

KIRAL

*A tool for aligning reads to KIR(Killer
hemoglobin-like receptors) Allele
Sequences*

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Input format:

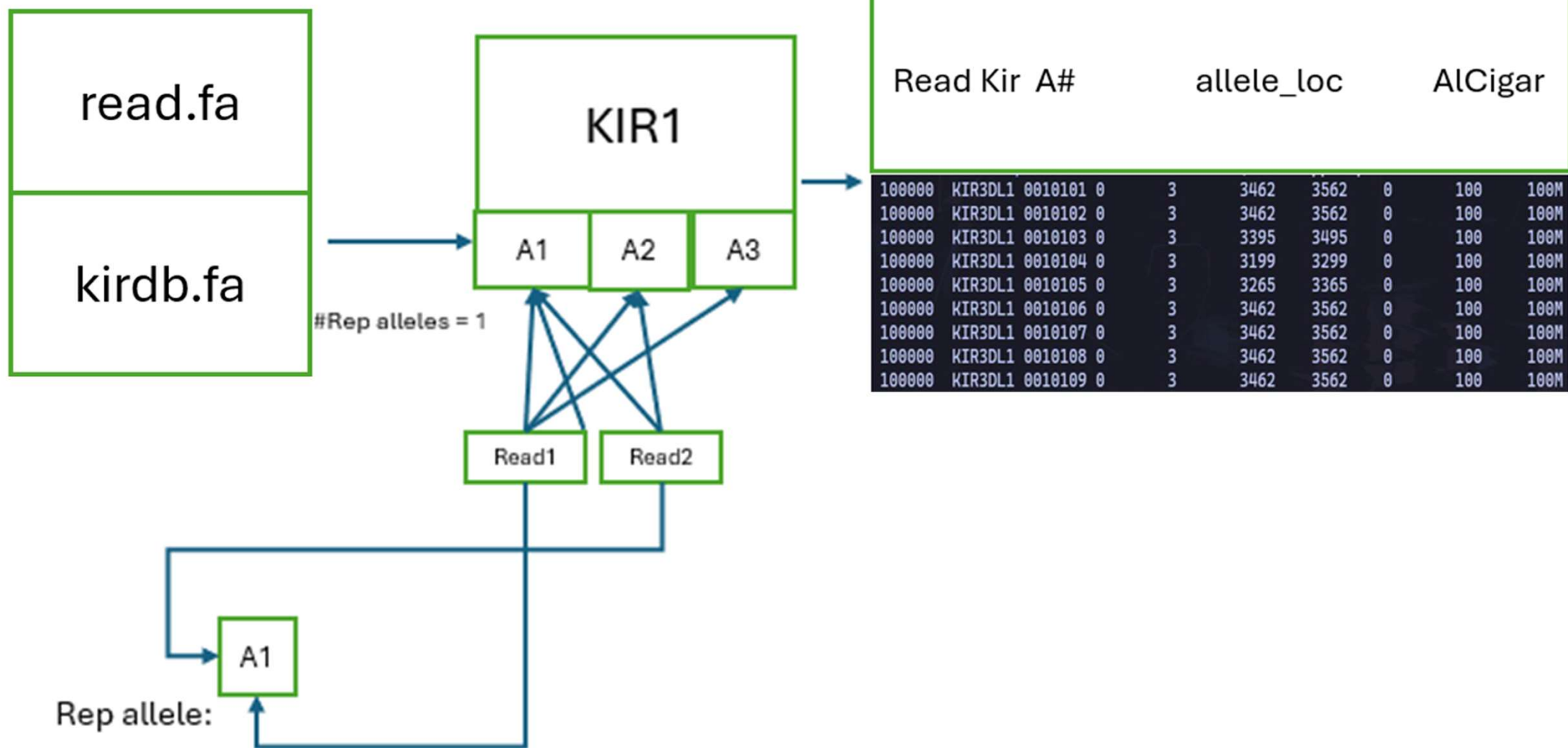


Naive:

#Rep alleles = 1



Categorical:



Regional:

#Rep alleles = 1

