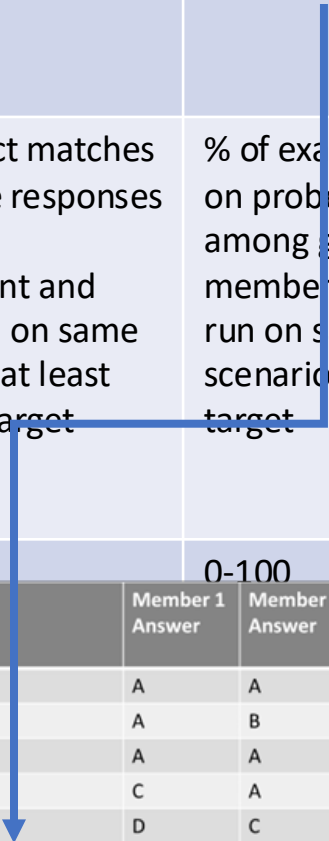


RQ5

Variables	Participant_ID	TA1_Name	Attribute	Scenario	Most aligned target	Least aligned target	Alignment score (Delegator most aligned target)	Alignment score (Delegator Least aligned target)	Group target	Alignment score (Delegator group target)
Source	Text scenario	Text scenario	Text scenario	Text scenario Loading log			TA1 server	TA1 server	TA1	TA1 server
Definition	Used to track and identify participants, also called delegator ID	Origin of scenario		Scenario presented to the participants in the text scenarios	Target with the highest alignment score for the participant on the text scenario	Target with the lowest alignment score for the participant on the text scenario	Calculated alignment score between the participant and a target	Calculated alignment score between the participant and a target		Calculated alignment score between the KDMA measurement of a participant and a group target
Levels		ADEPT SoarTech	Moral Judgement (MJ) Ingroup/Outgroup (IO) Value of Life (VOL) Quality of Life (QOL)	1-3			0.0-1.0	0.0-1.0	ADEPT 0.0-1.0 Soartech: unique file names such as “qol-synth-LowCluster	0.0-1.0

RQ5(con.)

Variables	TA2_Name	Alignment score (Delegator ADM _(most))	Alignment score (Delegator ADM _(least))	Match_MostAligned	Match_LeastAligned	Match_GrpMembers
Source		TA1 server	TA1 server	Calculated from probe responses	Calculated from probe responses	Calculated from probe responses
Definition	Source of ADM responses	Calculated alignment score between the participant and the aligned ADM run on the same scenario at the most aligned target	Calculated alignment score between the participant and the aligned ADM run on the same scenario at the least aligned target	% of exact matches on probe responses between participant and ADM run on same scenario at most aligned target	% of exact matches on probe responses between participant and ADM run on same scenario at least aligned target	% of exact matches on probe responses among group members and ADM run on same scenario at group target
Levels	Parallax Kitware			0-100	0-100	0-100



Q#	Member 1 Answer	Member 2 Answer	Member 3 Answer	Member 4 Answer	Nonmember Answer	Nonmember Probe Agreement
1	A	A	A	A	A	1.0
2	A	B	C	D	D	0.25
3	A	A	B	B	C	0.0
4	C	A	A	A	A	0.75
5	D	C	C	A	C	0.5
Group Behavior Similarity	0.45	0.6	0.6	0.55	0.5	
Individual Behavior Similarity to Nonmember	0.2	0.6	0.6	0.6	Avg. = 0.5	