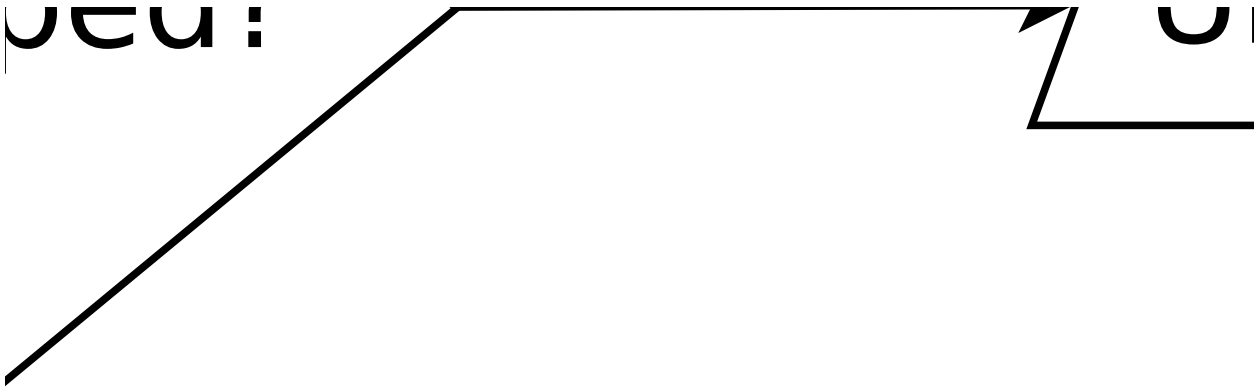




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unmapped pairs



Extract reads and
assembly (IDBA)

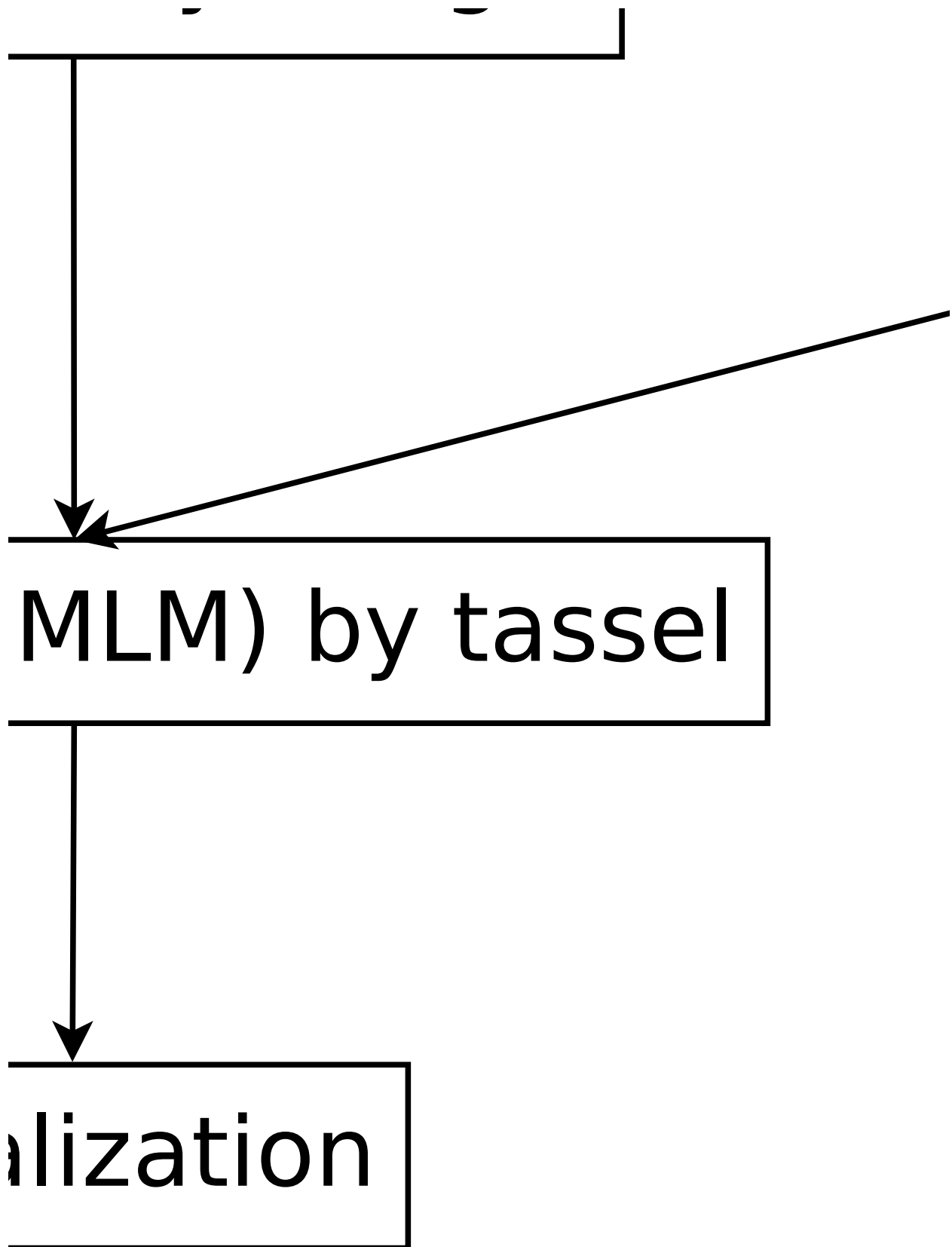


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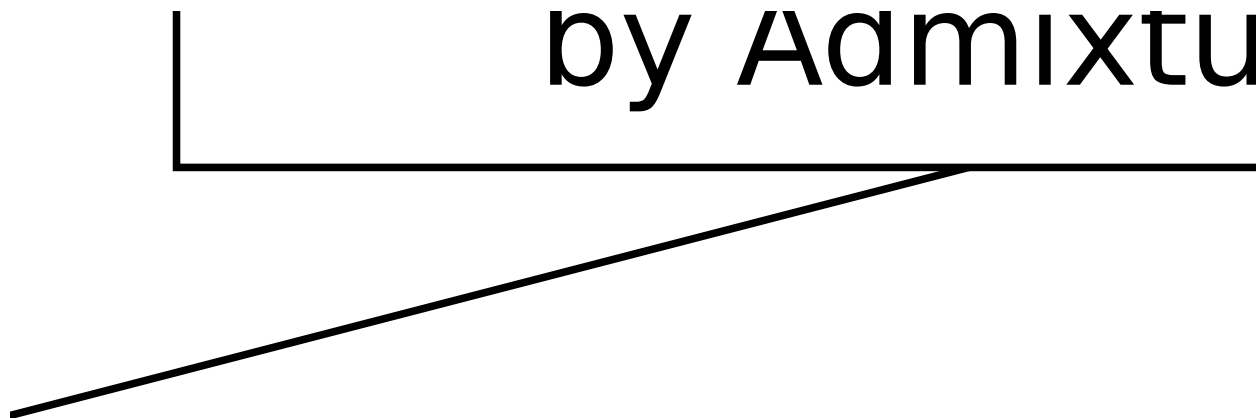
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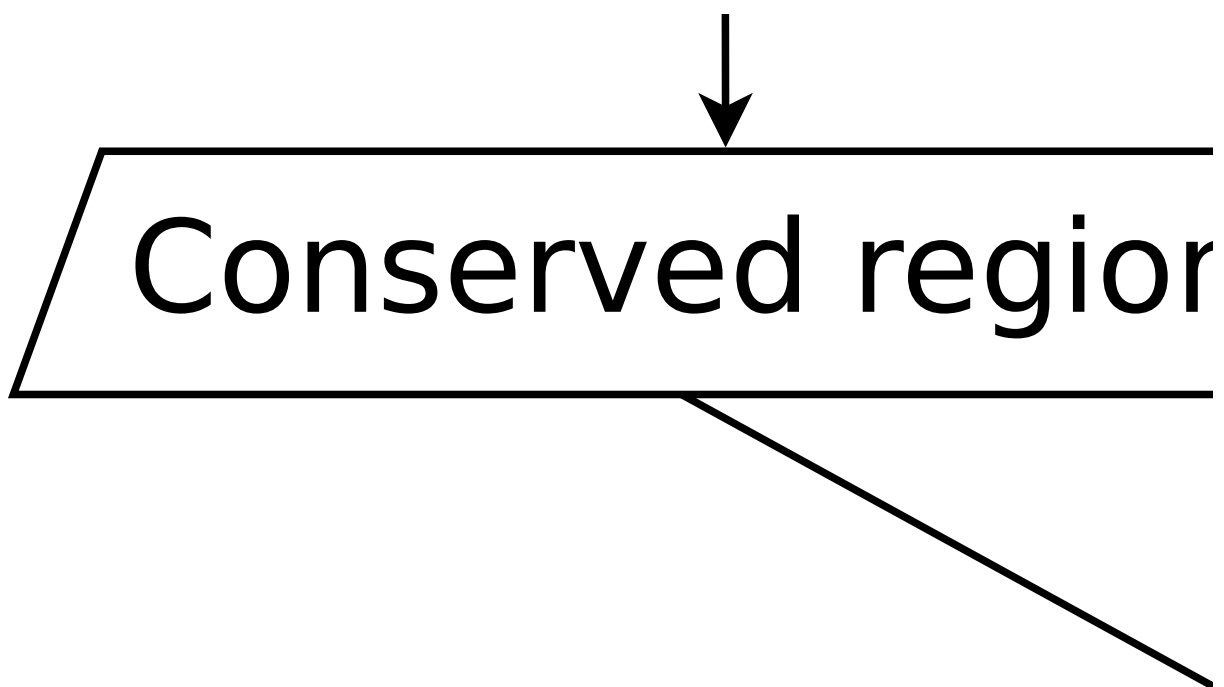


by *Admixtu*

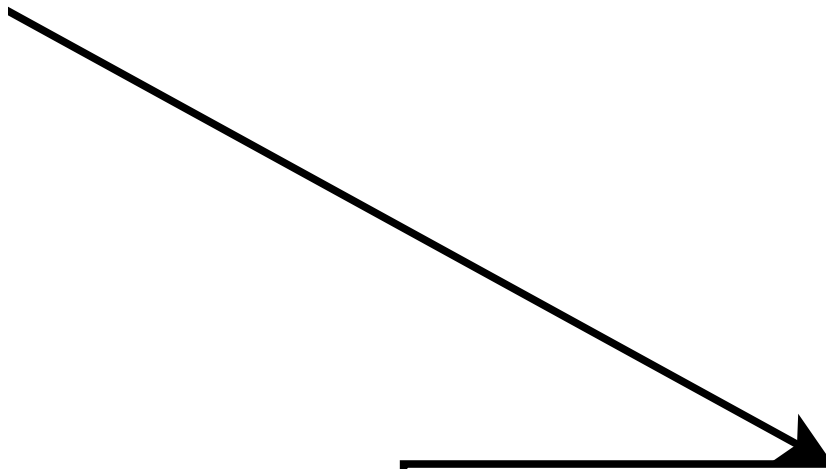


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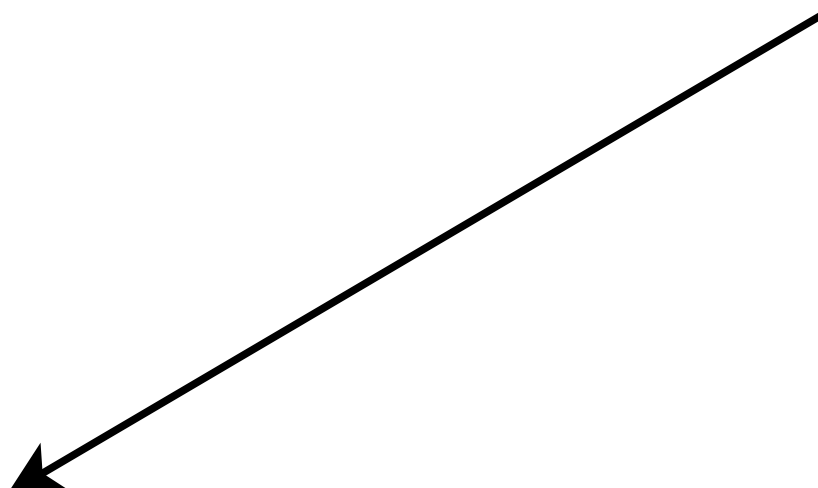


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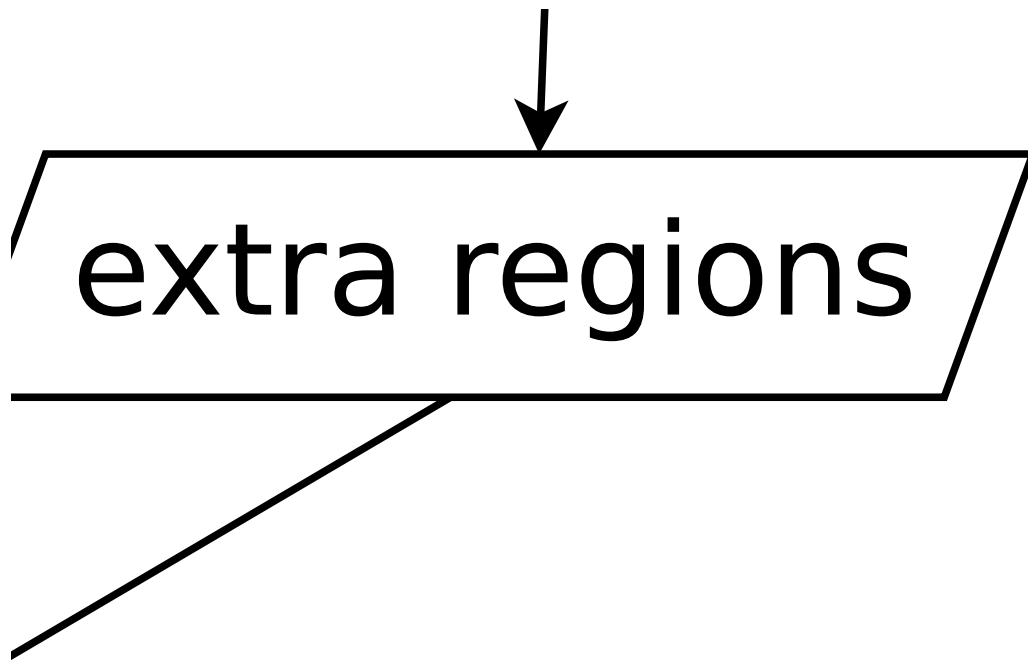


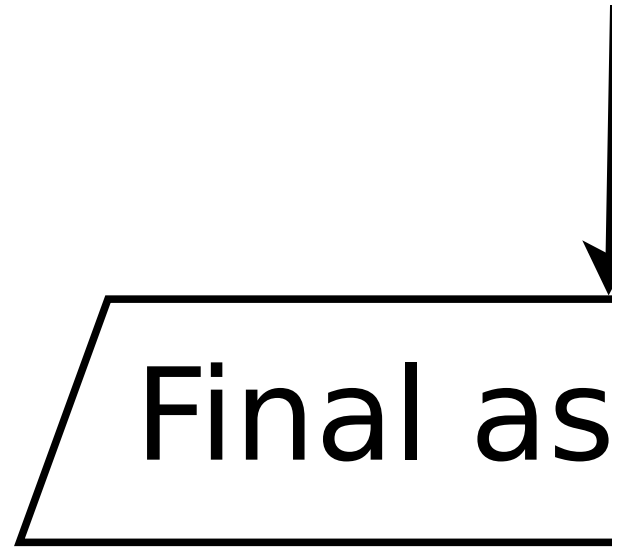
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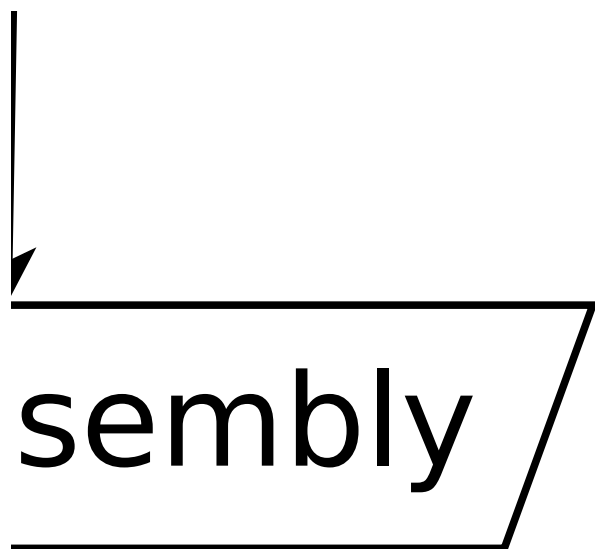
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