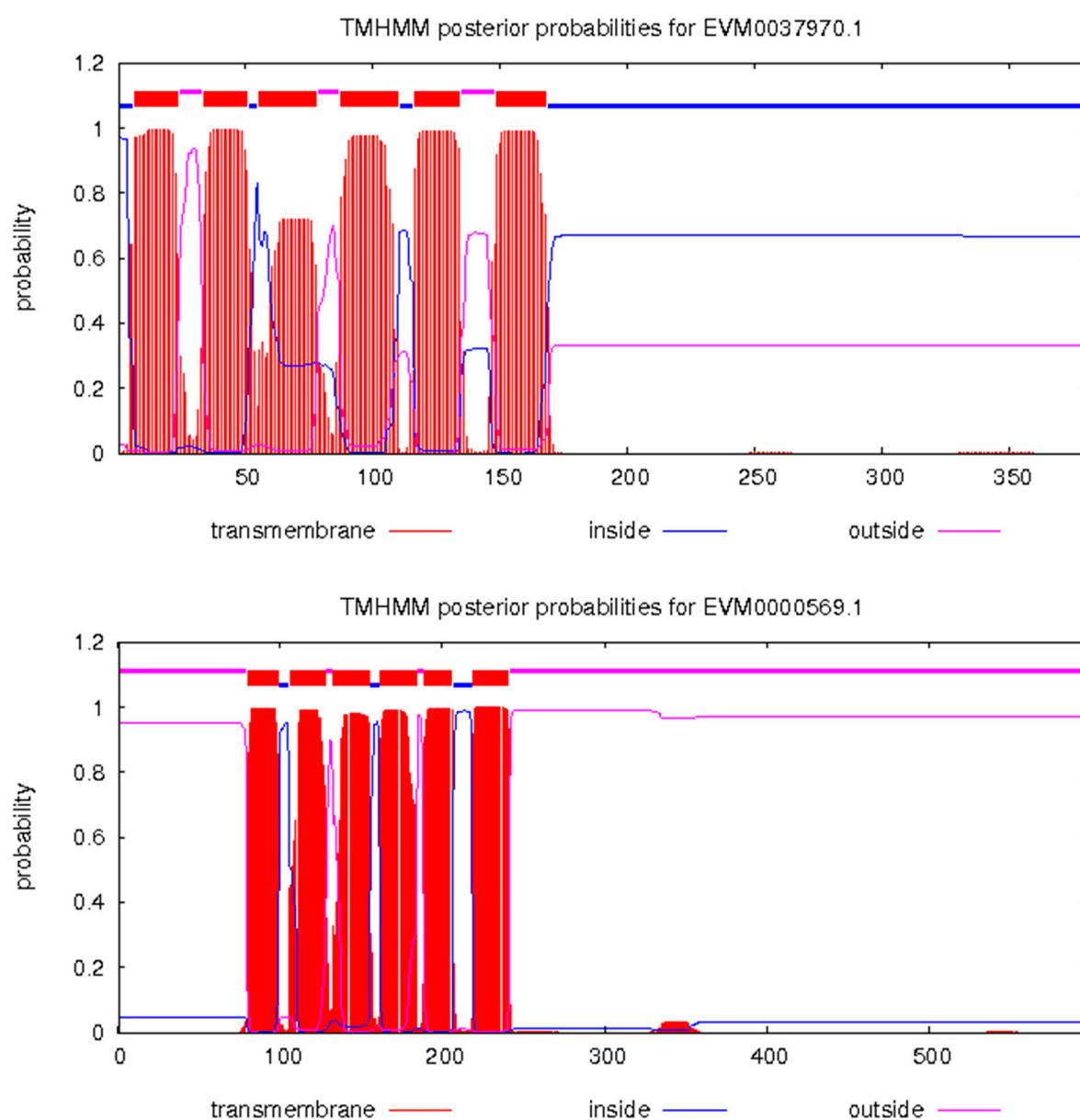
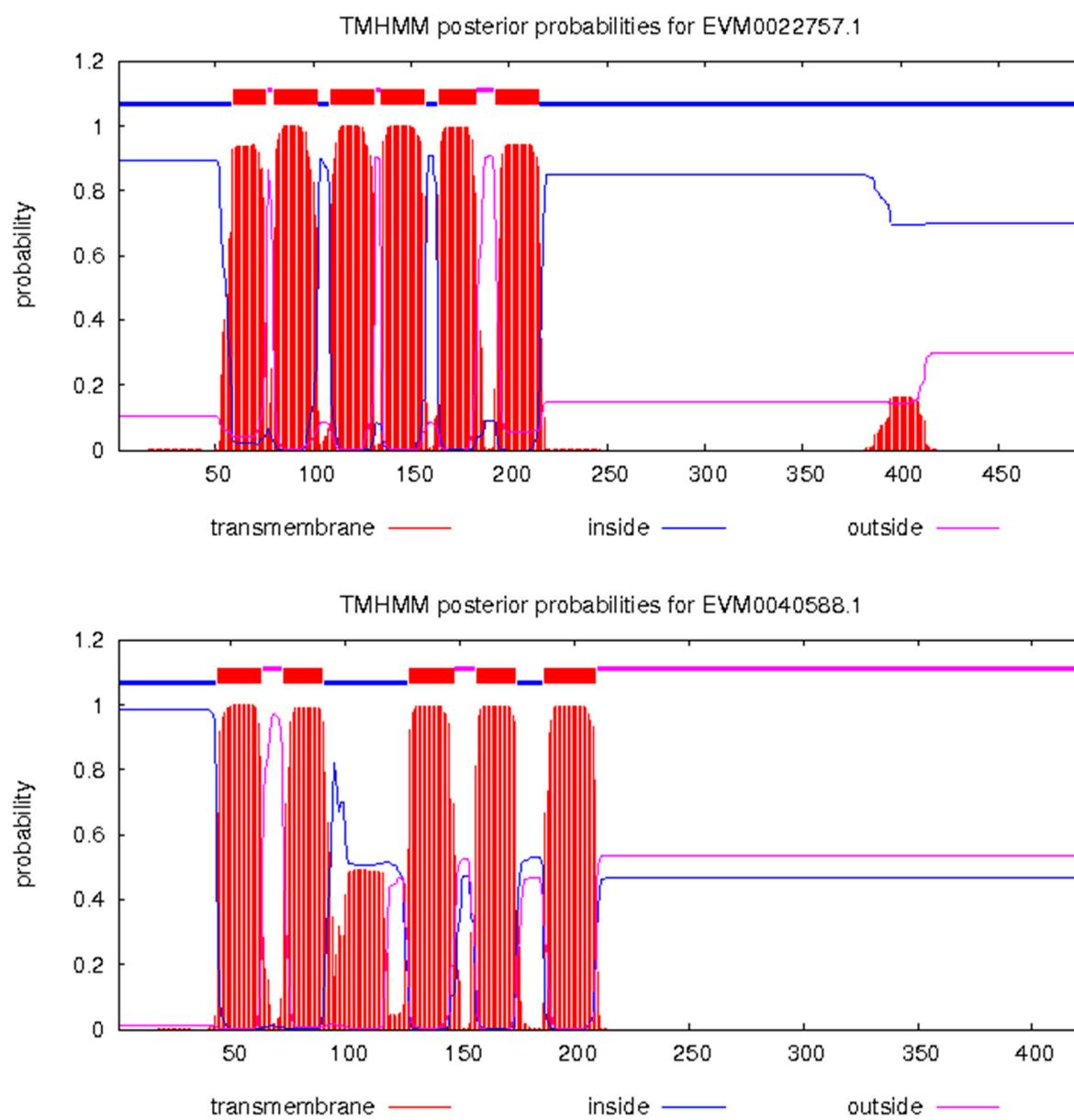


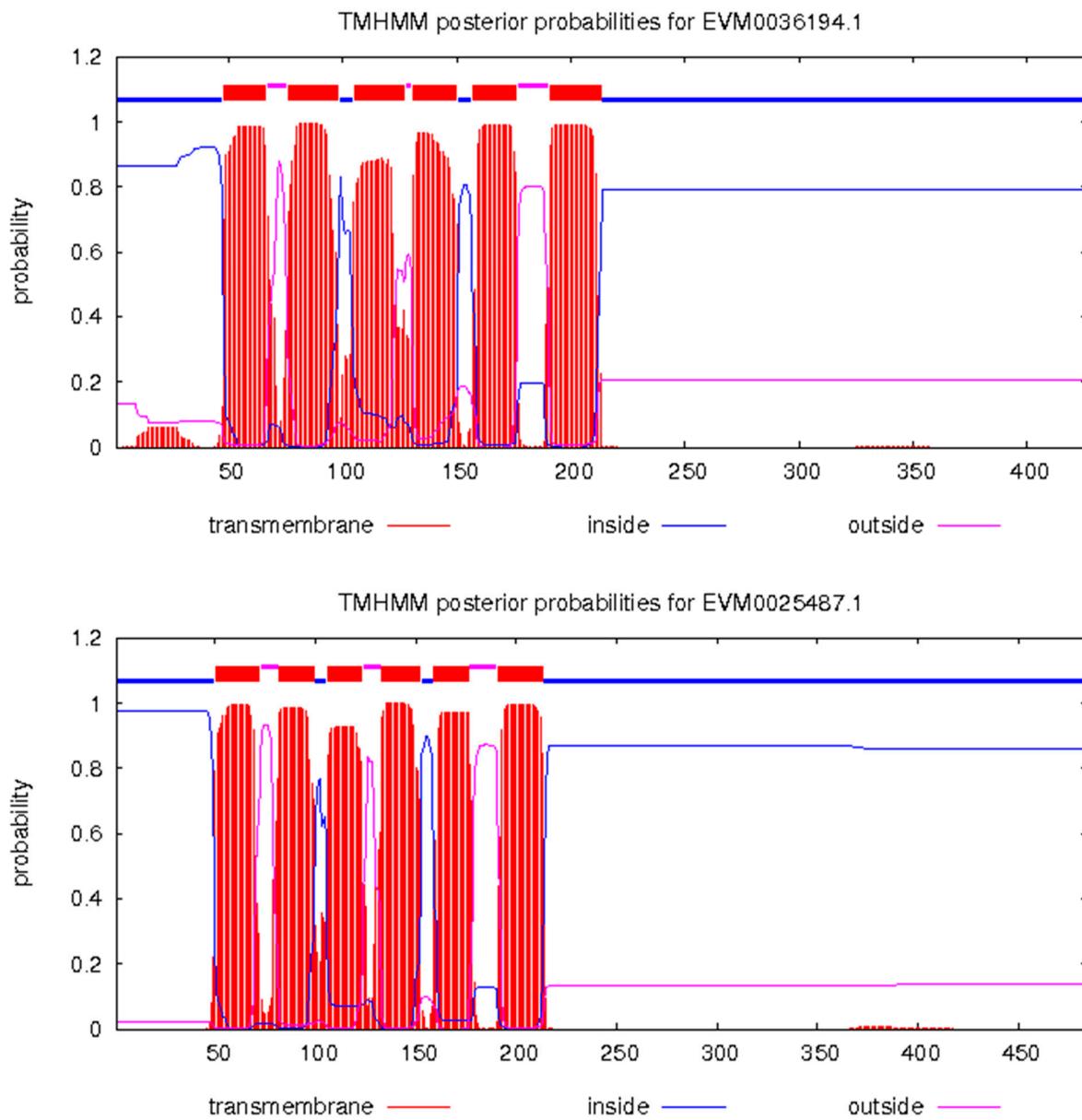
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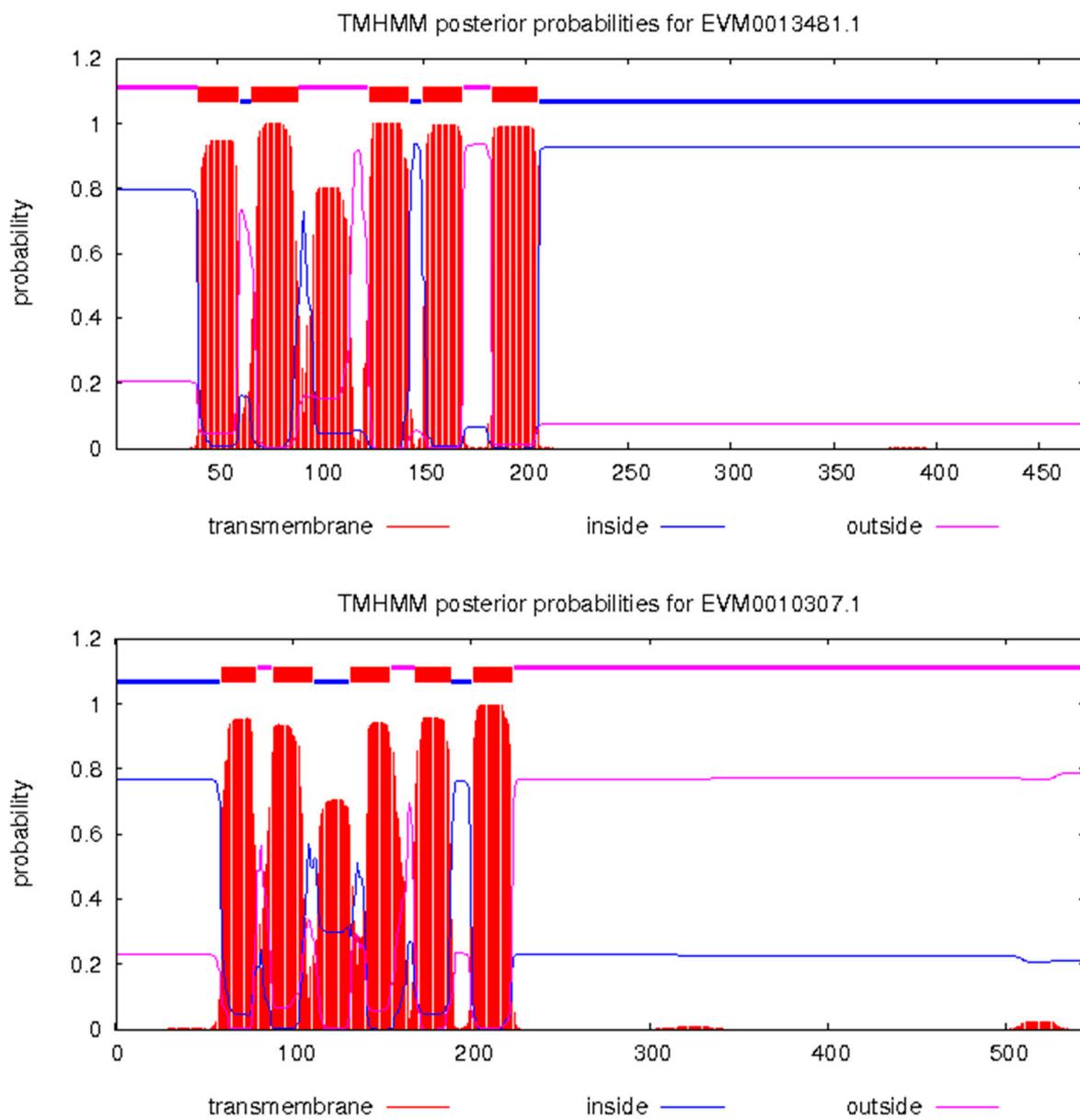
Supplementary Materials: Genome-Wide Identification, Characterization and Expression Profiling of Aluminum-Activated Maltate Transporters in *Eriobotrya japonica* Lindl.

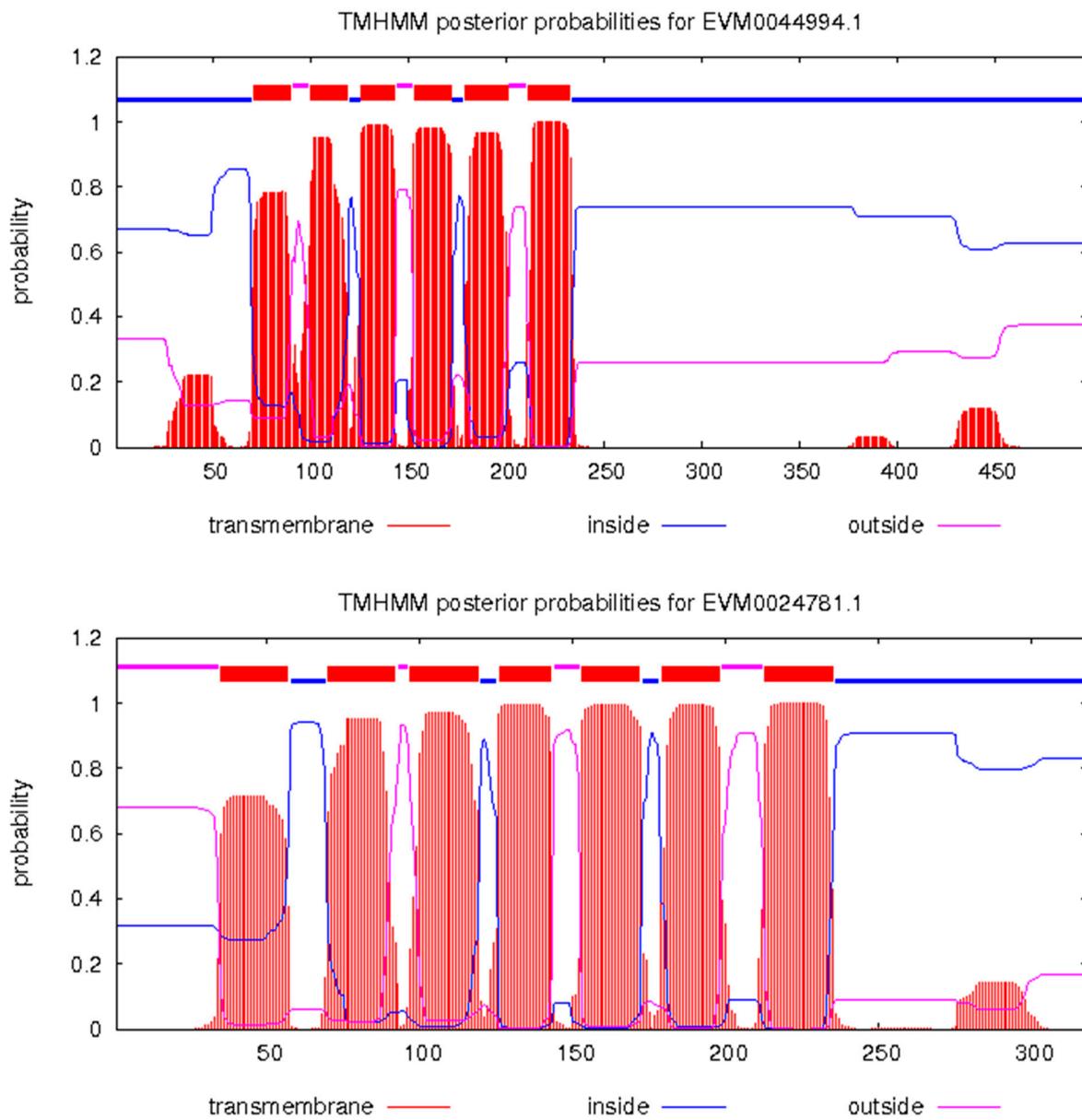
Muhammad Moaaz Ali ¹, Shariq Mahmood Alam ², Raheel Anwar ³, Sajid Ali ⁴, Meng Shi ¹, Dangdi Liang ¹, Zhimin Lin ^{5,*} and Faxing Chen ^{1,*}

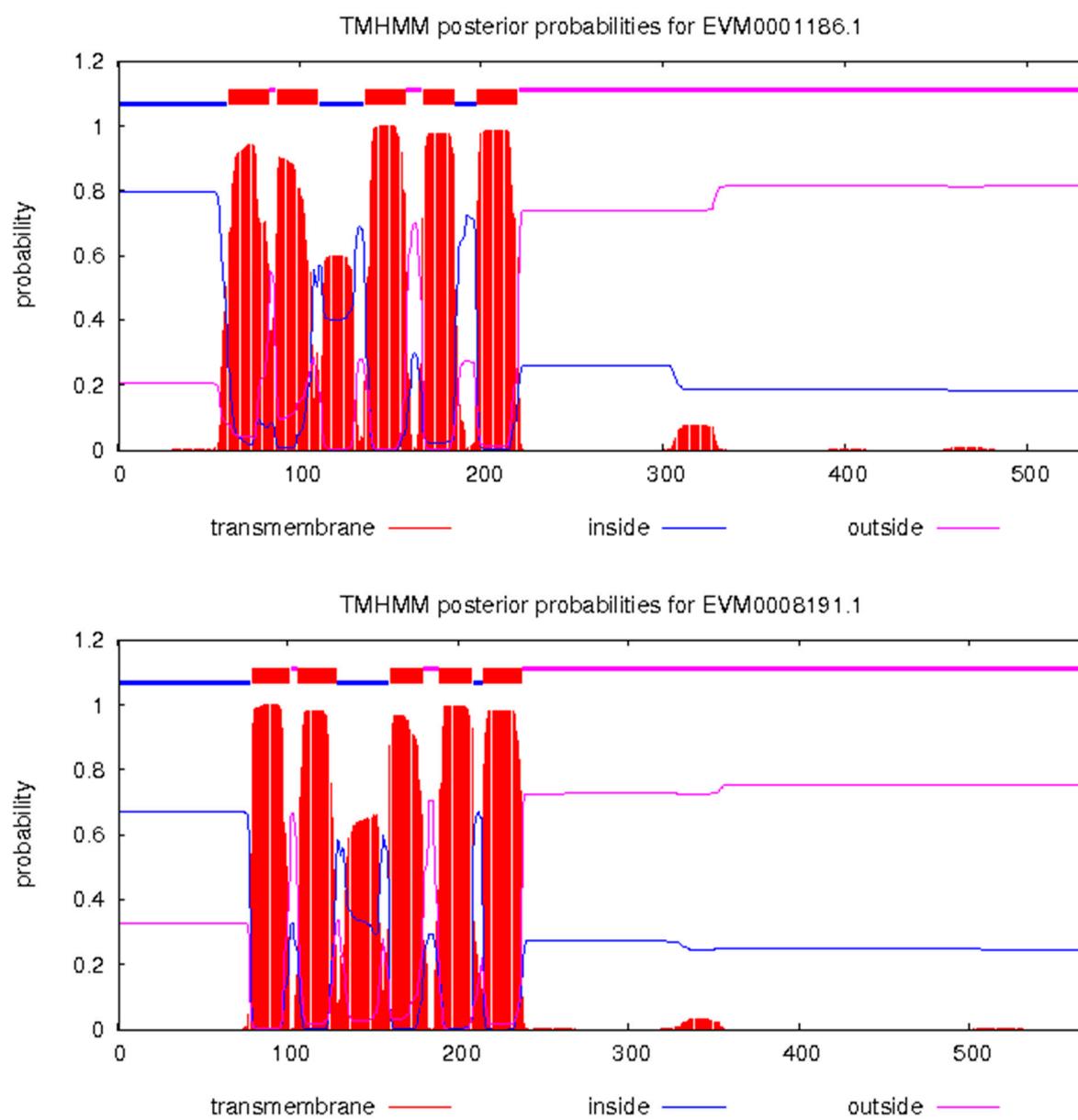


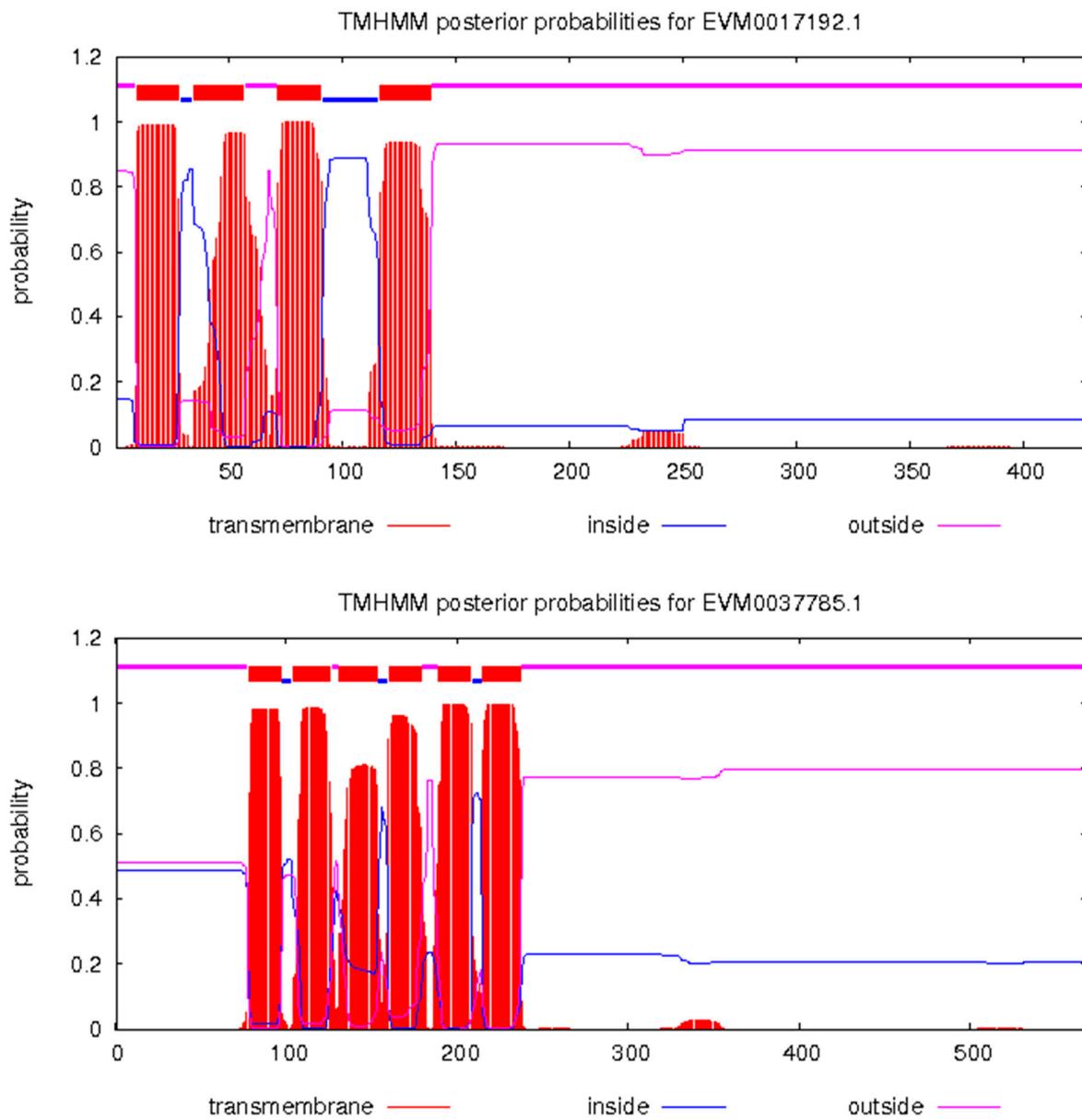


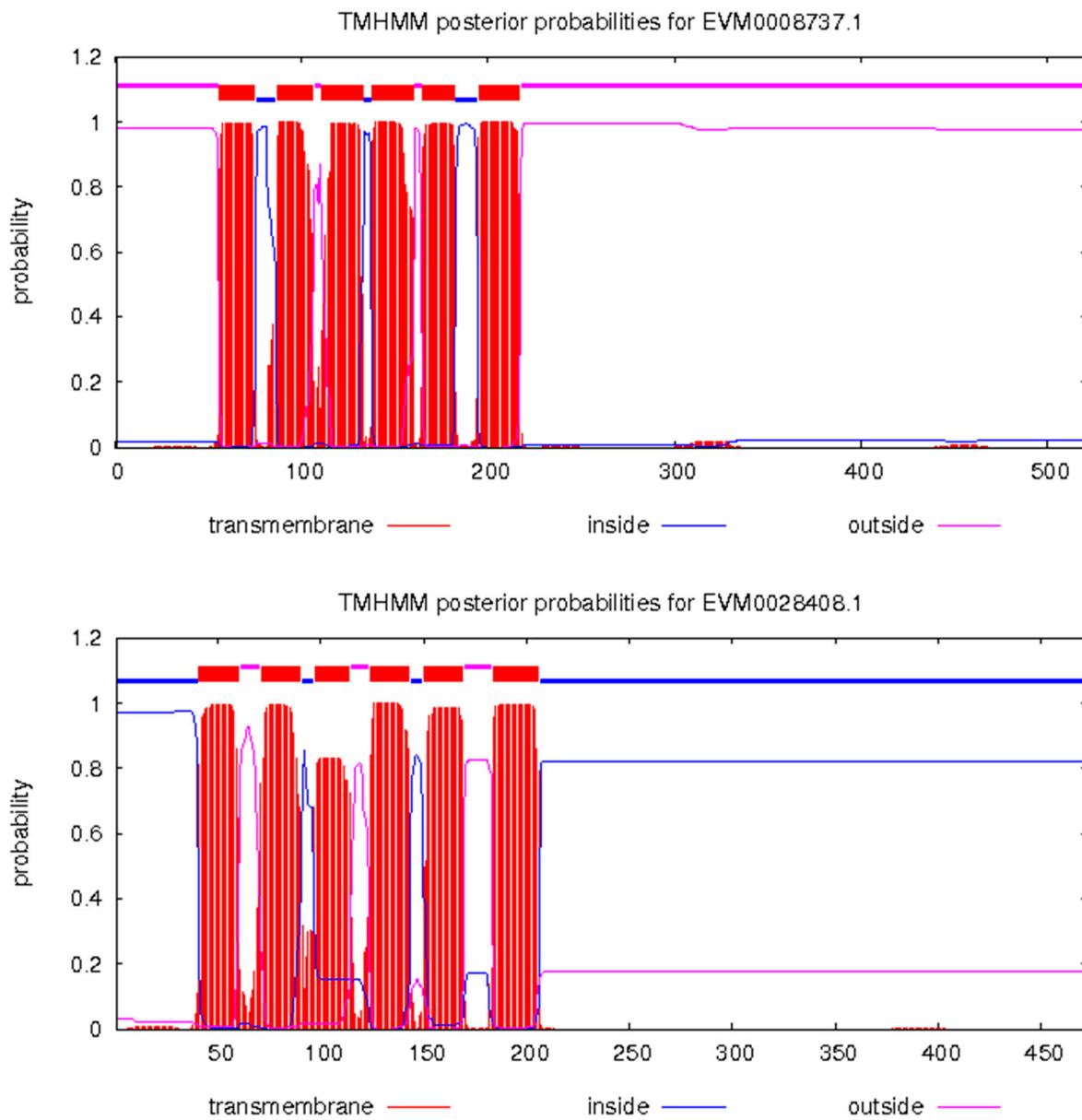


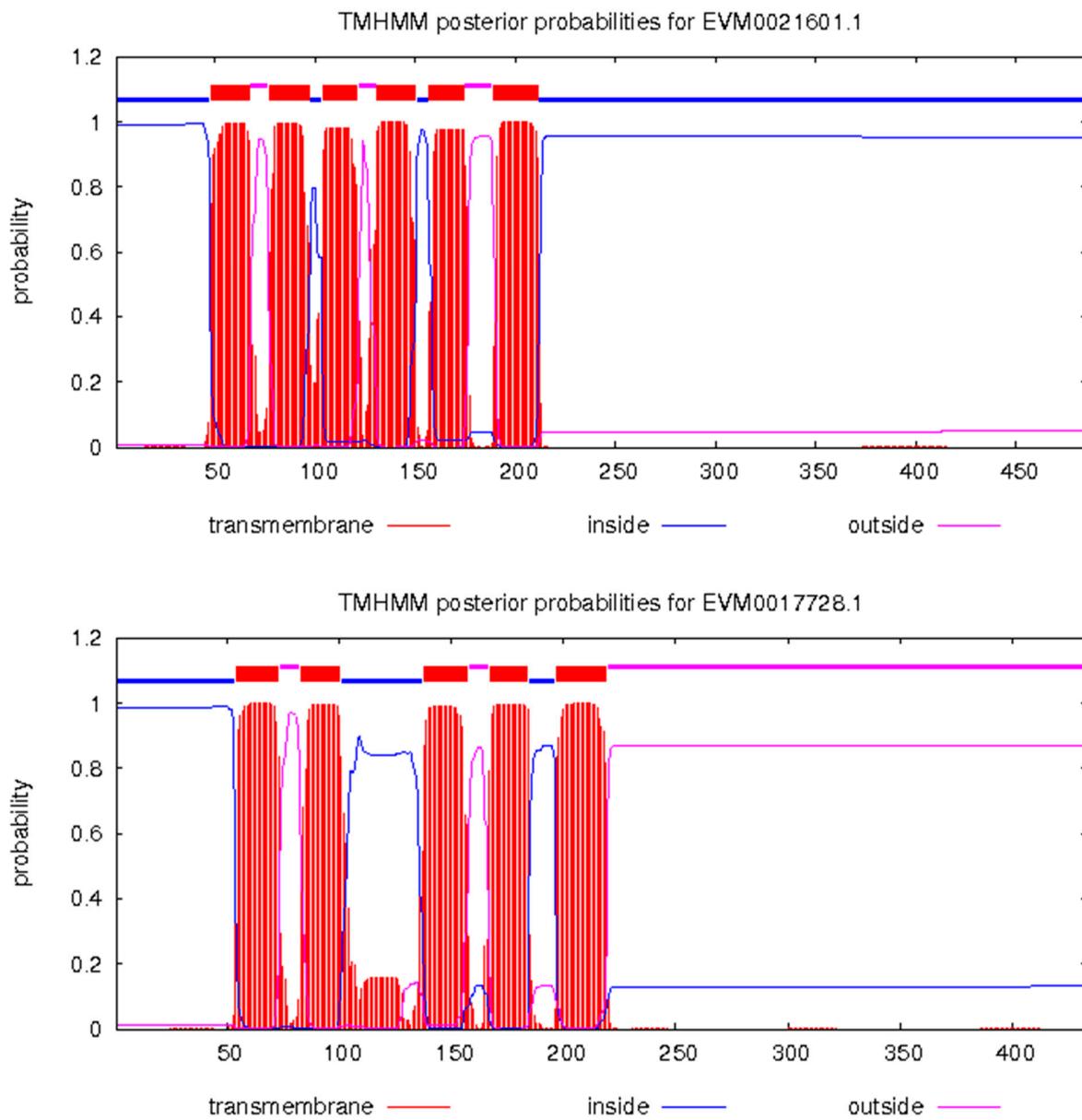


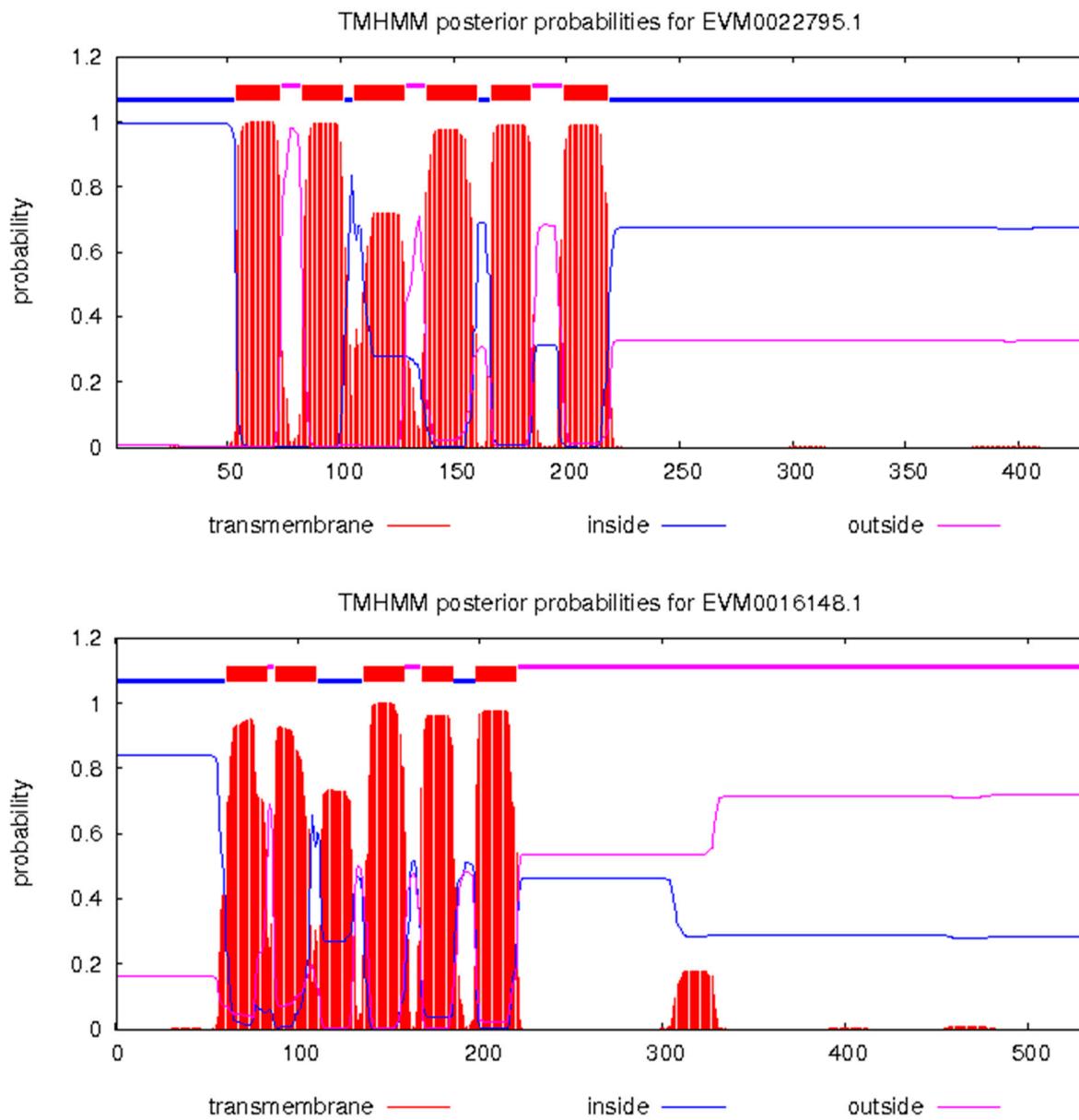


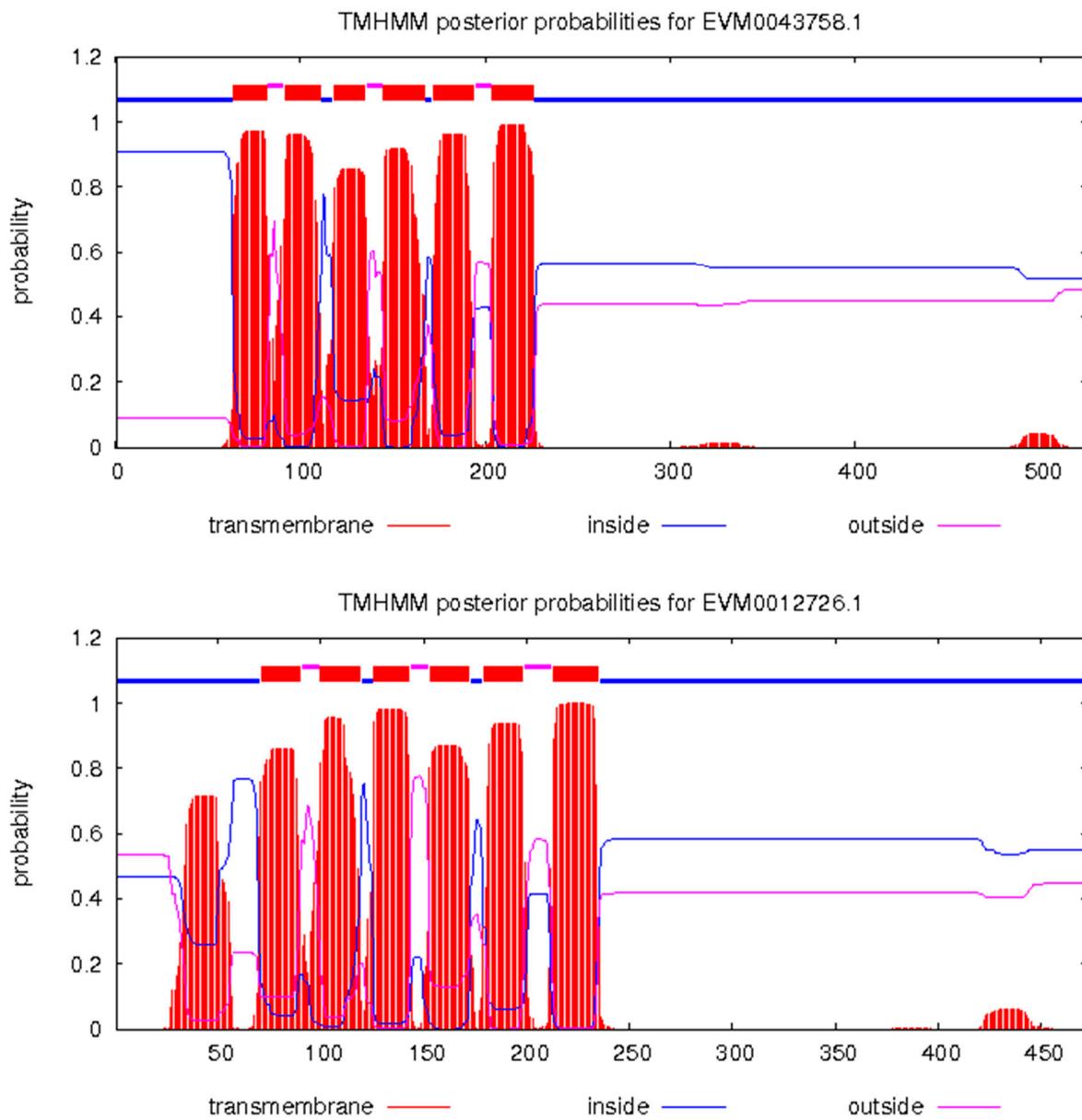












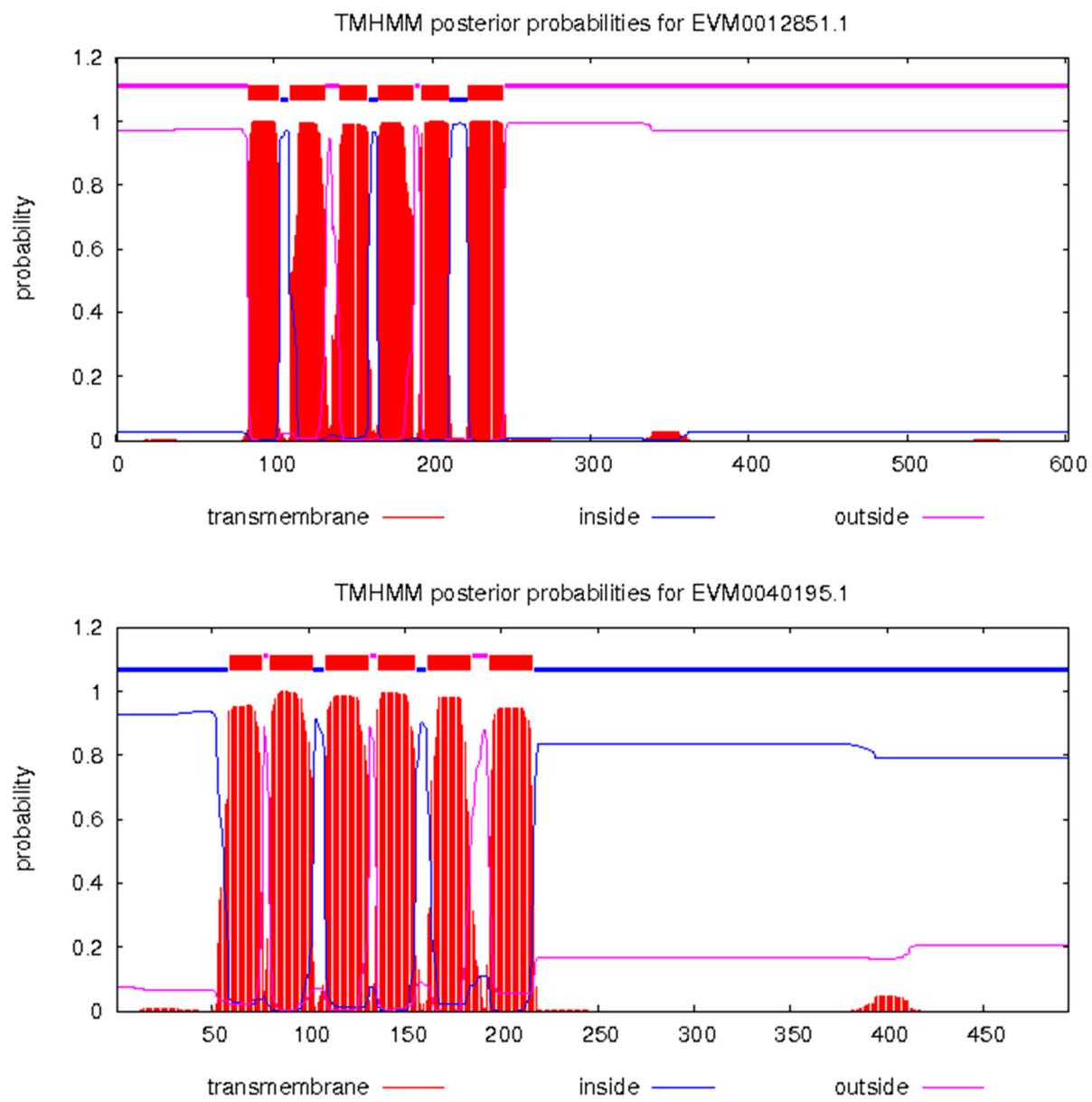


Figure S1. The predicted transmembrane structures of *EjALMT* genes.

Table S1. Summary of the information about transmembrane helices of EjALMT proteins.

Gene ID	Length (AAs)	Expected AAs	First 60	Predicted helices
EVM0037970.1	380	115.04	40.59	6
EVM0000569.1	597	123.83	0.00	6
EVM0022757.1	494	130.32	4.71	6
EVM0040588.1	423	111.87	16.55	5
EVM0036194.1	426	121.96	13.60	6
EVM0025487.1	484	121.26	10.78	6
EVM0013481.1	472	118.15	18.18	5
EVM0010307.1	544	116.12	0.94	5
EVM0044994.1	497	124.78	4.59	6
EVM0024781.1	318	139.29	15.87	7
EVM0001186.1	532	114.11	1.66	5
EVM0008191.1	568	119.09	0.00	5
EVM0017192.1	428	82.64	36.24	4
EVM0037785.1	568	120.29	0.00	6
EVM0008737.1	521	124.24	4.88	6
EVM0028408.1	472	118.71	19.42	6
EVM0021601.1	485	121.82	12.55	6
EVM0017728.1	433	104.91	6.61	5
EVM0022795.1	430	115.73	6.67	6
EVM0016148.1	532	119.03	1.55	5
EVM0043758.1	525	120.26	0.02	6
EVM0012726.1	472	132.02	15.49	6
EVM0012851.1	602	124.31	0.02	6
EVM0040195.1	494	125.81	4.79	6