Table S3 The secondary structure prediction of ZHD genes in grape

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| Gene name | Alpha helix (%) | Beta turn (%) | Random coil (%) |
| *VvZHD1* | 16.40 | 4.80 | 78.80 |
| *VvZHD2* | 11.95 | 1.59 | 86.45 |
| *VvZHD3* | 16.96 | 6.96 | 76.09 |
| *VvZHD4* | 10.23 | 20.45 | 69.32 |
| *VvZHD5* | 12.69 | 4.95 | 82.35 |
| *VvZHD6* | 9.78 | 17.39 | 72.83 |
| *VvZHD7* | 11.59 | 3.48 | 84.93 |
| *VvZHD8* | 15.30 | 3.20 | 81.49 |
| *VvZHD9* | 20.00 | 7.41 | 72.59 |
| *VvZHD10* | 11.35 | 4.29 | 84.36 |
| *VvZHD11* | 16.34 | 5.23 | 78.43 |
| *VvZHD12* | 16.31 | 4.26 | 79.43 |