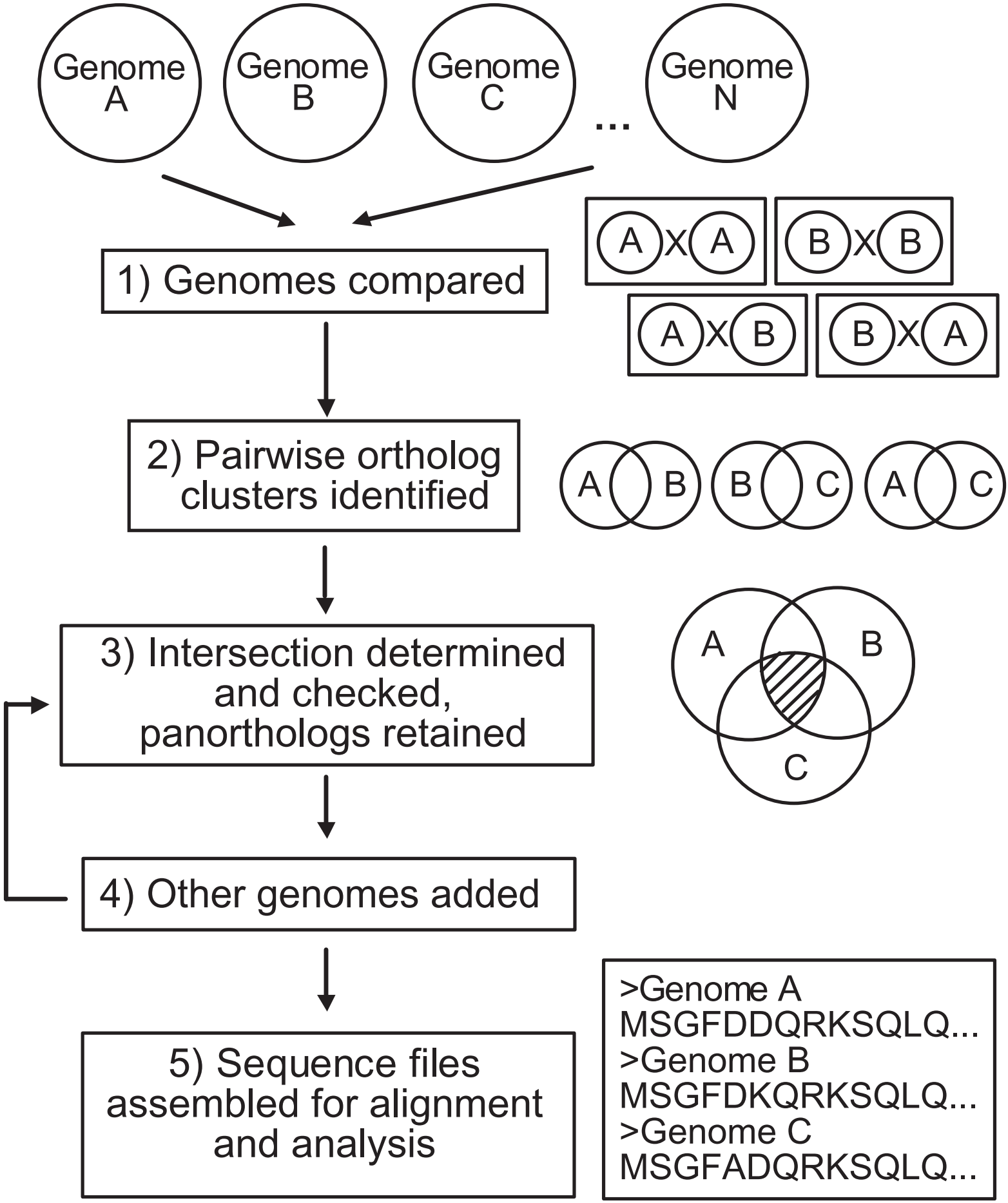
Genome A
Genome B
Genome C
...
Genome N
1) Genomes compared
A X A
B X B
A X B
B X A
2) Pairwise ortholog clusters identified
A B
B C
A C
3) Intersection determined and checked, panorthologs retained
A
B
C
4) Other genomes added
5) Sequence files assembled for alignment and analysis
>Genome A
MSGFDDQRKSQLQ...
>Genome B
MSGFDKQRKSQLQ...
>Genome C
MSGFADQRKSQLQ...