Clonality report

ARResT/Interrogate version 1.99.8

Date: 04/11/2023

Abundance filtering range: 0 - 100 %

Junction aa length range: 7 - 24

Junction class sets with no data:

DJ:Dh-Jh, VJ:Vk-Jk, Vk/intron-Kde

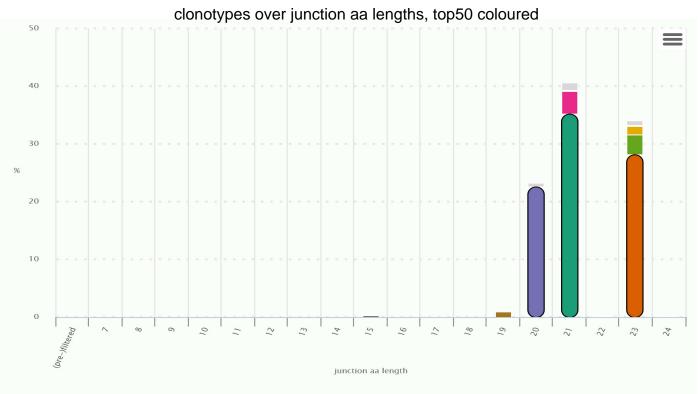
results.name	filtering	sample
2023_11_03IgHV	not pre-filtered	Marzena-Poli-259_S18_L001_R1_001

QC status	QC report	raw reads	w ju(nction)		species	short (<50nt)
pass		755417	673484	89.15%	7	0.29%

VJ:Vh-(Dh)-Jh junction class set

stats for junction class set

reads for normalisation	673484
filtered-in clonotypes	14564
clonotype 1 to mean of 5+6+7 ratio	17.64
clonotype 2 to mean of 5+6+7 ratio	14.07



top15 clonotypes over junction aa lengths

			2		
	genes and segmentation	seq	len	reads	norm.9
1	V5-51 -1/36/-0 J4	CARRGYSSSWPTHSLYYFDYW	21	237616	35.2
2	V4-39 -0/36/-11 J6	CARQAQKLRYFDCPG#YYYMDVW	23	189587	28.1!
3	V1-69=V1-69D -0/29/-0 J5	CAREGERFLEWSPYNWFDPW	20	151946	22.50
4	V3-30 -1/31/-10 J6	CARVSLGGEFLSFRYYGMDVW	21	25668	3.8:
5	V6-1 -1/38/-1 J2	CARGPPPLSGSYPTVGYWYFNLW	23	23703	3.57
6	V6-1 -1/38/-1 J2	CARGPPPLSGSYPTVGYWYFDLW	23	10095	1.50
7	V4-39 -0/37/-24 J6	CARQAQKLRYFDCPGA#VW	19	6619	0.98
8	V7-4-1 -2/19/-2 J3	CARPVPTTENAFDFW	15	1659	0.2!
9	V4-39 -0/36/-11 J6	CARQAQKLRYFDCPG#YYYMDVW	22	529	0.08
10	V3-30=V3-33 -1/31/-10 J6	CARVSLGGEFLSFRYYGMDVW	21	432	0.00
11	V5-51 -1/38/-2 J4	CARRGYSSSWPTHSLYYFDYW	21	418	0.00
12	V3-21 -0/47/-3 J5	CARDVTTTSNRREYNSRGEDWFDPW	25	257	0.04
13	V6-1 -1/38/-1 J2	CARGPPPLSGSYPTVGYWYFNLW	24	218	0.0
14	V3-30=V3-66 -1/31/-10 J6	CARVSLGGEFLSFRYYGMDVW	21	170	0.0
15	V5-51 -1/36/-0 J4	CARRGYSSSWHTHSLYYFDYW	21	148	0.0