

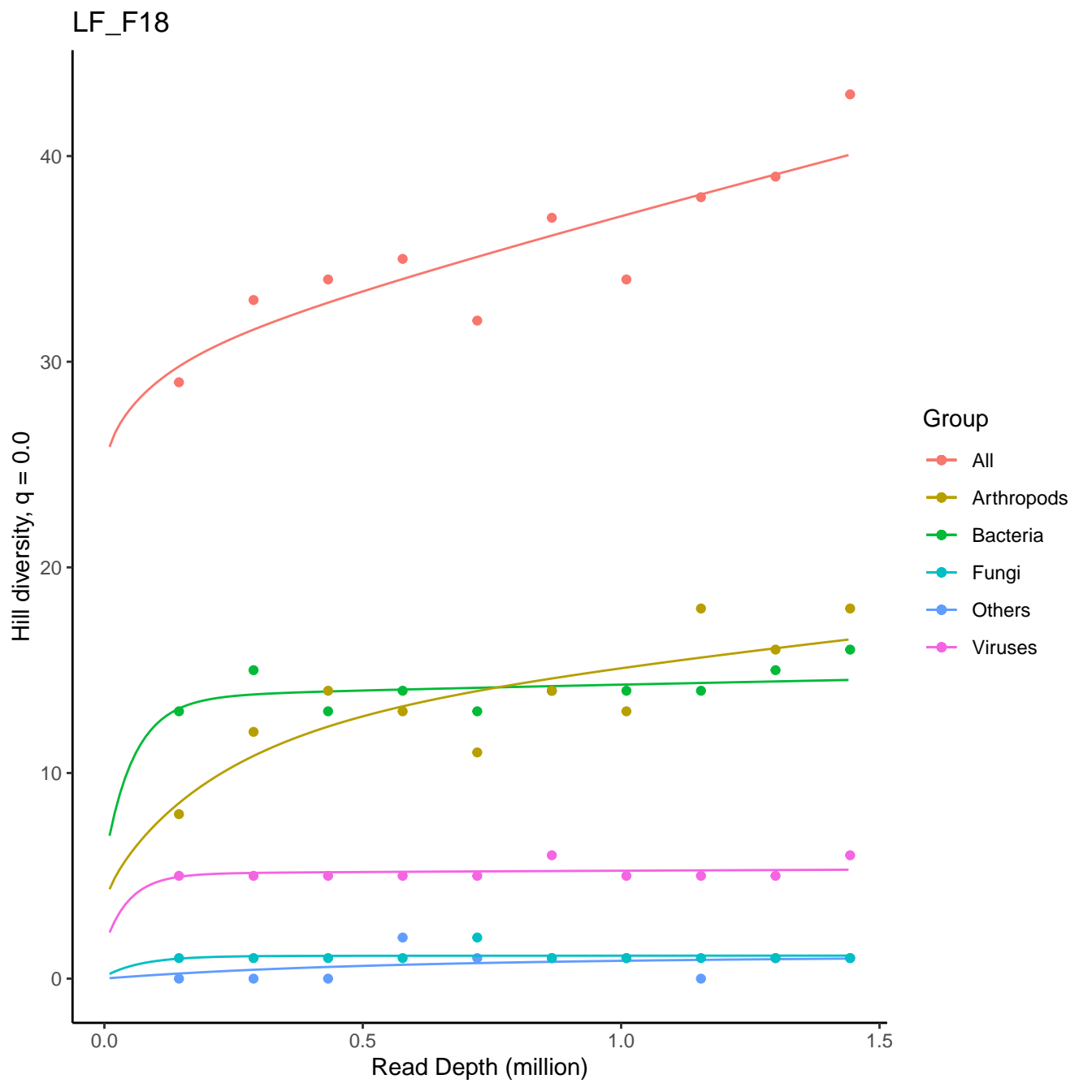
LF_F18

Hill diversity, $q = 0.0$

Group

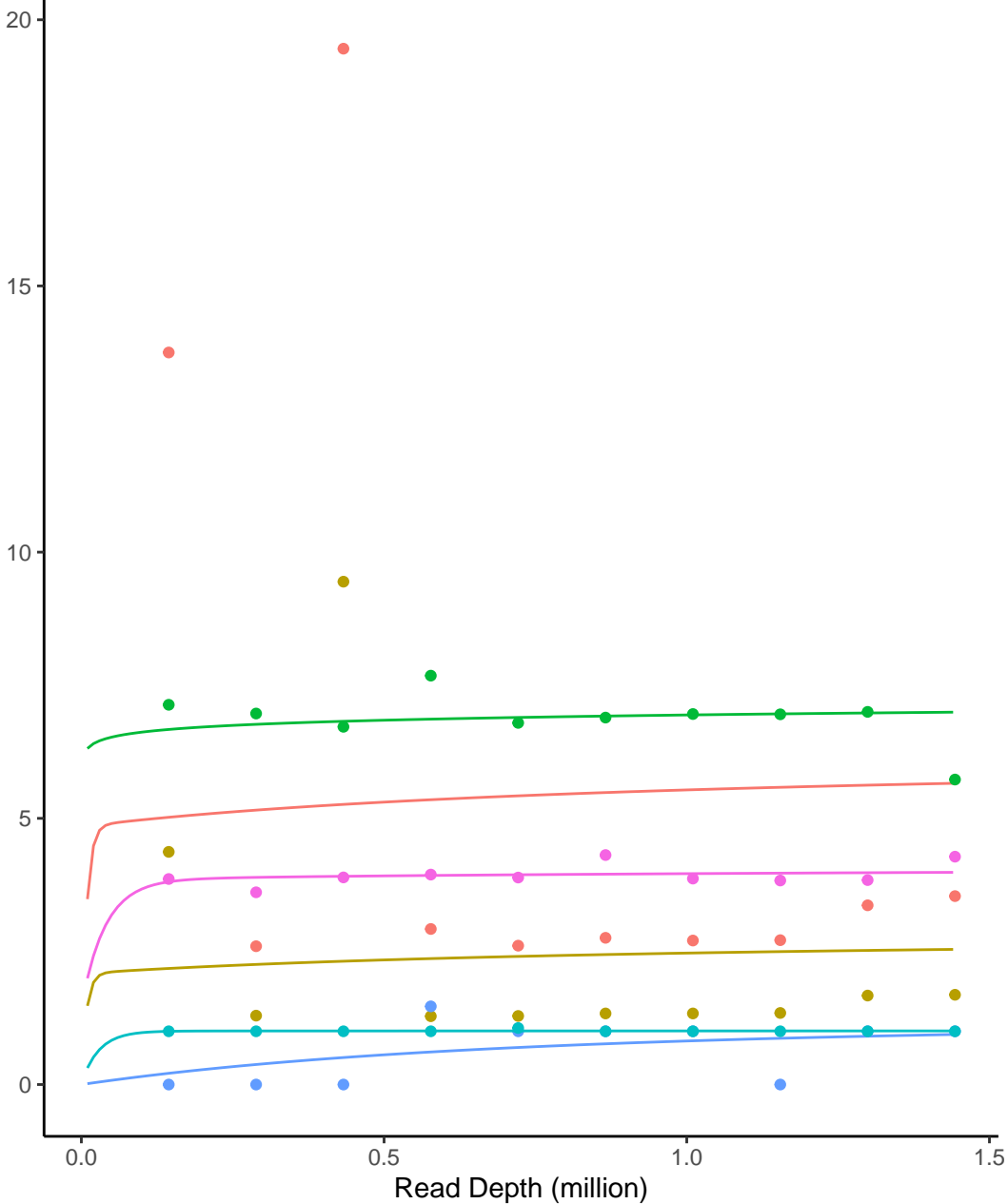
- All
- Arthropods
- Bacteria
- Fungi
- Others
- Viruses

Read Depth (million)



LF_F18

Hill diversity, $q = 1.0$



Group

- All
- Arthropods
- Bacteria
- Fungi
- Others
- Viruses

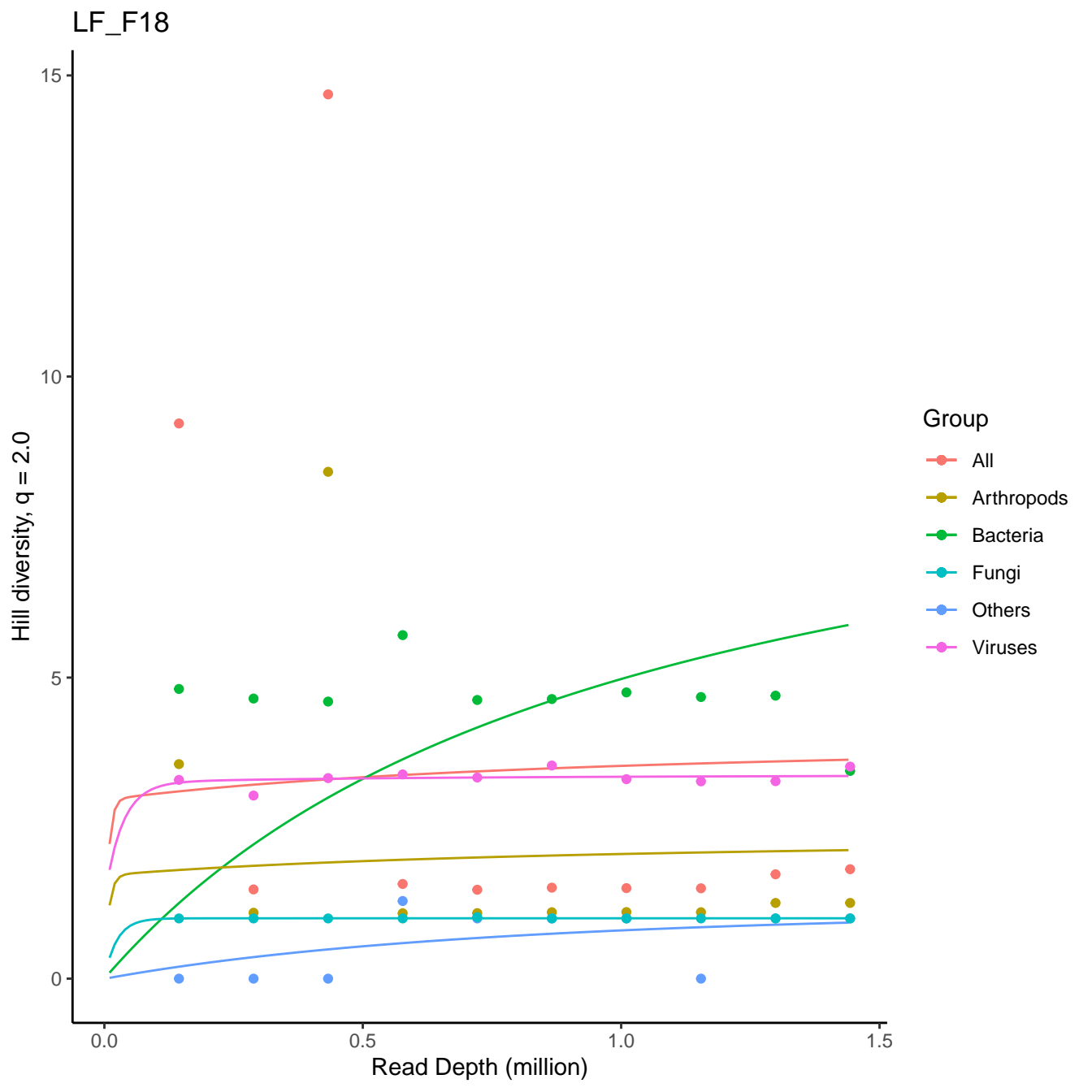
LF_F18

Hill diversity, $q = 2.0$

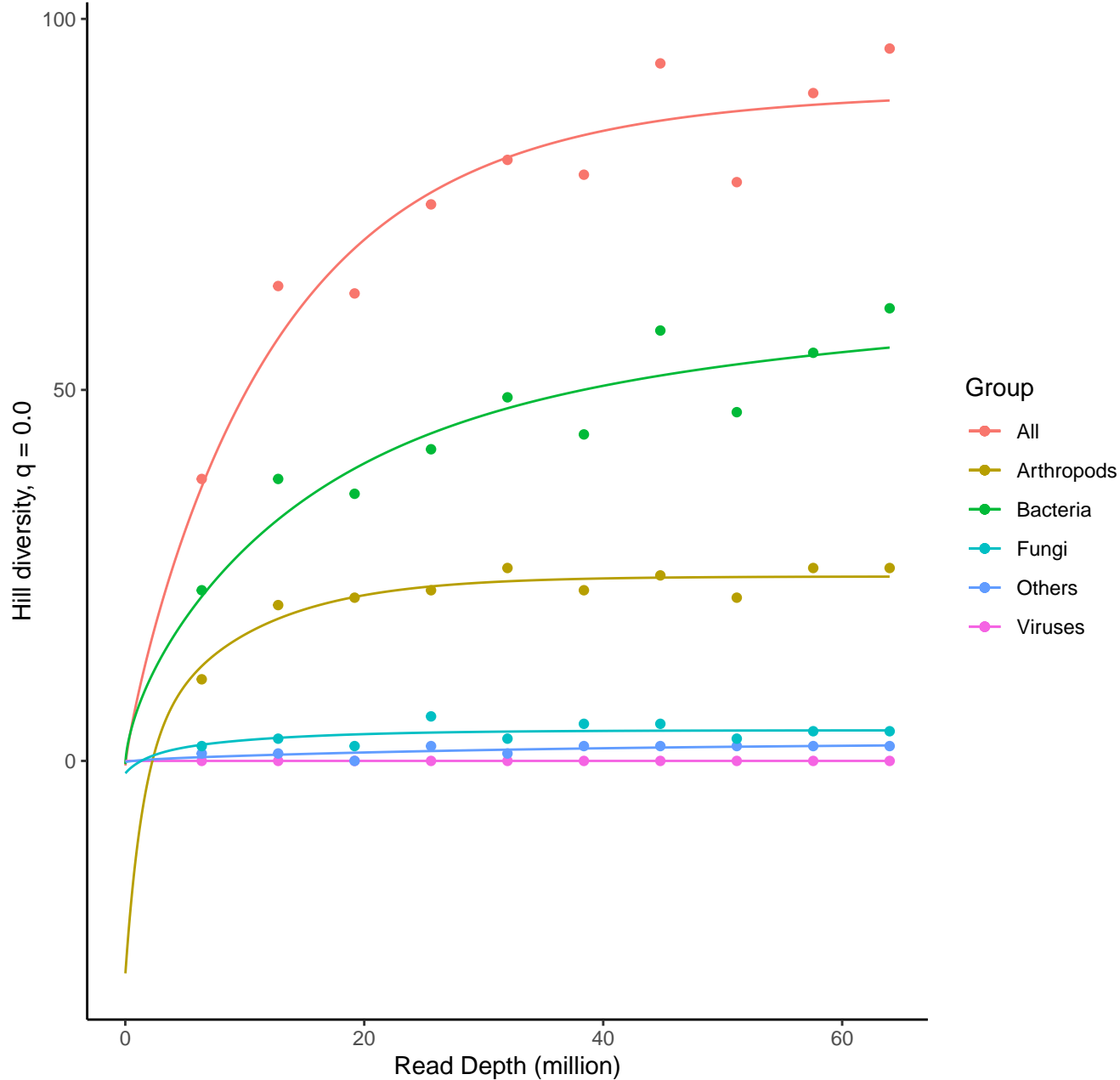
Group

- All
- Arthropods
- Bacteria
- Fungi
- Others
- Viruses

Read Depth (million)



PGcomcol3_Bimp



PGcomcol3_Bimp

Hill diversity, $q = 1.0$

Group

- All
- Arthropods
- Bacteria
- Fungi
- Others
- Viruses

Read Depth (million)

10

5

0

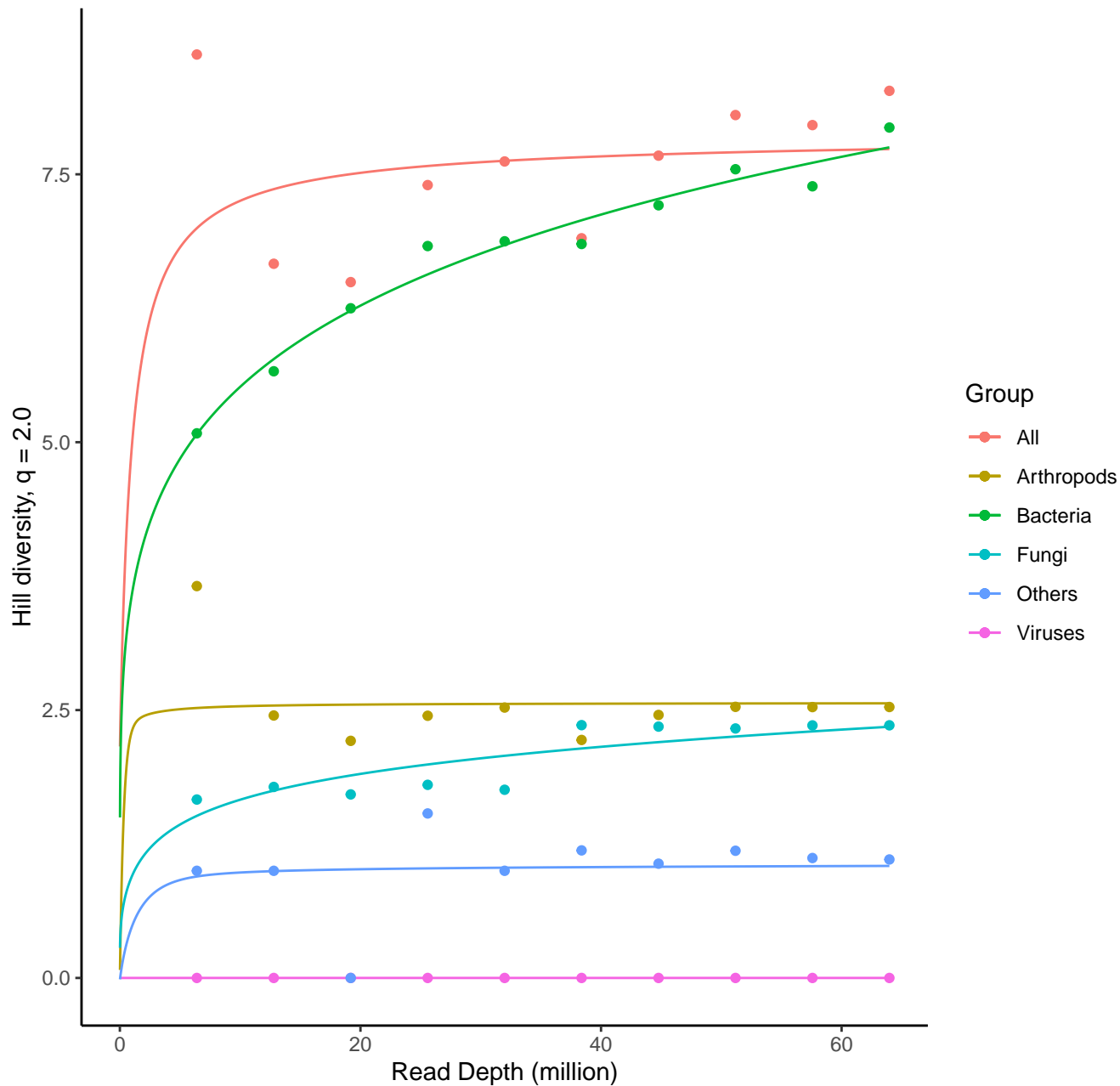
0

20

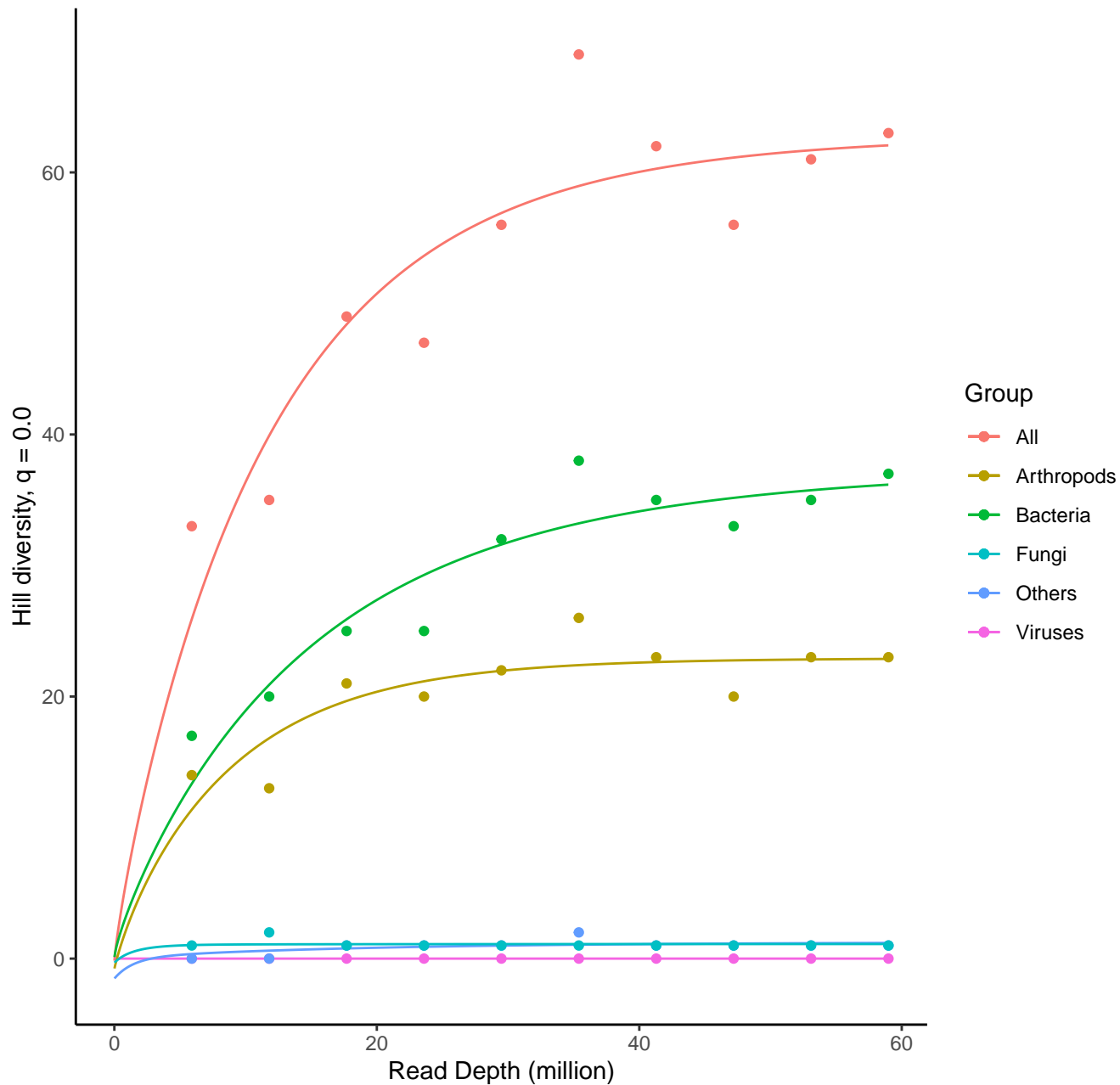
40

60

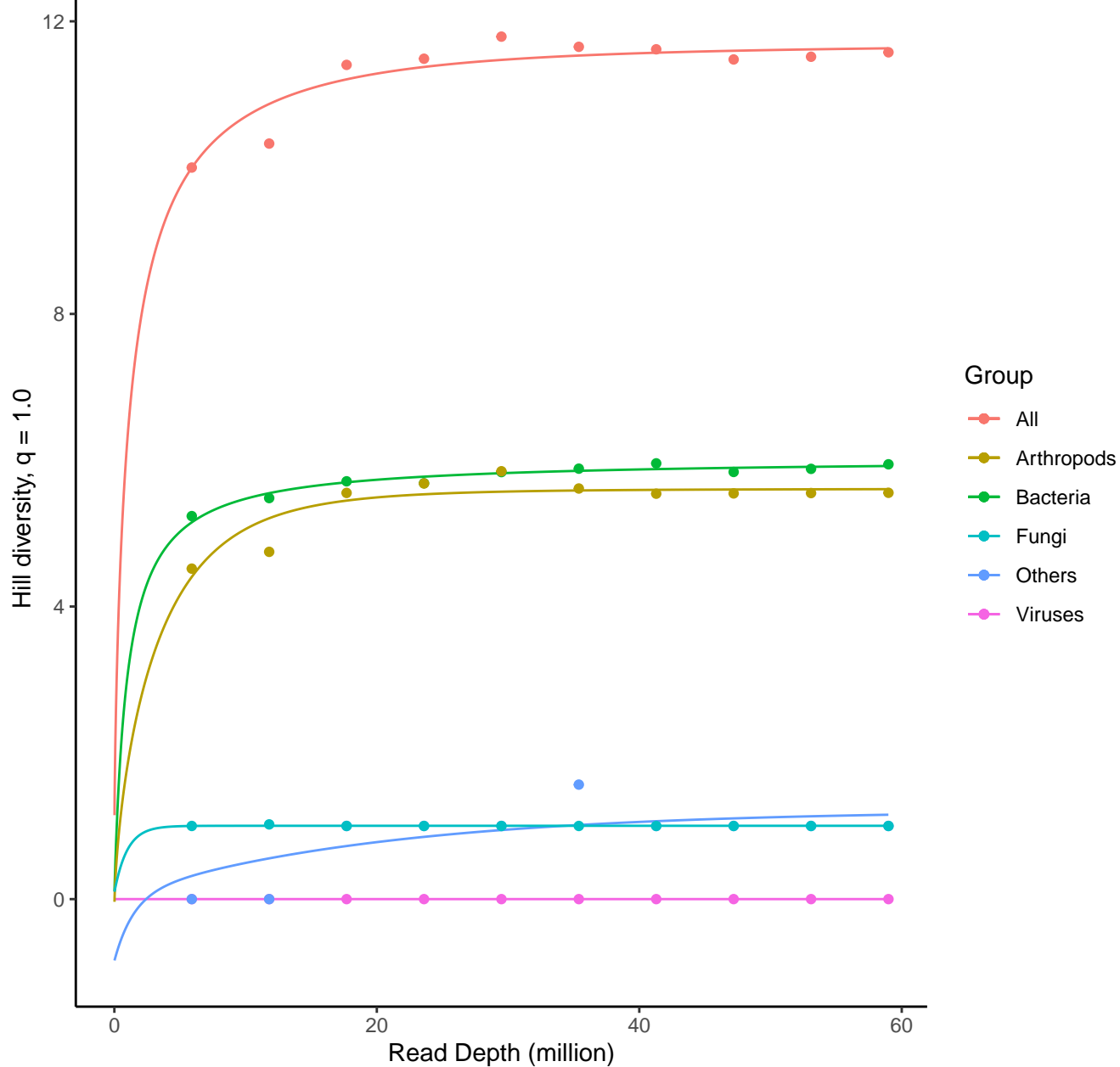
PGcomcol3_Bimp



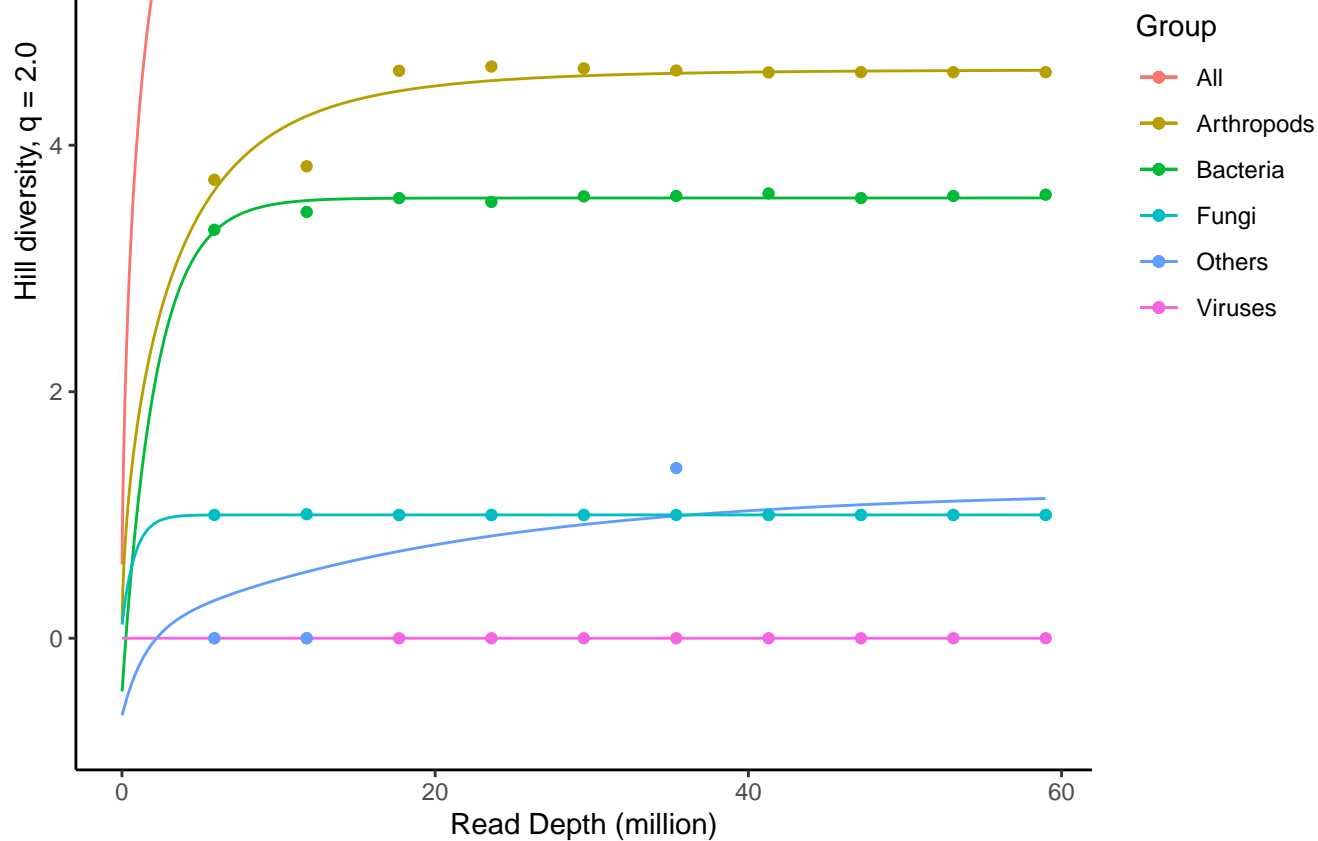
PGcomcol4a_Bimp



PGcomcol4a_Bimp

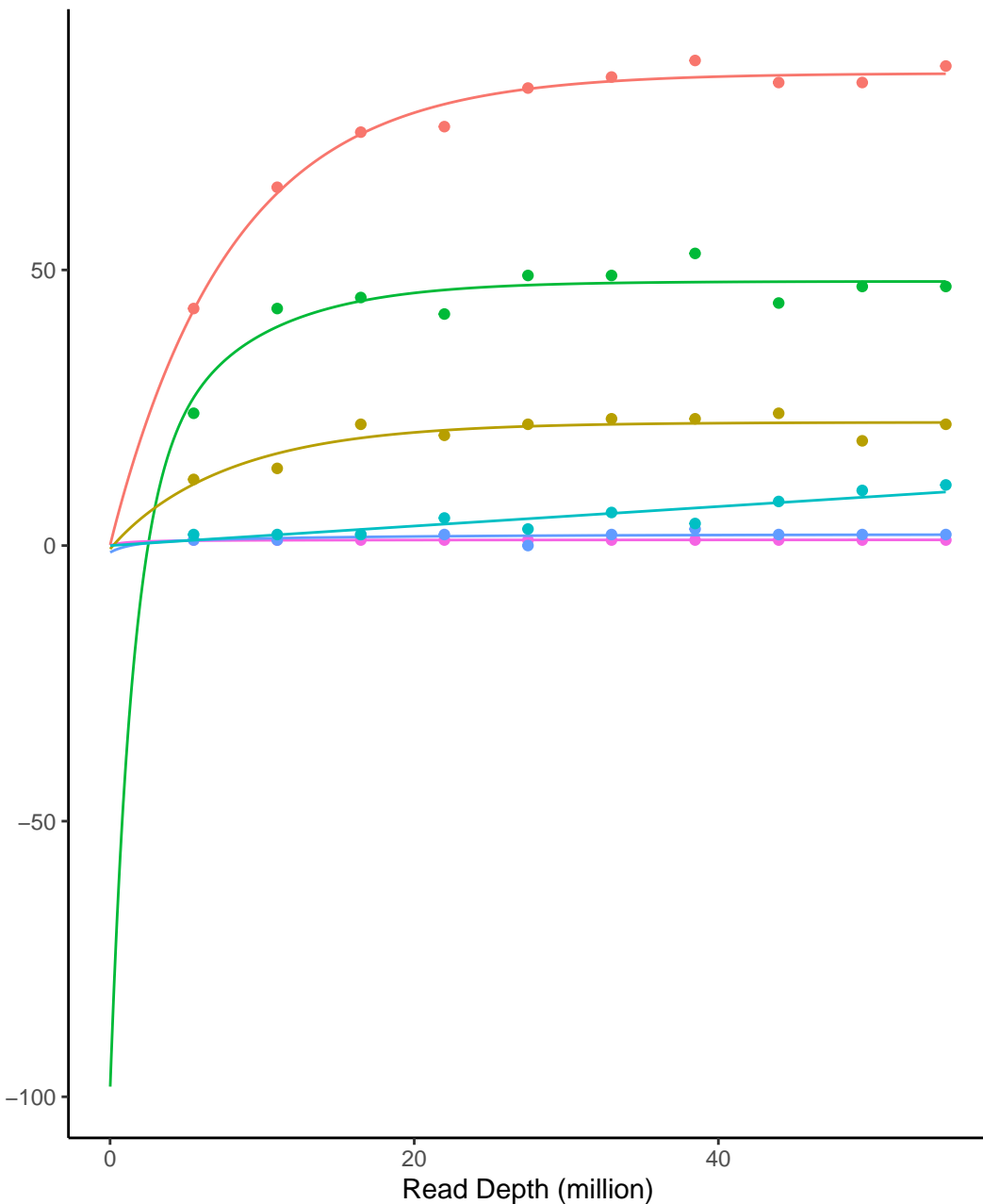


PGcomcol4a_Bimp



PGcomcol4b_Bimp

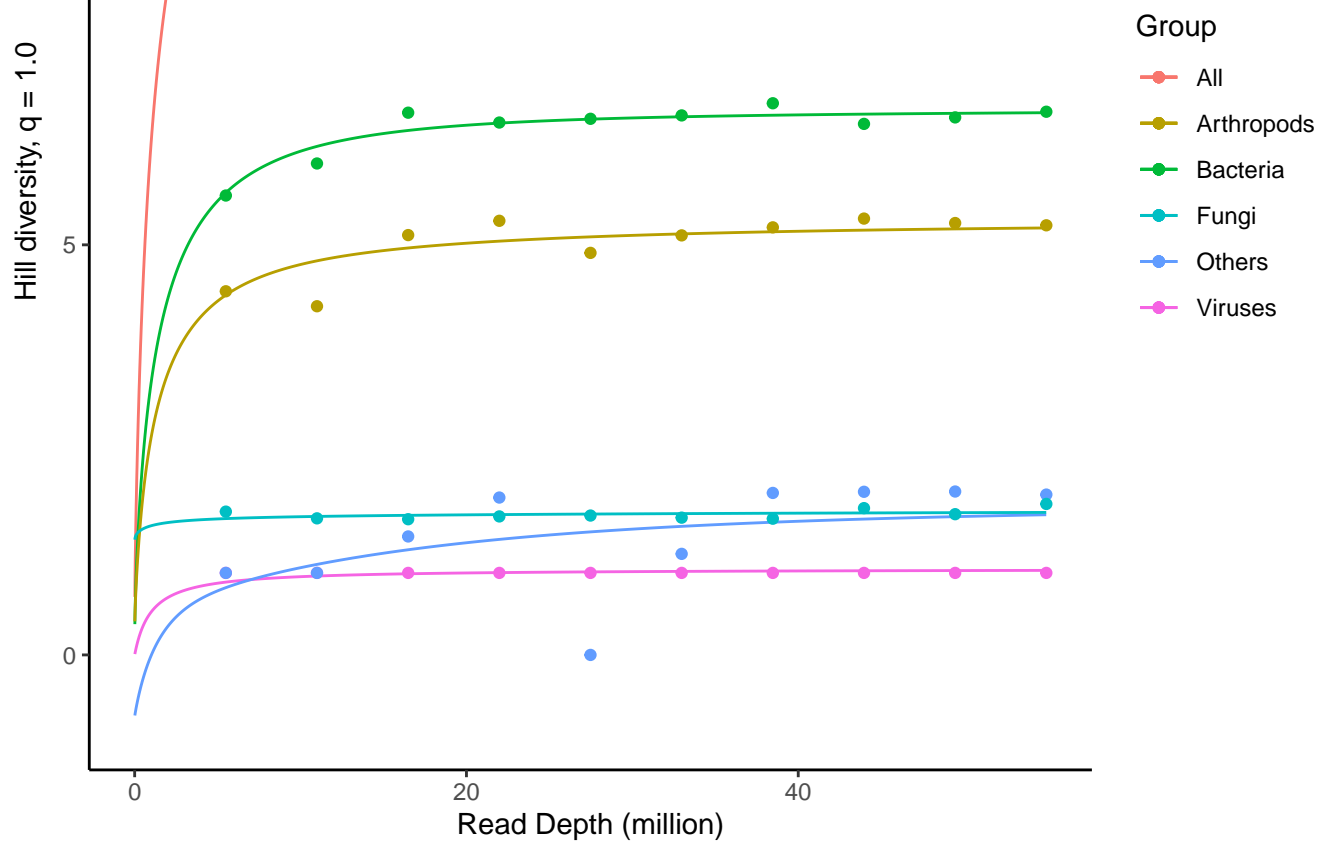
Hill diversity, $q = 0.0$



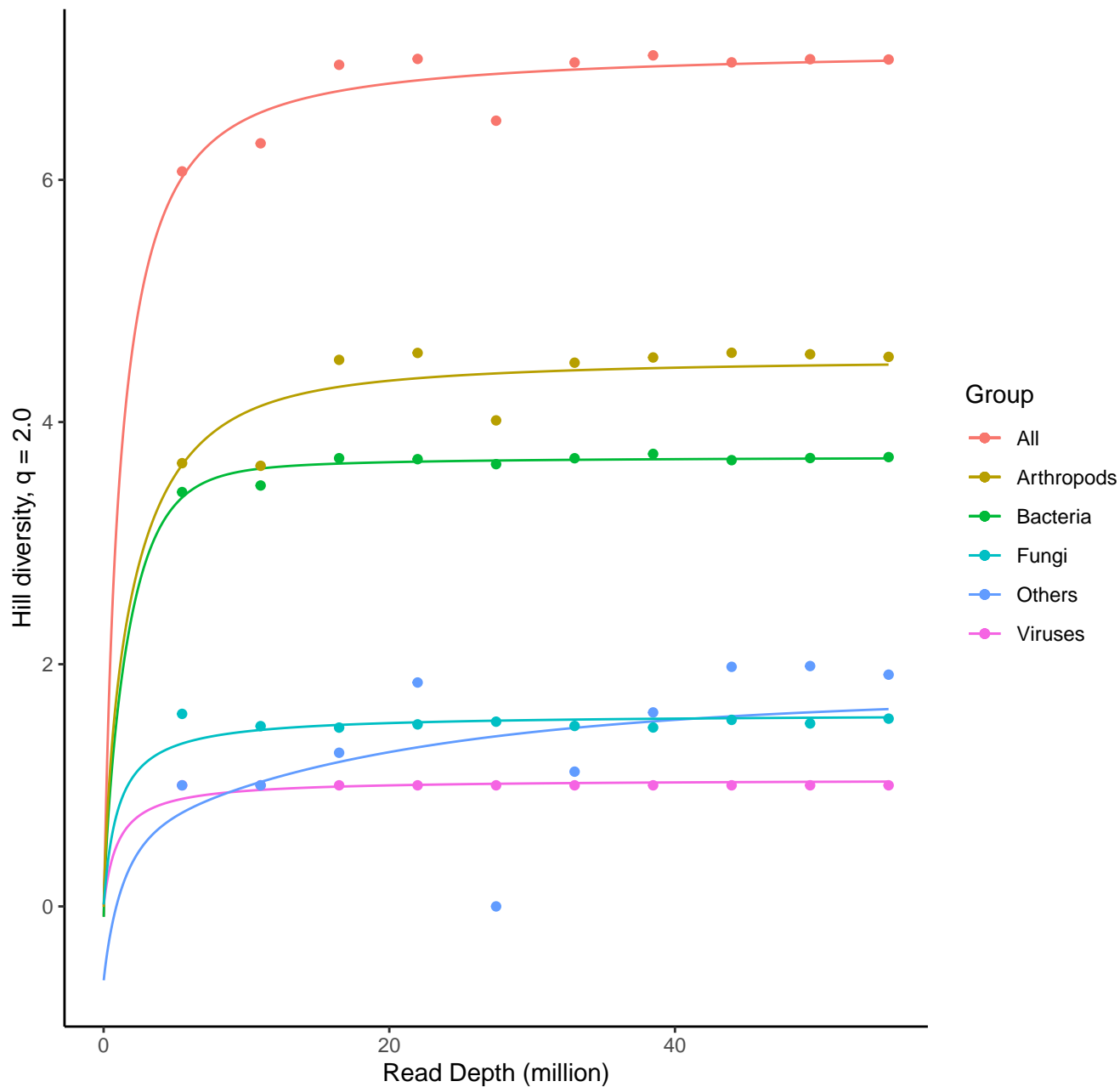
Group

- All
- Arthropods
- Bacteria
- Fungi
- Others
- Viruses

PGcomcol4b_Bimp

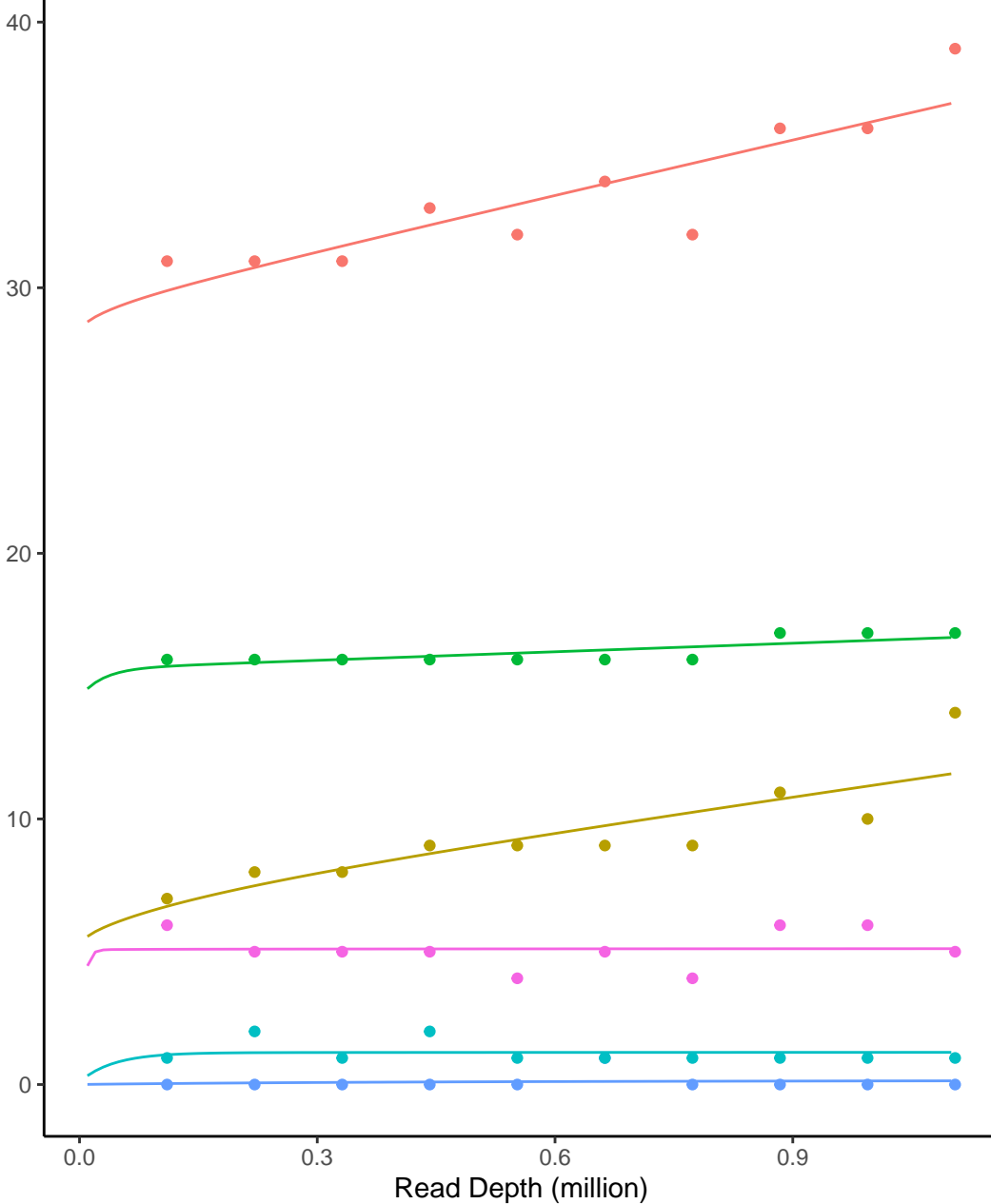


PGcomcol4b_Bimp



PGnos_high_hv13-1

Hill diversity, $q = 0.0$

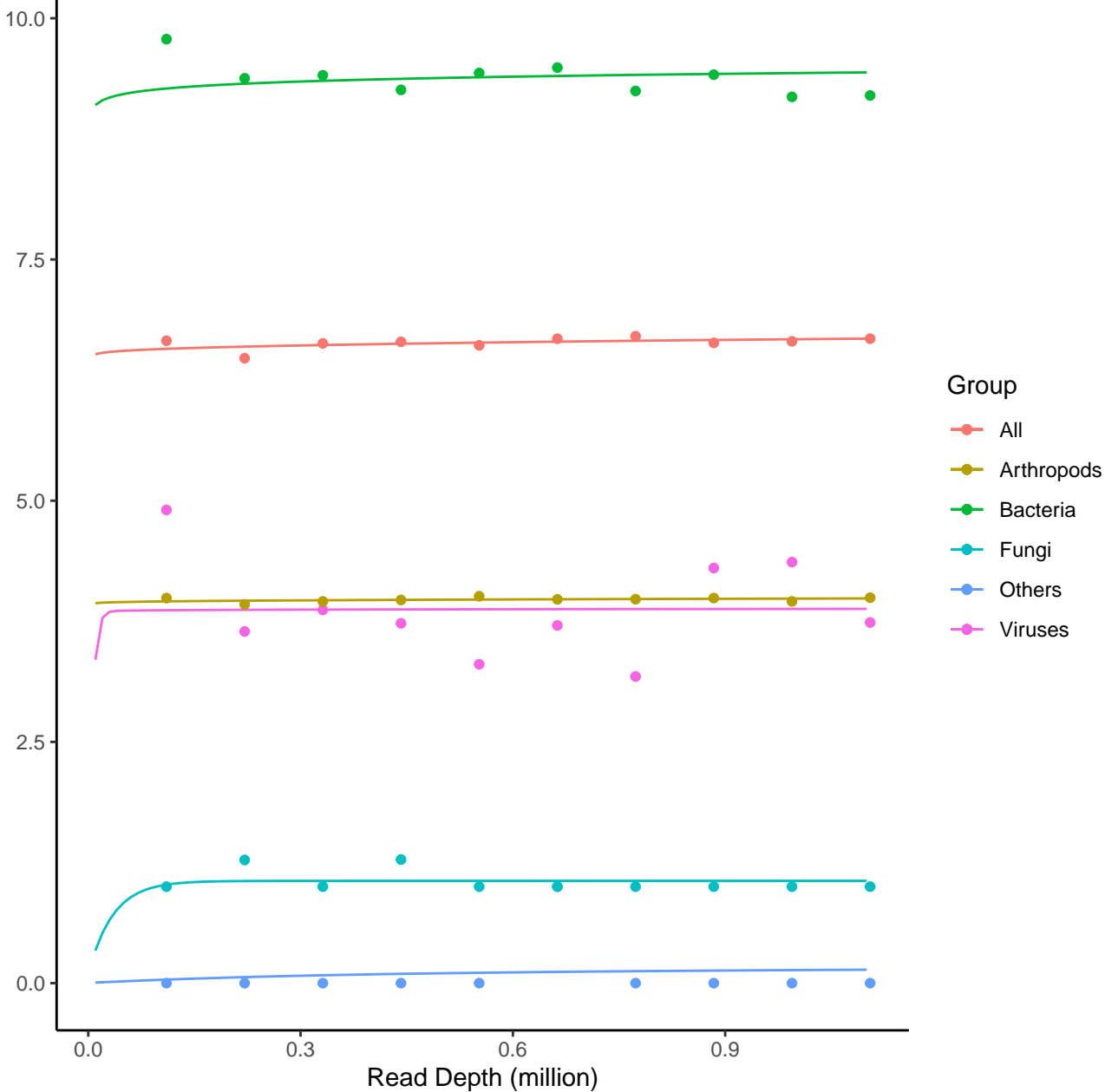


Group

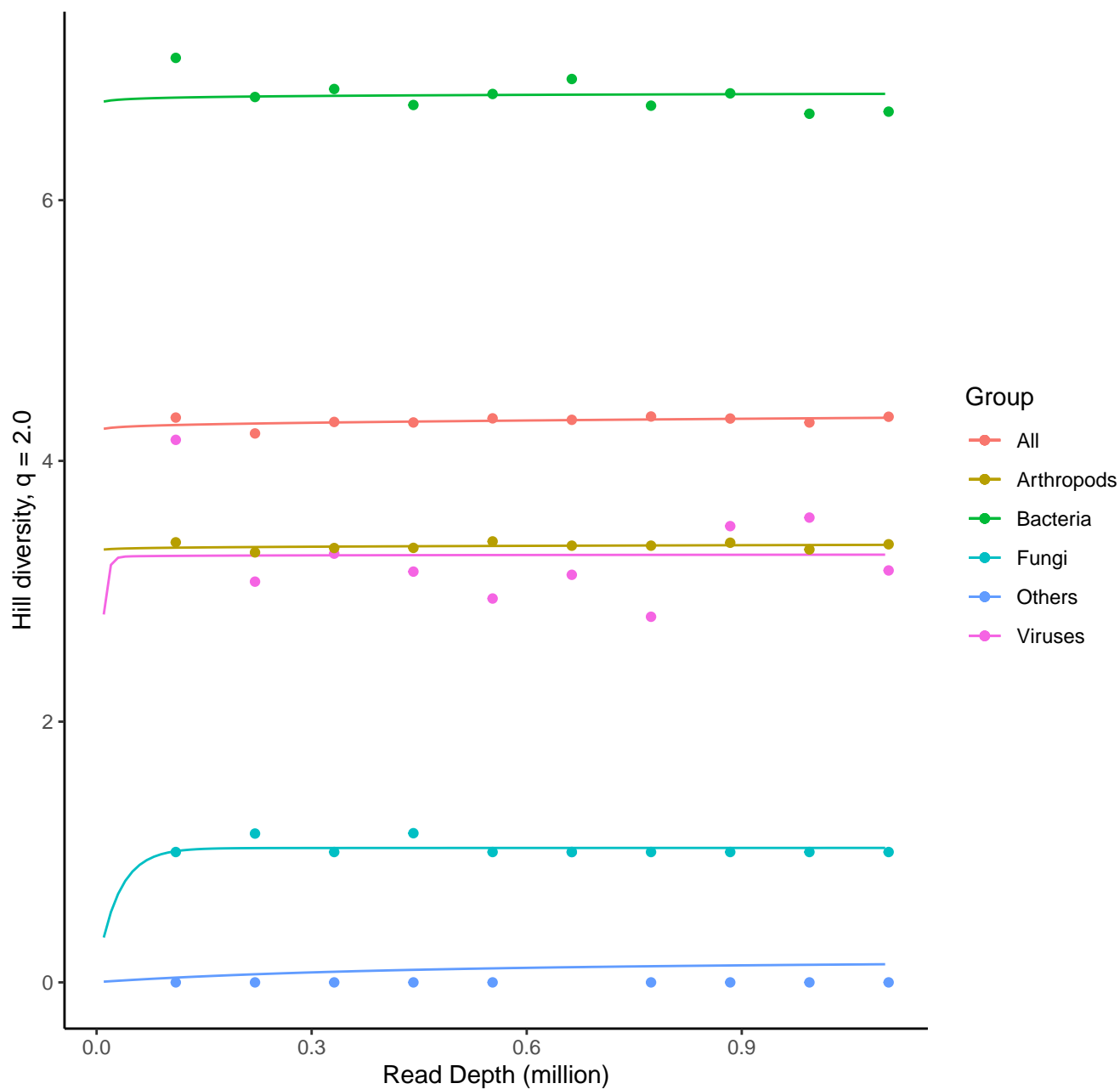
- All
- Arthropods
- Bacteria
- Fungi
- Others
- Viruses

PGnos_high_hv13-1

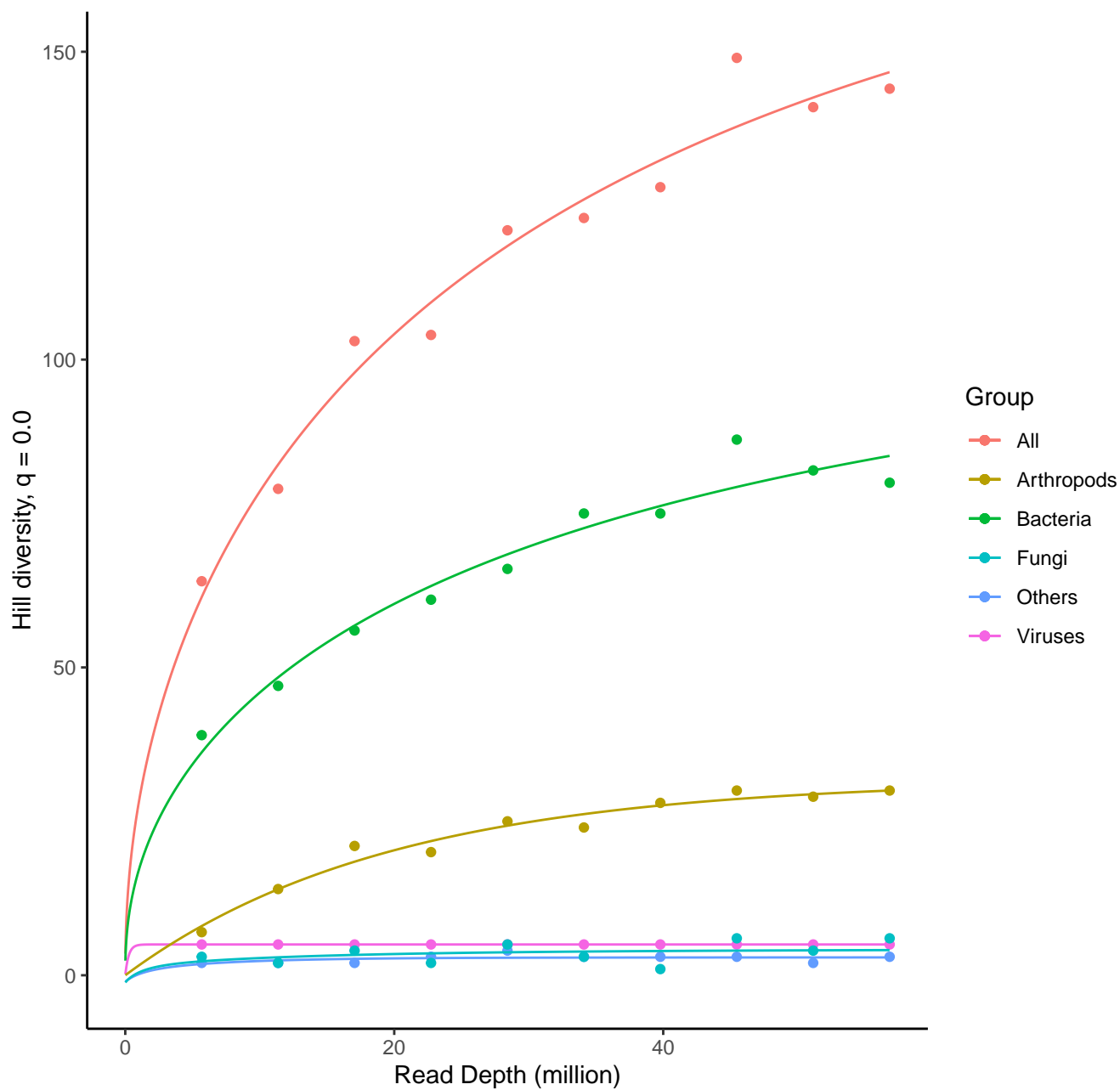
Hill diversity, $q = 1.0$

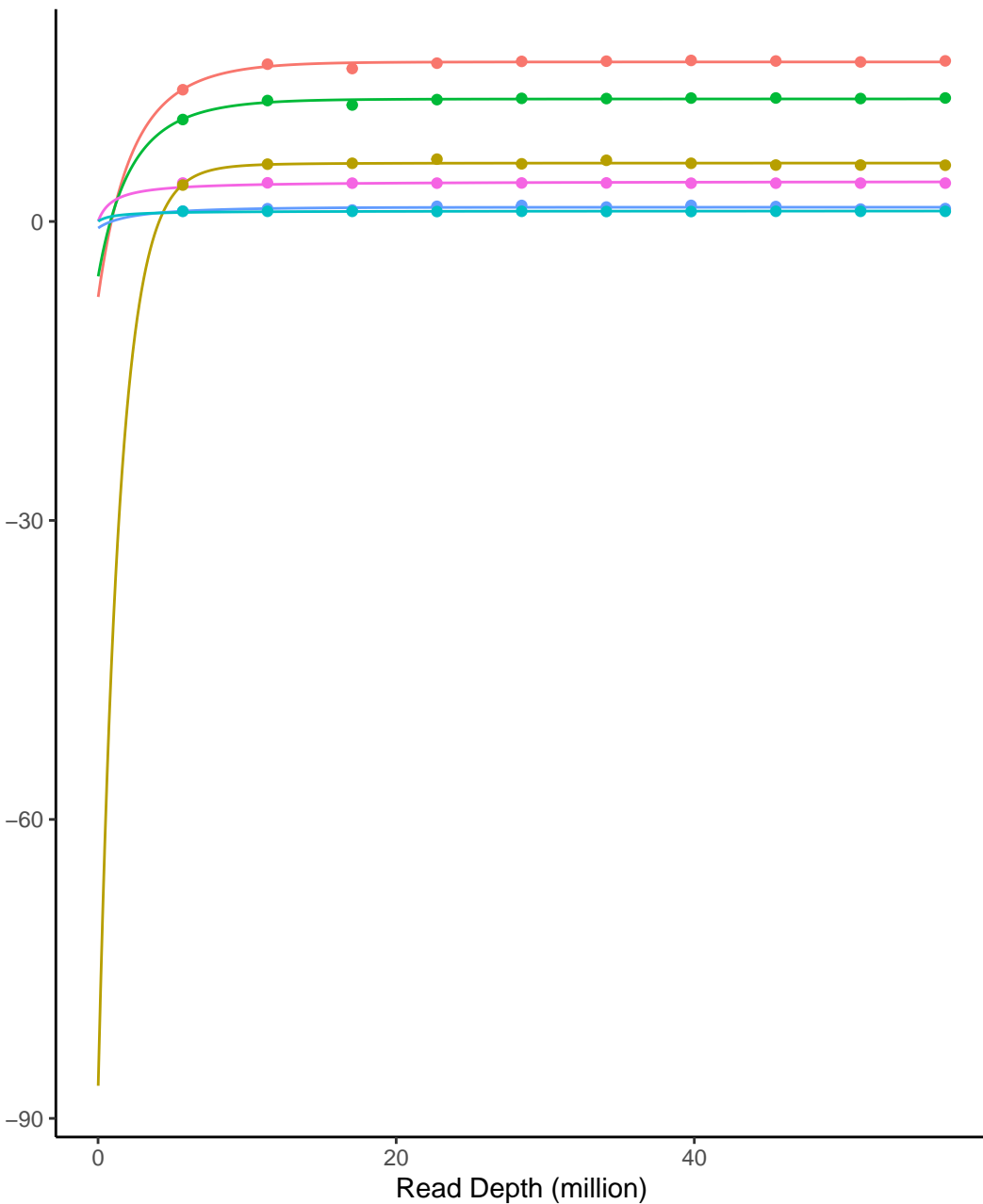


PGnos_high_hv13-1



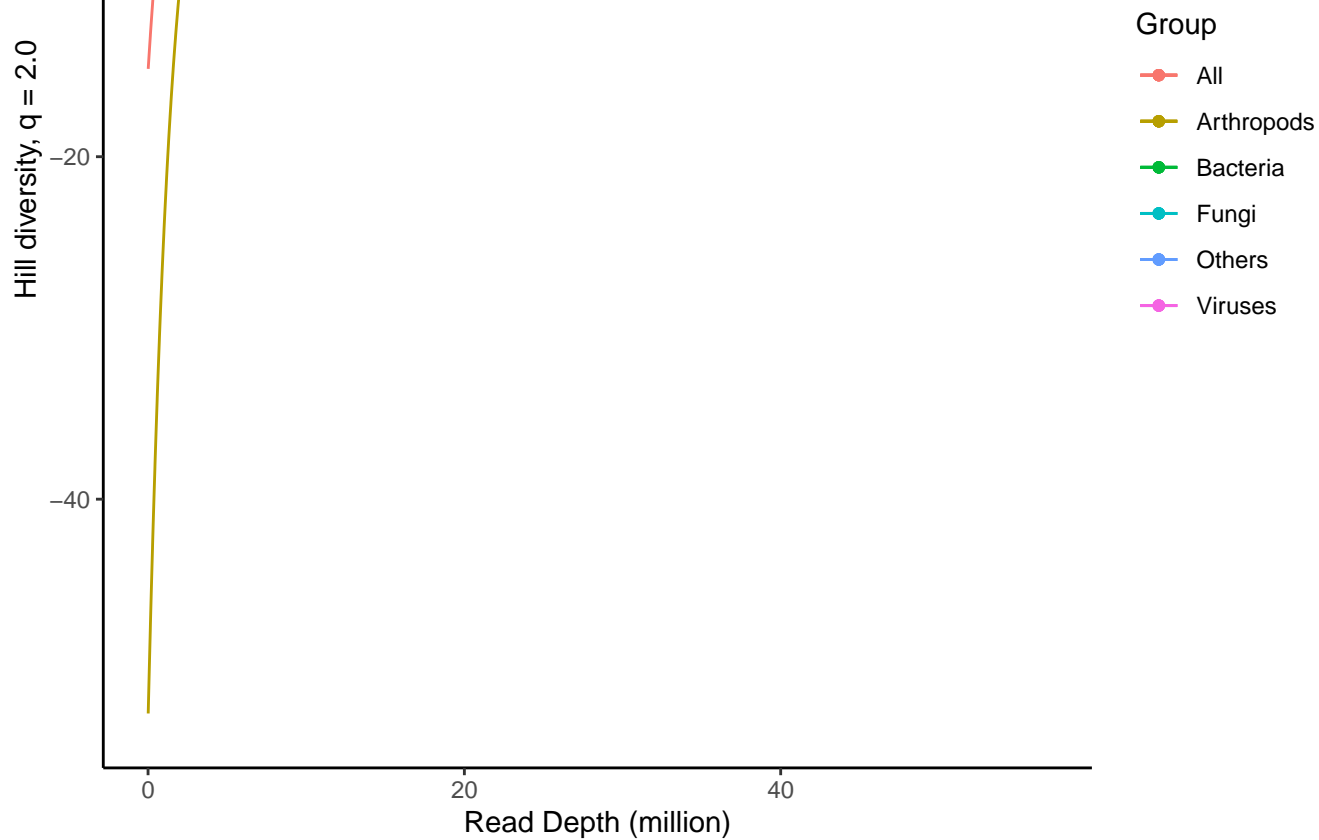
PGnos_high_hv13-2

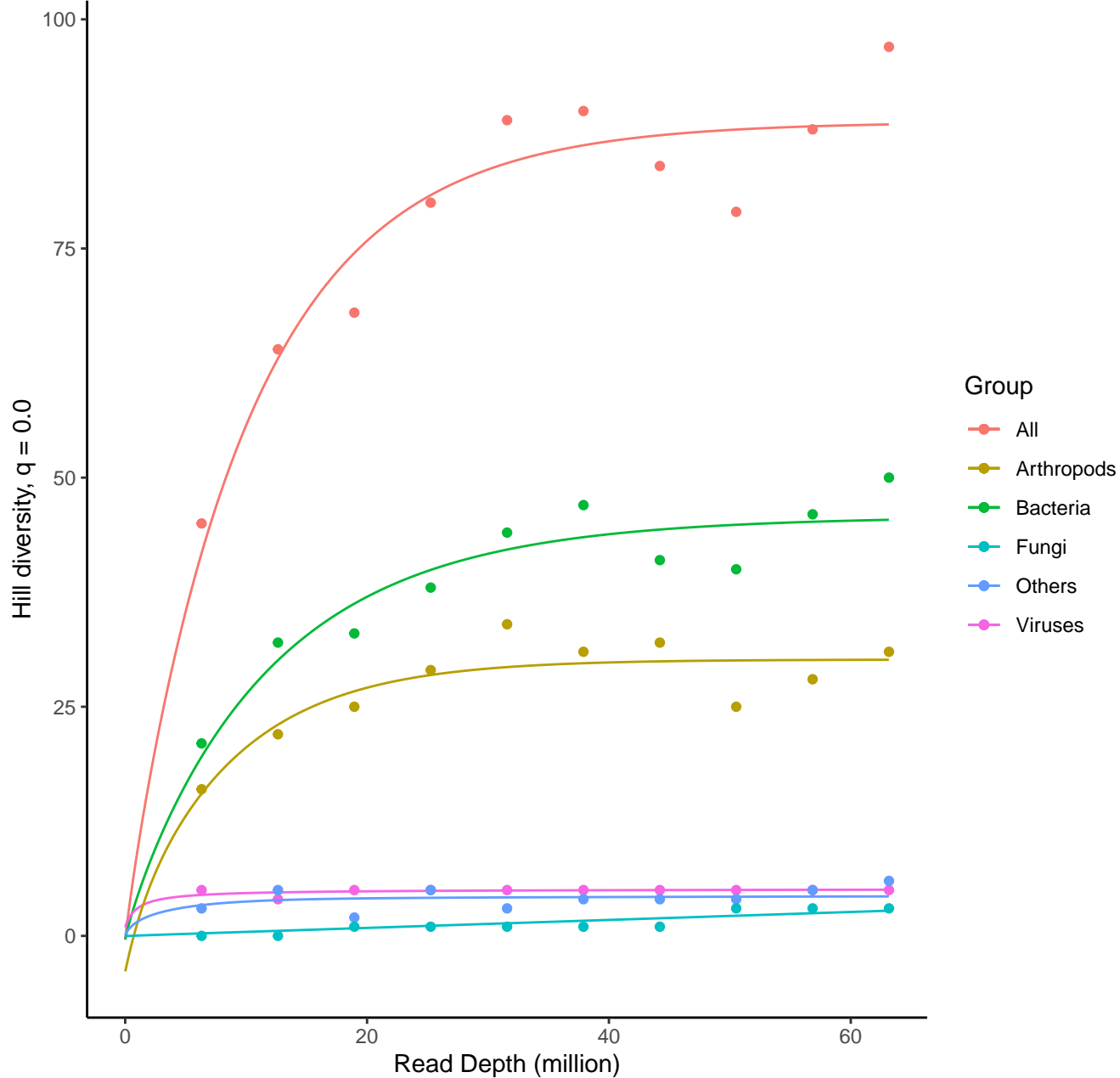


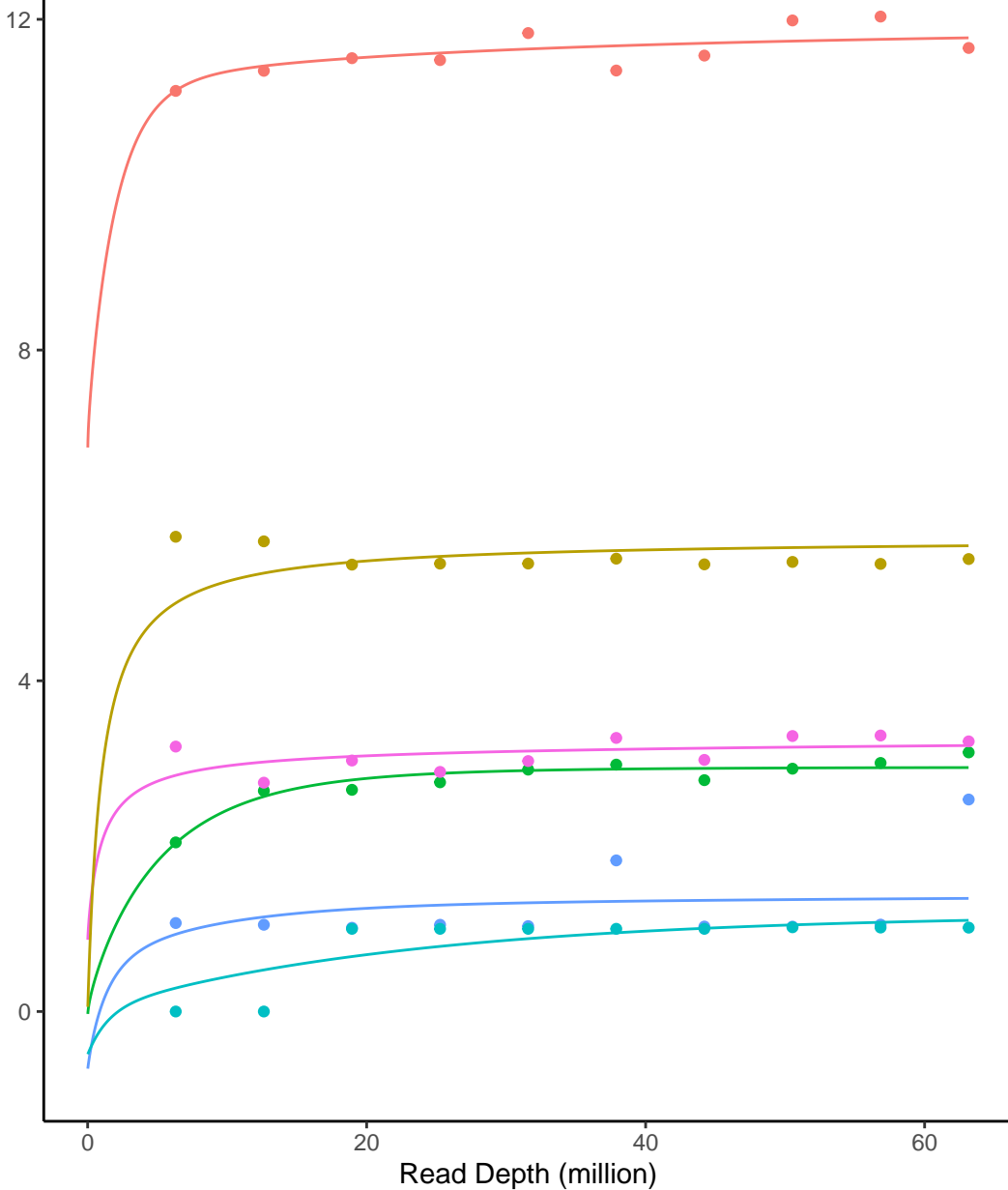
Hill diversity, $q = 1.0$ 

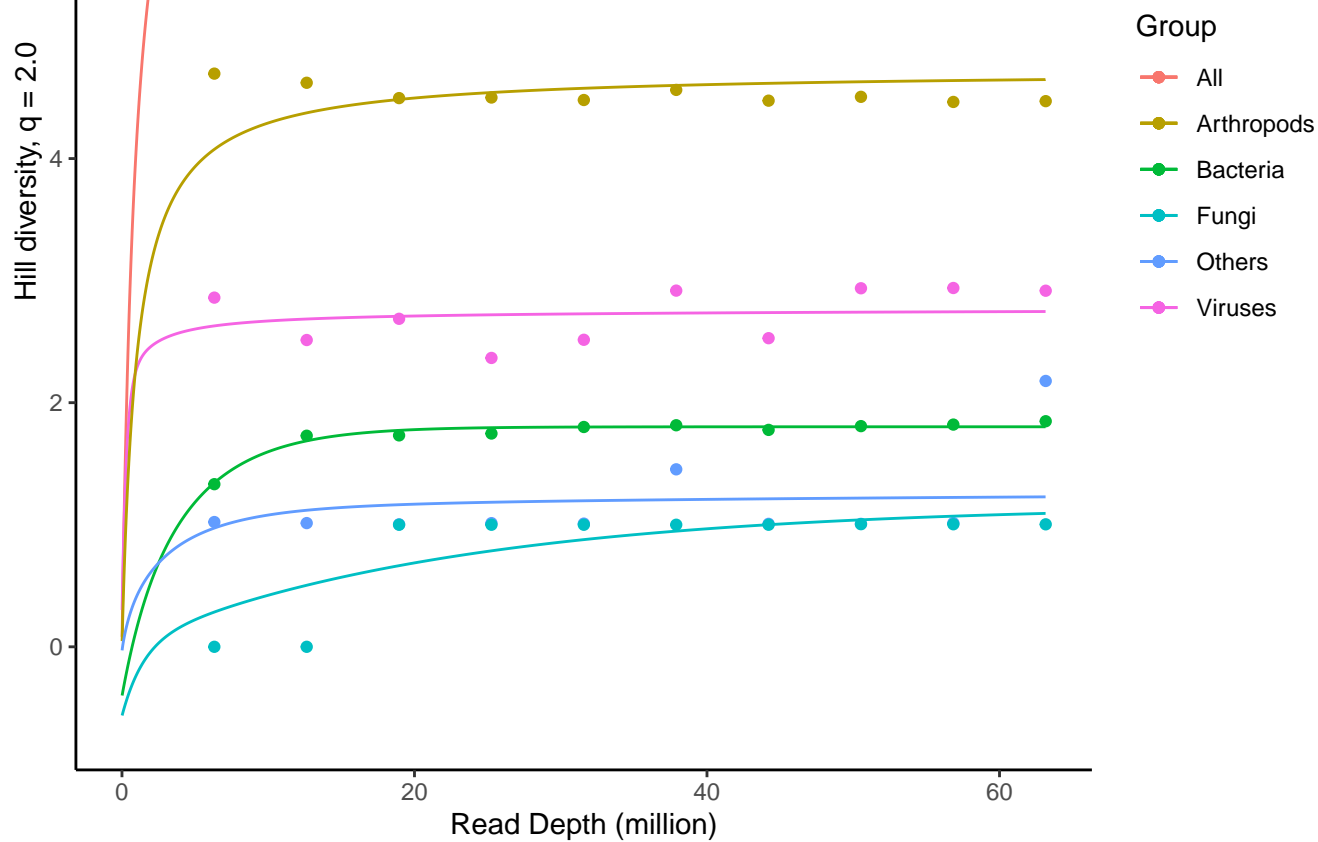
Group

- All
- Arthropods
- Bacteria
- Fungi
- Others
- Viruses



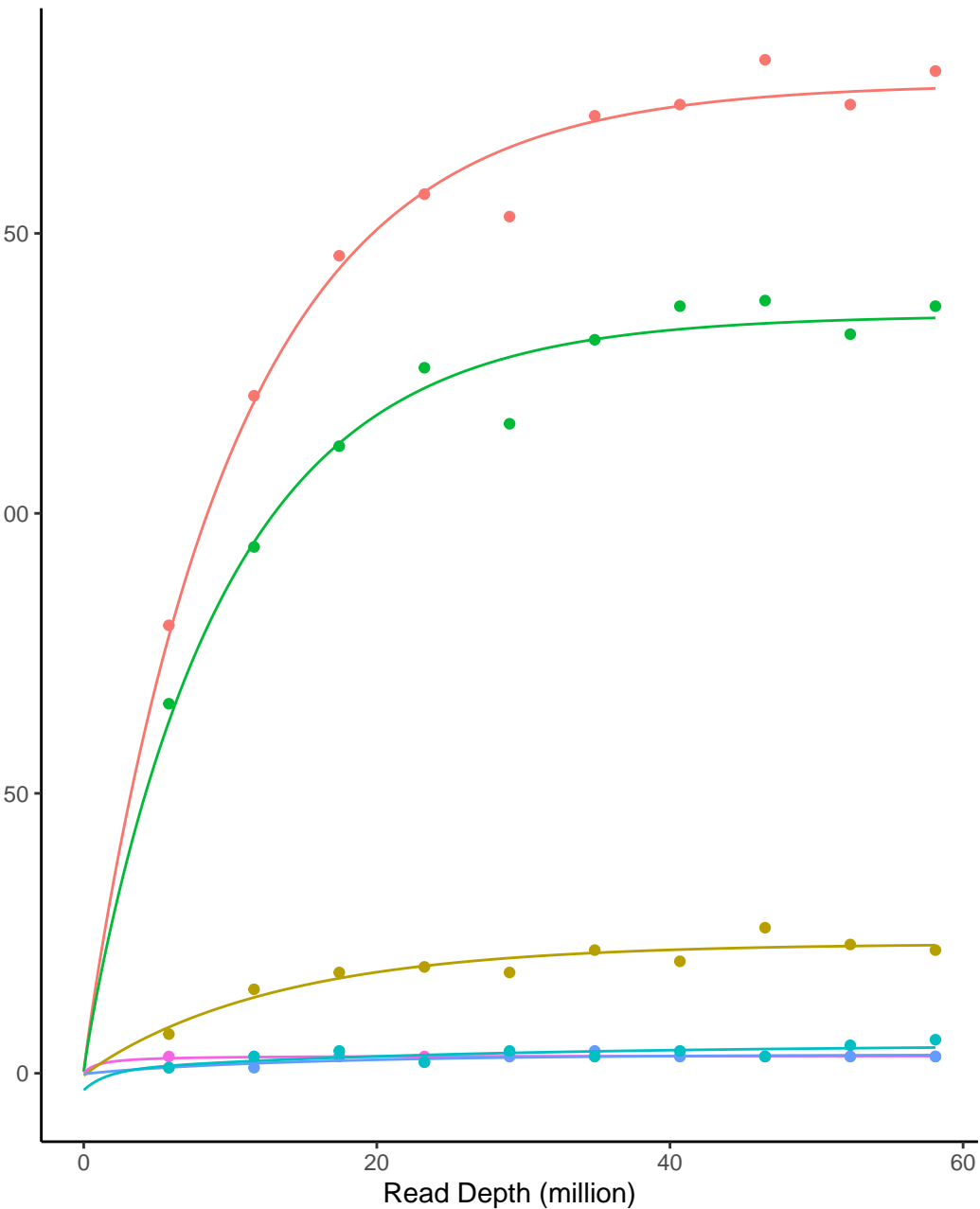


Hill diversity, $q = 1.0$ 



PGpollen_fresh

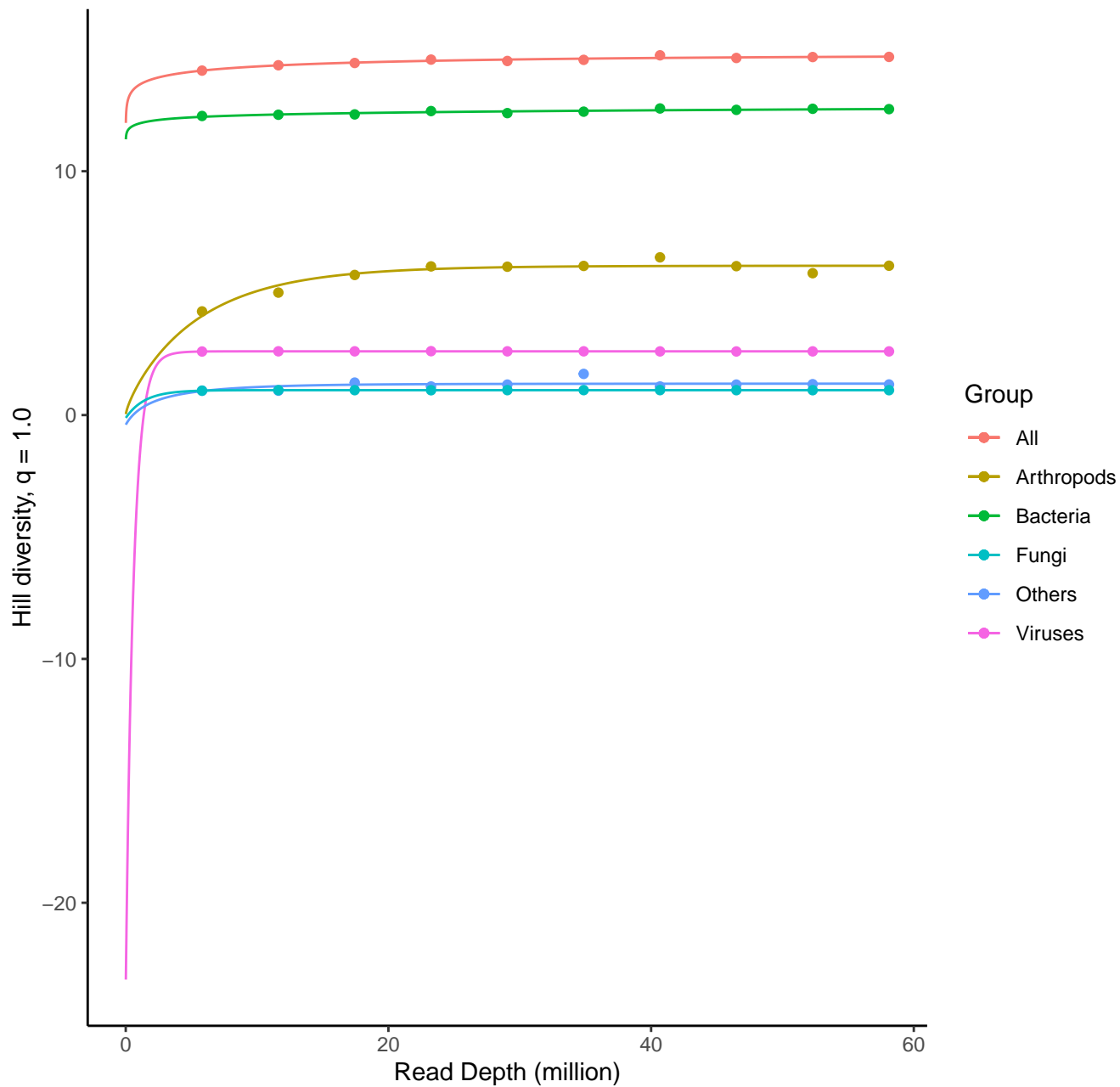
Hill diversity, $q = 0.0$



Group

- All
- Arthropods
- Bacteria
- Fungi
- Others
- Viruses

PGpollen_fresh



PGpollen_fresh

Hill diversity, $q = 2.0$

