Additional file 1 - Verify3D evaluation of protein models.

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| Model | Amino acid length | Percentage of amino acids that have reached acceptable 3D-1D score\* |
| Wsv249(ANK) | 208 | 75.48% |
| Wsv249(RING) | 56 | 26.79% |
| Wsv463a(FH2) | 339 | 37.17% |
| Wsv477(RRM) | 71 | 36.62% |
| Wsv479 | 418 | 53.35% |
| Wsv492 | 92 |  0.00%  |
| Wsv497 | 481 | 34.72% |

\*At least 80% of the amino acids have scored >= 0.2 in the 3D/1D profile.