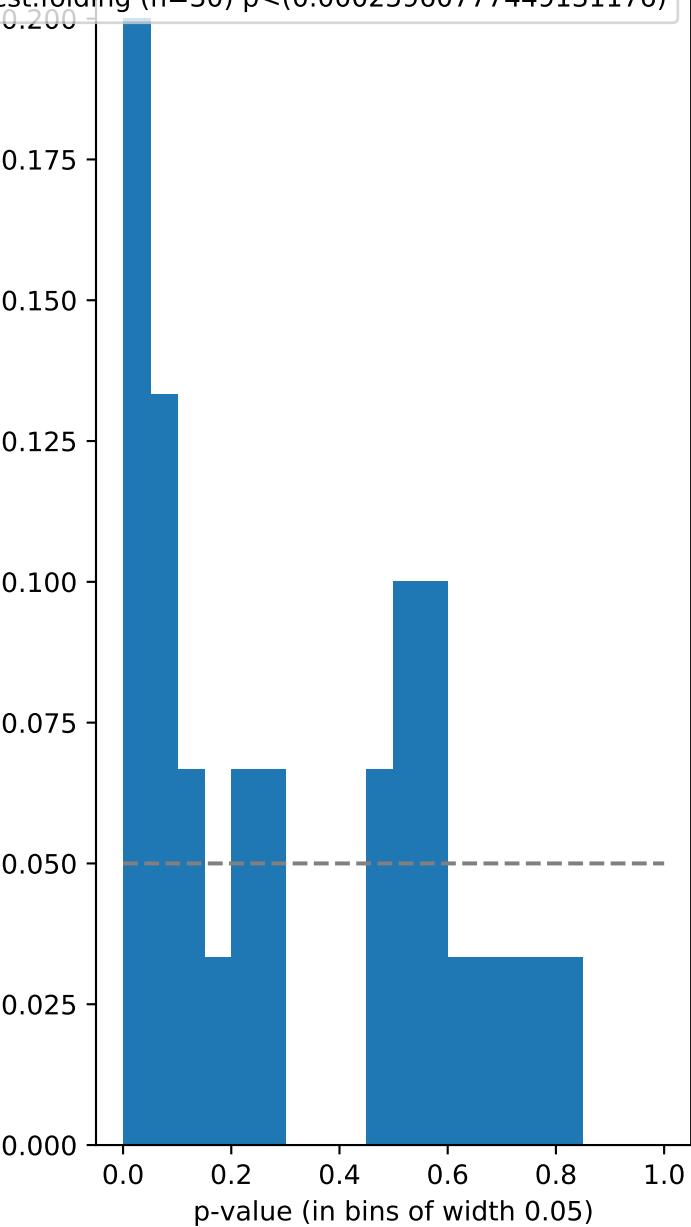


histogram for avgPairProbs

est.folding (n=30) $p < (0.00025960777449131176)$



histogram for helix-0-10

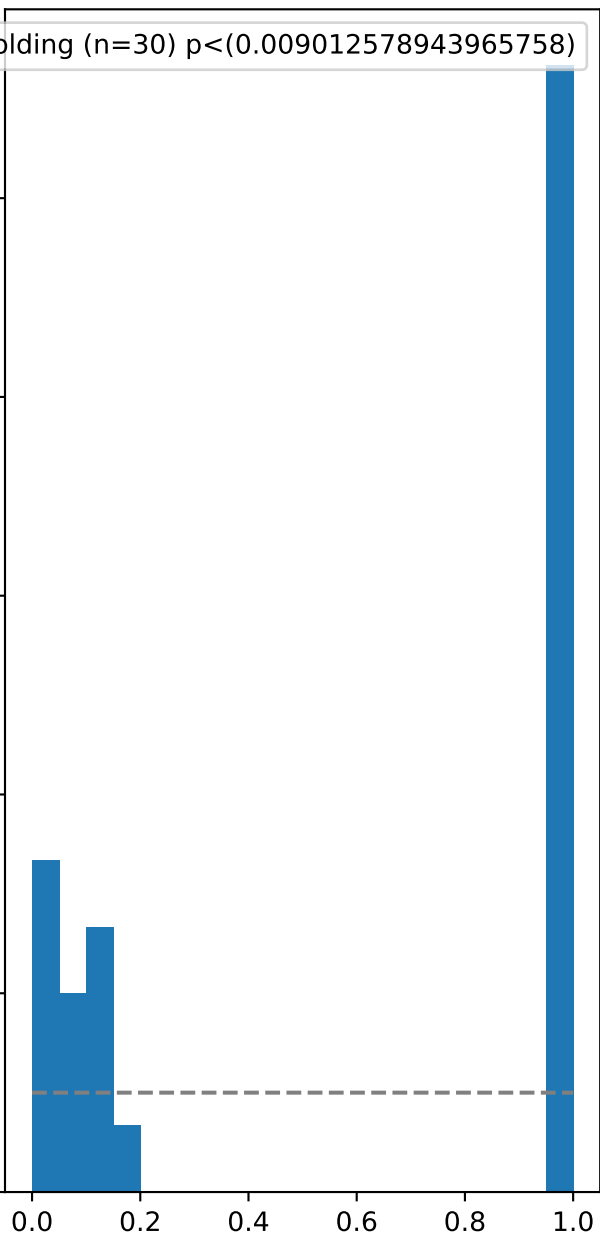
-test.folding (n=30) $p < (0.009012578943965758)$

fraction of repeats with p-value in this bin

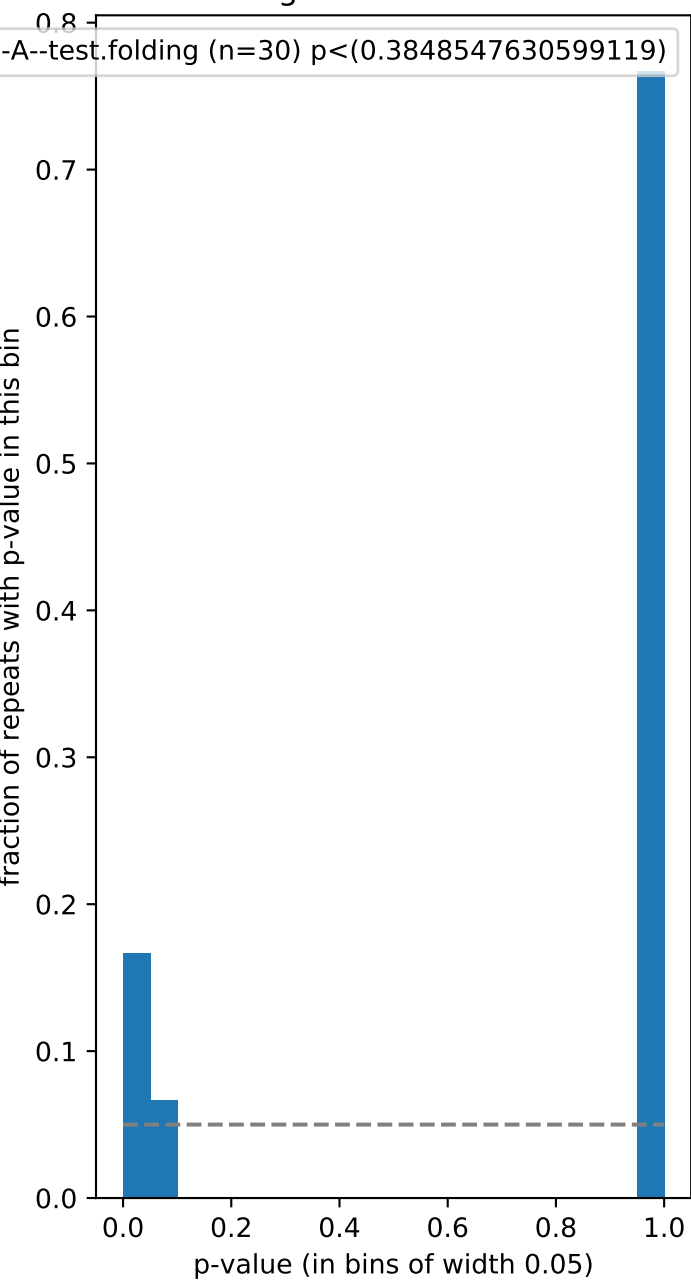
0.5
0.4
0.3
0.2
0.1
0.0

0.0 0.2 0.4 0.6 0.8 1.0

p-value (in bins of width 0.05)

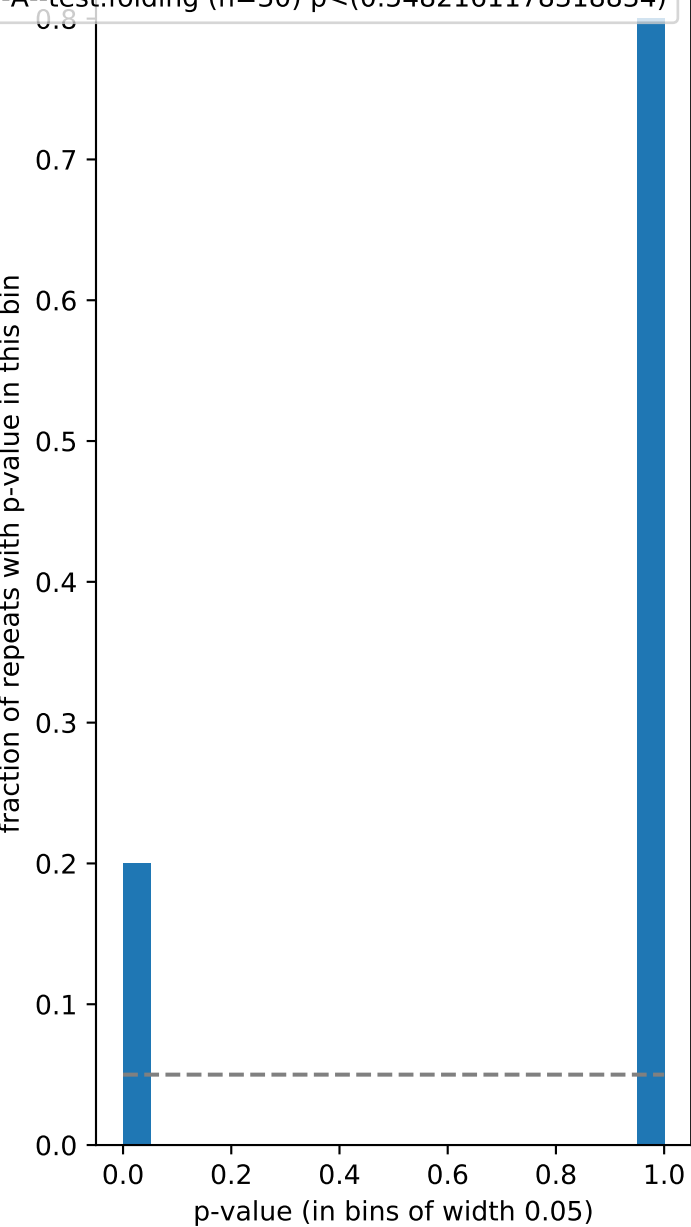


histogram for helix-0-11



histogram for helix-0-12

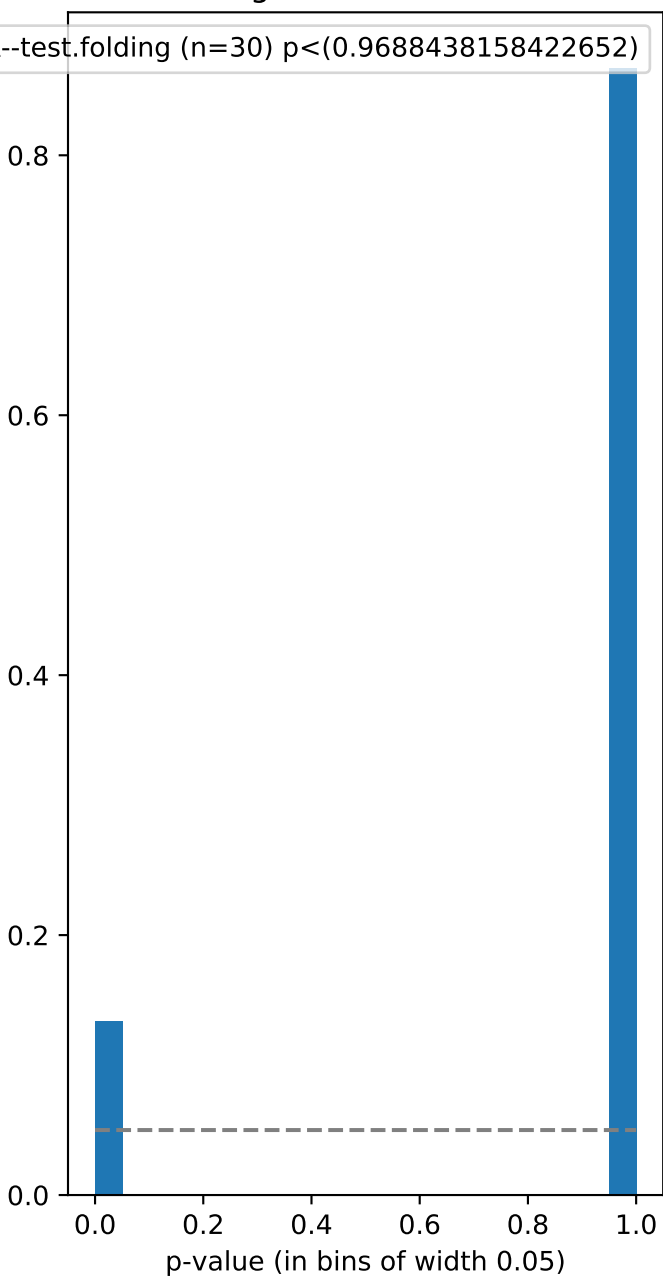
-A--test.folding (n=30) p<(0.5482161178318834)



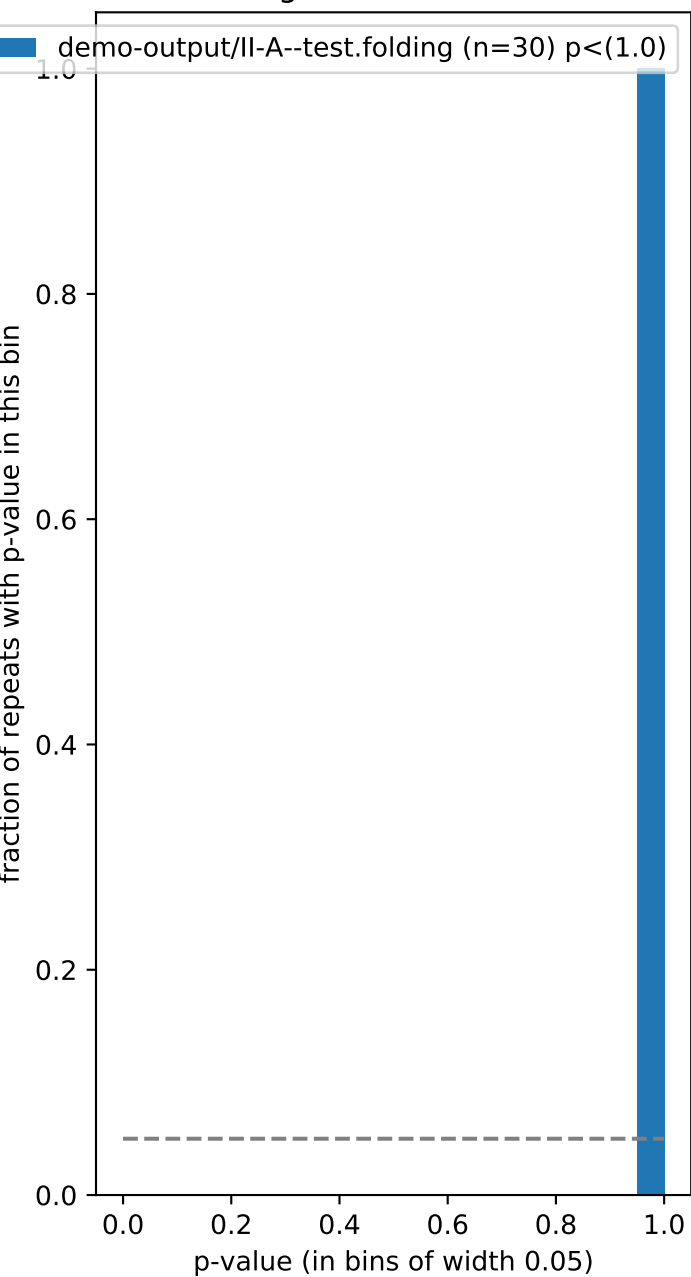
histogram for helix-0-13

-A--test.folding (n=30) $p < (0.9688438158422652)$

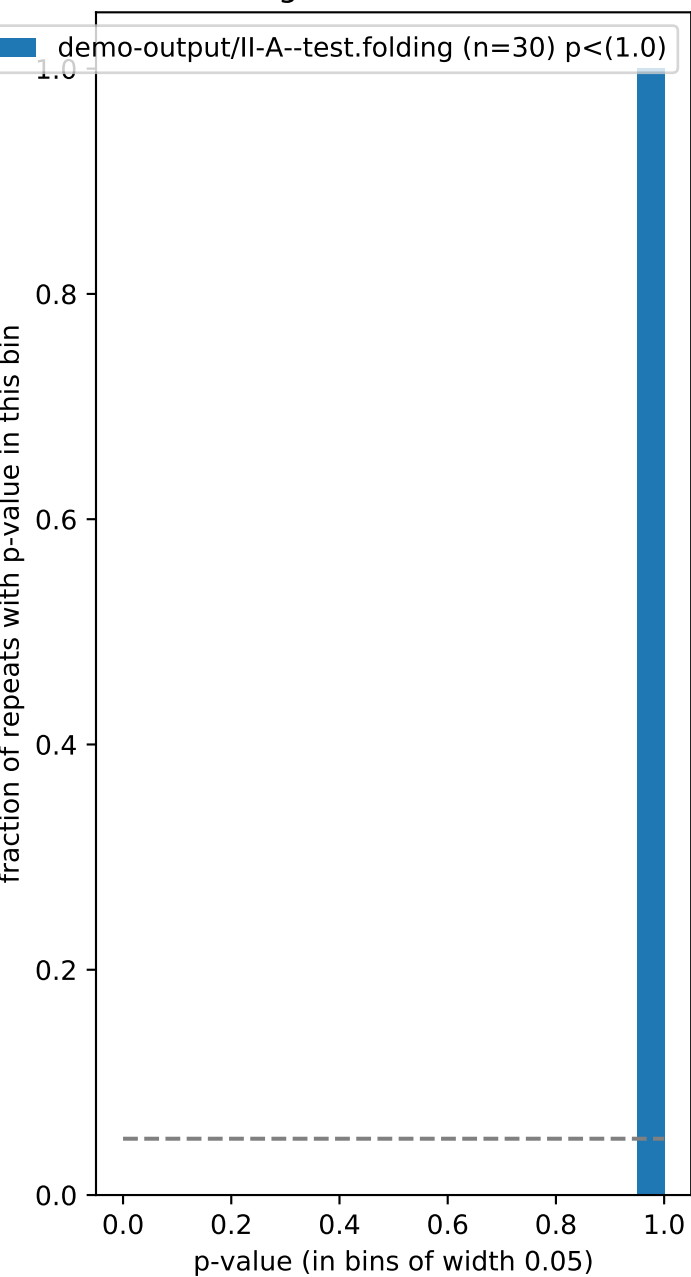
fraction of repeats with p-value in this bin



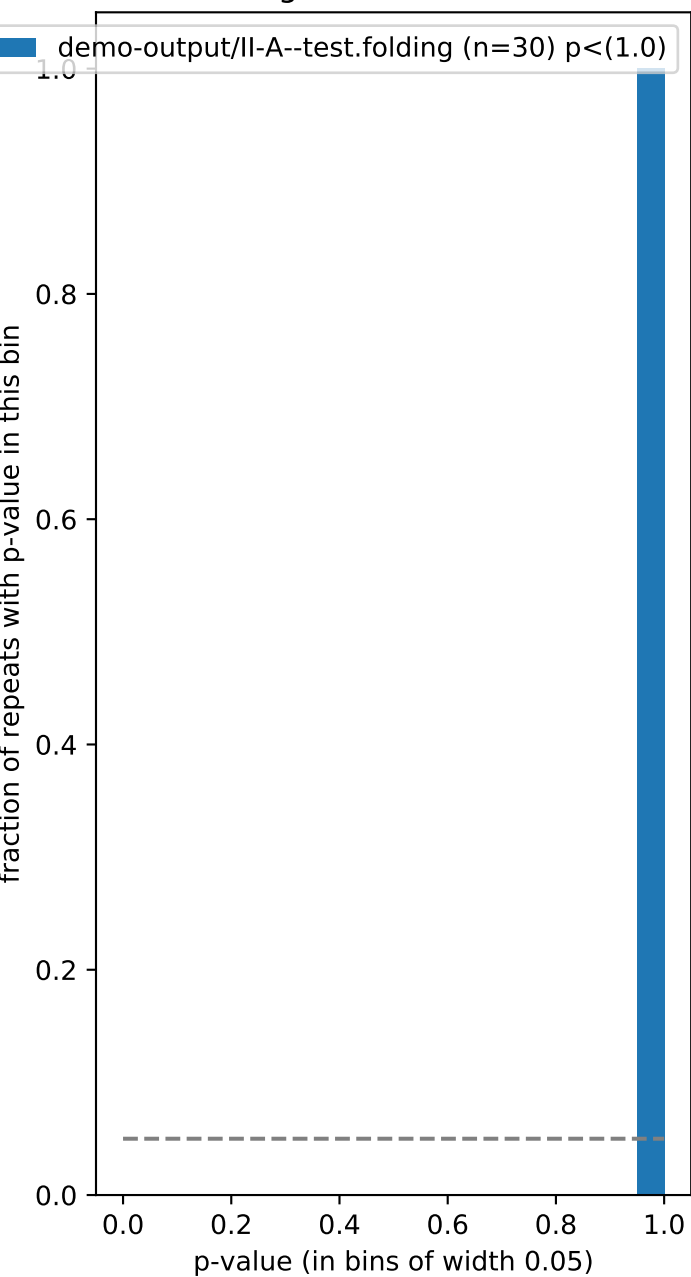
histogram for helix-0-14



histogram for helix-0-15



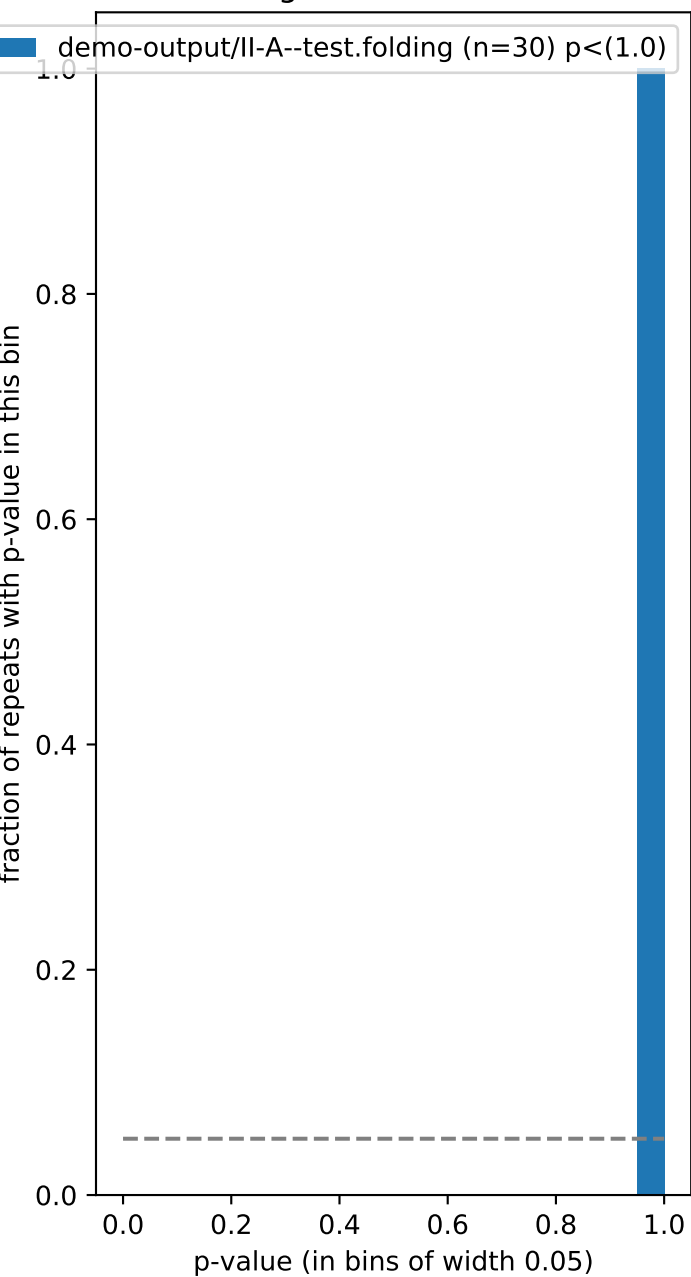
histogram for helix-0-16



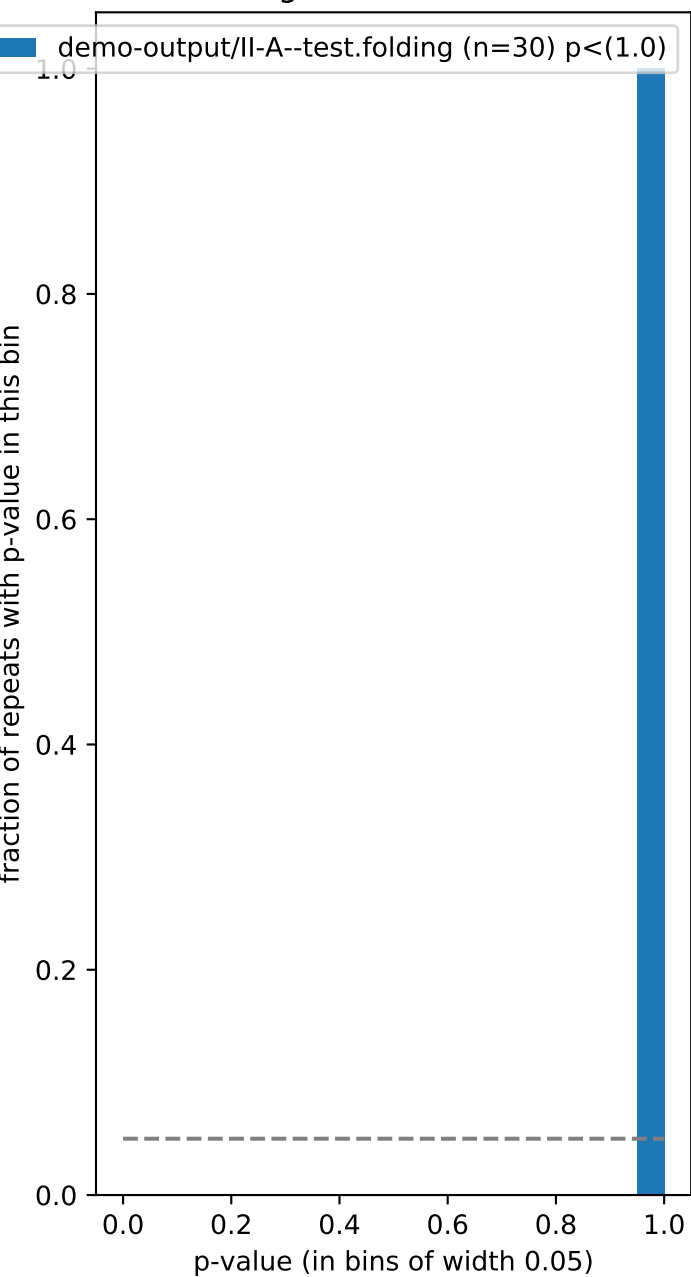
histogram for helix-0-17



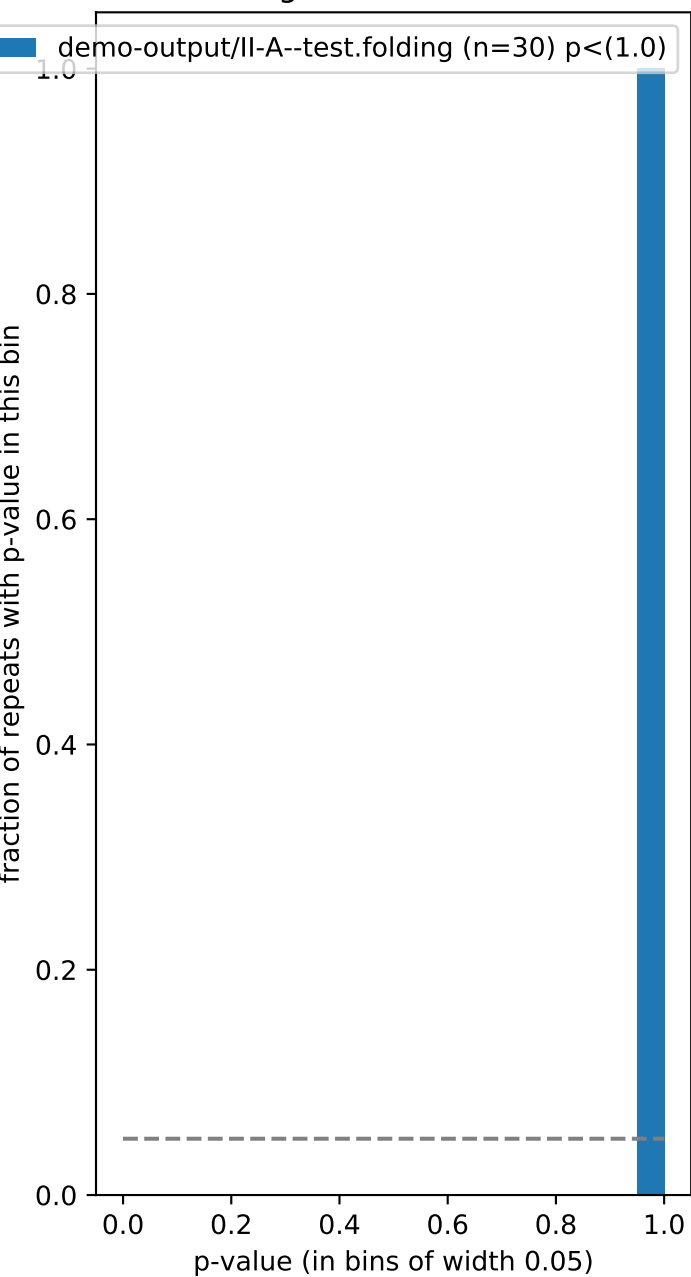
histogram for helix-0-18



histogram for helix-0-19

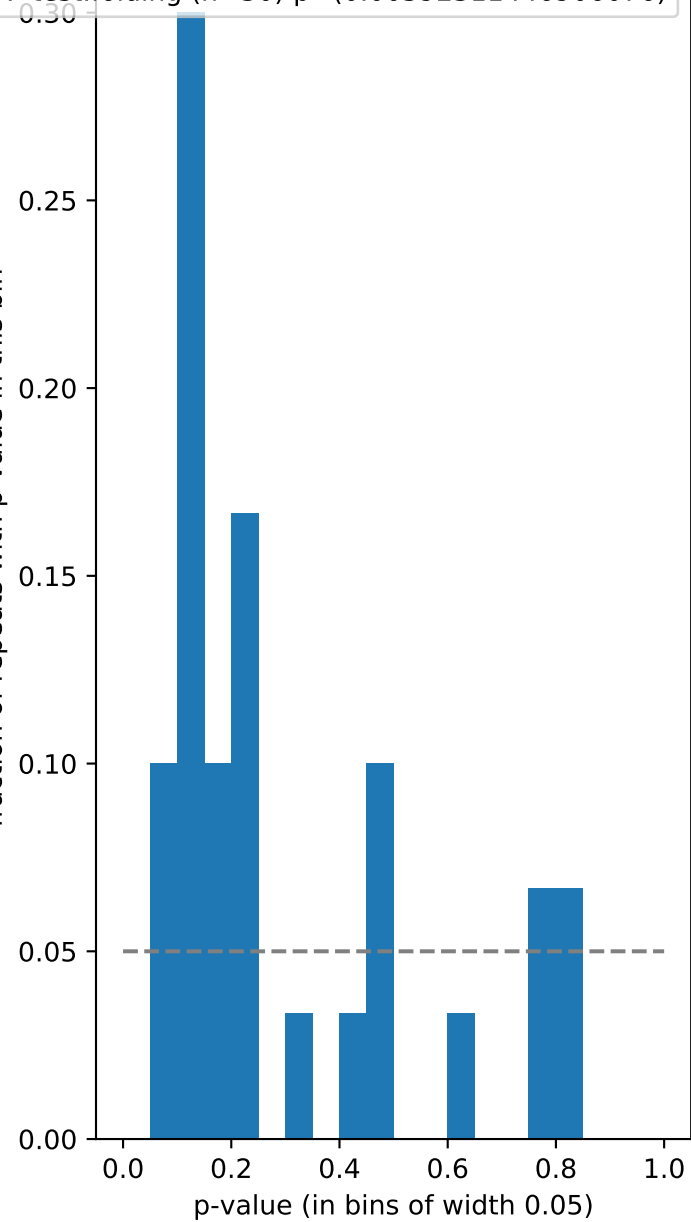


histogram for helix-0-20



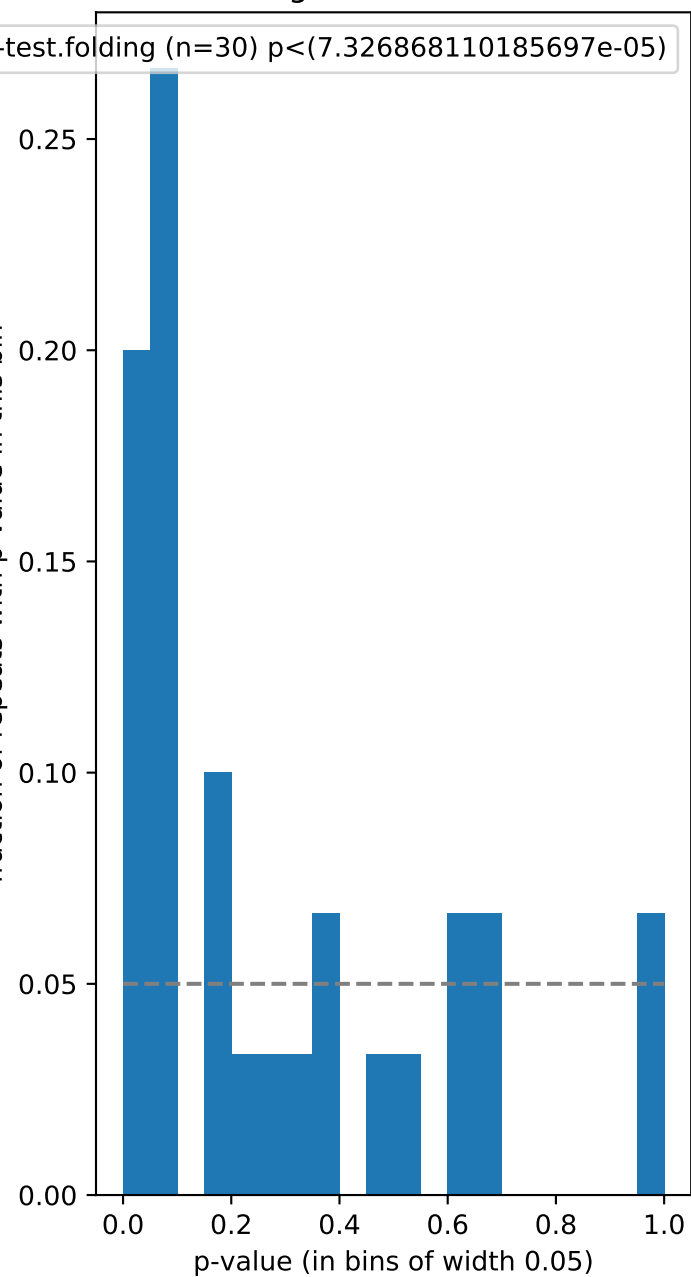
histogram for helix-0-5

A--test.folding (n=30) $p < (0.00592311446906076)$



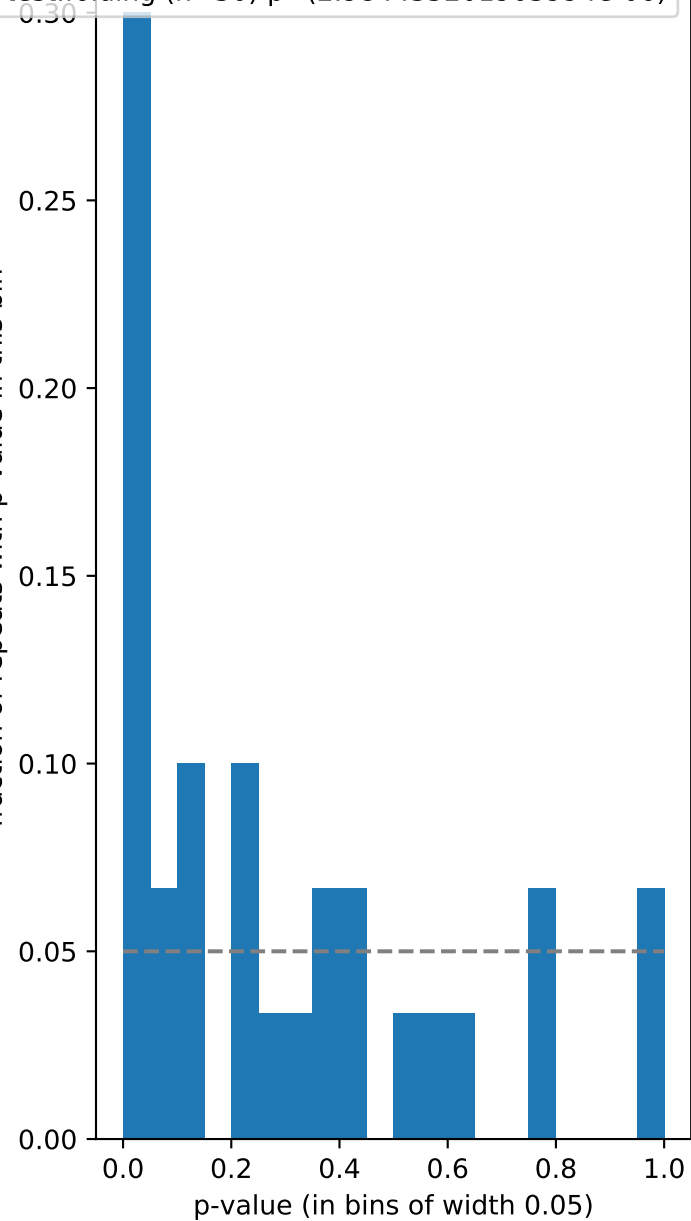
histogram for helix-0-6

test.folding (n=30) $p < (7.326868110185697e-05)$



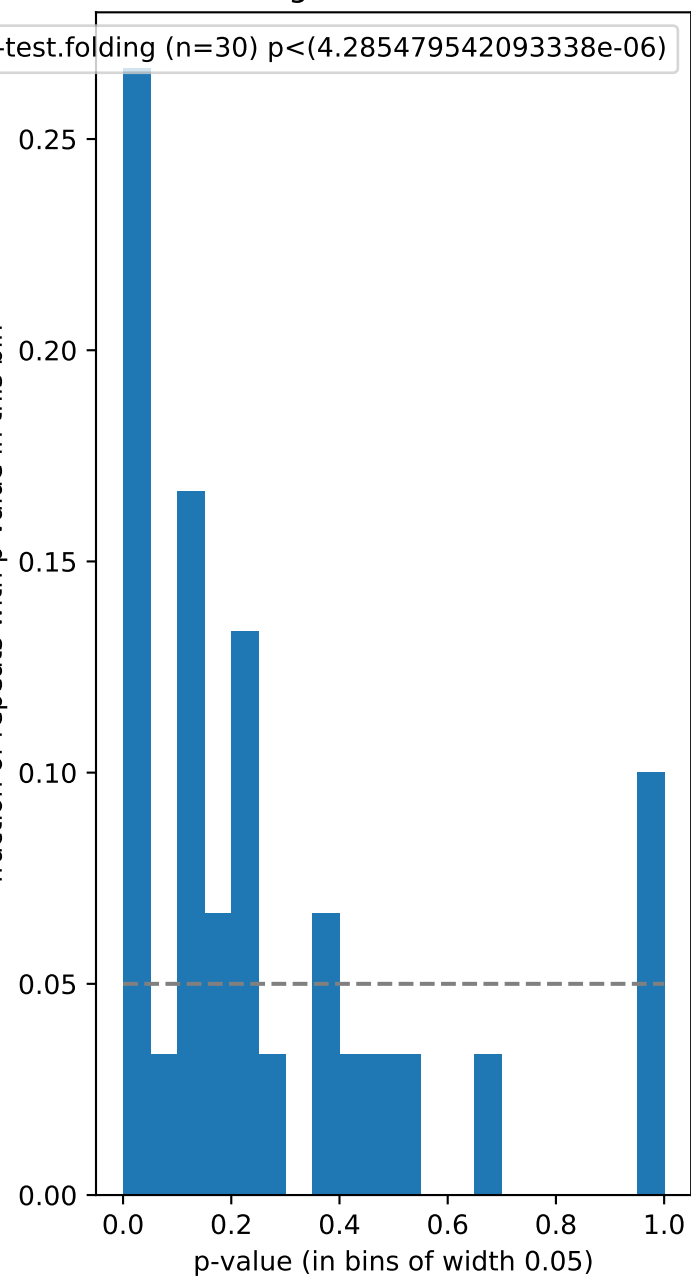
histogram for helix-0-7

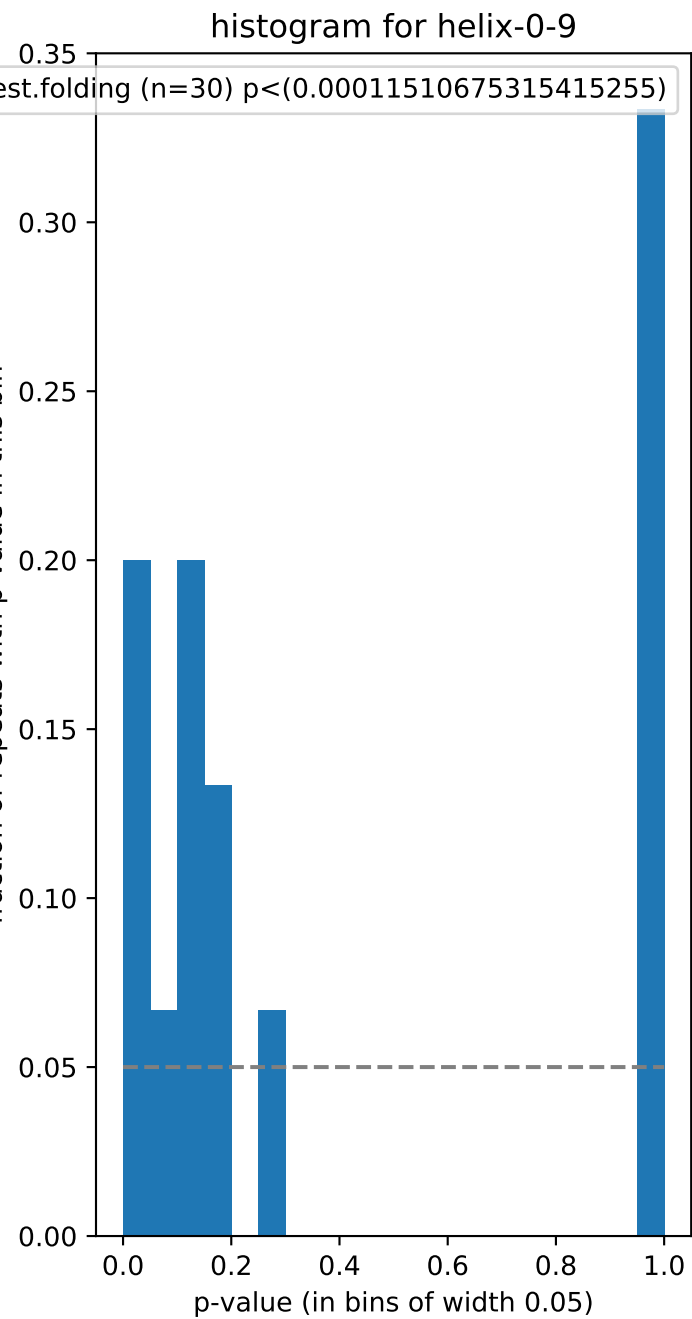
test.folding (n=30) $p < (2.984435201903994e-06)$



histogram for helix-0-8

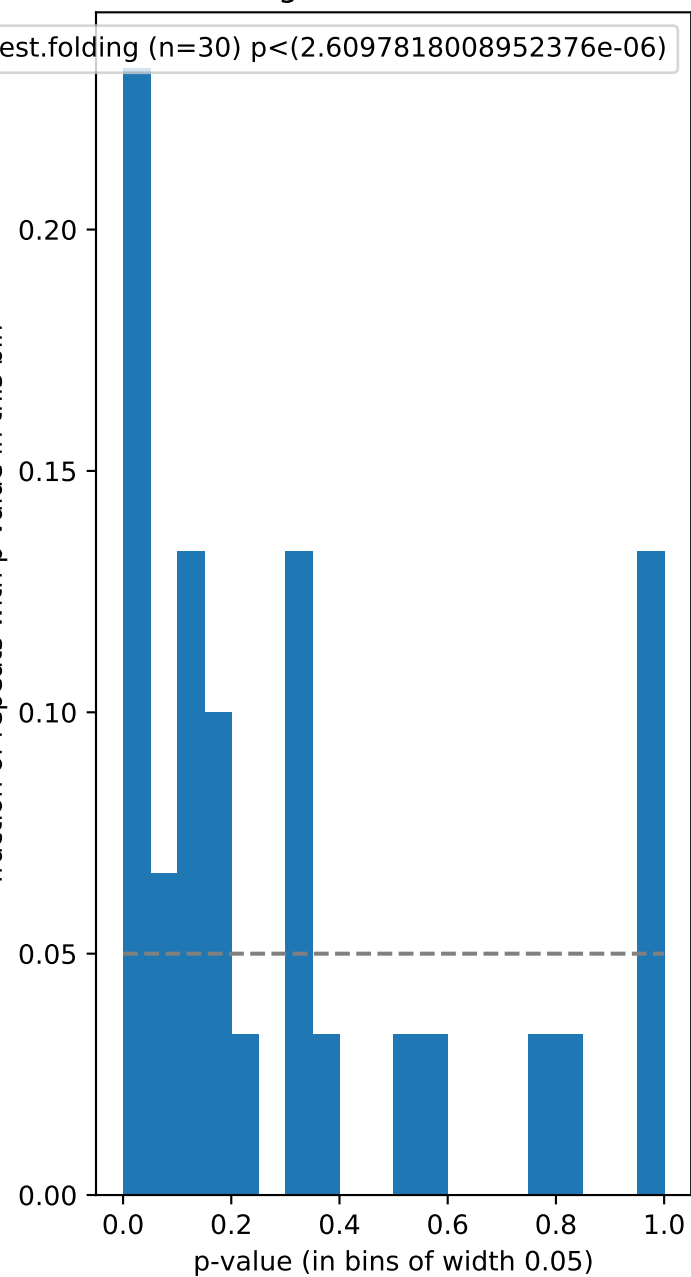
test.folding (n=30) $p < (4.285479542093338e-06)$





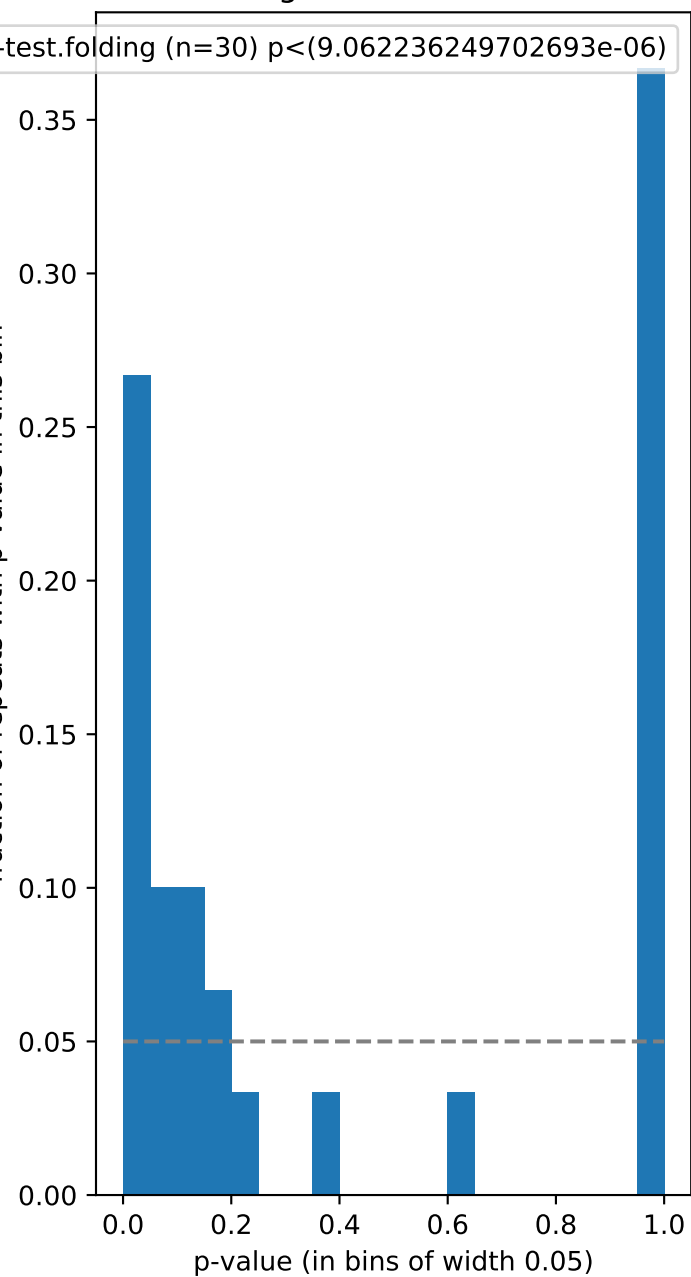
histogram for helix-1-10

est.folding (n=30) $p < (2.6097818008952376e-06)$



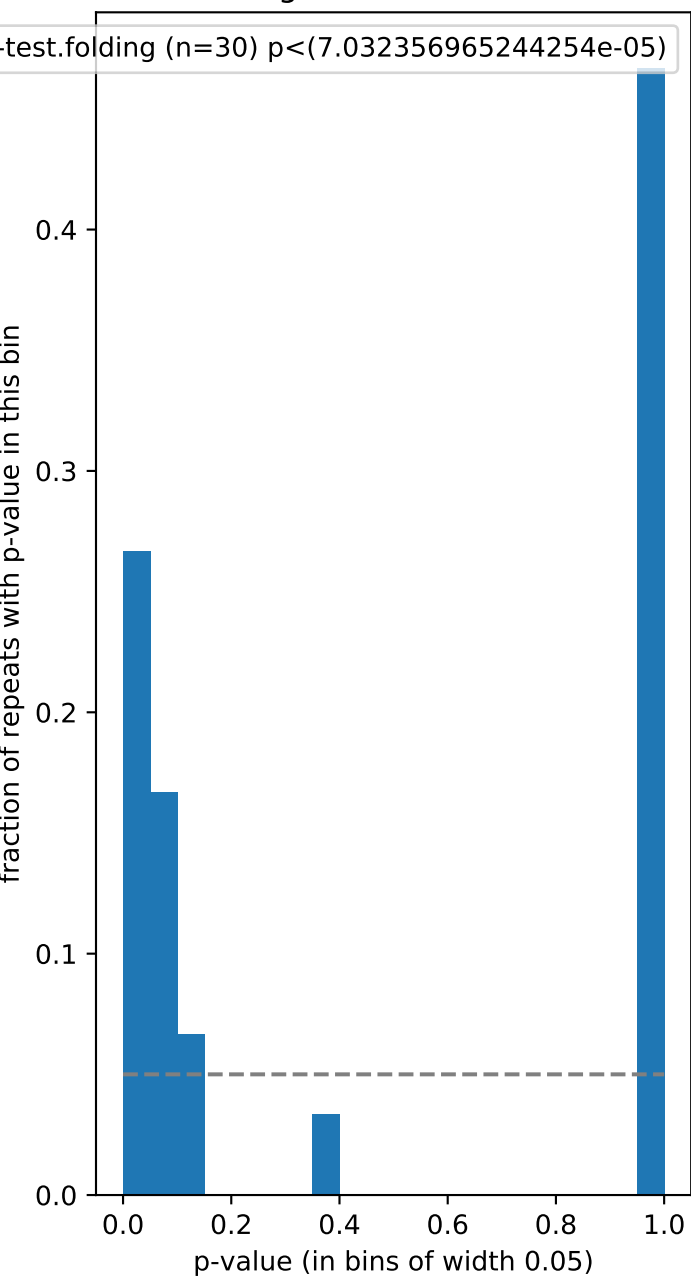
histogram for helix-1-11

test.folding (n=30) $p < (9.062236249702693e-06)$



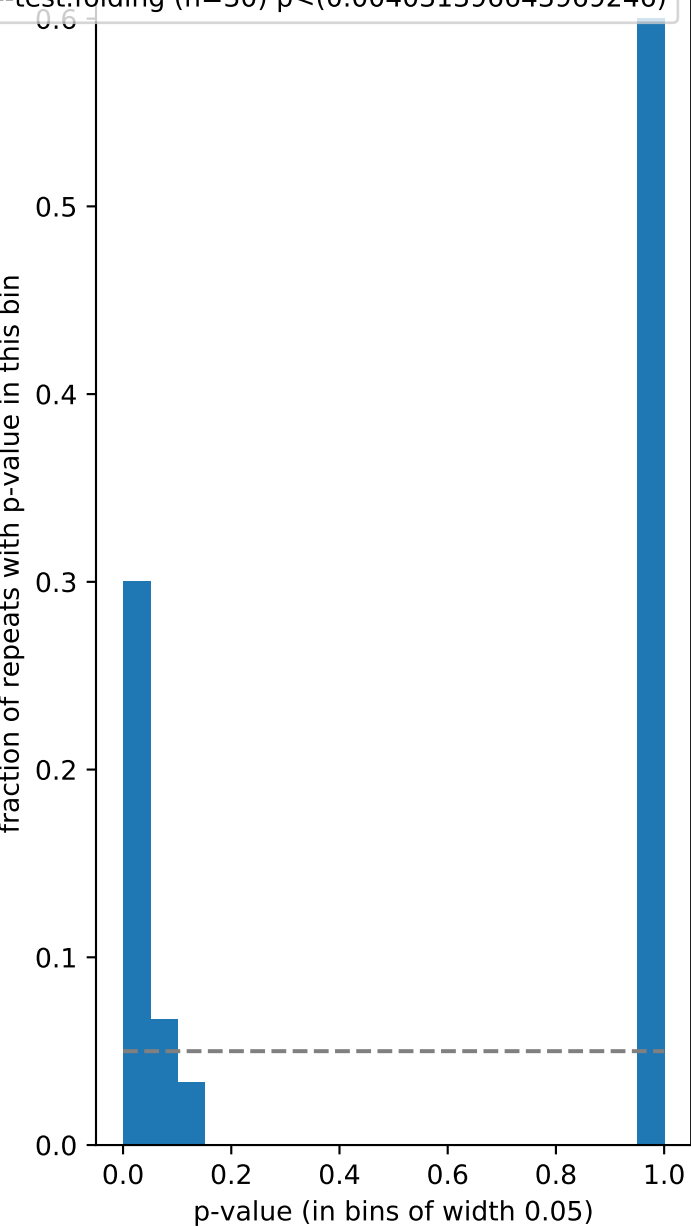
histogram for helix-1-12

test.folding (n=30) $p < (7.032356965244254e-05)$



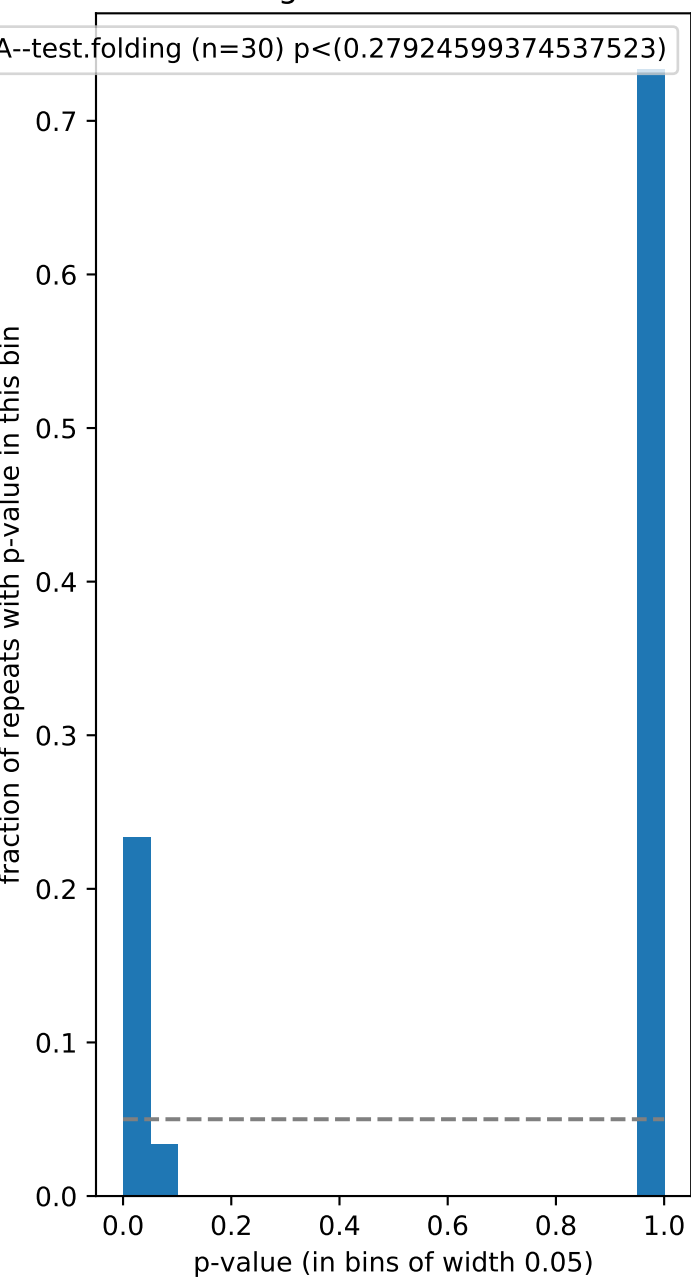
histogram for helix-1-13

-test.folding (n=30) $p < (0.004031396643969246)$



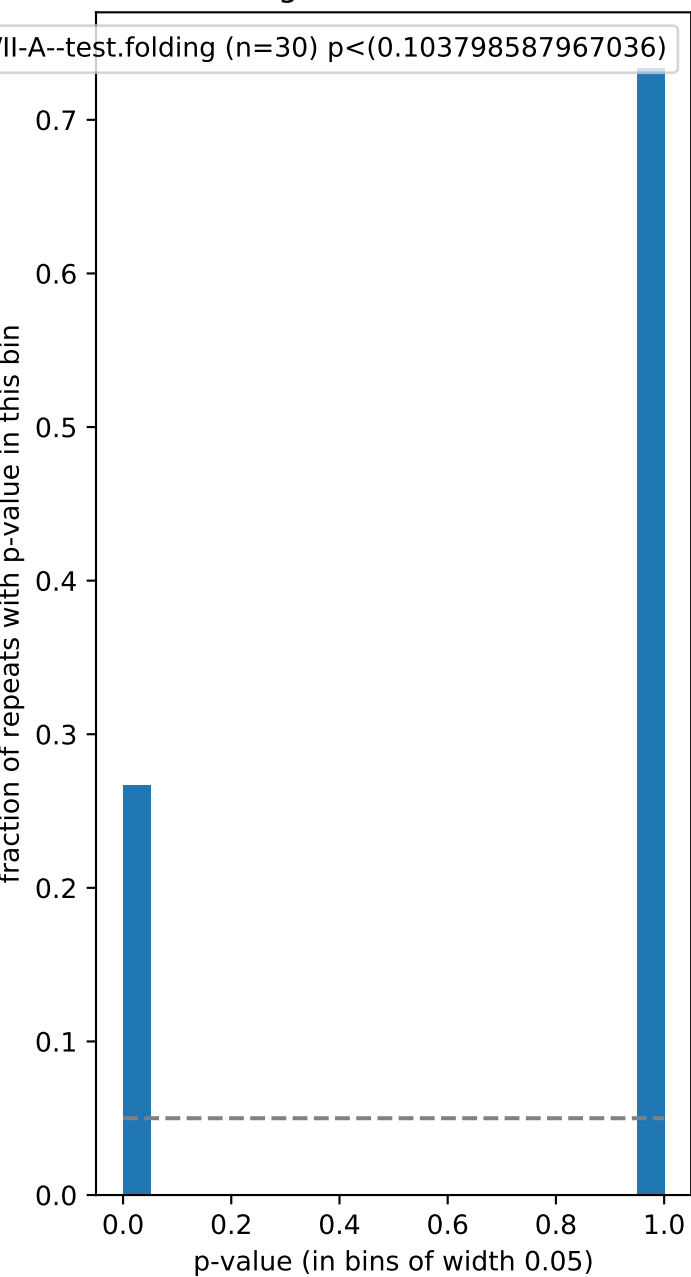
histogram for helix-1-14

A--test.folding (n=30) $p < (0.27924599374537523)$



histogram for helix-1-15

II-A--test.folding (n=30) $p < (0.103798587967036)$



histogram for helix-1-16

-A--test.folding (n=30) $p < (0.9998822688835856)$

fraction of repeats with p-value in this bin

0.8

0.6

0.4

0.2

0.0

0.0

0.2

0.4

0.6

0.8

1.0

p-value (in bins of width 0.05)

0.1

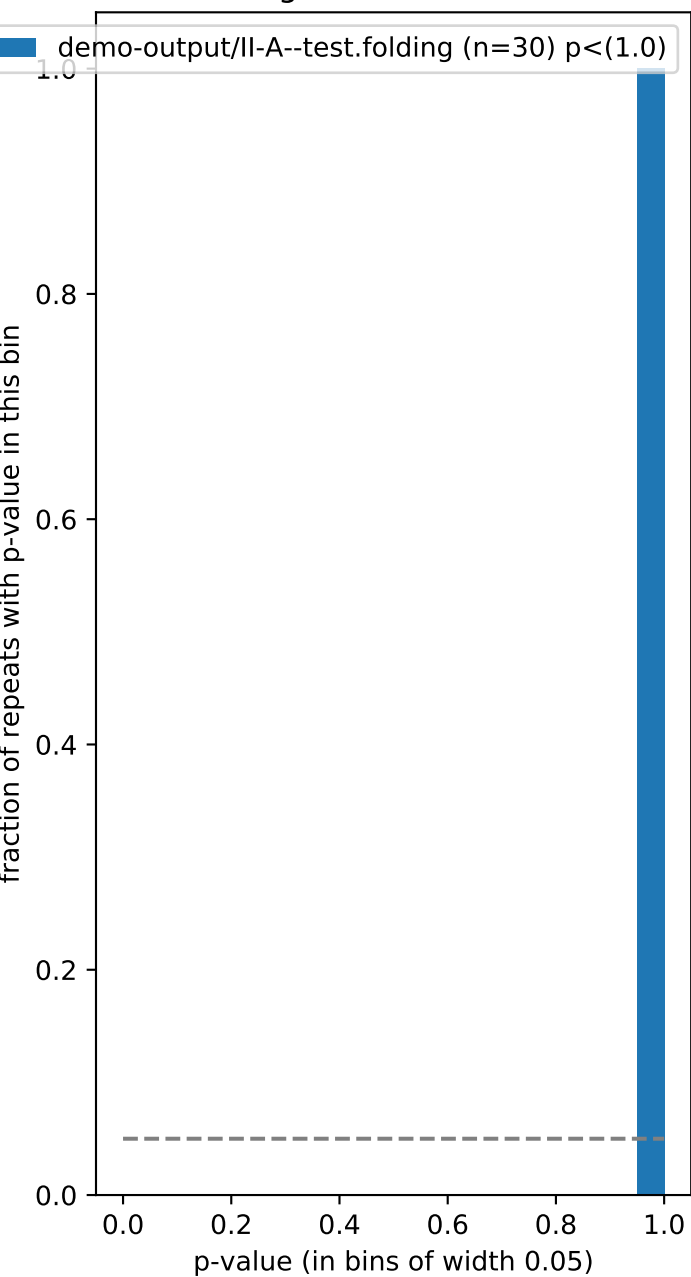
0.9

0.05 0.1 0.15 0.2 0.25 0.3 0.35 0.4 0.45 0.5 0.55 0.6 0.65 0.7 0.75 0.8 0.85 0.9 0.95 1.0

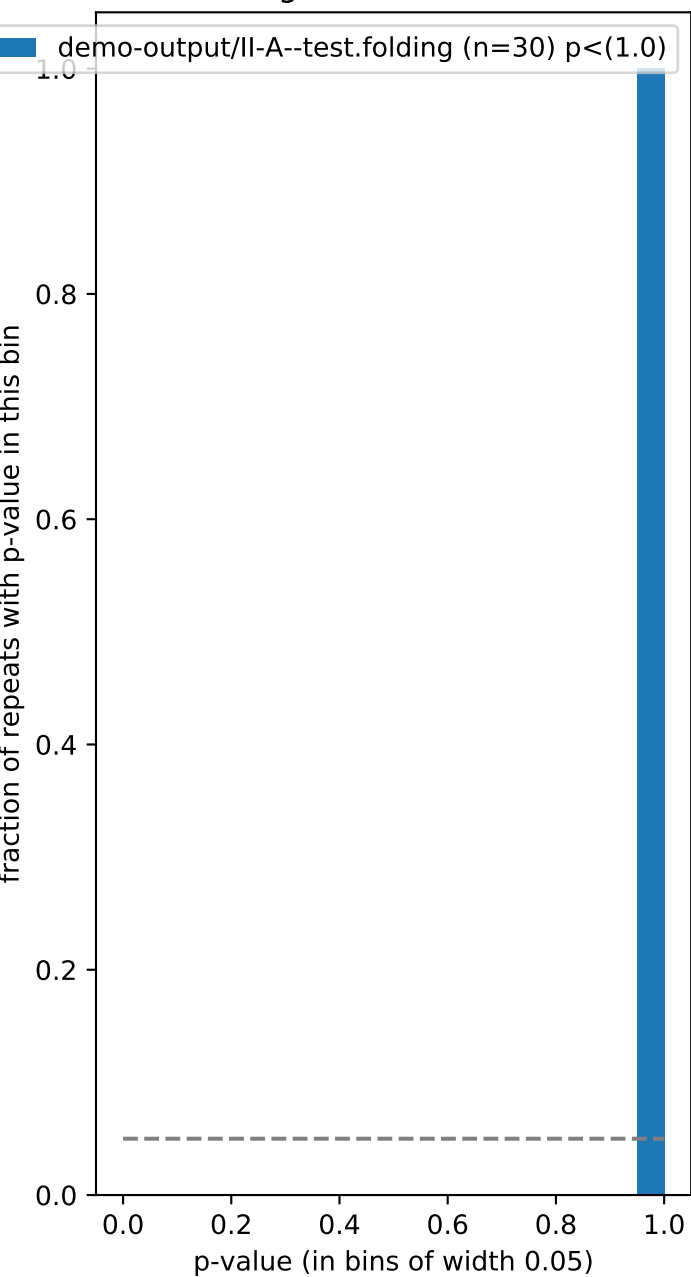
histogram for helix-1-17



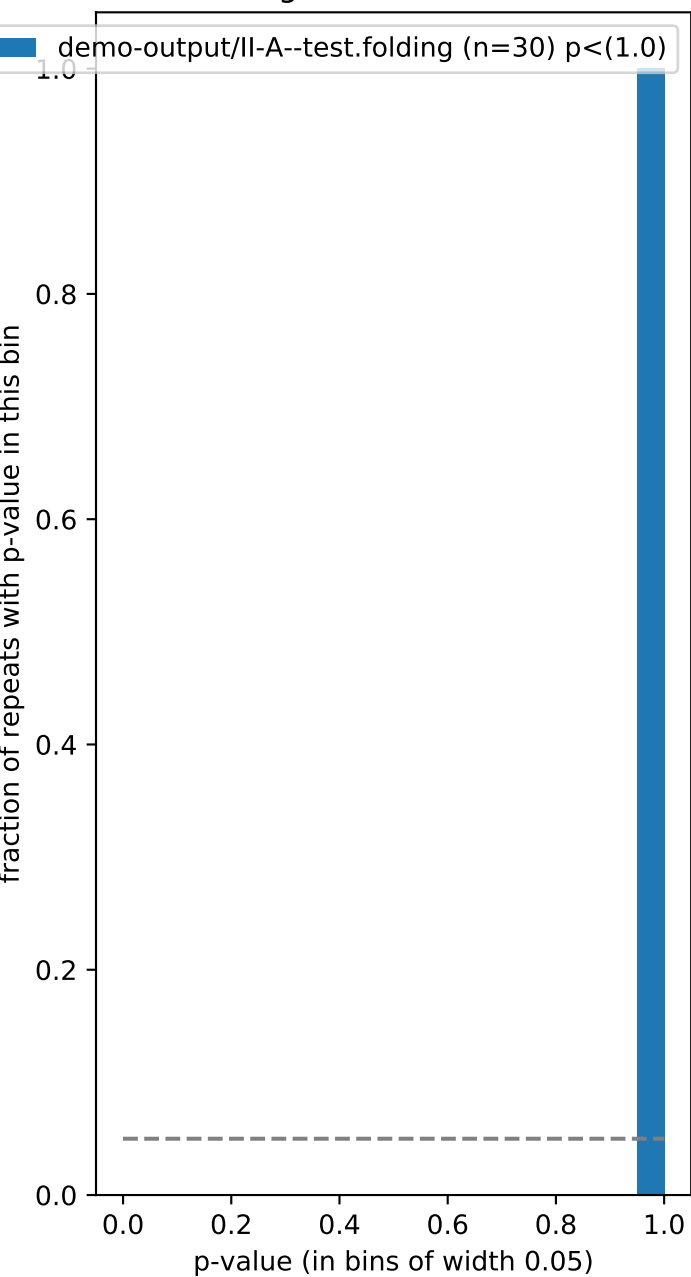
histogram for helix-1-18



histogram for helix-1-19

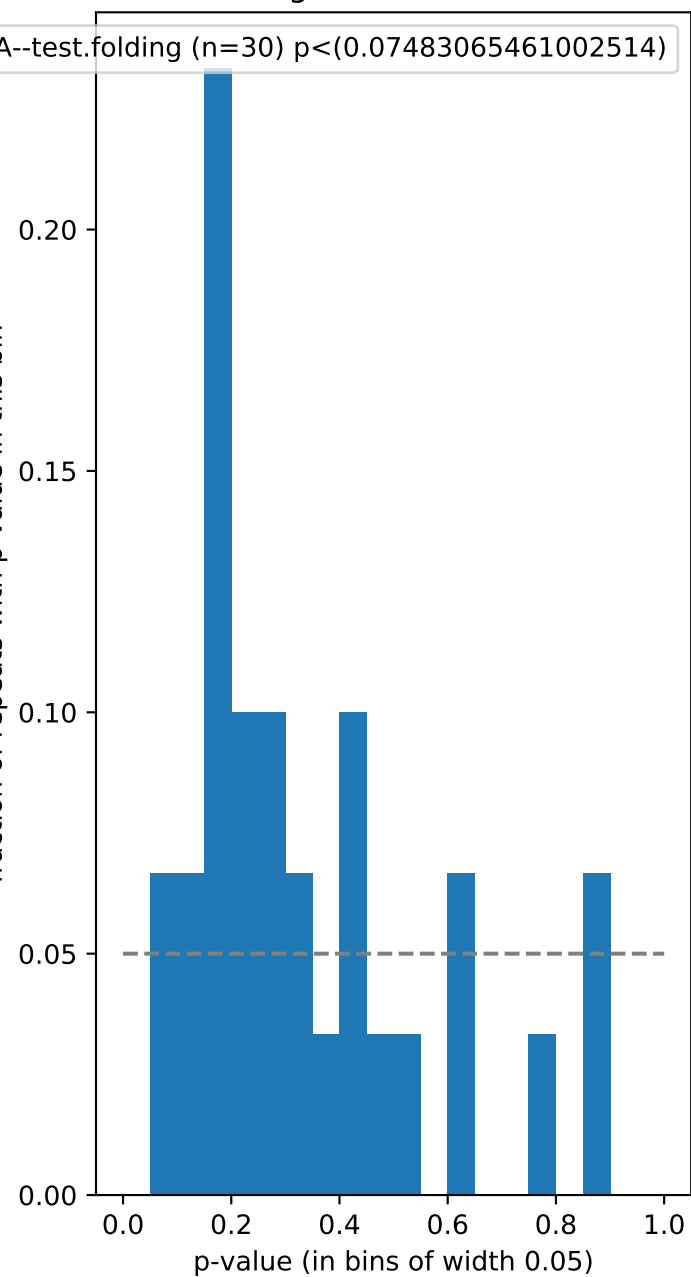


histogram for helix-1-20



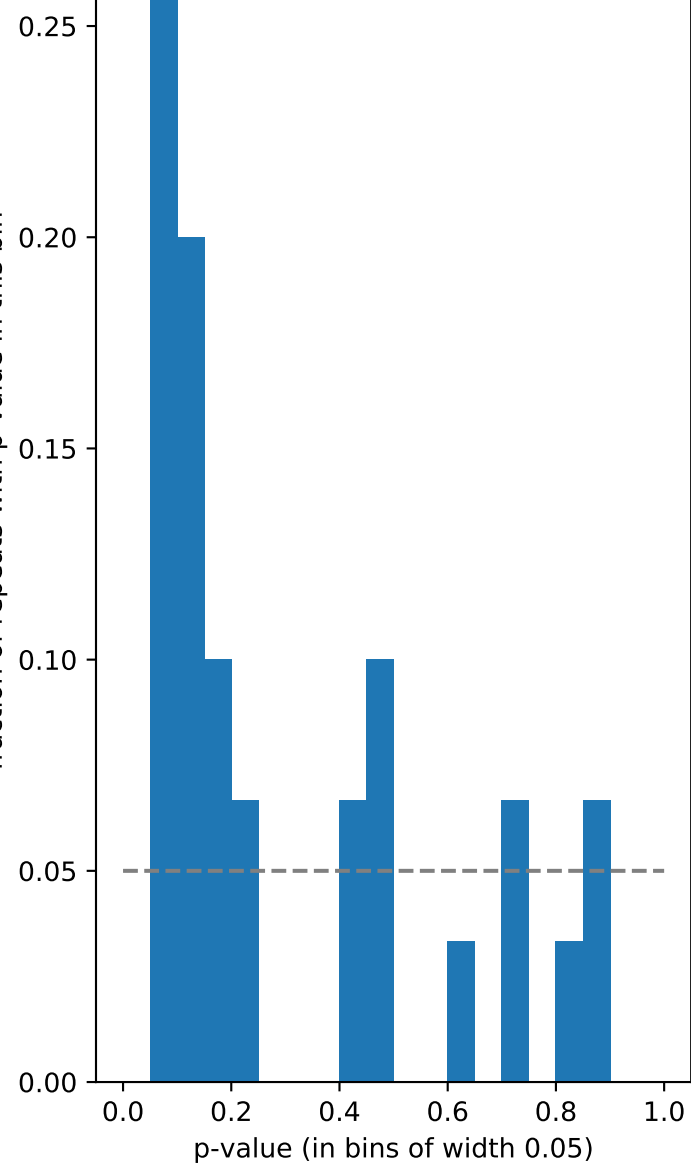
histogram for helix-1-5

A--test.folding (n=30) $p < (0.07483065461002514)$

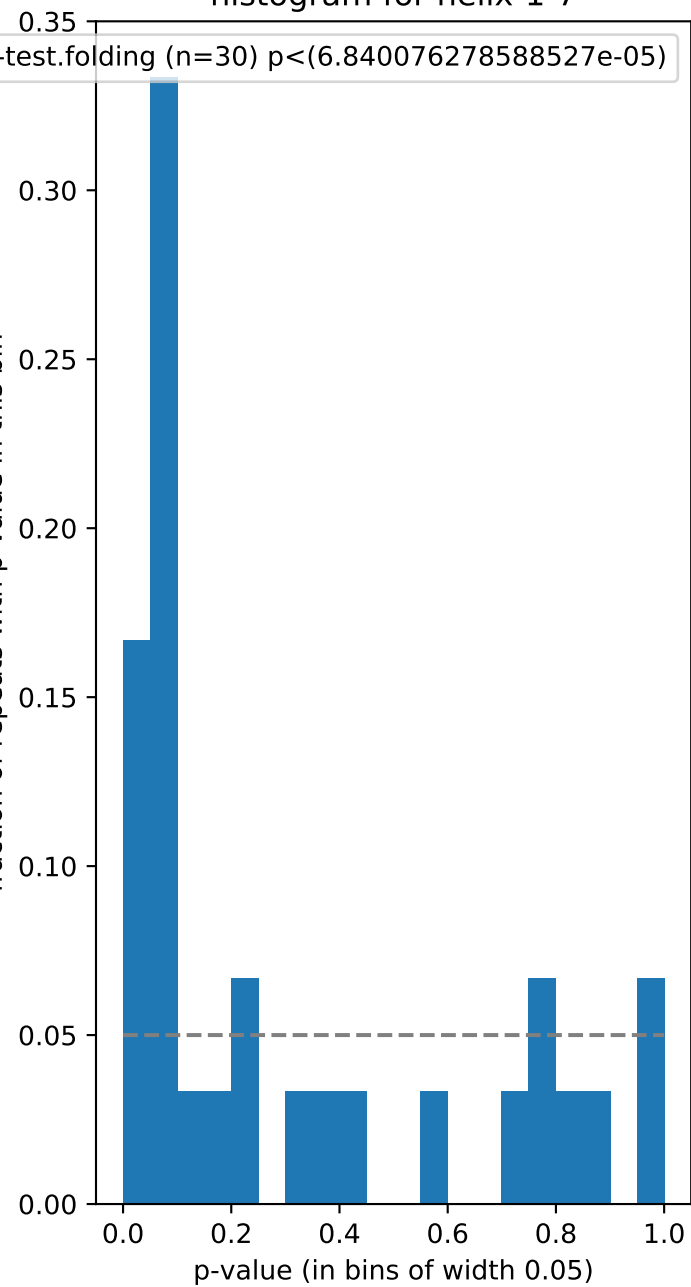


histogram for helix-1-6

-test.folding (n=30) $p < (0.004132449282940288)$

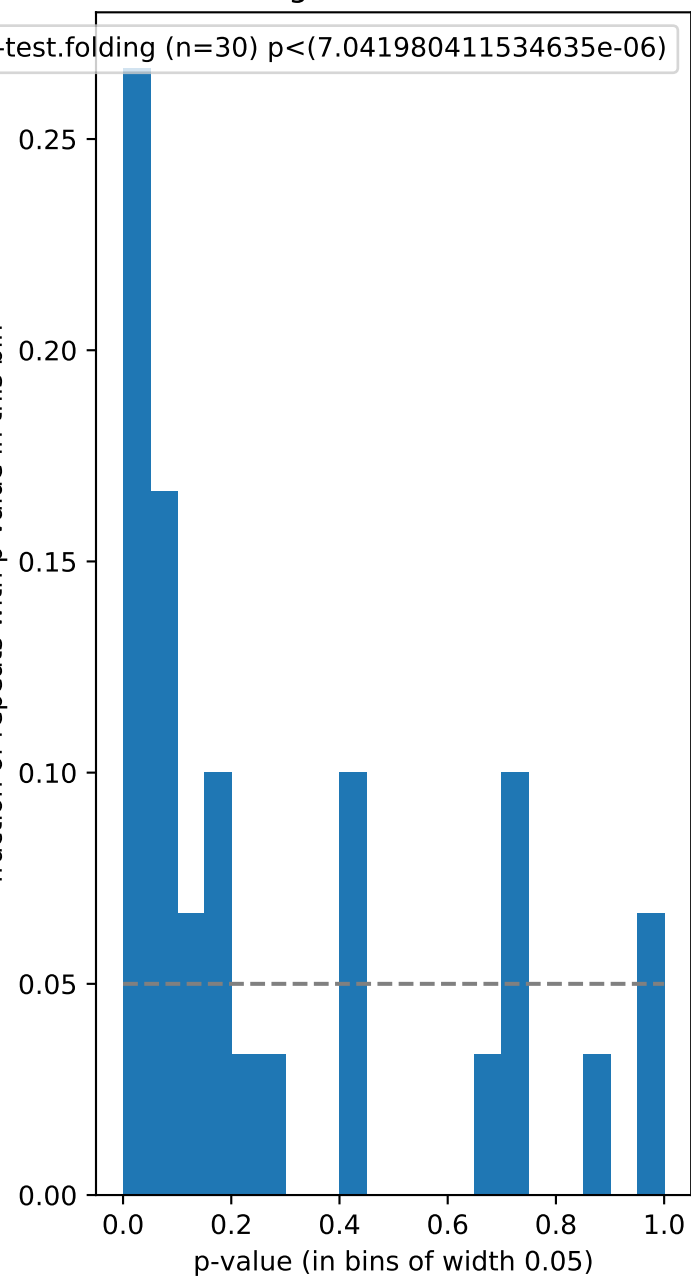


histogram for helix-1-7



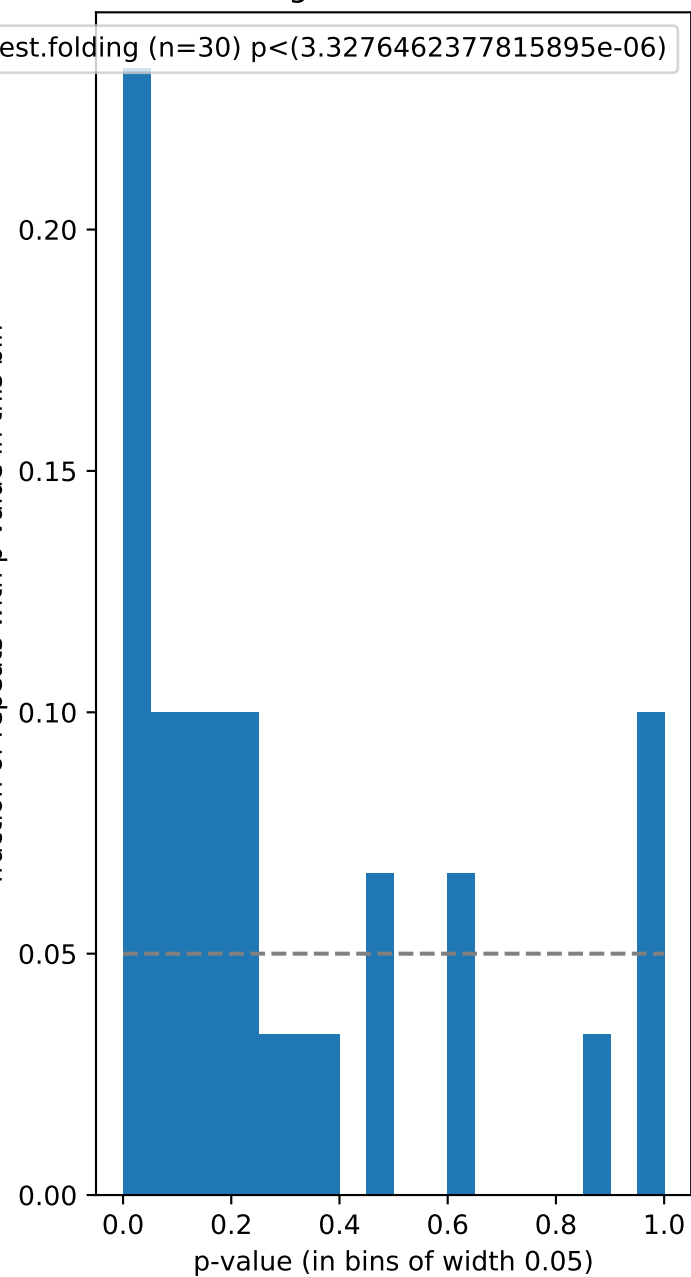
histogram for helix-1-8

test.folding (n=30) $p < (7.041980411534635e-06)$



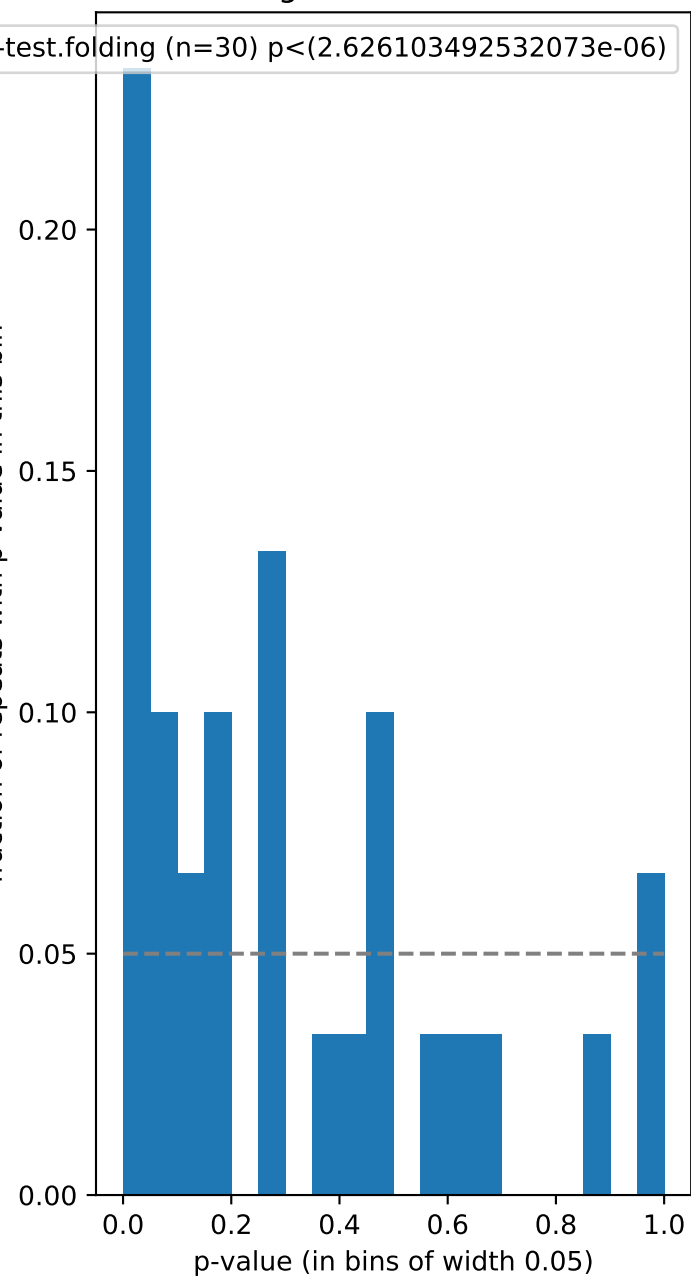
histogram for helix-1-9

est.folding (n=30) $p < (3.3276462377815895e-06)$



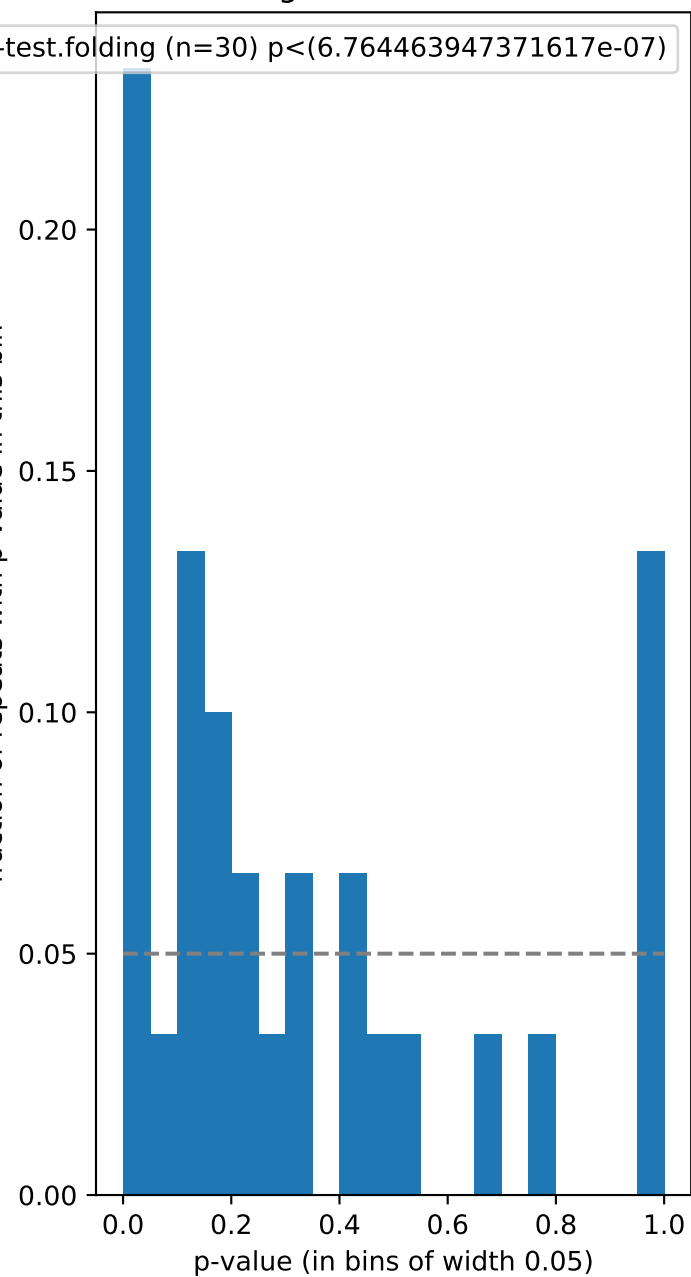
histogram for helix-2-10

test.folding (n=30) $p < (2.626103492532073e-06)$



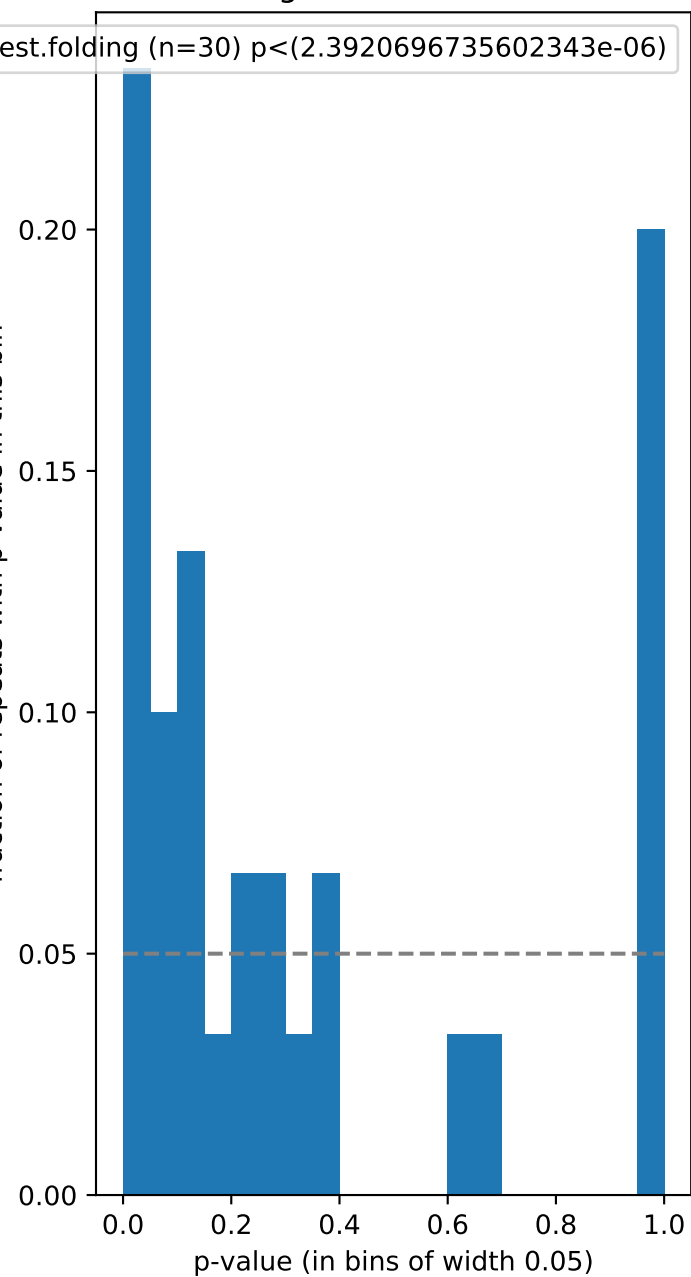
histogram for helix-2-11

test.folding (n=30) $p < (6.764463947371617e-07)$



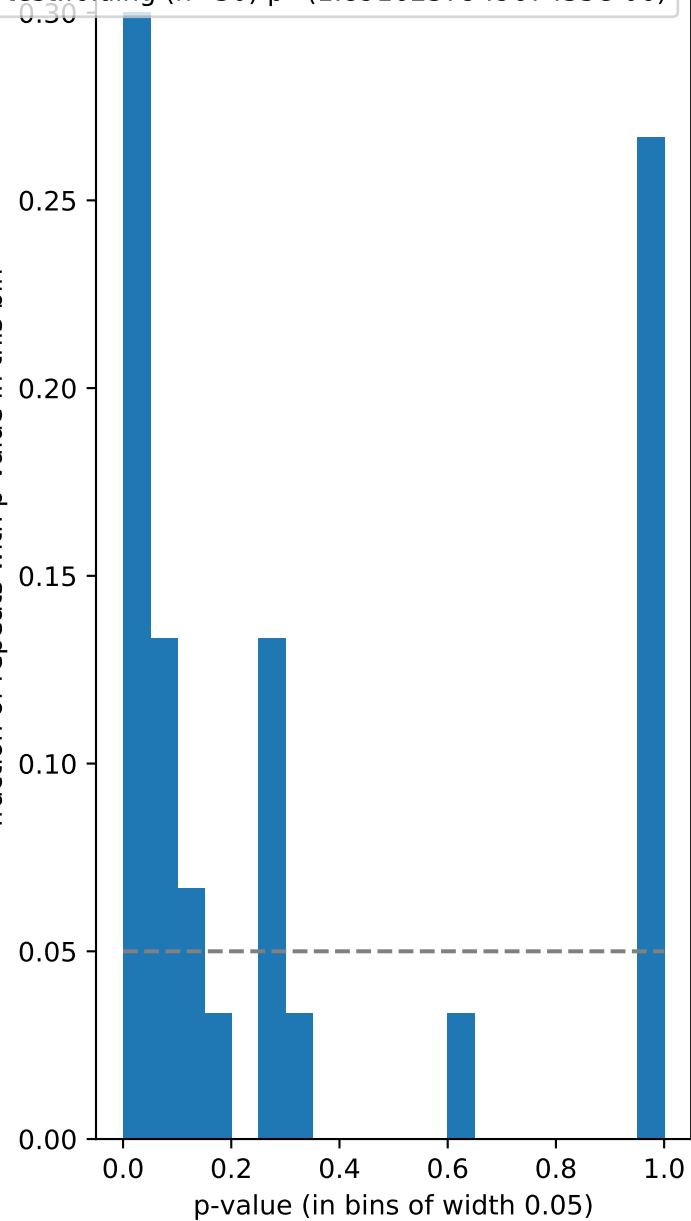
histogram for helix-2-12

est.folding (n=30) $p < (2.3920696735602343e-06)$



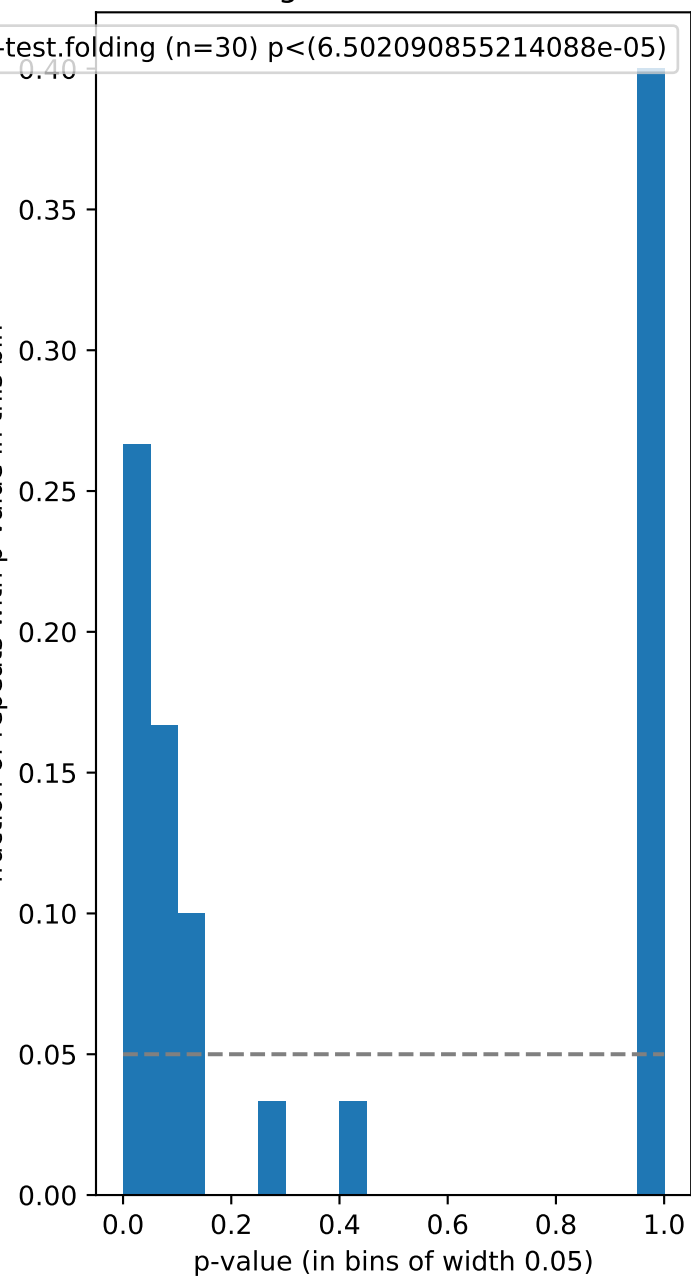
histogram for helix-2-13

test.folding (n=30) $p < (1.891623794967455e-06)$



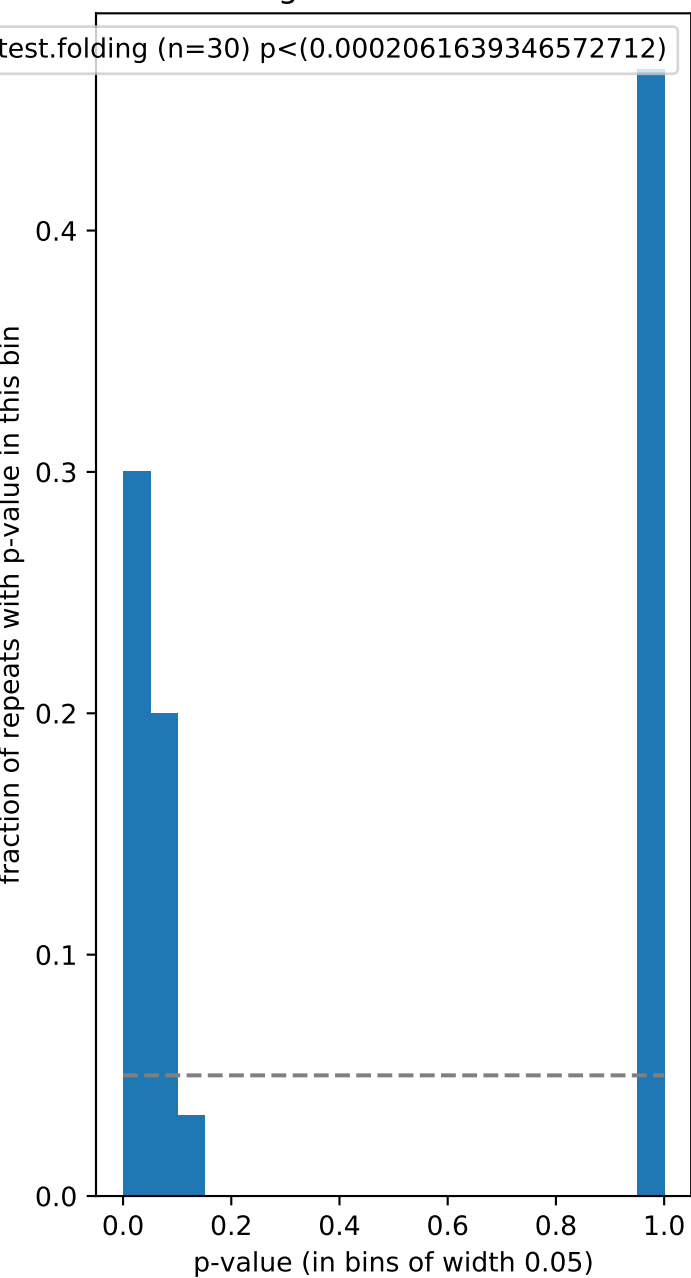
histogram for helix-2-14

test.folding (n=30) $p < (6.502090855214088e-05)$



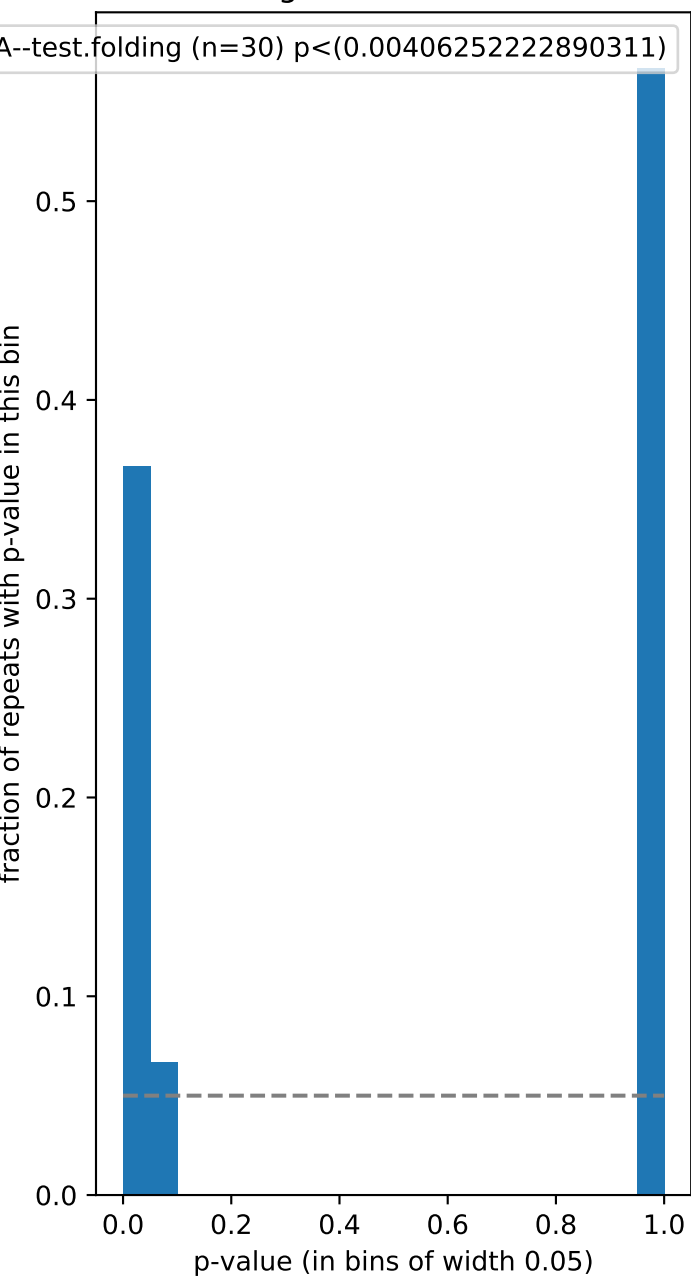
histogram for helix-2-15

test.folding (n=30) $p < (0.0002061639346572712)$



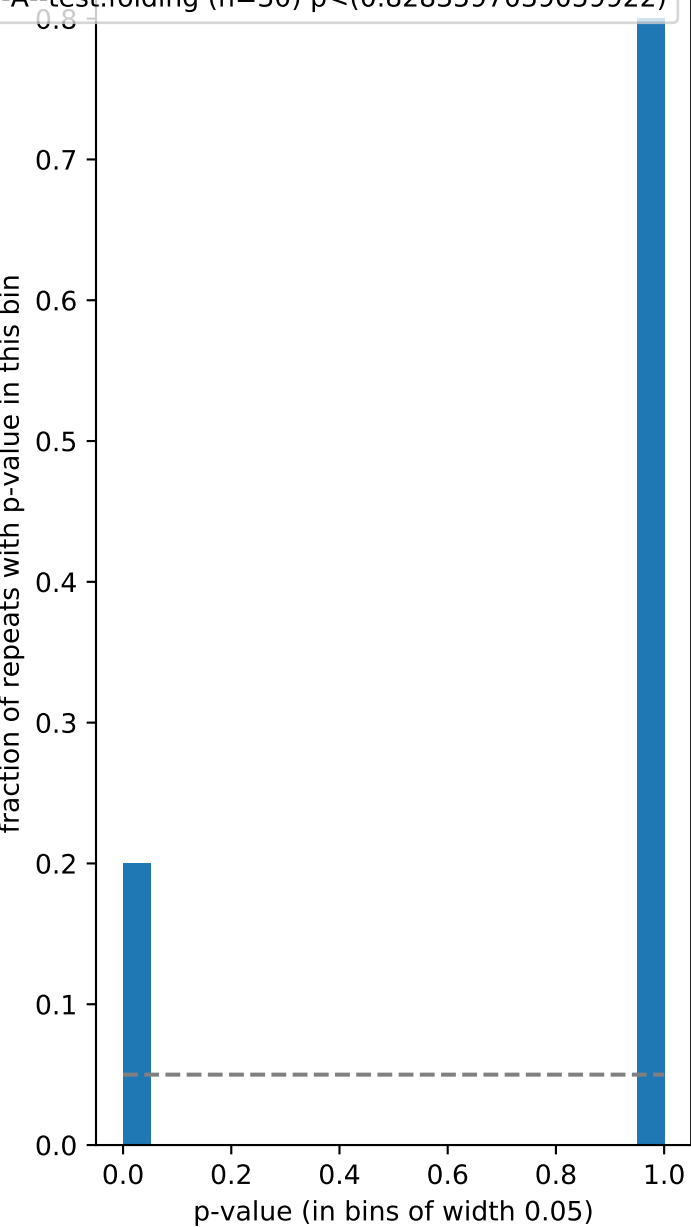
histogram for helix-2-16

A--test.folding (n=30) $p < (0.00406252222890311)$



histogram for helix-2-17

-A--test.folding (n=30) $p < (0.8283597039059922)$



histogram for helix-2-18

-A--test.folding (n=30) $p < (0.9999511175171857)$

fraction of repeats with p-value in this bin

0.8

0.6

0.4

0.2

0.0

0.0

0.2

0.4

0.6

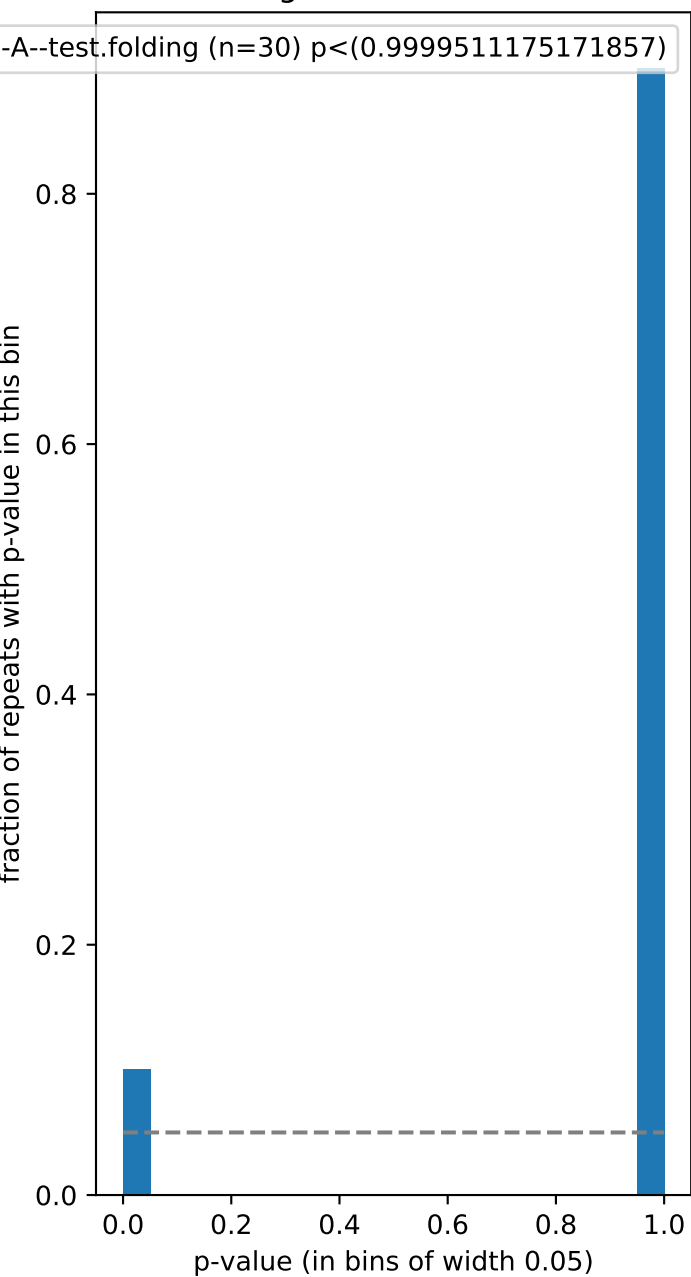
0.8

1.0

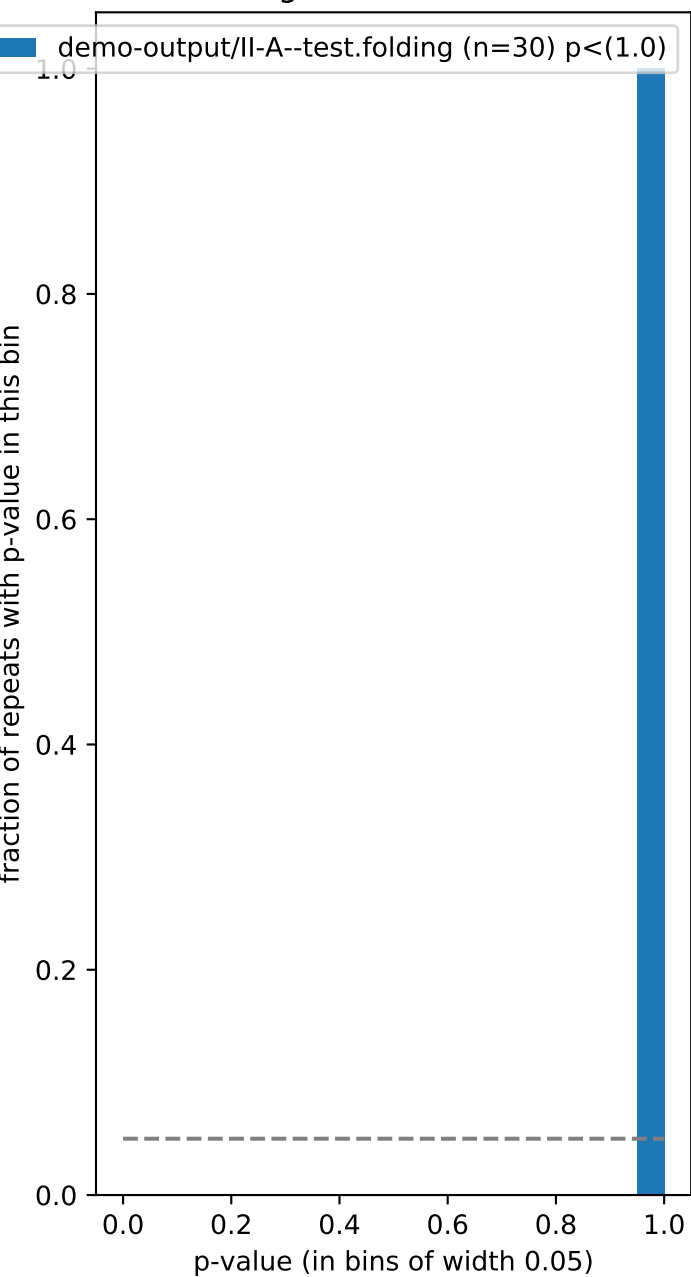
p-value (in bins of width 0.05)

0.1

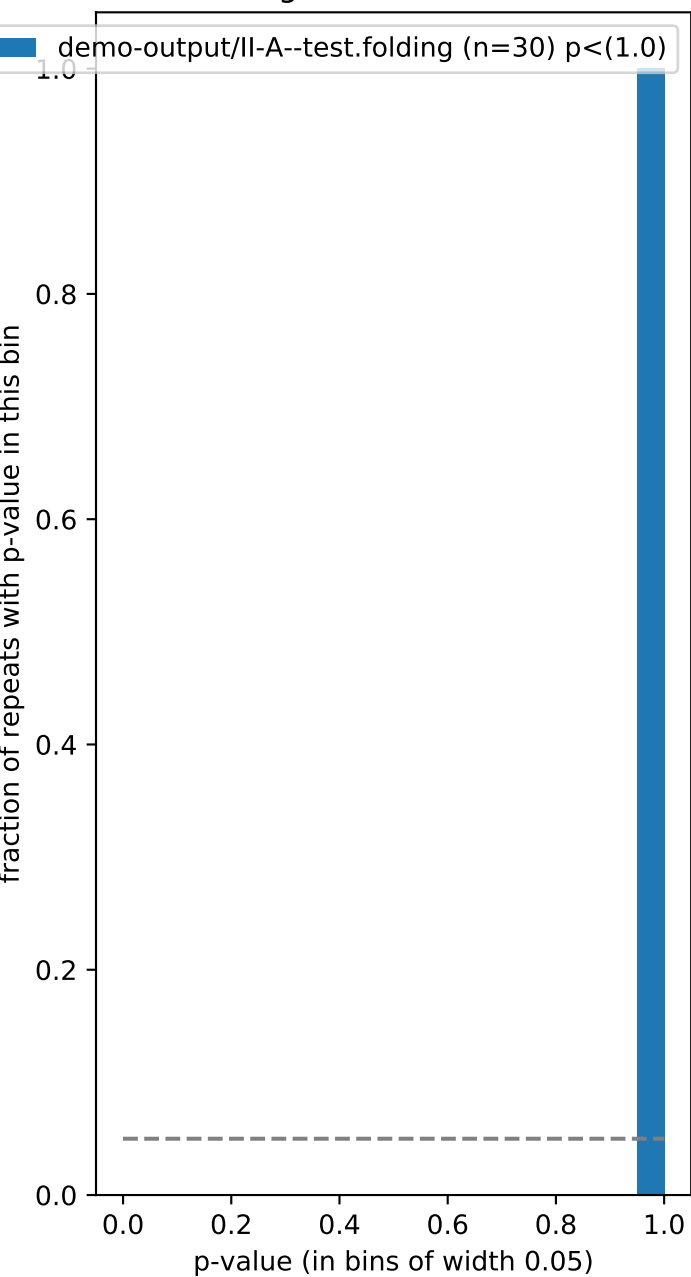
0.9

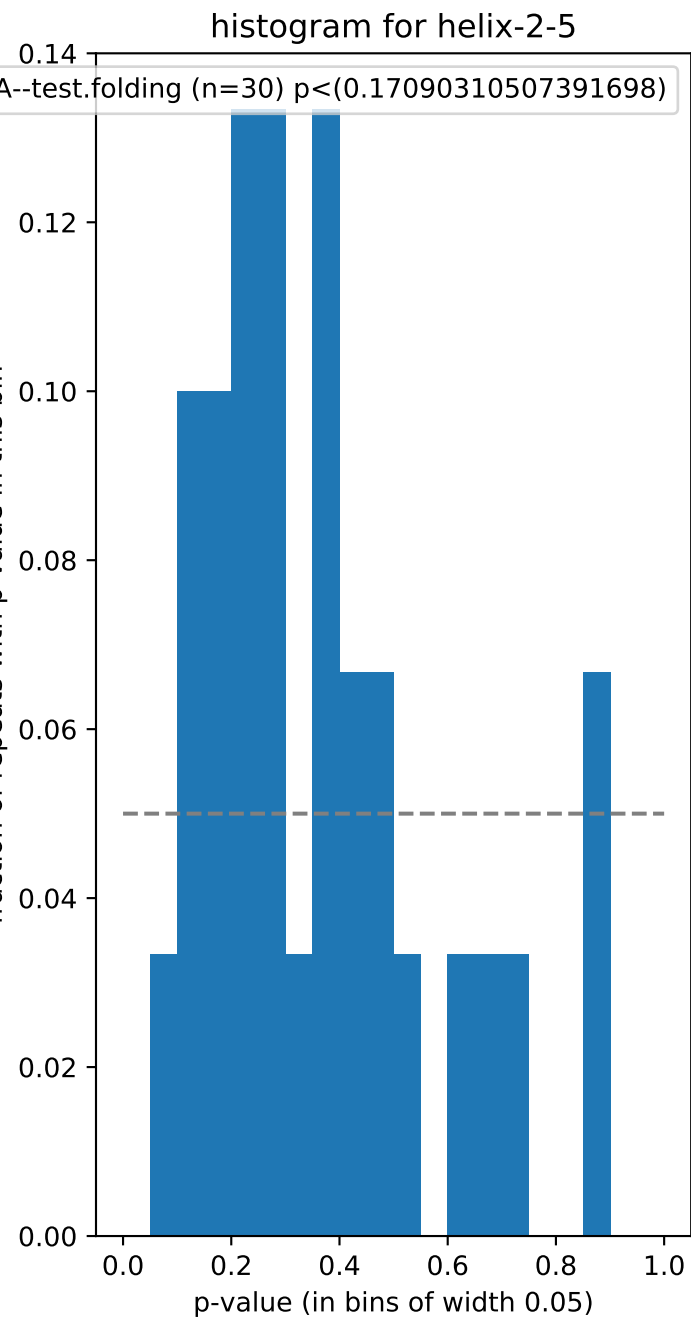


histogram for helix-2-19



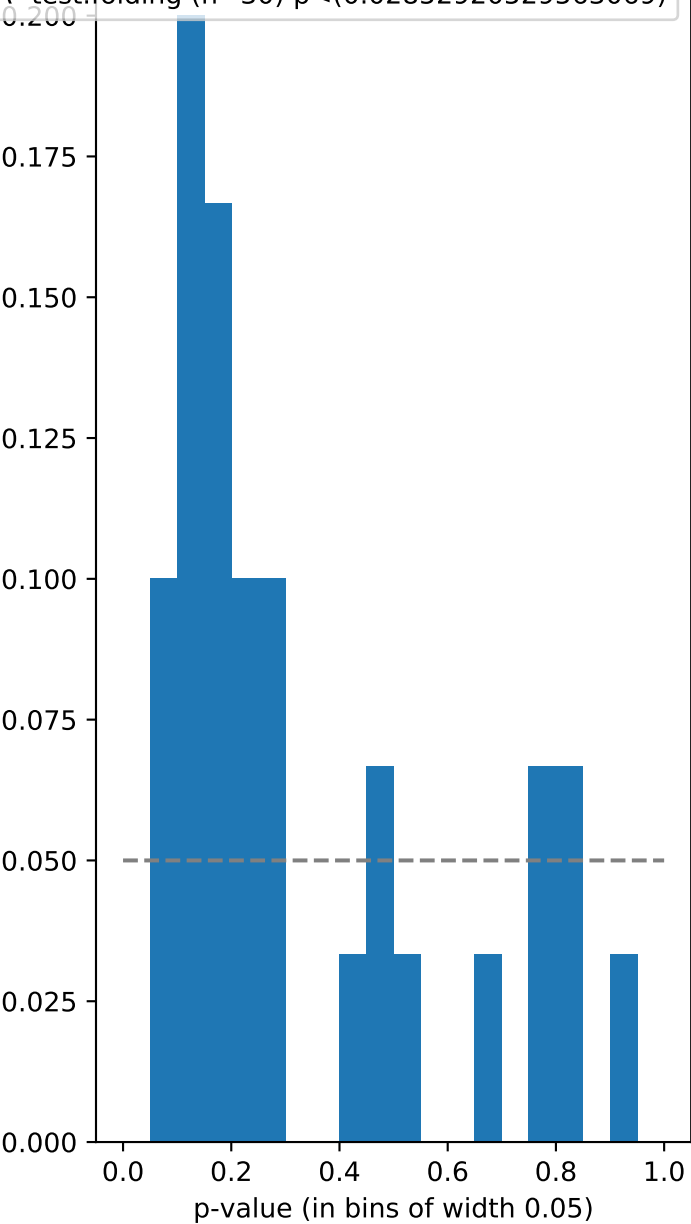
histogram for helix-2-20





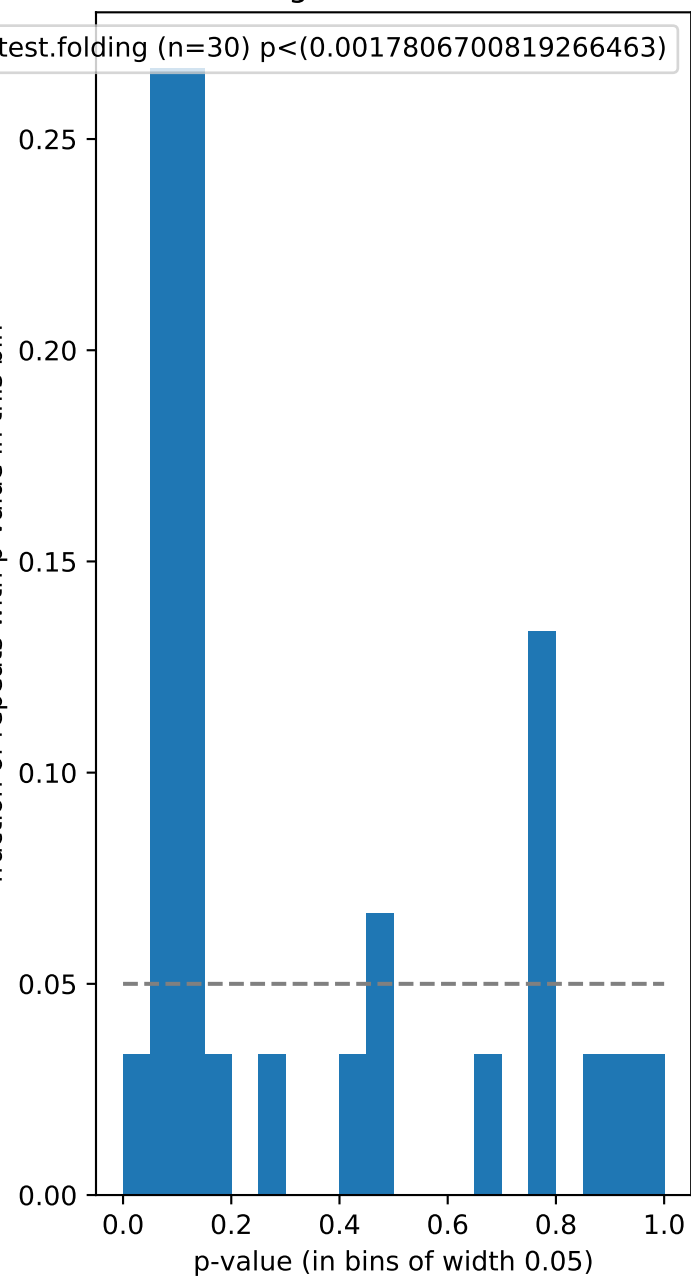
histogram for helix-2-6

A--test.folding (n=30) $p < (0.02852920529563069)$



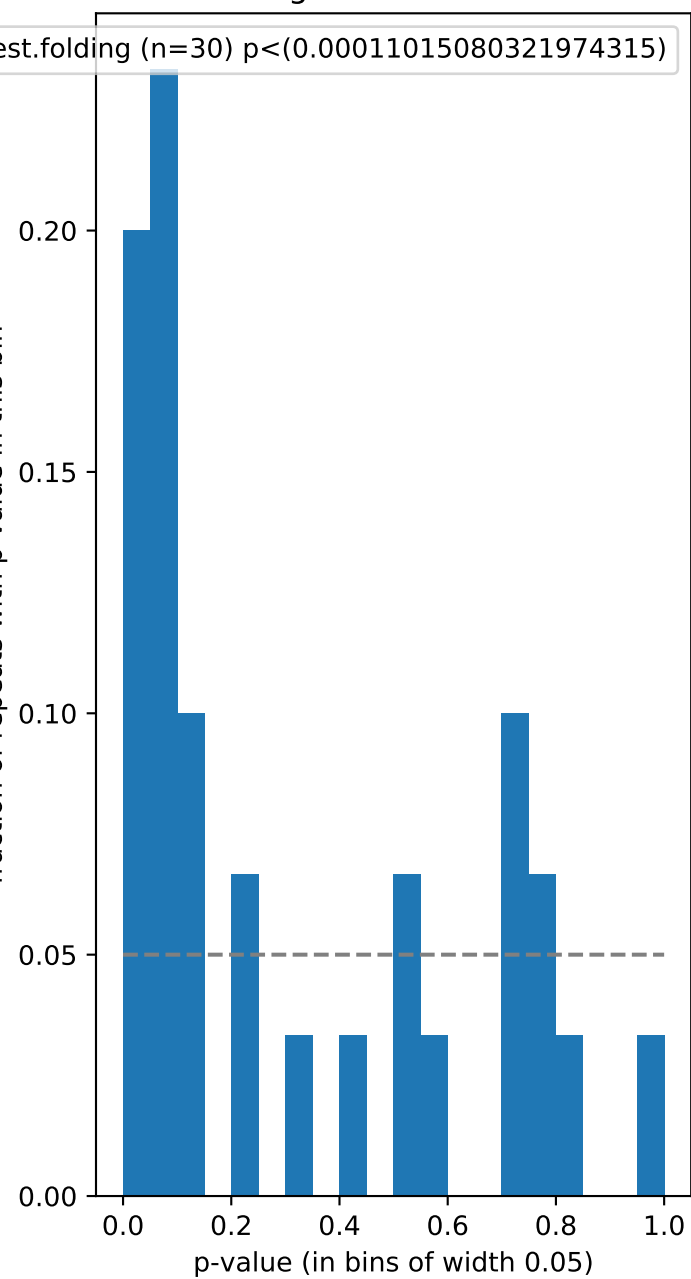
histogram for helix-2-7

test.folding (n=30) $p < (0.0017806700819266463)$



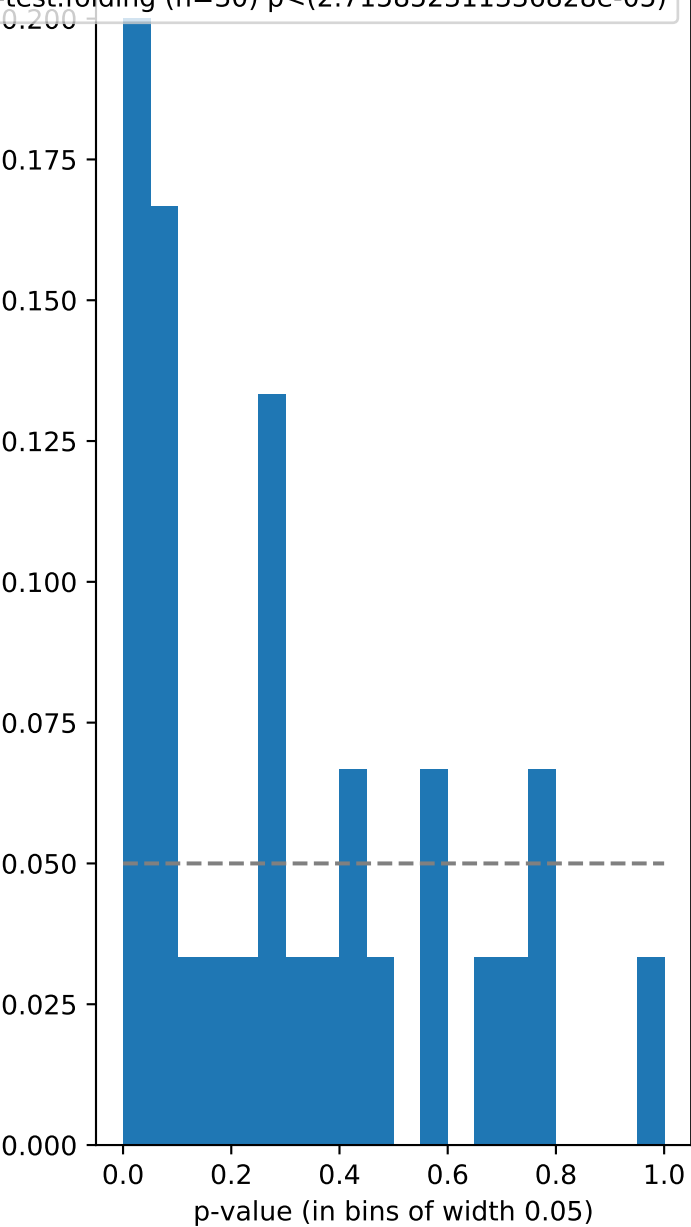
histogram for helix-2-8

est.folding (n=30) $p < (0.00011015080321974315)$



histogram for helix-2-9

test.folding (n=30) $p < (2.715852311336828e-05)$



histogram for logGeomMeanFullyUnpaired

--test.folding (n=30) $p < (0.037139582586920884)$

