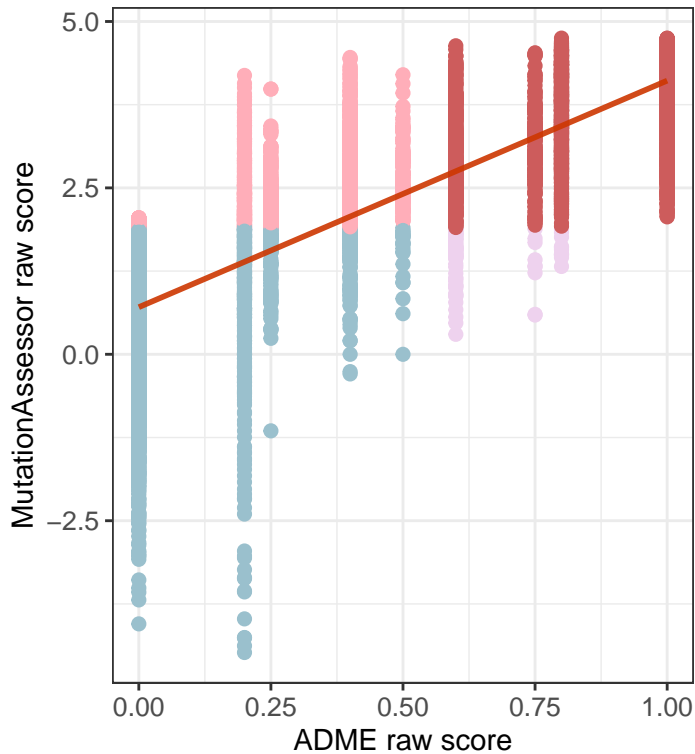


Corr: 0.81; Agreement: 73%



Predicted effect

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