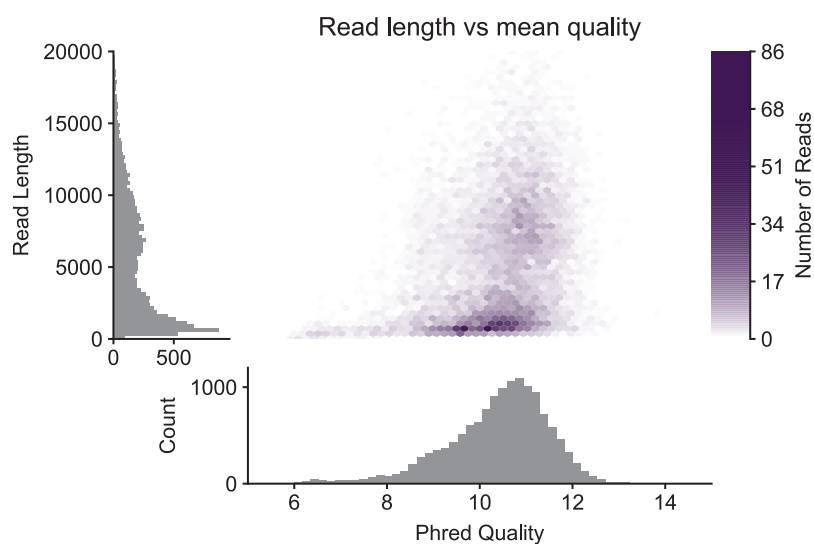
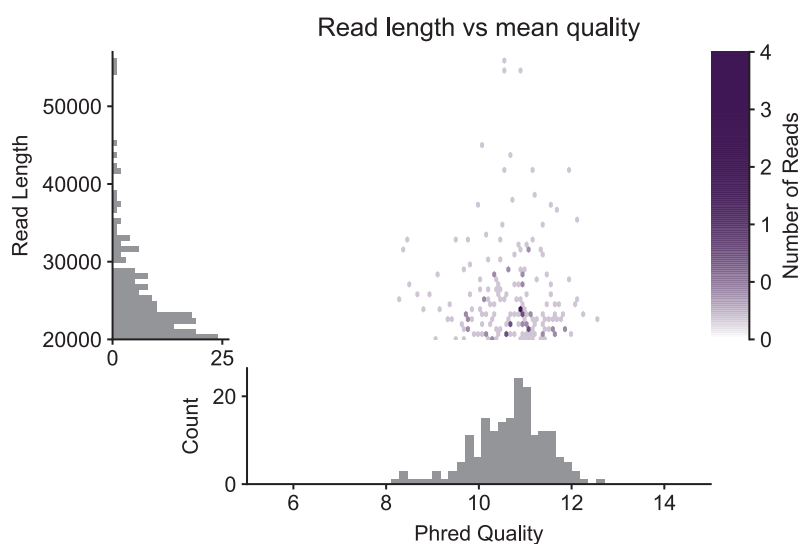


Phred Scores (Q Scores) for Sequencing Reads

Short-Read
Illumina

Paired-End Reads	% Bases Q20	% Bases Q30
R1	83.7	75.8
R2	88.8	83.3

Long-Read
MinION

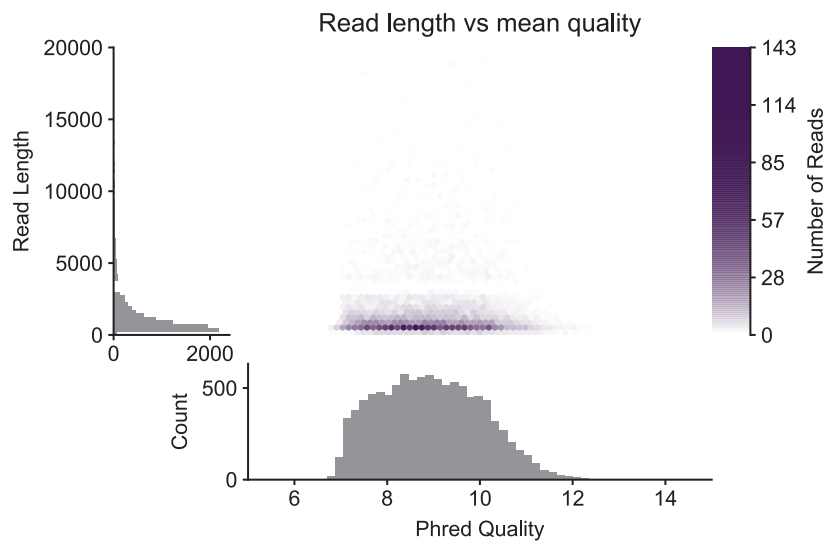
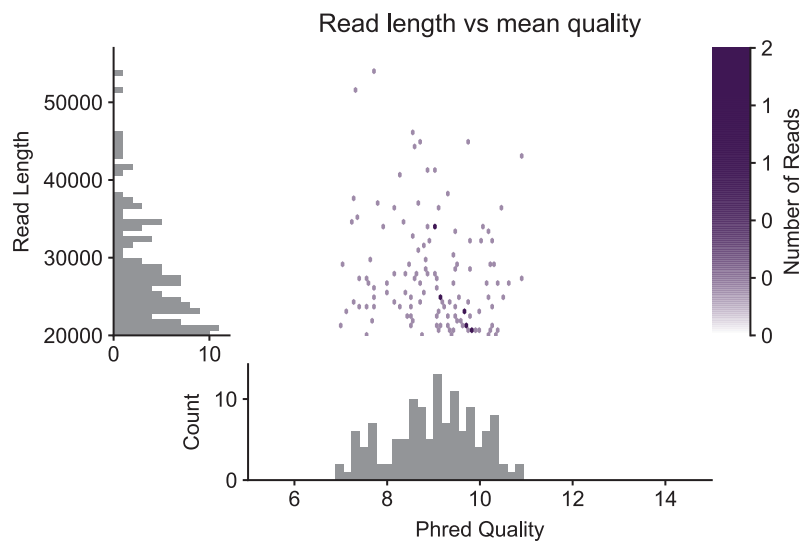


Phred Scores (Q Scores) for Sequencing Reads

Short-Read
Illumina

Paired-End Reads	% Bases Q20	% Bases Q30
R1	94.5	91.5
R2	92.4	88.4

Long-Read
MinION

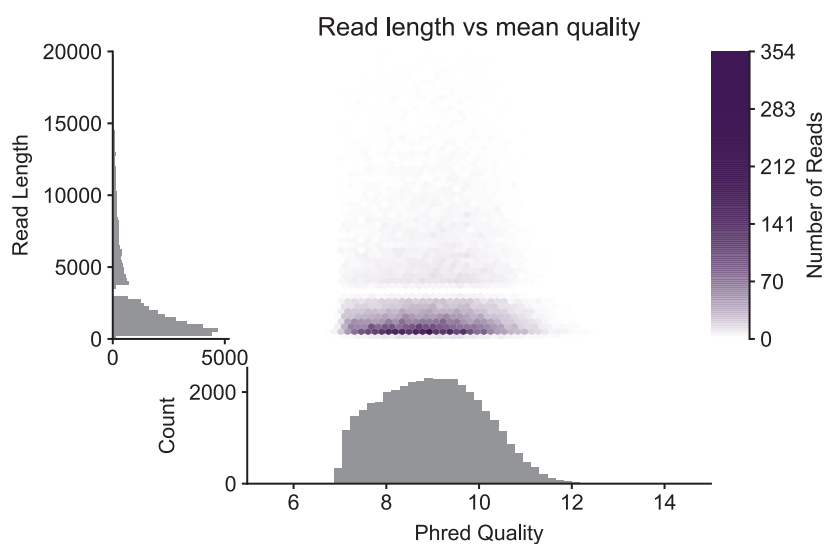
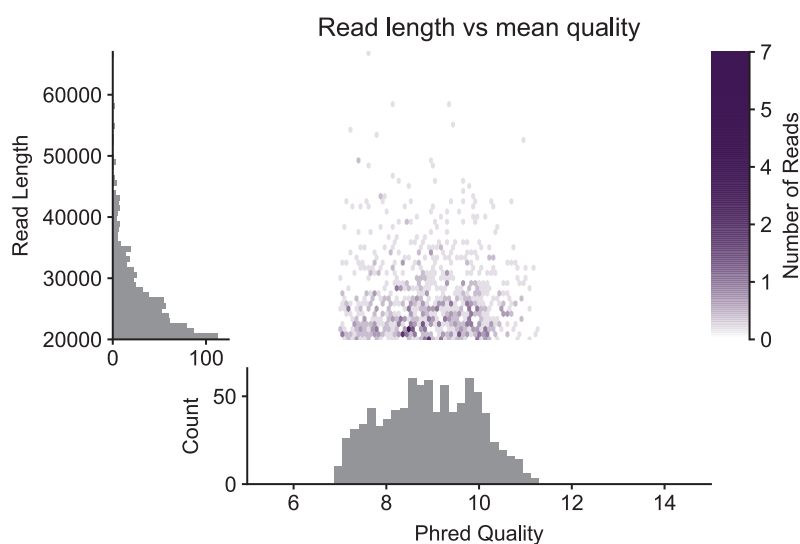


Phred Scores (Q Scores) for Sequencing Reads

Short-Read
Illumina

Paired-End Reads	% Bases Q20	% Bases Q30
R1	94.6	91.6
R2	92.8	88.9

Long-Read
MinION

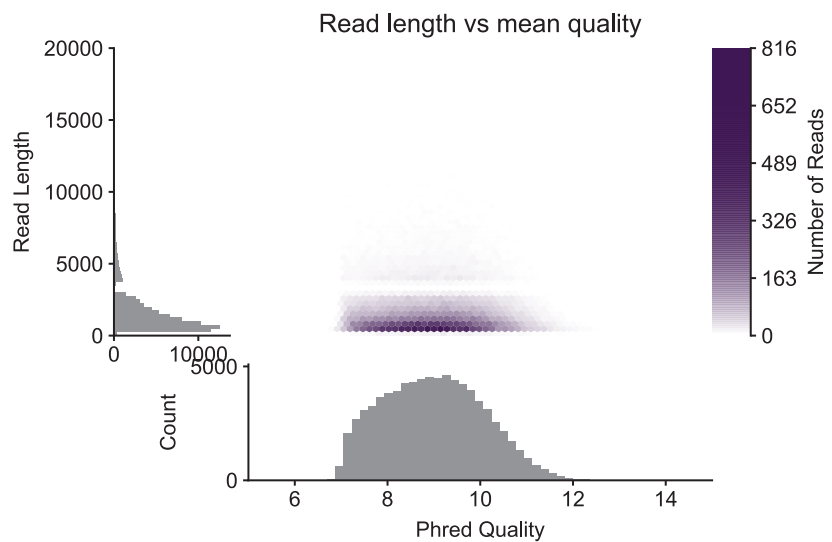
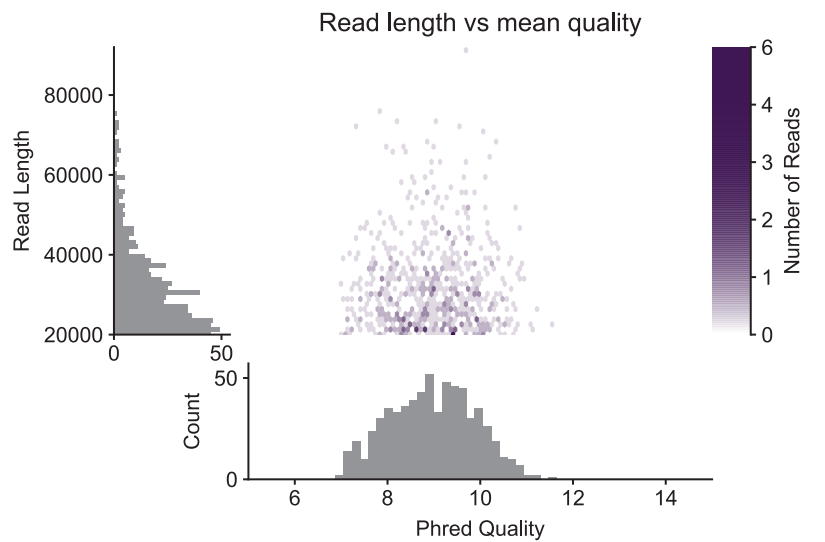


Phred Scores (Q Scores) for Sequencing Reads

Short-Read
Illumina

Paired-End Reads	% Bases Q20	% Bases Q30
R1	92.7	89.0
R2	90.3	85.7

Long-Read
MinION

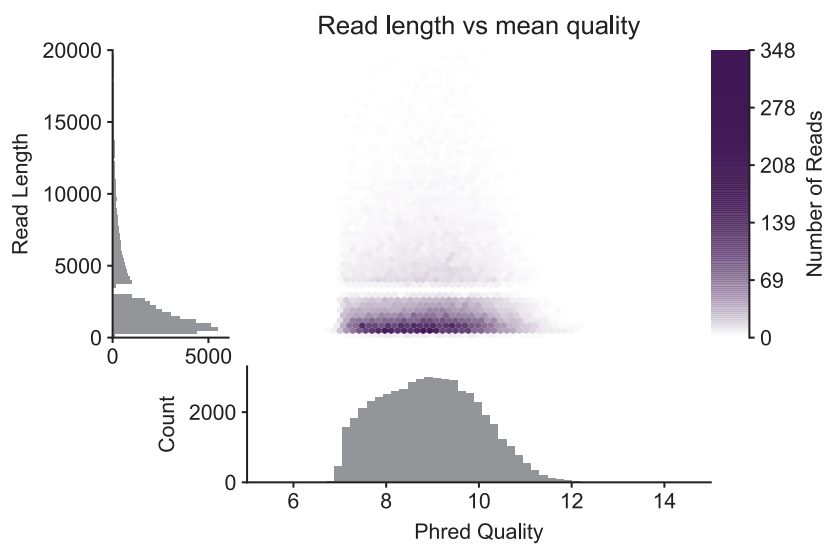
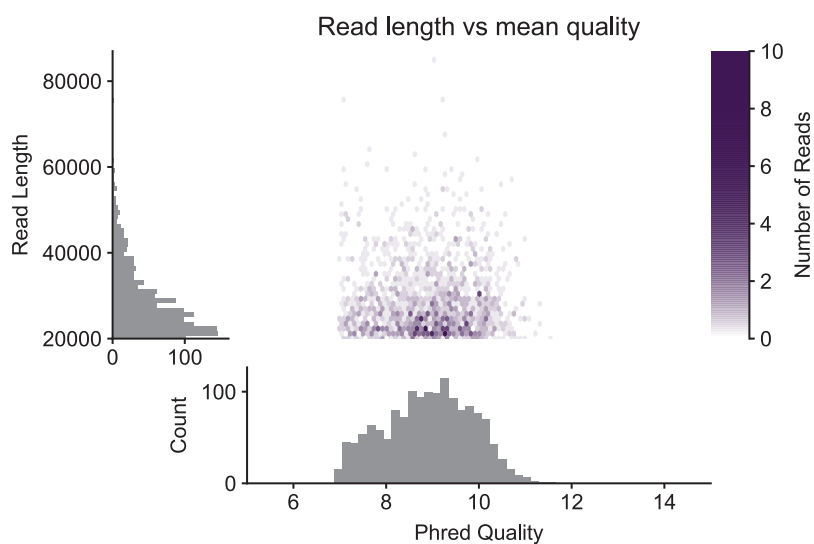


Phred Scores (Q Scores) for Sequencing Reads

Short-Read
Illumina

Paired-eEd reads	% Bases Q20	% Bases Q30
R1	94.8	92.0
R2	93.0	89.1

Long-Read
MinION

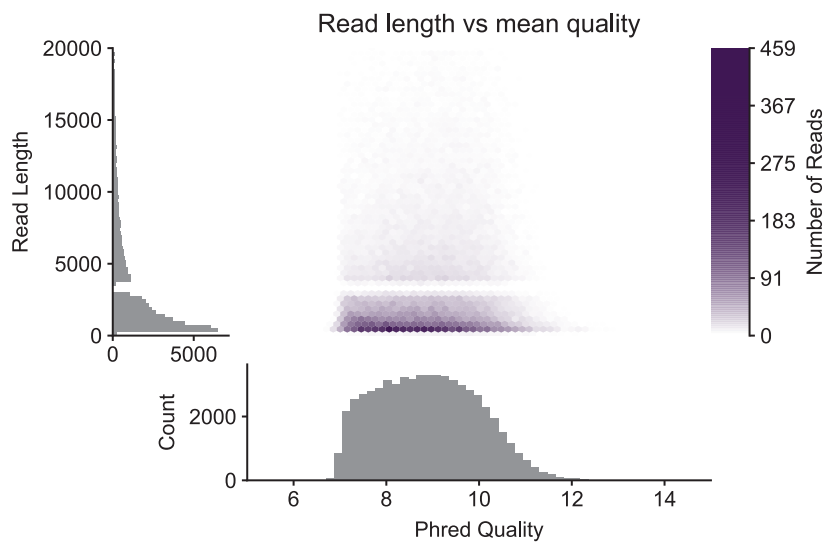
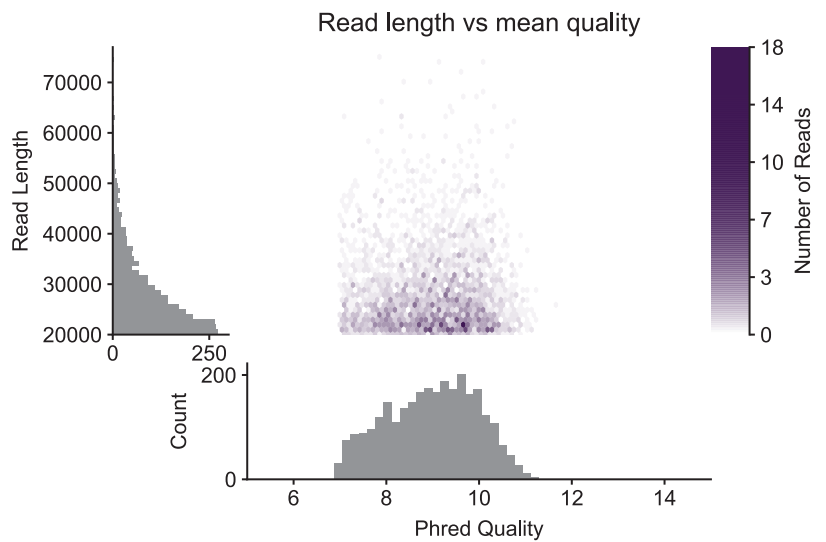


Phred Scores (Q Scores) for Sequencing Reads

Short-Read
Illumina

Paired-End reads	% Bases Q20	% Bases Q30
R1	96.0	93.5
R2	95.7	92.7

Long-Read
MinION

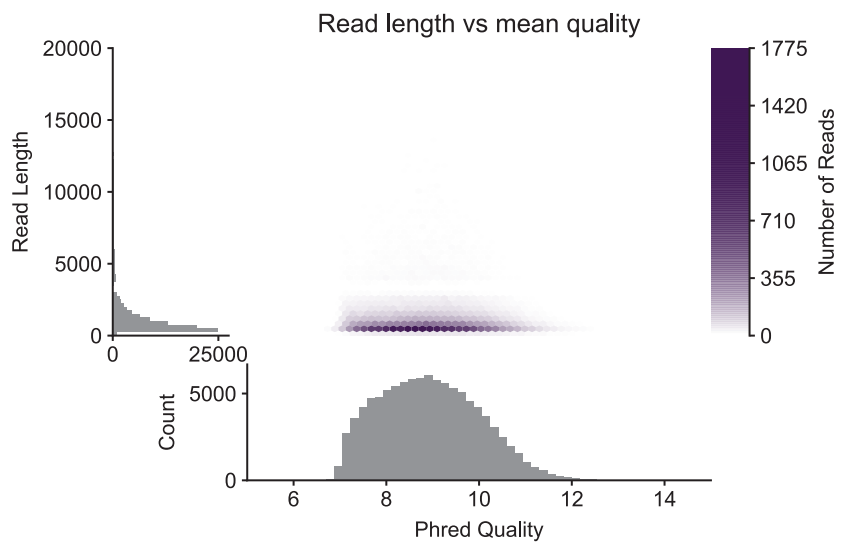
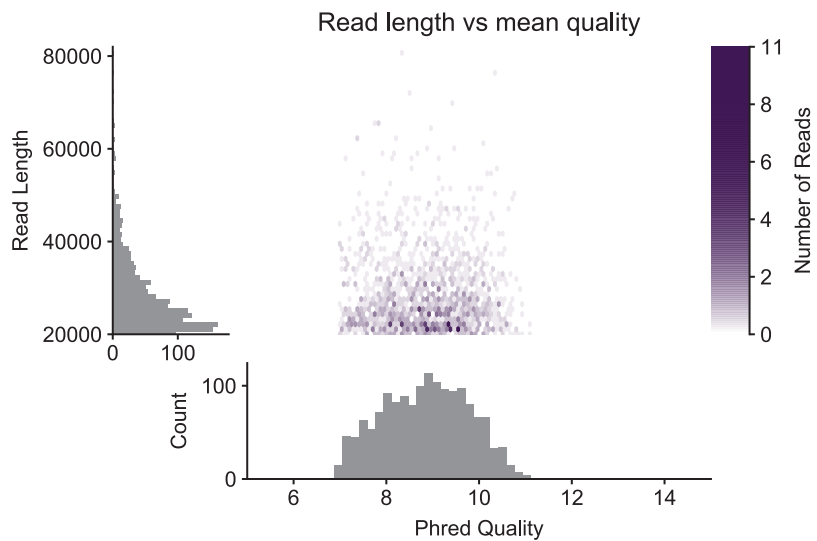


Phred Scores (Q Scores) for Sequencing Reads

Short-Read
Illumina

Paired-End Reads	% Bases Q20	% Bases Q30
R1	93.0	89.5
R2	91.0	86.5

Long-Read
MinION



Phred Scores (Q Scores) for Sequencing Reads

Short-Read
Illumina

Paired-End Reads	% Bases Q20	% Bases Q30
R1	98.3	96.0
R2	97.3	93.8

Long-Read
MinION

