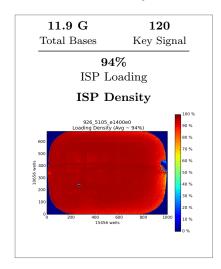
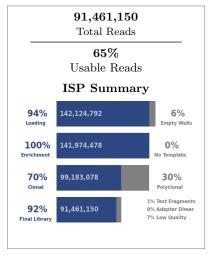
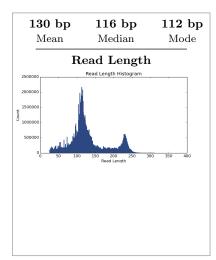
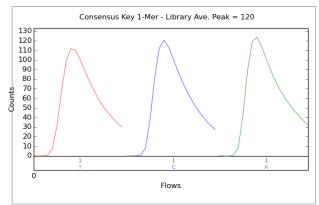
## Run Summary









Addressable Wells	$151,\!539,\!288$	
With ISPs	142,124,792	93.8%
Live	141,974,478	99.9%
Test Fragment	564,969	00.4%
Library	141,409,509	99.6%
Library ISPs	141,409,509	
Filtered: Polyclonal	42,791,400	30.3%
<u> </u>	· · ·	30.3% 04.9%
Filtered: Polyclonal	42,791,400 6,907,711	00.0,0

Barcode Name	Sample	Bases	$\geq Q20$	Reads	Mean Read Length	Read Length Histogram
No barcode	none	659,730,284	505,956,976	5,512,028	119 bp	0 50 100 150 200 250 300 350
$IonCode\_0155$	18084225	83,406,527	71,218,462	435,922	191 bp	0 50 100 150 200 250 300 350
$IonCode\_0156$	18084226	89,022,567	75,658,947	457,487	194 bp	0 50 100 150 200 250 300 350
$IonCode\_0157$	18084227	77,240,473	65,659,637	407,860	189 bp	0 50 100 150 200 250 300 350
IonCode_0158	18084229	110,077,796	94,027,149	560,761	196 bp	0 50 100 150 200 250 300 350
$IonCode\_0159$	18084251	115,787,060	97,794,707	618,147	187 bp	0 50 100 150 200 250 300 350
IonCode_0160	18084278	393,390,393	339,333,639	2,235,835	175 bp	0 50 100 150 200 250 300 350
IonCode_0161	18084281	513,194,794	432,602,728	2,778,631	184 bp	0 50 100 150 200 250 300 350

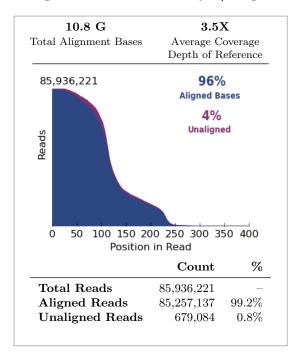


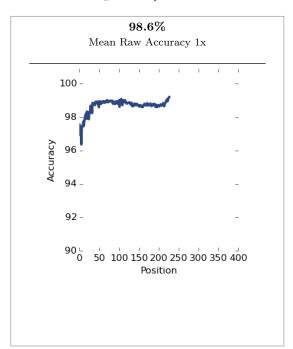
IonCode_0162	18084282	430,645,027	360,082,787	2,364,537	182 bp	0 50 100 150 200 250 300 350
IonCode_0163	18084284	619,793,587	527,842,177	3,365,103	184 bp	0 50 100 150 200 250 300 350
IonCode_0164	18084285	568,608,133	487,435,833	3,132,503	181 bp	0 50 100 150 200 250 300 350
IonCode_0308	18073151	278,814,232	259,335,679	2,408,648	115 bp	0 50 150 200 250 300 350
IonCode_0412	18084274	218,288,044	204,841,608	2,725,856	80 bp	0 5 100 150 200 250 300 350
IonCode_0413	18084282	380,525,864	336,410,199	2,445,759	155 bp	0 50 100 150 200 250 300 350
IonCode_0414	18084274	753,594,906	693,877,834	7,050,267	106 bp	0 50 10 150 200 250 300 350
IonCode_0415	18084282	609,458,837	546,337,749	5,043,786	120 bp	0 50 10 150 200 250 300 350
IonCode_0416	18084281	341,279,283	301,468,102	2,118,890	161 bp	0 50 100 150 200 250 300 350
IonCode_0417	18084278	317,955,043	281,109,349	1,996,411	159 bp	0 50 100 150 200 250 300 350
IonCode_0418	18084281	634,580,012	566,076,182	5,442,927	116 bp	0 50 100 150 200 250 300 350
IonCode_0419	18084284	319,201,186	283,570,081	2,198,967	145 bp	0 50 100 15 200 250 300 350
IonCode_0424	18084284	778,209,209	697,564,107	6,503,961	119 bp	0 50 100 150 200 250 300 350
IonCode_0426	18084285	803,175,371	735,108,355	7,189,947	111 bp	0 50 10 150 200 250 300 350
IonCode_0432	18084278	696,691,776	643,873,762	6,355,513	109 bp	0 50 100 150 200 250 300 350
IonCode_0470	18084252	1,871,092,933	1,731,590,588	16,292,283	114 bp	0 50 100 150 200 250 300 350
IonCode_0475	18084250	247,539,955	225,094,237	1,806,220	137 bp	0 50 100 15 200 250 300 350

Test Fragment	Reads	Percent 50AQ17	Read Length Histogram
$\mathbf{TF}_{-}\mathbf{C}$	387,682	91	. Такиментумысты сермен комуческый комуческый при самых терместы терместы стамментого муссы с сомобытым.
${f TF}_{-}{f 1}$	123,661	92	6 30 20 00 05 50 00 70 00 90 100 110 120 110 100 100 100 100 100 10



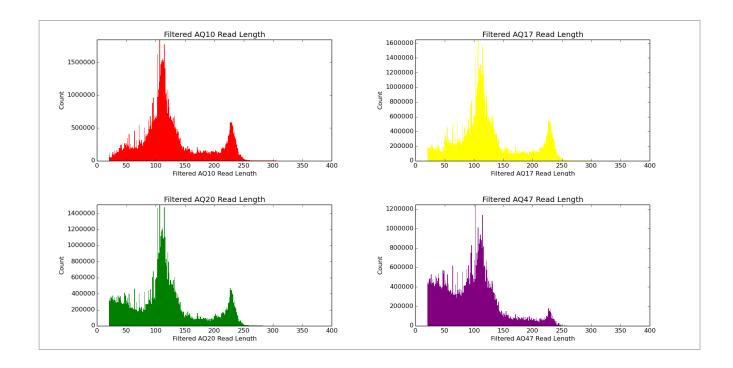
## Alignment Summary (aligned to Homo sapiens)





	AQ17	AQ20	Perfect
Total Number of Bases [Mbp]	9.37 G	8.34 G	6.65 G
Mean Length [bp]	123	116	97
Longest Alignment [bp]	347	345	336
Mean Coverage Depth	3.0	2.7	2.1







#### ${\bf coverage Analysis}$

**Library type:** AmpliSeq DNA **Target regions:** Barcode specific

gg					
Barcode Name	Sample	Mapped Reads	On Target	Mean Depth	Uniformity
IonCode_0155	18084225	432,903	93.78%	885.2	90.34%
IonCode_0156	18084226	454,862	95.21%	951.2	92.21%
IonCode_0157	18084227	403,477	93.49%	818.7	92.59%
IonCode_0158	18084229	557,076	95.67%	1,182	93.27%
IonCode_0159	18084251	614,693	93.54%	1,226	92.04%
IonCode_0160	18084278	2,219,316	89.37%	2,959	89.60%
IonCode_0161	18084281	2,760,120	91.37%	3,884	90.82%
IonCode_0162	18084282	2,343,290	90.12%	3,257	91.51%
IonCode_0163	18084284	3,336,450	90.46%	4,685	93.62%
IonCode_0164	18084285	3,102,139	90.45%	4,315	91.53%
IonCode_0308	18073151	2,401,002	67.22%	7,357	95.25%
IonCode_0412	18084274	2,643,887	9.20%	34.09	0.71%
IonCode_0413	18084282	2,434,092	92.12%	2,160	66.41%
IonCode_0414	18084274	6,968,813	0.08%	2.636	50.63%
IonCode_0415	18084282	5,016,964	0.20%	9.434	1.20%
IonCode_0416	18084281	2,109,717	93.93%	1,969	66.90%
IonCode_0417	18084278	1,982,497	92.86%	1,820	70.48%
IonCode_0418	18084281	5,380,844	0.73%	34.99	0.86%
IonCode_0419	18084284	2,188,108	89.04%	1,739	45.67%
IonCode_0424	18084284	6,427,449	0.57%	31.7	0.67%
IonCode_0426	18084285	7,143,784	0.07%	4.928	4.27%
IonCode_0432	18084278	6,324,602	0.24%	12.79	1.20%
IonCode_0470	18084252	16,212,136	99.30%	1,040	88.15%
lonCode_0475	18084250	1,798,916	90.06%	9,785	89.83%
H 1 1 P H	5	items per page			1 - 24 of 24 item



#### sampleID

Barcode Name	Sample	Sample ID	Coverage at 20x	Tracking Reads
lonCode_0155	18084225	M-TGRTRSRA	100.0%	4.99%
lonCode_0156	18084226	M-TTGTRSAW	100.0%	3.57%
lonCode_0157	18084227	F-TKRYRSRA	100.0%	4.45%
lonCode_0158	18084229	F-YKRCGGGT	100.0%	3.06%
lonCode_0159	18084251	F-CKAYRSGT	100.0%	4.76%
lonCode_0160	18084278	M-TGATRSRA	100.0%	3.59%
lonCode_0161	18084281	M-YGRYRCGW	100.0%	1.98%
lonCode_0162	18084282	M-YTRYRGRA	100.0%	2.25%
IonCode_0163	18084284	M-CKRTRCAW	100.0%	2.22%
IonCode_0164	18084285	F-YKRCRSRT	100.0%	2.62%
IonCode_0308	18073151	M-YGRCRCAT	100.0%	0.56%
IonCode_0412	18084274	N/A	0.0%	0.00%
IonCode_0413	18084282	M-YTRYRGRA	100.0%	5.21%
IonCode_0414	18084274	N/A	0.0%	0.00%
IonCode_0415	18084282	N/A	0.0%	0.00%
IonCode_0416	18084281	M-YGRYRCGW	100.0%	3.89%
IonCode_0417	18084278	M-TGATRSRA	100.0%	5.07%
IonCode_0418	18084281	N/A	0.0%	0%
IonCode_0419	18084284	M-CKRTRCAW	100.0%	6.32%
IonCode_0424	18084284	N/A	0.0%	0%
IonCode_0426	18084285	N/A	0.0%	0.00%
IonCode_0432	18084278	N/A	0.0%	0%
IonCode_0470	18084252	M-YKRYASRA	100.0%	0.11%
IonCode_0475	18084250	F-YKRYRSAW	100.0%	8.62%
M 1 P M 5		▼ items per page		1 - 24 of 24 items

 $\mathbf{HID}_{S}NP_{G}enotyper$ 

Click on the 'HID\_SNP\_Genotyper.html' link above to view results



#### **Analysis Details**

Run Name R\_2018\_08\_29\_13\_18\_26\_DGL-01-278-DGL\_926\_5105\_e1400e0

**Run Date** Aug. 28, 2018, 4:58 p.m.

Run Flows 520

Projects commercial

Sample | 18084285, 18084251, 18084284, 18084250, 18084278, 18084229, 18084274,

18073151, 18084252, 18084282, 18084281, 18084225, 18084226, 18084227

Reference

Instrument DGL-01

Operation Mode Customer mode

Flow Order TACGTACGTCTGAGCATCGATCGATGTACAGC

Library Key TCAG
TF Key ATCG

Chip Barcode

Chip Check Passed
Chip Type P1.1.17
Chip Data tiled
Chip Lot Number QPW135
Chip Wafer 24
Barcode Set IonCode

Analysis Name Auto\_DGL-01-278-DGL\_926\_5105\_e1400e0\_470

**Analysis Date** Aug. 30, 2018, 1:21 a.m.

Analysis Flows

runID XVEKJ

BeadFind ArgsjustBeadFind -args-json /opt/ion/config/args\_P1.1.17\_beadfind.jsonAnalysis ArgsAnalysis -args-json /opt/ion/config/args\_P1.1.17\_analysis.jsonPre-BaseCallerBaseCaller -barcode-filter-minreads 10 -phasing-residual-filter=2.0

-max-phasing-levels 2

Calibration Args Calibration

BaseCaller Args BaseCaller –barcode-filter-minreads 10 –phasing-residual-filter=2.0

-max-phasing-levels 2 -num-unfiltered 1000 -barcode-filter-postpone 1

Alignment Args | tmap mapall ... stage1 map4

IonStats Args ionstats alignment

Analysis Parameters | default

### Chef Summary

Ion Chef was not used for this run



# Software Version

$Torrent\_Suite$	5.10.0
host	4RZ9ZV1
ion-analysis	5.10.10-1
ion-dbreports	5.10.26-1
ion-gpu	5.10.0-1
ion-pipeline	5.10.8-1
ion-torrentpy	5.10.8-1
ion-torrentr	5.10.9 - 1
Script	2.1.45
LiveView	2389
DataCollect	3602
OIA	51000
OS	35
Graphics	97

