Task 2: Alignment

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HISAT2 on Galaxy was used for alignment. There are totally 12 samples (6 fetal vs. 6 adult) available. However, due to the resource limitation, only half of them (3 fetal vs. 3 adult) were chosen to be analyzed.

Fetal

SRR1554537

55133946 reads; of these: 55133946 (100.00%) were paired; of these: 474170 (0.86%) aligned concordantly 0 times 52684816 (95.56%) aligned concordantly exactly 1 time 1974960 (3.58%) aligned concordantly >1 times

SRR1554538

68026190 reads; of these: 68026190 (100.00%) were paired; of these: 549305 (0.81%) aligned concordantly 0 times 65058884 (95.64%) aligned concordantly exactly 1 time 2418001 (3.55%) aligned concordantly >1 times

SRR1554541

69278357 reads; of these: $69278357 \ (100.00\%) \ were \ paired; of these: \\ 561488 \ (0.81\%) \ aligned \ concordantly \ 0 \ times \\ 66402692 \ (95.85\%) \ aligned \ concordantly \ exactly \ 1 \ time \\ 2314177 \ (3.34\%) \ aligned \ concordantly \ >1 \ times$

Adult

SRR1554535

38063721 reads; of these: $38063721\ (100.00\%)\ \text{were paired; of these:}$ $533477\ (1.40\%)\ \text{aligned concordantly 0 times}$ $36345262\ (95.49\%)\ \text{aligned concordantly exactly 1 time}$ $1184982\ (3.11\%)\ \text{aligned concordantly} > 1\ \text{times}$

SRR1554536

21450348 reads; of these: 21450348 (100.00%) were paired; of these: 141926 (0.66%) aligned concordantly 0 times 20543060 (95.77%) aligned concordantly exactly 1 time 765362 (3.57%) aligned concordantly >1 times

SRR1554539

33742728 reads; of these: 33742728 (100.00%) were paired; of these: 503288 (1.49%) aligned concordantly 0 times 32420443 (96.08%) aligned concordantly exactly 1 time 818997 (2.43%) aligned concordantly >1 times