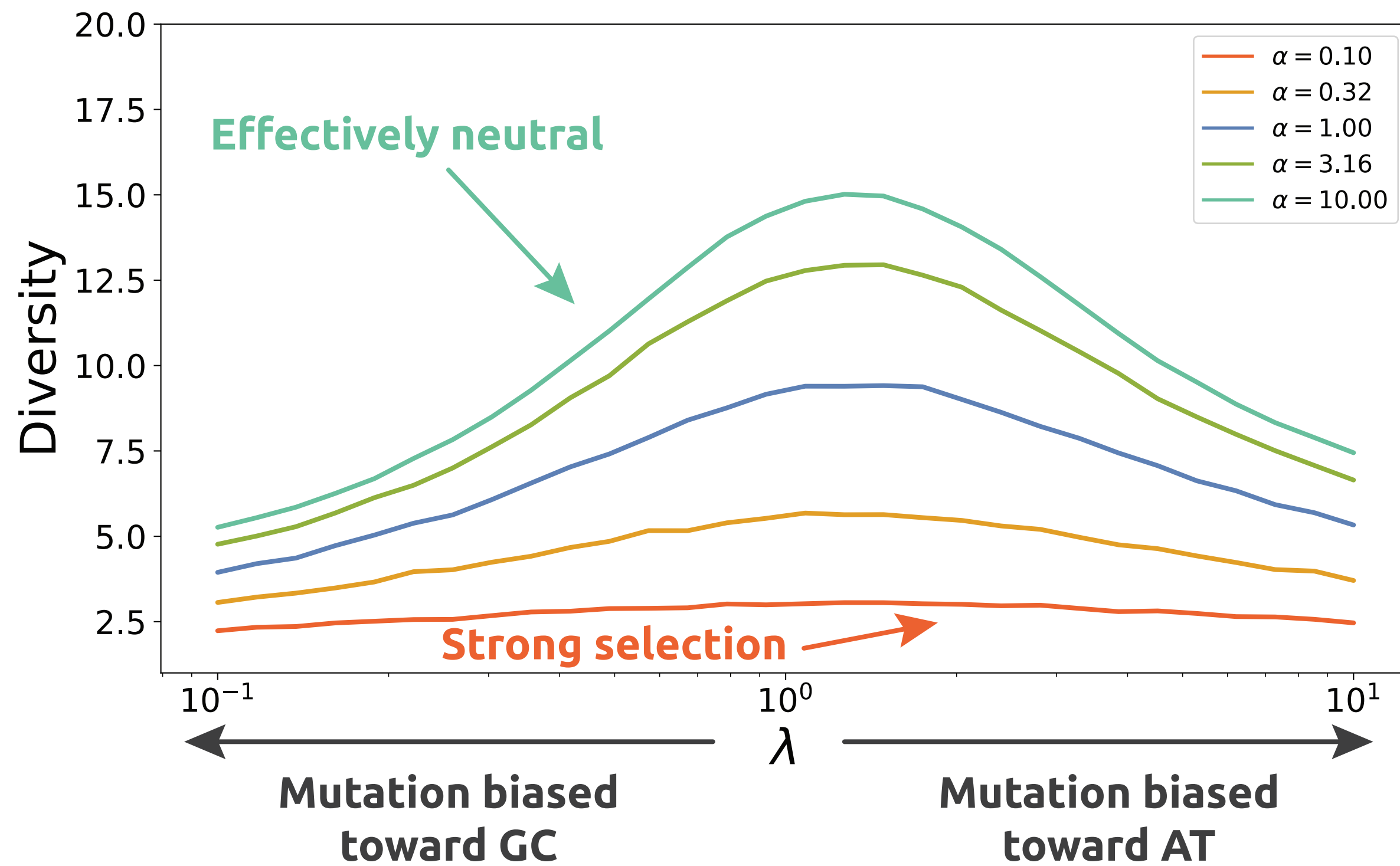


Diversity of amino-acids for each site of the sequence (average)



Diversity of amino-acids for the whole sequence

