

@SFHMR : 01926 : 02303
Read ID

@*SFHMR* : 01926 : 02303

Read ID

GAGACCCGCATCGCCATCTTGCGCAAGAAAGCACAGATG

DNA sequence

Fastq file format

@*SFHMR* : 01926 : 02303

Read ID

GAGACCCGCATCGCCATCTTGCGCAAGAAAGCACAGATG

DNA sequence

+

A plus

Fastq file format

@*SFHMR* : 01926 : 02303

Read ID

GAGACCCGCATCGCCATCTTGCGCAAGAAAGCACAGATG

DNA sequence

+

A plus

?? >> 3*D* <; < @@*BB*9??@; 6 > 999?9*CC* : *DCB* >>? *BBBB* <

Nucleotide qualities

Fastq file format

Quality

Dna sequence G A G A C C C G C A T C
Quality ? ? > > 3 D < ; < @ @ B

Nucleotide G has a quality of ?

Deciphering of quality "?"

Symbol \rightarrow Encoded value \rightarrow Quality score \rightarrow Probability of error

Deciphering of quality "?"

Symbol \rightarrow Encoded value

Deciphering of quality "?"

Symbol → Encoded value

<http://www.ascii-code.com/>

ASCII printable characters (character code 32-127)

Codes 32-127 are common for all the different variations of the ASCII table, they are called printable characters, represent letters, digits, punctuation marks, and a few miscellaneous symbols. You will find almost every character on your keyboard. Character 127 represents the command DEL.

DEC	OCT	HEX	BIN	Symbol	HTML Number	HTML Name	Description
32	040	20	00100000		 		Space
33	041	21	00100001	!	!		Exclamation mark
34	042	22	00100010	"	"	"	Double quotes (or speech marks)
35	043	23	00100011	#	#		Number
62	076	3E	00111110	>	>	>	Greater than (or close angled bracket)
63	077	3F	00111111	?	?		Question mark
64	100	40	01000000	@	@		At symbol

Deciphering of quality "?"

Encoded value → Quality score

By looking at ascii table we found out that symbol ? is encoded as a decimal value 63 Symbols used for quality score encoding (Sanger) start with symbol ! (encoded as decimal value 33) To obtain the quality score (also known as Phred quality score) we need to subtract smallest possible value (33) from the obtained value 63. $63 - 33 = 30$

Deciphering of quality "?"

Quality score \rightarrow Probability of error

$$p = 10^{-\frac{Q}{10}}$$

Our $Q = 30$, therefore

$$p = 10^{-\frac{30}{10}} = 10^{-3} = 0.001$$

Probability that nucleotide G is an error = 0.001

Fastq quality control

Assessment of quality

<http://www.bioinformatics.babraham.ac.uk/projects/fastqc/>

Fastq quality control

Trimming and filtering of reads

<http://www.bioinformatics.babraham.ac.uk/projects/fastqc/>

Fastq quality control

Repeated assessment of quality

<http://www.bioinformatics.babraham.ac.uk/projects/fastqc/>

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:4EH>;A;?>@?7?CBBCLC=BCB<@>;+:;:;>B<>>66*:66;;>D?7/<;4<;99525.(/(66/)445;5*99;==7;,:AAGCBB=B=F?B=FB@FMD>
@HLUIT:03001:01228
AACACATTCTGACCAGAAATGCTGTTATTAATTTGGTCATAAAAATTTCATGATAATGCCAACAAAAATAAGTAAGGCATGTTGTTATGGATTGACCAACCGTATTTCATCAATATACTCGTGGG
+
255--'-':569;/:;@5@CCD>8163-8+>@4<;94388*8884::8=863)449<59584*9999;;A?@;C=CB BBB<B?9?7B<?79BB@A<A;AAAA;AABBB<;:BAA<??@3??
@HLUIT:03099:01750
TATAAAATTCATGATAAATGCCAACAAAAATAAGTAAGGCATGTTGTTATGGATTGACCAACCGTATTTCATCAATATACTGTGGGGCTTGACATTTCGATGTCATTGATTGAGCCCTATAGATGGCCCTCTCGTGGTTTATTGCATTGGCTCCAGA
AACTGCCCATAGTTGGAGCAGACTTCGGAATTATTACATCAGCATCAATTTAGAACAATTTGCGATCCATCAACGTGGTTATAGCTTAAATCTAGACC
+
BB4FEFM?DCC@BB9A9B@79C=DC4ICCCA:@?2:3FCIDB>B8?>;2::5@:999B<@;@@>7?@A@A<EE::.A@>./.@A@>9?::==A>DCDBCCCF?>A<BCB=B9B@?>>7652.(...//<8;66(5555/55:5'-1'-1'-8--(
66663(337:880.(//;<=>7:2999?::?><B@@AAAFCCBC?3:::@;@A:AAA@CE>BCCC>CCFB@B<BBBA?@?F;F@>>DDCCD>E
@HLUIT:03375:01864
GTAAATTACCTCGGTAACAGGAGCTTCAGATTTTAGATCTTAGGCAGCATCACGGCGGTATAAGCTACTTATTCGATTTAATTTGGACCGATTGCACCTATAACACATTTCTGACCCGAAATGCTGTTATTAATTTGGTCATAAAAATTCATGATAAT
GCCAACAAAAATAAGTAAGGCATGTTGTTATGGATTGACCAACCGTATTTCATCAATATACTGTGGGGCTTGACATTTCGATGTCATTGATTGAGCCCTATAGATGGCCCTCTCGTGGTTTAT
+
558>5?@?:@?>>8>9475.(./-/44<<5(88854767/886??>==;;;:@@9??75??@?>;4;7?>?6HA:A@::;6A@9>;9.'..99=8?::7(88/?BCG9EEC9CA@@>6==;>4>>6>8A@A9A3AA@?:C?@AB@9AAA
:AB>E;A:::;;@AB=<>NEFMC<C@>CC;7::<BCAB<B;B<C@9699CC=BBCFFFFB9DFAB<C?>>AC>BCCFDABB<DCB;BBC;@6@==?BBB@A;??7?ABA@A;A5AABBB
@HLUIT:01436:01892
AAAAATGCCAACAAAAATAAGTAAGGCATGTTGTTATGGATTGACCAACCGTATTTCATCAATATACTGTGGGGCTTGACATTTCGATGTCATTGATTGAGCCCTATAGATGGCCCTCTCGTGGTTTATTGCATGGCTTCCAGAAACTGCCCATAGTTGG
GAGCAGACTTCGGAATTATTACATCAGCATCAATTTAGAACAATTTGCGATCCATCAACGTGGTTATAGCTTAAATGCTAGACCACAAAGCCGCTTTCATTACGCTTTA
+
AA9::;9@:::;*:@?:@?:?9FFDK?FC@BCC=BCCABBB>D@D=BBBC>CCEDD=BBBCBEBBAAA5AA<C>CBBC=BDCCC@A?>:><6<99/48068<566?E<ED4@?<?97-666'9B9@?B=82788-:3599:1@A@@3>?>77.44
'1115557244-1-1389588@A=;<:???:@7A??:ABAA;BA=?@394;<;>:=>5=8;888688-44414111.3456-56145:>29--'-46:5C
@HLUIT:03184:02349
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CATCTCGGAATTATTACATCAGCATCAATTTAGAACAATTTGCGATCCATCAACGTGGTTATAGCTTAAATGCTAGACCACAAAGCCGCTTTCATTACGCTTTA
+
@?<B>;@@?2D>BBB<B=BCDDC=CC=AAA9?7B;CDE=D=B<C>>>CBBAE<CBB>><;94667'7=6B=CEJG=CCCCDHBC=BCF@F?>?<@C:DDCCCB=CF=A@>ECC;AAA6AC=CECC=BCF@H=?@B;BCBA9CCCAA;AE;CBC
BBBBB;AB;C;B;AA;ACCC@C@BDDC=CH<C?>@FFF;AACCC?CCCB<AAAA<A;??::CCCC8@BDBDEC;CAAA5?@::;@F:AAA;@@?@?@?
@HLUIT:02849:02599
AACACATAAATAAGTAAGGCATGTTGTTATGGATTGACCAACCGTATTTCATCAATATACTGTGGGGCTTGACATTTCGATGTCATTGATTGAGCCCTATAGATGGCCCTCTCGTGGTTTATTGCATGGCTTCCAGAAACTGCCCATAGTTGGAGCAGC
TTTCGGAATTATTACATCAGCATCAATTTAGAACAATTTGCGATCCATCAACGTGGTTATAGCTTAAATGCTAGACCACAAAGCCGCTTTCATTACGCTTTA
+
6>;*::::;E;3A;ACBBC>FF=CC@9?9?>::2:7A;CA>::@?@?;;>;?7BB@3DDDA;B<BB@A9@ABBBCECC=DF?9?>>7;::ACBEB=68-4-...>@<?/>=:8;>>97.>5(54-888/987;;5>><..'. '1..9A@?>
A@:@;@;AA<BBBCBBDCCB<C<CCCD?EDB<CCCBBAE>DDCB=CCCB;??9?A?ACBB;DD@>@AAB=BBB;BBC=CC@9?7?@?>@>C:1881D
@HLUIT:03305:01951
AACAAAAATAAGTAAGGCATGTTGTTATGGATTGACCAACCGTATTTCATCAATATACTGTGGGGCTTGACATTTCGATGTCATTGATTGAGCCCTATAGATGGCCCTCTCGTGGTTTATTGCATGGCTTCCAGAAACTGCCCATAGTTGGAGCAGC
TTTCGGAATTATTACATCAGCATCAATTTAGAACAATTTGCGATCCATCAACGTGGTTATAGCTTAAATGCTAGACCACAAAGCCGCTTTCATTACGCTTTA
+
9C::*::::8C::4B<CBBA=B@:AA:3::9AAA@99A;@?@?:@?>>>CB BBBB@C?73CD@<A9?7A?>?7CFBBCEA;@B<BBB<3;::?>>><69)978>>A>>8;+99857<99<4-43(77-::3(33393(779===592:9DA?A99
8CC=B@<@<AA=BBBCBDDCCCC=B:BBCA::;+999A??<BBBA<A>@<C=A>==AA@6AAA@BDBC<BBA:AAA8@BB<ECBE@BBCCC<CC<H
@HLUIT:03305:01951

```