Thesis work

Table peptides 2-50 AAR - SwissProt (555426 proteins)

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Introduction

I attach the table with the results obtained based on the file **uniprot_sprot.fasta** (updated in September - 555426 proteins). This table is similar to Table 1 in the file Searching the protein sequence database, with the updated results:

| Length of peptide (N) | Total possible N-peptides (20^N) | Number found | Percentage found (%) |
|-------------------------|------------------------------------|--------------|----------------------------|
| 1 | 20 | 20 | 100 |
| 2 | 400 | 400 | 100 |
| 3 | 8000 | 8000 | 100 |
| 4 | 160000 | 159999 | 99.99938 |
| 5 | 3200000 | 3113509 | 97.29716 |
| 6 | 64000000 | 32921109 | 51.43923 |
| 7 | 1280000000 | 84118859 | 6.571786 |
| 8 | 25600000000 | 100896814 | 0.3941282 |
| 9 | 512000000000 | 105834330 | 0.02067077 |
| 10 | 10240000000000 | 108976567 | 0.001064224 |
| 11 | 204800000000000 | 111551595 | 0.00005446855 |
| 12 | 4.096000×10^{15} | 113751287 | 2.777131×10^{-6} |
| 13 | 8.192000×10^{16} | 115660117 | 1.411867×10^{-7} |
| 14 | 1.638400×10^{18} | 117330023 | 7.161256×10^{-9} |
| 15 | 3.276800×10^{19} | 118797220 | 3.625403×10^{-10} |
| 16 | 6.553600×10^{20} | 120090802 | 1.832440×10^{-11} |
| 17 | 1.310720×10^{22} | 121236766 | 9.249631×10^{-13} |
| 18 | 2.621440×10^{23} | 122255013 | 4.663659×10^{-14} |
| 19 | 5.242880×10^{24} | 123160391 | 2.349098×10^{-15} |
| 20 | 1.048576×10^{26} | 123966286 | 1.182235×10^{-16} |
| 21 | 2.097152×10^{27} | 124685091 | 5.945448×10^{-18} |
| 22 | 4.194304×10^{28} | 125325735 | 2.987998×10^{-19} |
| 23 | 8.388608×10^{29} | 125896961 | 1.500809×10^{-20} |
| 24 | 1.677722×10^{31} | 126406287 | 7.534402×10^{-22} |
| 25 | 3.355443×10^{32} | 126859712 | 3.780714×10^{-23} |
| 26 | 6.710886×10^{33} | 127262721 | 1.896362×10^{-24} |
| 27 | 1.342177×10^{35} | 127619577 | 9.508399×10^{-26} |
| 28 | 2.684355×10^{36} | 127934582 | 4.765935×10^{-27} |
| 29 | 5.368709×10^{37} | 128211241 | 2.388120×10^{-28} |
| 30 | 1.073742×10^{39} | 128452495 | 1.196307×10^{-29} |
| 31 | 2.147484×10^{40} | 128661754 | 5.991280×10^{-31} |
| 32 | 4.294967×10^{41} | 128841734 | 2.999830×10^{-32} |
| 33 | 8.589935×10^{42} | 128994975 | 1.501699×10^{-33} |
| 34 | 1.717987×10^{44} | 129123214 | 7.515960×10^{-35} |
| 35 | 3.435974×10^{45} | 129227999 | 3.761030×10^{-36} |

| Length of peptide (N) | Total possible N-peptides (20^N) | Number found | Percentage found (%) |
|-----------------------|------------------------------------|--------------|----------------------------|
| 36 | 6.871948×10^{46} | 129311151 | 1.881725×10^{-37} |
| 37 | 1.374390×10^{48} | 129374023 | 9.413199×10^{-39} |
| 38 | 2.748779×10^{49} | 129418106 | 4.708203×10^{-40} |
| 39 | 5.497558×10^{50} | 129444962 | 2.354590×10^{-40} |
| 40 | 1.099512×10^{52} | 129455435 | 1.177390×10^{-42} |
| 41 | 2.199023×10^{53} | 129450779 | 5.886740×10^{-44} |
| 42 | 4.398047×10^{54} | 129431855 | 2.942940×10^{-45} |
| 43 | 8.796093×10^{55} | 129399352 | 1.471100×10^{-46} |
| 44 | 1.759219×10^{57} | 129354043 | 7.352926×10^{-48} |
| 45 | 3.518437×10^{58} | 129296497 | 3.674827×10^{-49} |
| 46 | 7.036874×10^{59} | 129227572 | 1.836434×10^{-50} |
| 47 | 1.407375×10^{61} | 129147725 | 9.176498×10^{-52} |
| 48 | 2.814750×10^{62} | 129057687 | 4.585050×10^{-53} |
| 49 | 5.629500×10^{63} | 128957792 | 2.290751×10^{-54} |
| 50 | 1.125900×10^{65} | 128848422 | 1.144404×10^{-55} |

I also attach the Table 1 of the file mentioned in the beginning, so that you can see the notorious differences of the results obtained.

TABLE I

| Length of peptide | Total possible N-peptides | Number found | Percentage found |
|-------------------|------------------------------|-----------------|---------------------|
| N | 20**N | | |
| 2 | 400 | 400 | 100.0 |
| 3 | 8,000 | 7,995 | 99.9 |
| 4 | 160,000 | 115,817 | 72.4 |
| 5 | 3,200,000 | 333,965 | 10.4 |
| 6 | 64,000,000 | 387,925 | 0.61 |
| 7 | 1,280,000,000 | 399,330 | 0.031 |
| 8 | 25,600,000,000 | 405,682 | 0.0016 |

FAB LATEX 2