Corr: 0.81; Agreement: 73% 5.0 -MutationAssessor raw score

0.50 (
ADME raw score

0.00

0.25

Predicted effect

- D in ADME; N in MutationAssessor
- D in both
- D in MutationAssessor; N in ADME
- N in both

1.00

0.75