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Iso-Seq mapping to L1HS/PA2 consensus sequence

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Protocol status: Working

We use this protocol and it's working

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Abstract

The protocol describes the steps to map HiFi reads to a consensus sequence and retrieve density plots

Troubleshooting



Mapping to L1HS/PA2 consensus sequence

- 1 A L1HS and L1PA2 consensus sequence was used to create a minimap2 (version 2.24; RRID:SCR_018550) index

Command

L1 consensus index minimap2

L1 consensus index minimap2

```
minimap2 -d L1consensus.mmi L1consensus.fa
```

to map FLNC reads (HiFi reads).

- 2 The density of mapped reads was visualized in the Integrative Genomics Viewer (IGV) (version 2.12.3; RRID:SCR_011793)
- 3 The number of mapped reads in the L1s 5' UTR was retrieved using samtools view (-c) (version 1.9; RRID:SCR_002105), specifying the first 900 bp of the consensus sequence as the coordinates of interest.